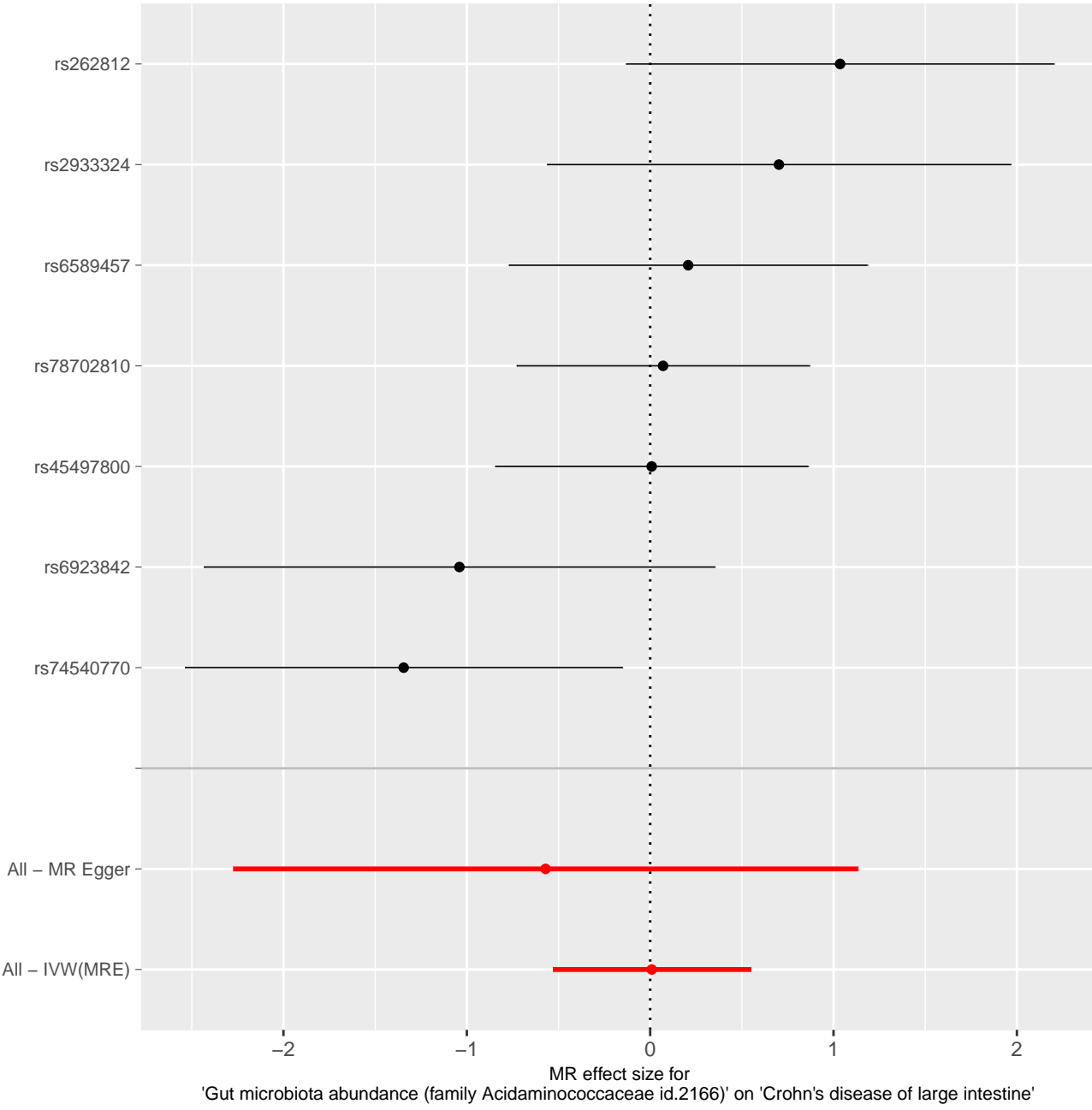
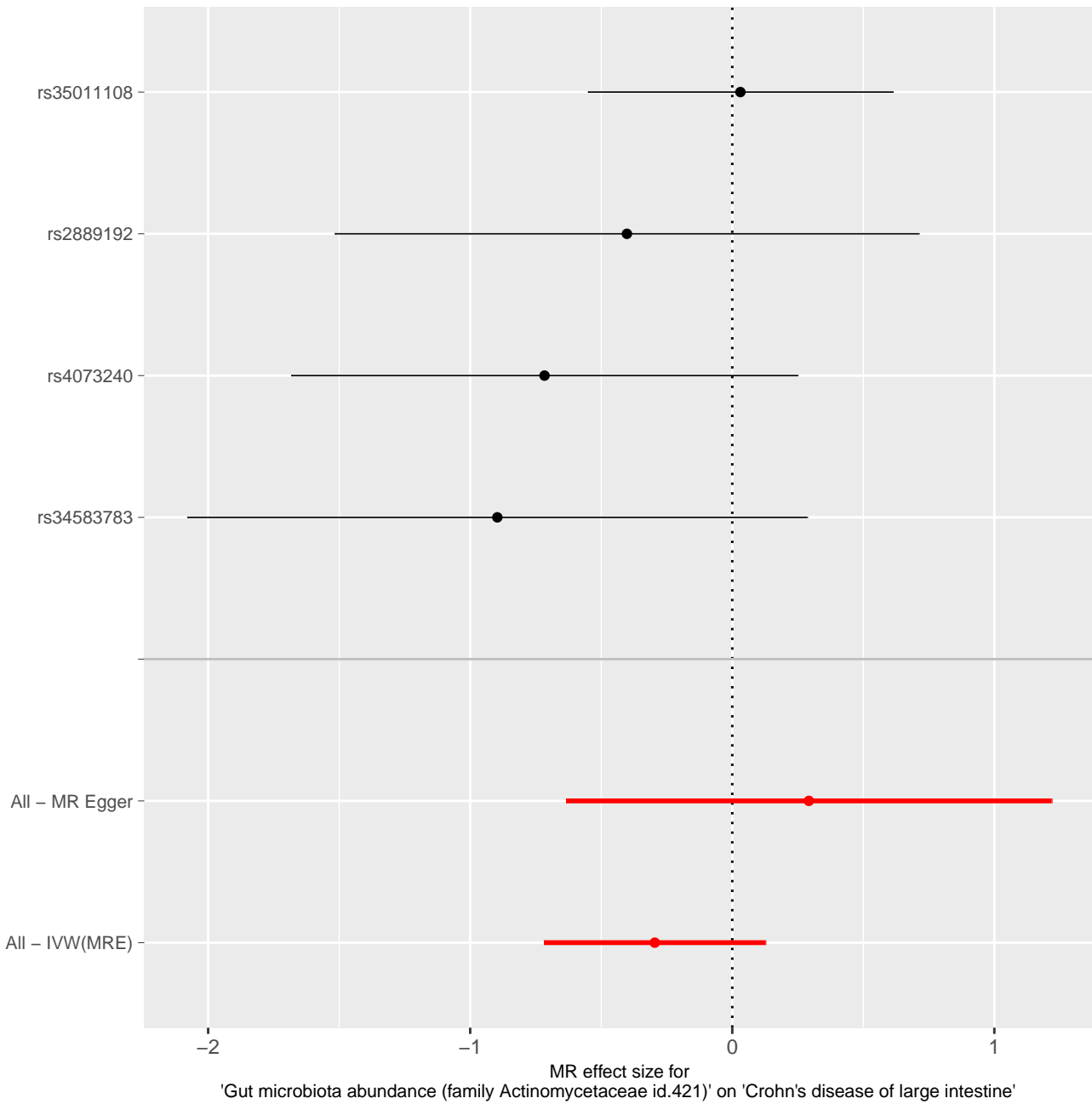


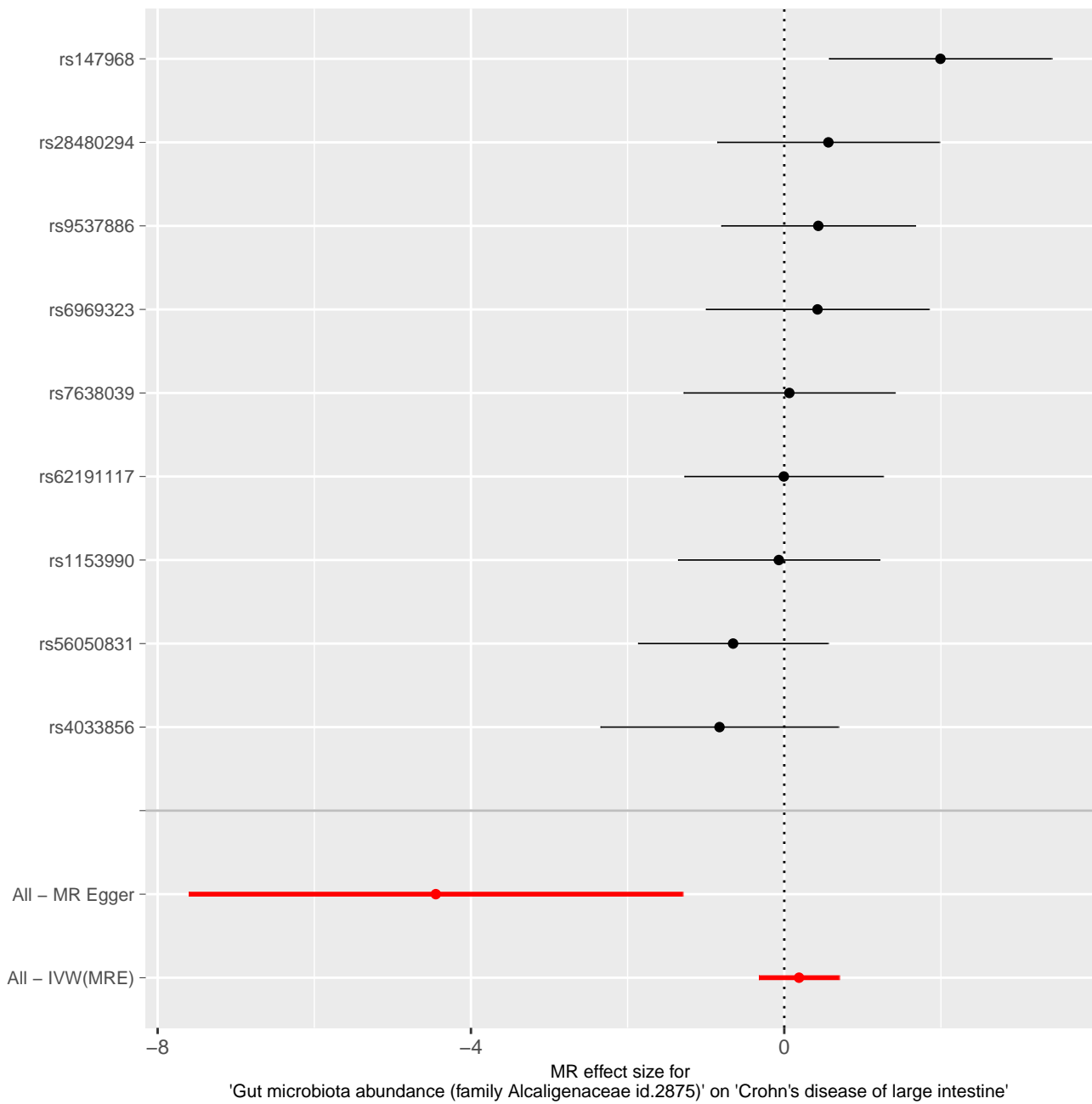
Batch 1 : Gut microbiota abundance (family Acidaminococcaceae id.2166) on Crohn's disease of large intestine



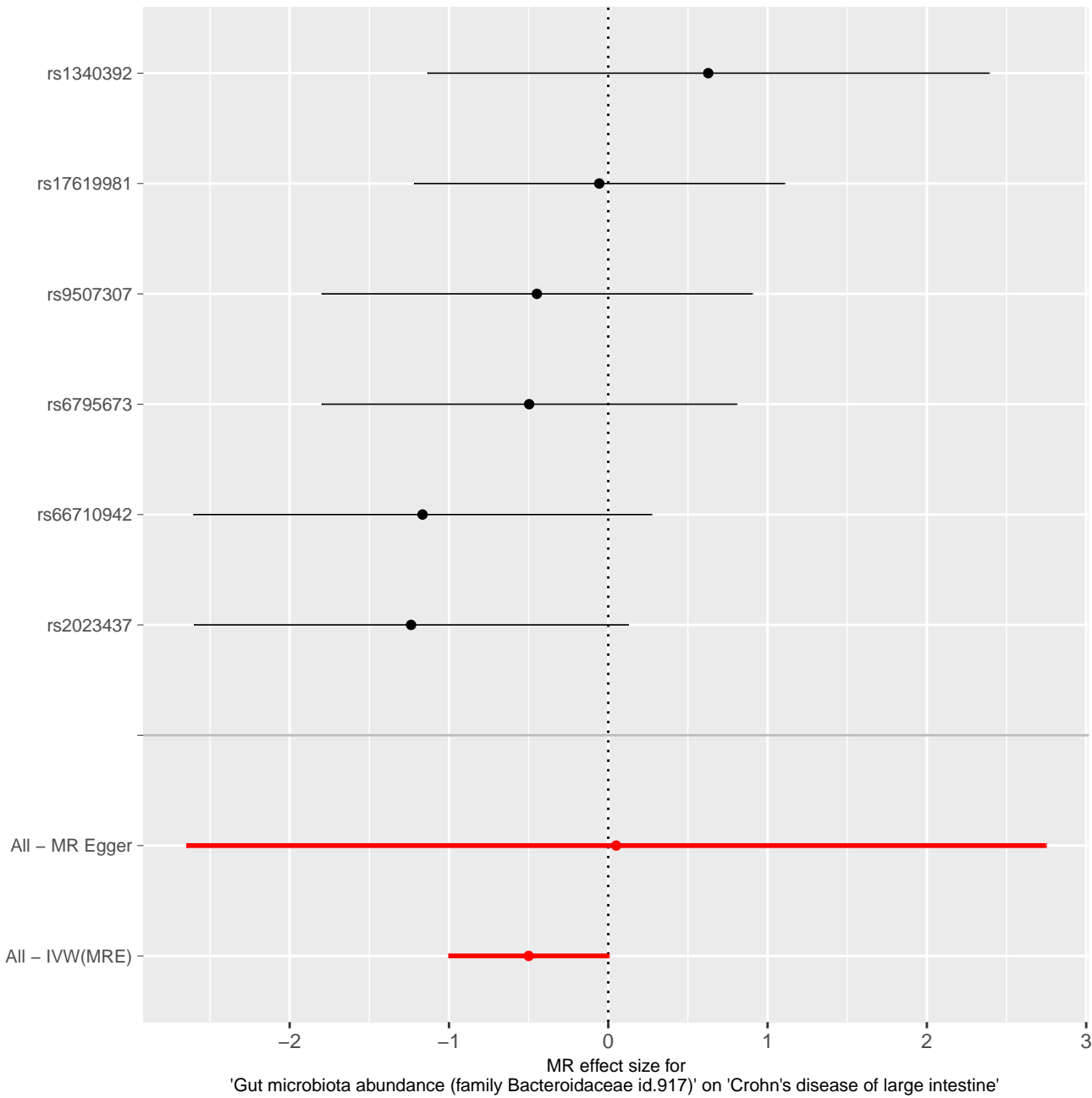
Batch 2 : Gut microbiota abundance (family Actinomycetaceae id.421) on Crohn's disease of large intestine



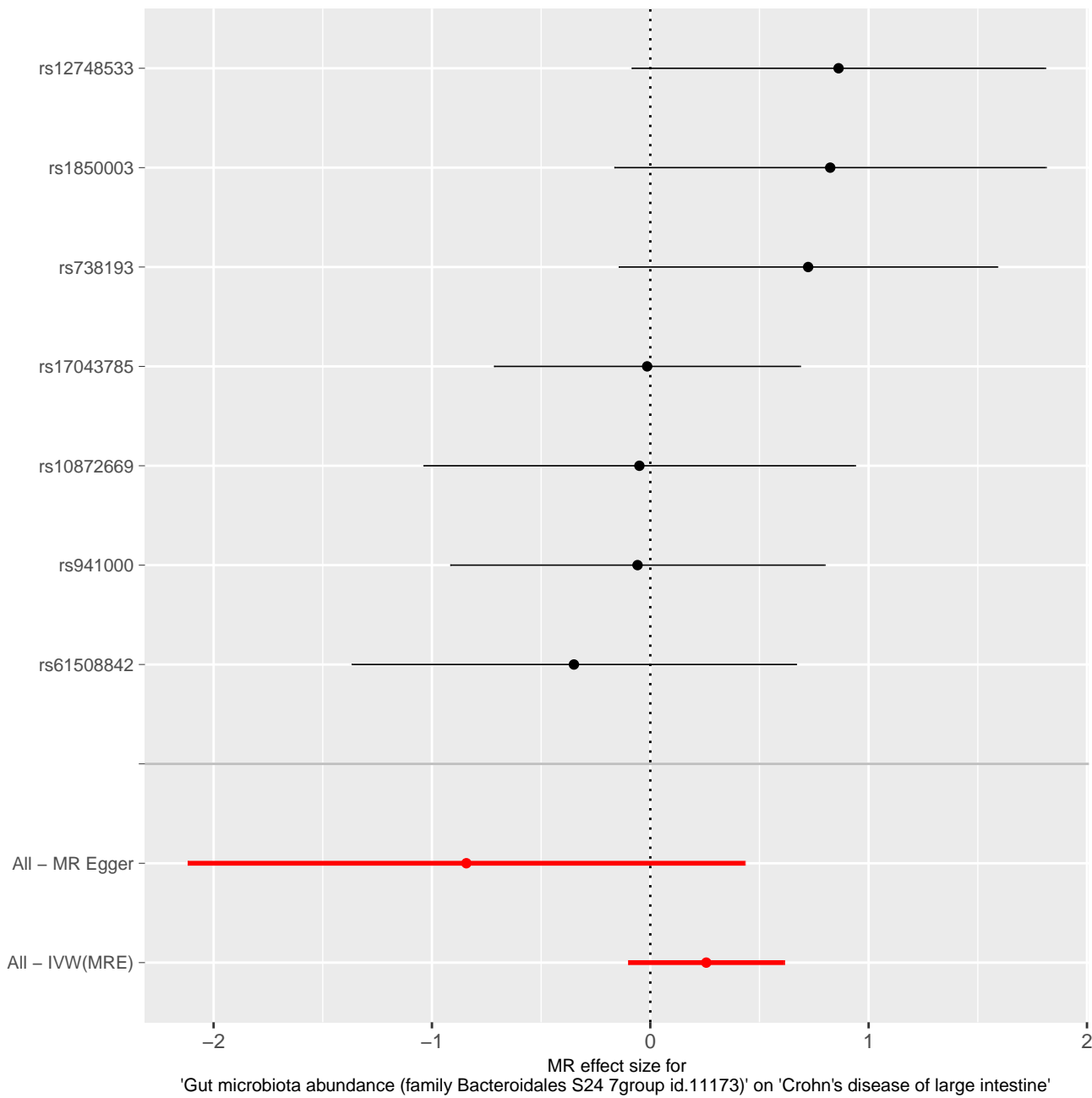
Batch 3 : Gut microbiota abundance (family Alcaligenaceae id.2875) on Crohn's disease of large intestine



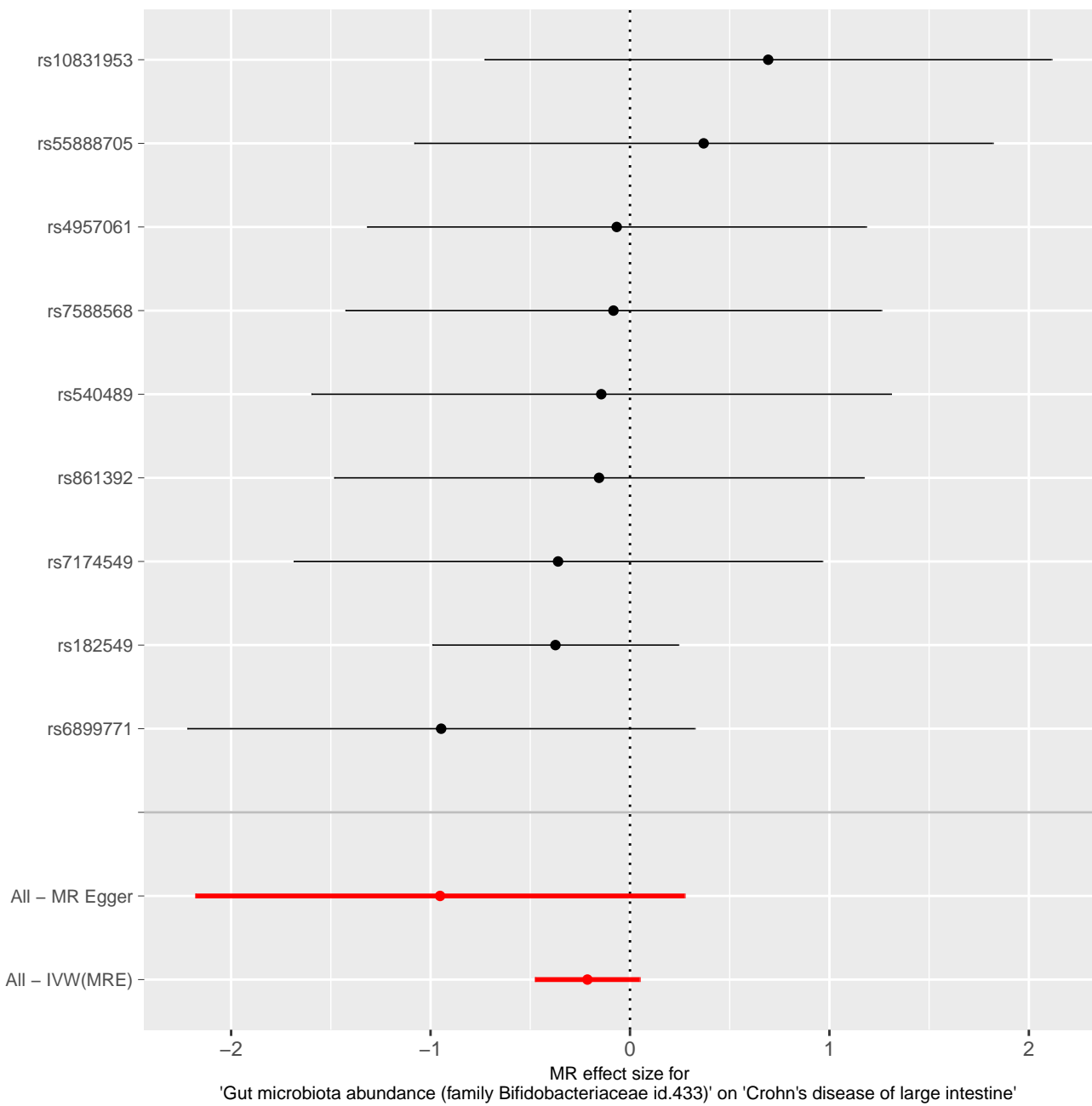
Batch 4 : Gut microbiota abundance (family Bacteroidaceae id.917) on Crohn's disease of large intestine



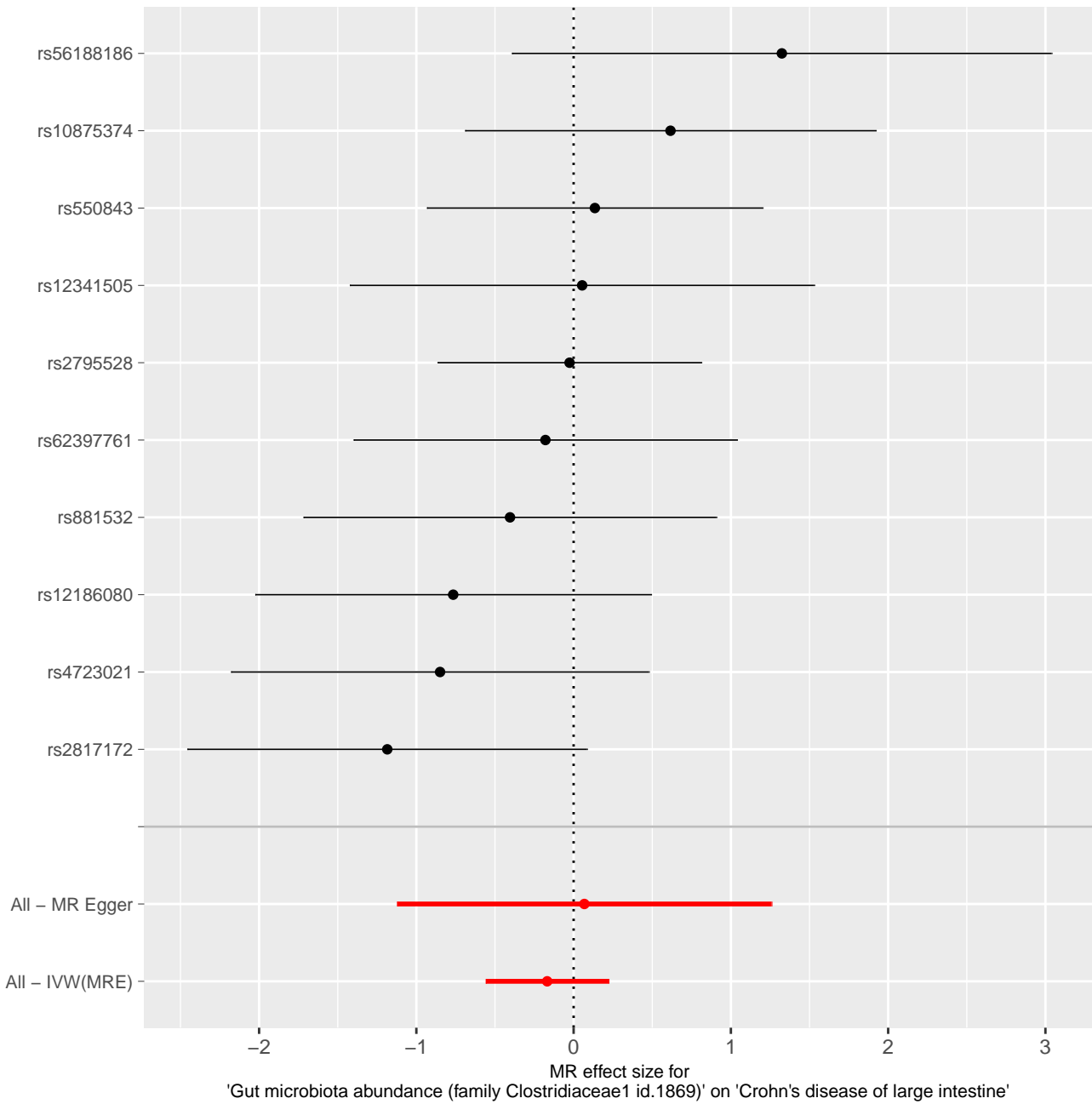
Batch 5 : Gut microbiota abundance (family Bacteroidales S24 7group id.11173) on Crohn's disease of large intestine



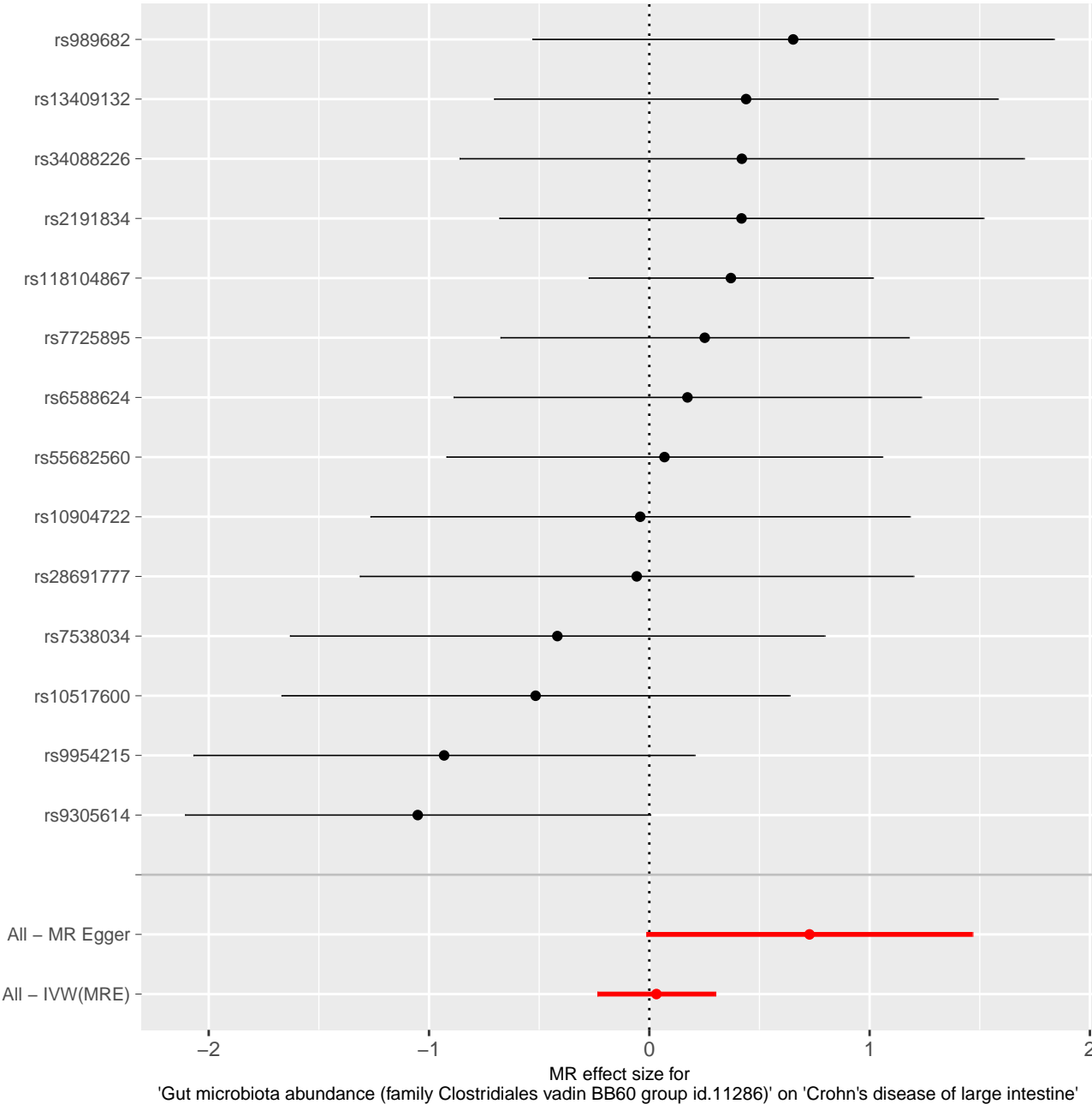
Batch 6 : Gut microbiota abundance (family Bifidobacteriaceae id.433) on Crohn's disease of large intestine



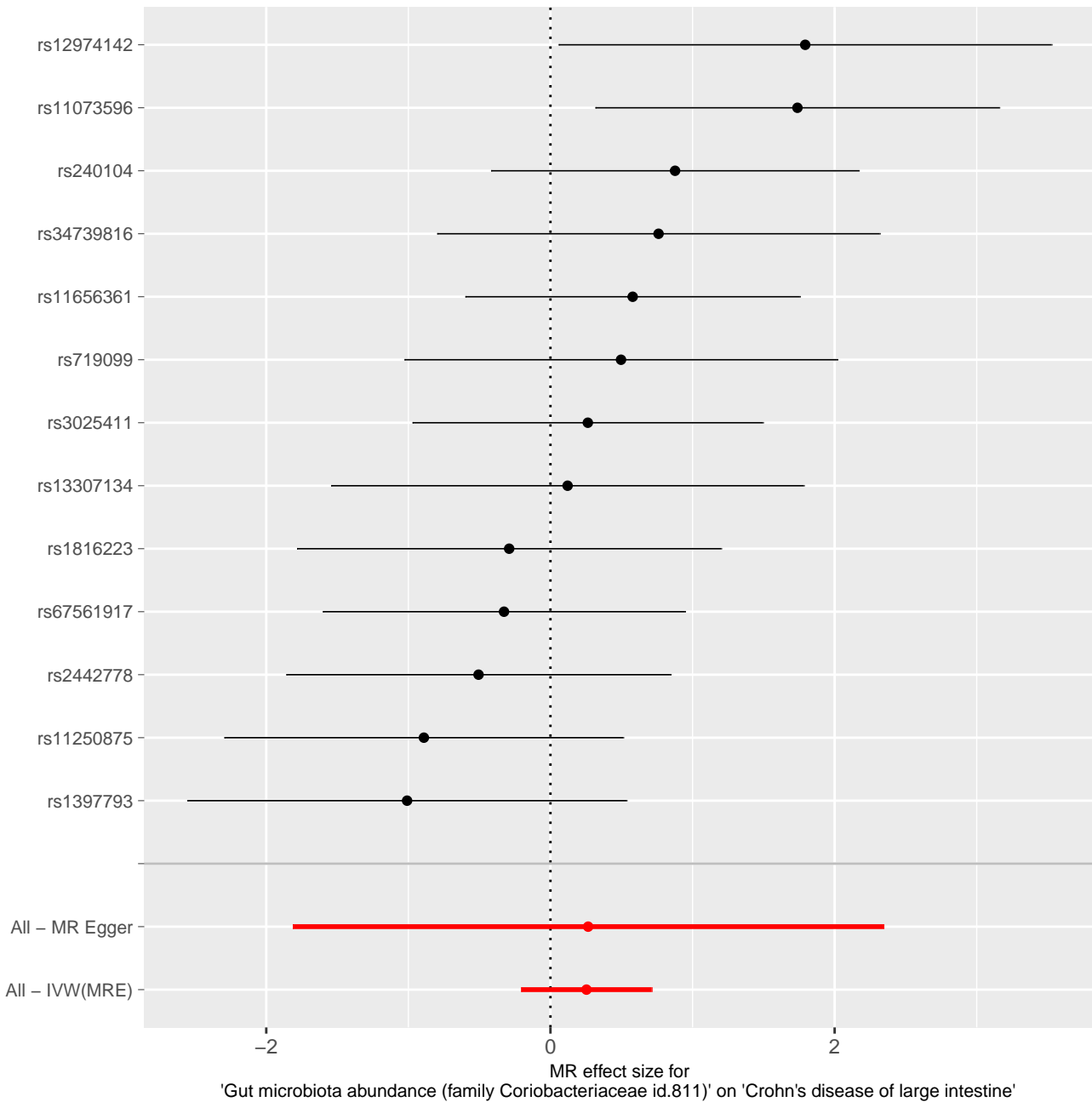
Batch 8 : Gut microbiota abundance (family Clostridiaceae1 id.1869) on Crohn's disease of large intestine



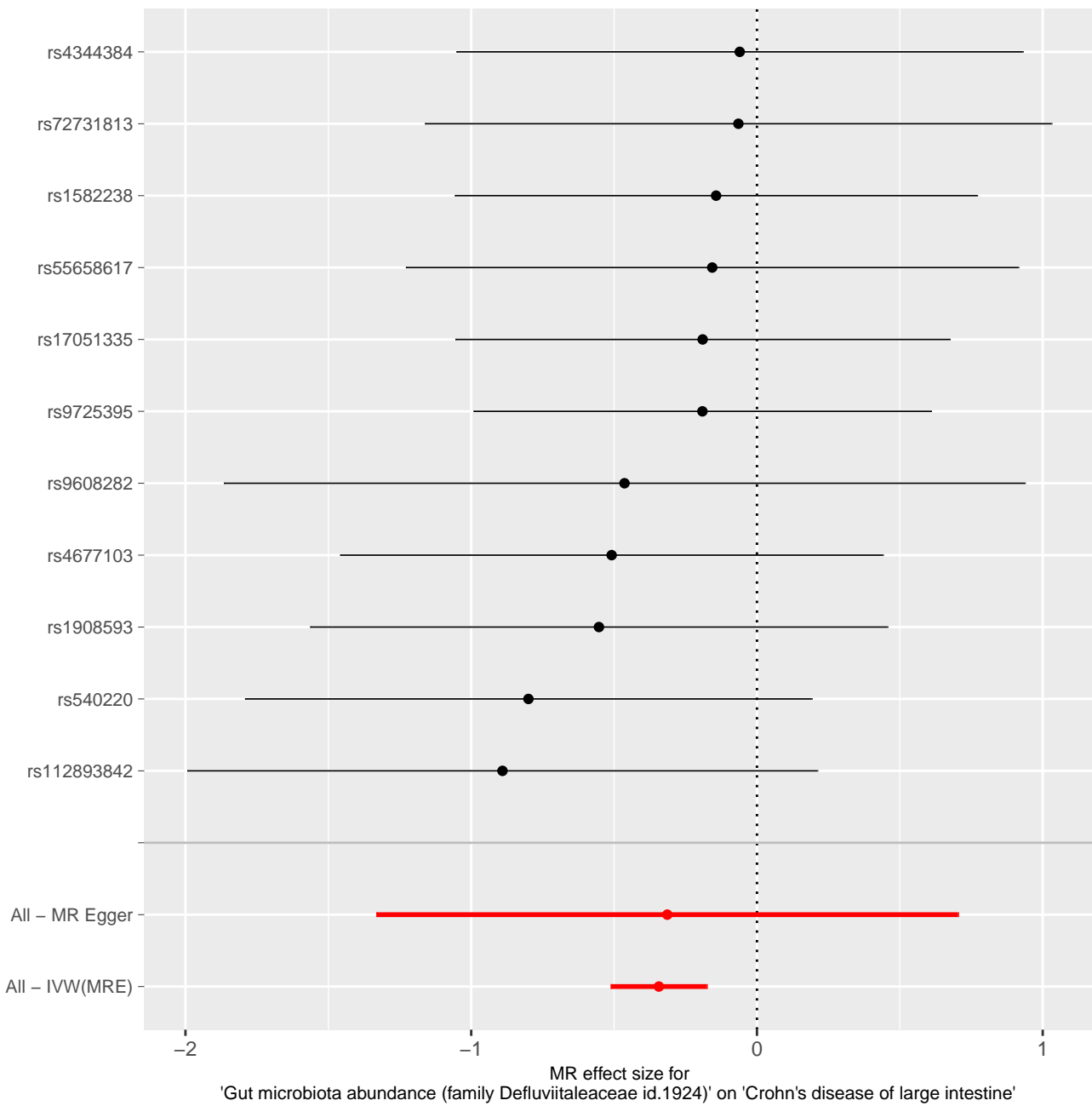
Batch 9 : Gut microbiota abundance (family Clostridiales vadin BB60 group id.11286) on Crohn's disease of large intestine



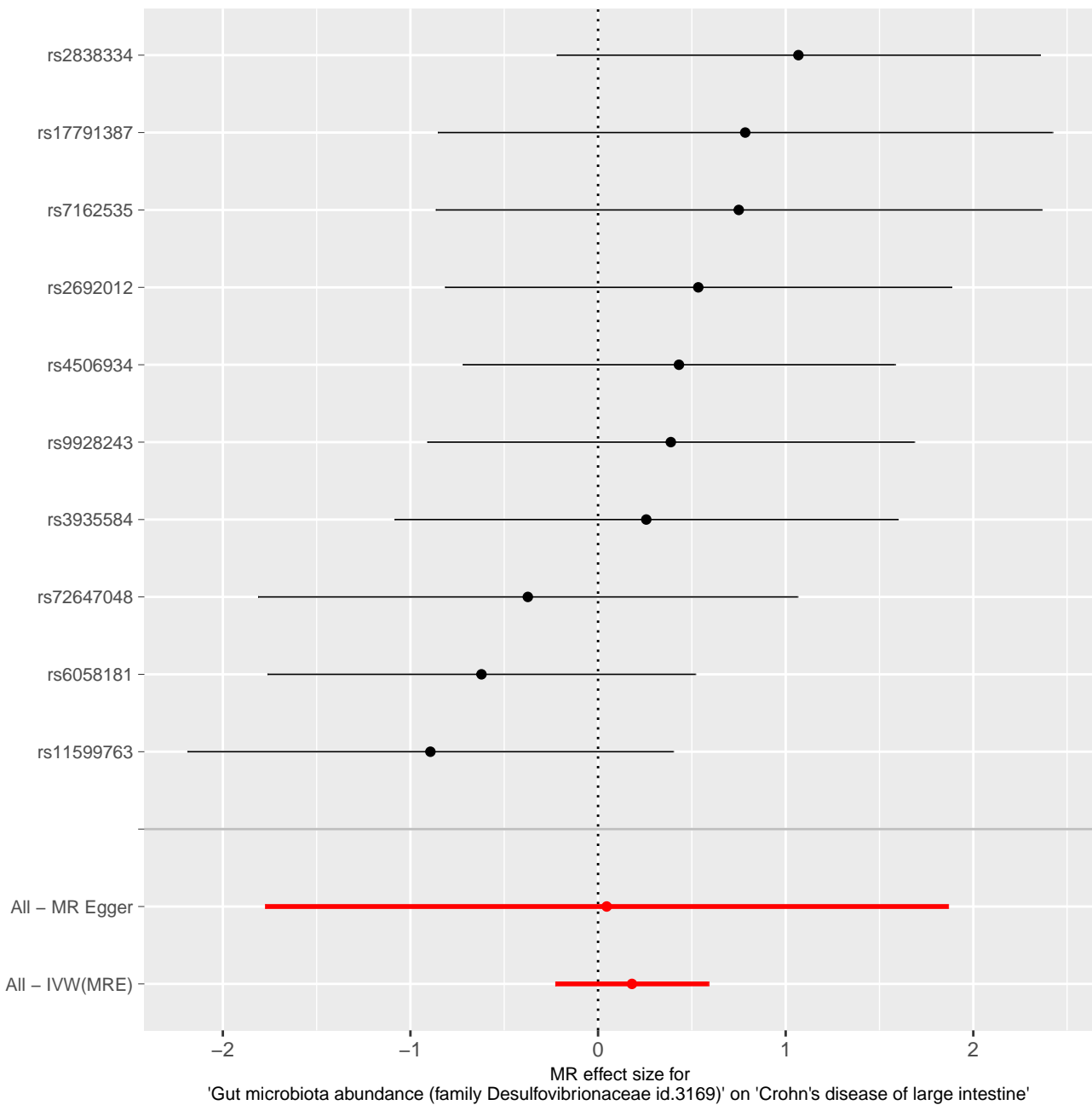
Batch 10 : Gut microbiota abundance (family Coriobacteriaceae id.811) on Crohn's disease of large intestine

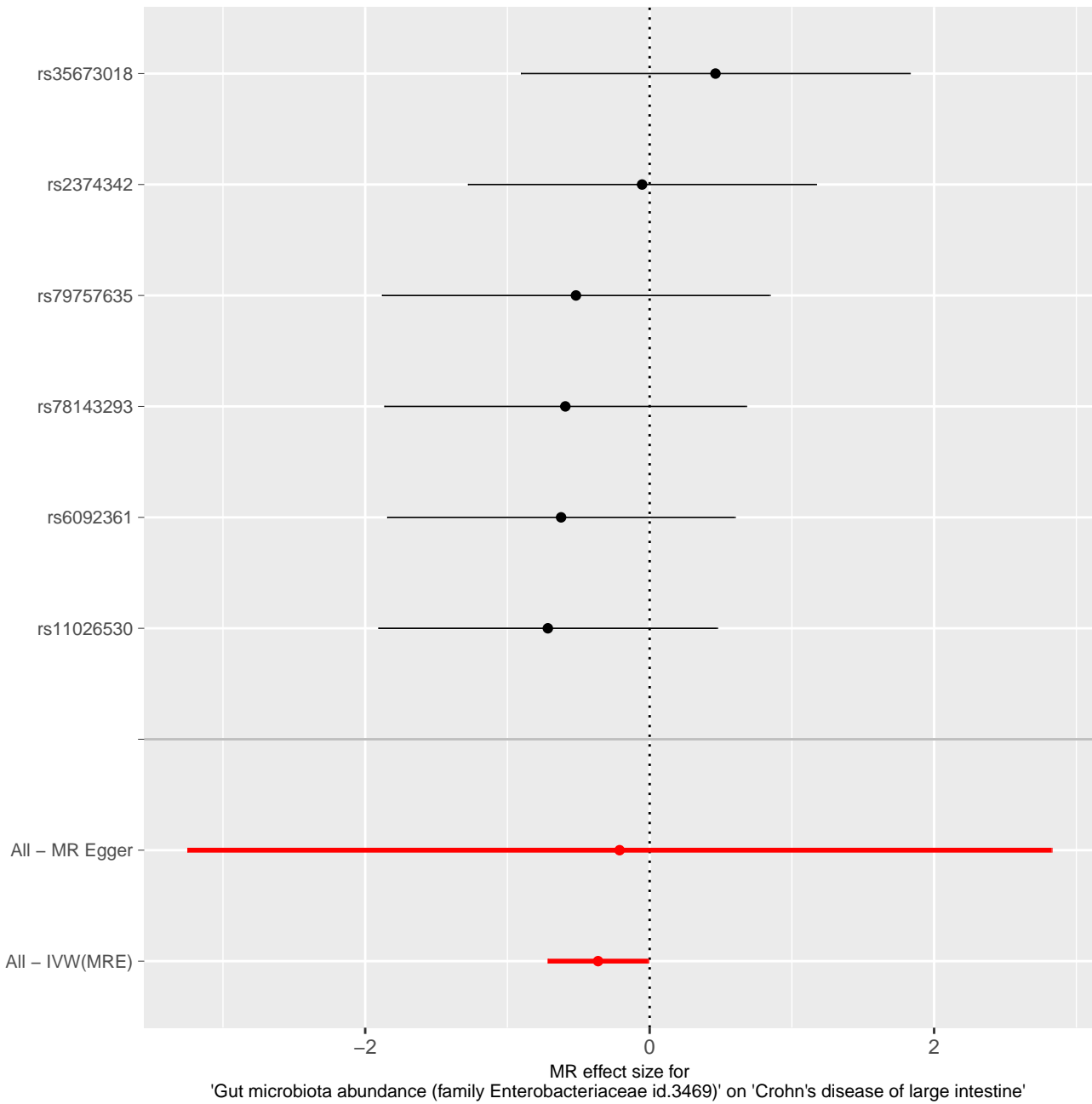


Batch 11 : Gut microbiota abundance (family Defluviitaleaceae id.1924) on Crohn's disease of large intestine

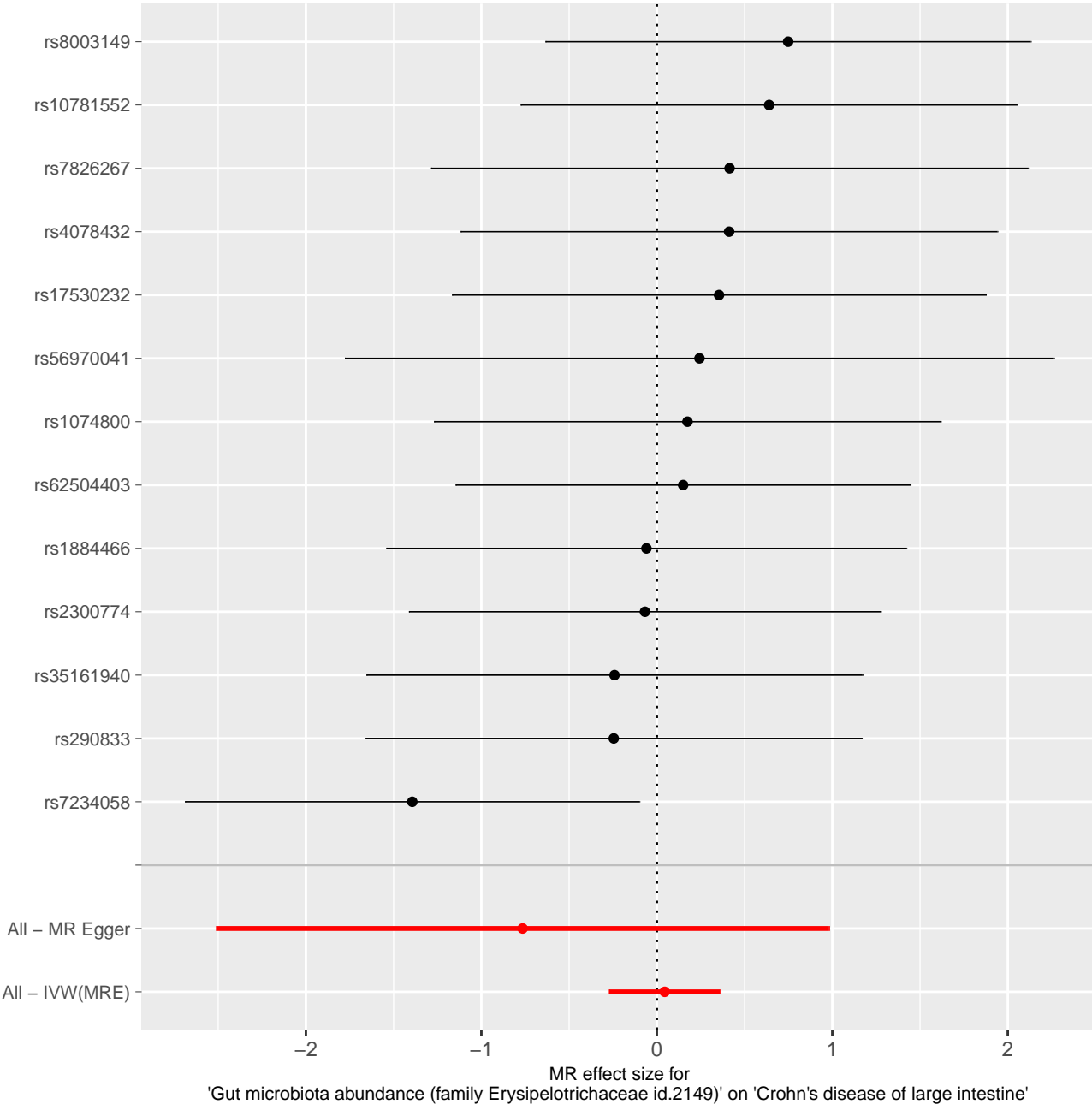


Batch 12 : Gut microbiota abundance (family Desulfovibrionaceae id.3169) on Crohn's disease of large intestine

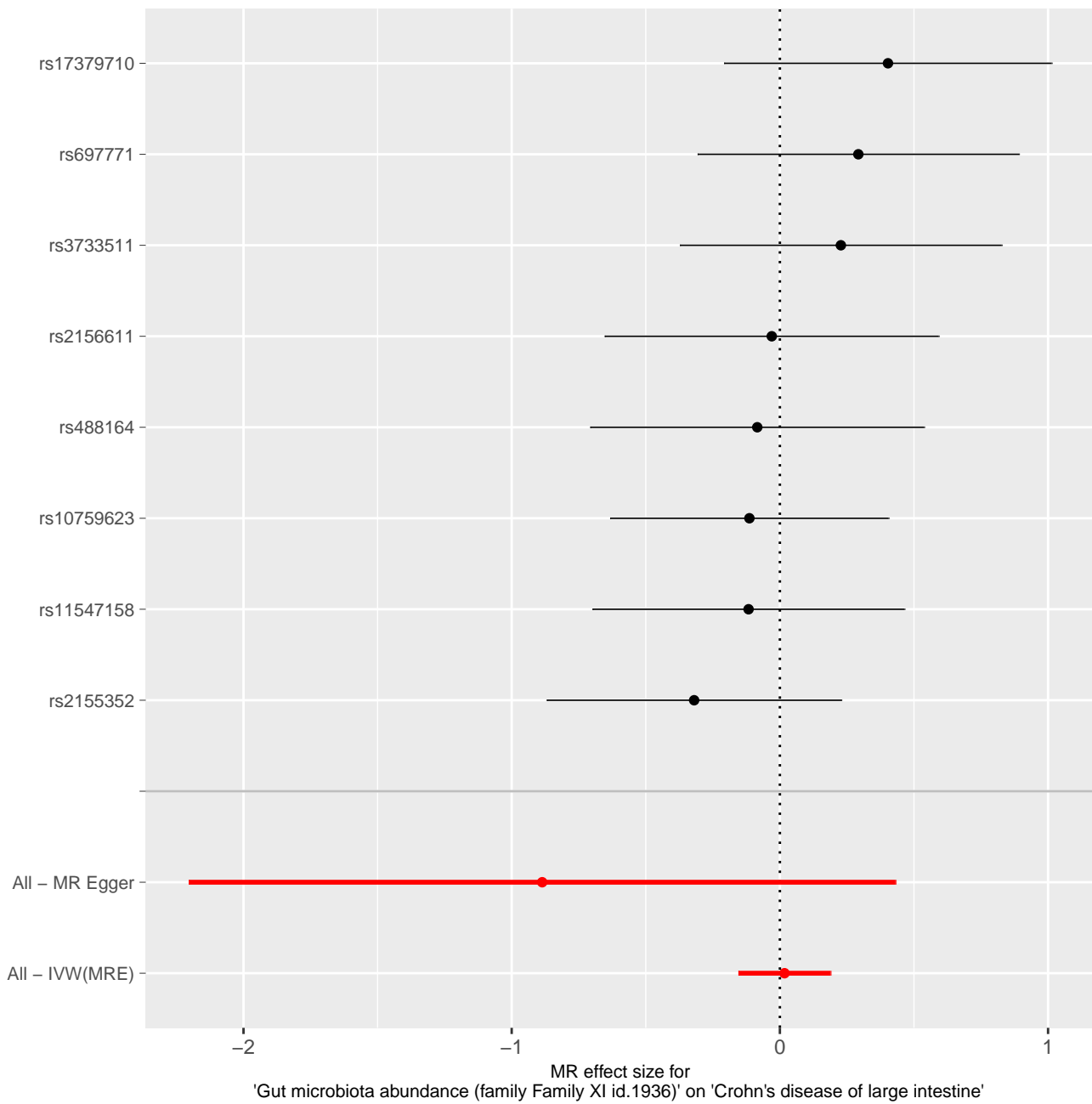


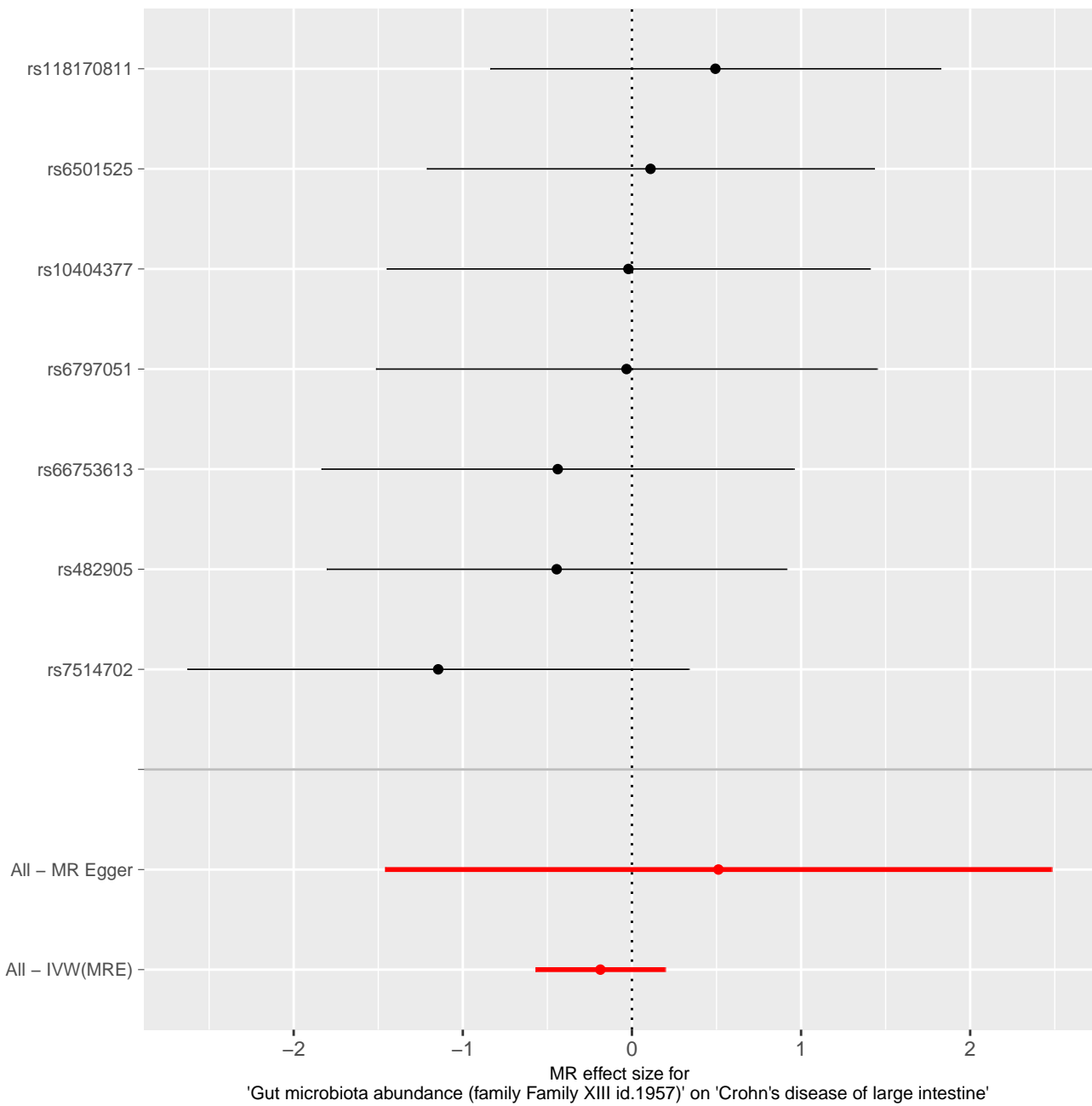


Batch 14 : Gut microbiota abundance (family Erysipelotrichaceae id.2149) on Crohn's disease of large intestine

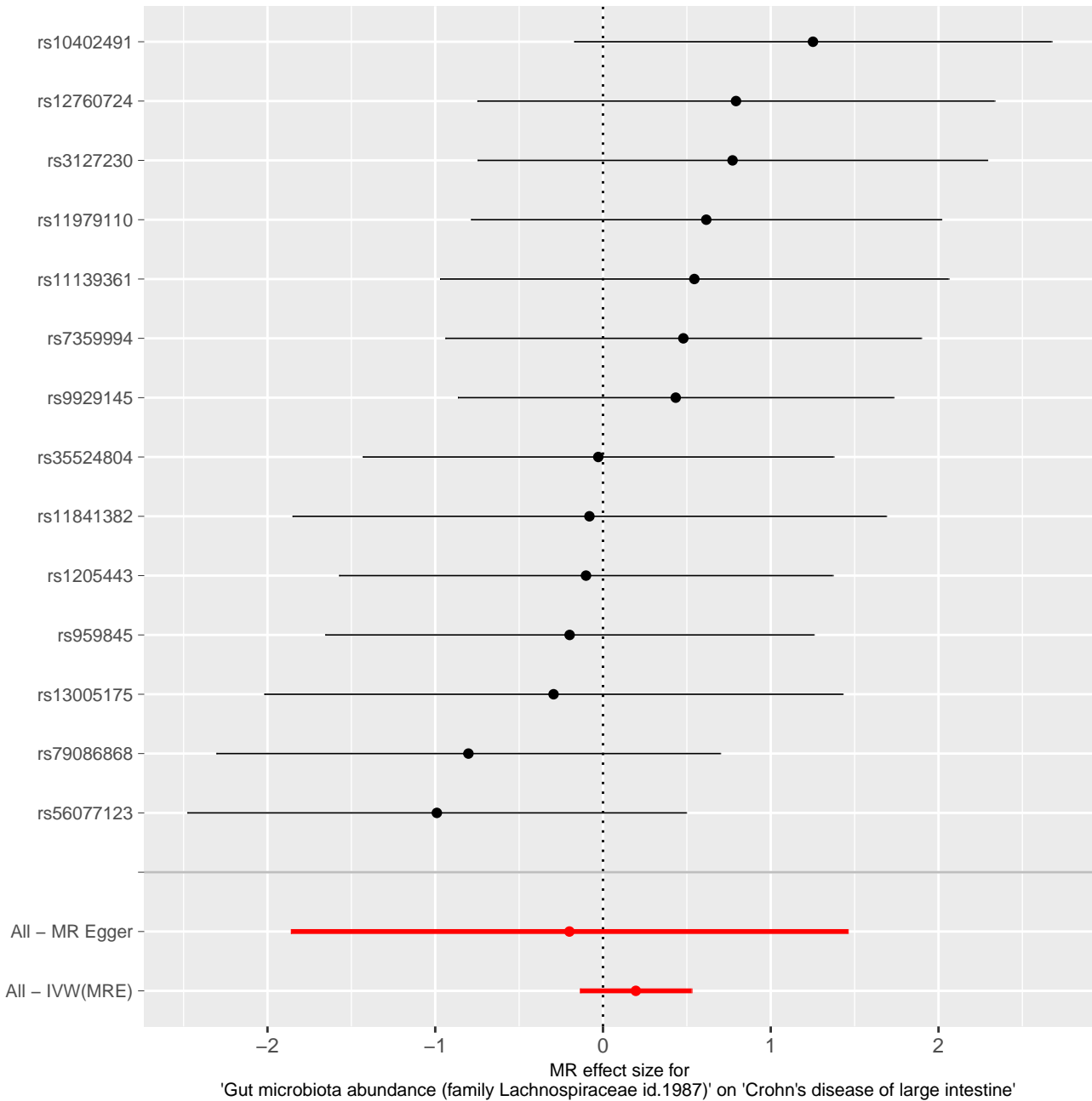


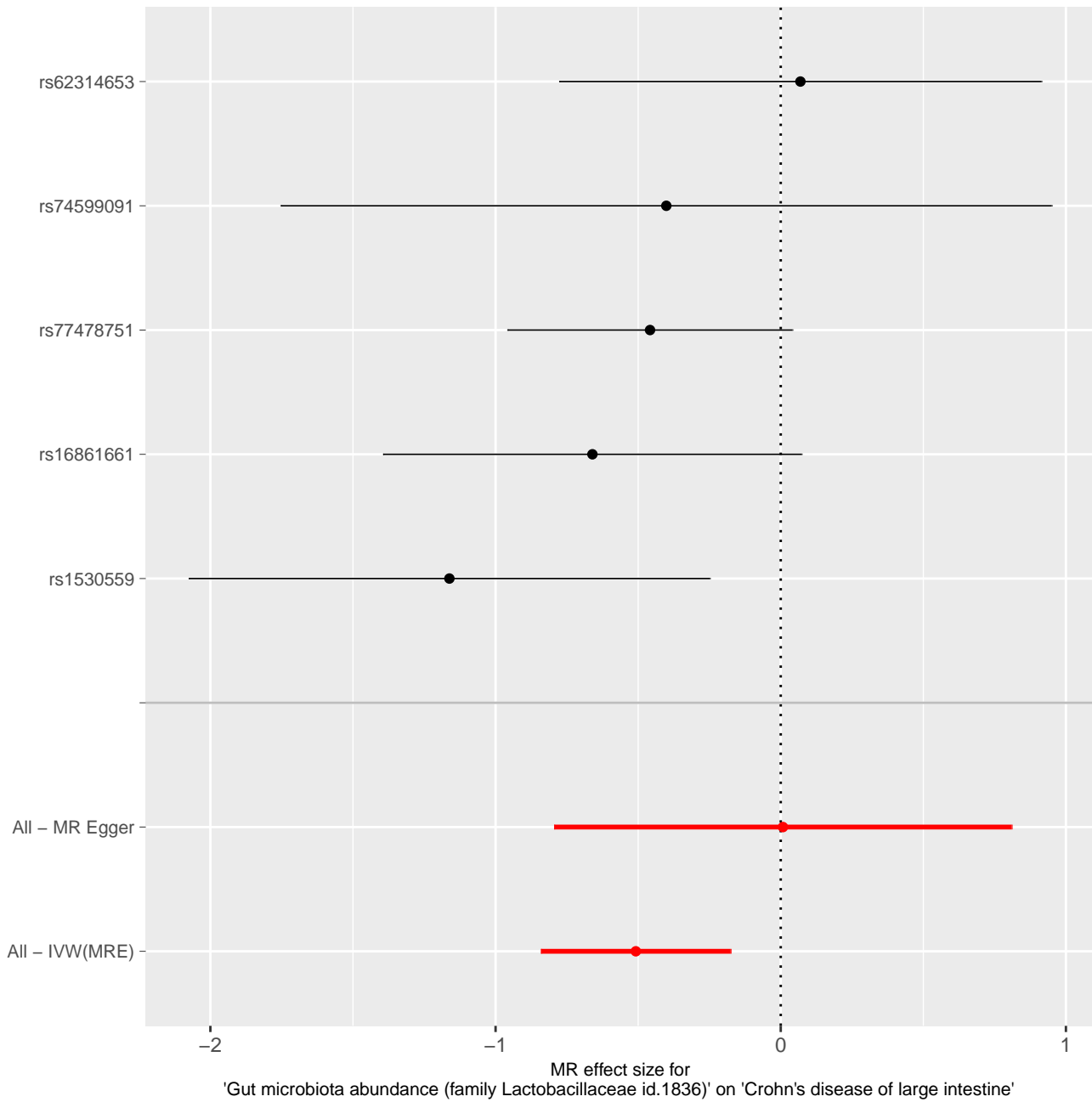
Batch 15 : Gut microbiota abundance (family Family XI id.1936) on Crohn's disease of large intestine



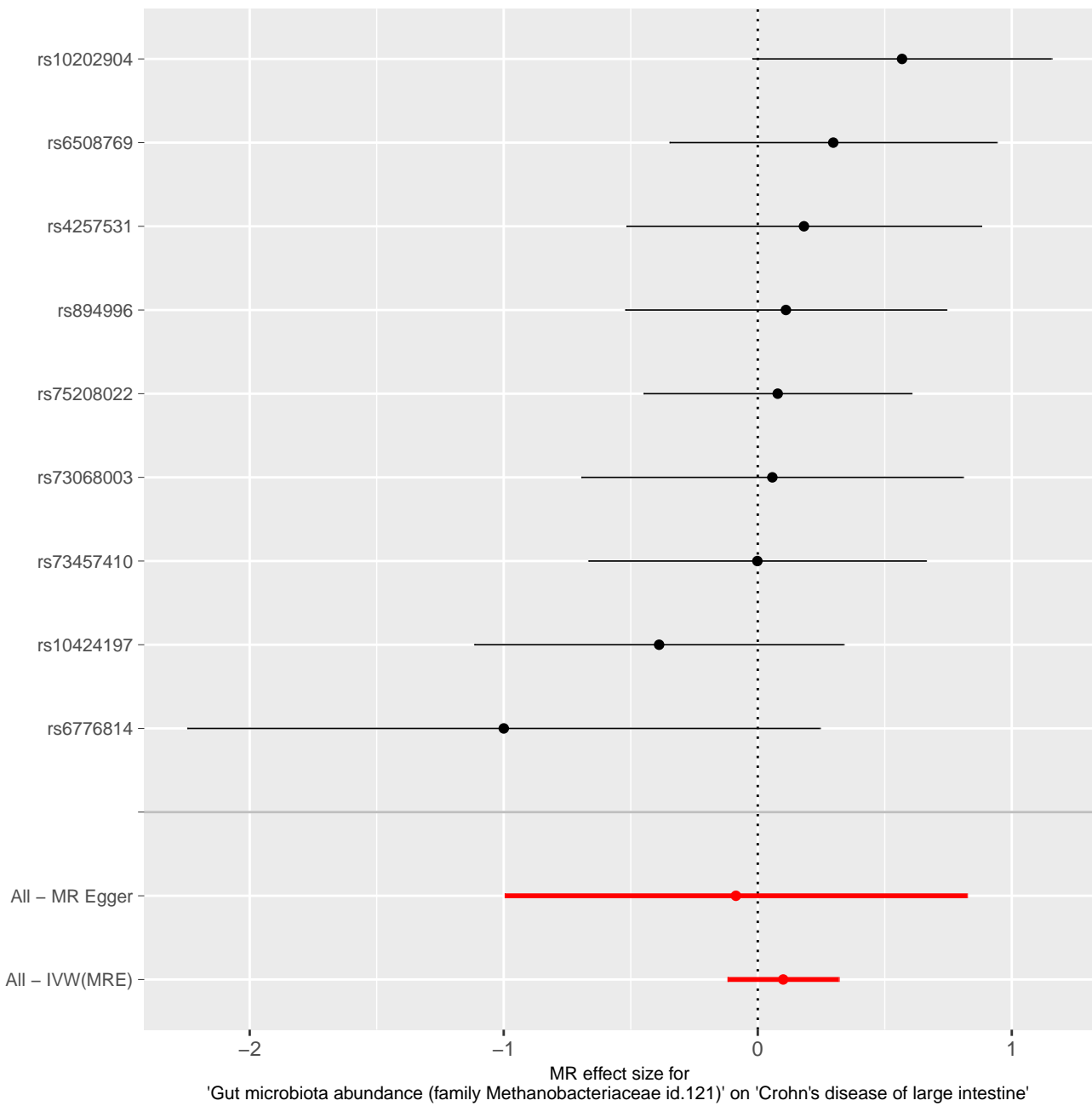


Batch 17 : Gut microbiota abundance (family Lachnospiraceae id.1987) on Crohn's disease of large intestine

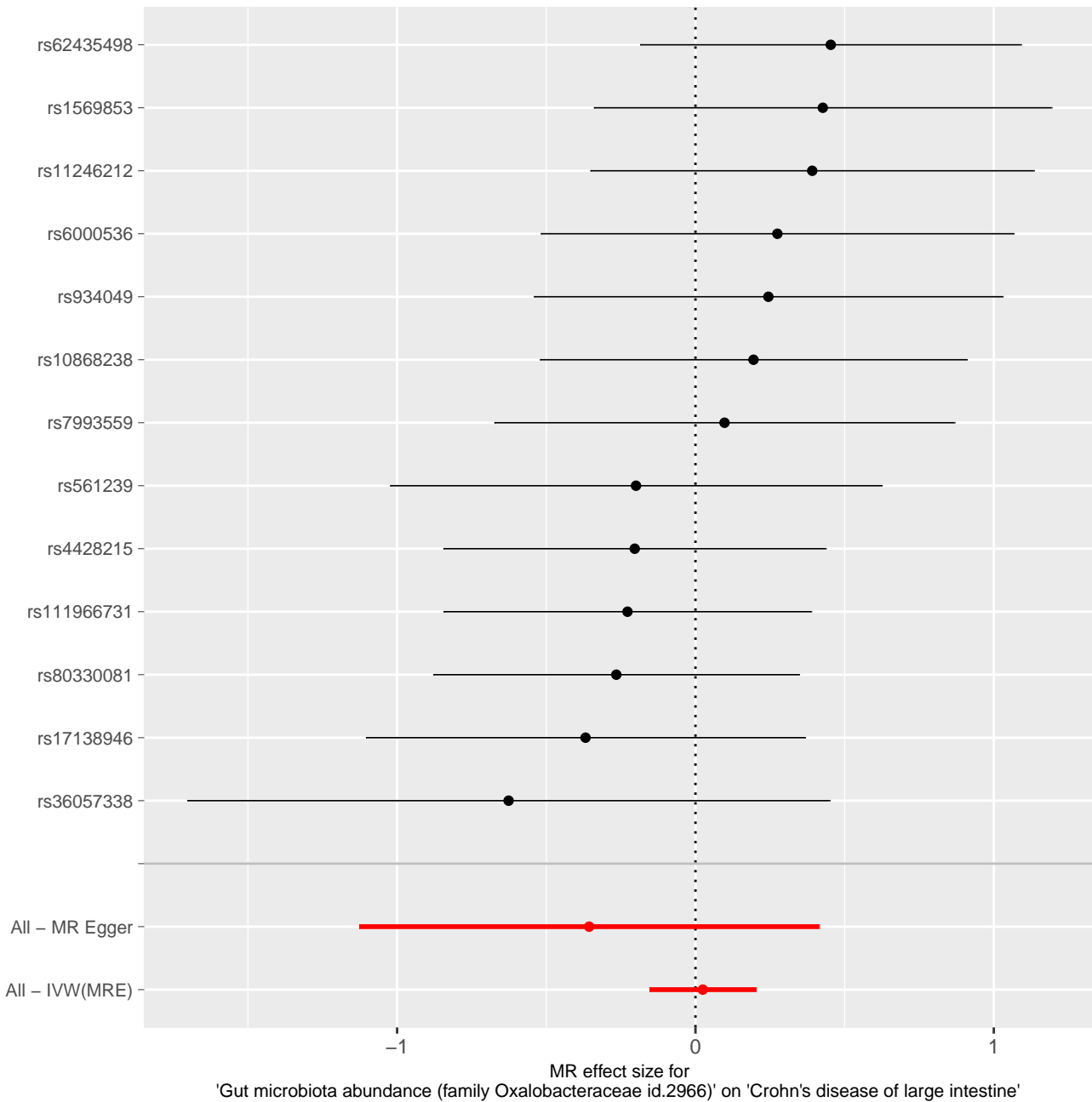




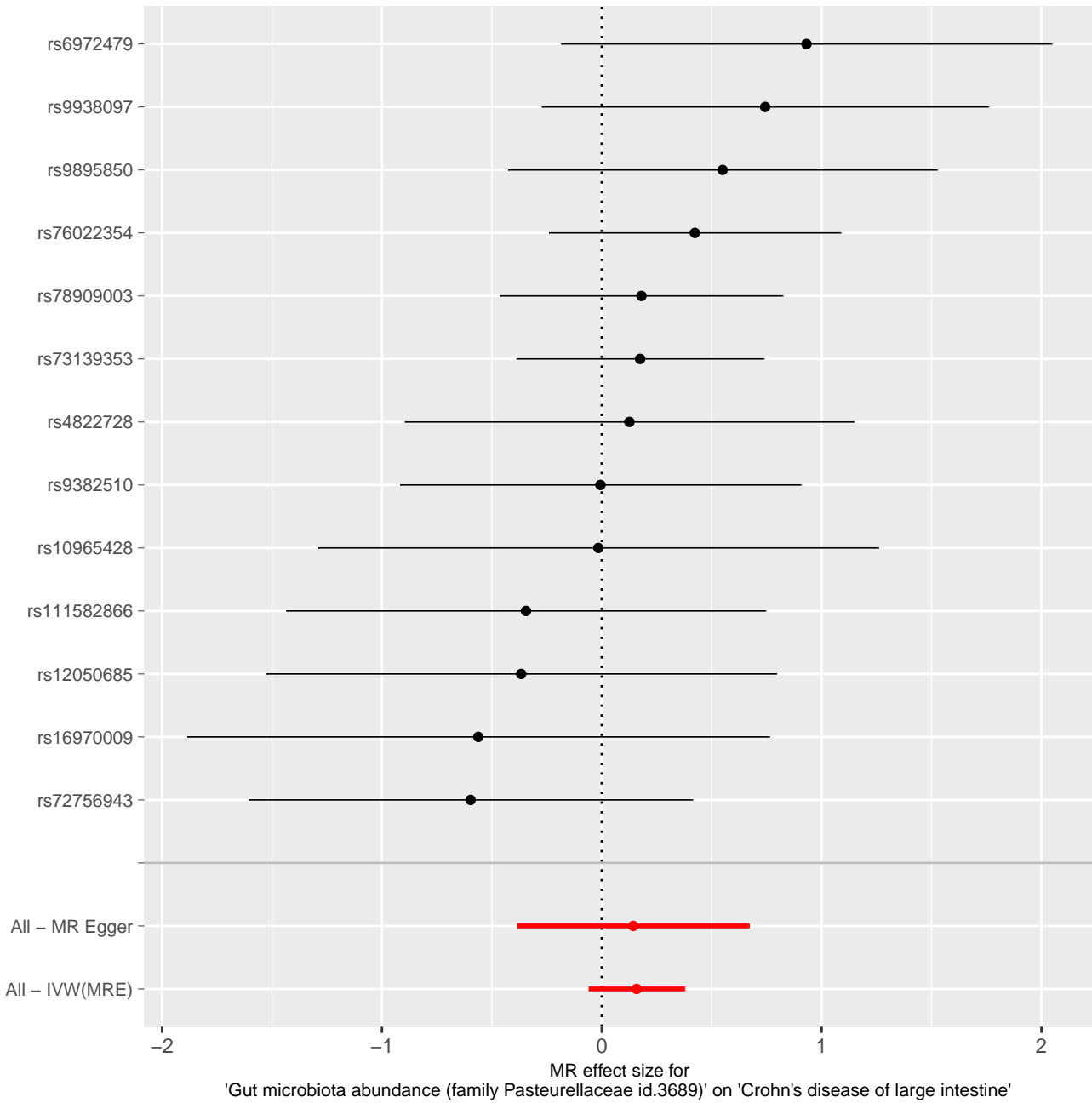
Batch 19 : Gut microbiota abundance (family Methanobacteriaceae id.121) on Crohn's disease of large intestine

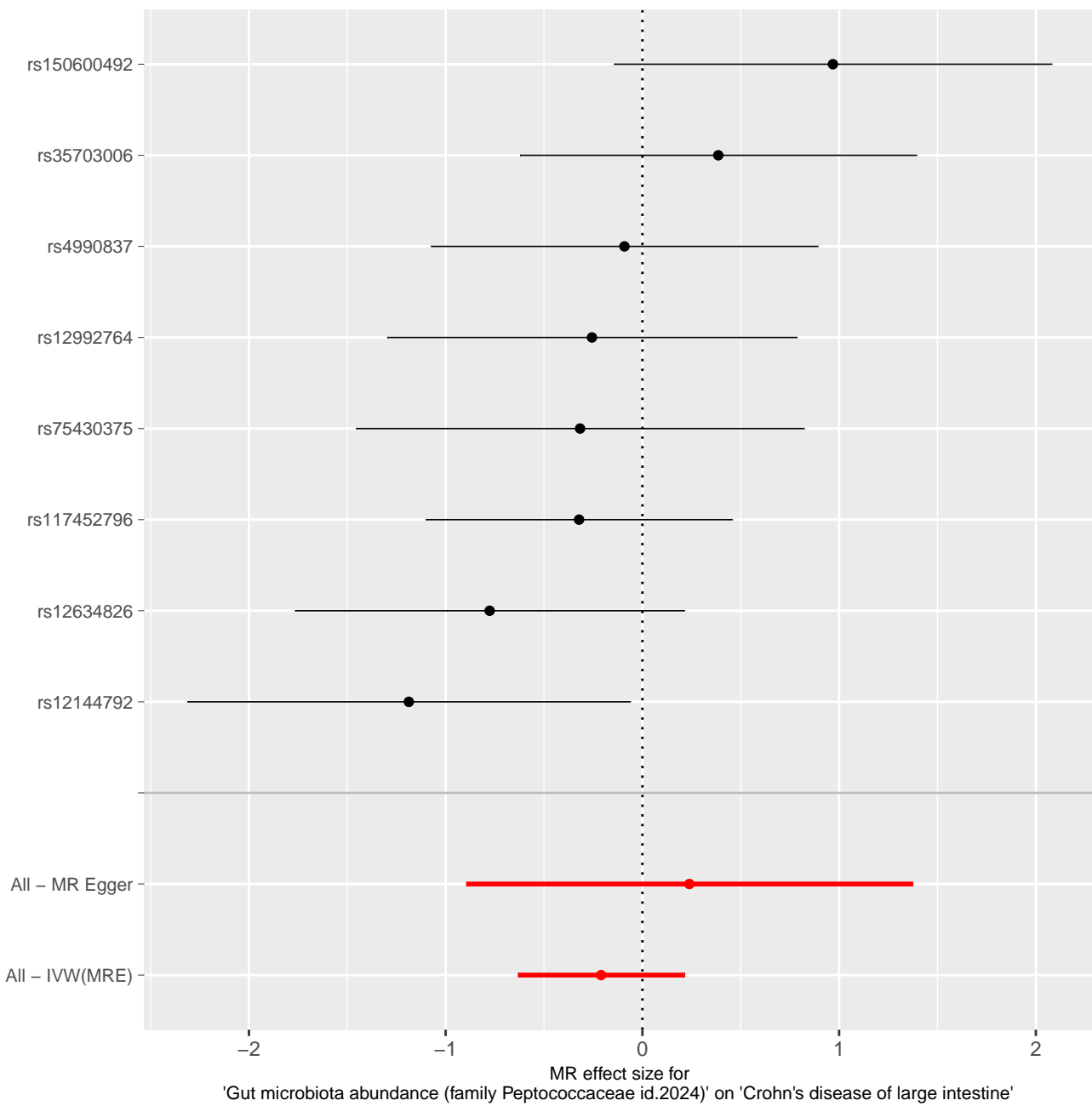


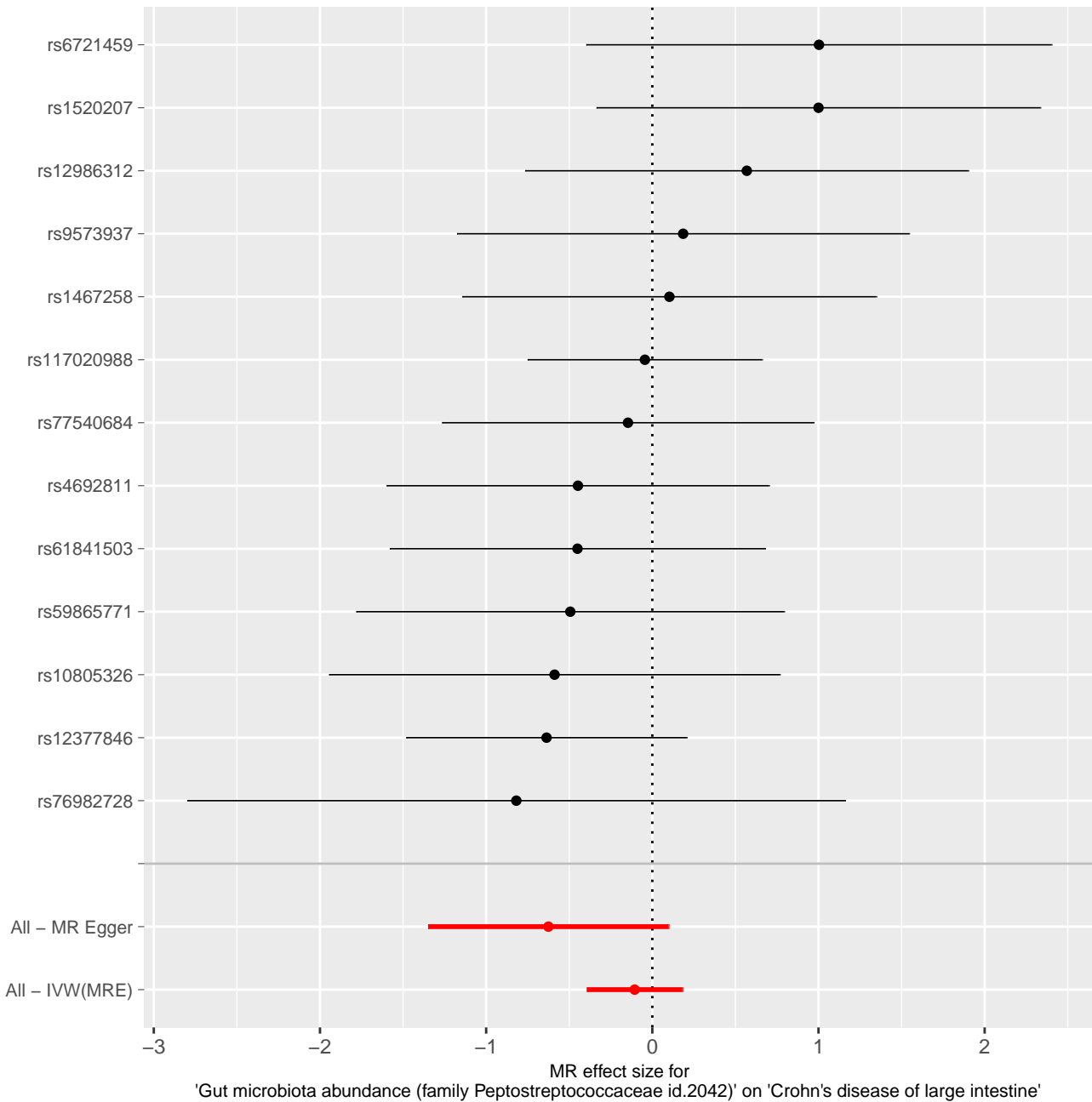
Batch 20 : Gut microbiota abundance (family Oxalobacteraceae id.2966) on Crohn's disease of large intestine

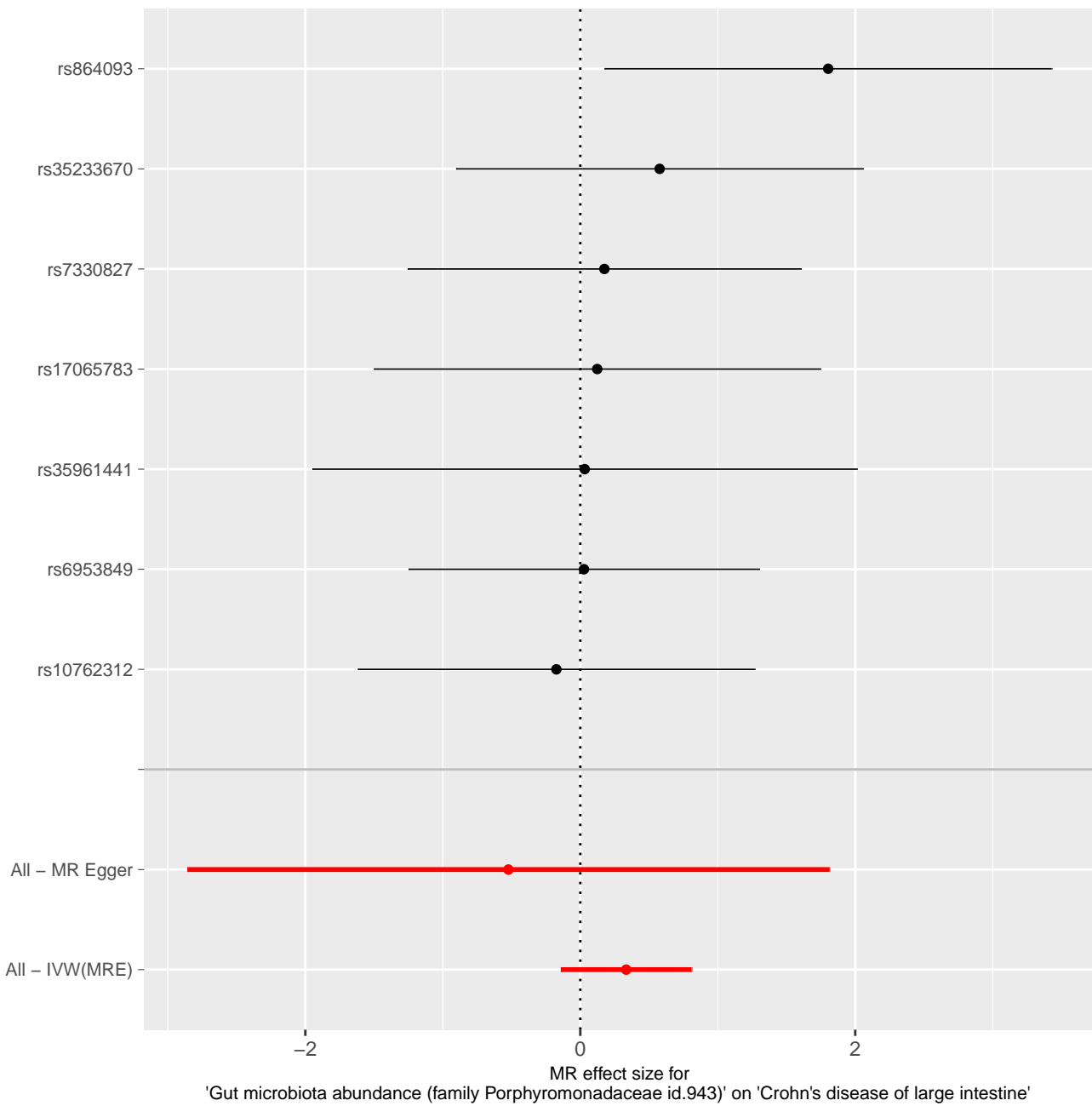


Batch 21 : Gut microbiota abundance (family Pasteurellaceae id.3689) on Crohn's disease of large intestine

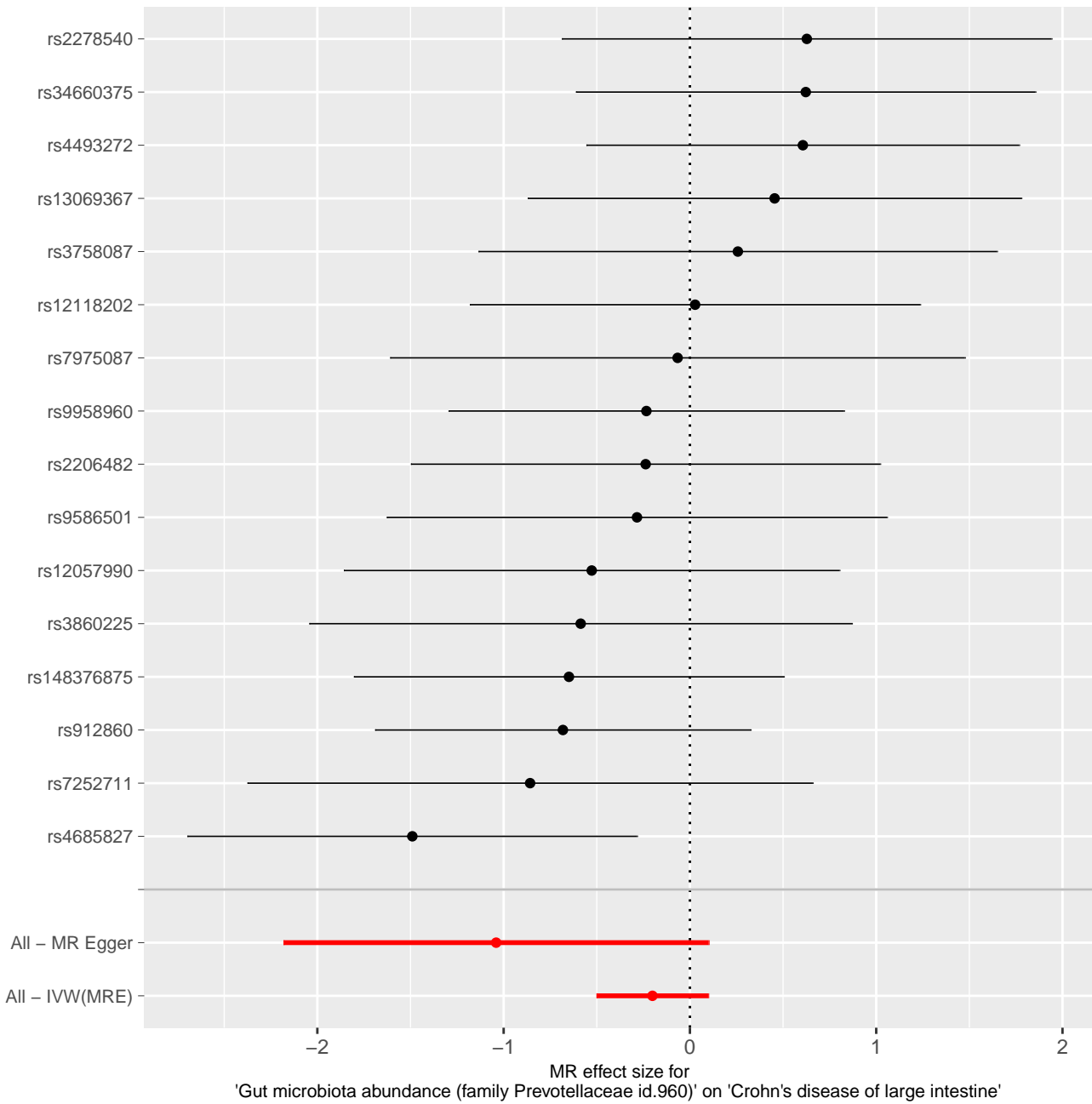




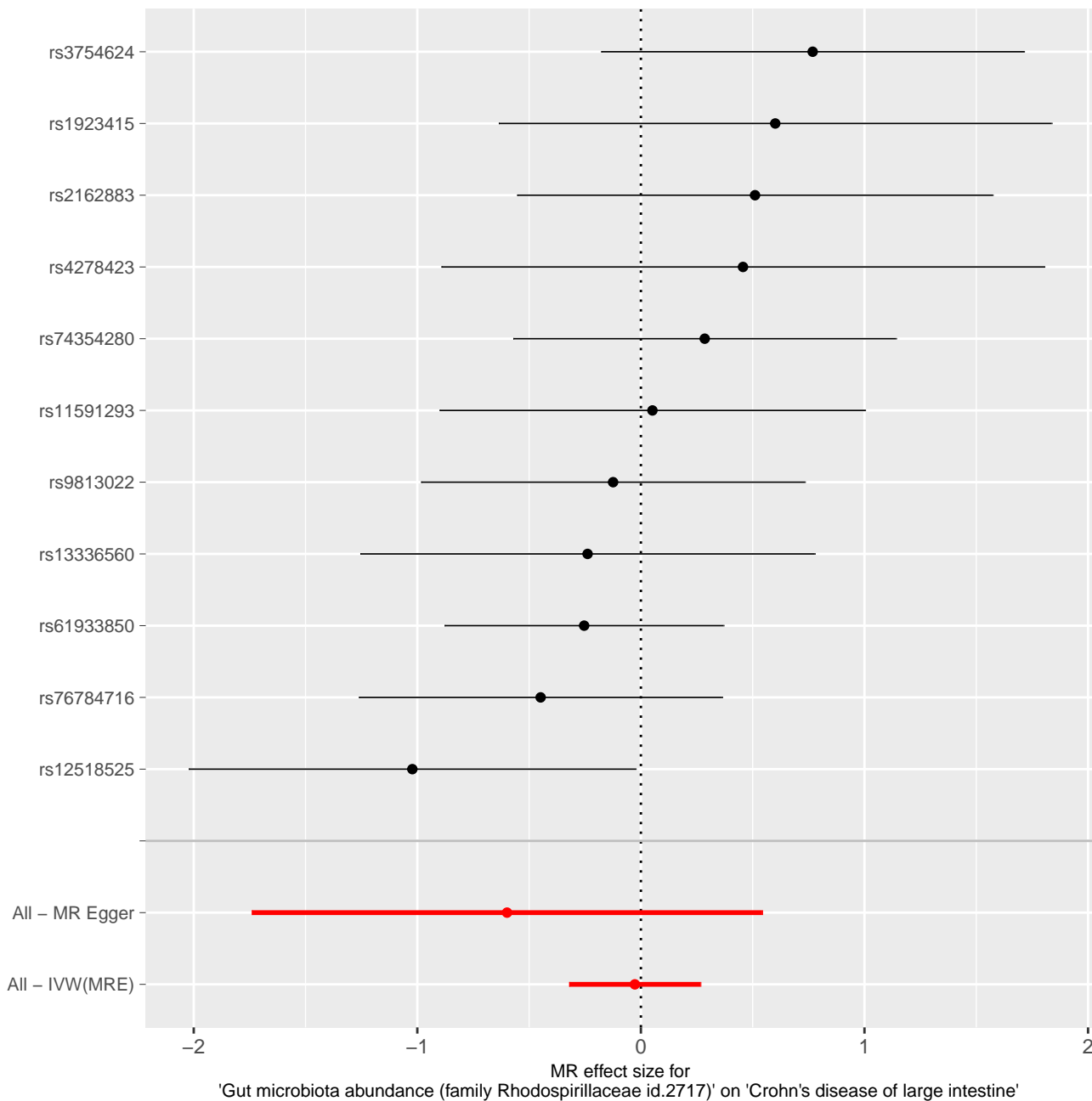




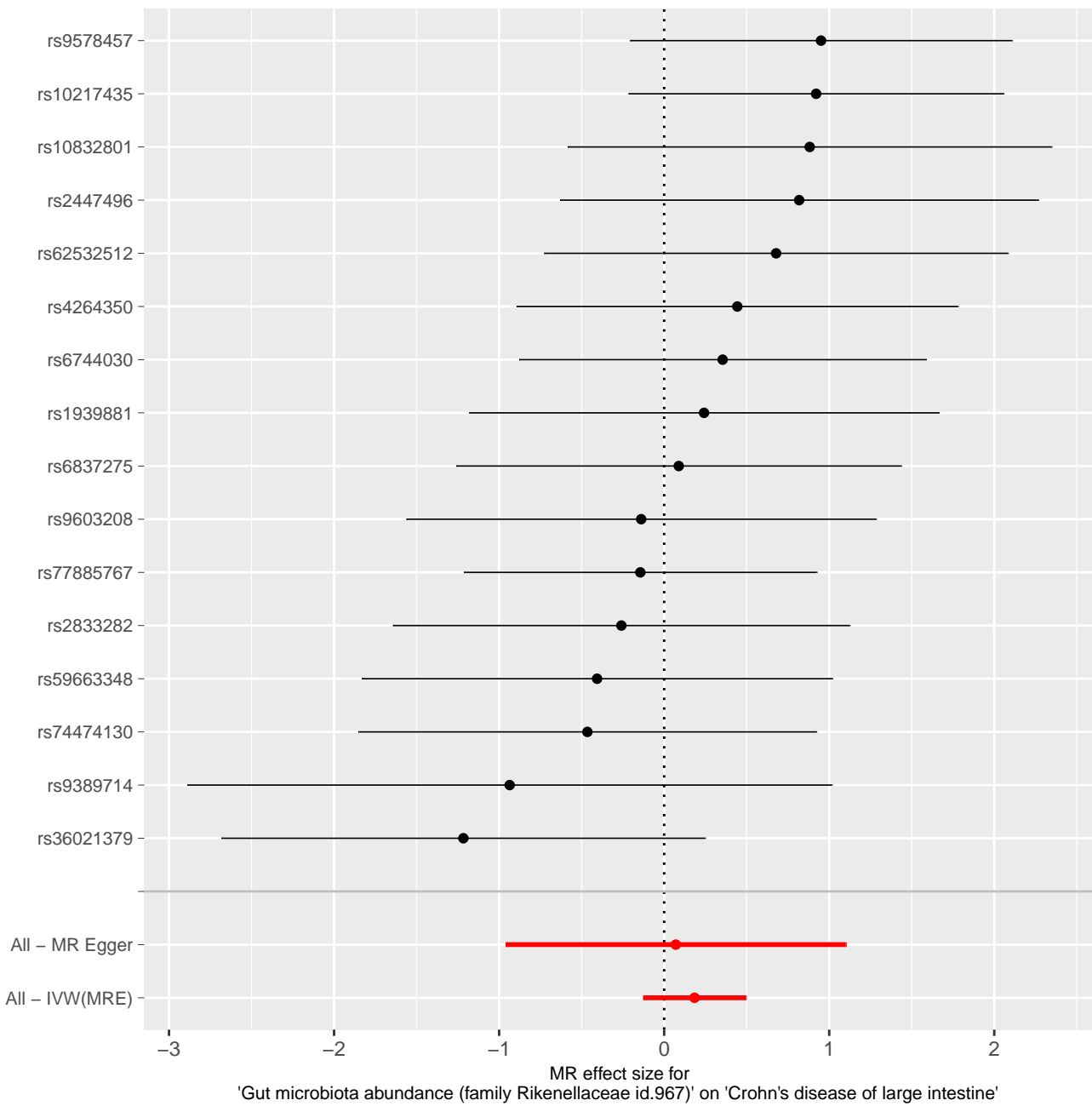
Batch 25 : Gut microbiota abundance (family Prevotellaceae id.960) on Crohn's disease of large intestine

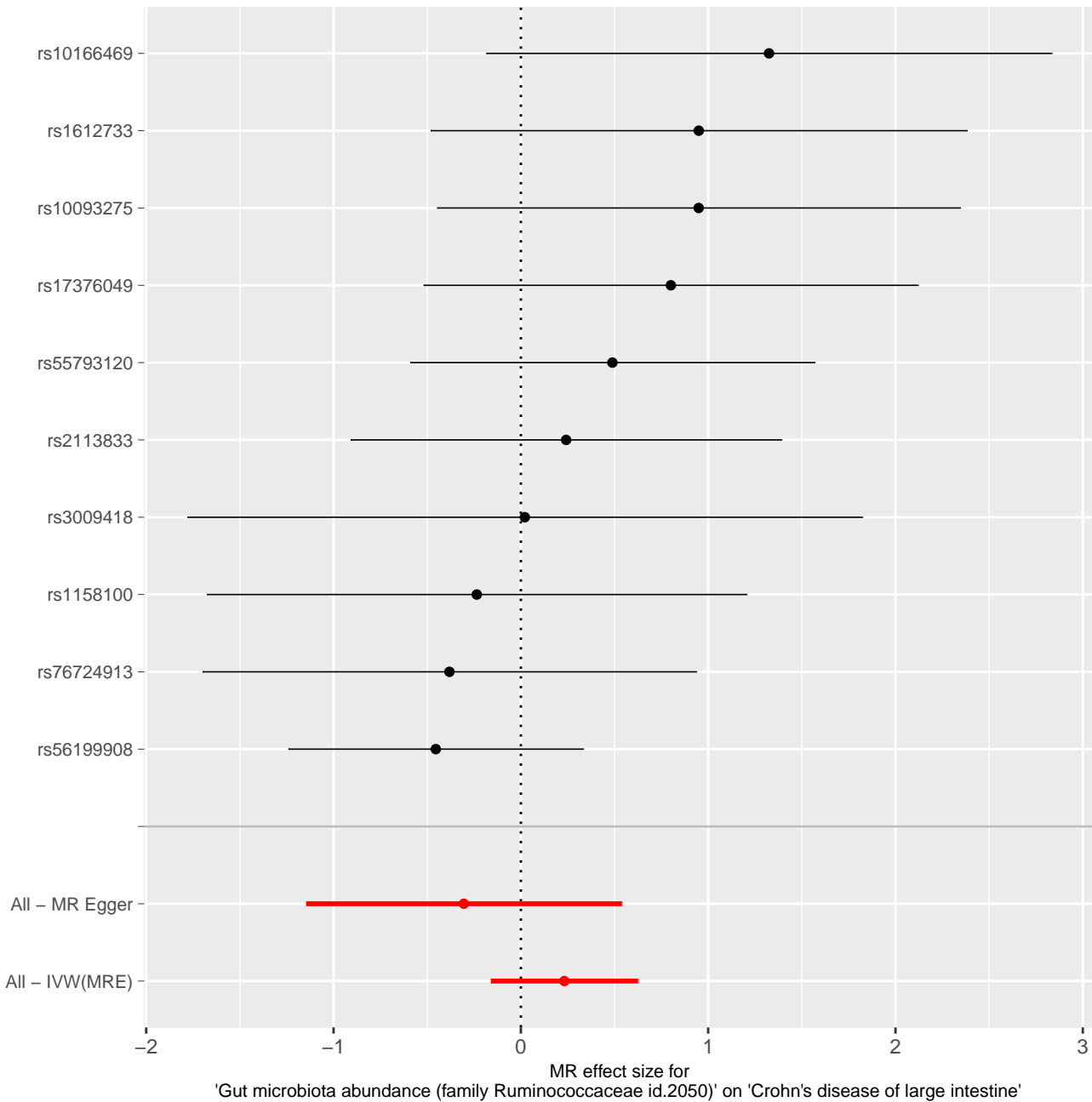


Batch 26 : Gut microbiota abundance (family Rhodospirillaceae id.2717) on Crohn's disease of large intestine

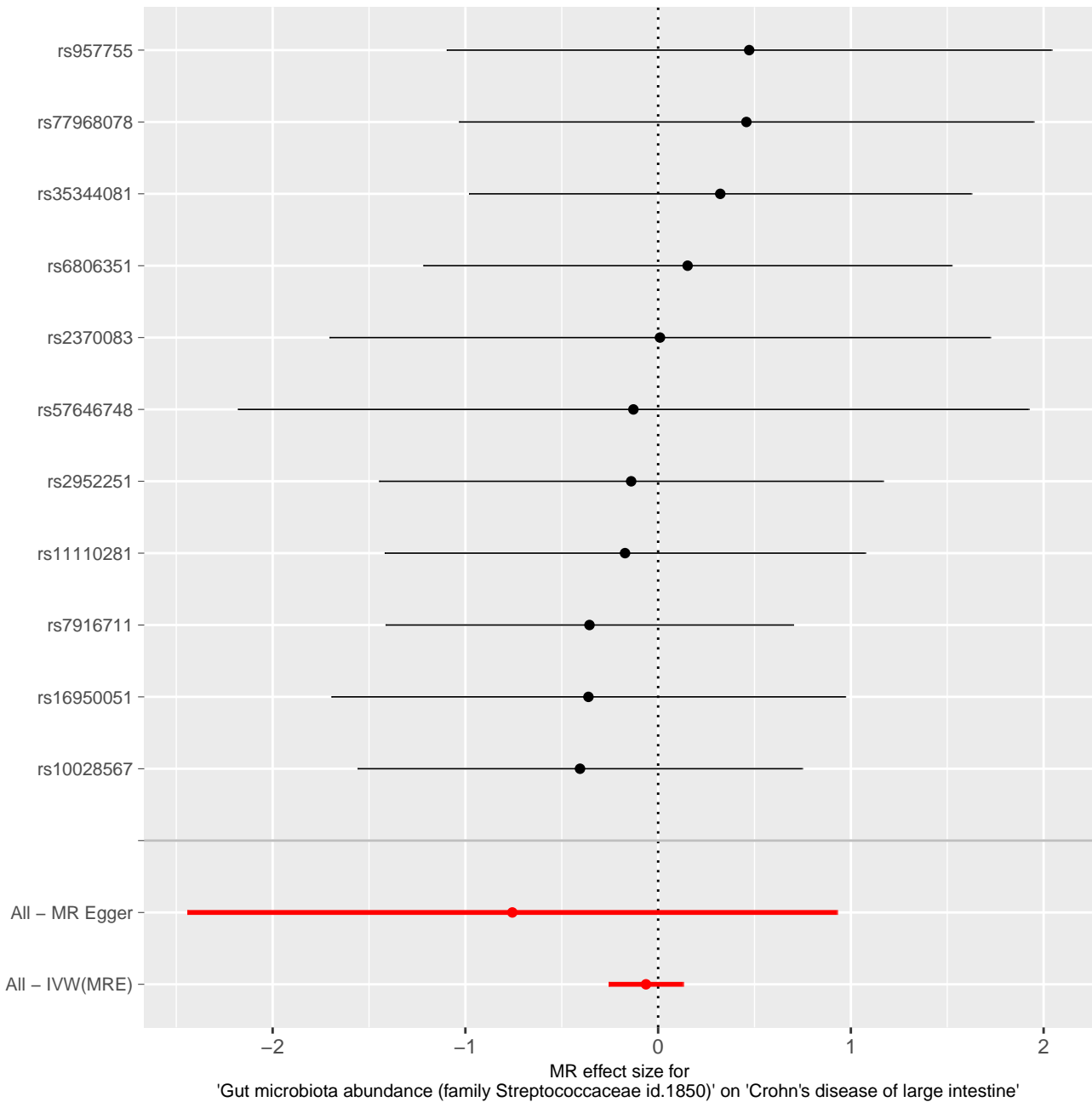


Batch 27 : Gut microbiota abundance (family Rikenellaceae id.967) on Crohn's disease of large intestine

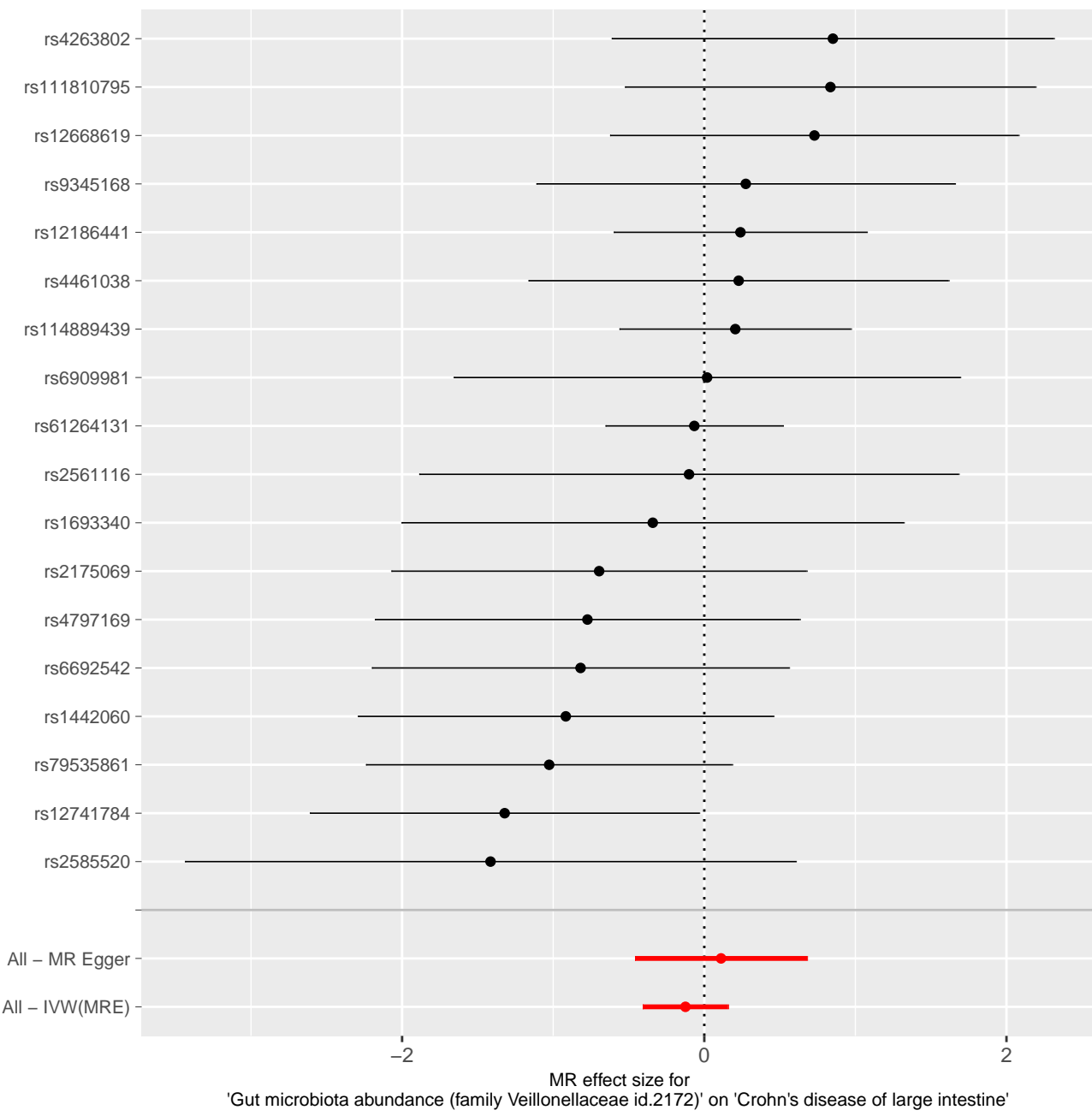




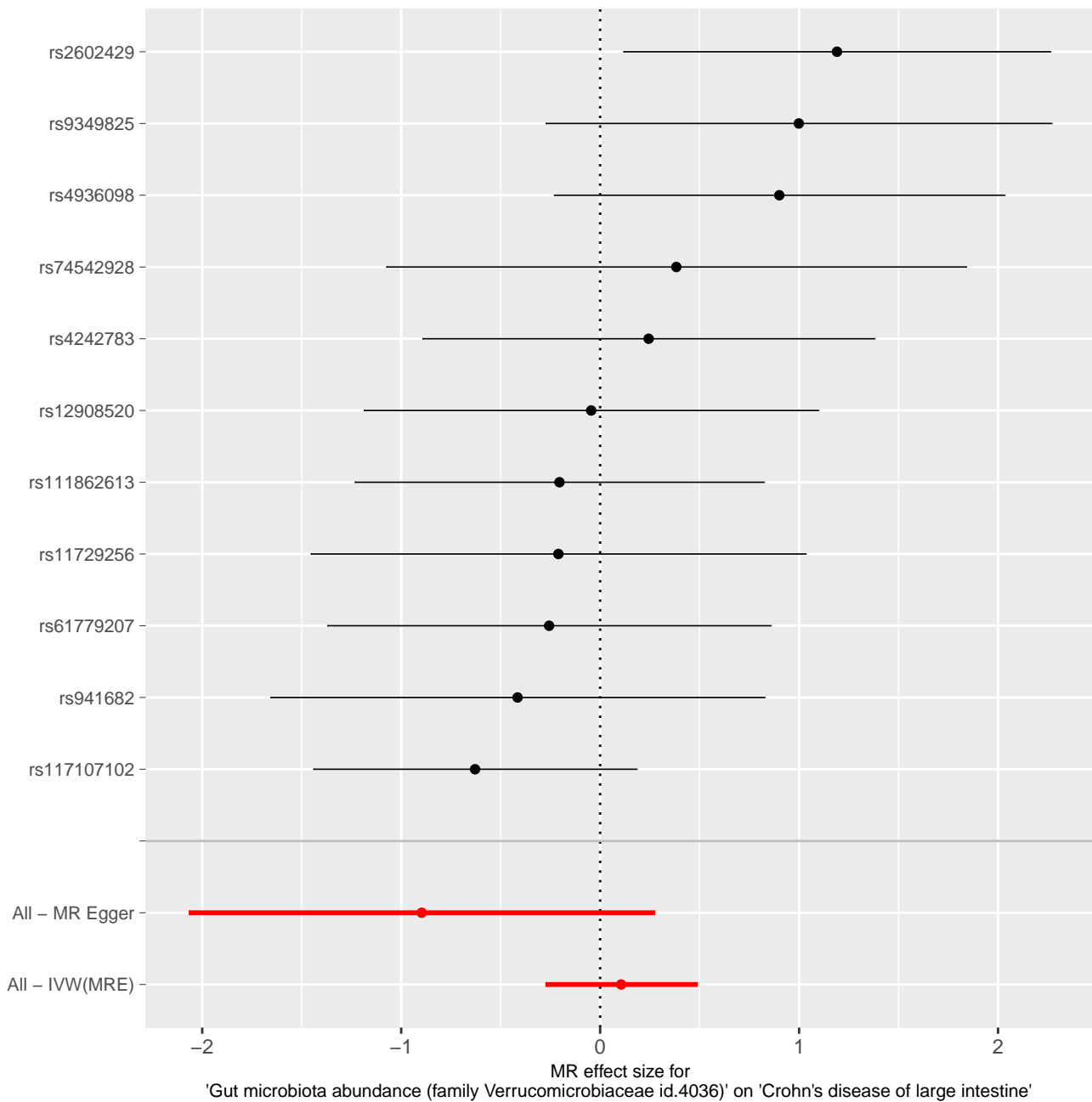
Batch 29 : Gut microbiota abundance (family Streptococcaceae id.1850) on Crohn's disease of large intestine



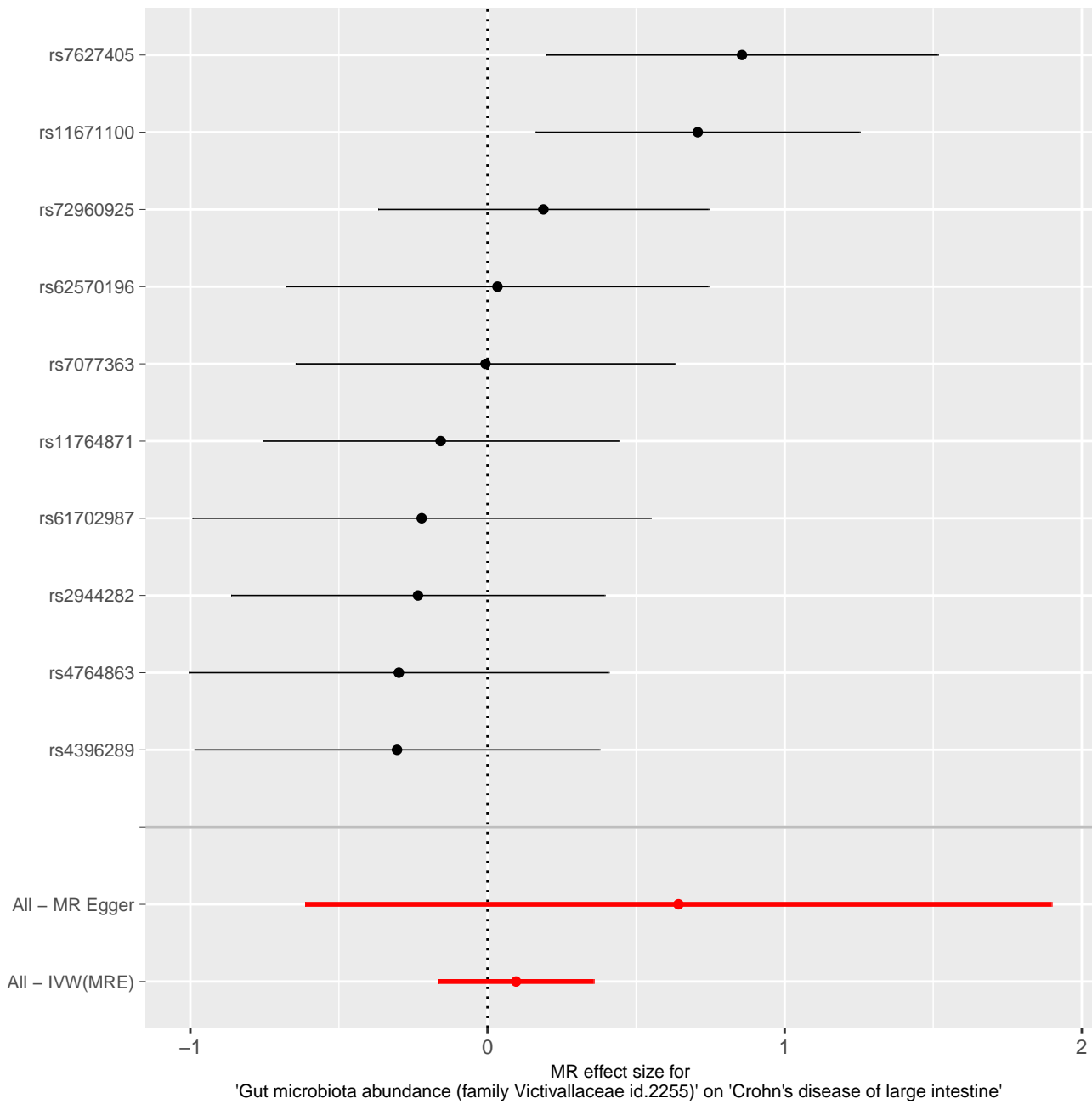
Batch 30 : Gut microbiota abundance (family Veillonellaceae id.2172) on Crohn's disease of large intestine

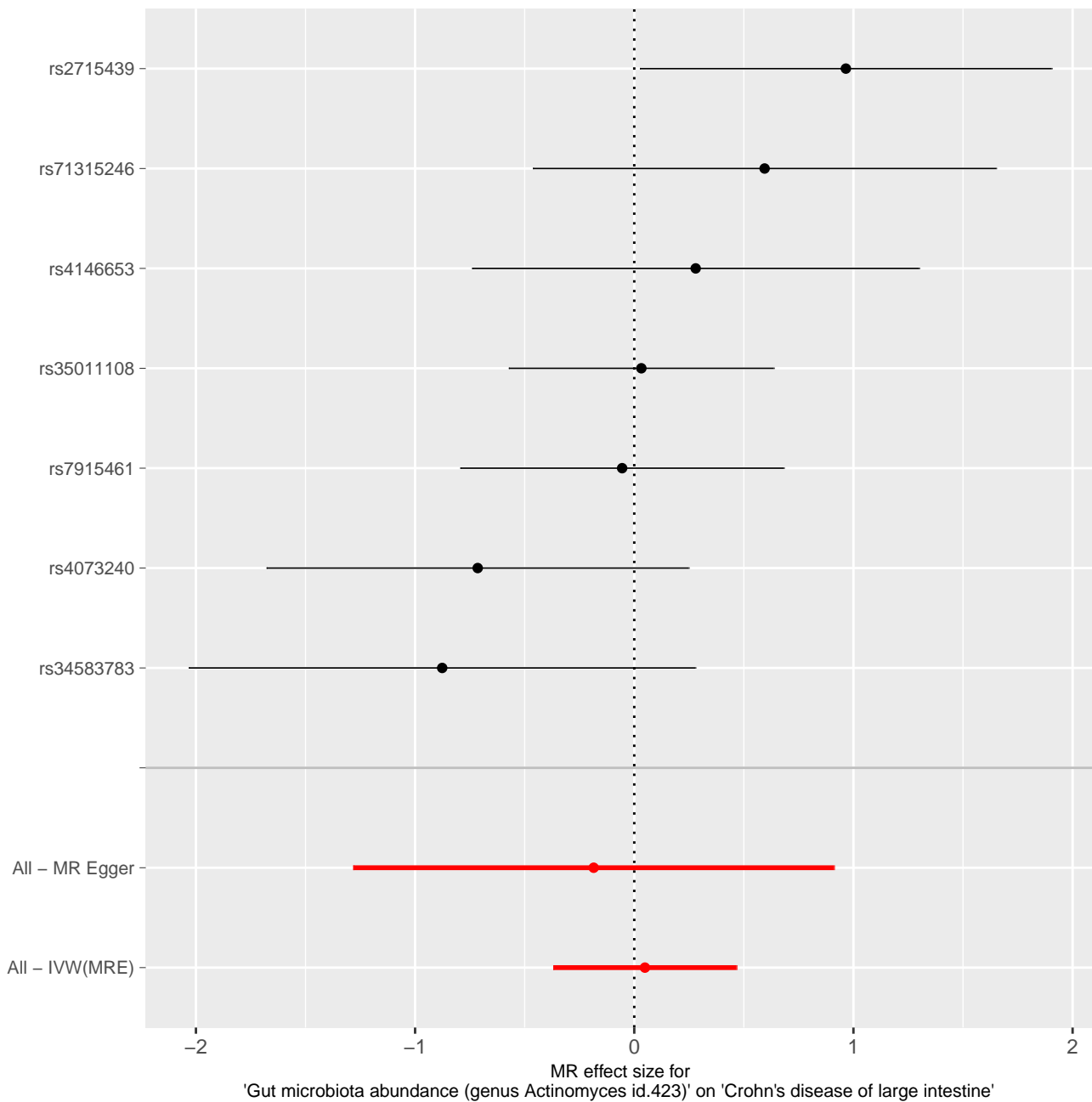


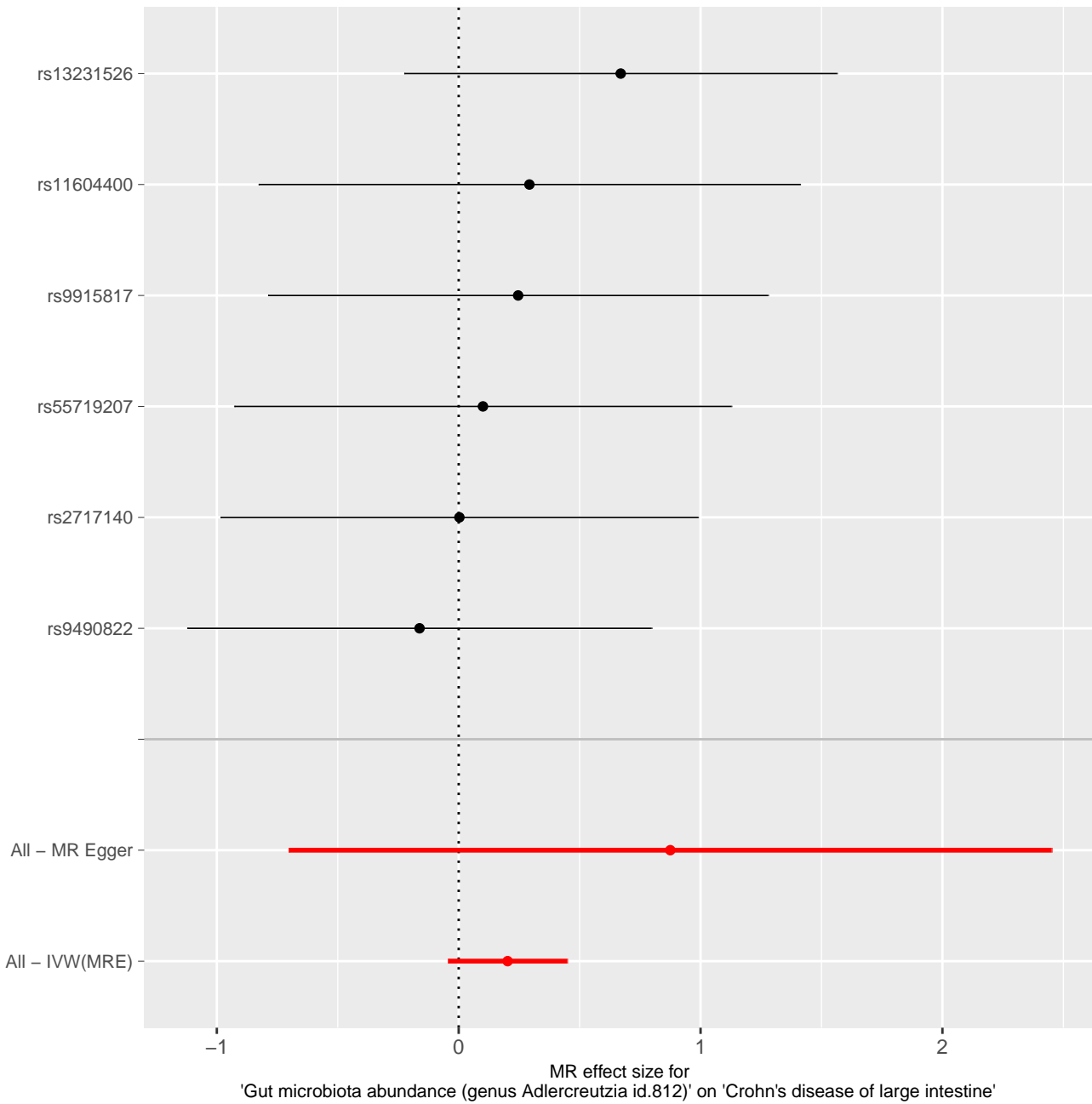
Batch 31 : Gut microbiota abundance (family Verrucomicrobiaceae id.4036) on Crohn's disease of large intestine



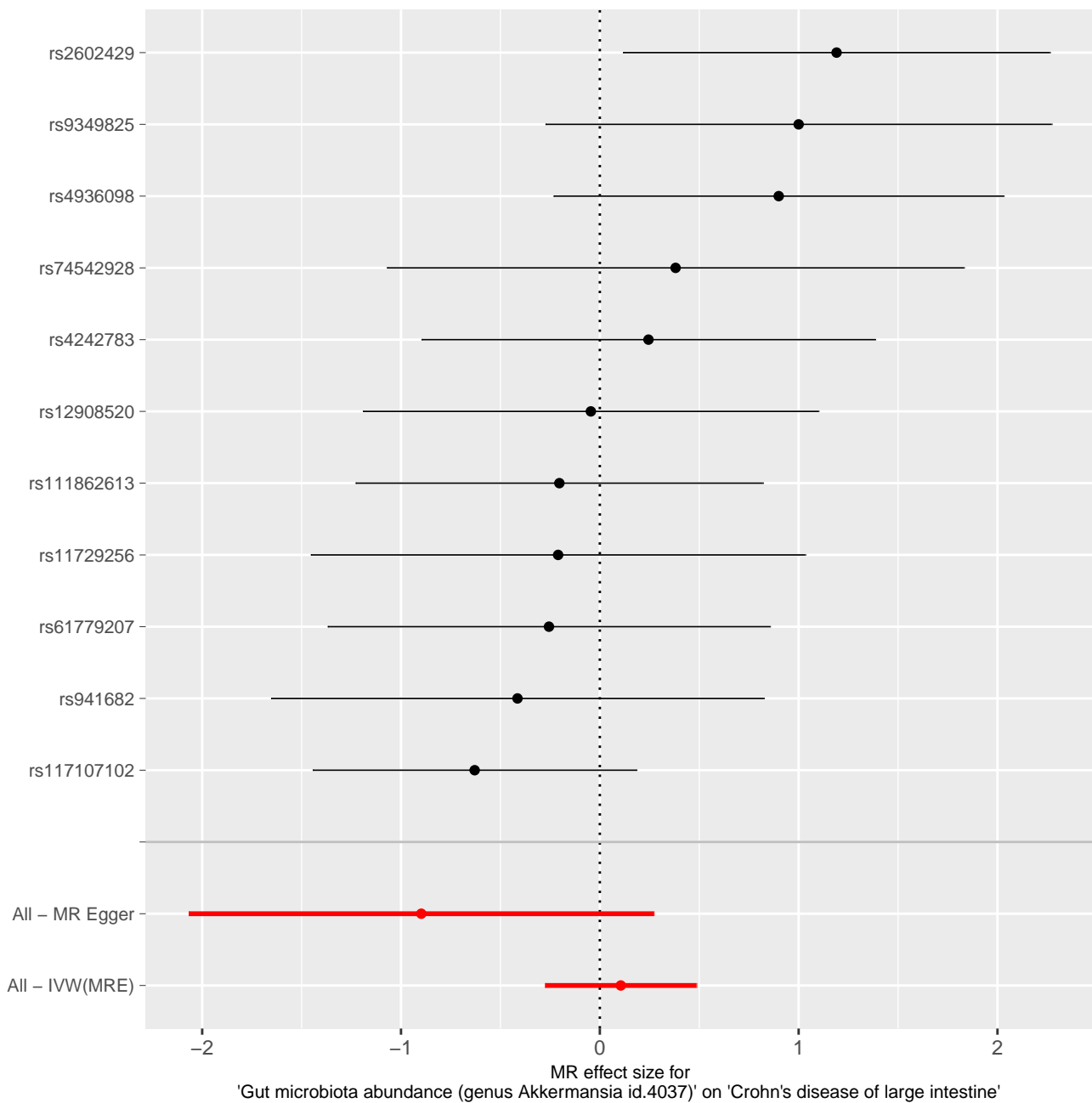
Batch 32 : Gut microbiota abundance (family Victivallaceae id.2255) on Crohn's disease of large intestine



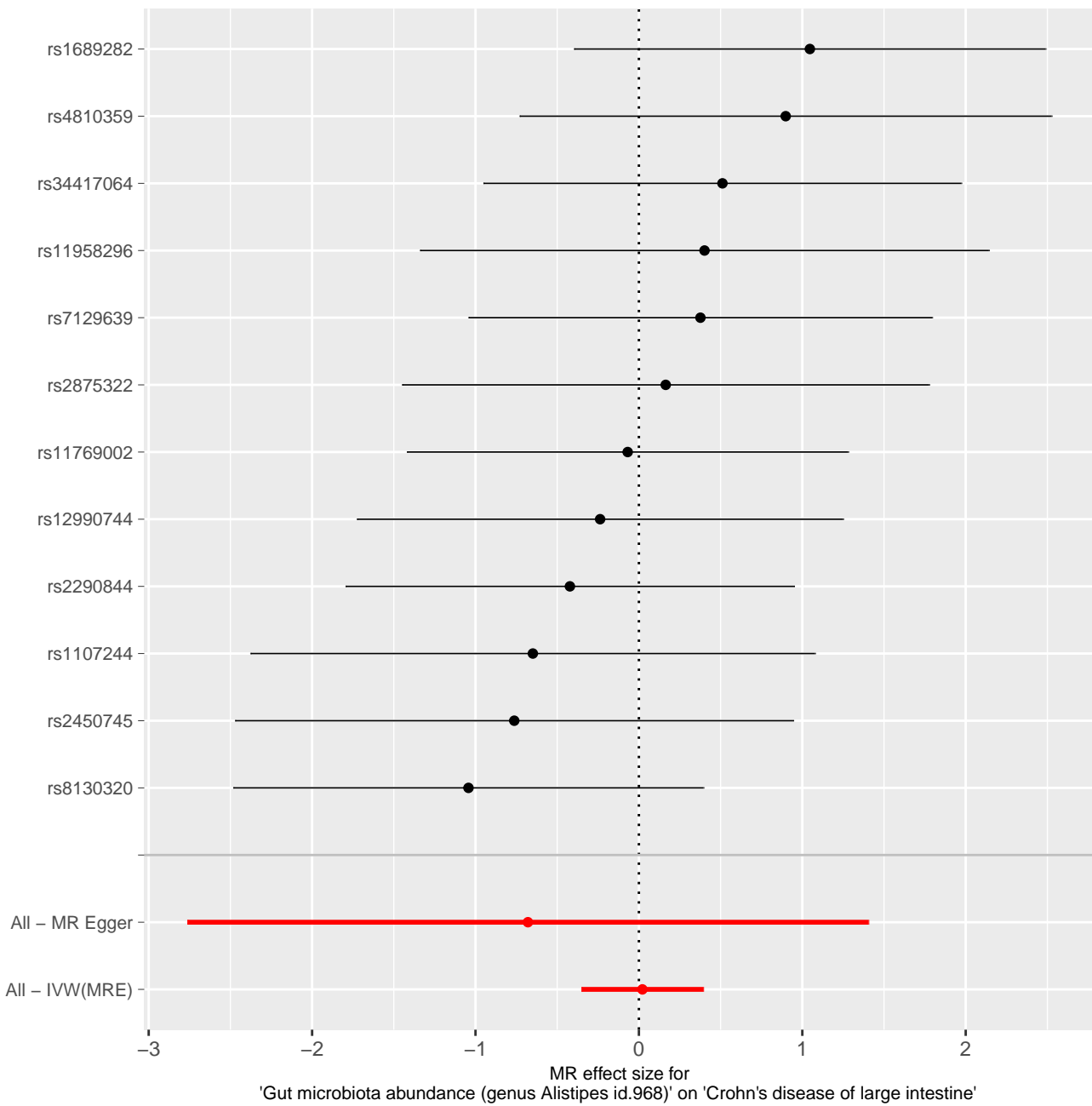




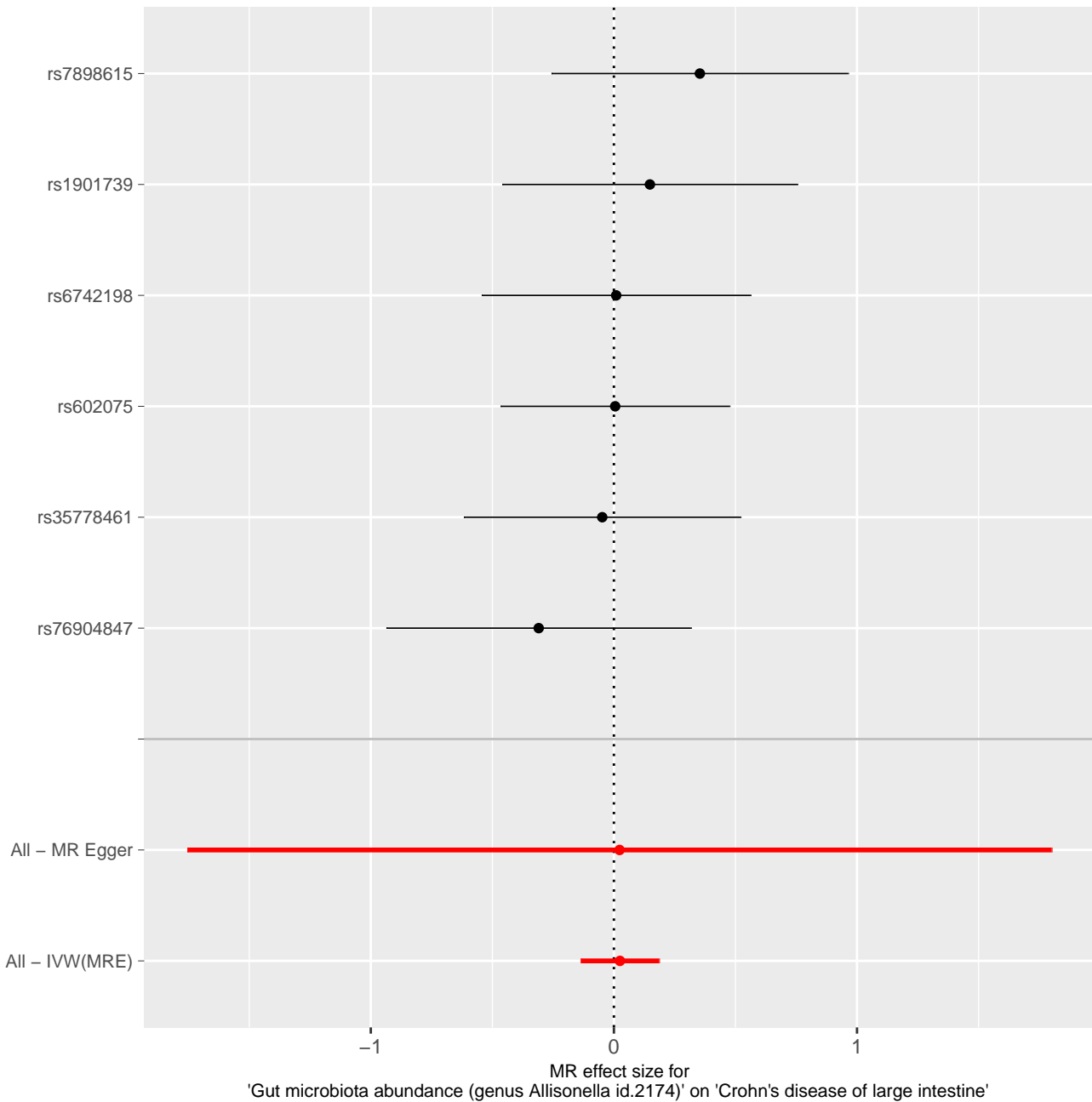
Batch 35 : Gut microbiota abundance (genus Akkermansia id.4037) on Crohn's disease of large intestine

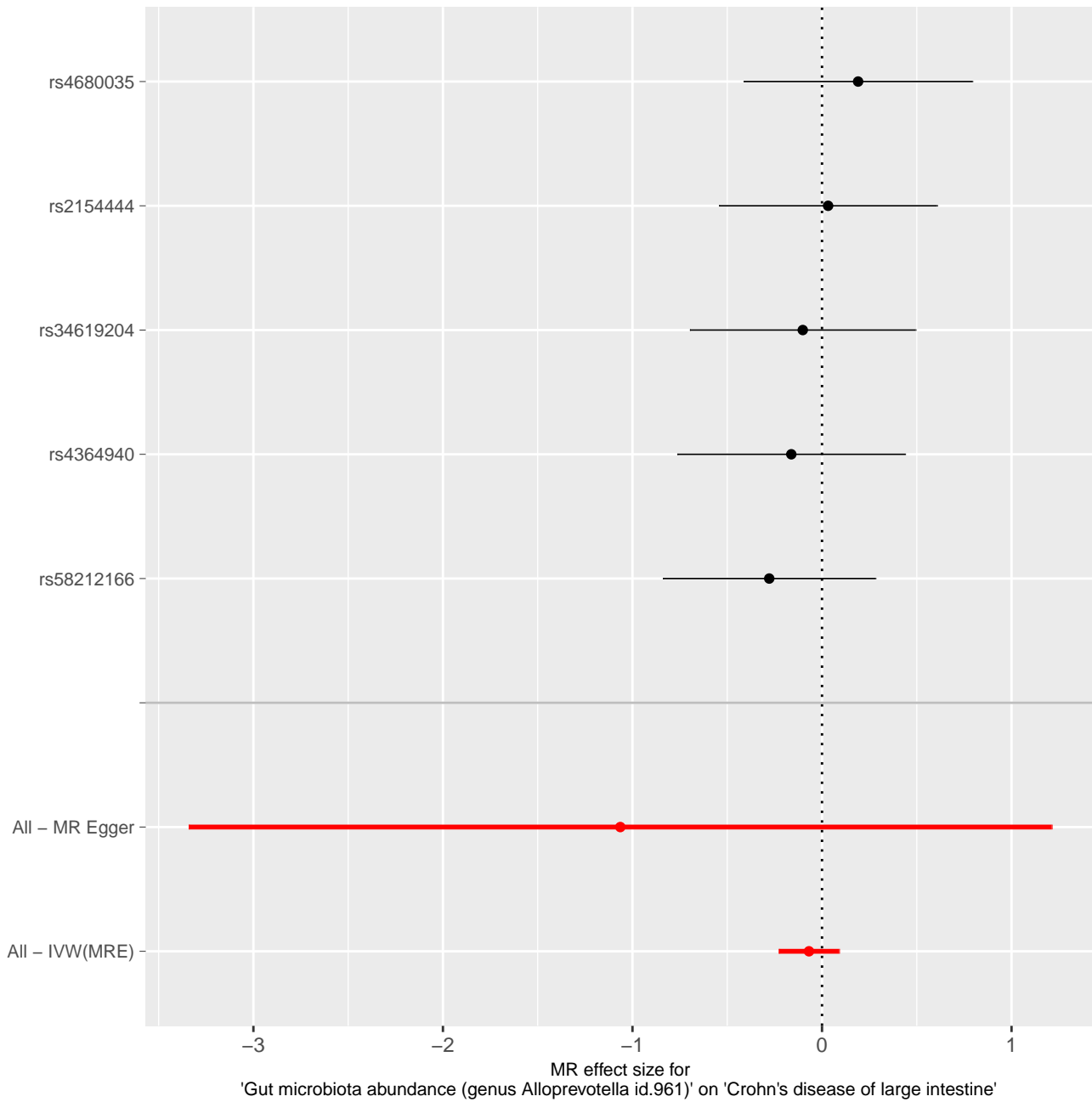


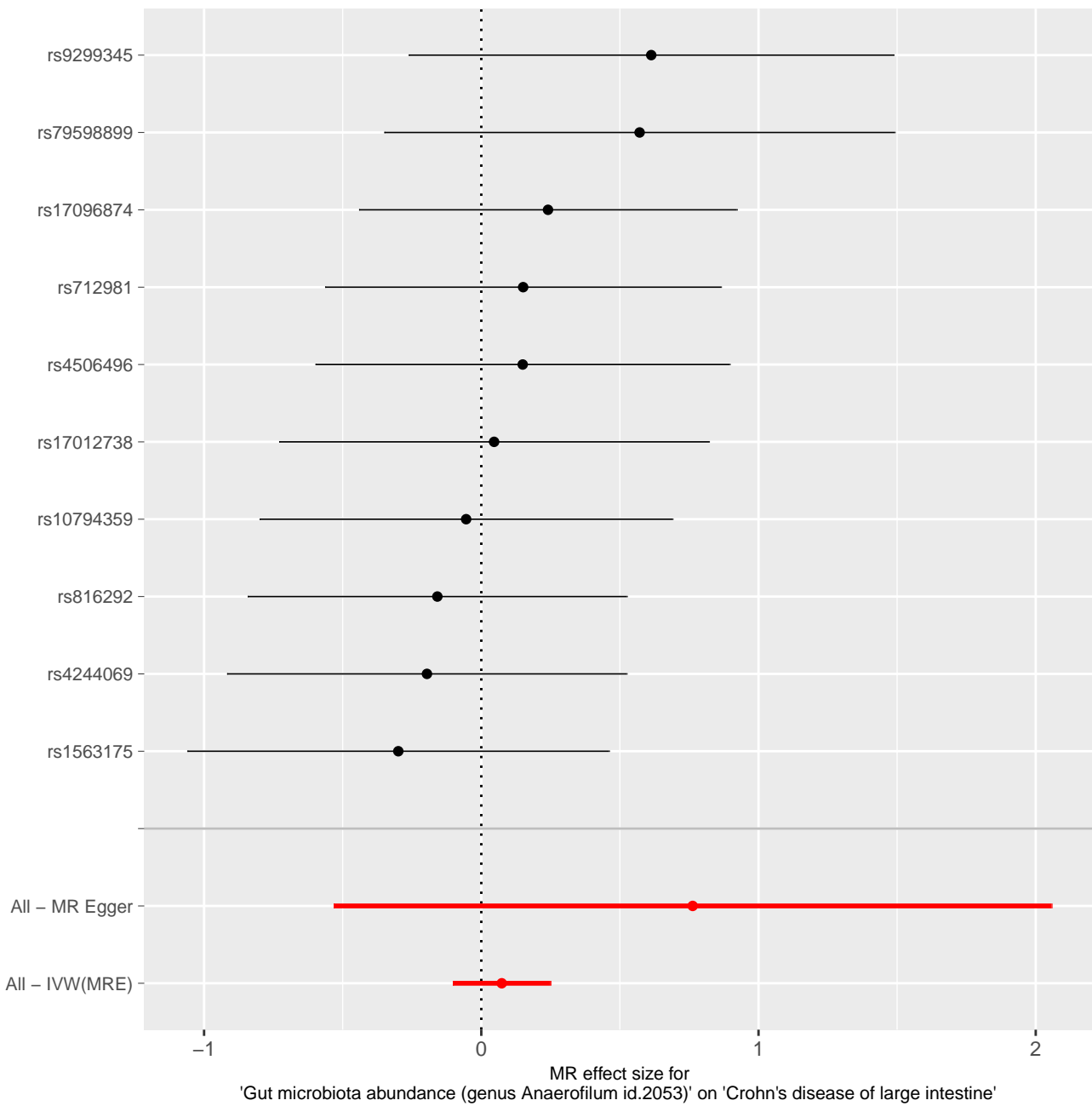
Batch 36 : Gut microbiota abundance (genus Alistipes id.968) on Crohn's disease of large intestine



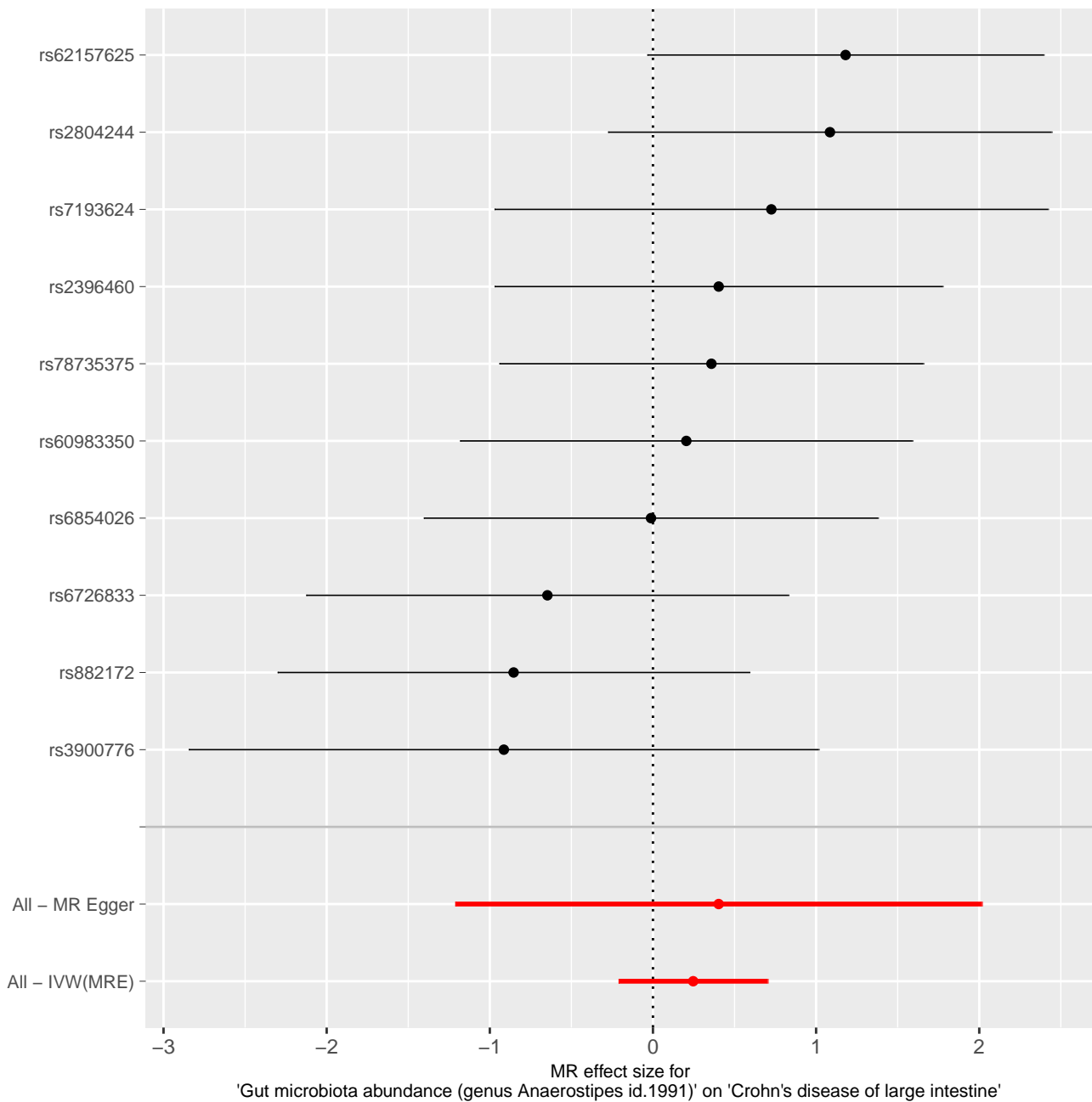
Batch 37 : Gut microbiota abundance (genus Allisonella id.2174) on Crohn's disease of large intestine



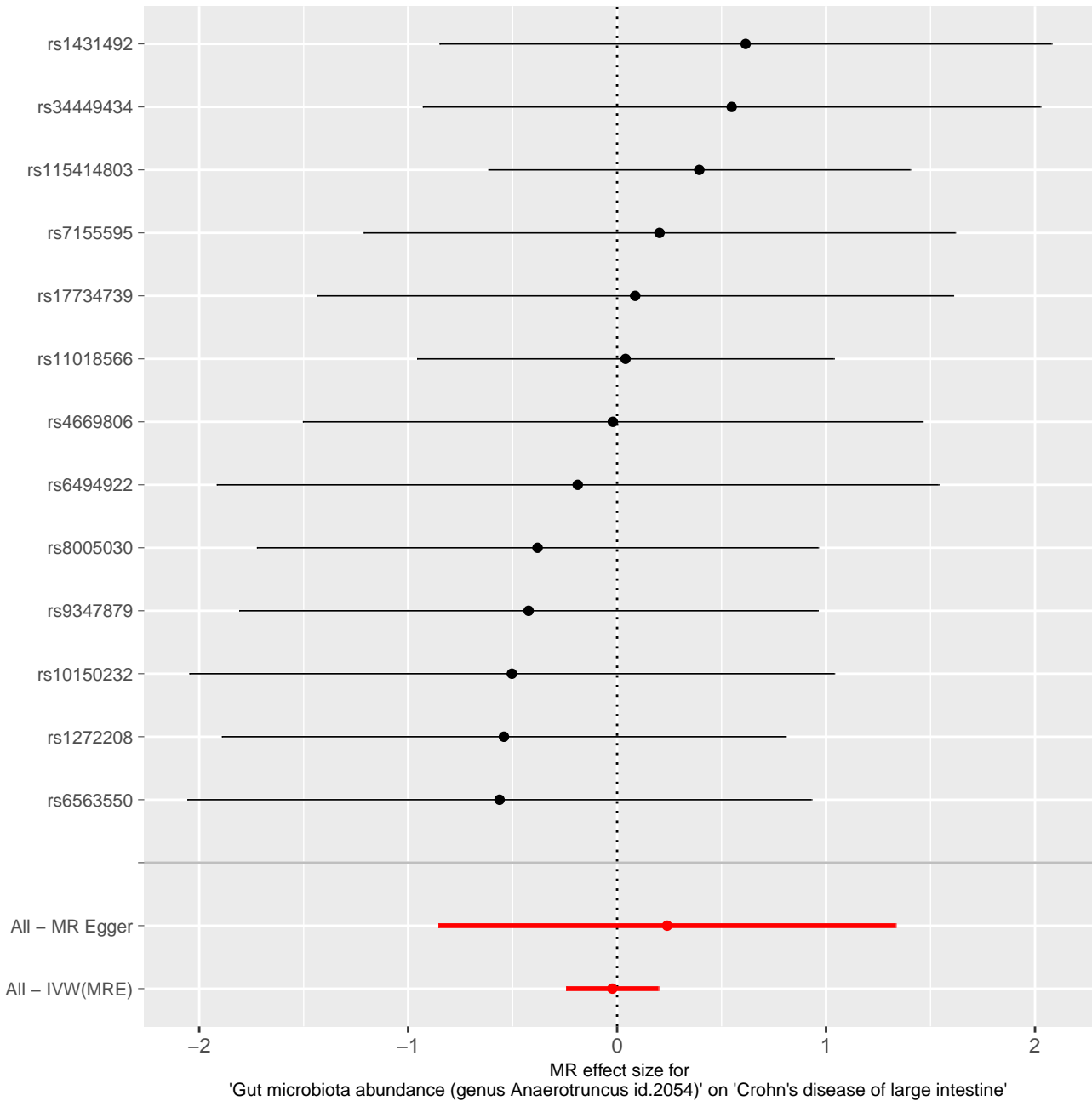




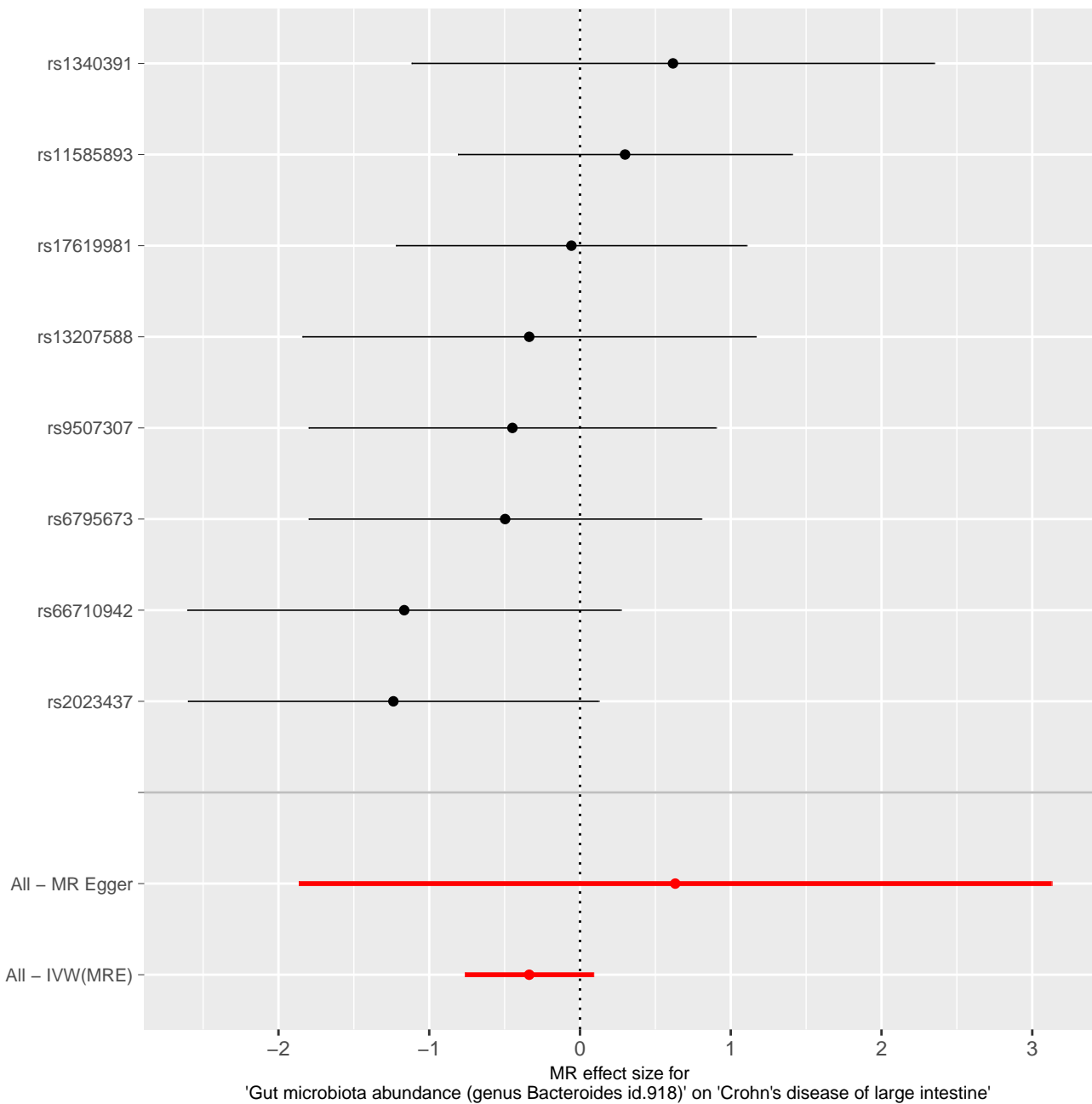
Batch 40 : Gut microbiota abundance (genus Anaerostipes id.1991) on Crohn's disease of large intestine



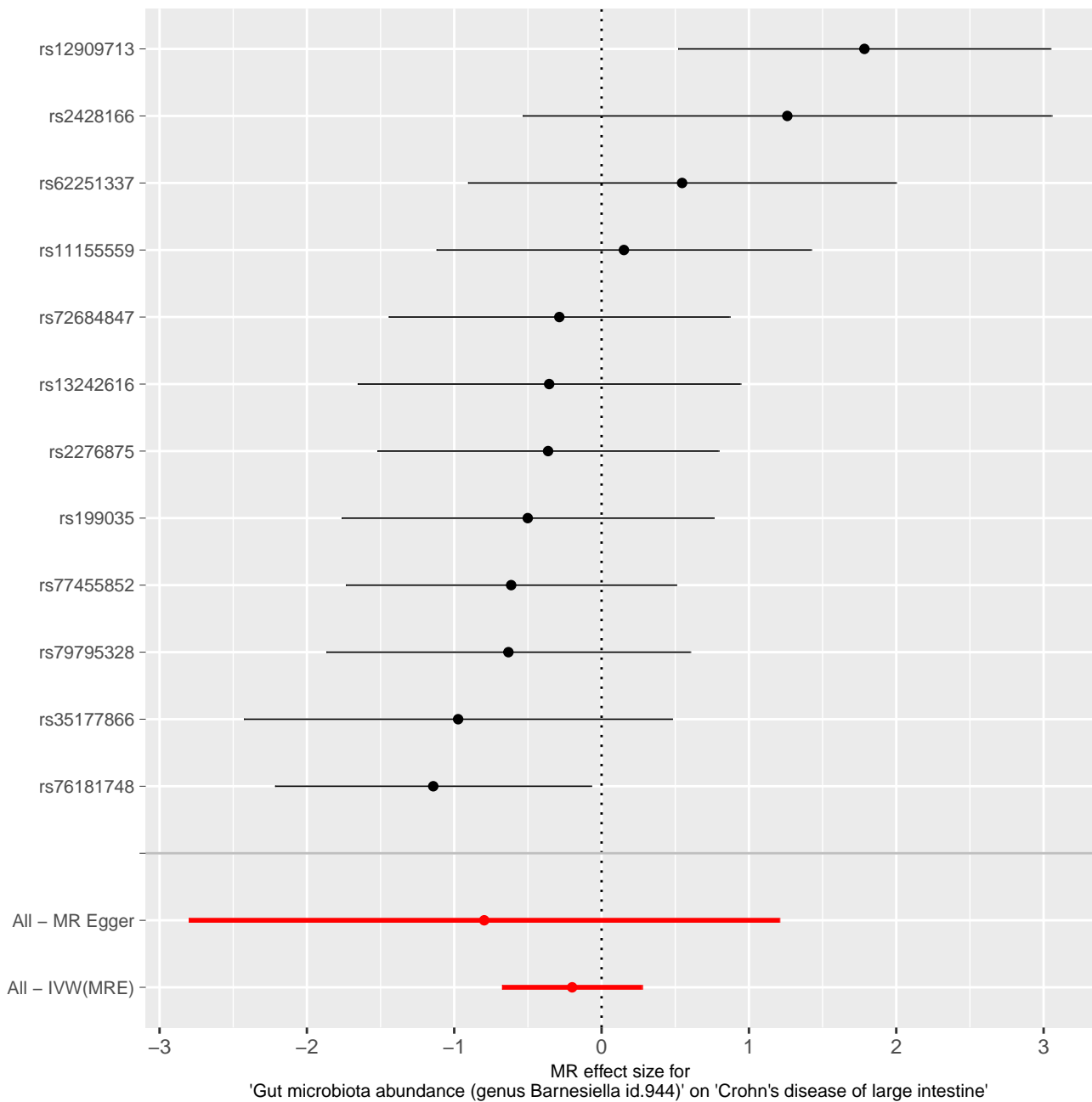
Batch 41 : Gut microbiota abundance (genus Anaerotruncus id.2054) on Crohn's disease of large intestine

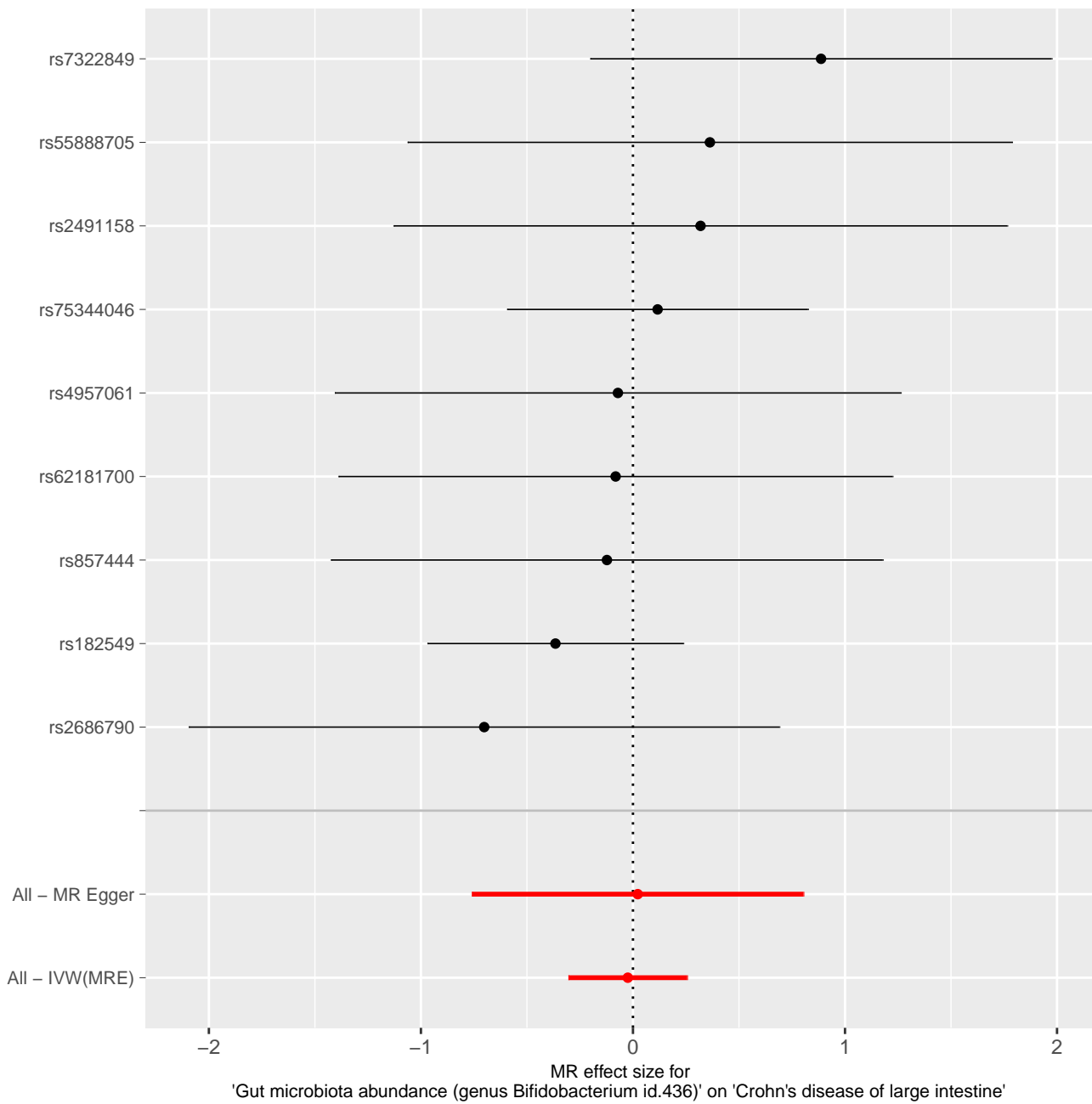


Batch 42 : Gut microbiota abundance (genus Bacteroides id.918) on Crohn's disease of large intestine

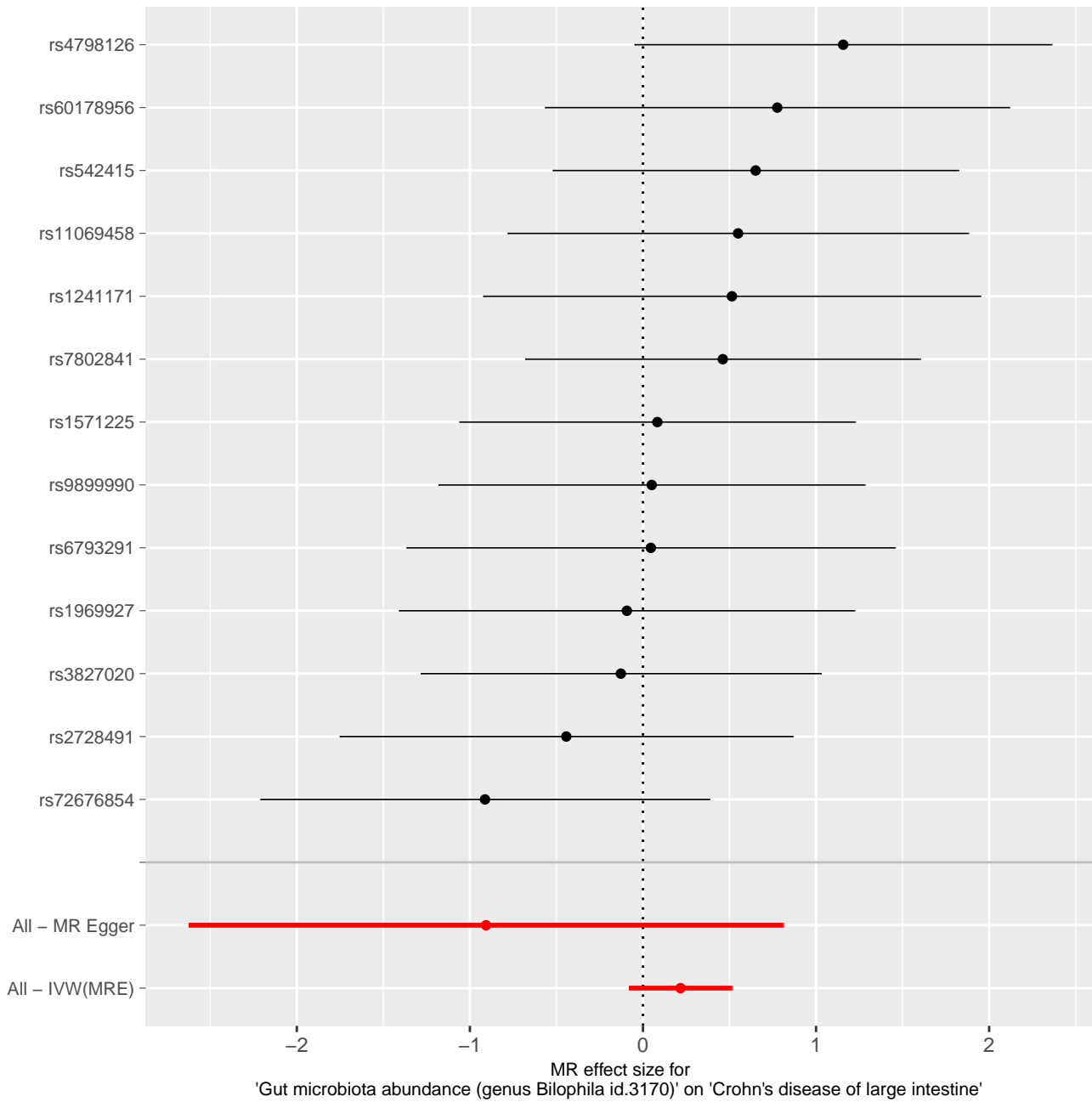


Batch 43 : Gut microbiota abundance (genus *Barnesiella* id.944) on Crohn's disease of large intestine

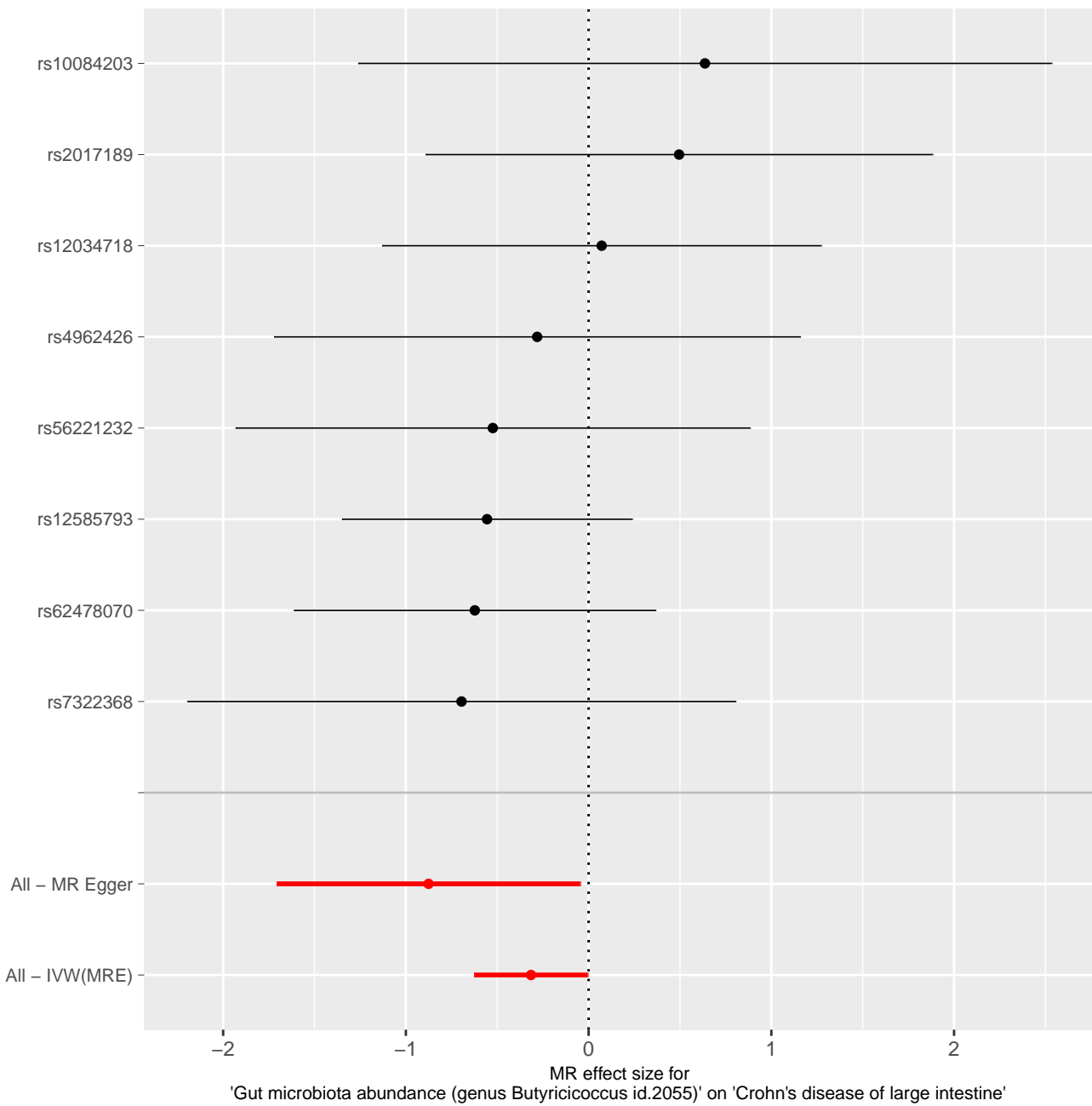




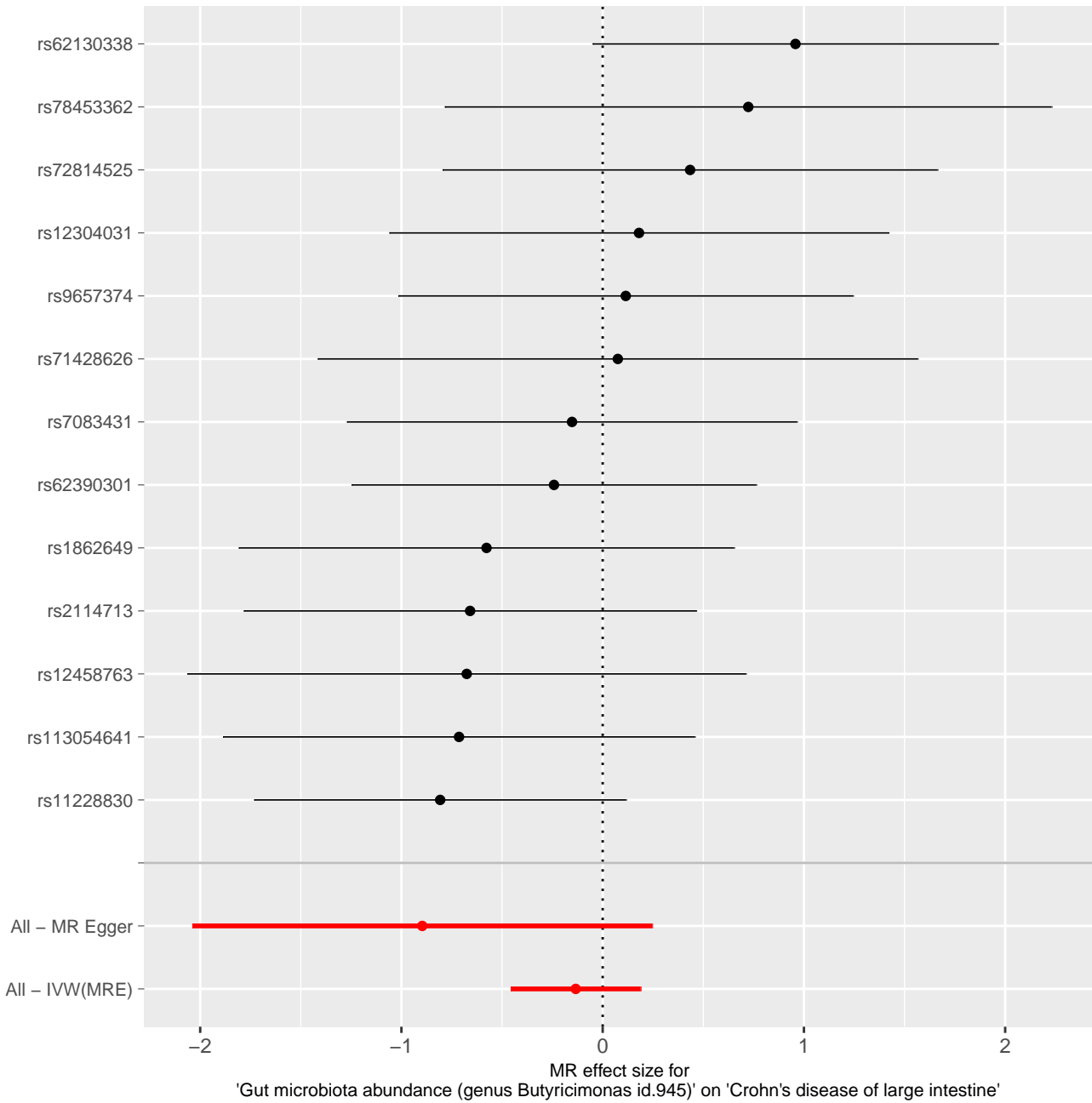
Batch 45 : Gut microbiota abundance (genus Bilophila id.3170) on Crohn's disease of large intestine



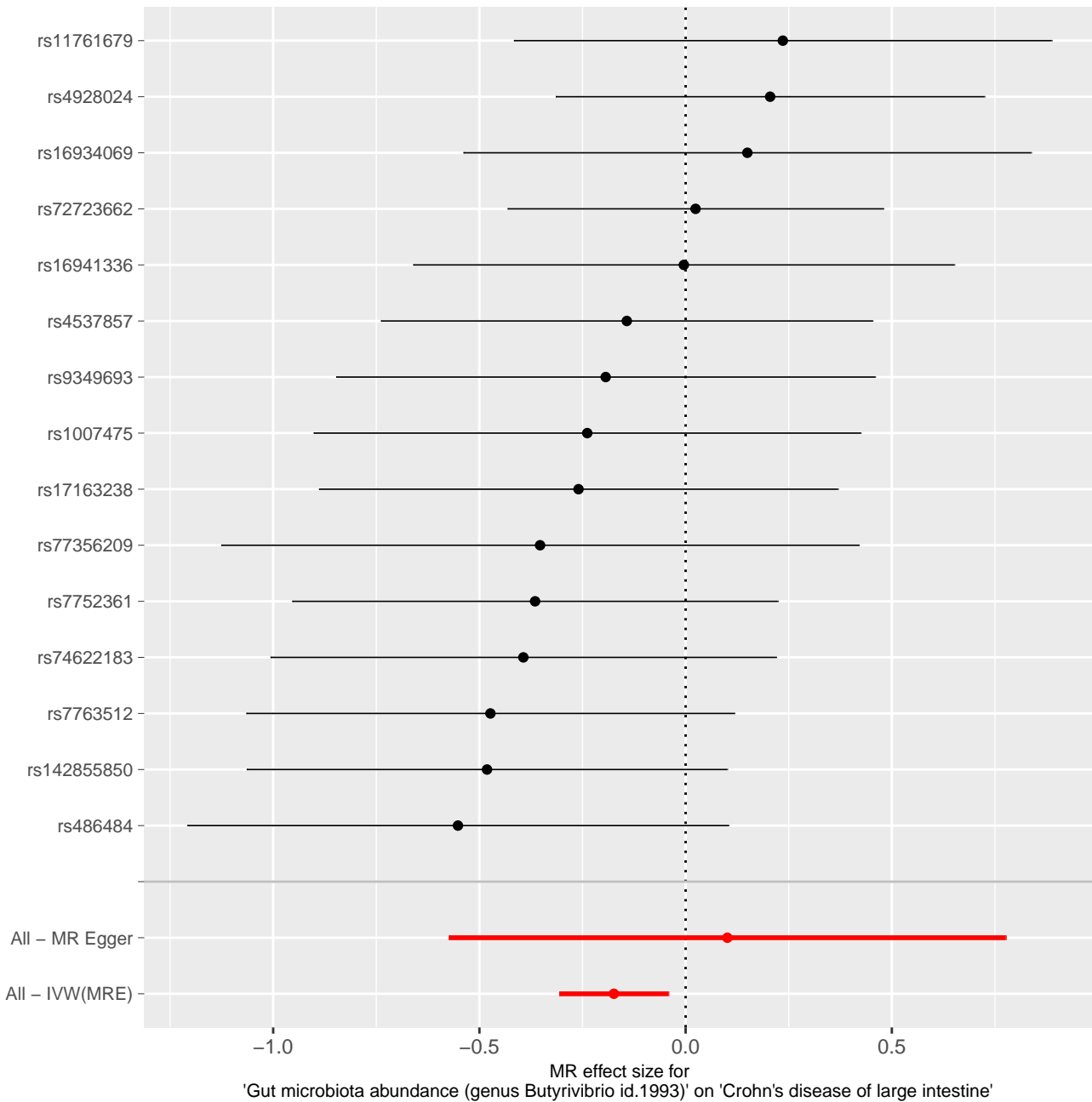
Batch 47 : Gut microbiota abundance (genus Butyricicoccus id.2055) on Crohn's disease of large intestine

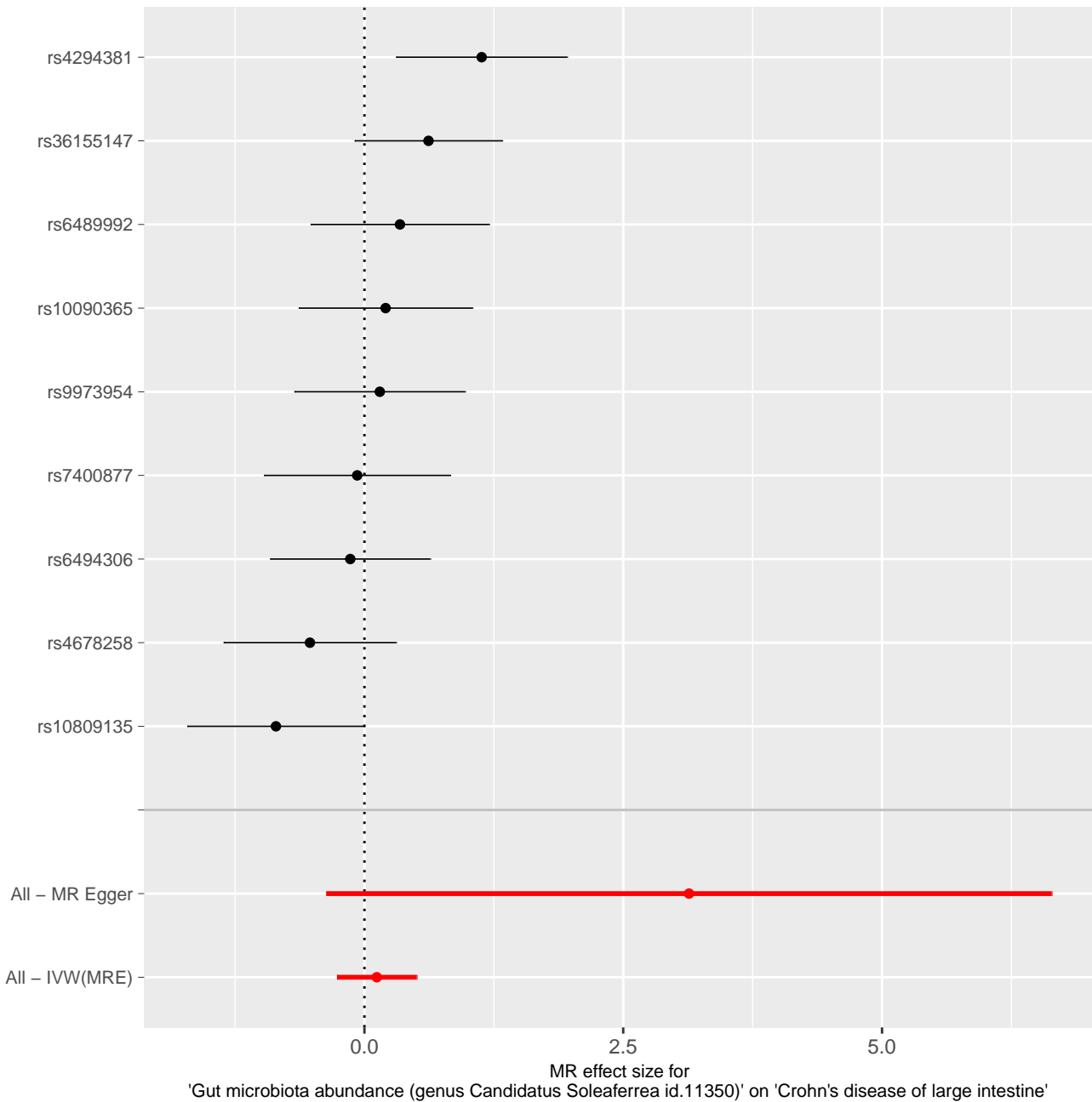


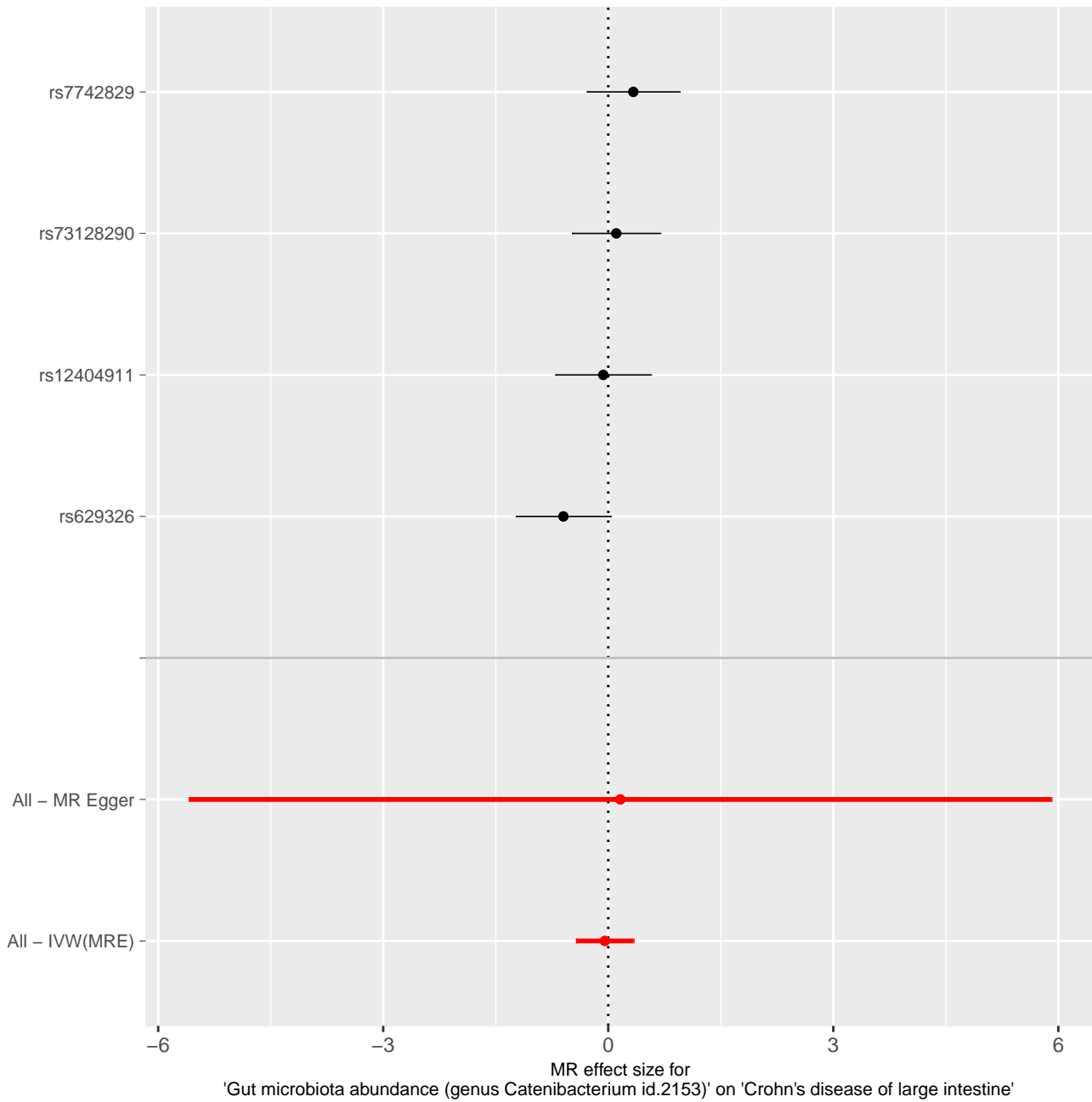
Batch 48 : Gut microbiota abundance (genus *Butyricimonas* id.945) on Crohn's disease of large intestine

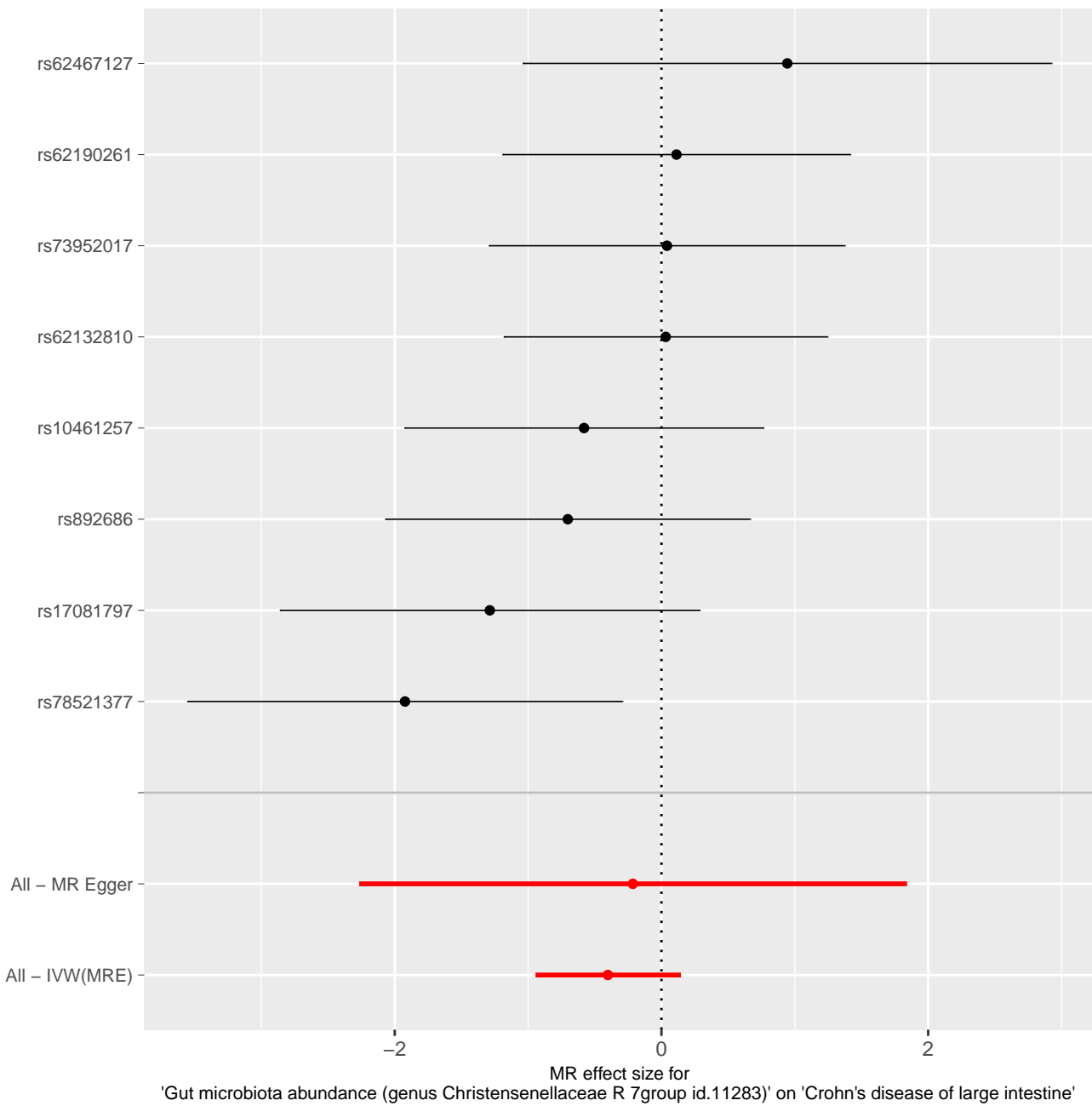


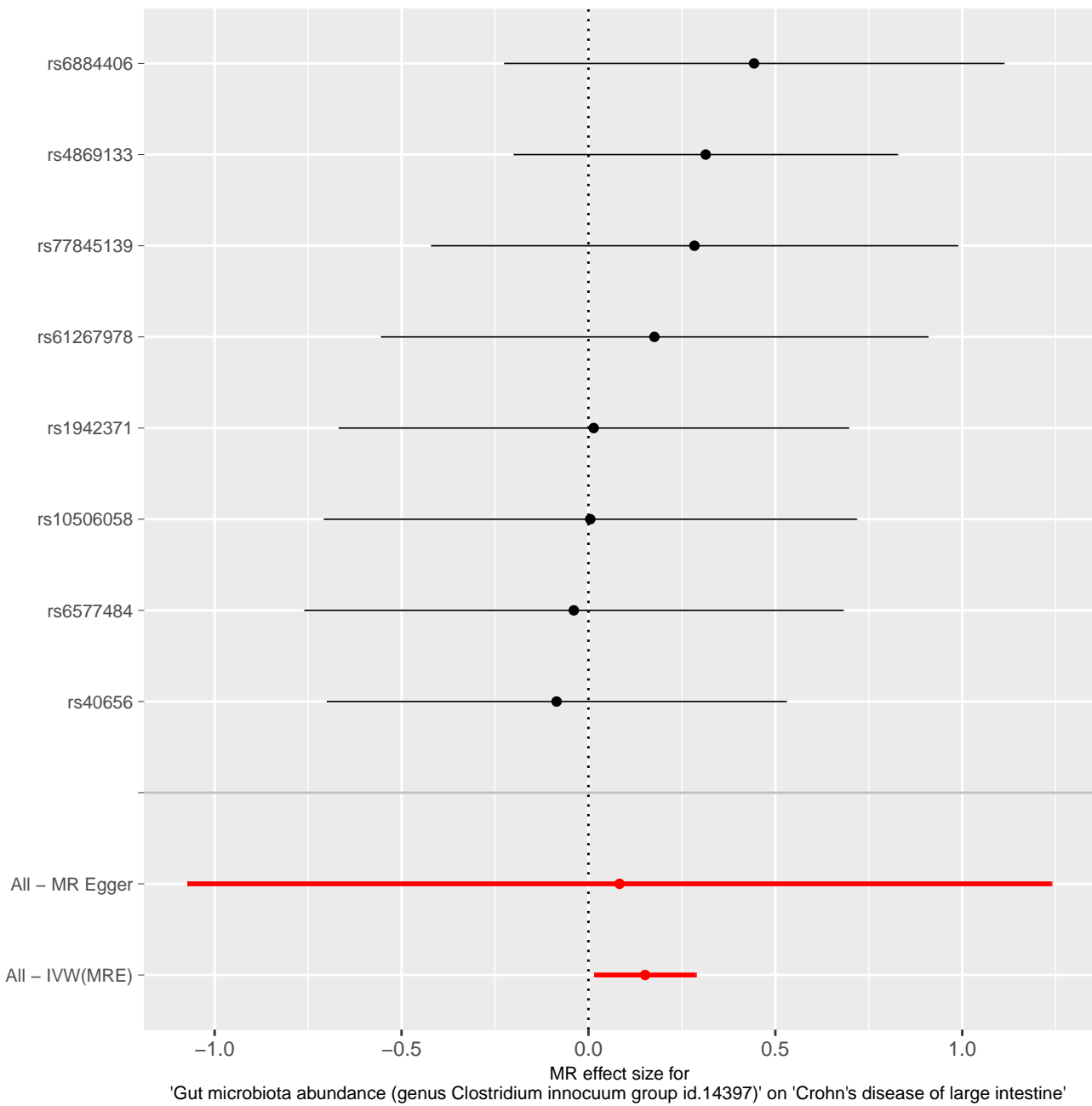
Batch 49 : Gut microbiota abundance (genus Butyrivibrio id.1993) on Crohn's disease of large intestine

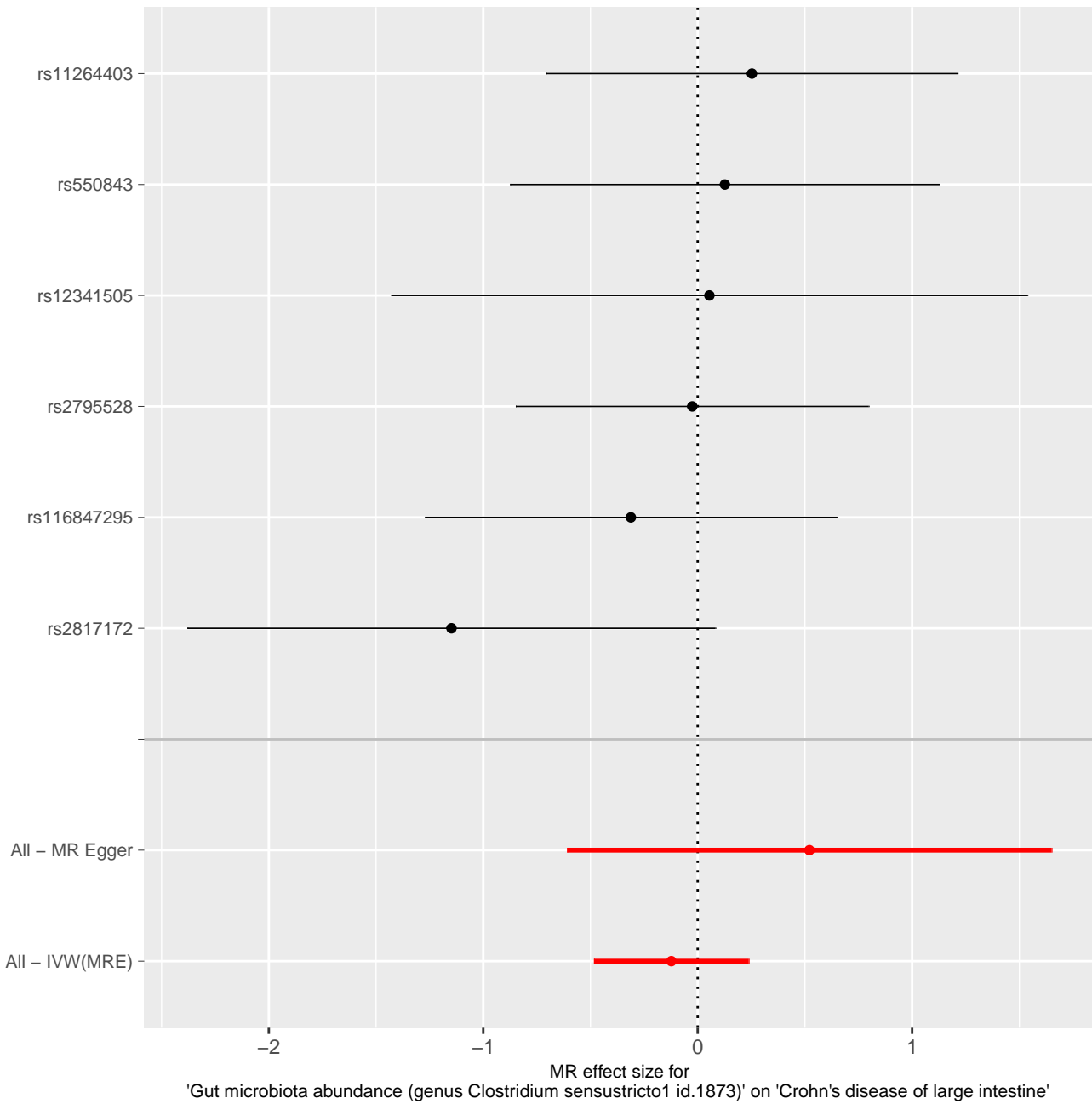


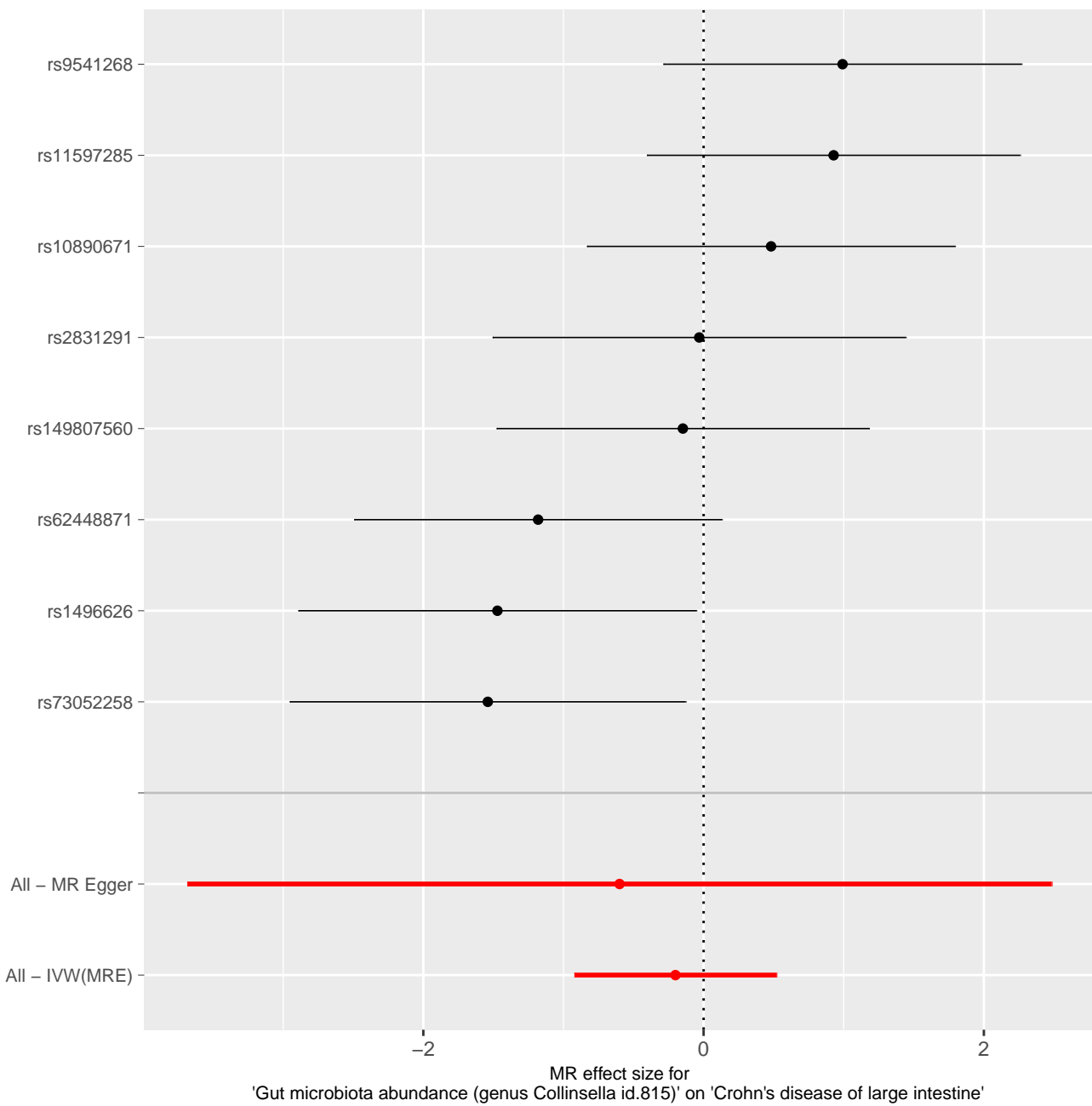




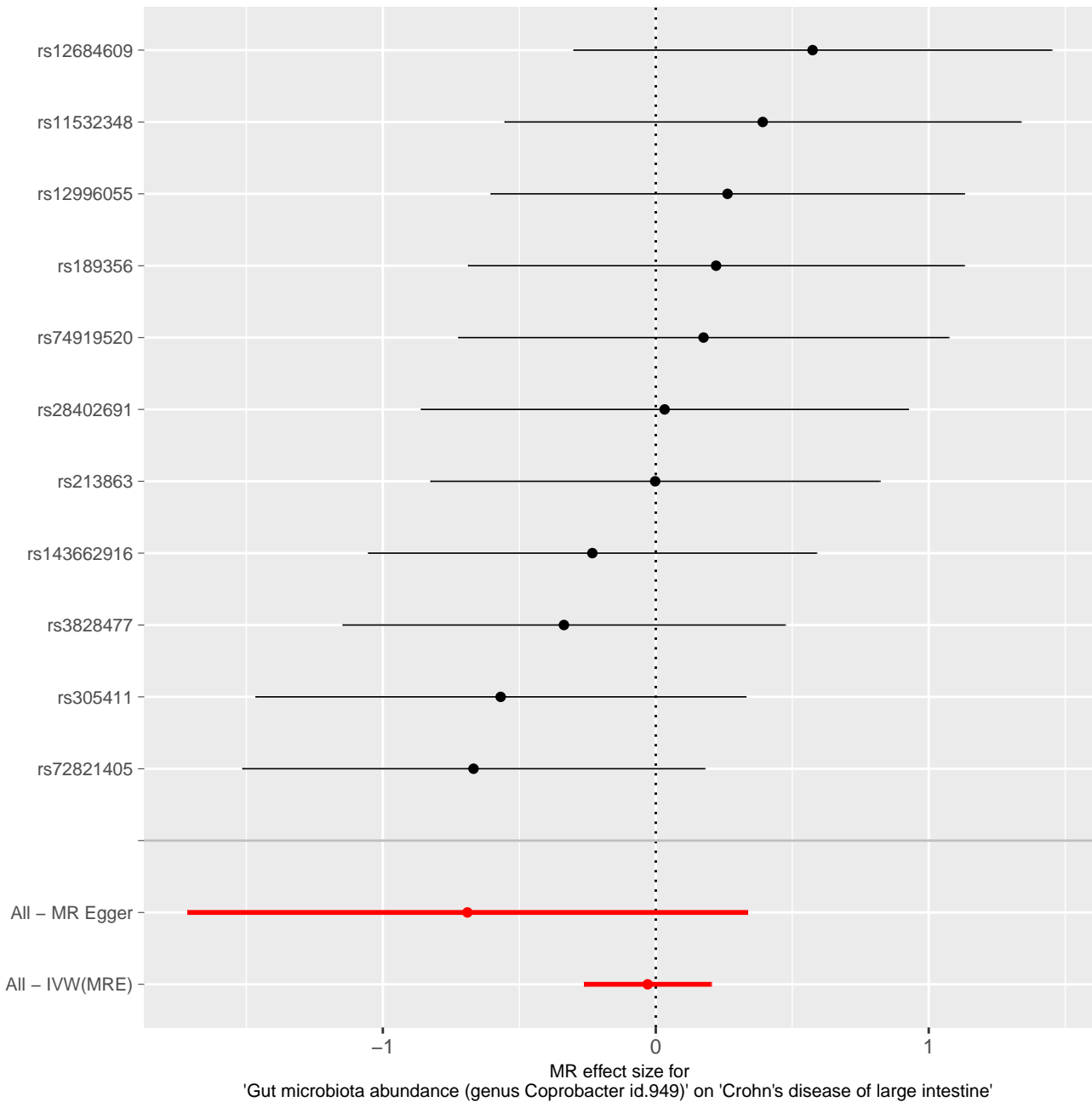




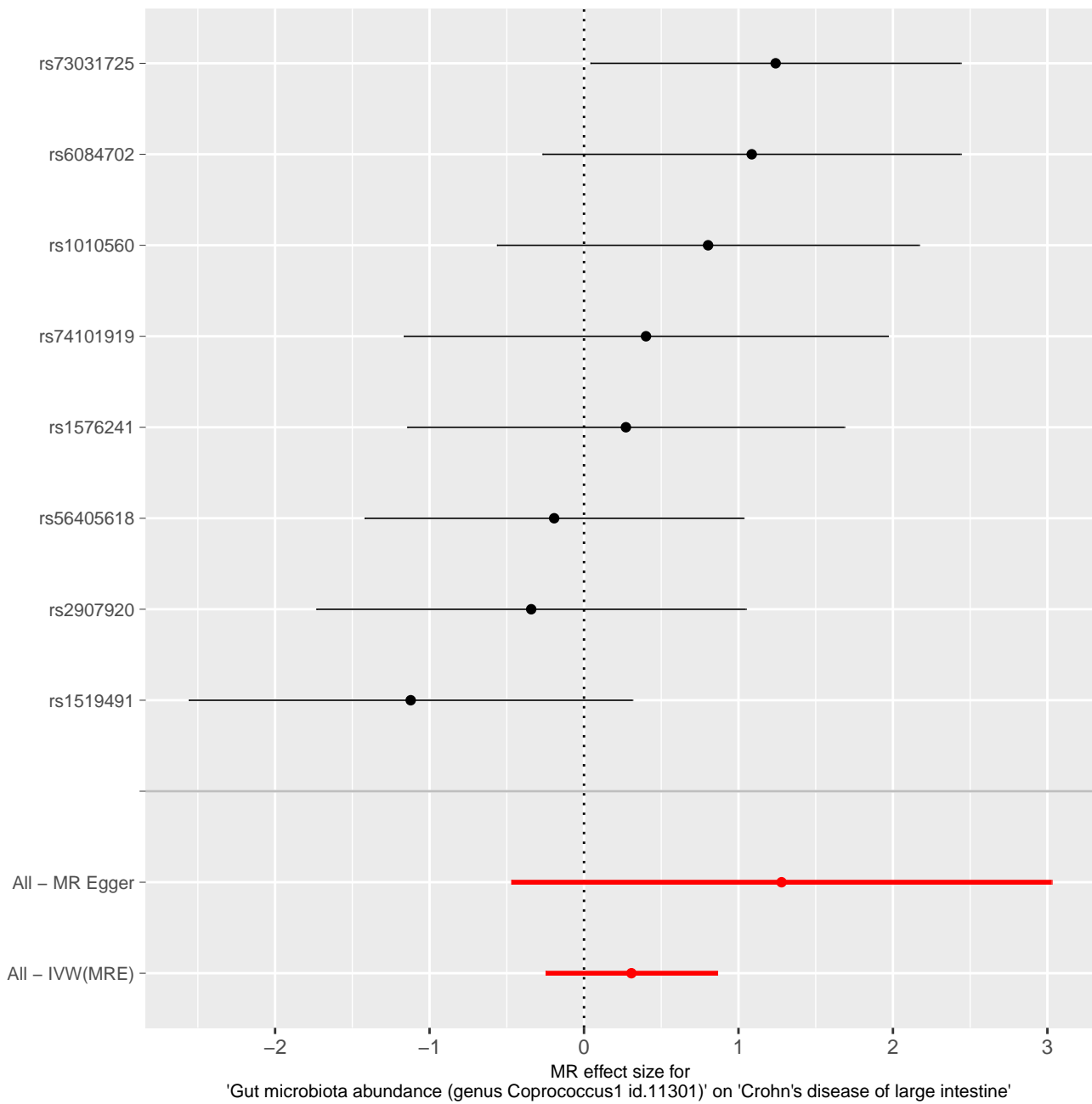


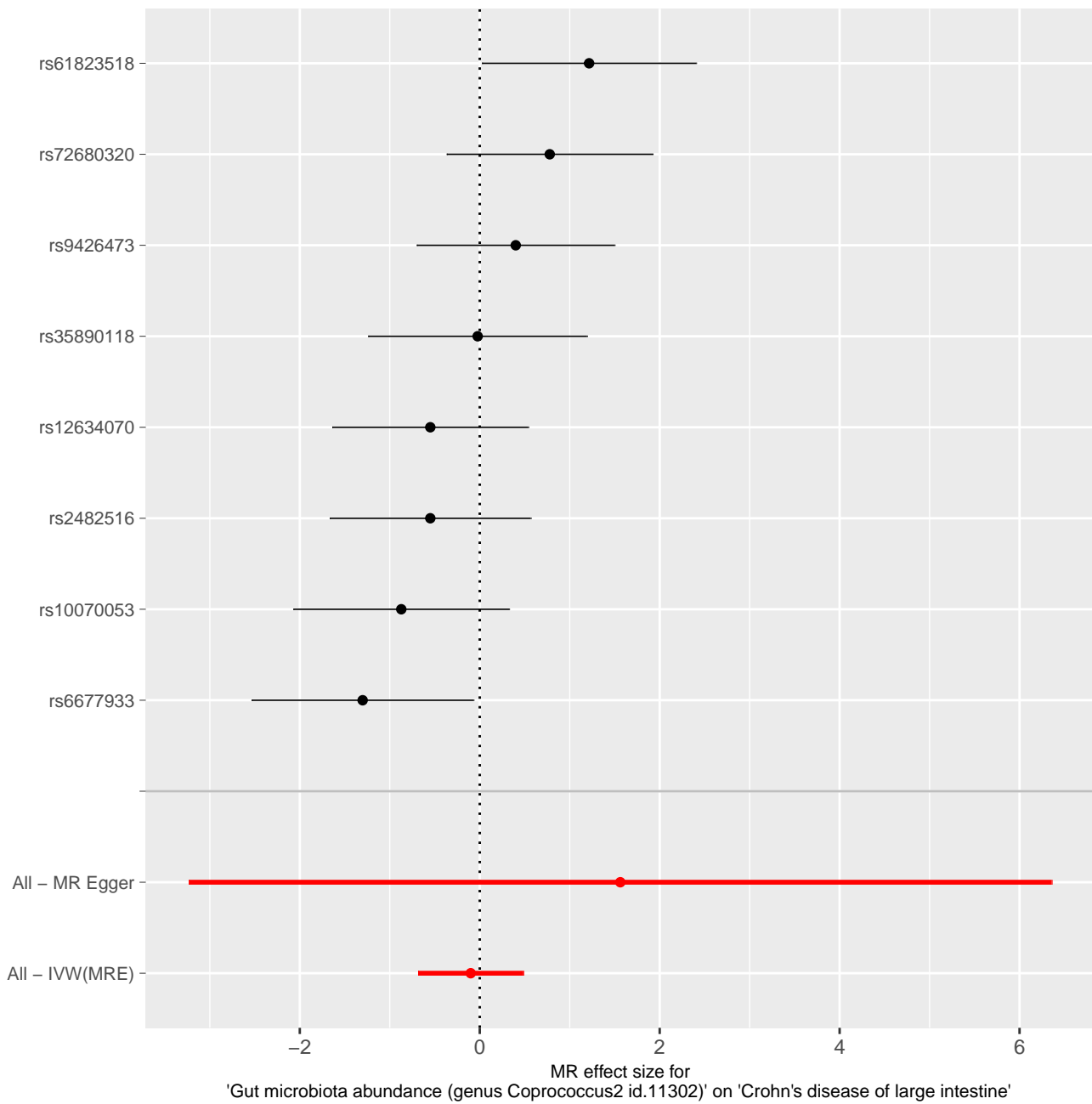


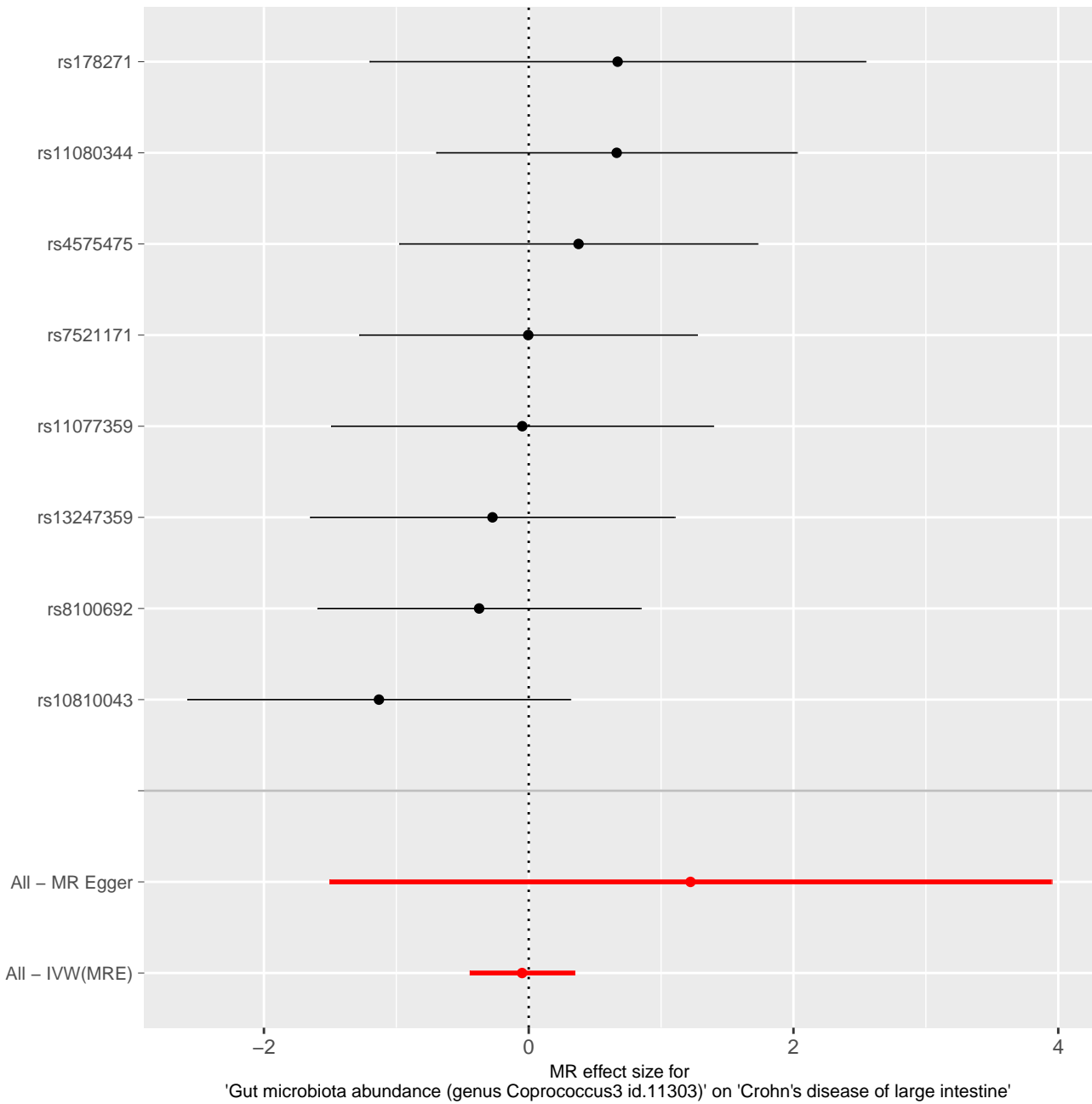
Batch 56 : Gut microbiota abundance (genus Coprobacter id.949) on Crohn's disease of large intestine



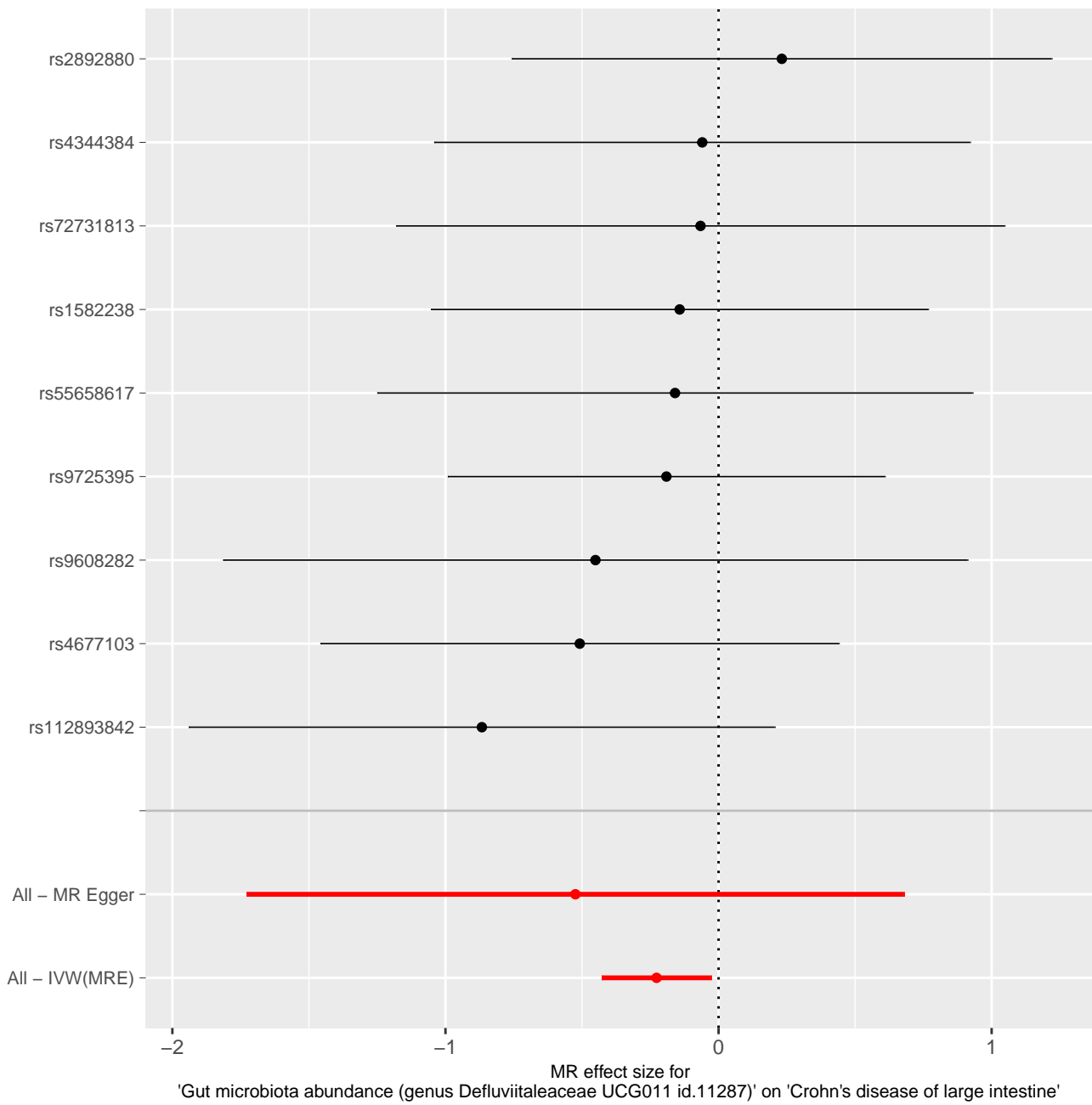
Batch 57 : Gut microbiota abundance (genus Coprococcus1 id.11301) on Crohn's disease of large intestine



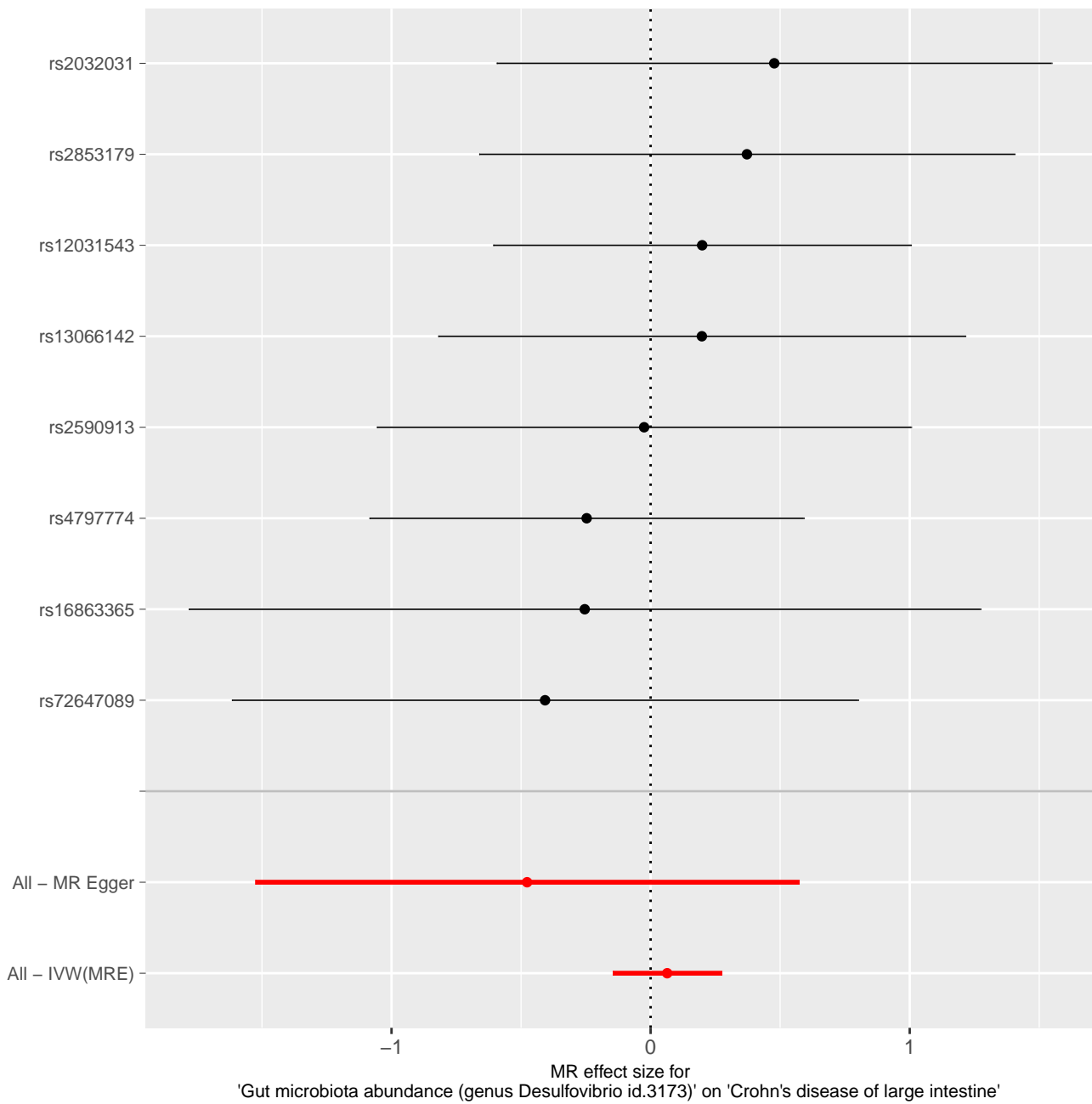




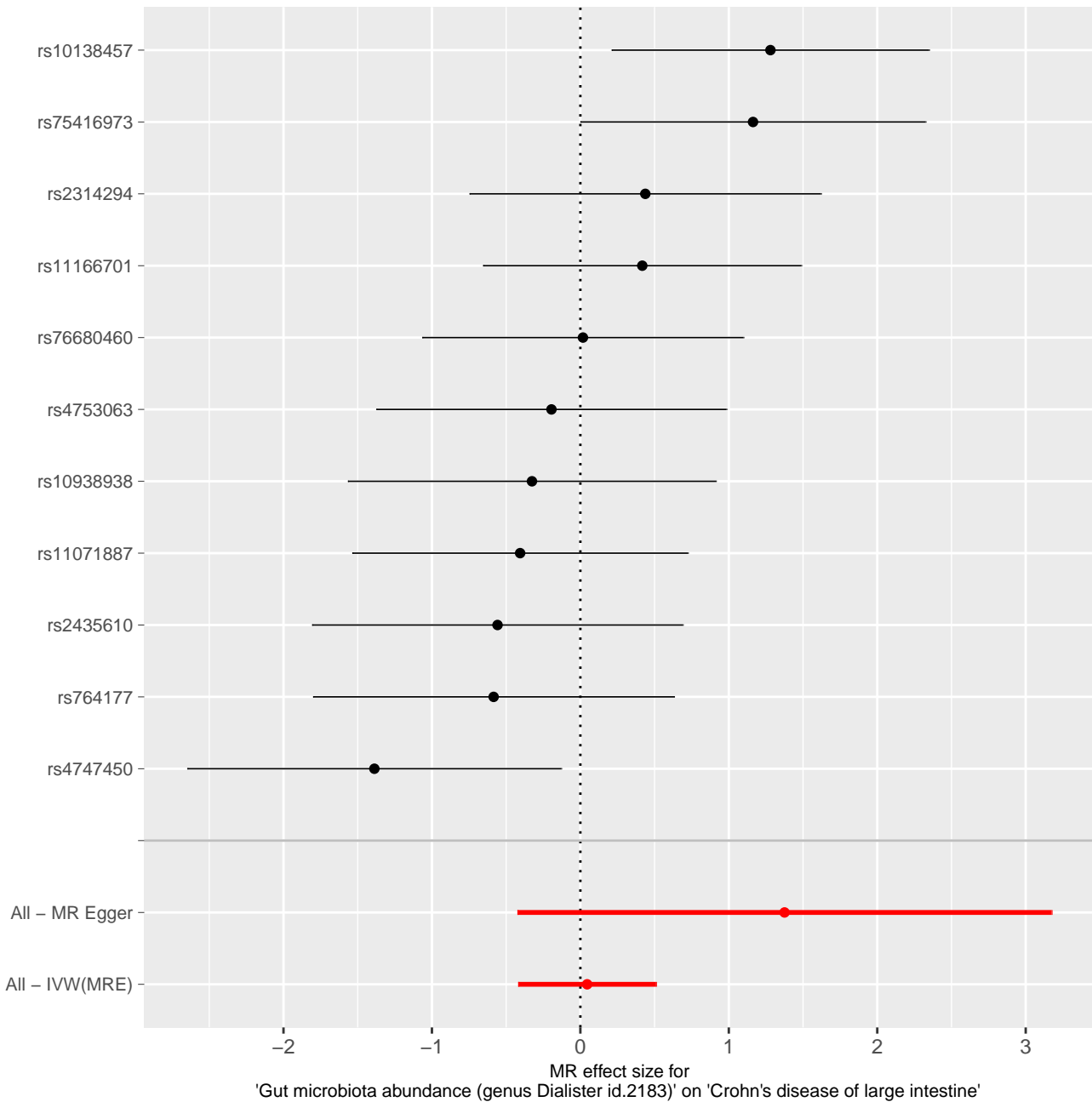
Batch 60 : Gut microbiota abundance (genus Defluviitaleaceae UCG011 id.11287) on Crohn's disease of large intestine



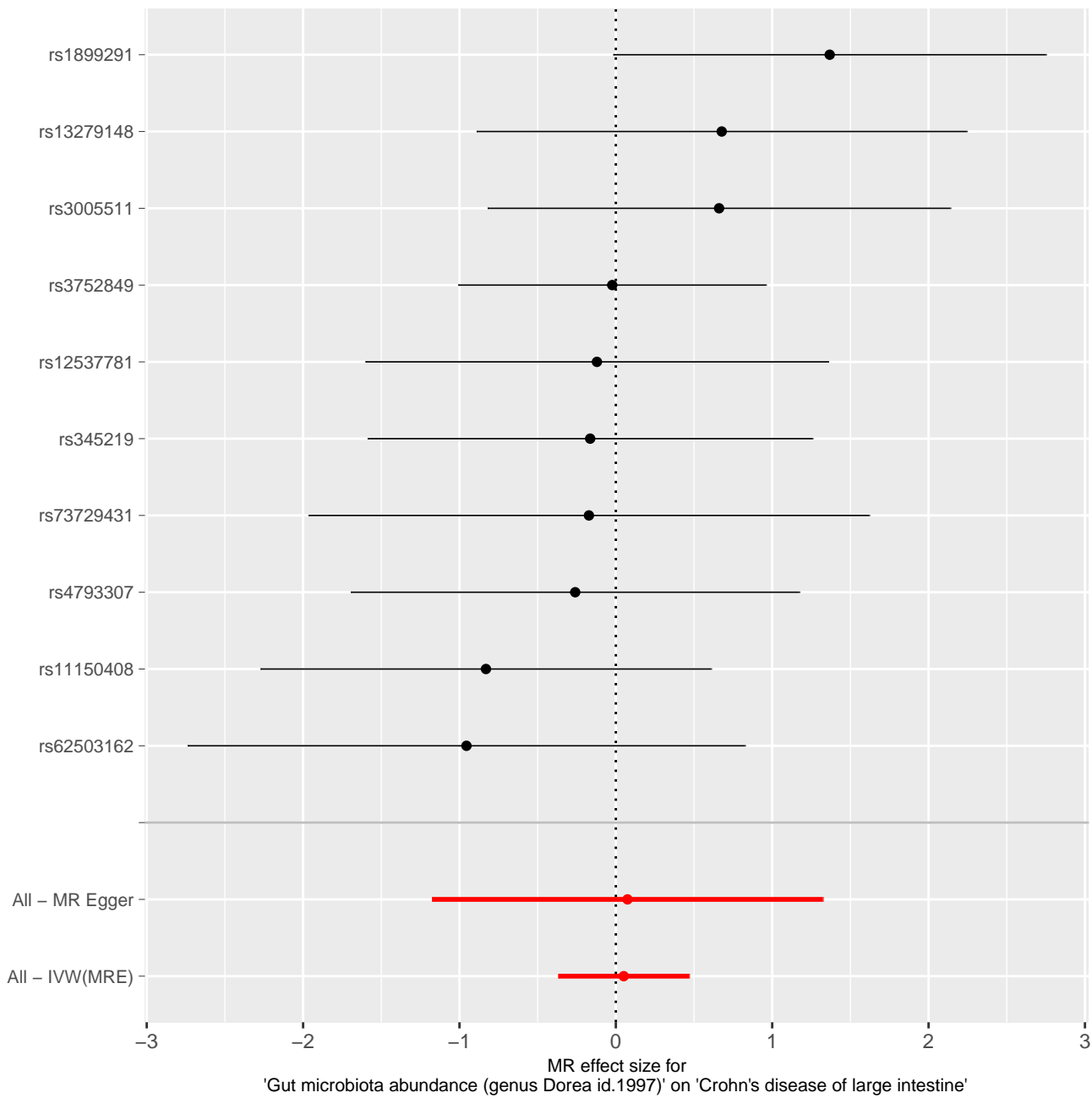
Batch 61 : Gut microbiota abundance (genus Desulfovibrio id.3173) on Crohn's disease of large intestine



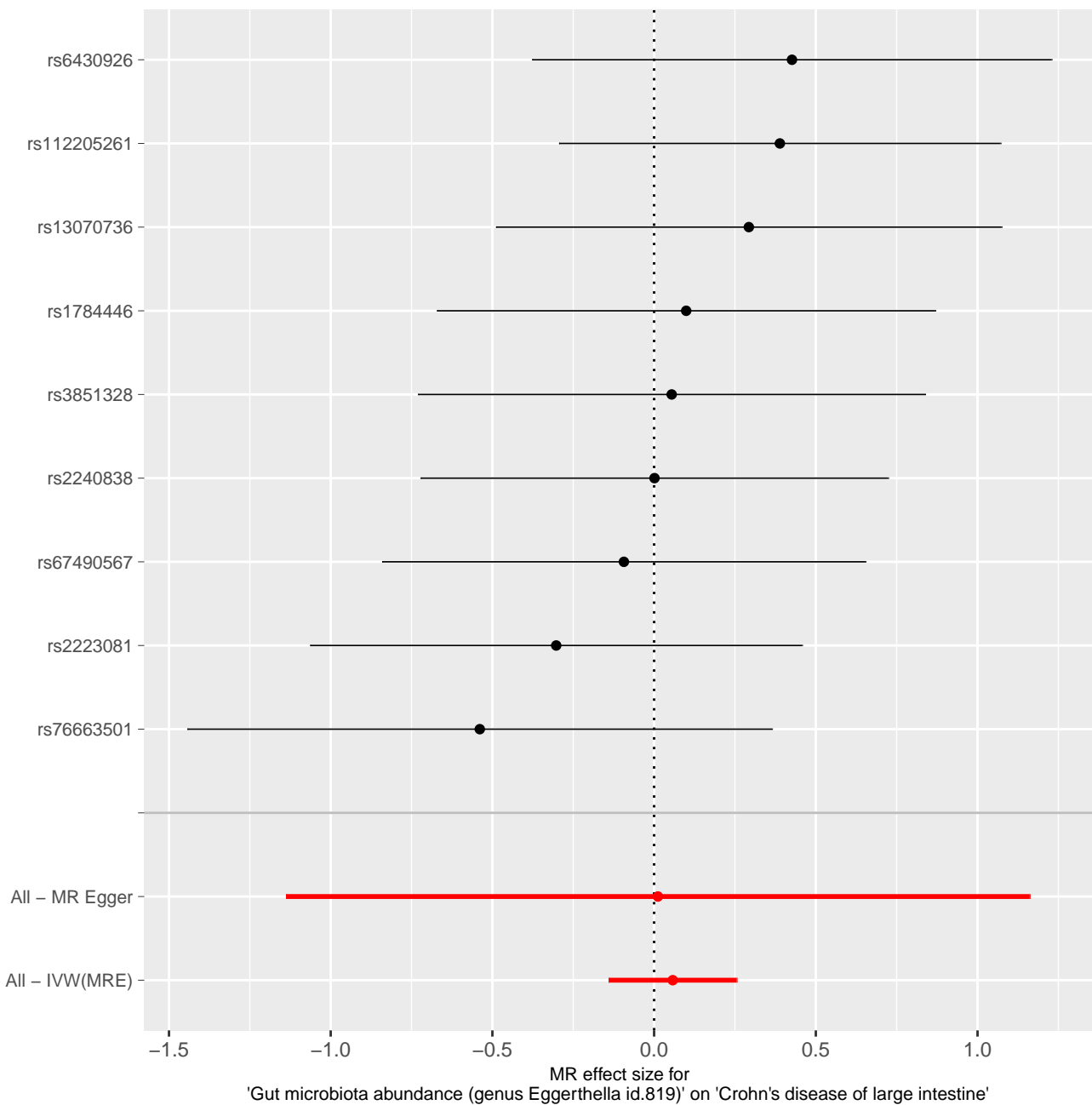
Batch 62 : Gut microbiota abundance (genus Dialister id.2183) on Crohn's disease of large intestine



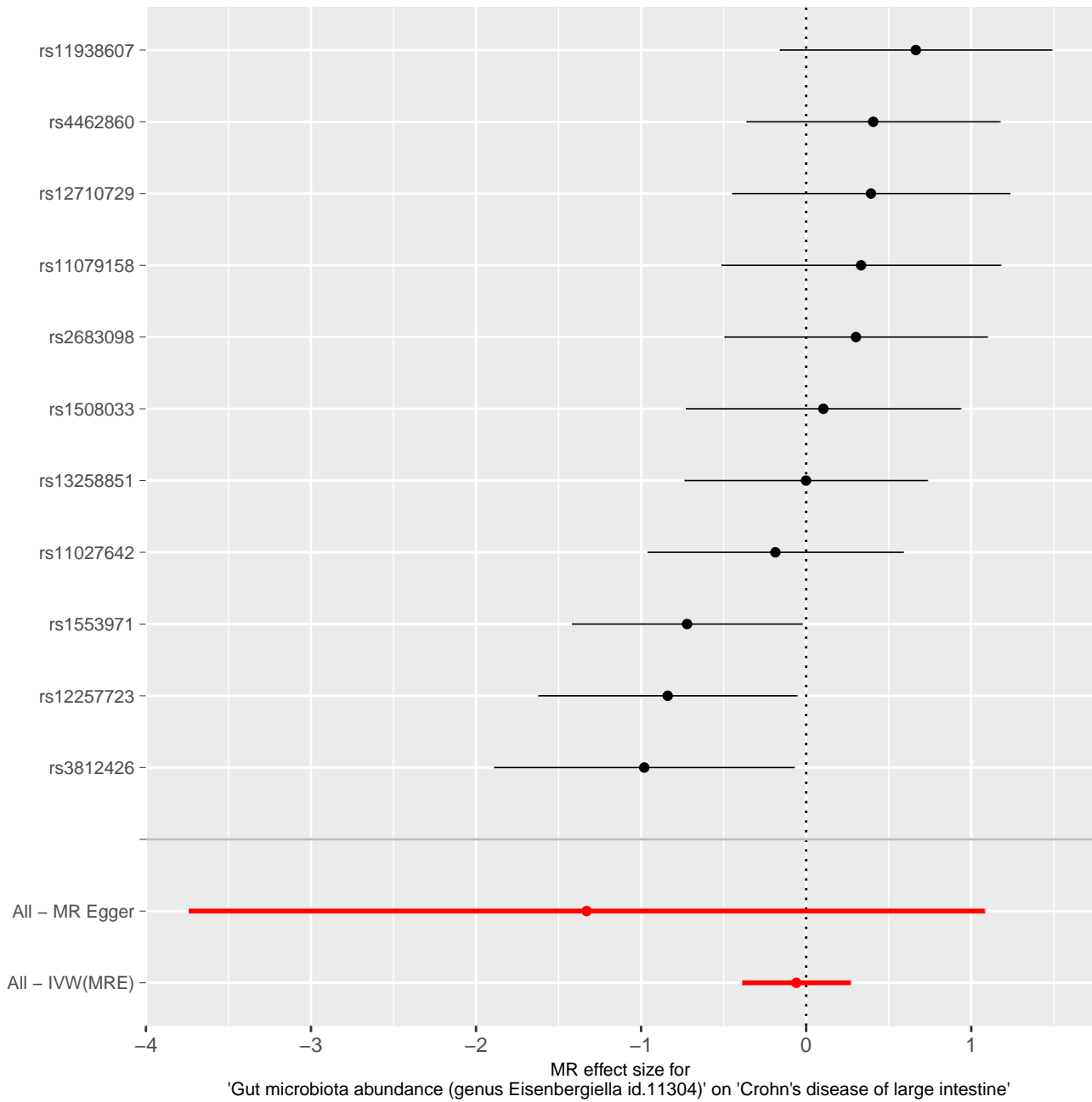
Batch 63 : Gut microbiota abundance (genus Dorea id.1997) on Crohn's disease of large intestine



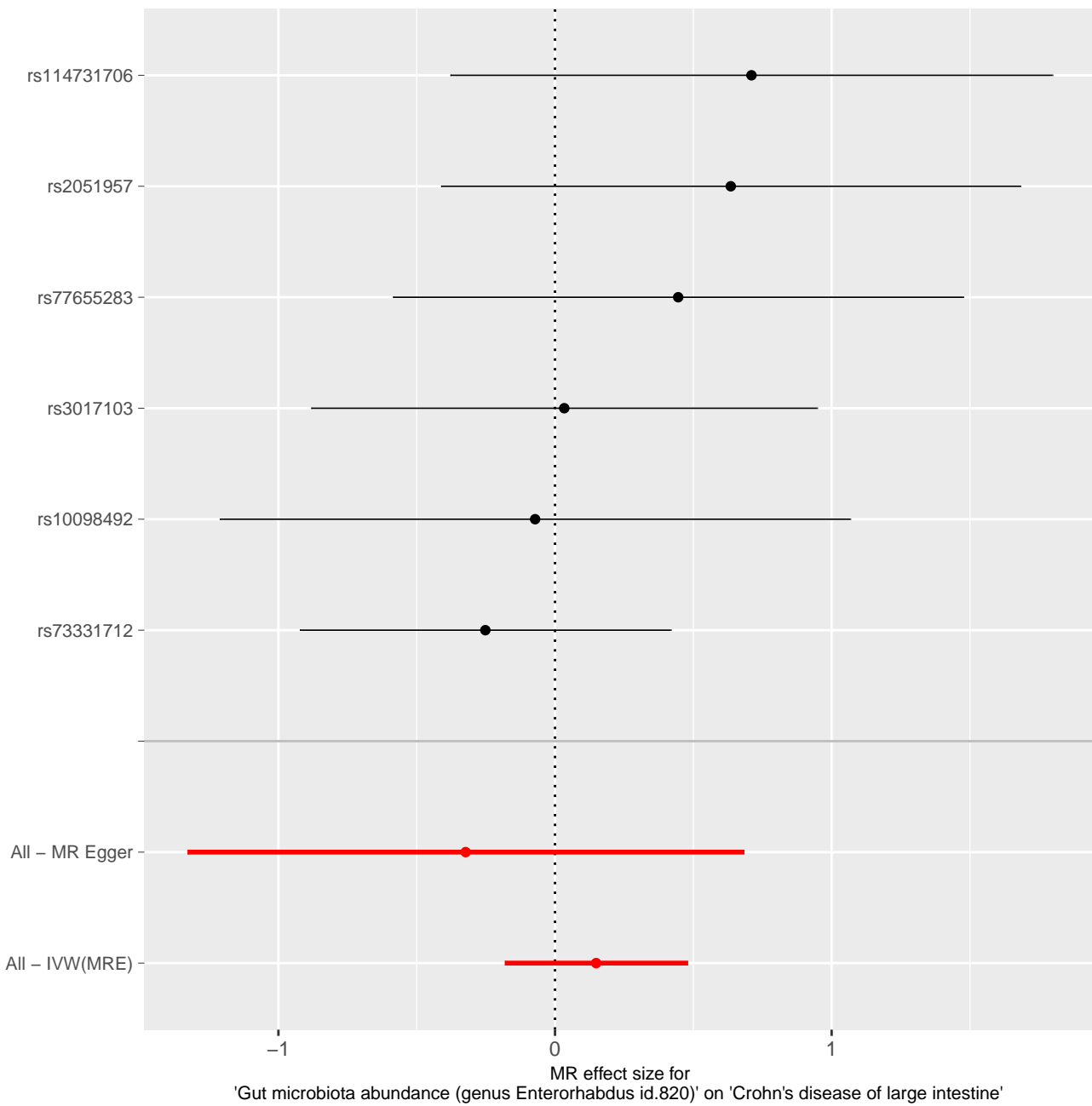
Batch 64 : Gut microbiota abundance (genus Eggerthella id.819) on Crohn's disease of large intestine

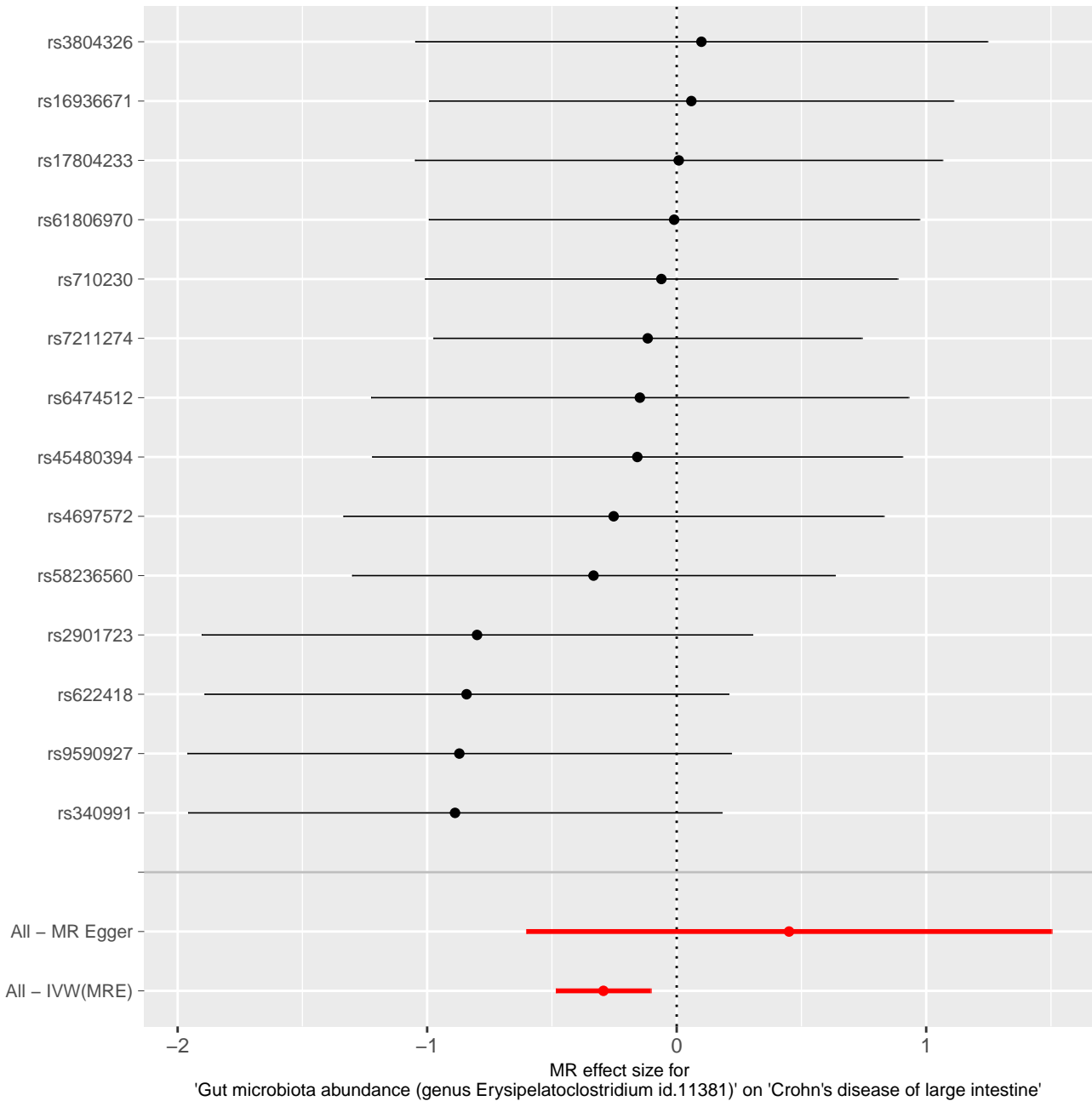


Batch 65 : Gut microbiota abundance (genus Eisenbergiella id.11304) on Crohn's disease of large intestine

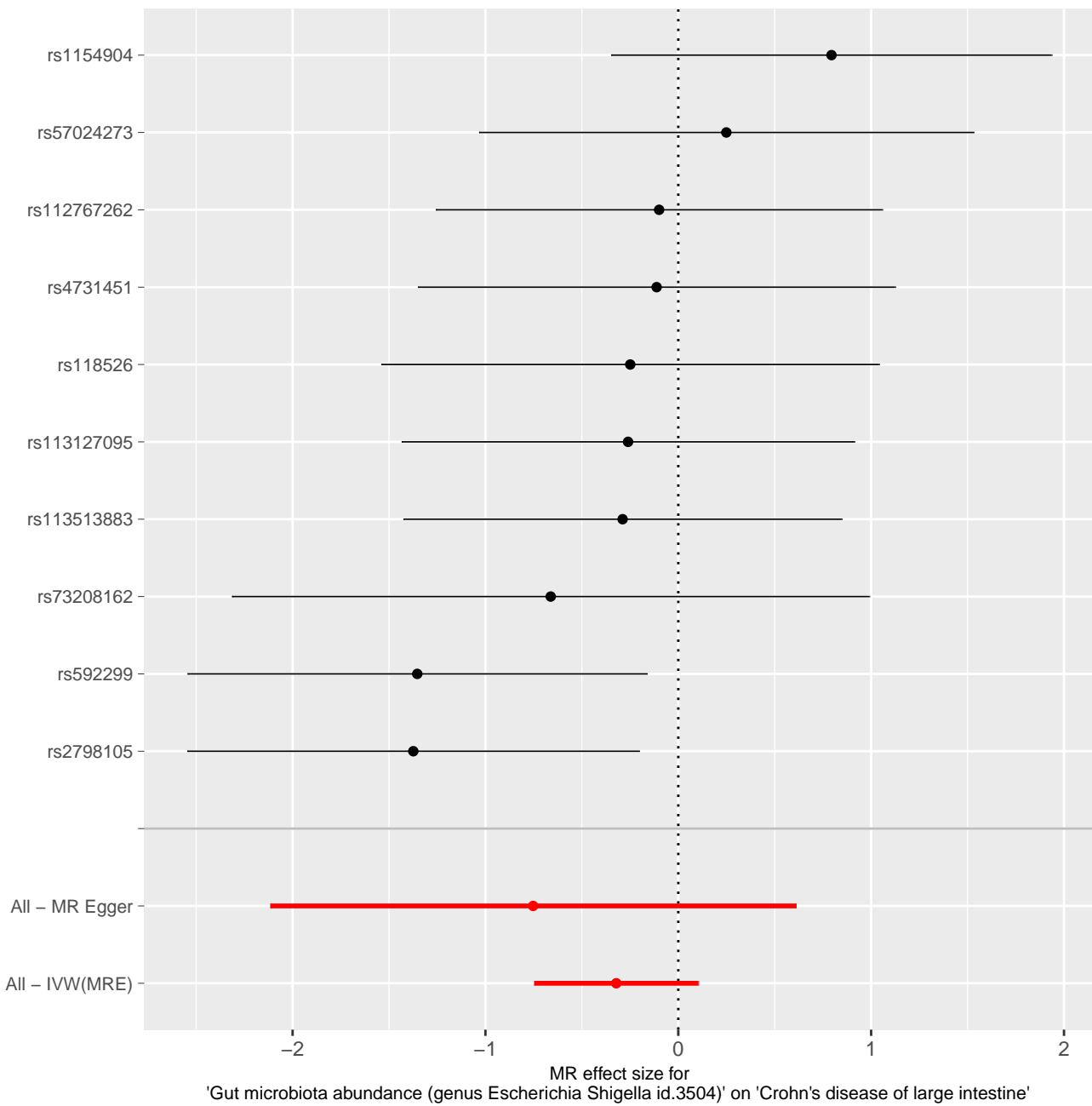


Batch 66 : Gut microbiota abundance (genus Enterorhabdus id.820) on Crohn's disease of large intestine

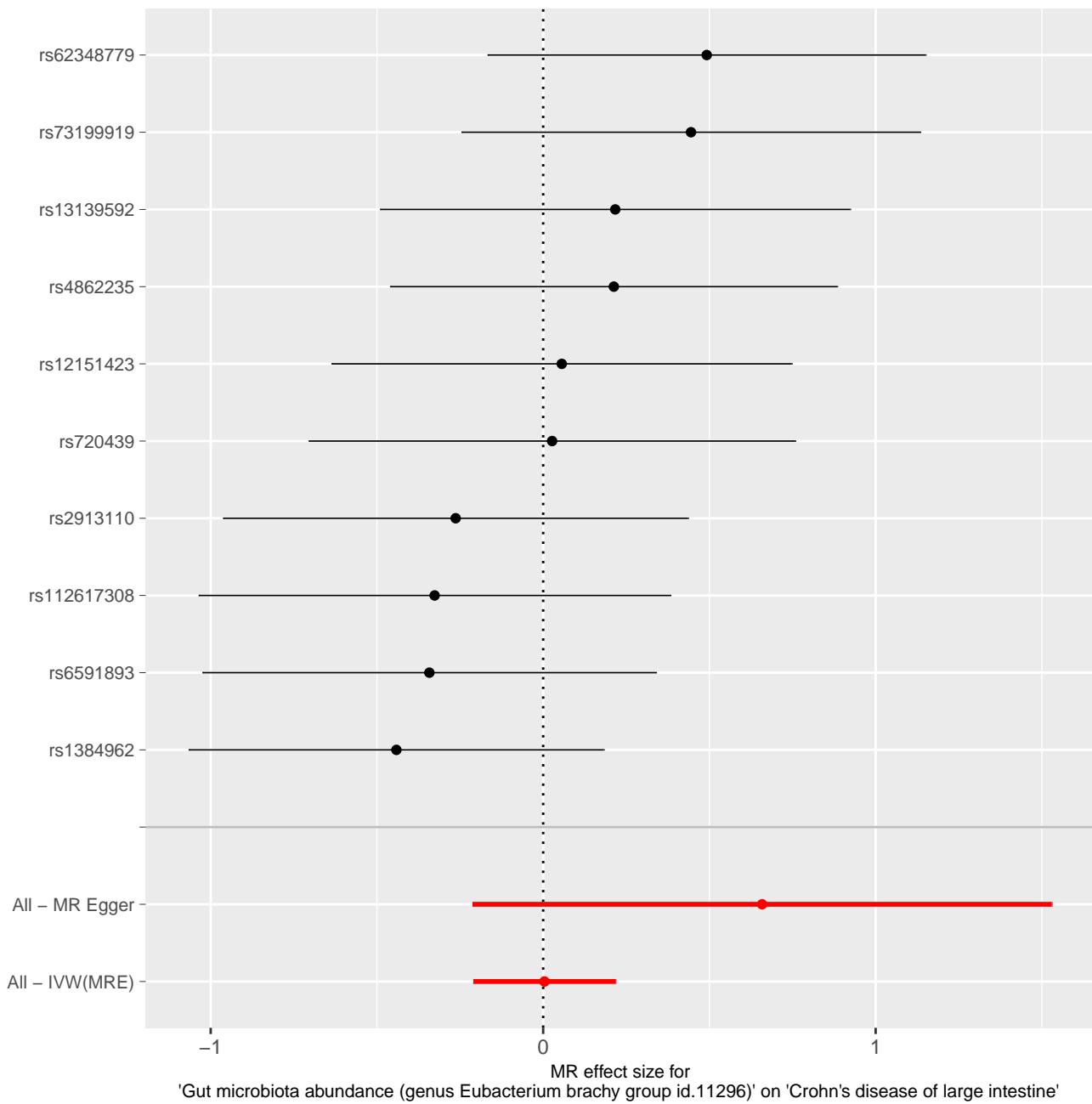




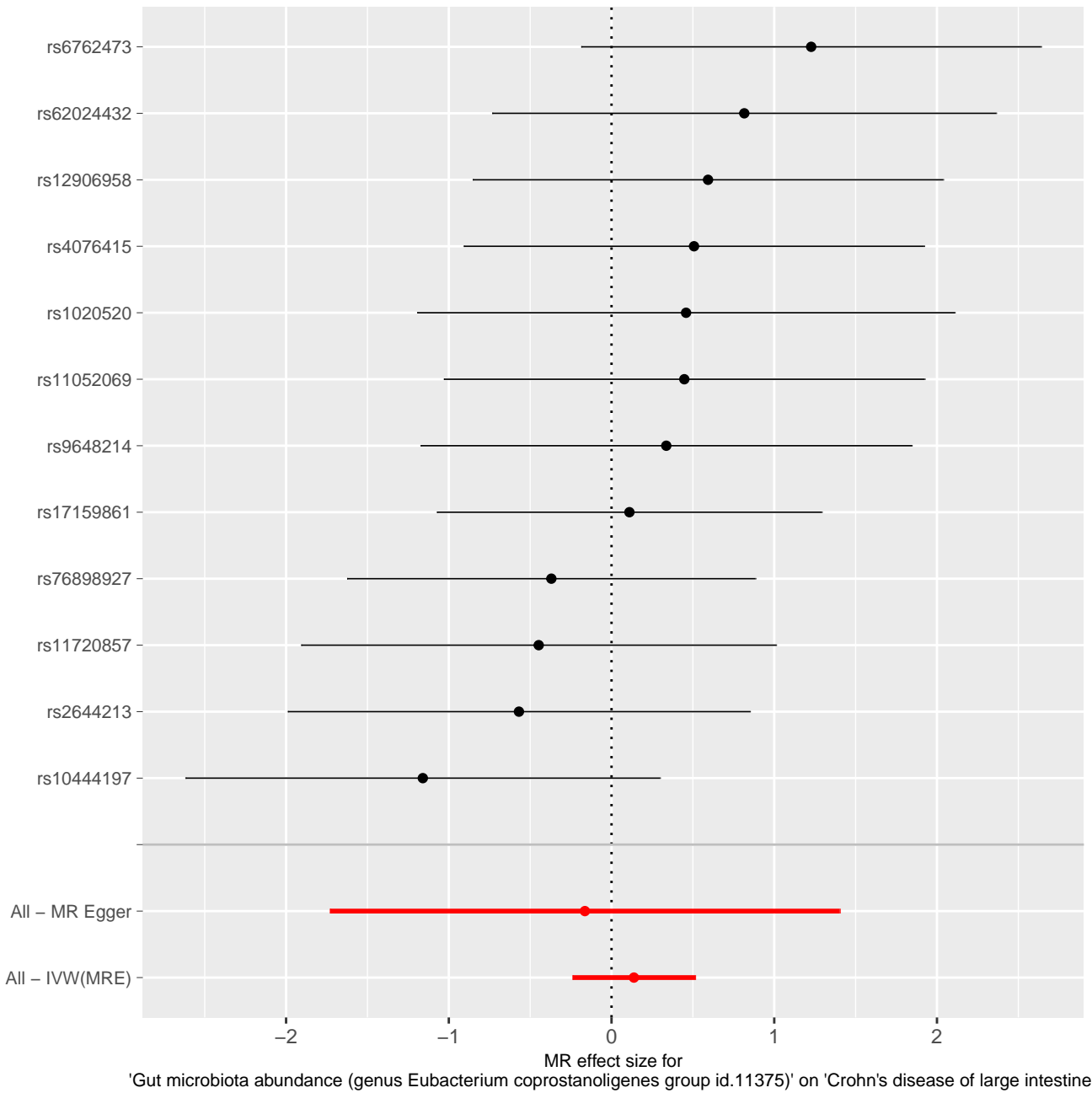
Batch 69 : Gut microbiota abundance (genus Escherichia Shigella id.3504) on Crohn's disease of large intestine

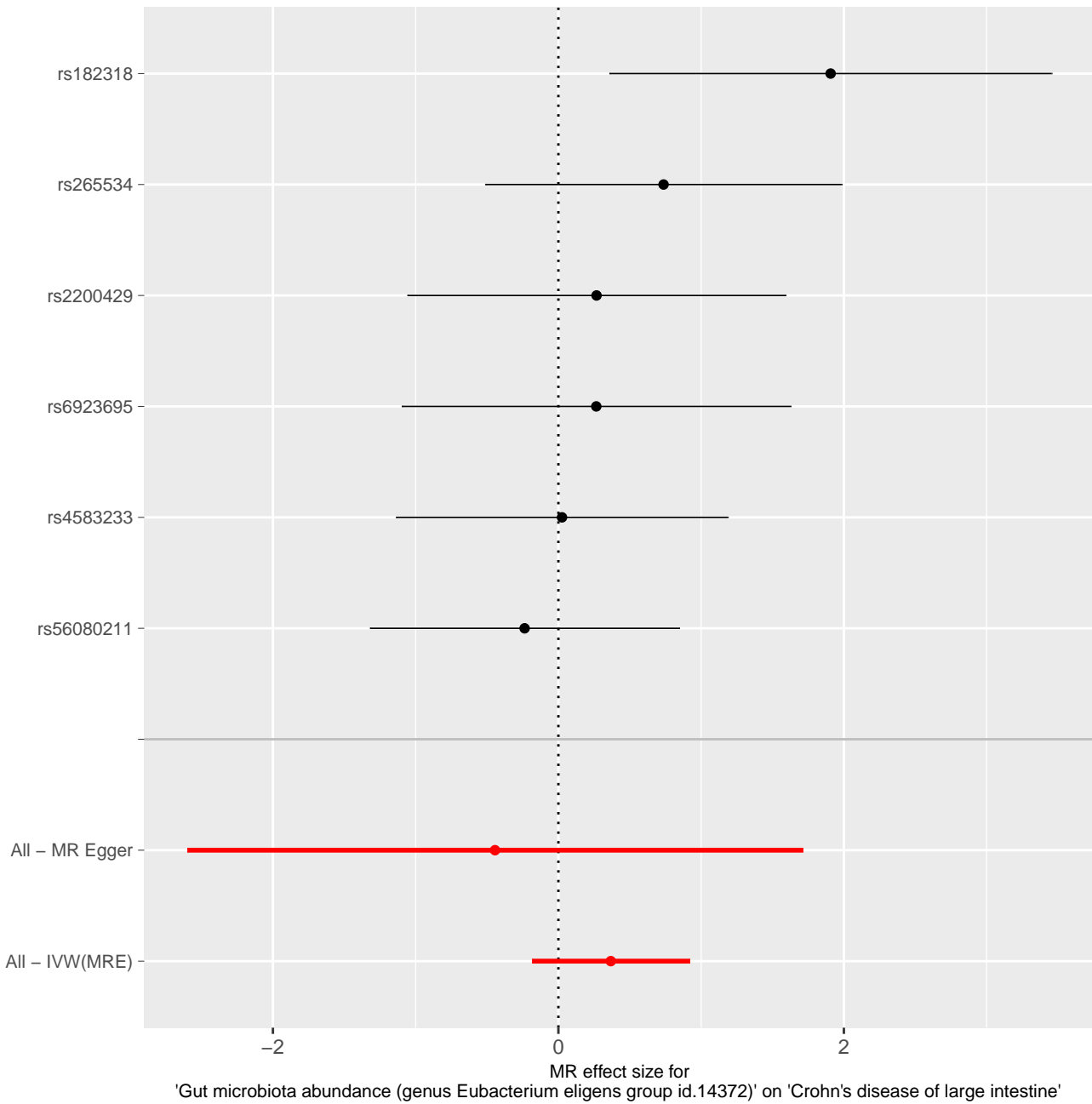


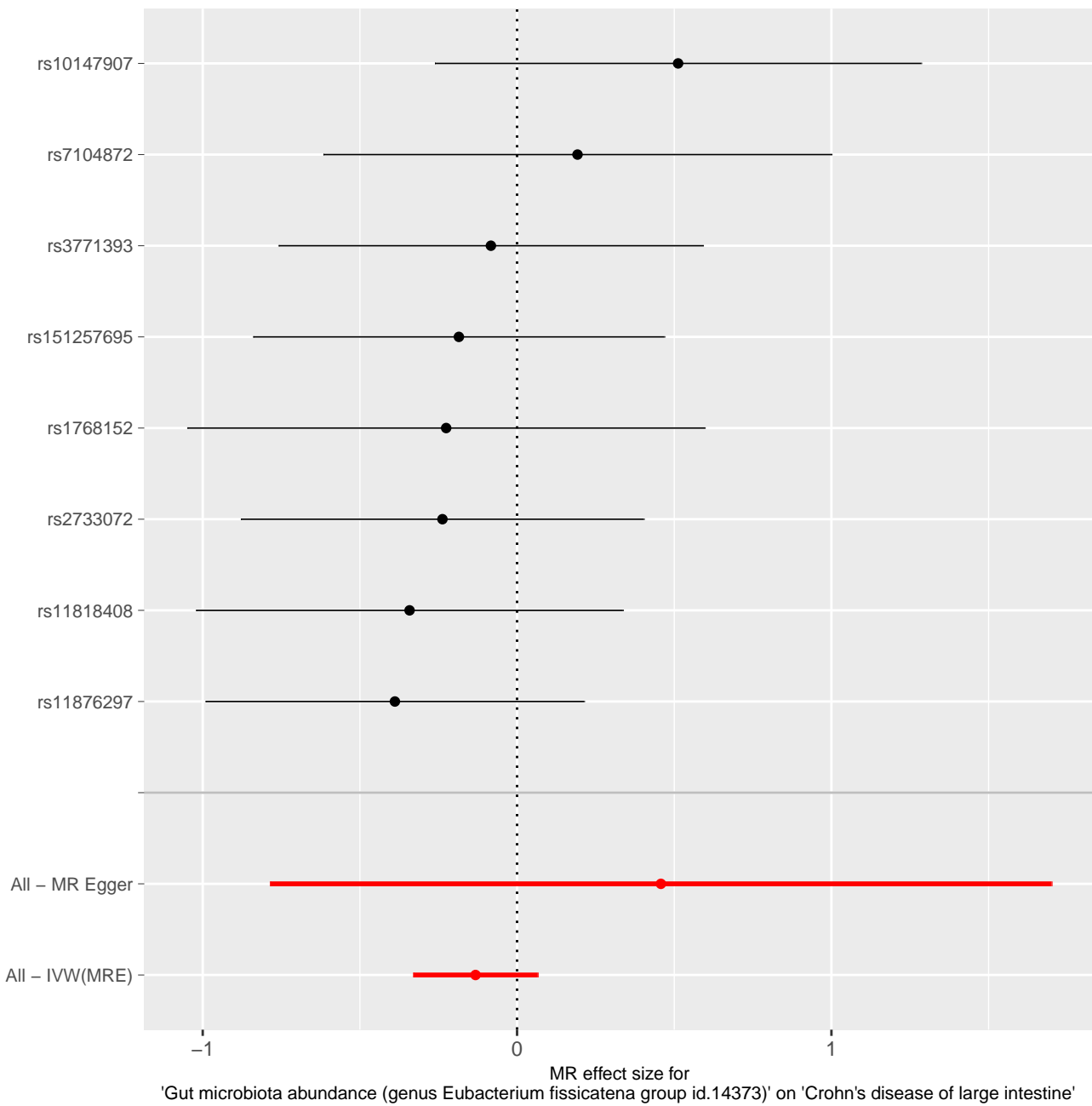
Batch 70 : Gut microbiota abundance (genus Eubacterium brachy group id.11296) on Crohn's disease of large intestine



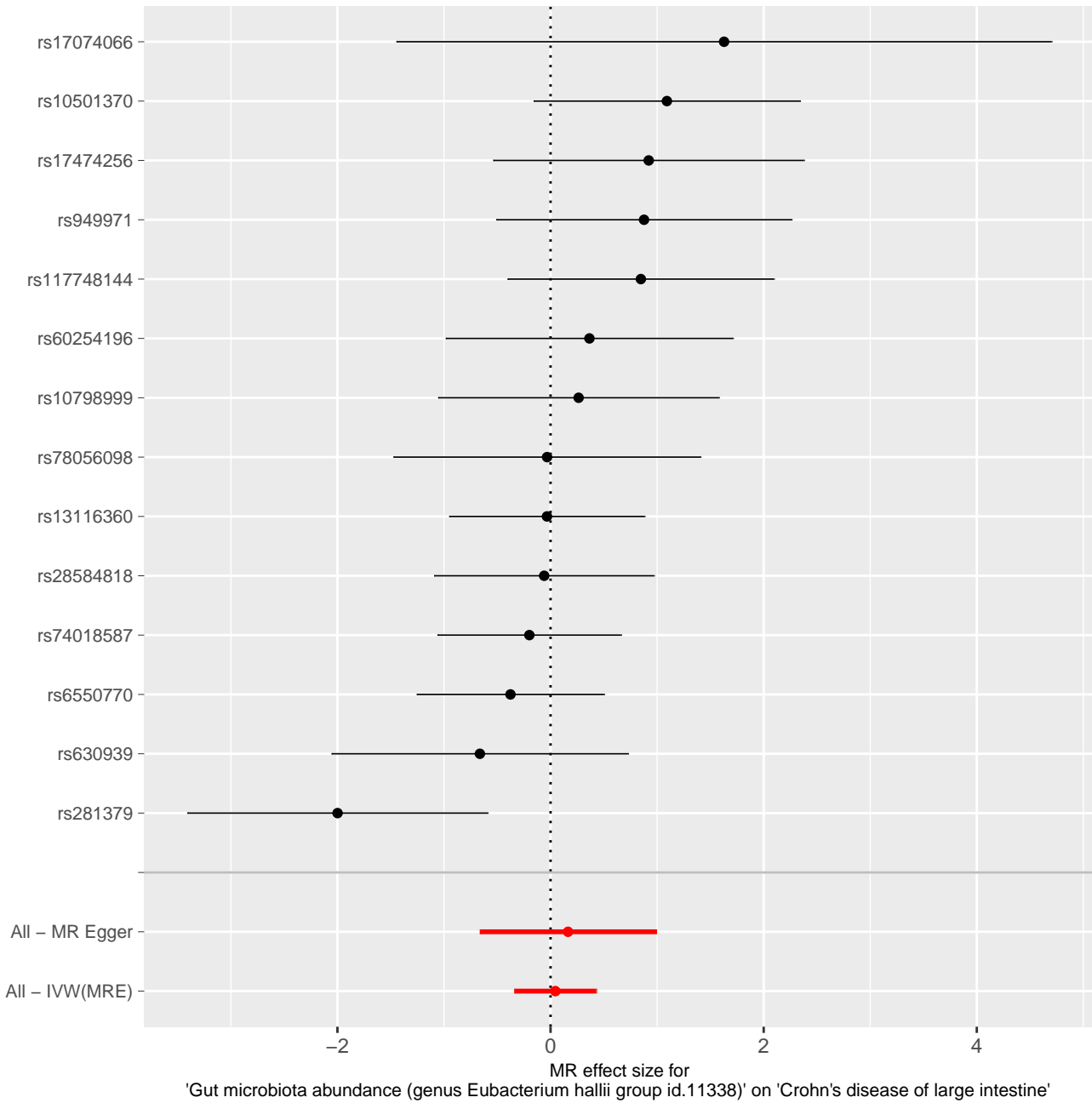
Batch 71 : Gut microbiota abundance (genus Eubacterium coprostanoligenes group id.11375) on Crohn's disease of large intestine



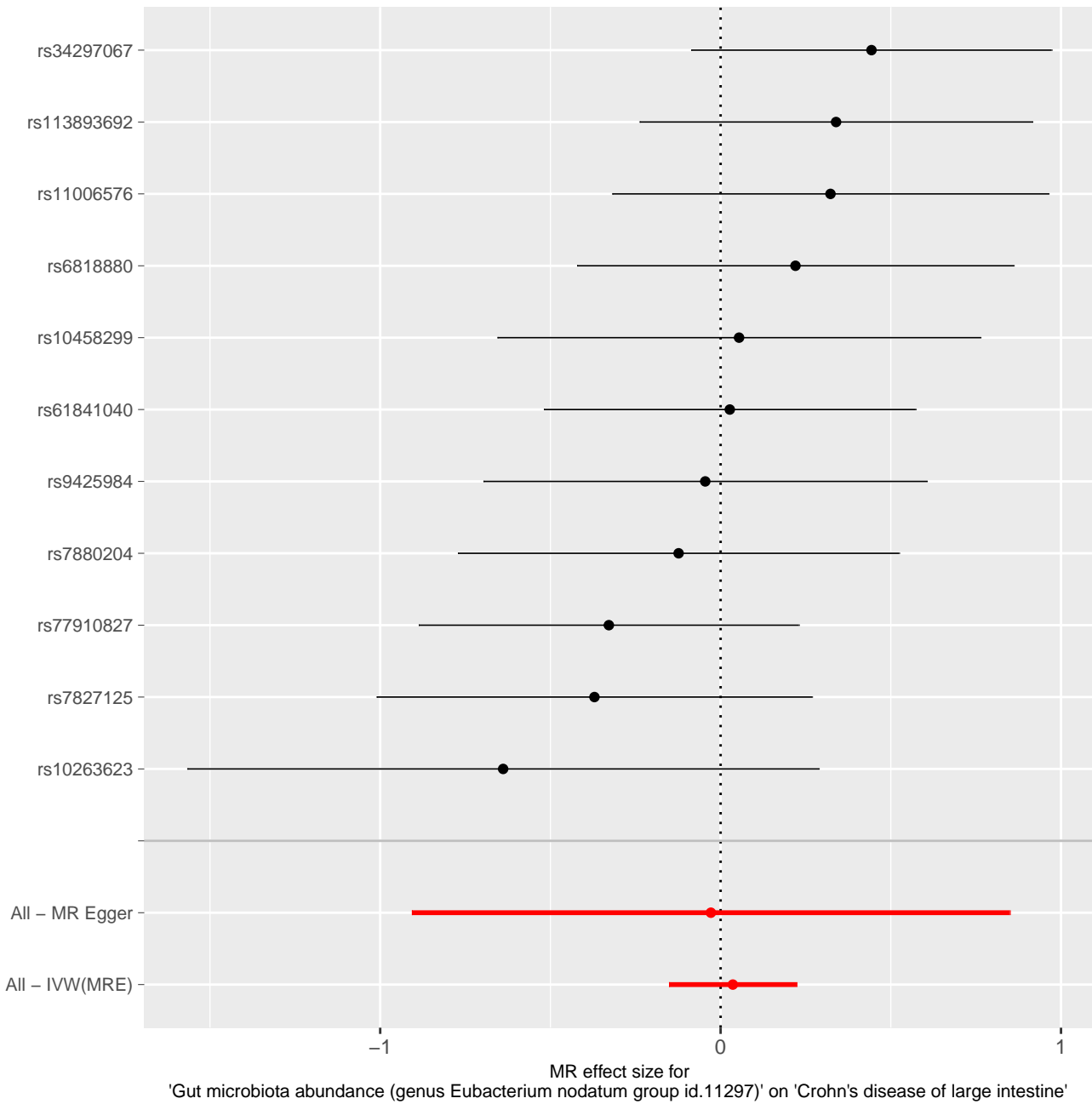


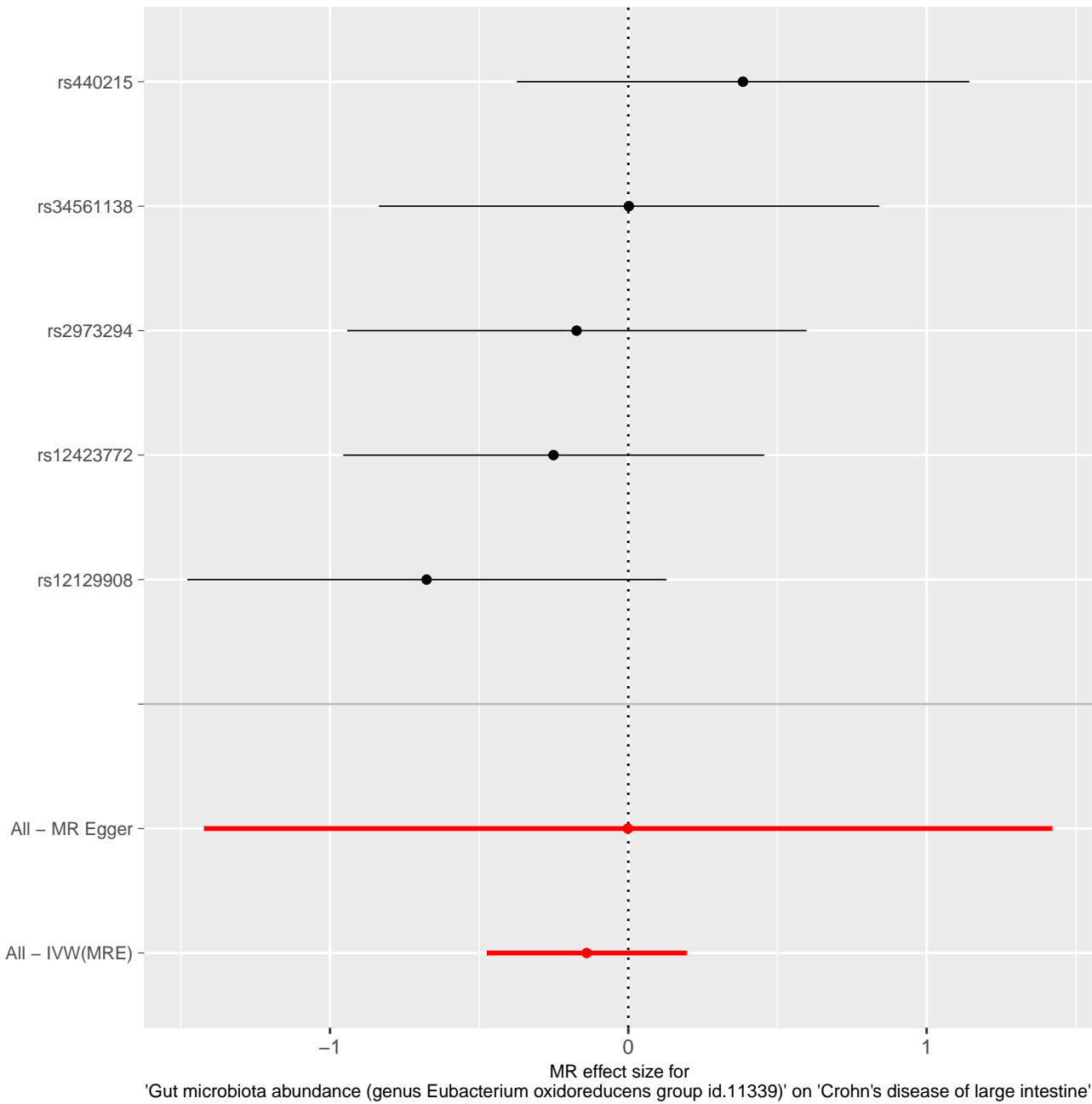


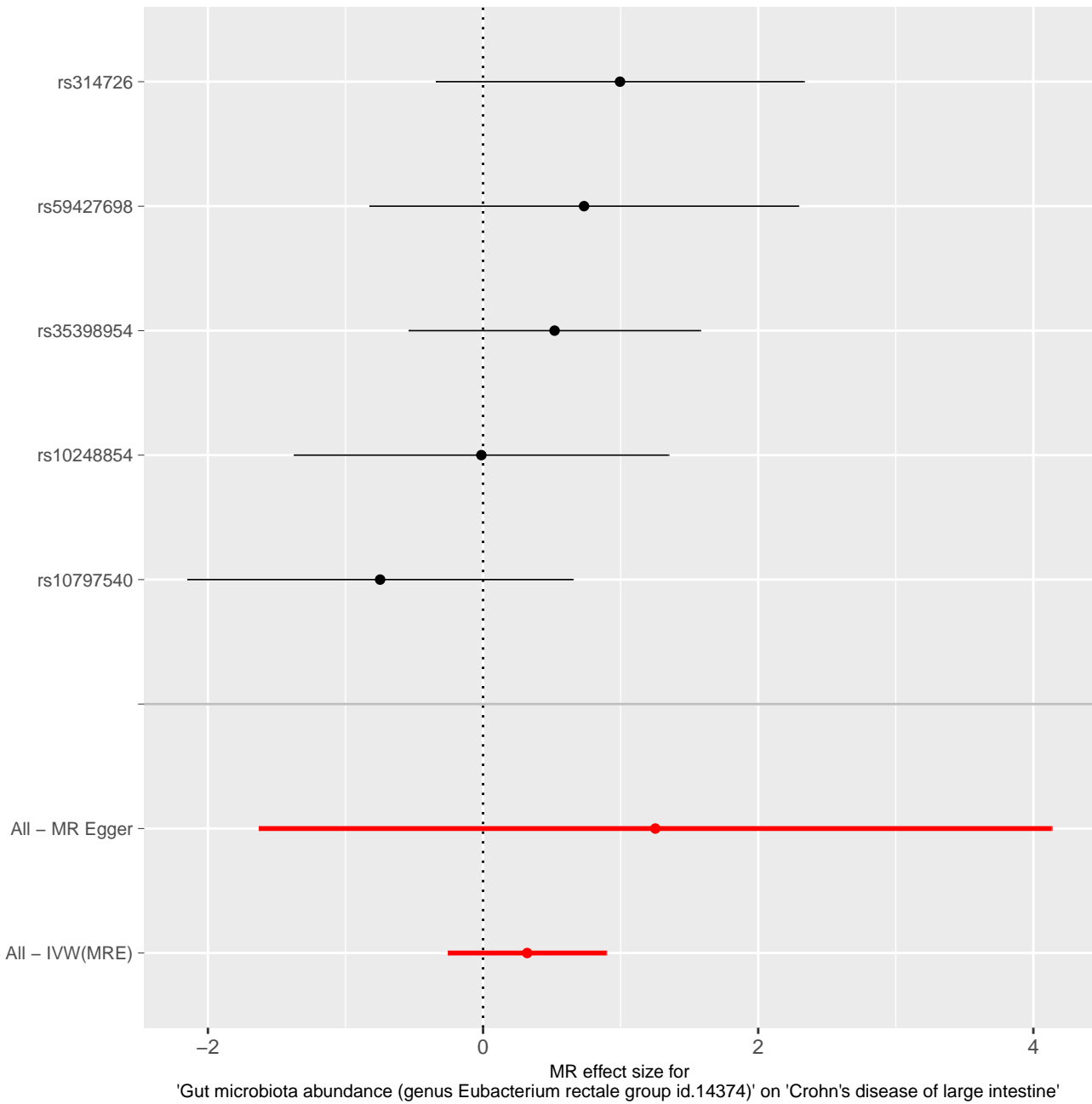
Batch 74 : Gut microbiota abundance (genus Eubacterium hallii group id.11338) on Crohn's disease of large intestine



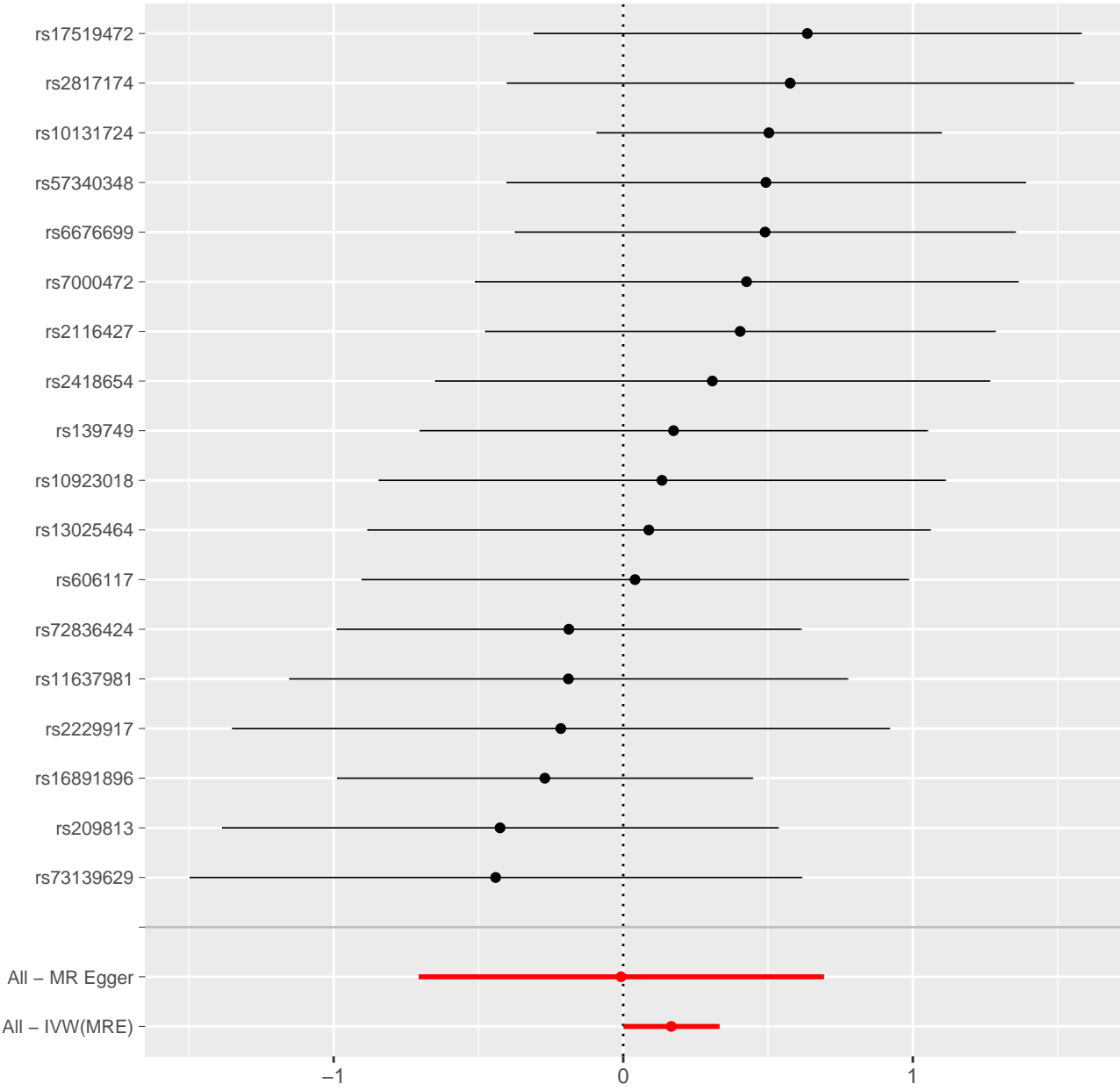
Batch 75 : Gut microbiota abundance (genus Eubacterium nodatum group id.11297) on Crohn's disease of large intestine





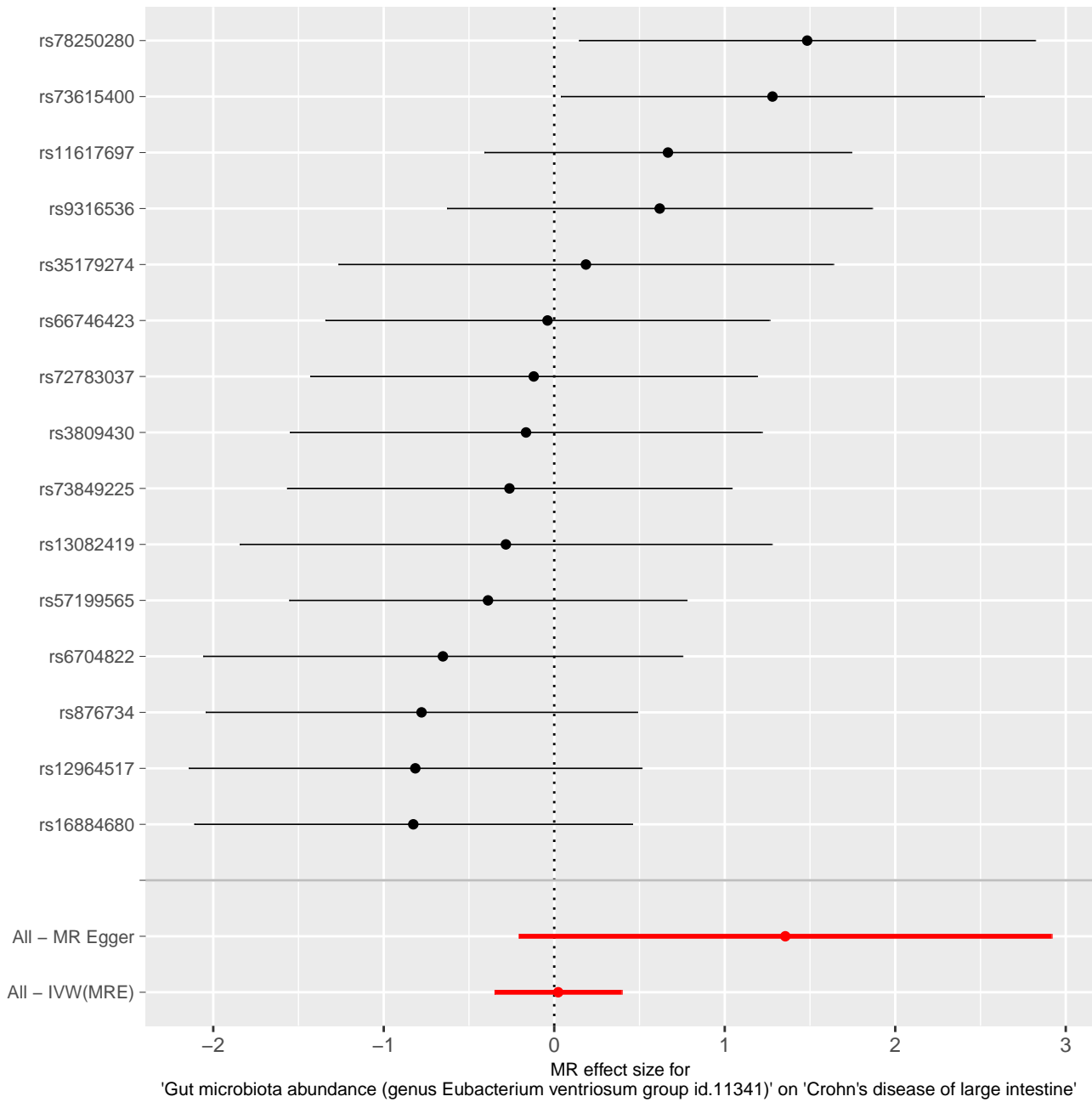


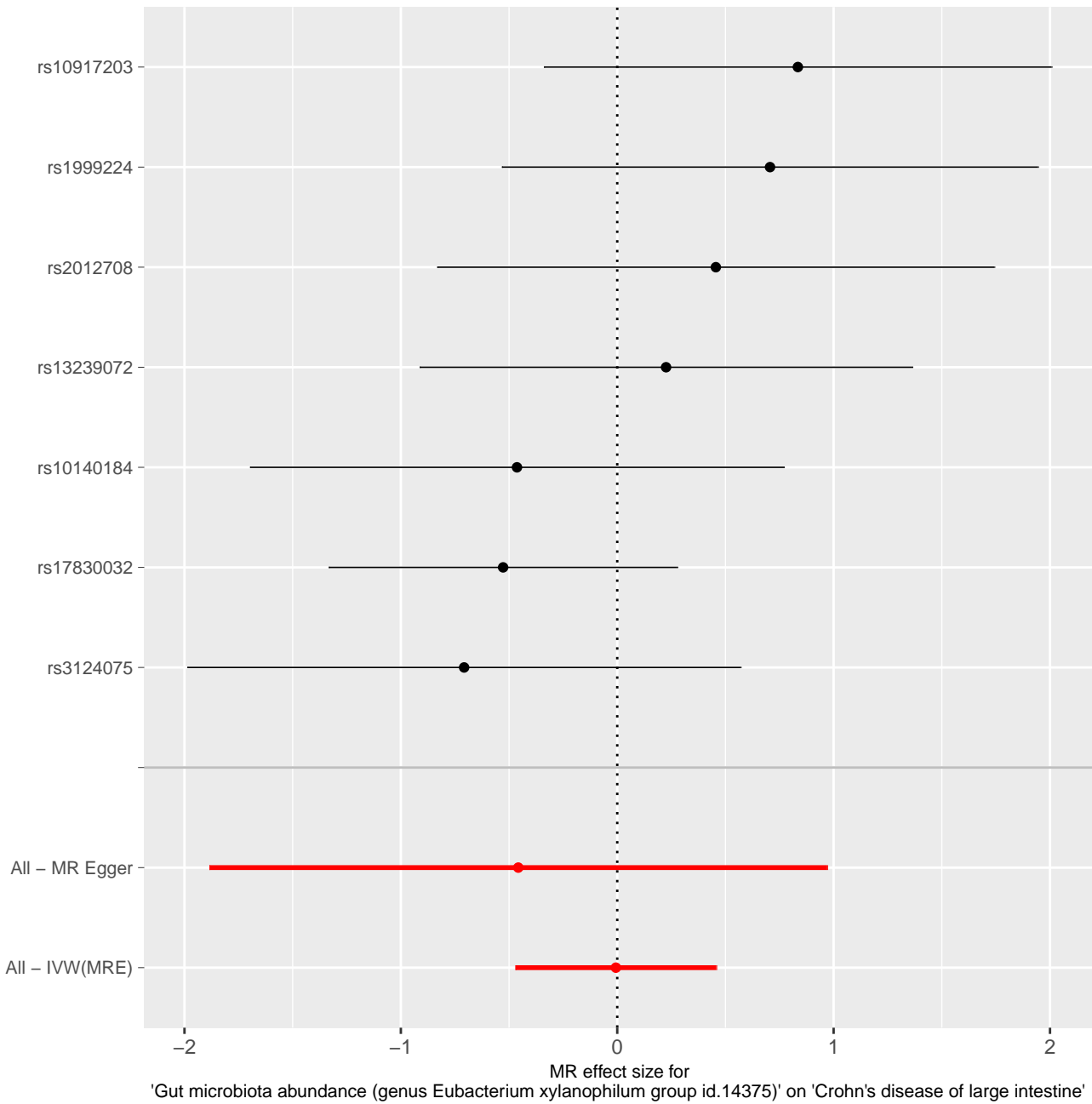
Batch 78 : Gut microbiota abundance (genus Eubacterium ruminantium group id.11340) on Crohn's disease of large intestine



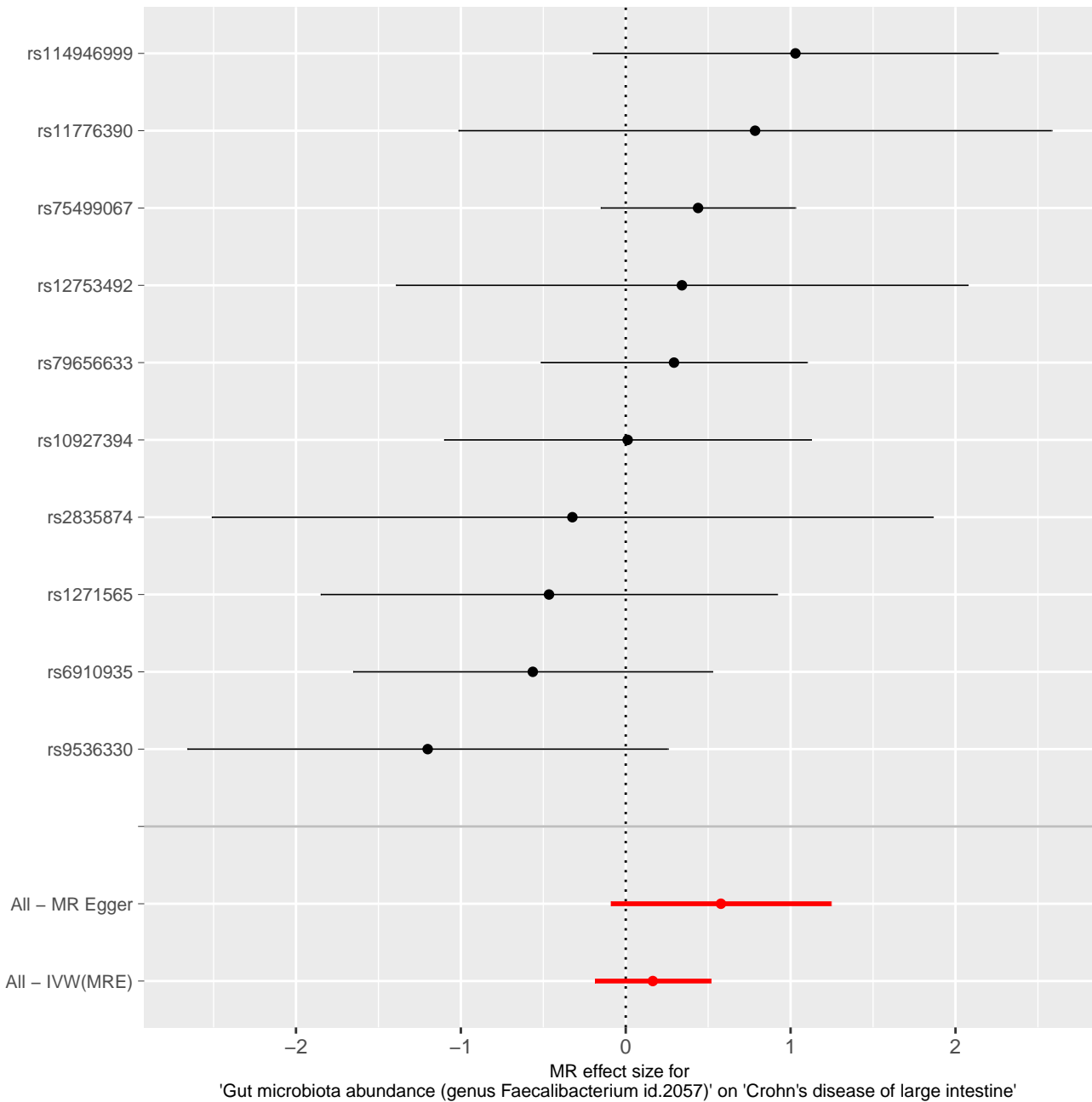
MR effect size for 'Gut microbiota abundance (genus Eubacterium ruminantium group id.11340)' on 'Crohn's disease of large intestine'

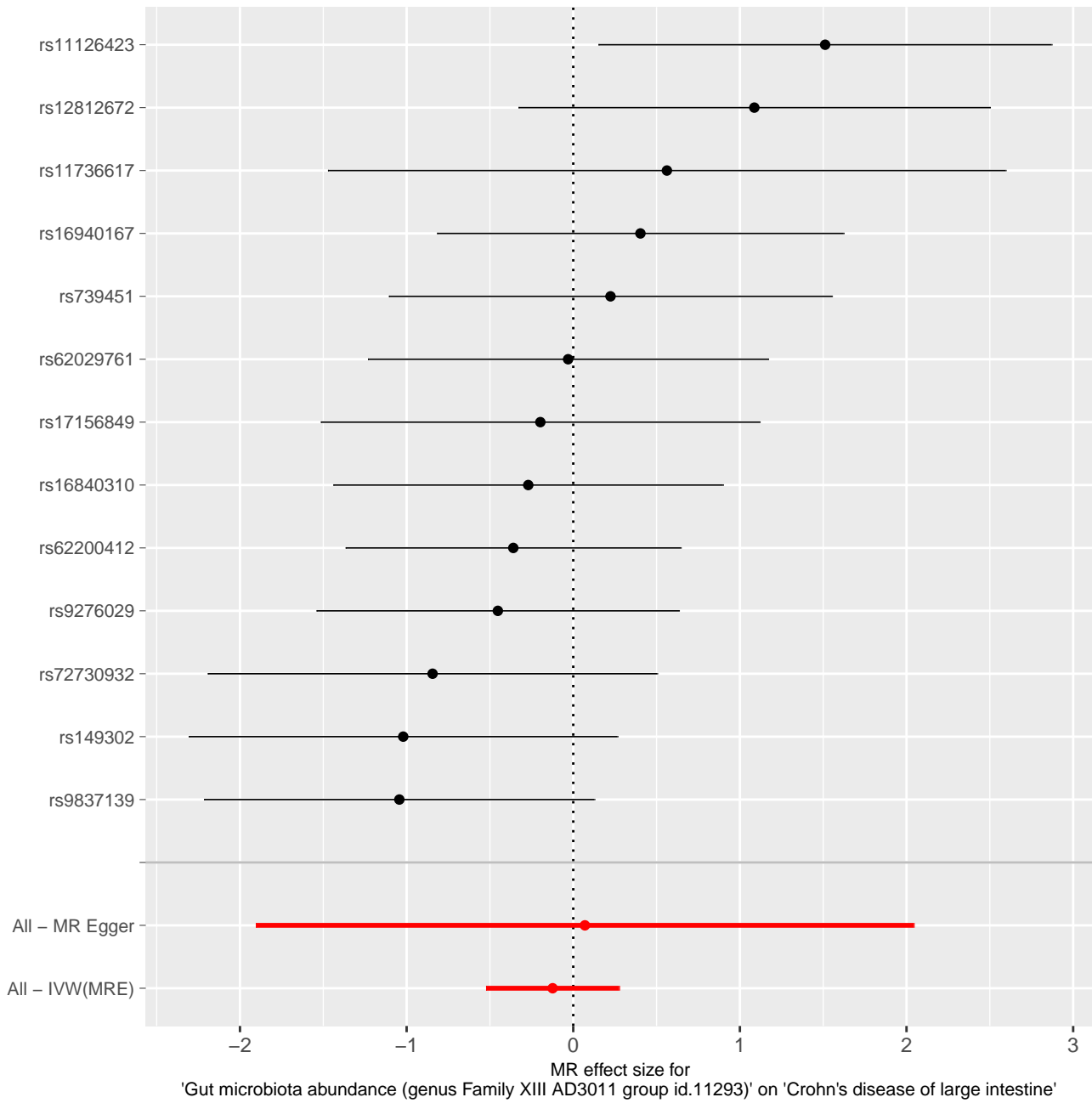
Batch 79 : Gut microbiota abundance (genus Eubacterium ventriosum group id.11341) on Crohn's disease of large intestine



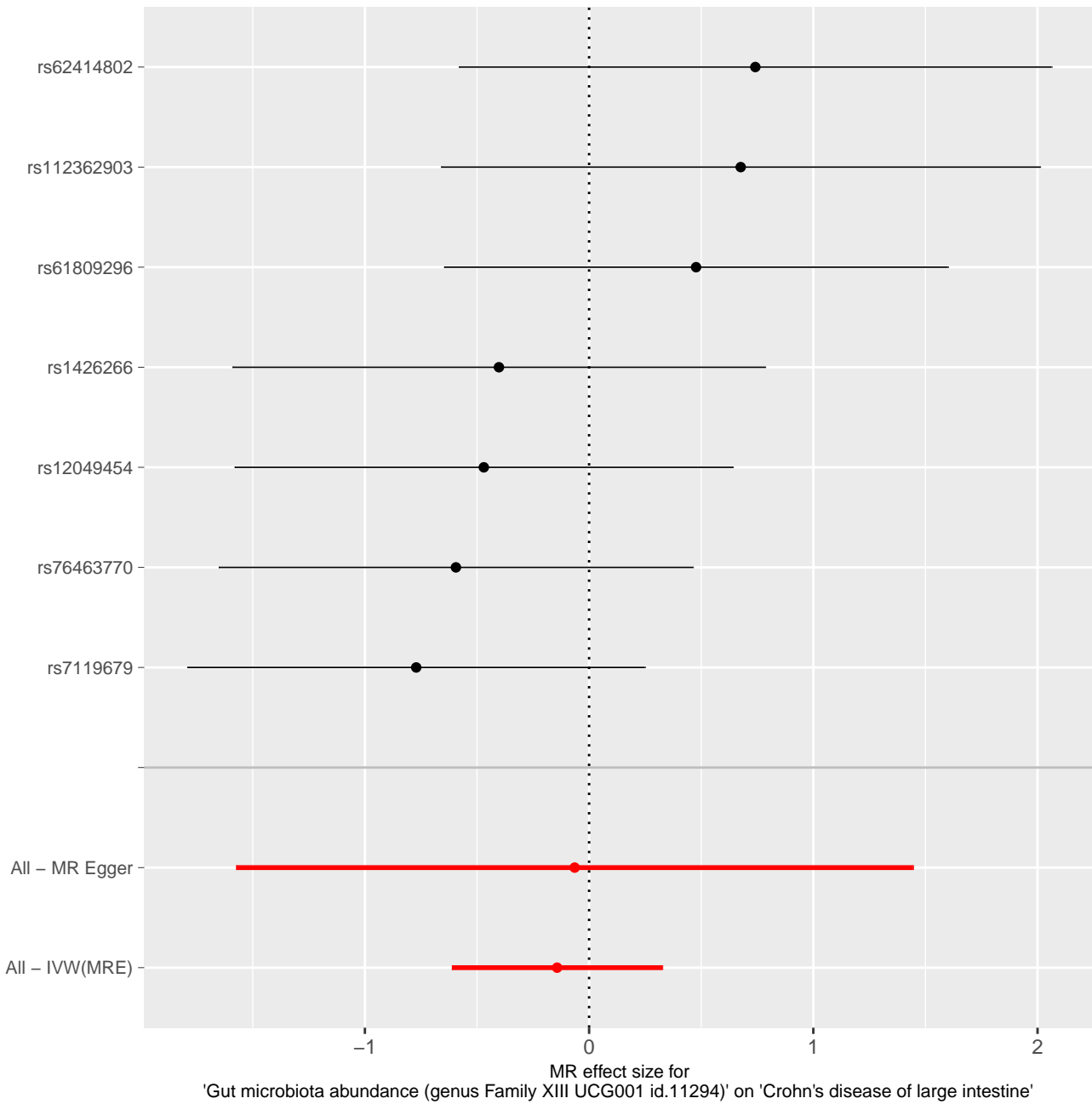


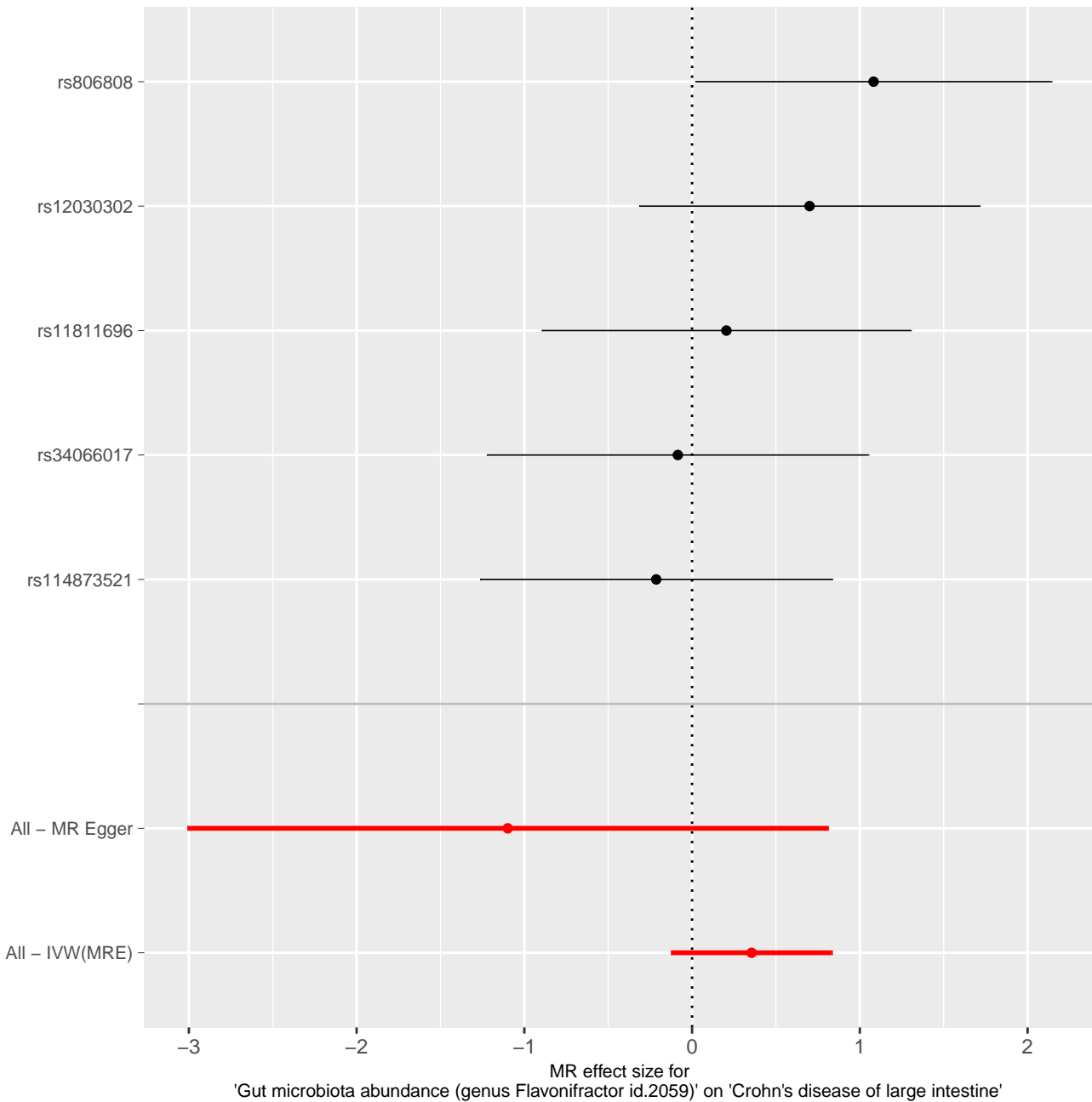
Batch 81 : Gut microbiota abundance (genus Faecalibacterium id.2057) on Crohn's disease of large intestine



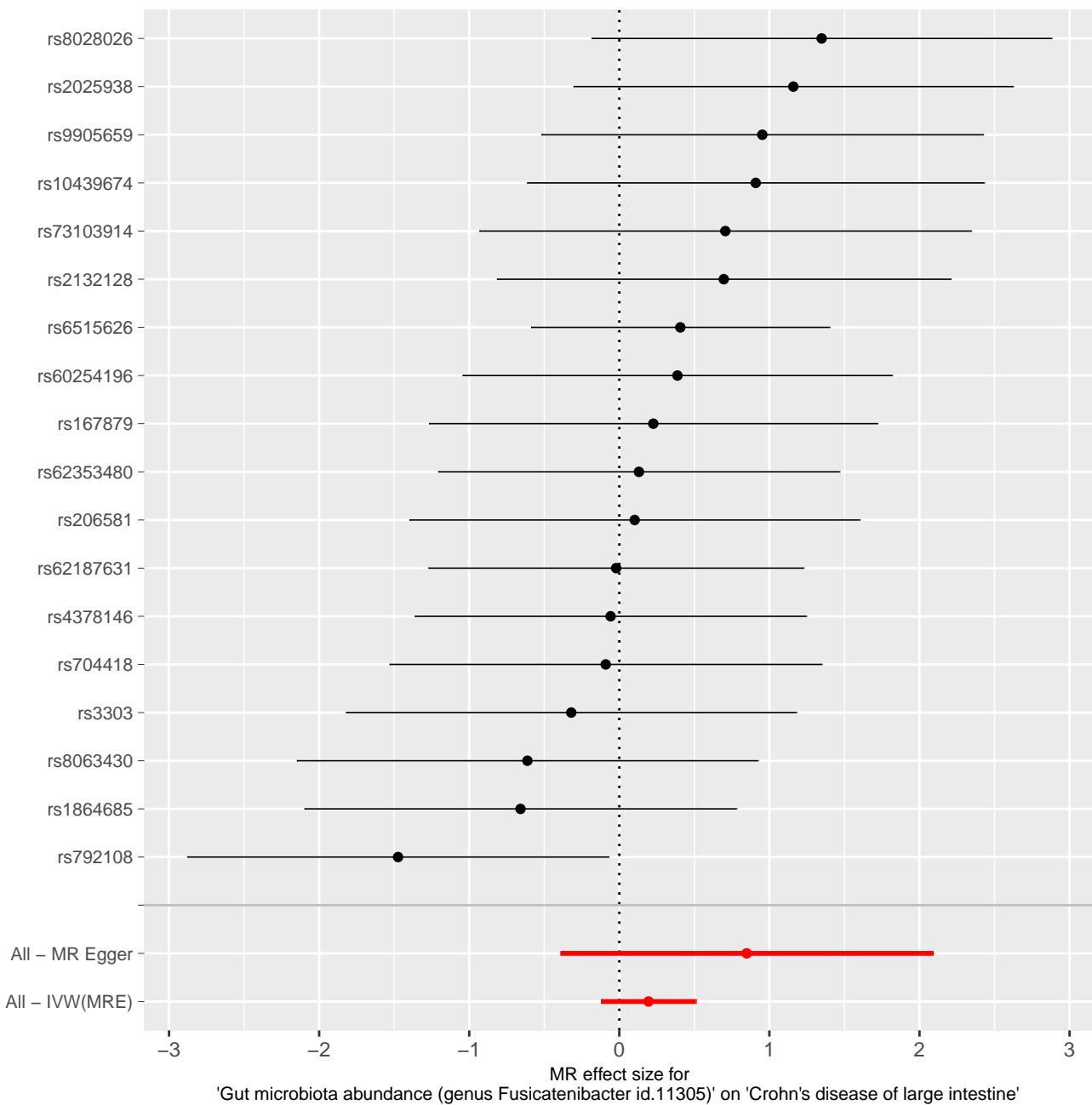


Batch 83 : Gut microbiota abundance (genus Family XIII UCG001 id.11294) on Crohn's disease of large intestine

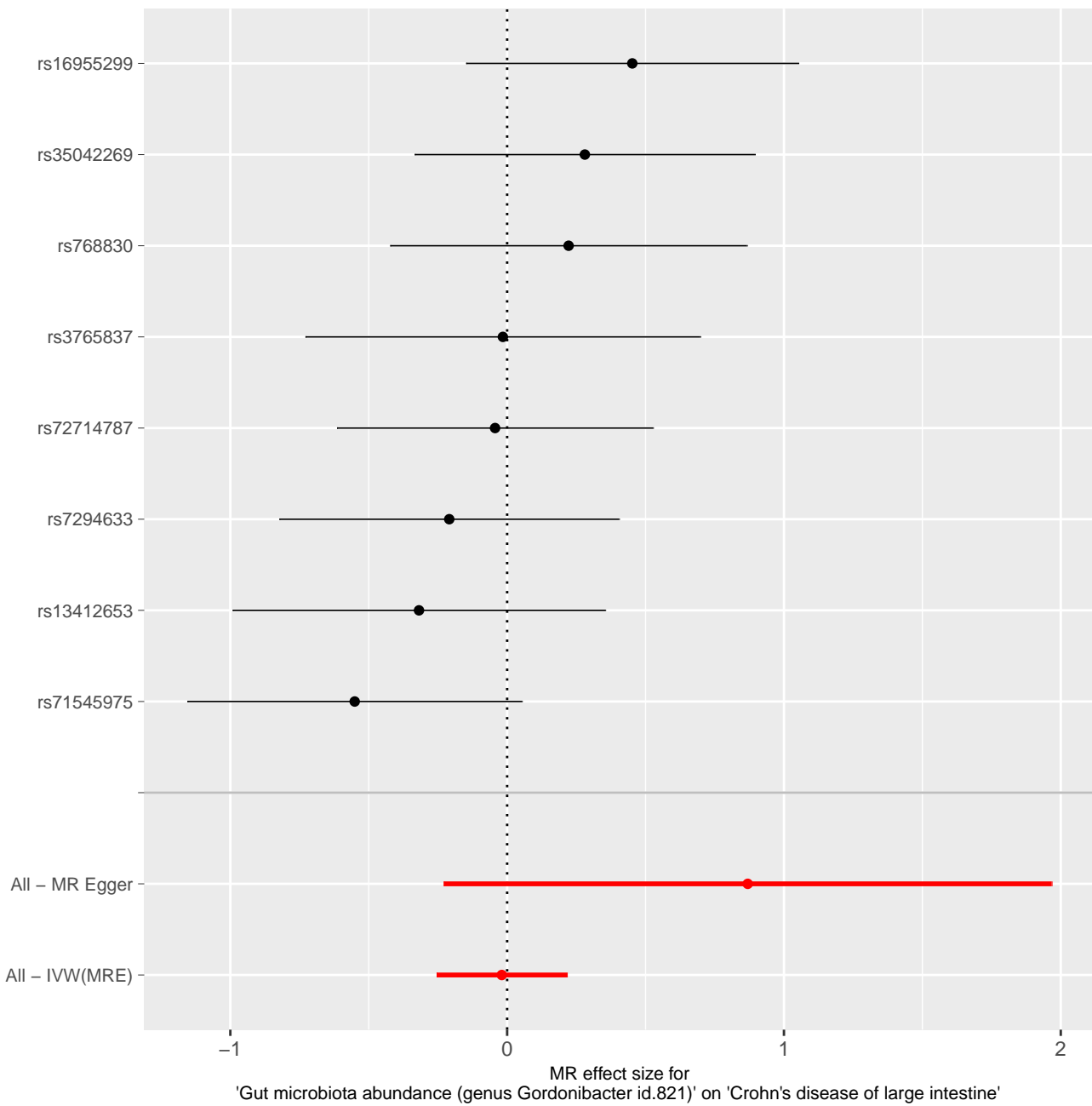




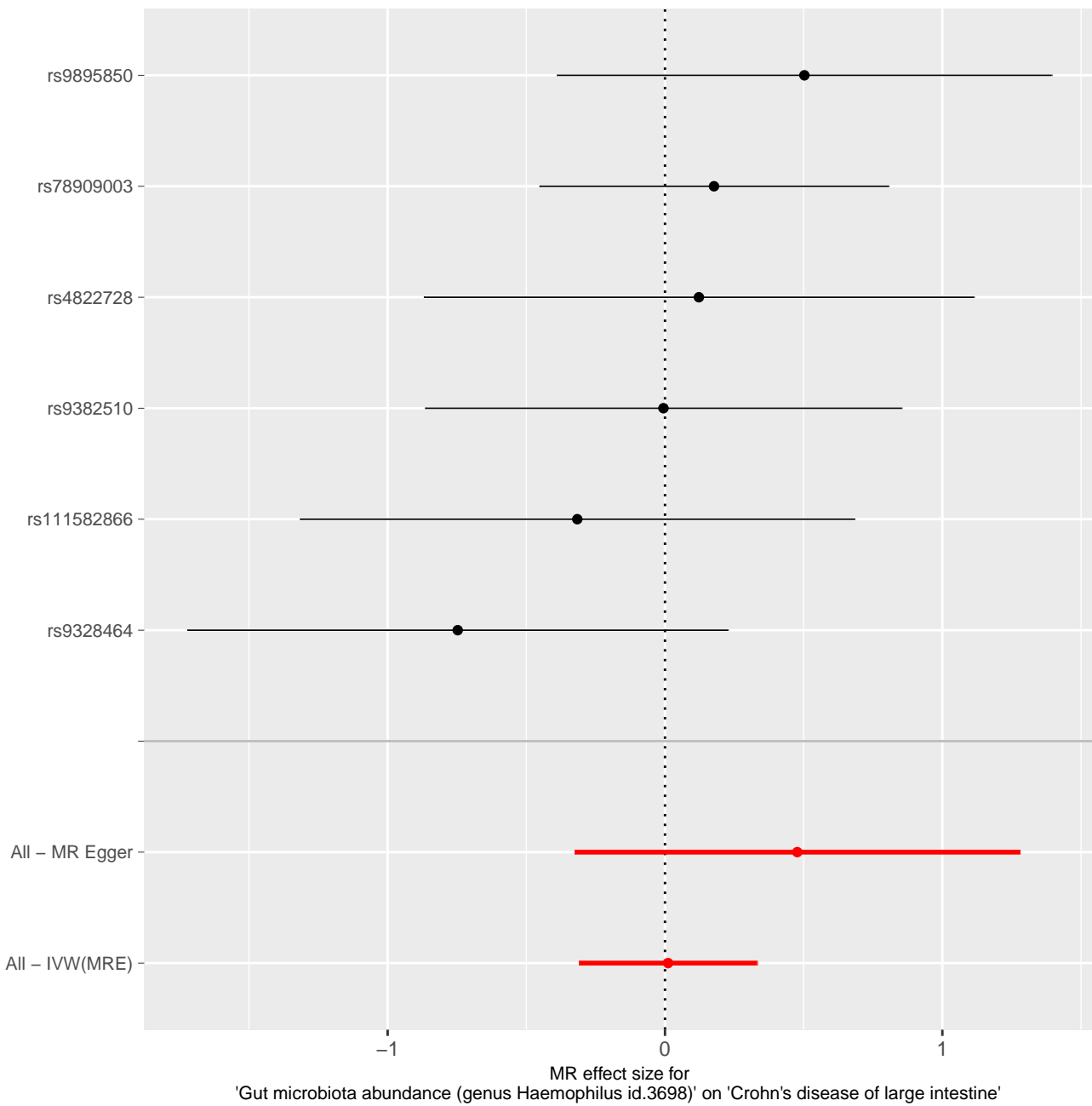
Batch 85 : Gut microbiota abundance (genus Fusicatenibacter id.11305) on Crohn's disease of large intestine



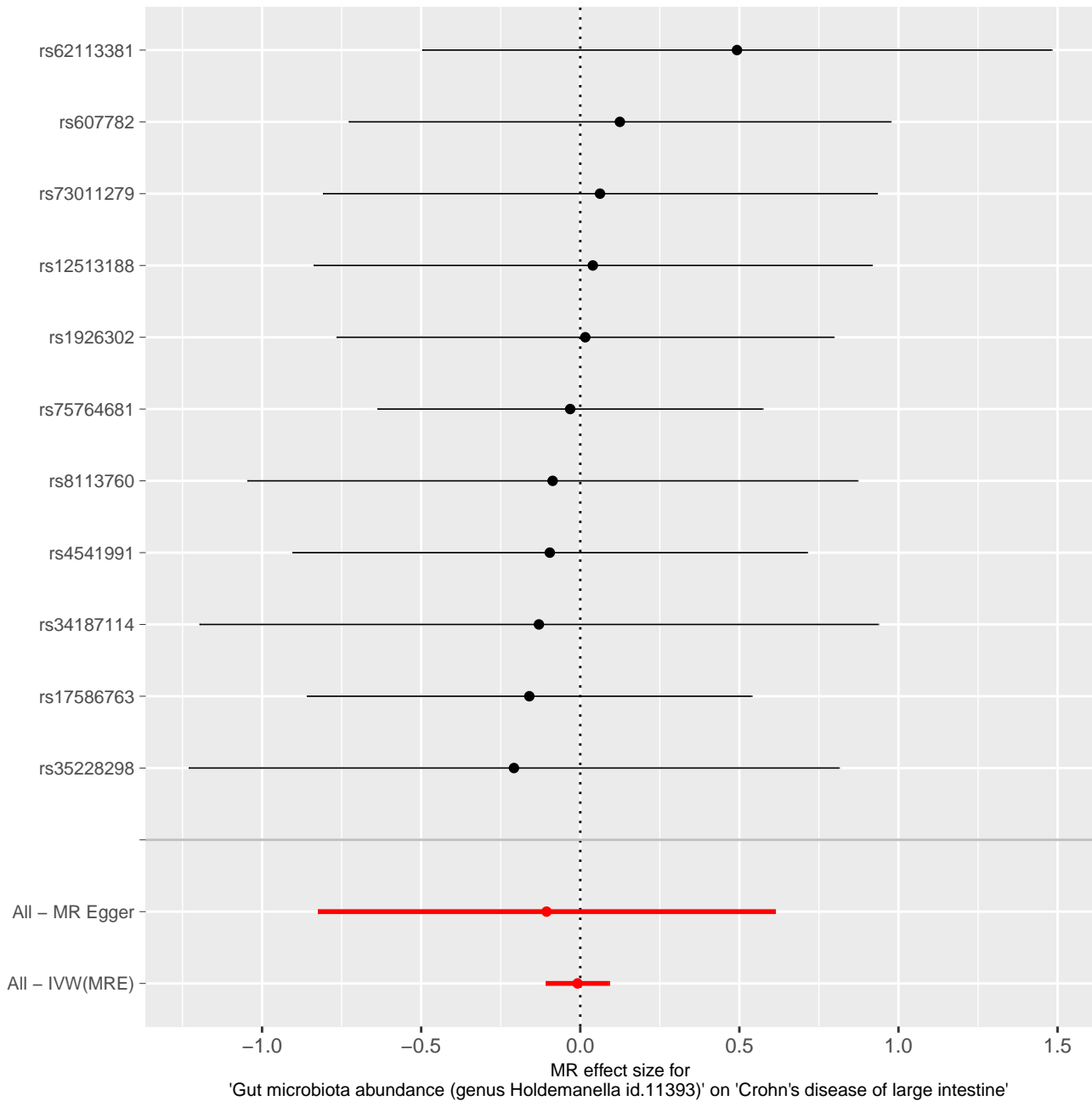
Batch 86 : Gut microbiota abundance (genus Gordonibacter id.821) on Crohn's disease of large intestine



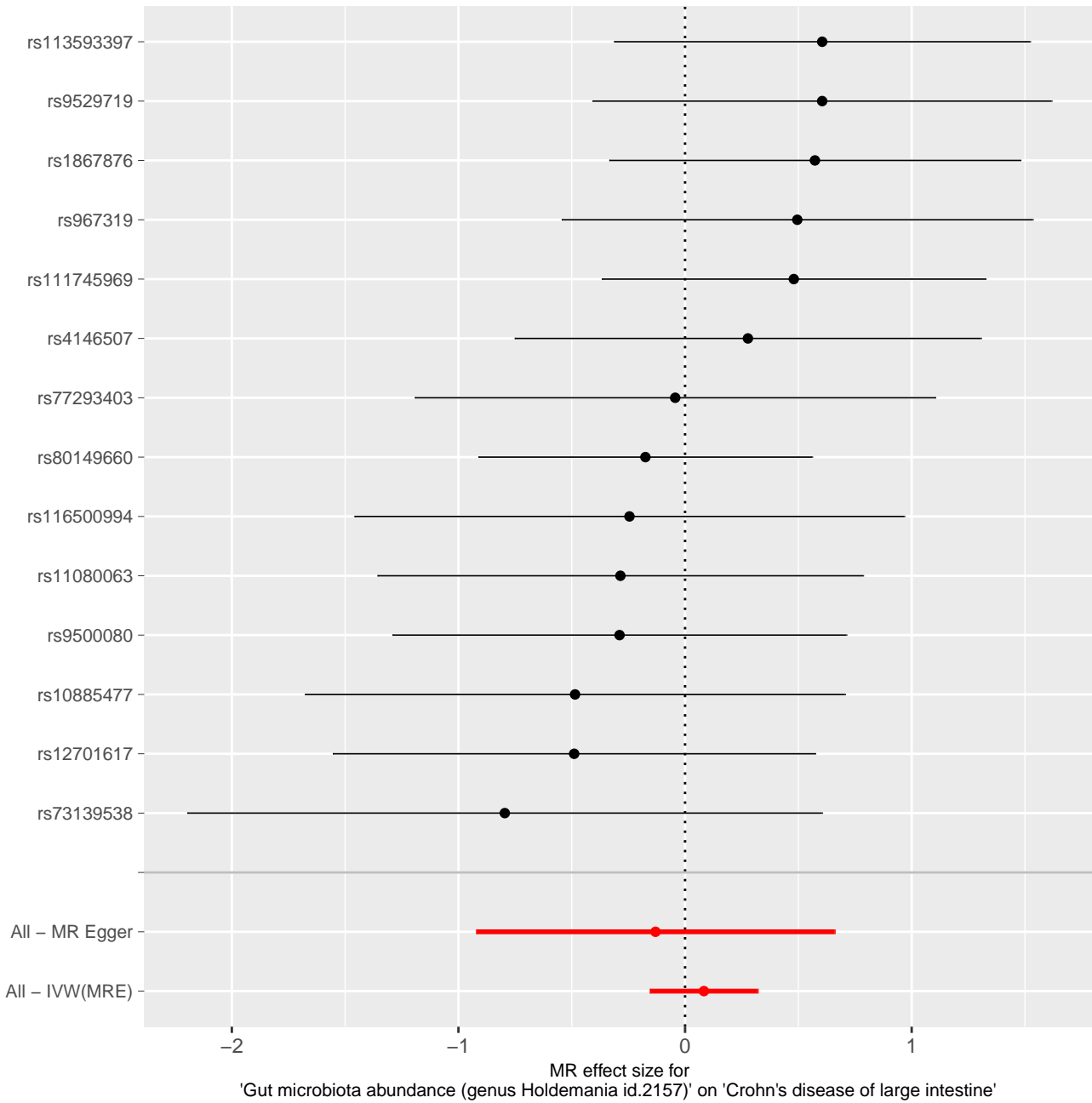
Batch 87 : Gut microbiota abundance (genus Haemophilus id.3698) on Crohn's disease of large intestine



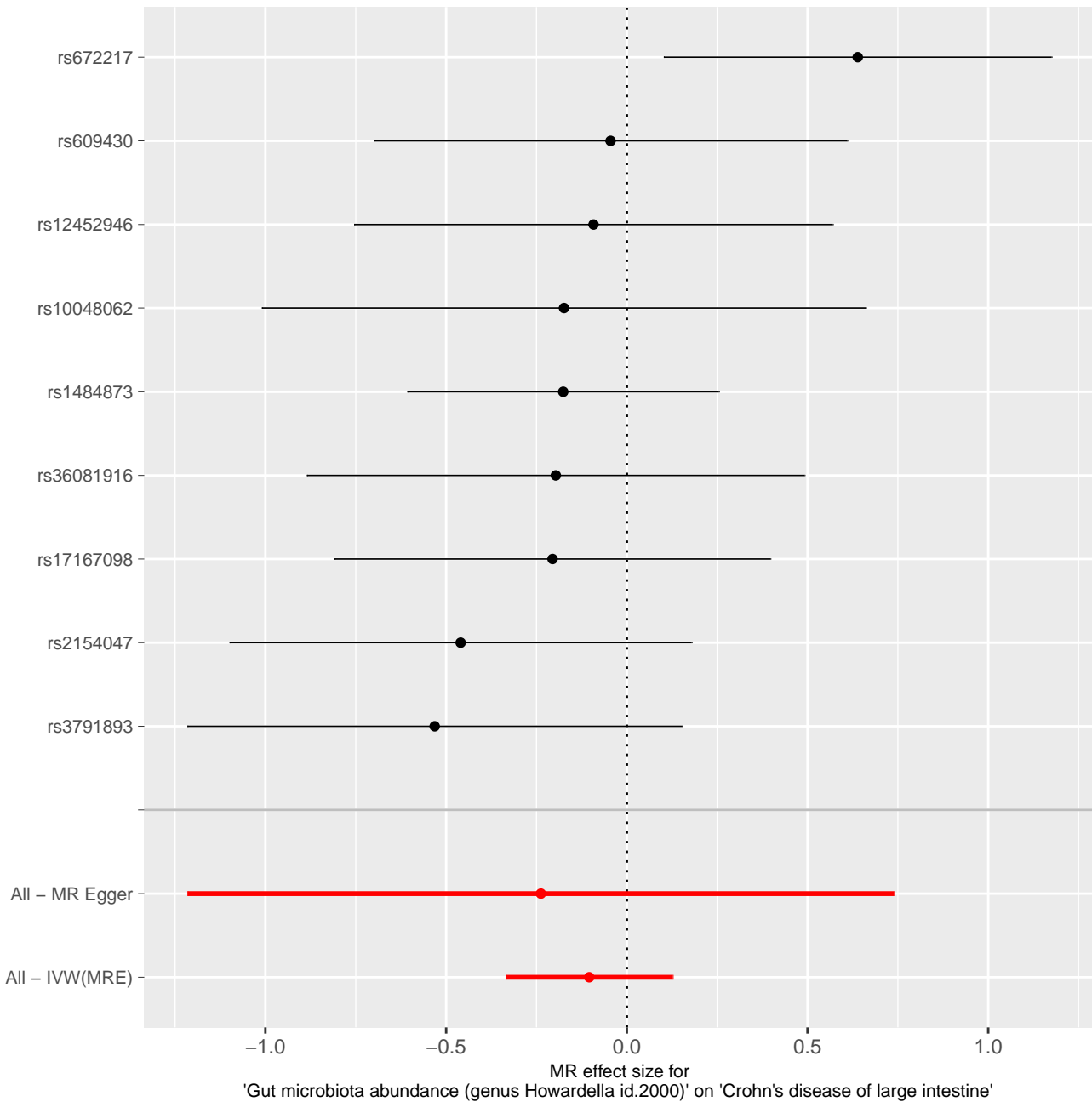
Batch 88 : Gut microbiota abundance (genus Holdemanella id.11393) on Crohn's disease of large intestine

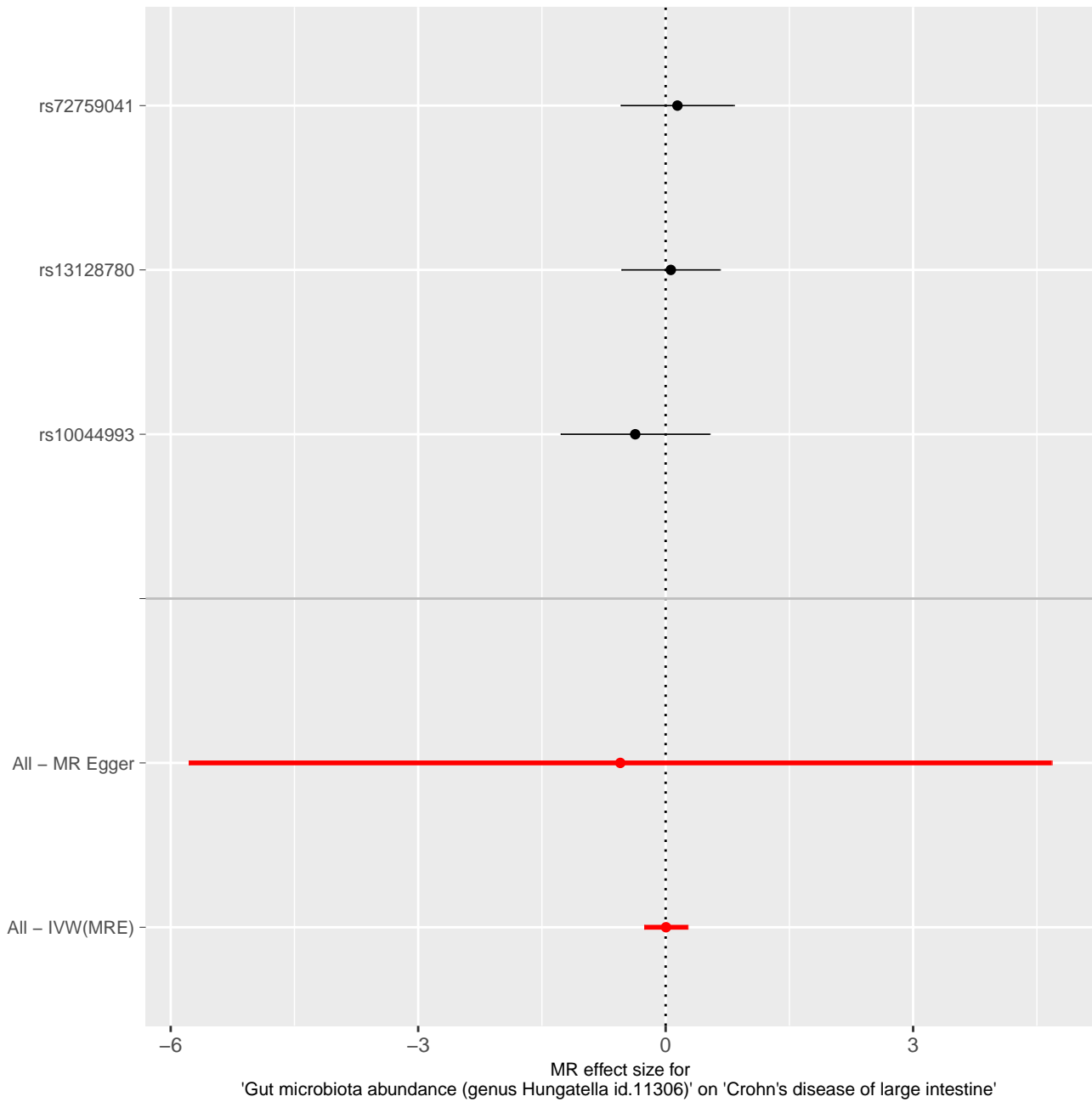


Batch 89 : Gut microbiota abundance (genus Holdemania id.2157) on Crohn's disease of large intestine

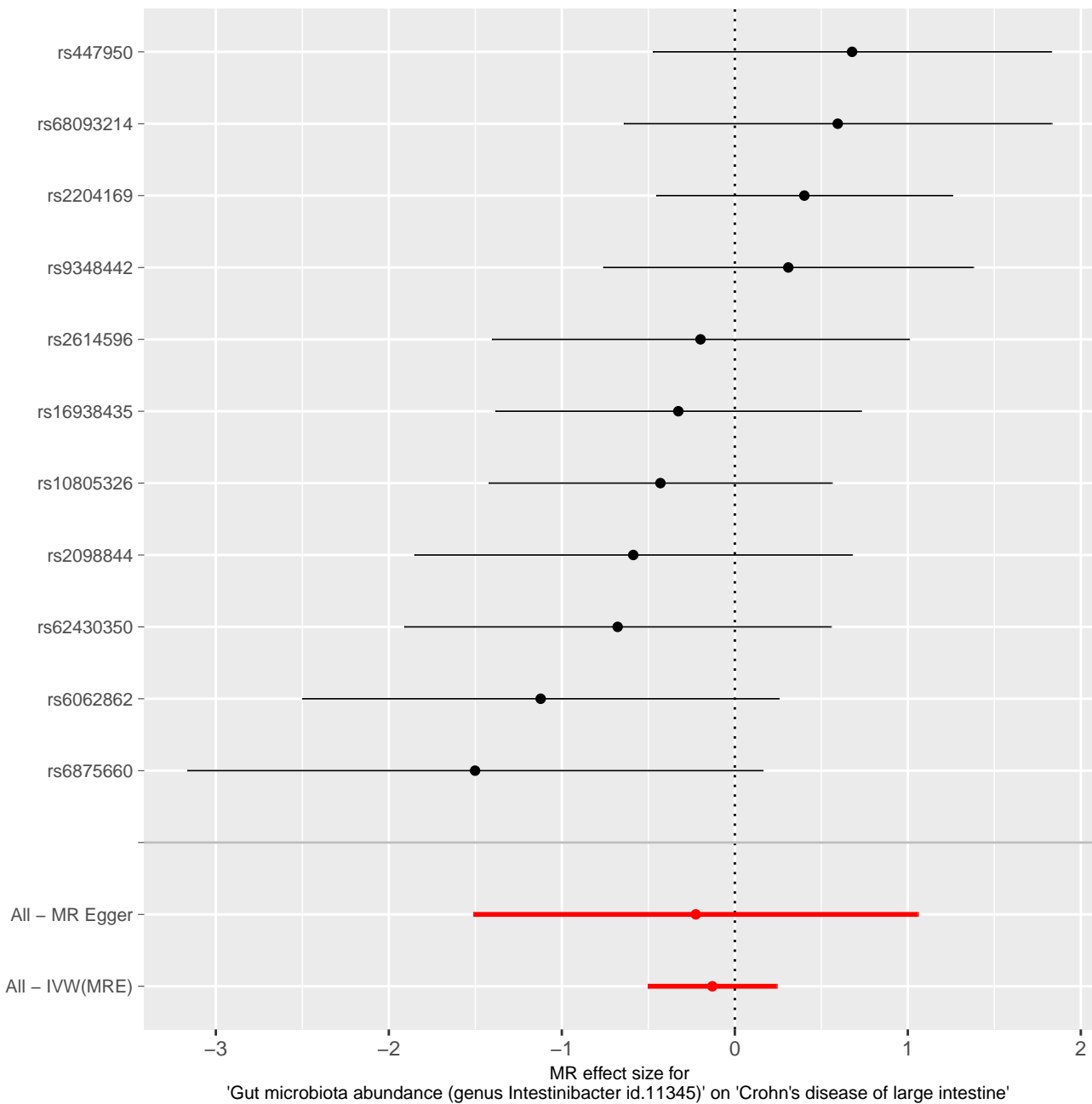


Batch 90 : Gut microbiota abundance (genus Howardella id.2000) on Crohn's disease of large intestine

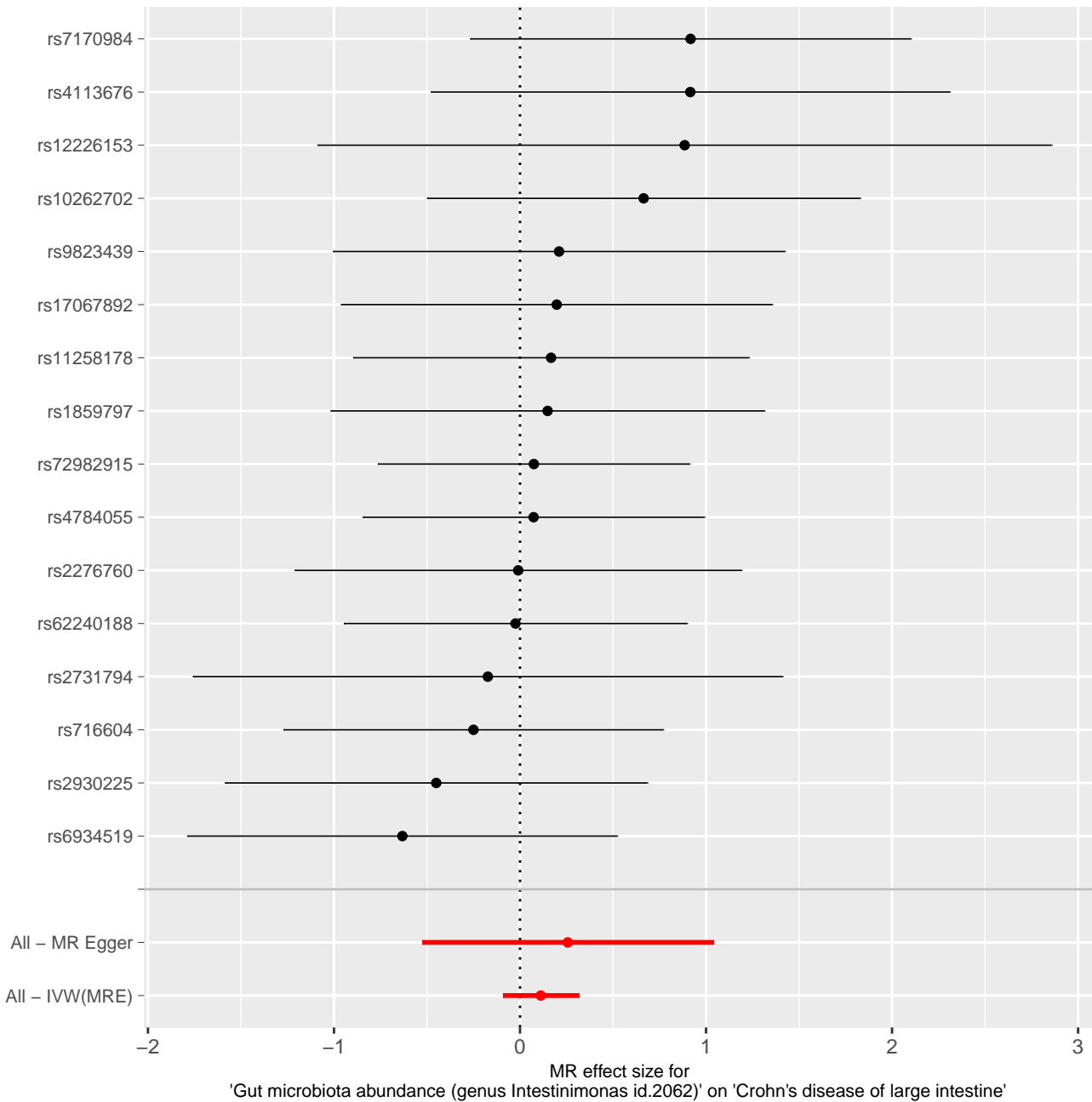




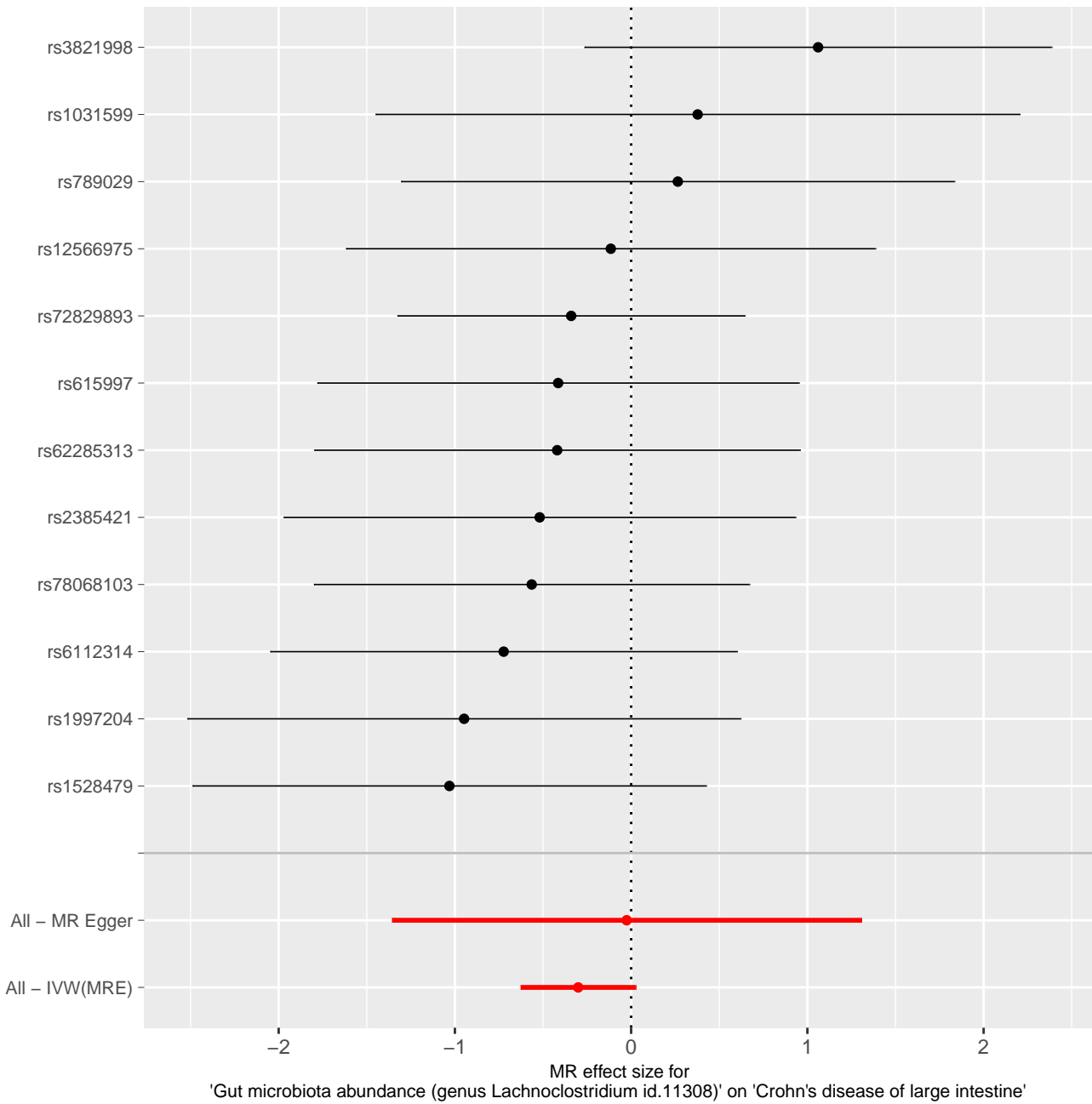
Batch 92 : Gut microbiota abundance (genus Intestinibacter id.11345) on Crohn's disease of large intestine

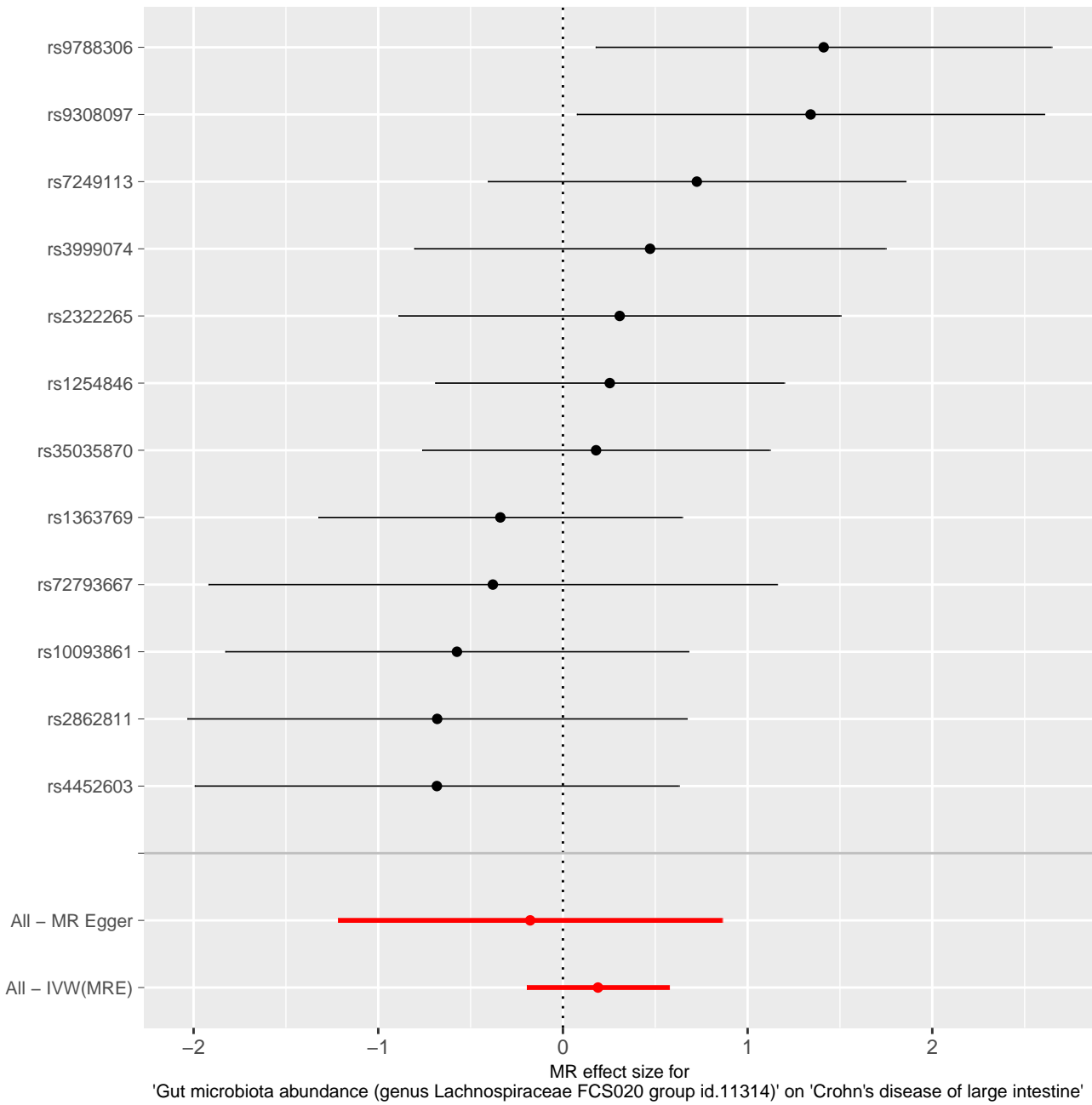


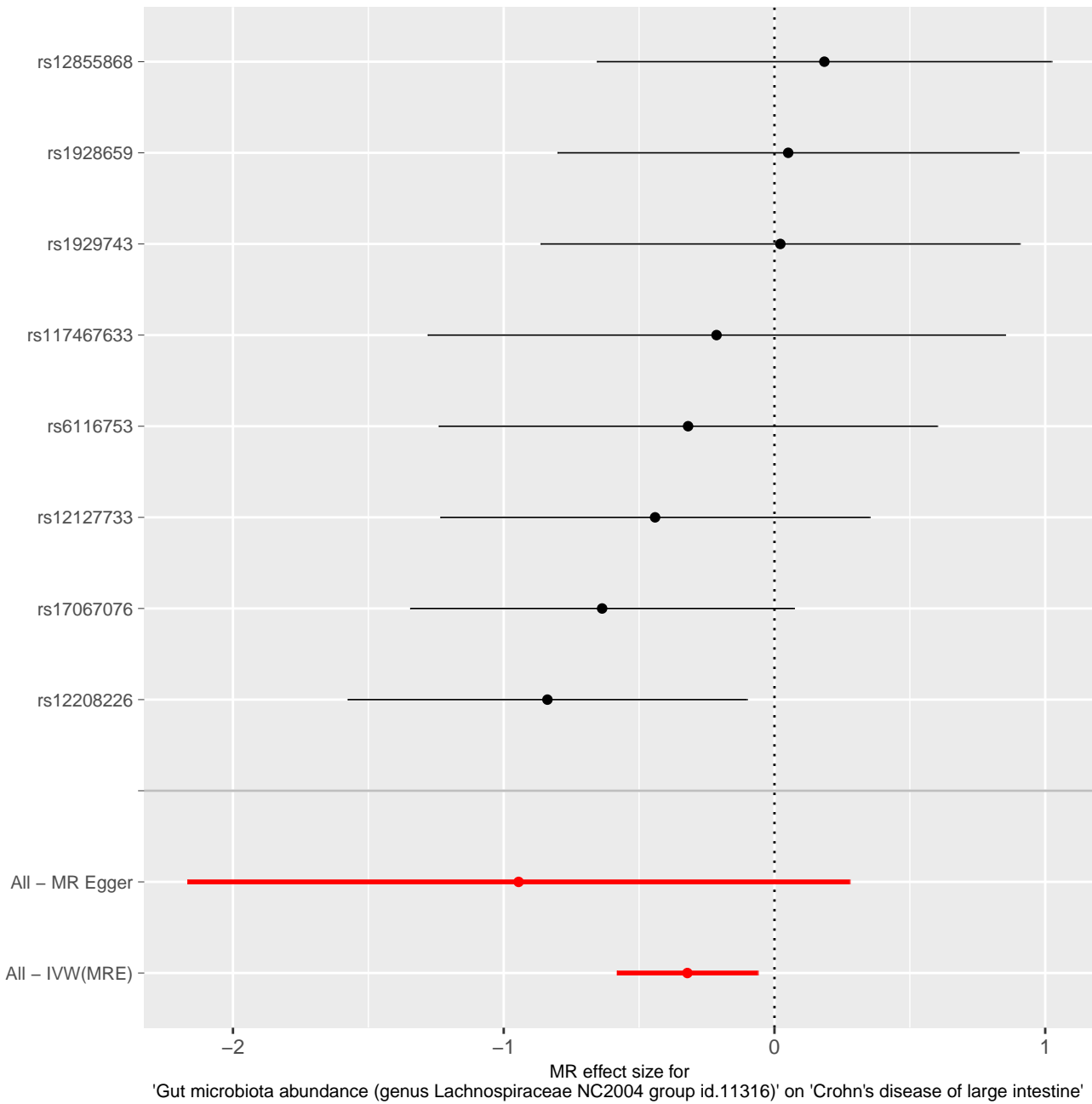
Batch 93 : Gut microbiota abundance (genus Intestinimonas id.2062) on Crohn's disease of large intestine



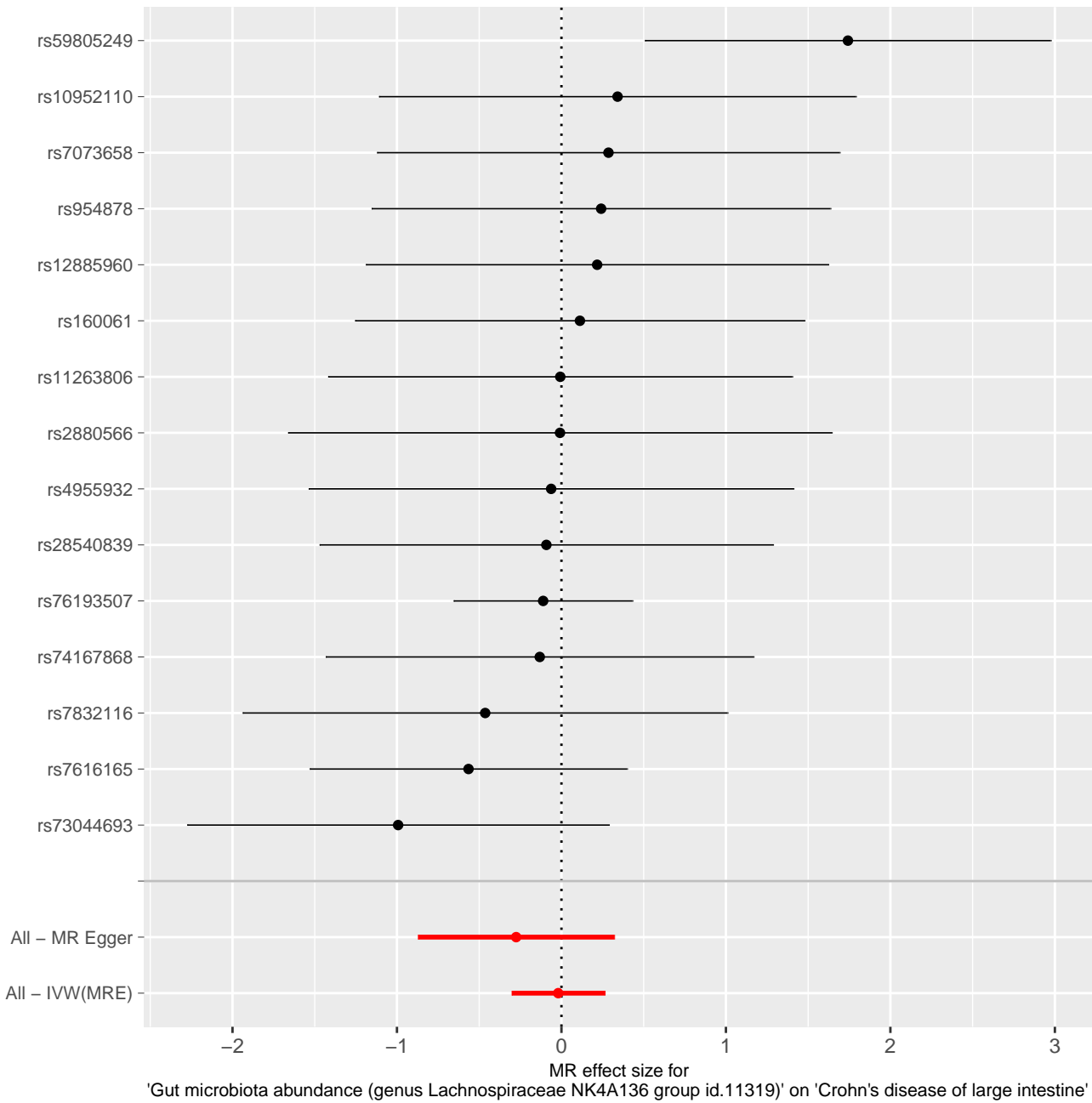
Batch 94 : Gut microbiota abundance (genus Lachnoclostridium id.11308) on Crohn's disease of large intestine

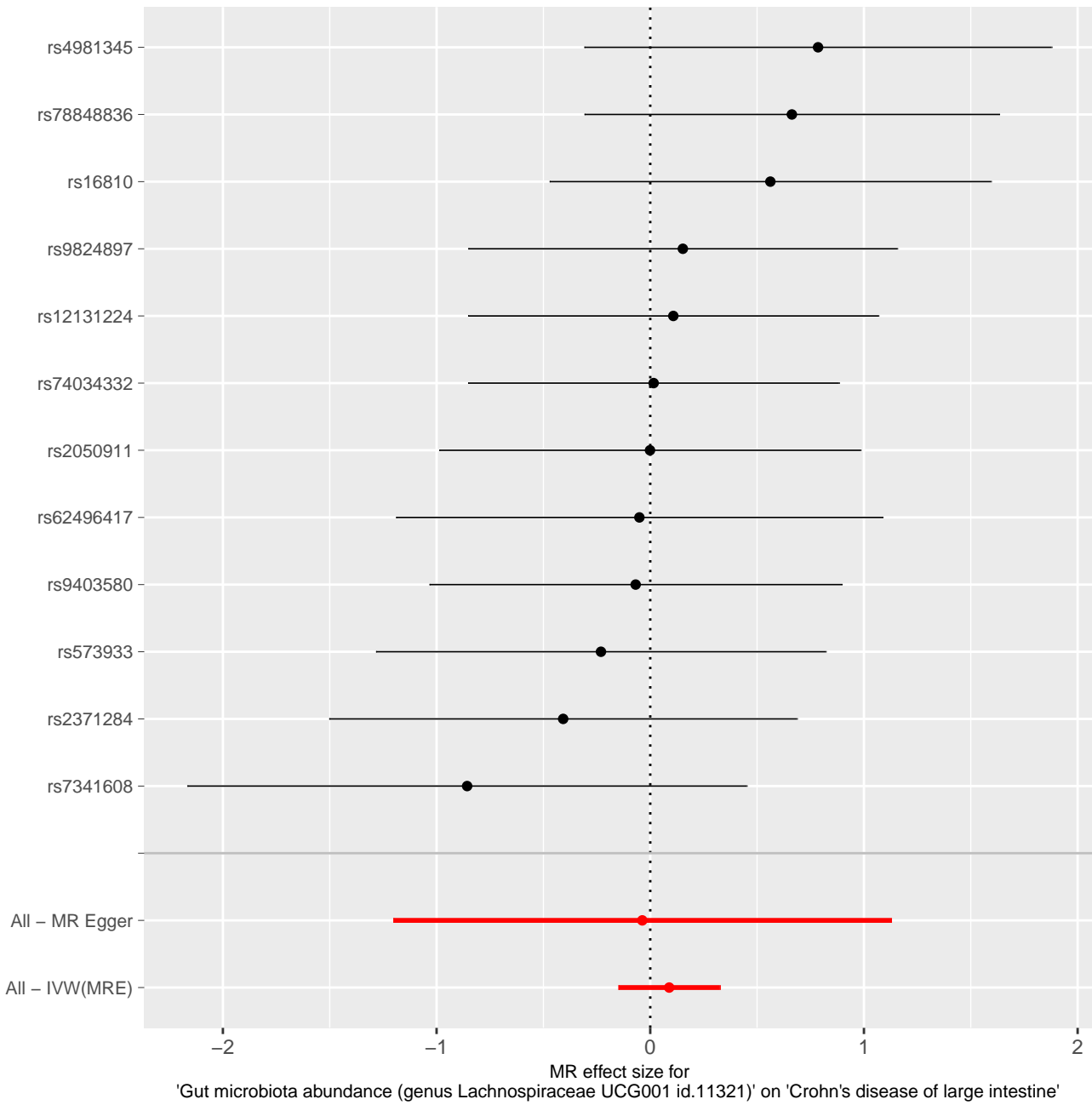


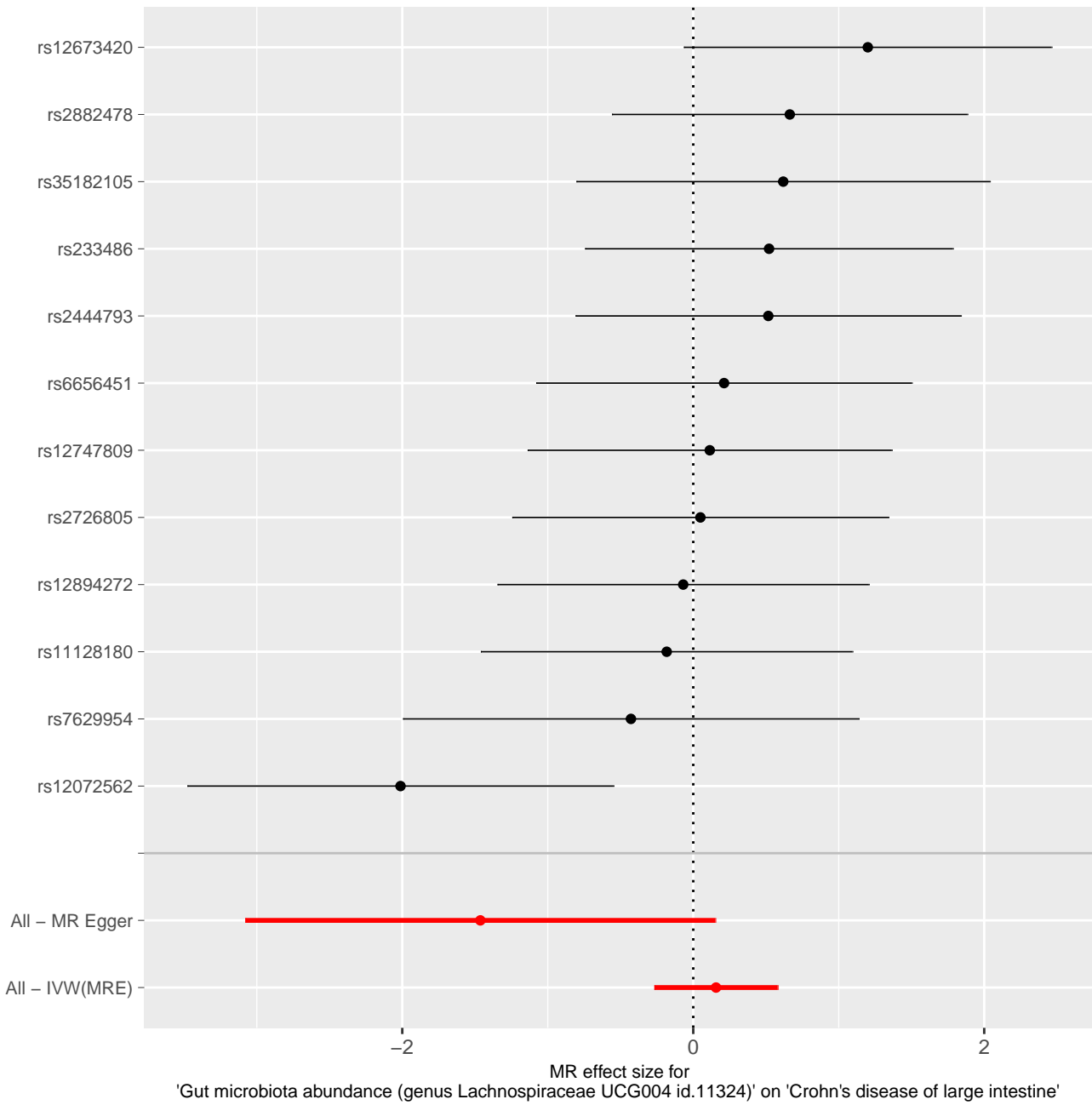


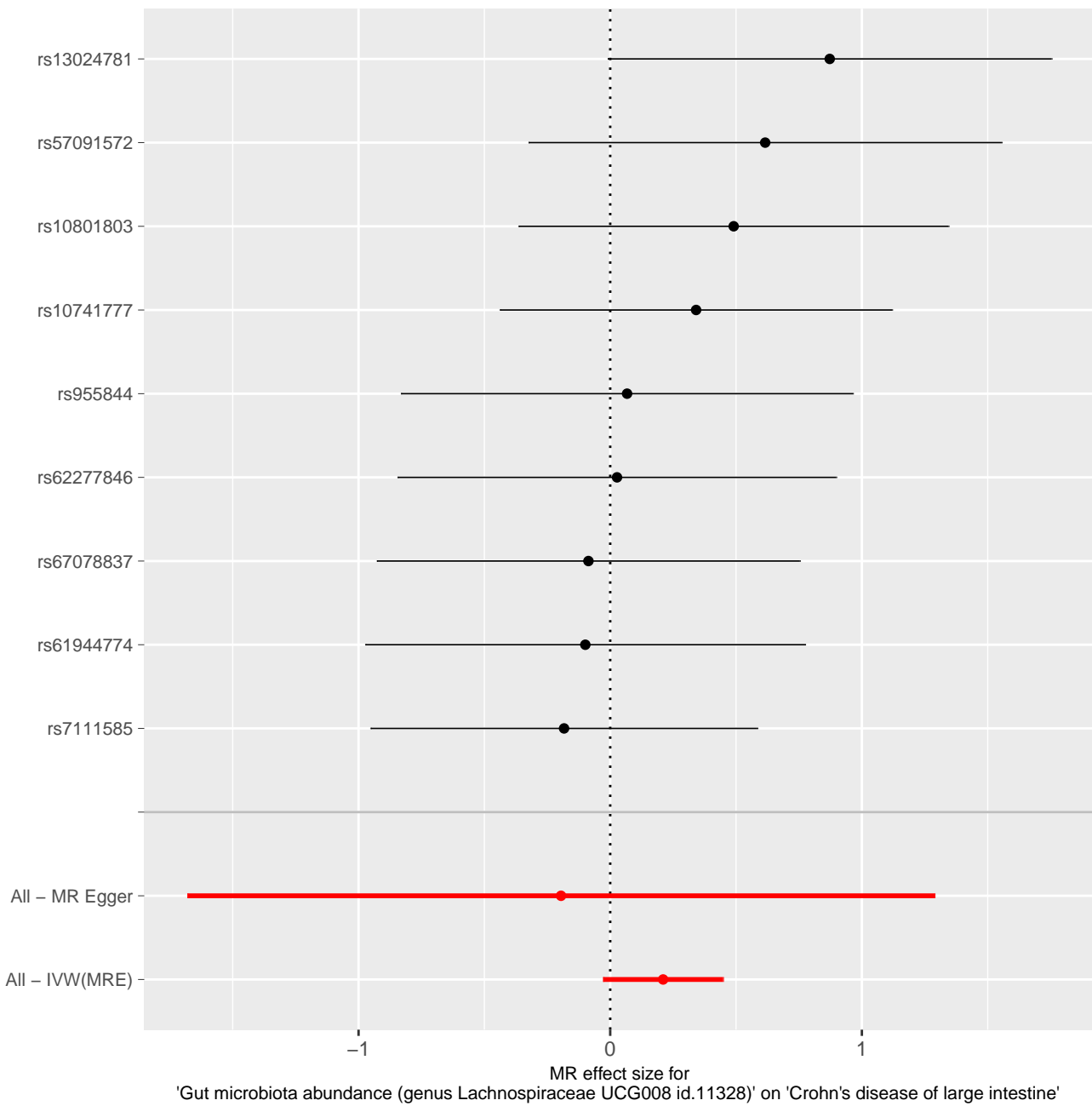


Batch 98 : Gut microbiota abundance (genus Lachnospiraceae NK4A136 group id.11319) on Crohn's disease of large intestine



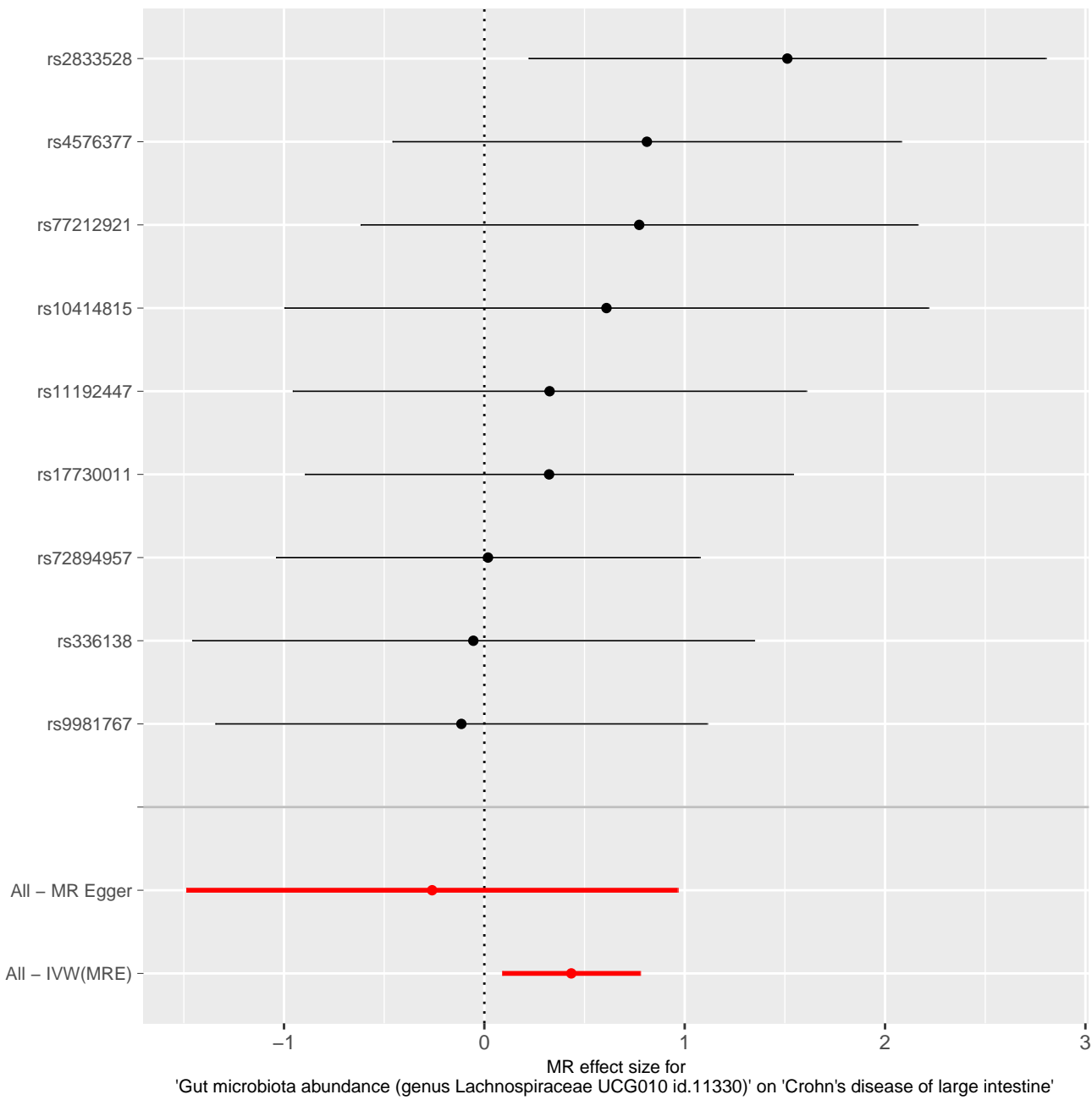


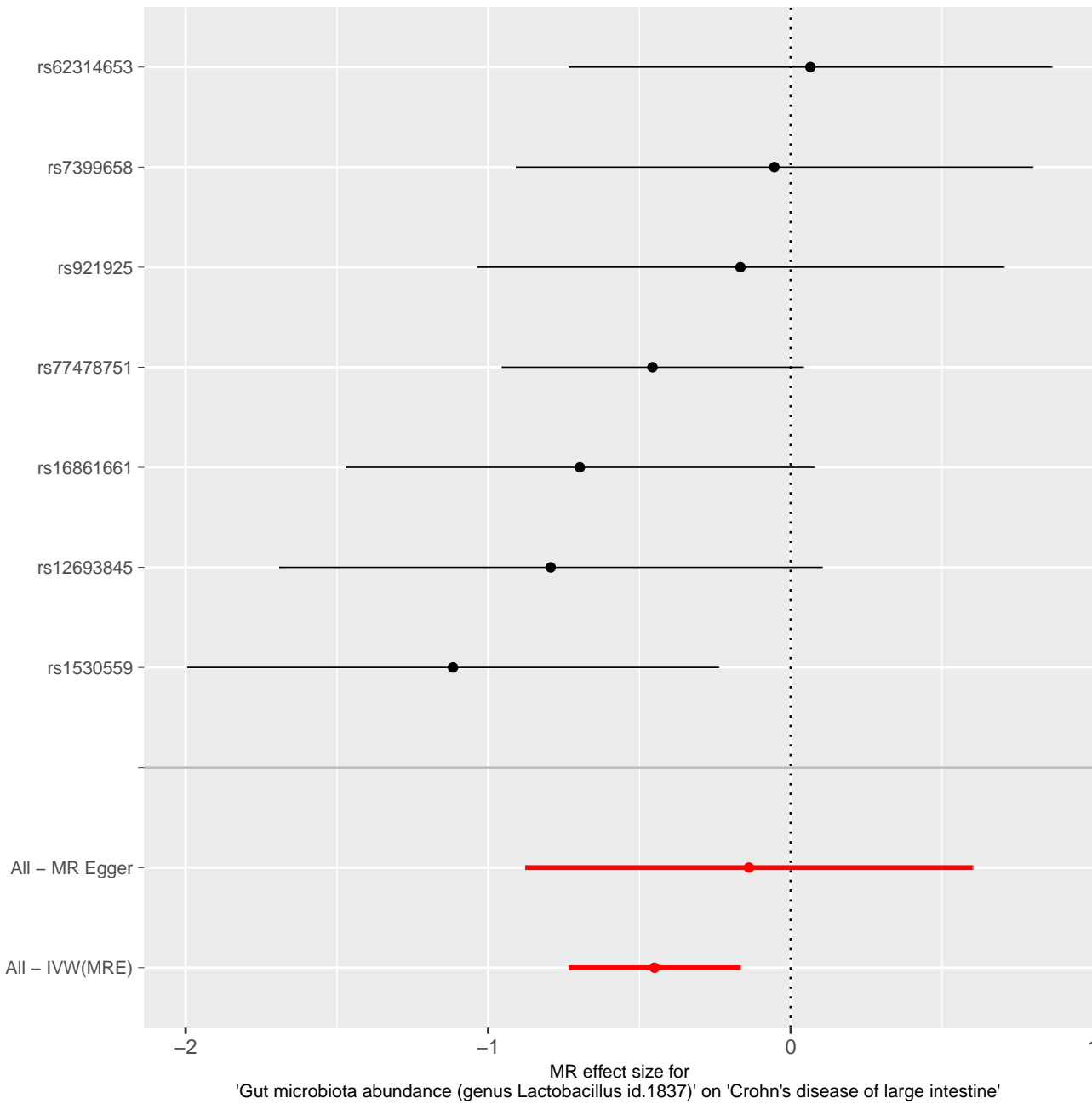




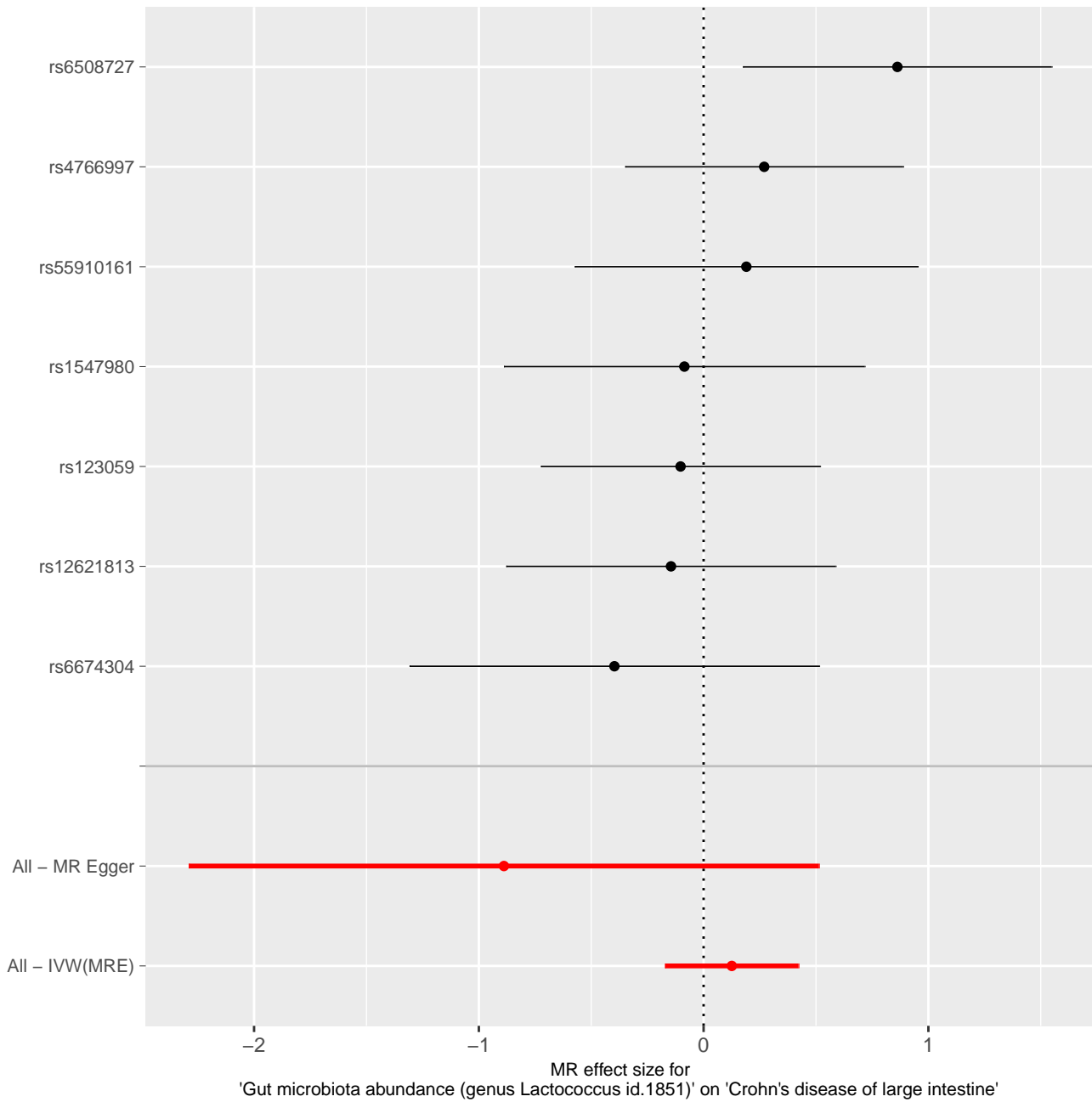
MR effect size for 'Gut microbiota abundance (genus Lachnospiraceae UCG008 id.11328)' on 'Crohn's disease of large intestine'

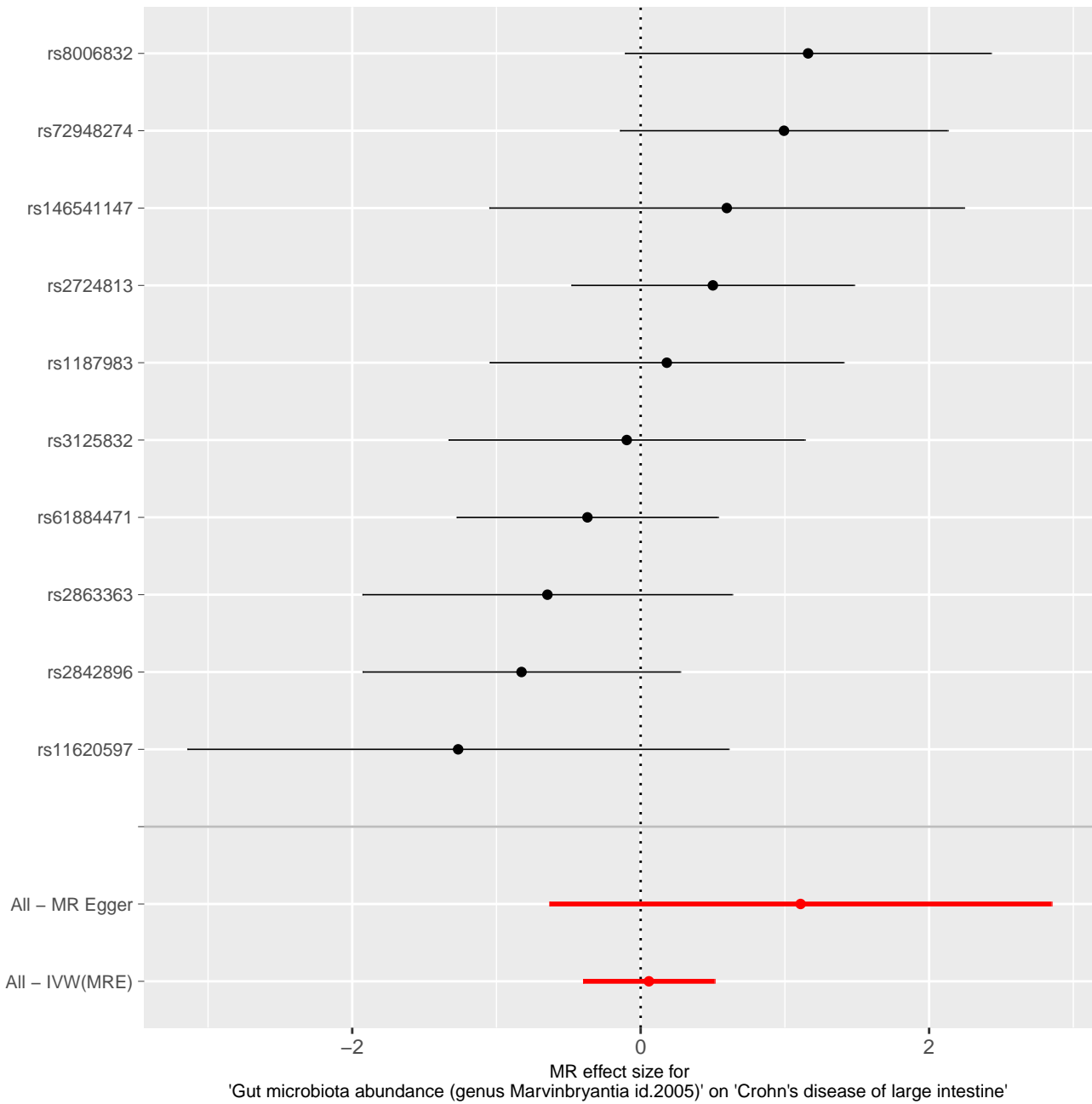
Batch 102 : Gut microbiota abundance (genus Lachnospiraceae UCG010 id.11330) on Crohn's disease of large intestine

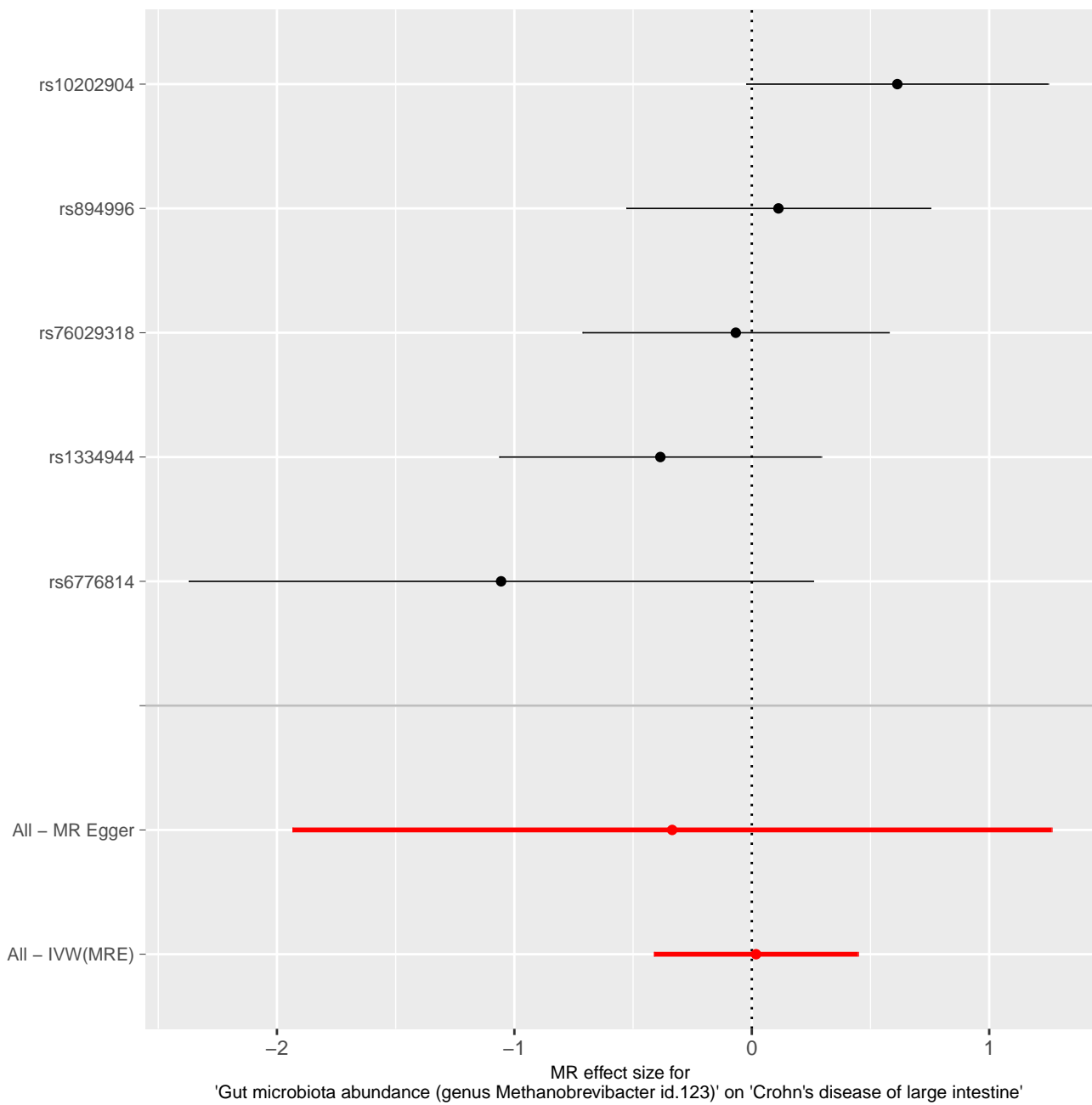




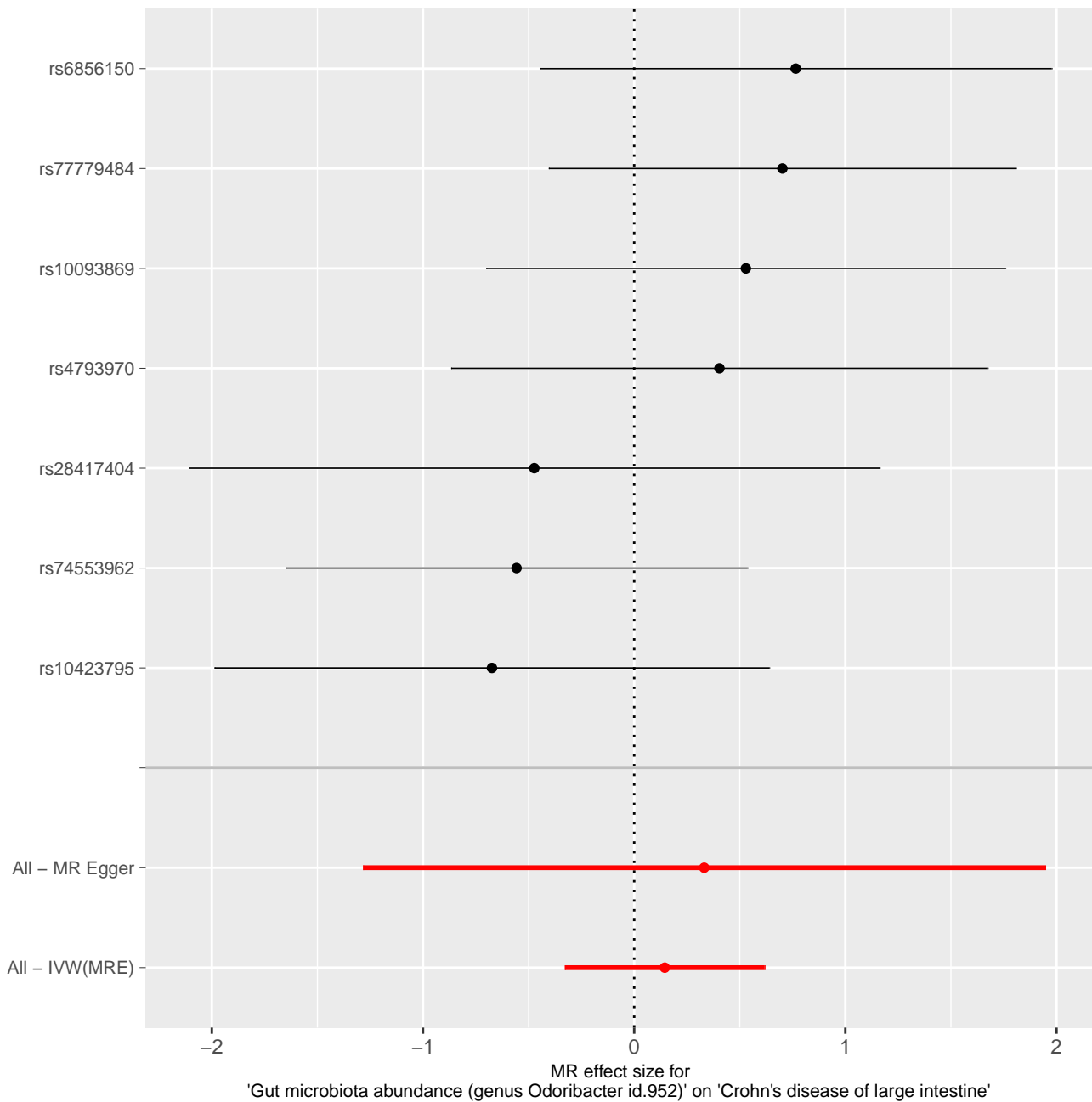
Batch 105 : Gut microbiota abundance (genus Lactococcus id.1851) on Crohn's disease of large intestine



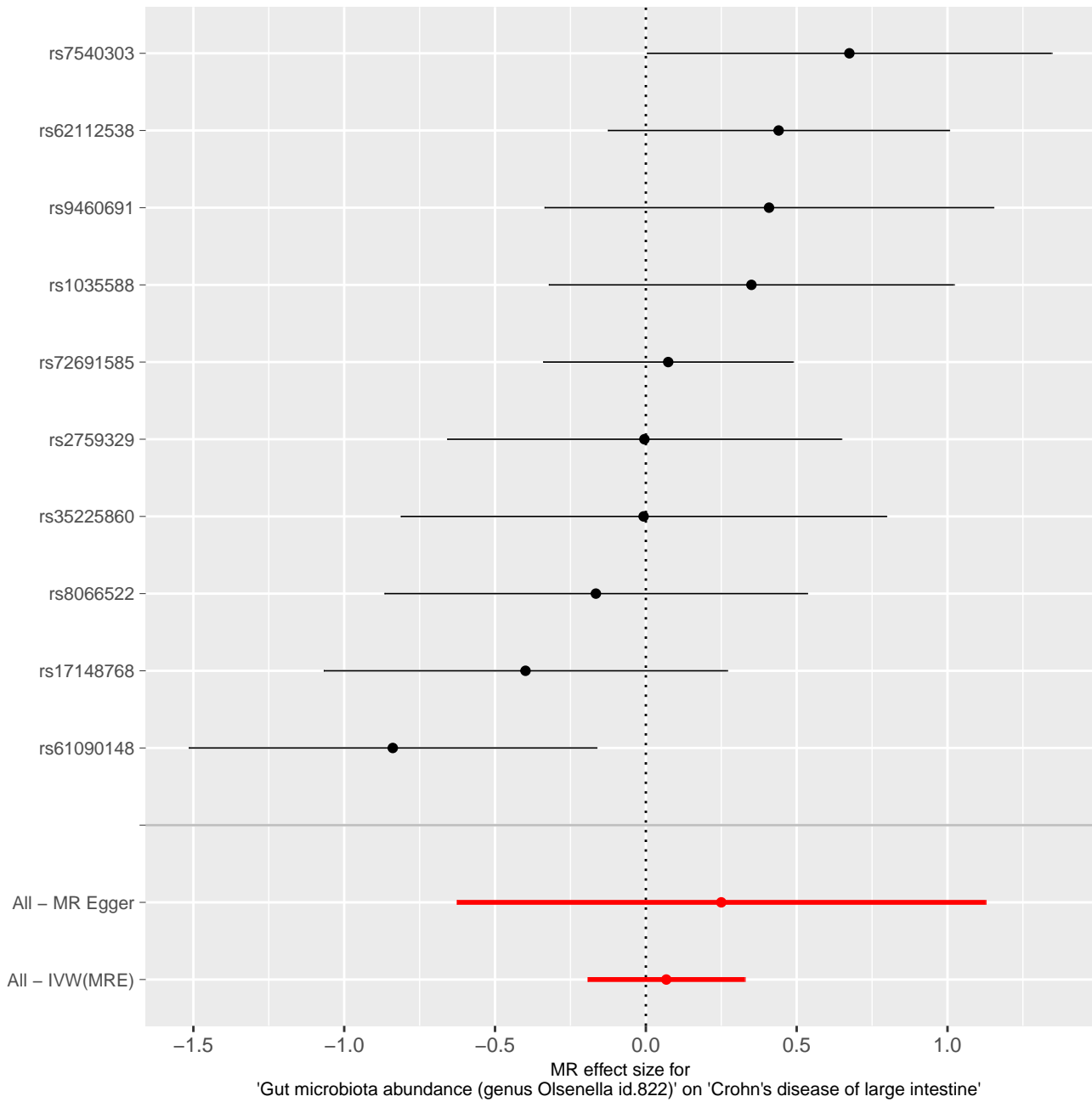




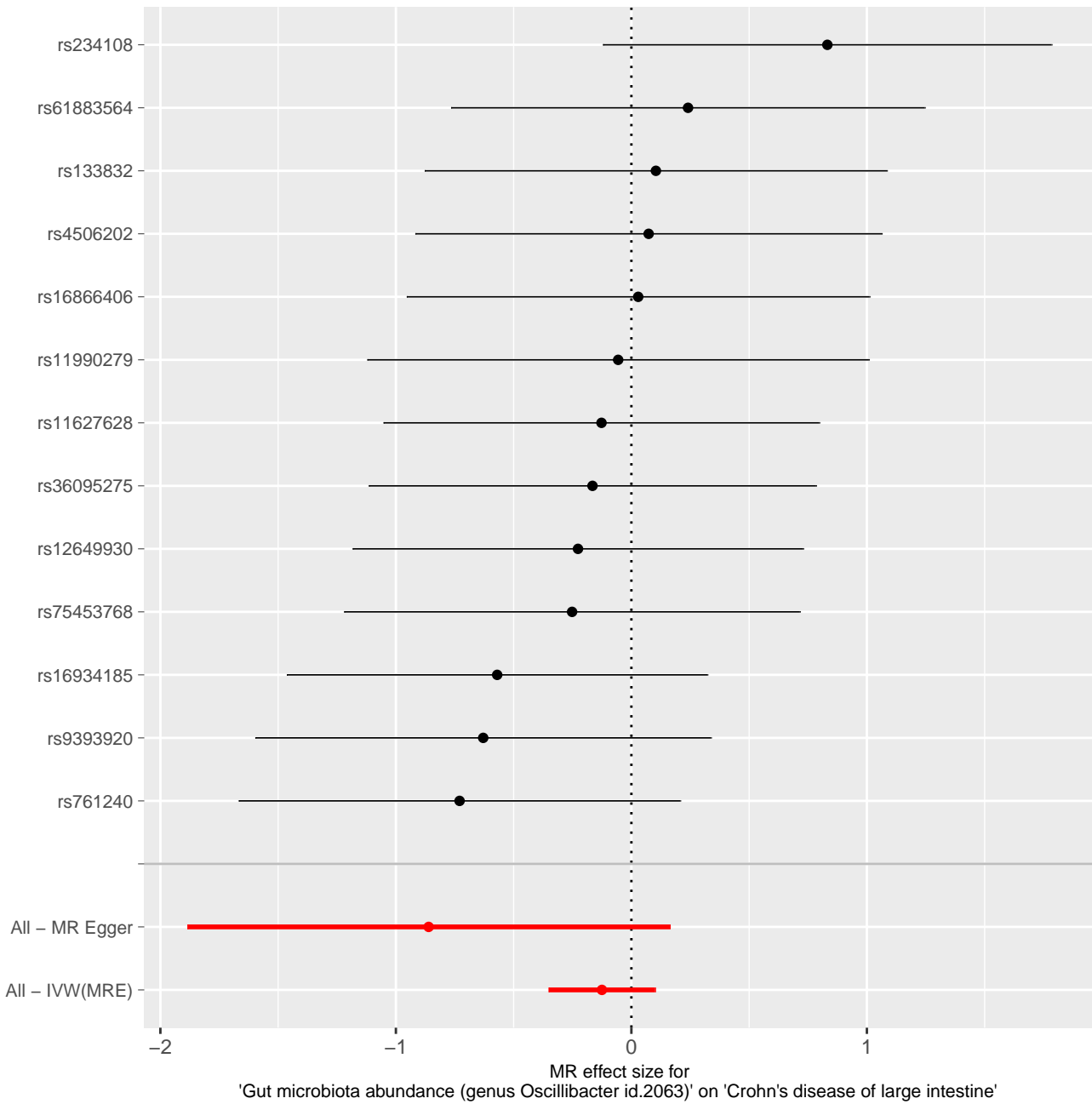
Batch 108 : Gut microbiota abundance (genus Odoribacter id.952) on Crohn's disease of large intestine



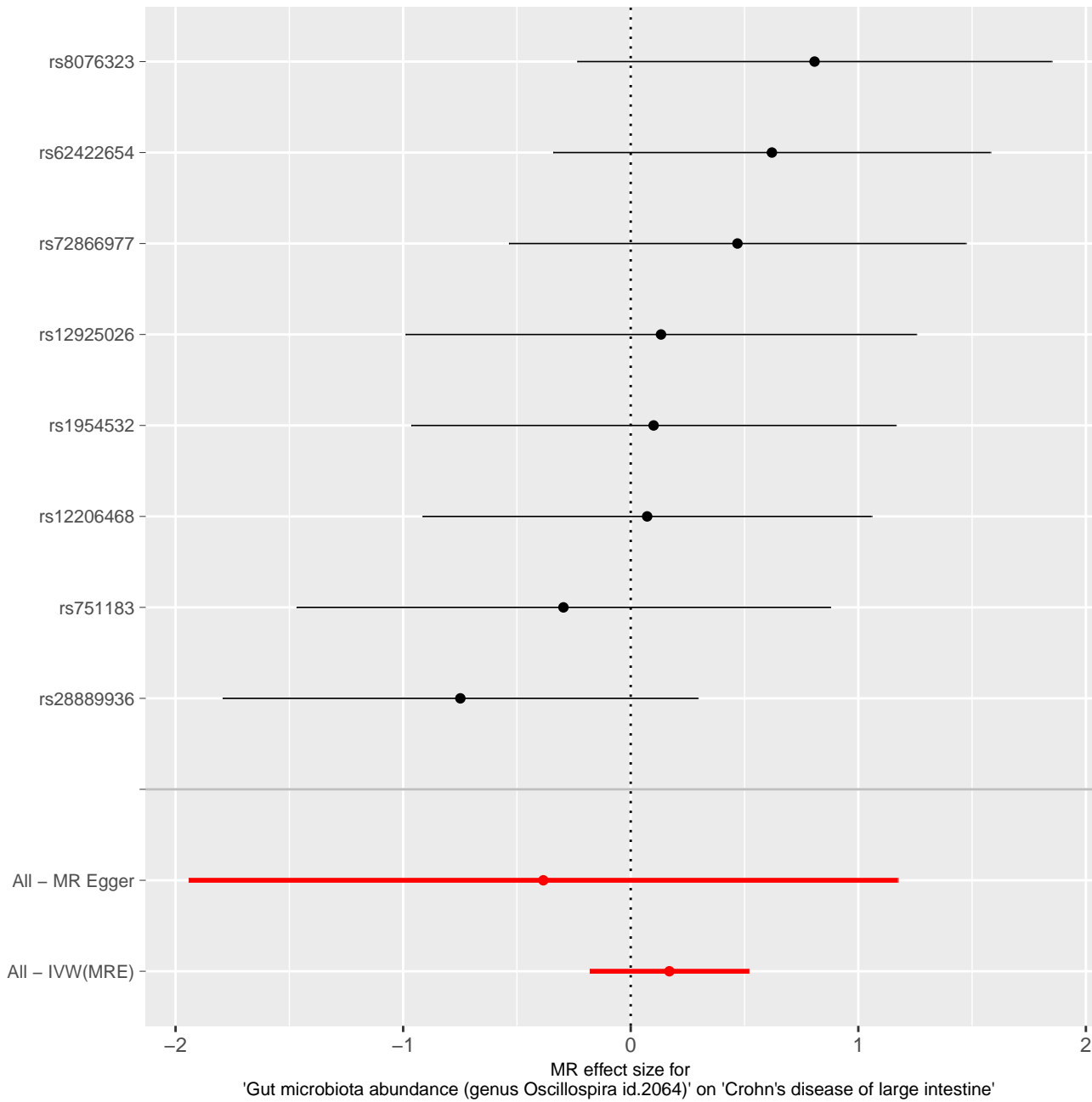
Batch 109 : Gut microbiota abundance (genus Olsenella id.822) on Crohn's disease of large intestine



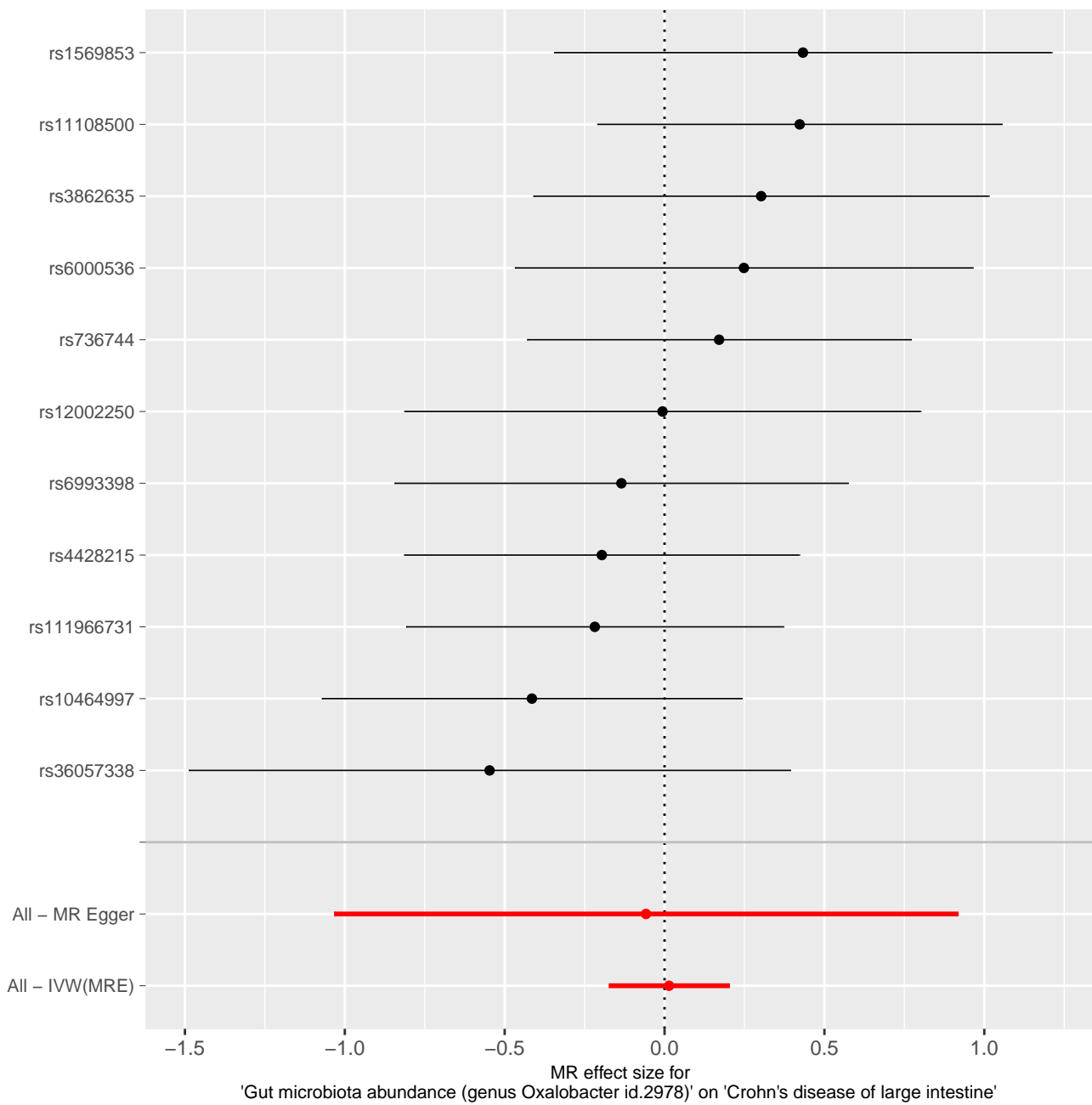
Batch 110 : Gut microbiota abundance (genus Oscillibacter id.2063) on Crohn's disease of large intestine

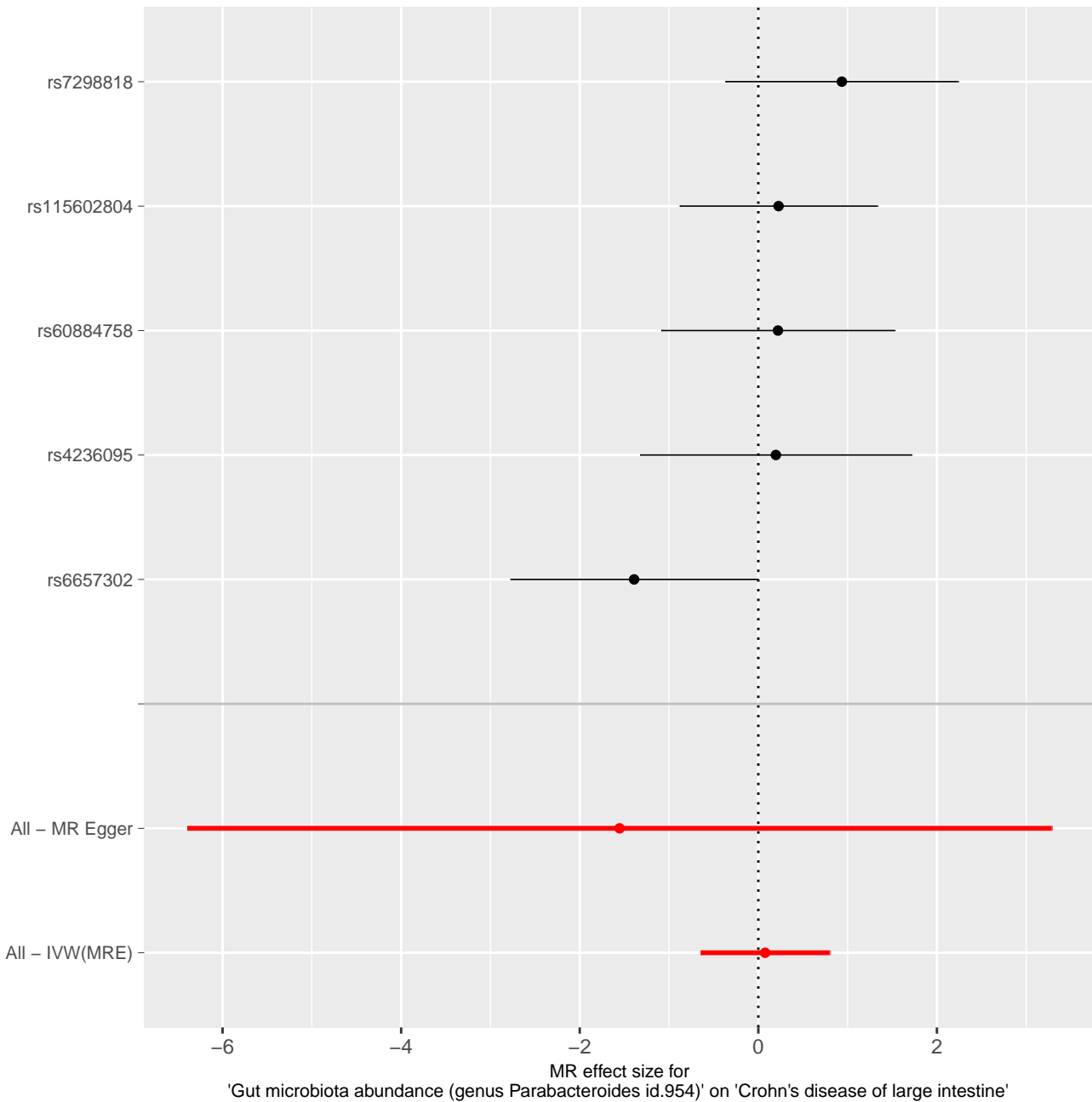


Batch 111 : Gut microbiota abundance (genus Oscillospira id.2064) on Crohn's disease of large intestine

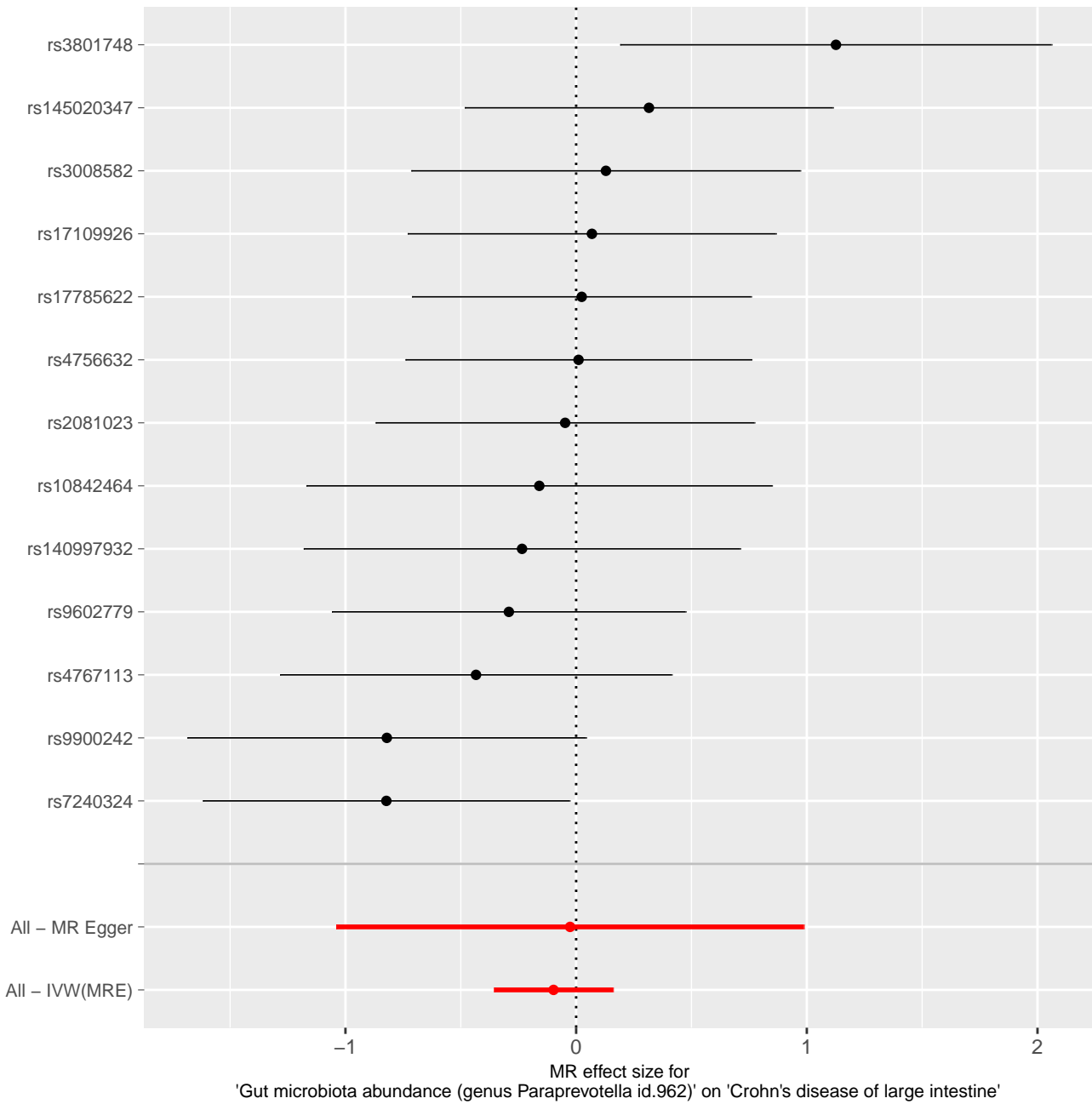


Batch 112 : Gut microbiota abundance (genus Oxalobacter id.2978) on Crohn's disease of large intestine

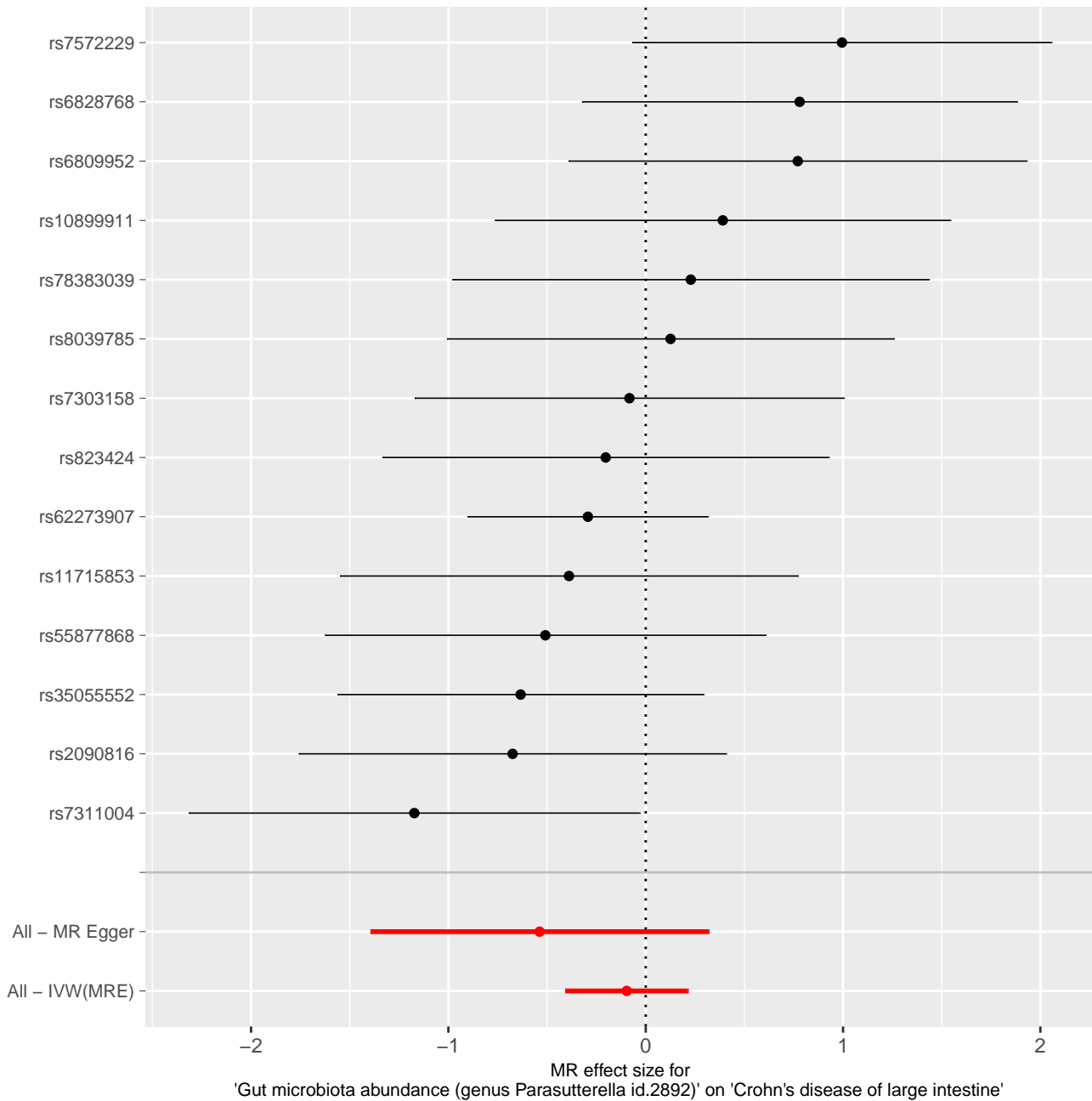


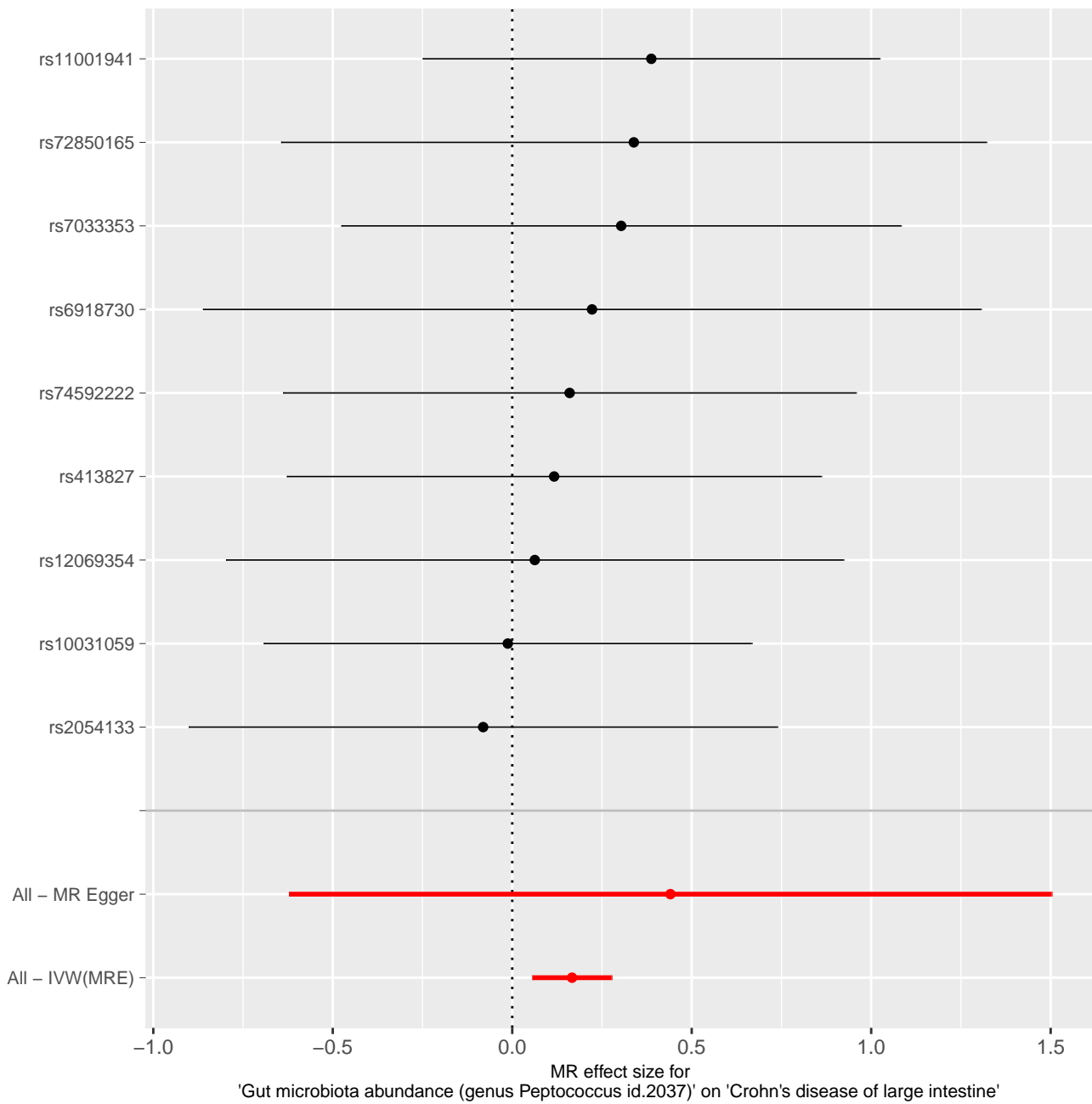


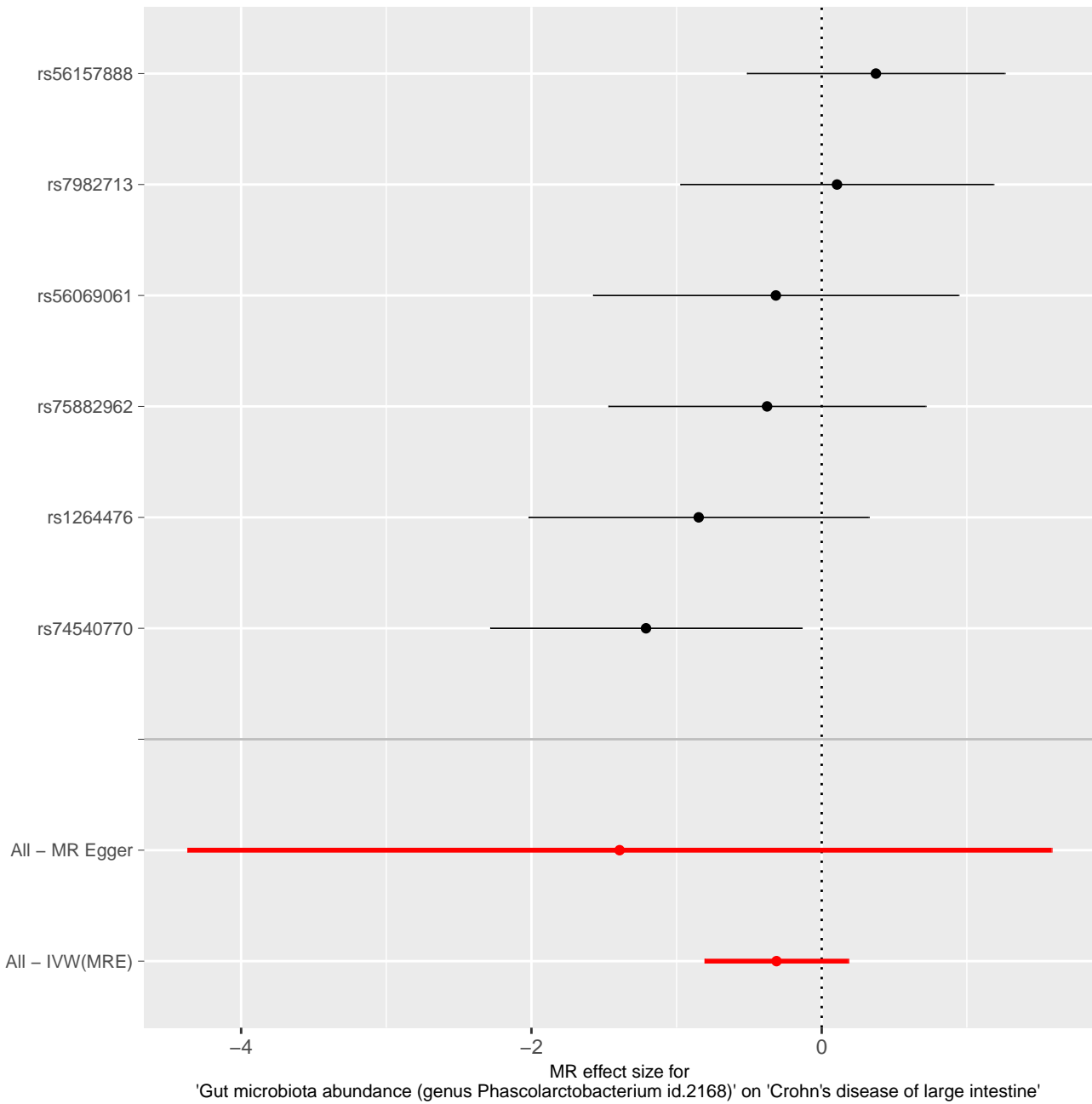
Batch 114 : Gut microbiota abundance (genus Paraprevotella id.962) on Crohn's disease of large intestine



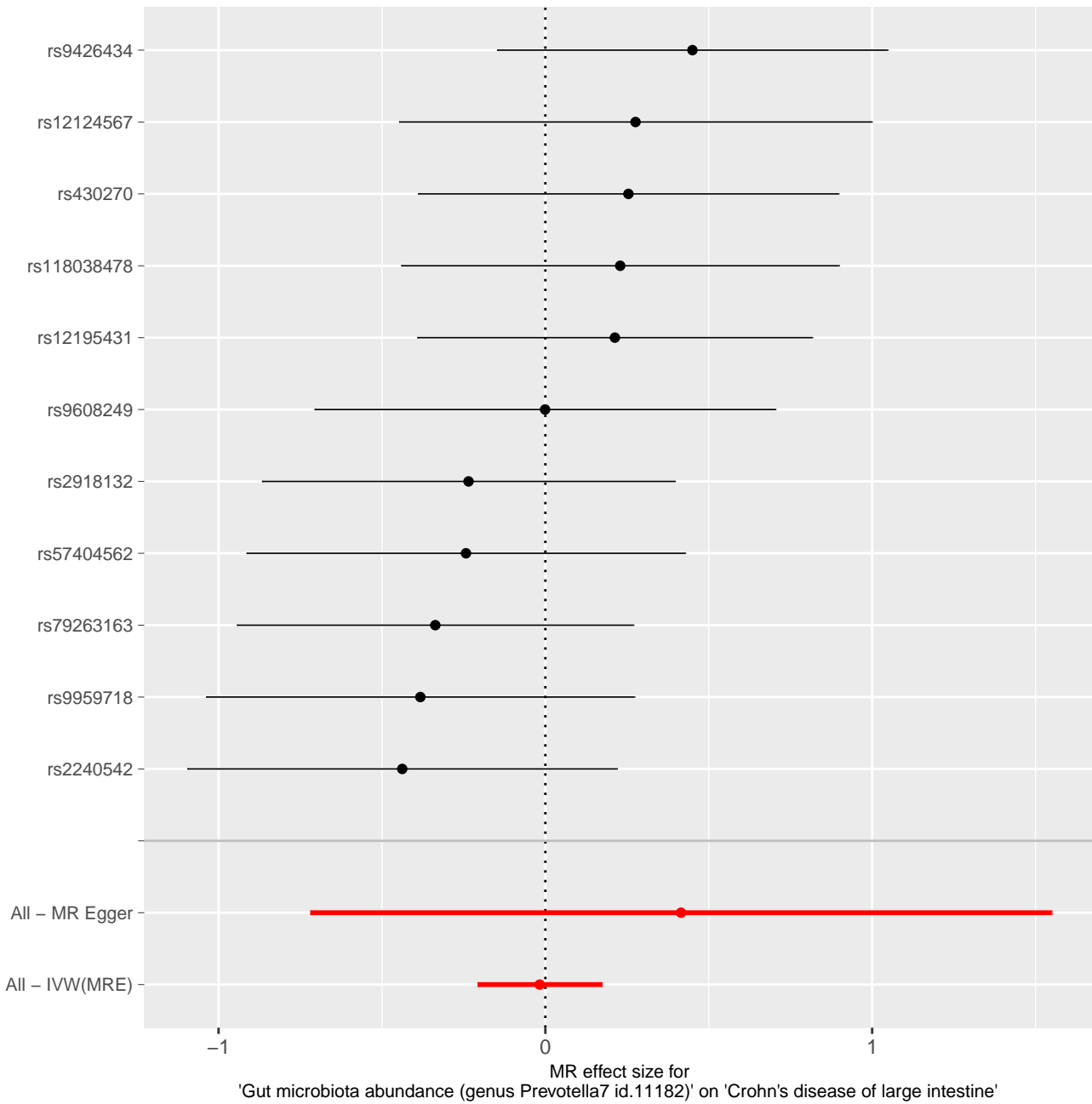
Batch 115 : Gut microbiota abundance (genus Parasutterella id.2892) on Crohn's disease of large intestine



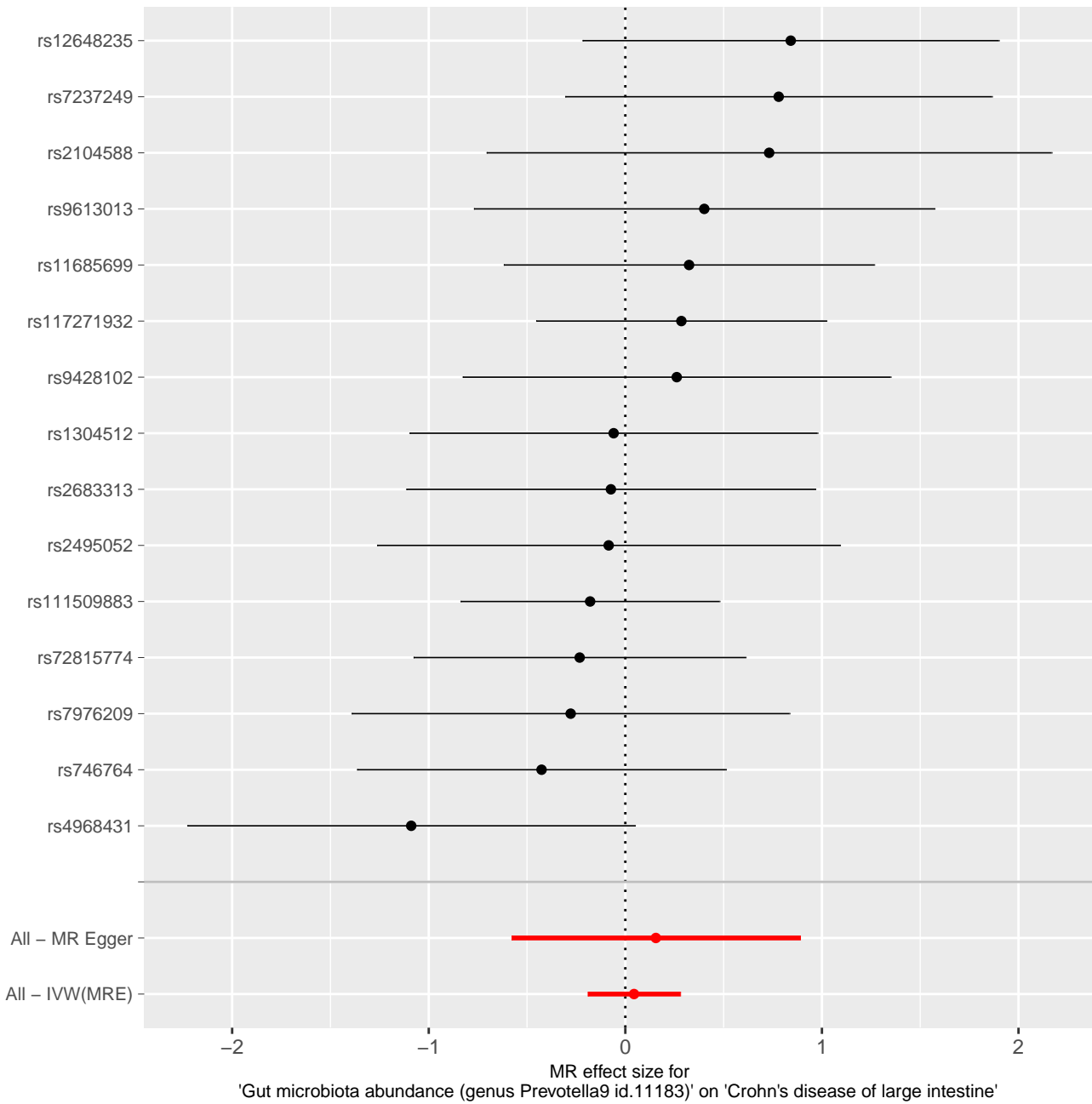


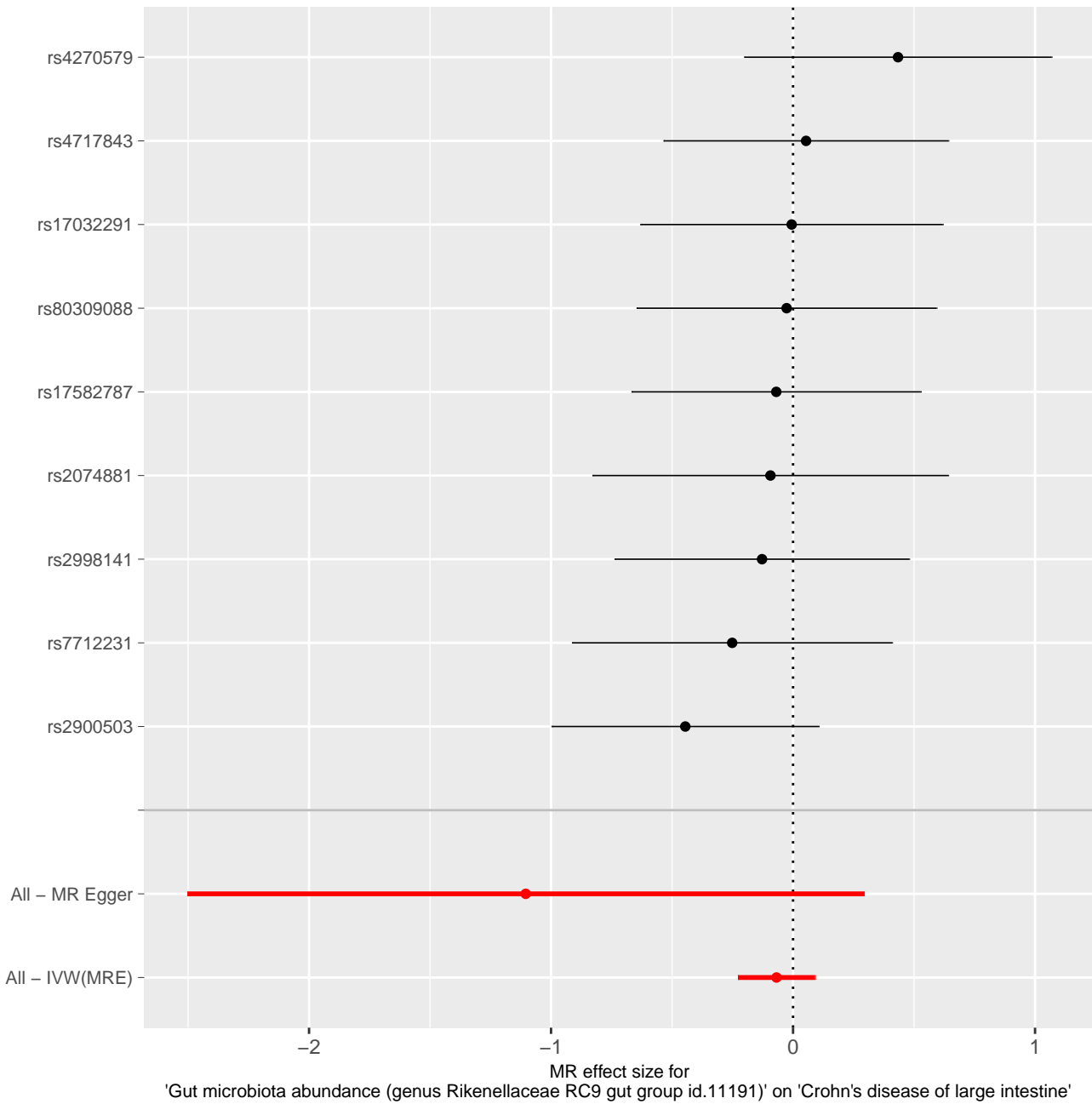


Batch 118 : Gut microbiota abundance (genus Prevotella7 id.11182) on Crohn's disease of large intestine

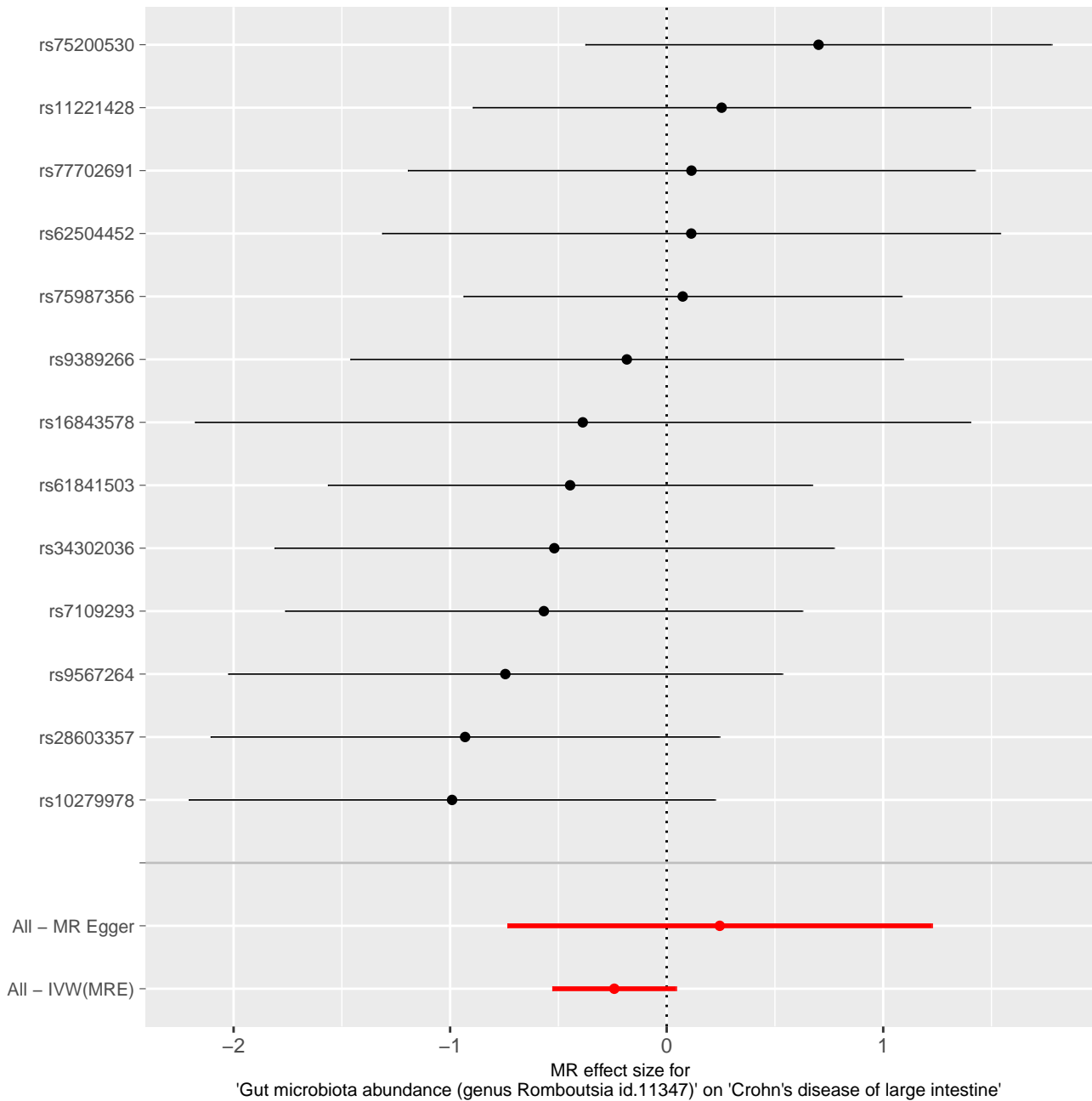


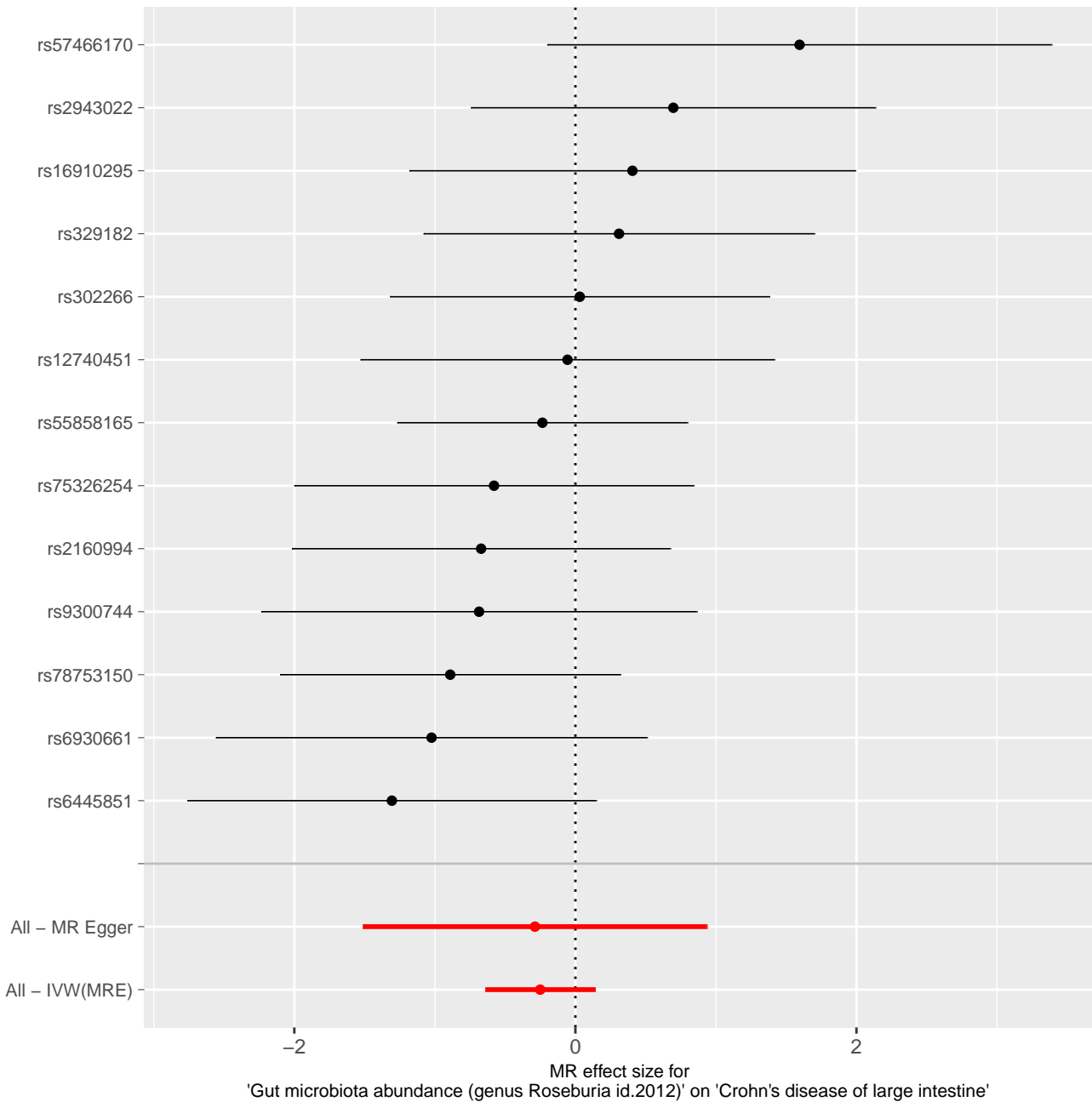
Batch 119 : Gut microbiota abundance (genus Prevotella9 id.11183) on Crohn's disease of large intestine



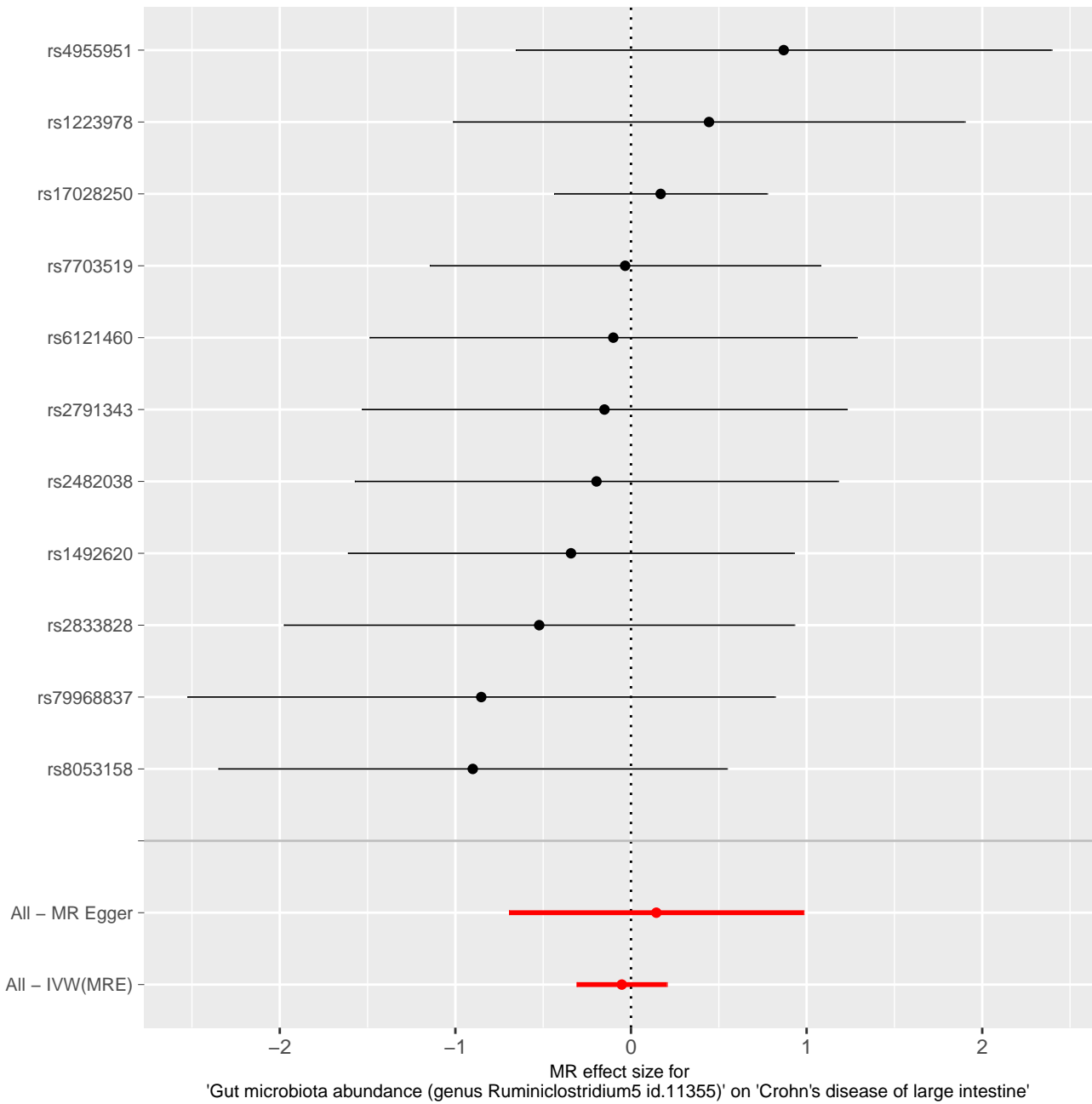


Batch 121 : Gut microbiota abundance (genus Romboutsia id.11347) on Crohn's disease of large intestine

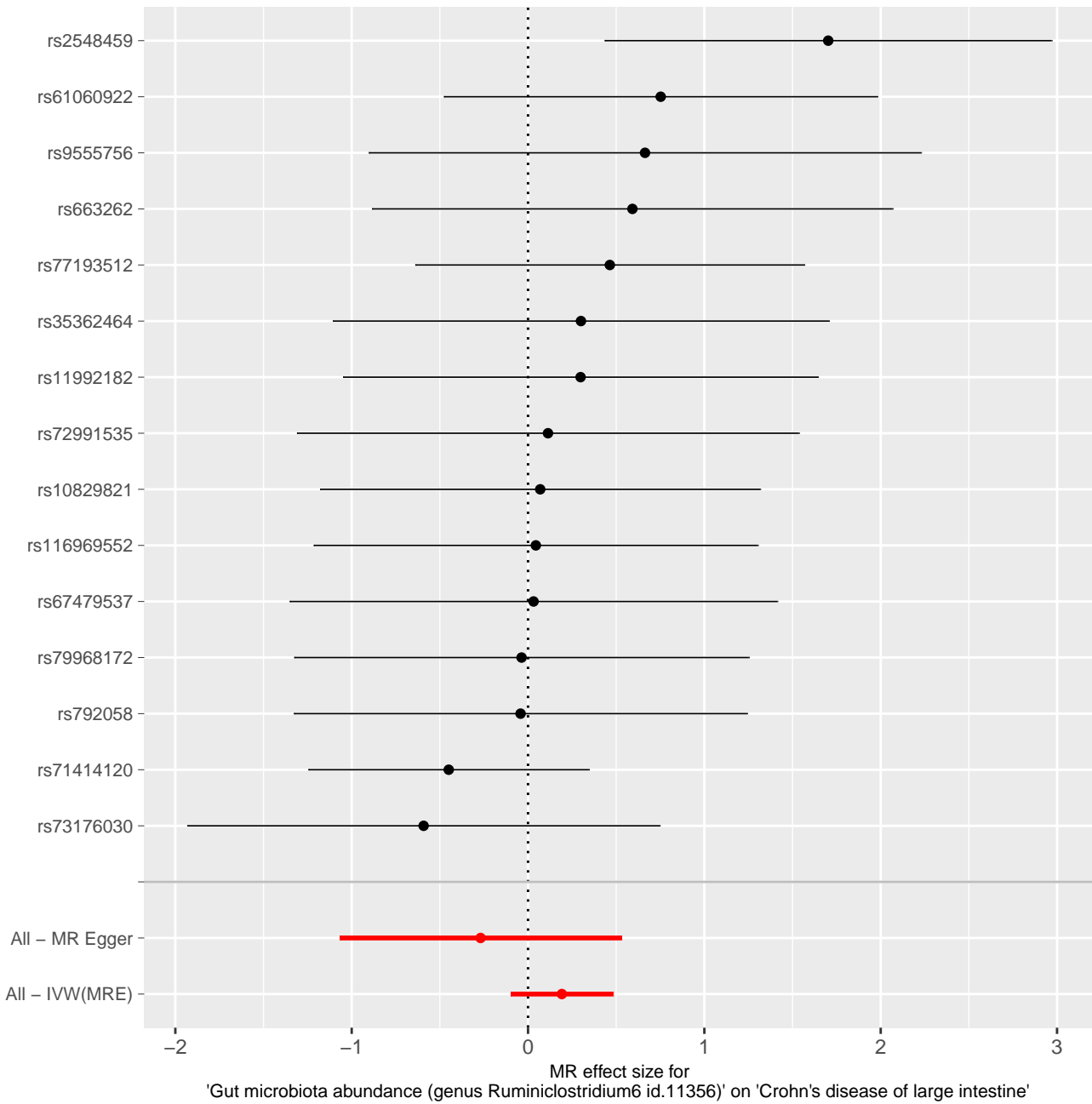




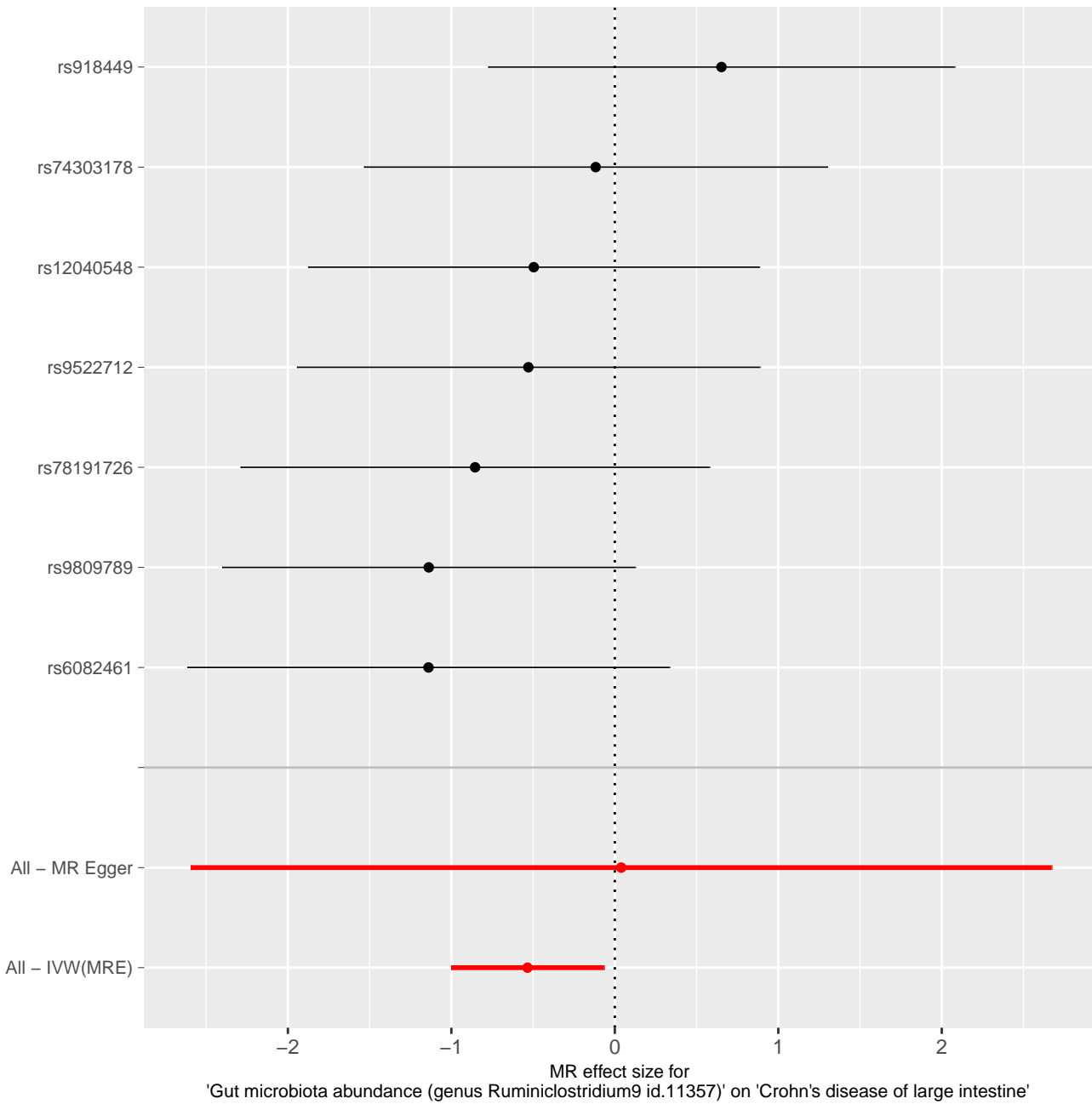
Batch 123 : Gut microbiota abundance (genus Ruminiclostridium5 id.11355) on Crohn's disease of large intestine



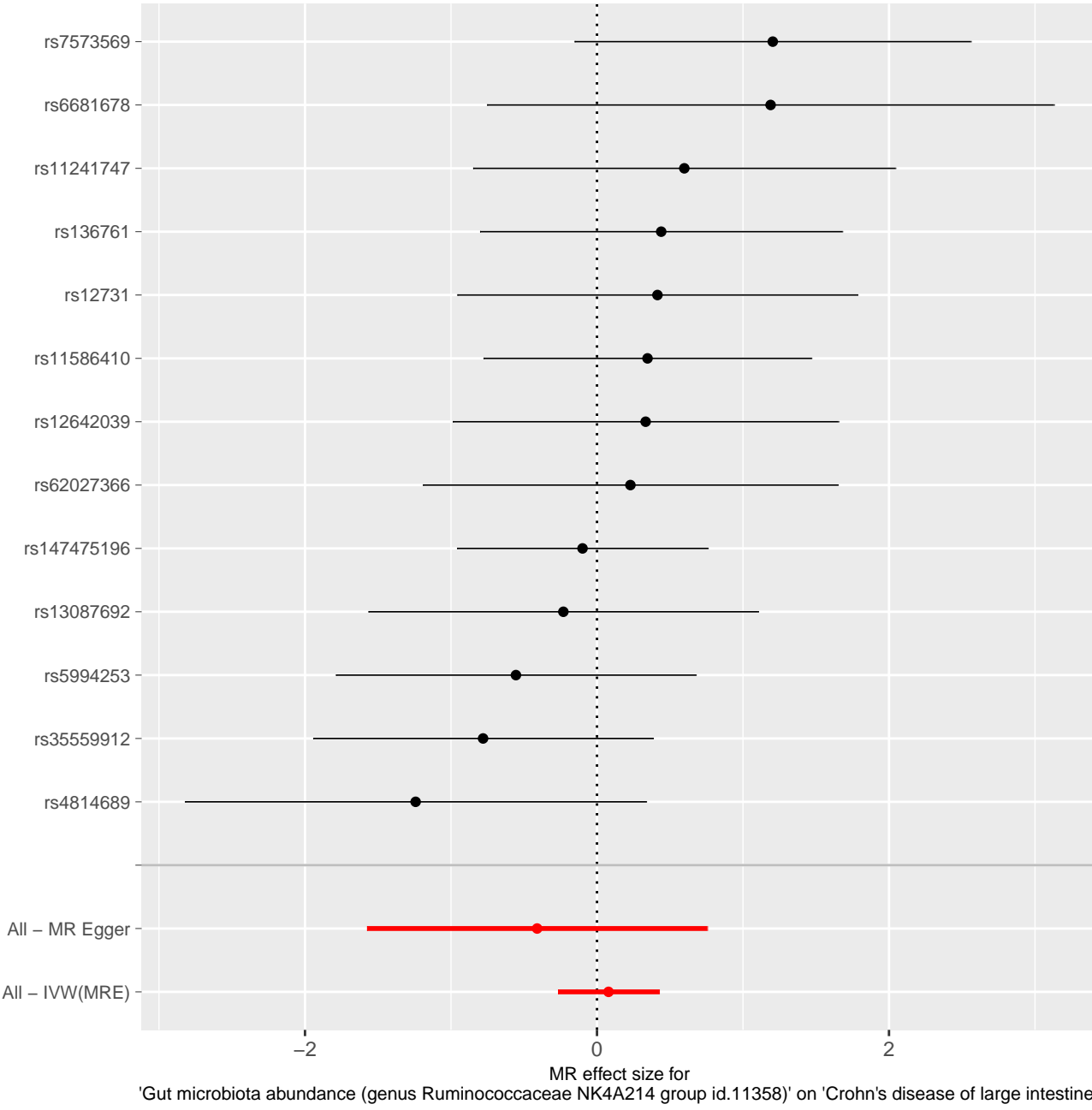
Batch 124 : Gut microbiota abundance (genus Ruminiclostridium6 id.11356) on Crohn's disease of large intestine



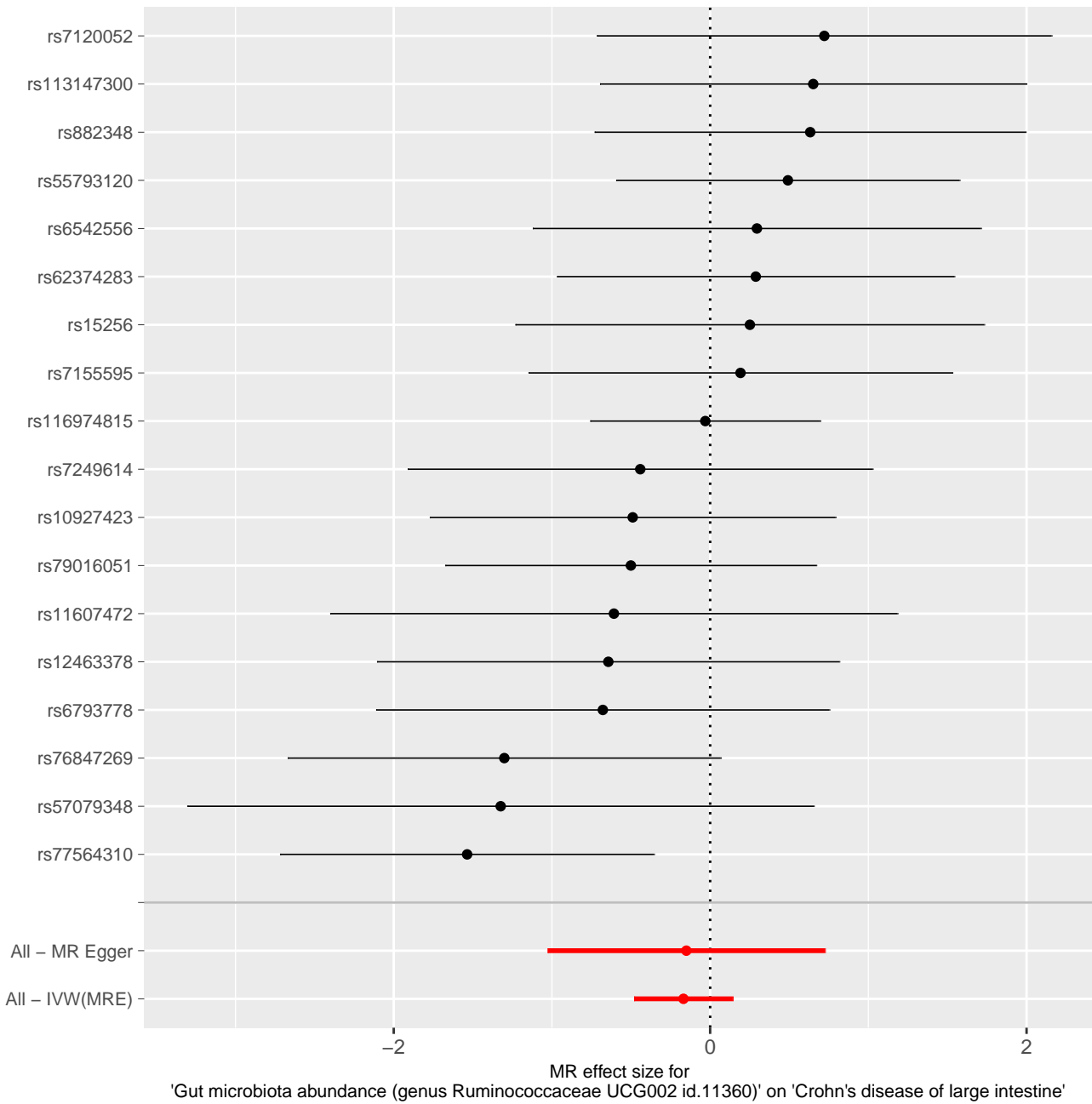
Batch 125 : Gut microbiota abundance (genus Ruminiclostridium9 id.11357) on Crohn's disease of large intestine

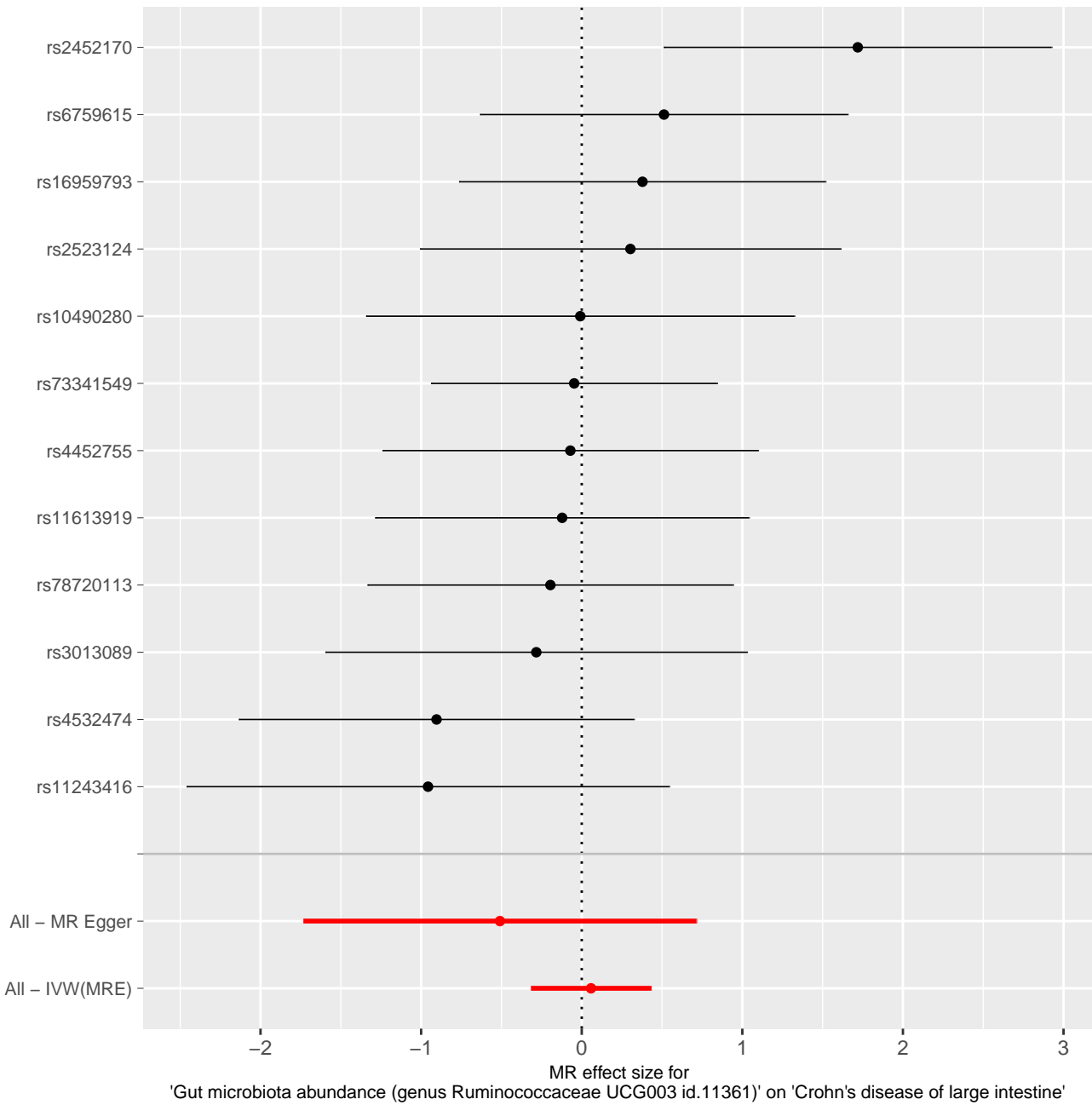


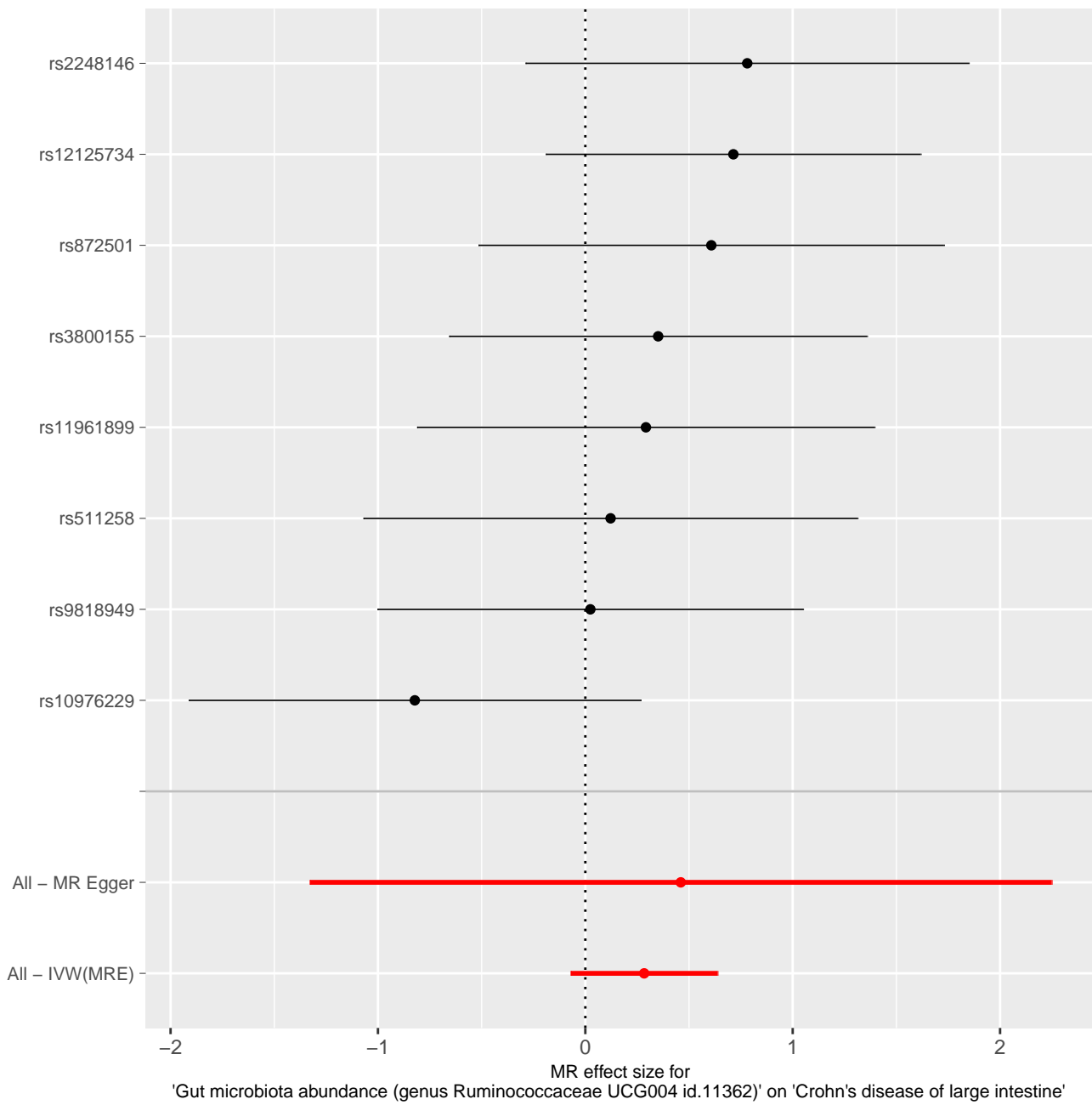
Batch 126 : Gut microbiota abundance (genus Ruminococcaceae NK4A214 group id.11358) on Crohn's disease of large intestine

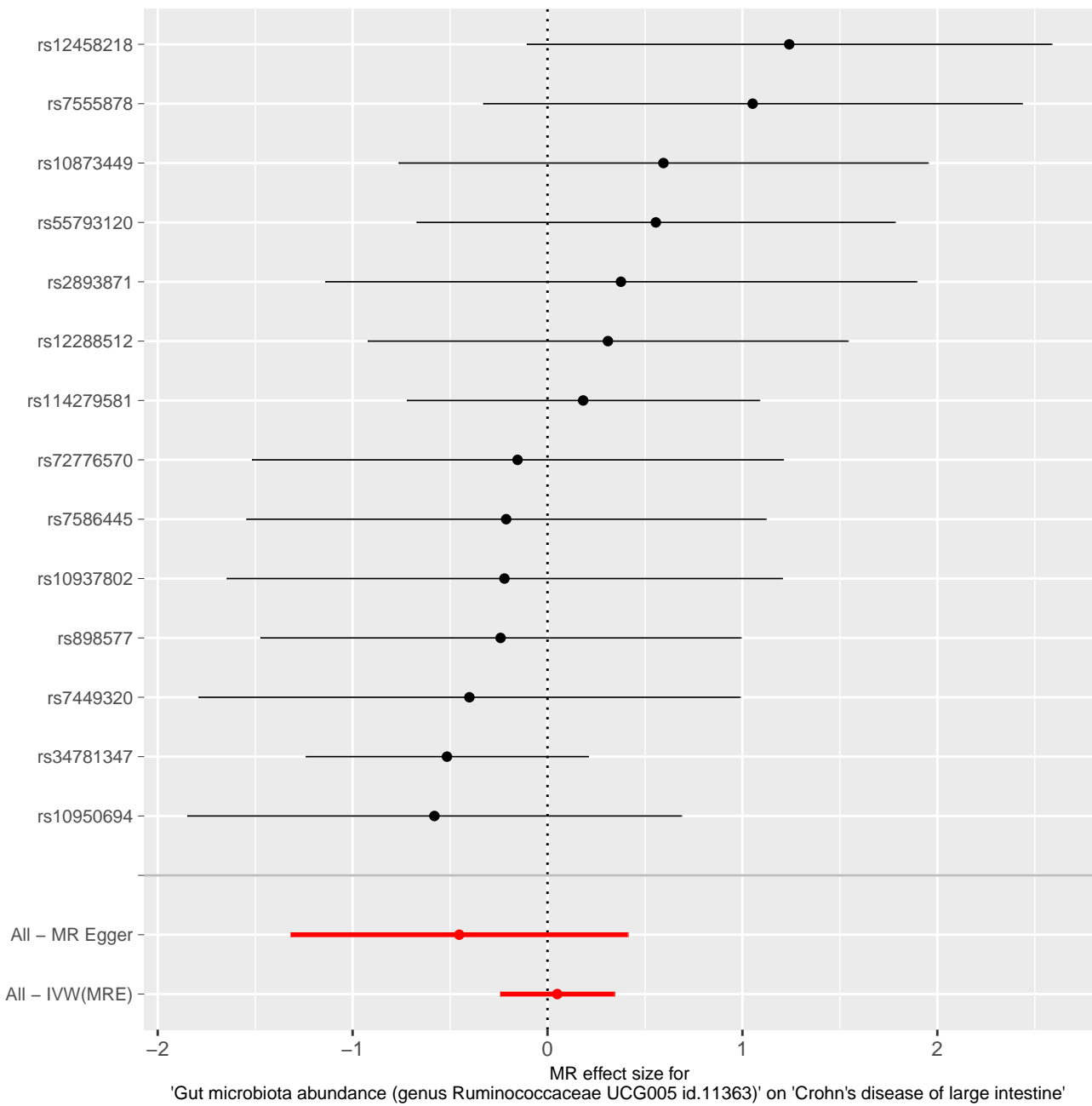


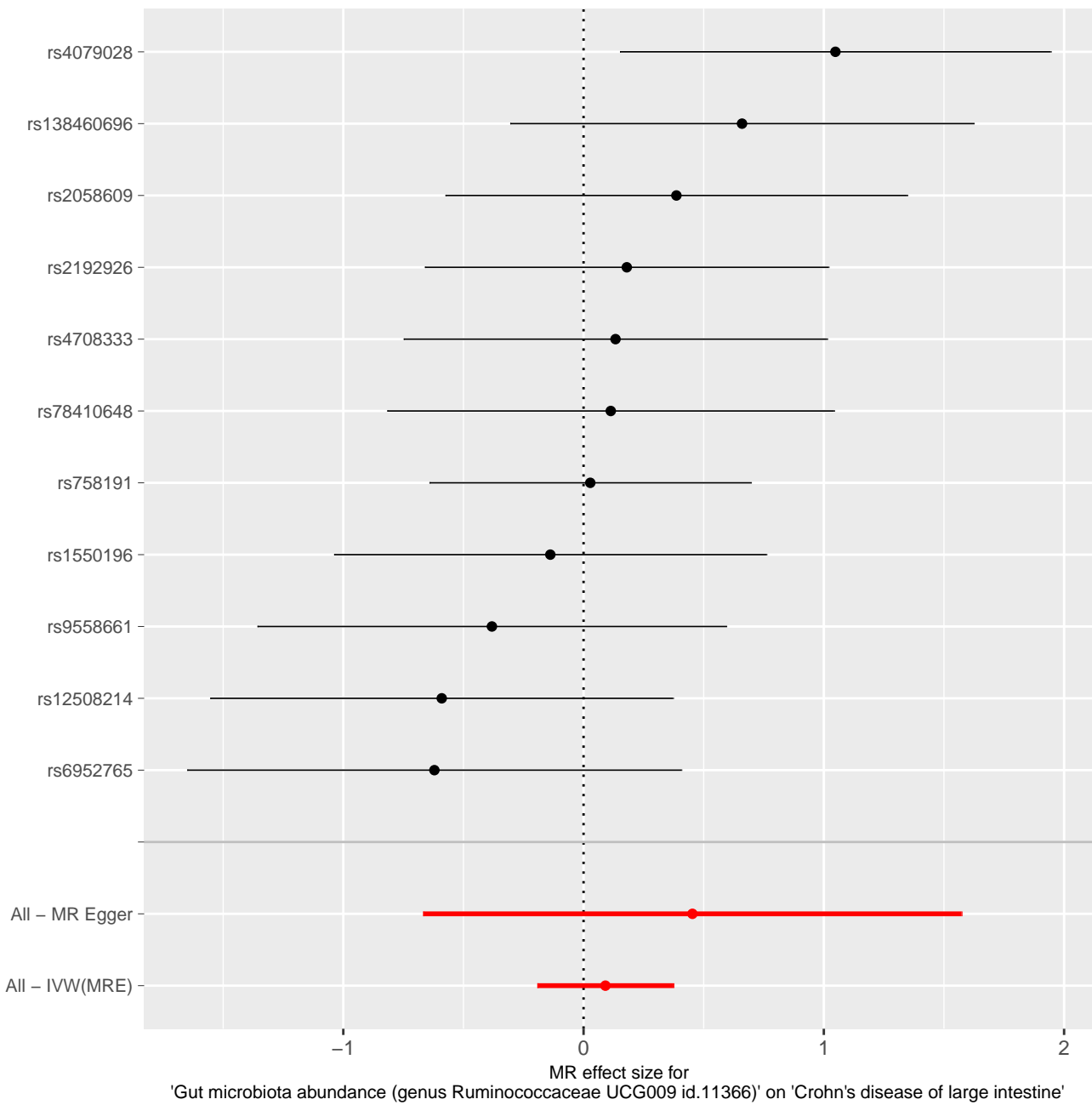
Batch 127 : Gut microbiota abundance (genus Ruminococcaceae UCG002 id.11360) on Crohn's disease of large intestine

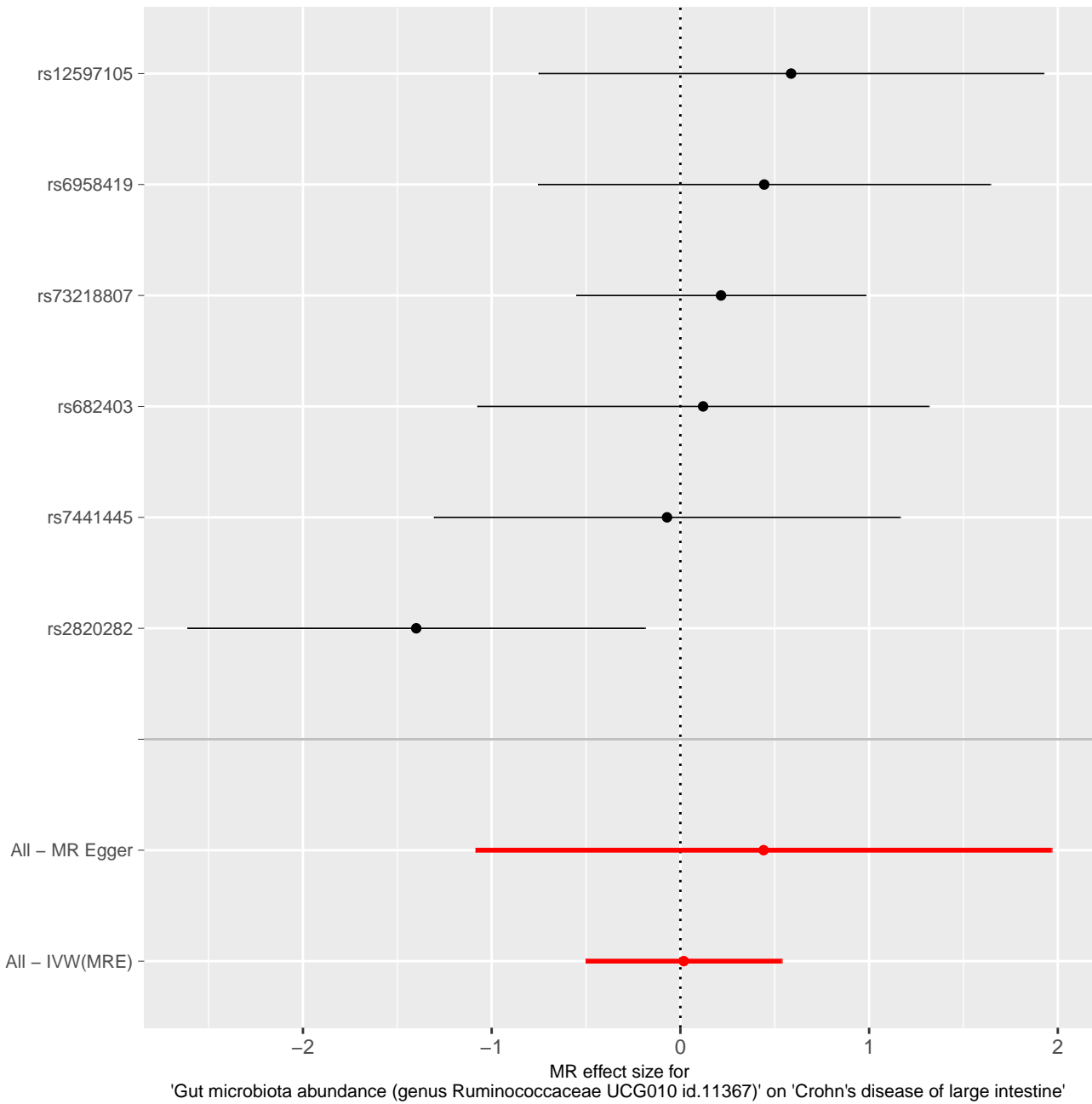


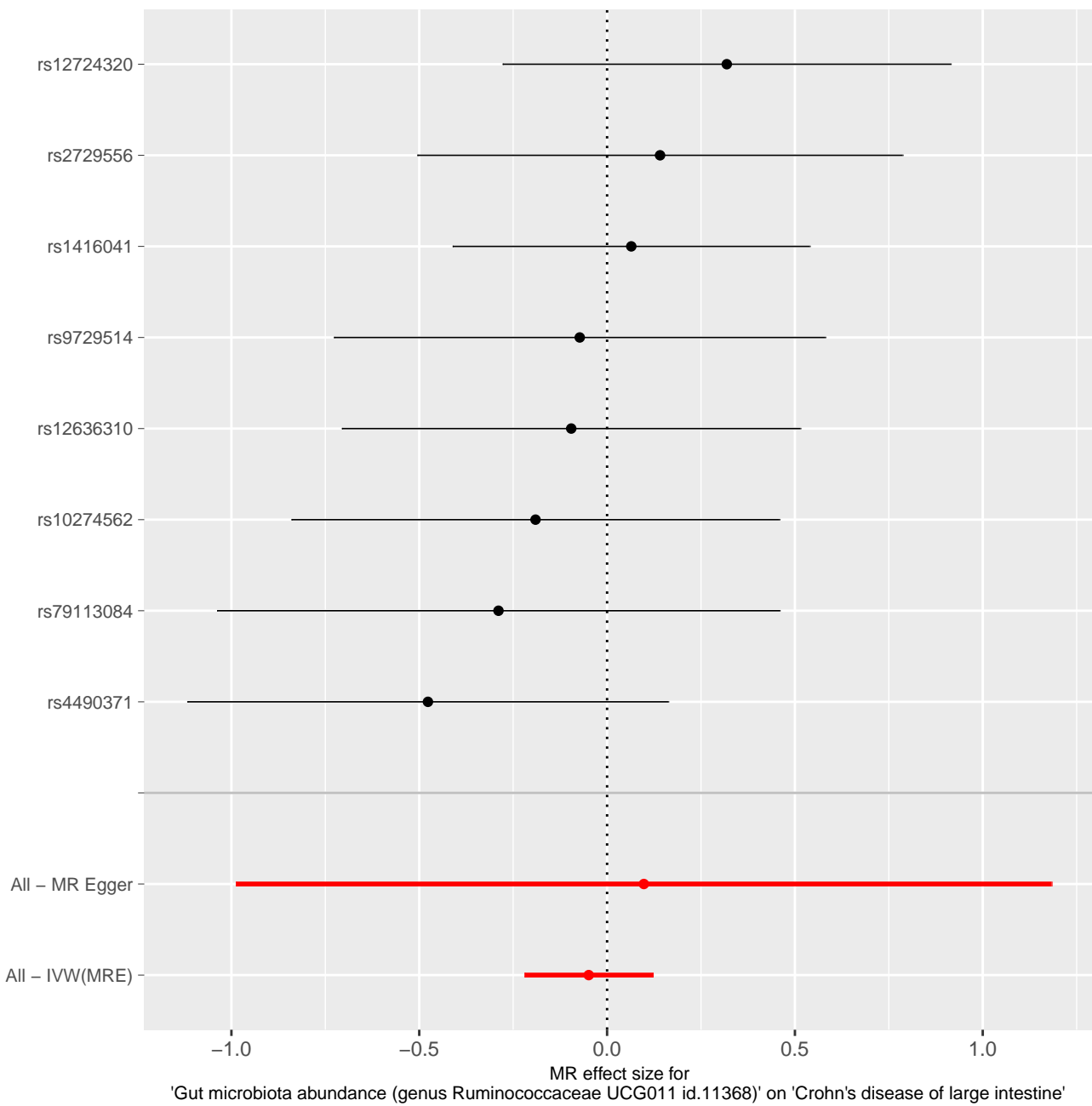




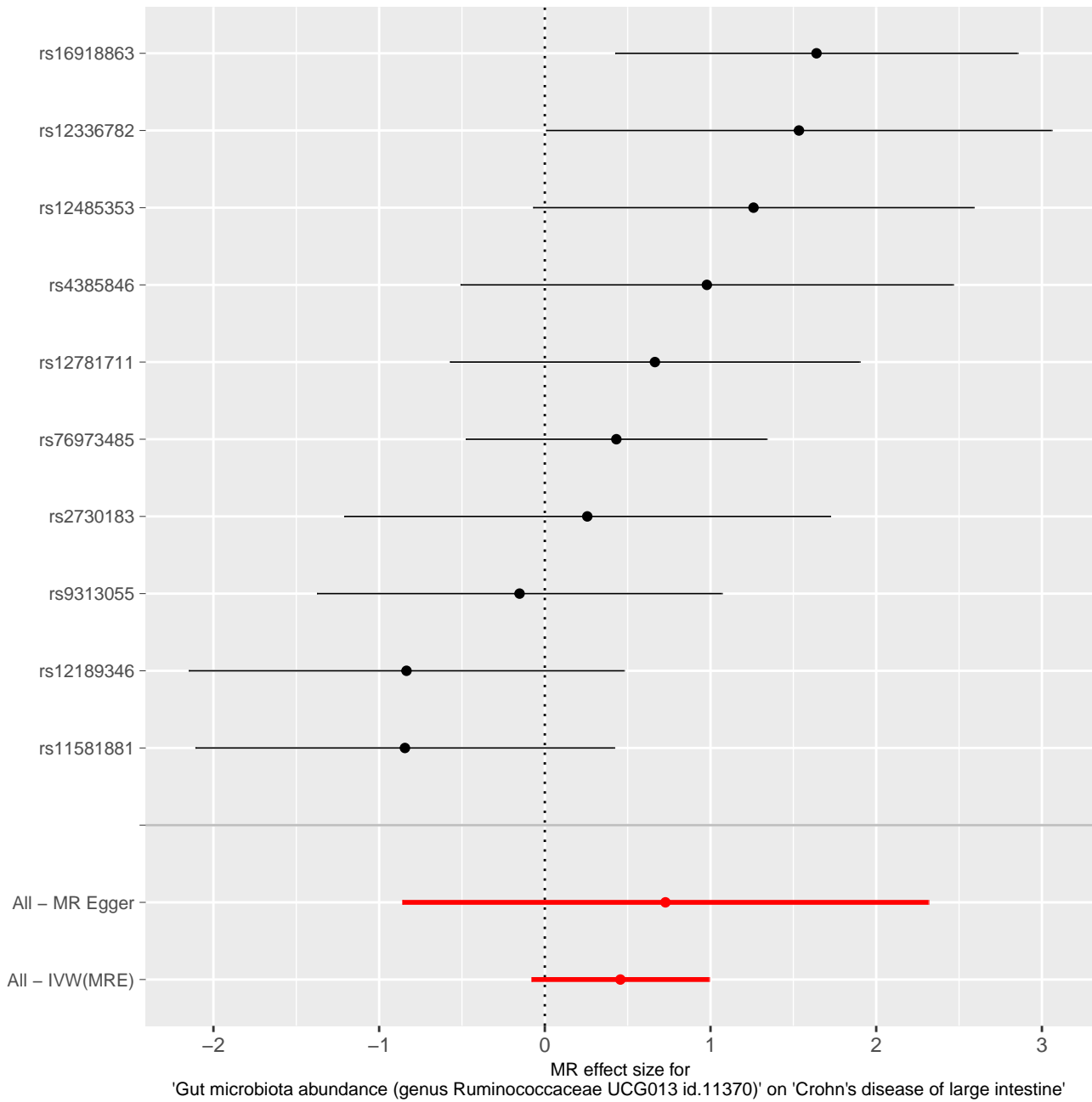


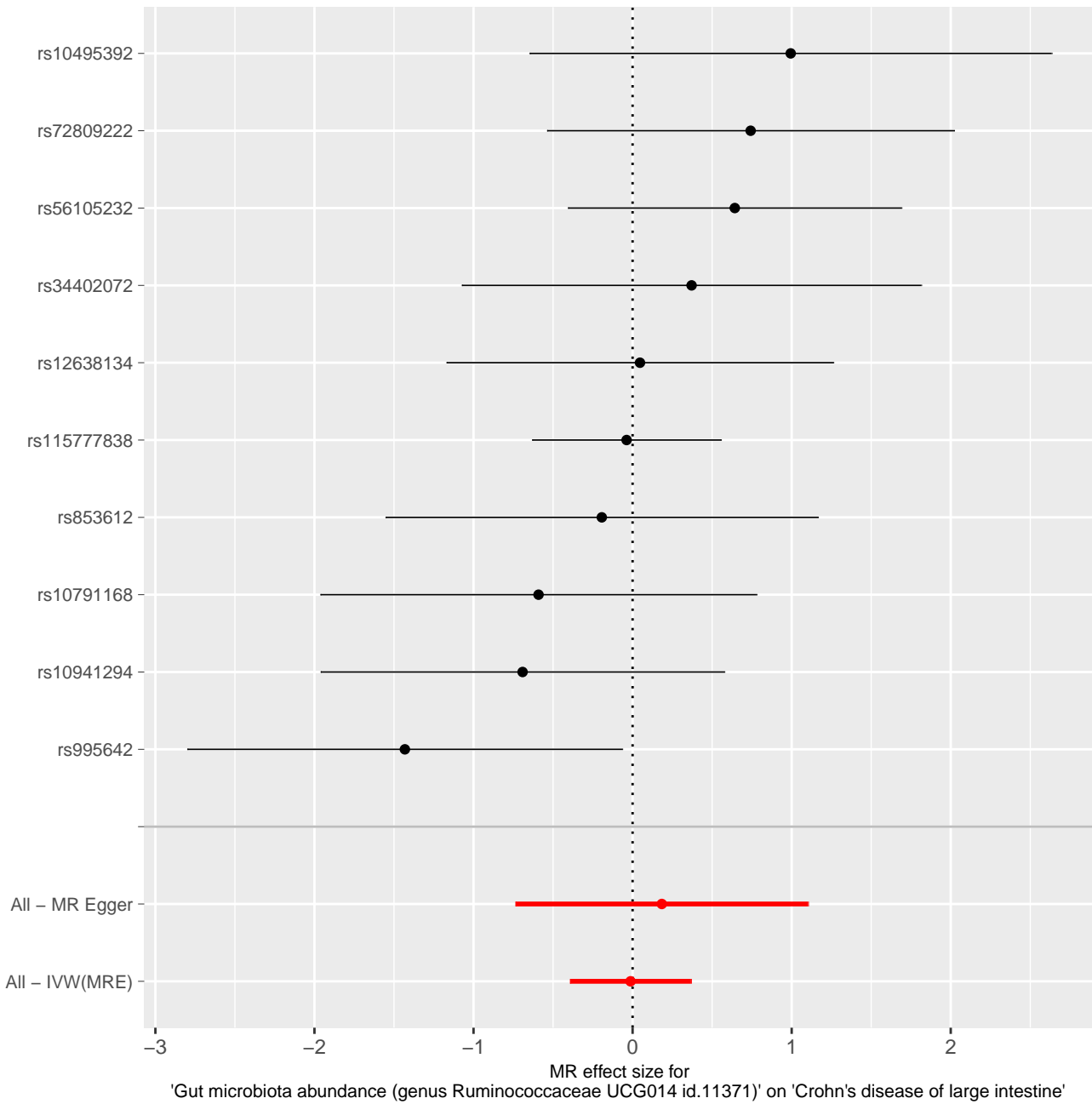




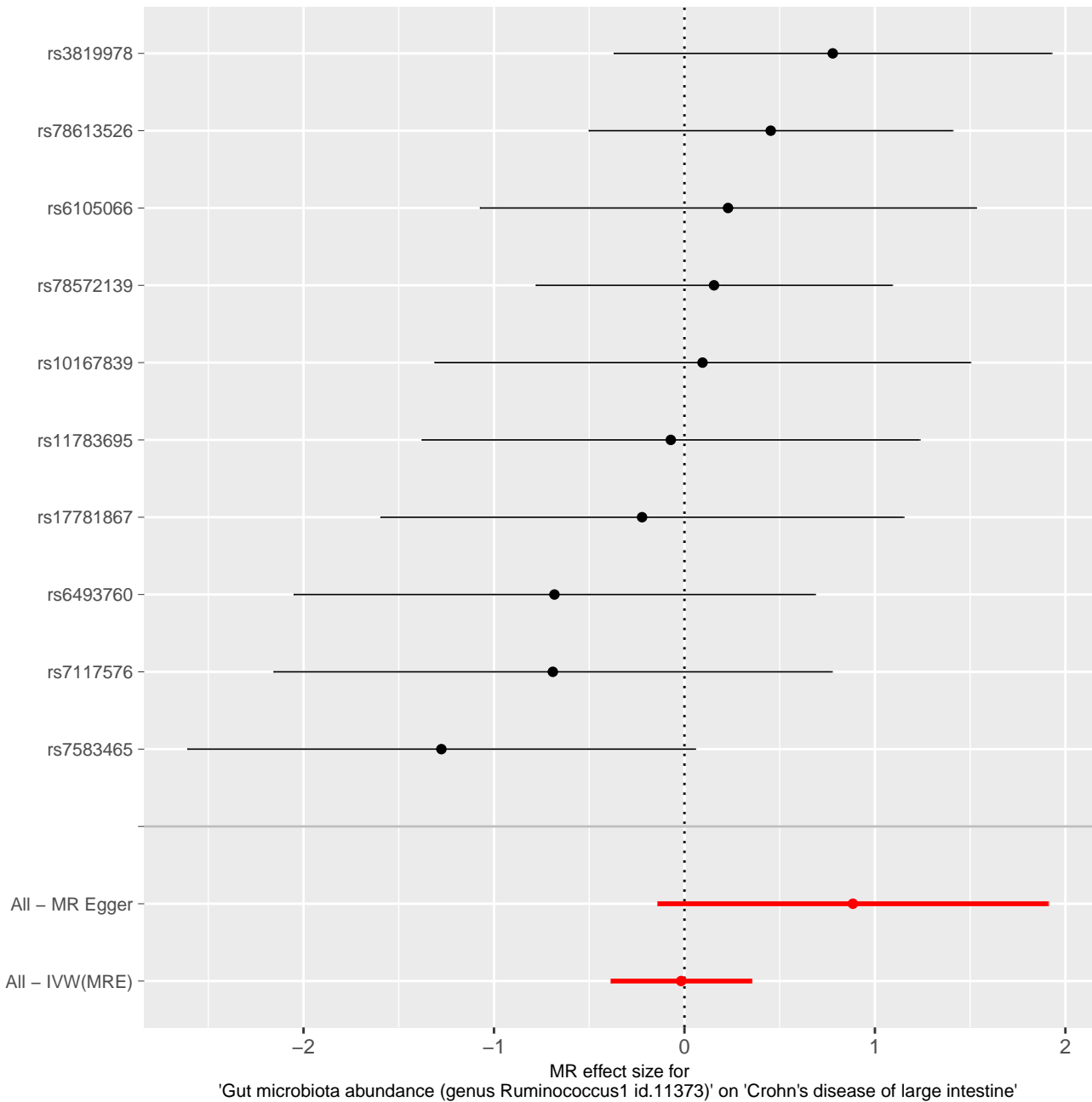


Batch 134 : Gut microbiota abundance (genus Ruminococcaceae UCG013 id.11370) on Crohn's disease of large intestine

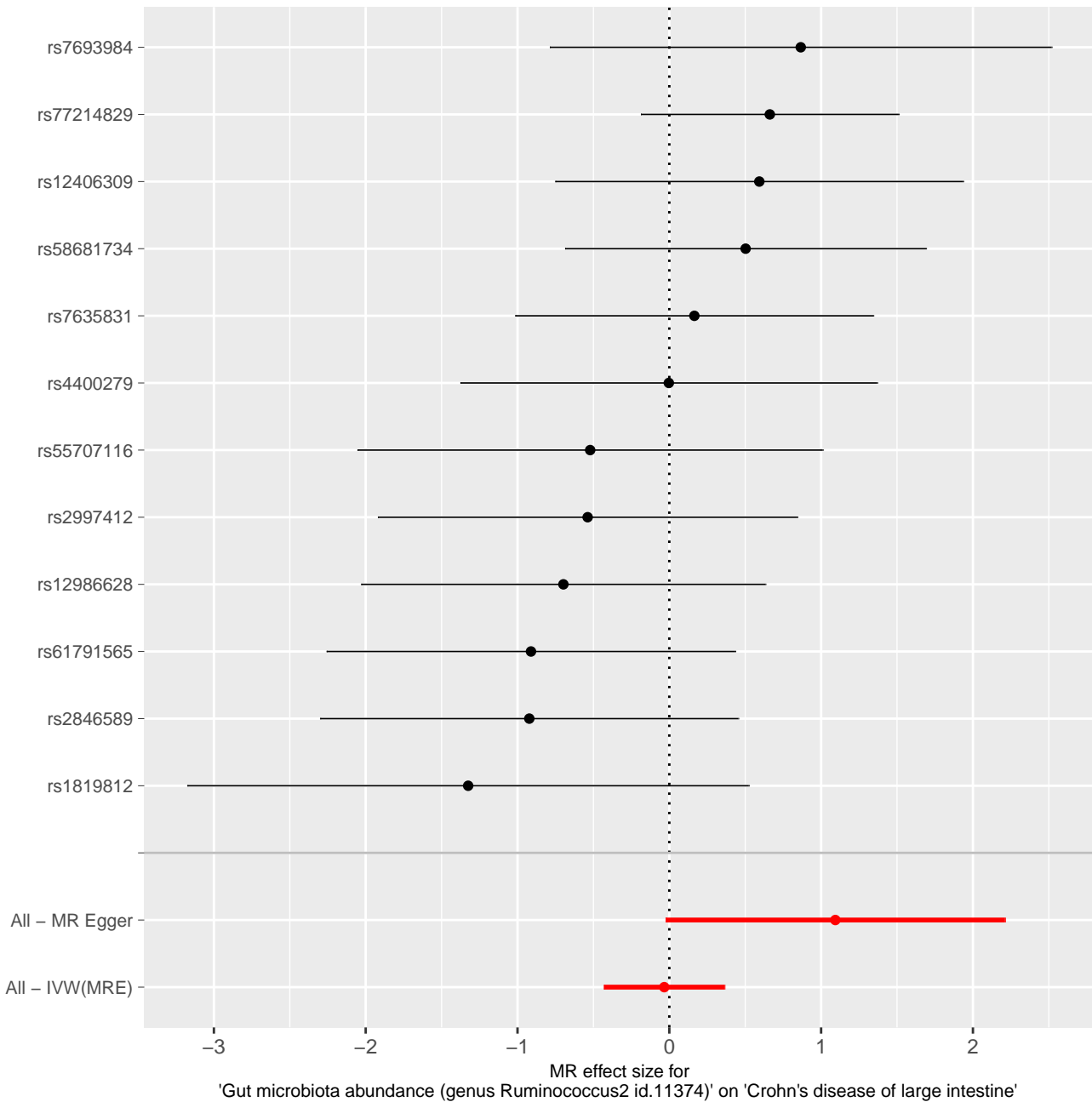


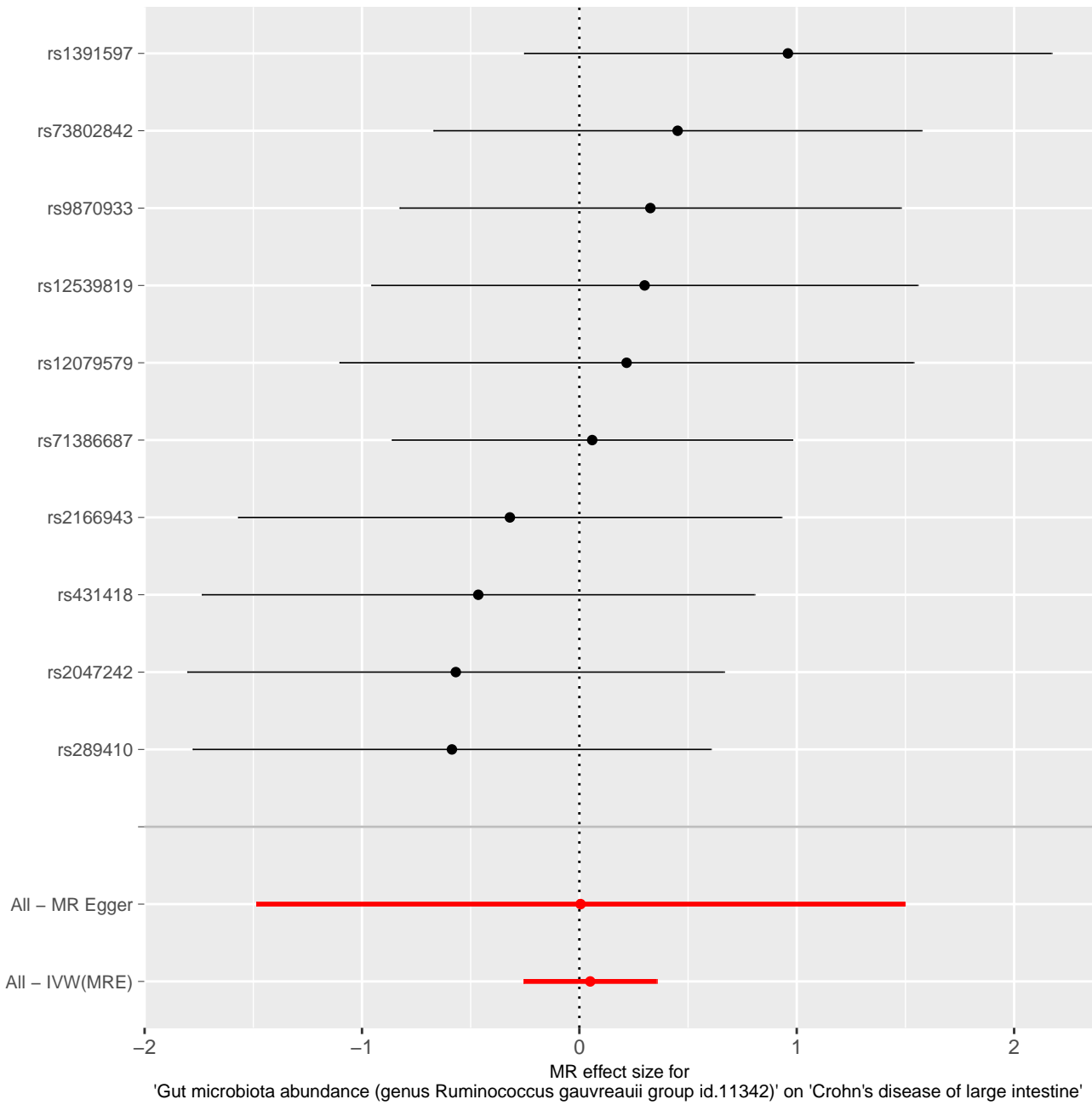


Batch 136 : Gut microbiota abundance (genus Ruminococcus1 id.11373) on Crohn's disease of large intestine

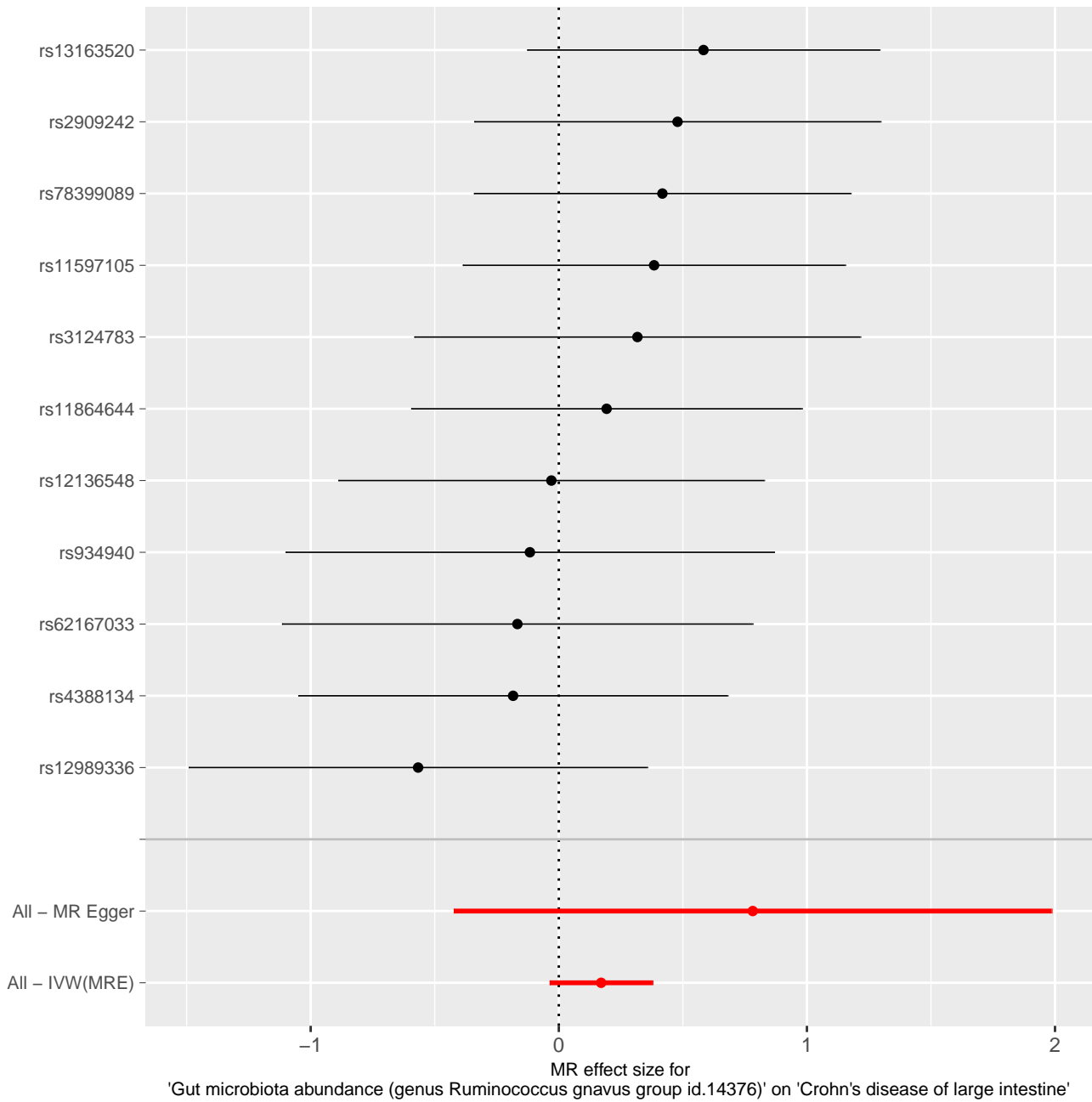


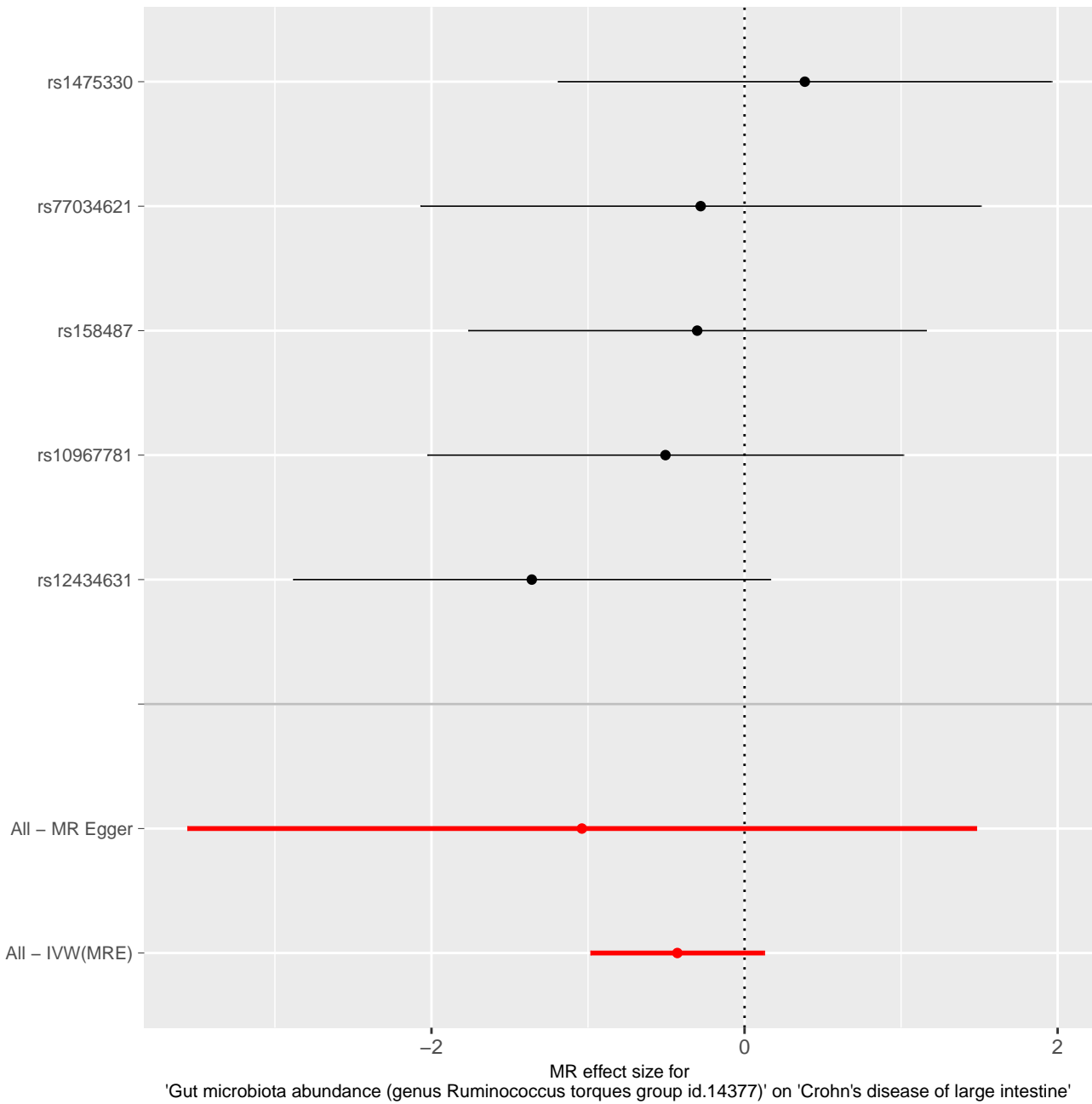
Batch 137 : Gut microbiota abundance (genus Ruminococcus2 id.11374) on Crohn's disease of large intestine

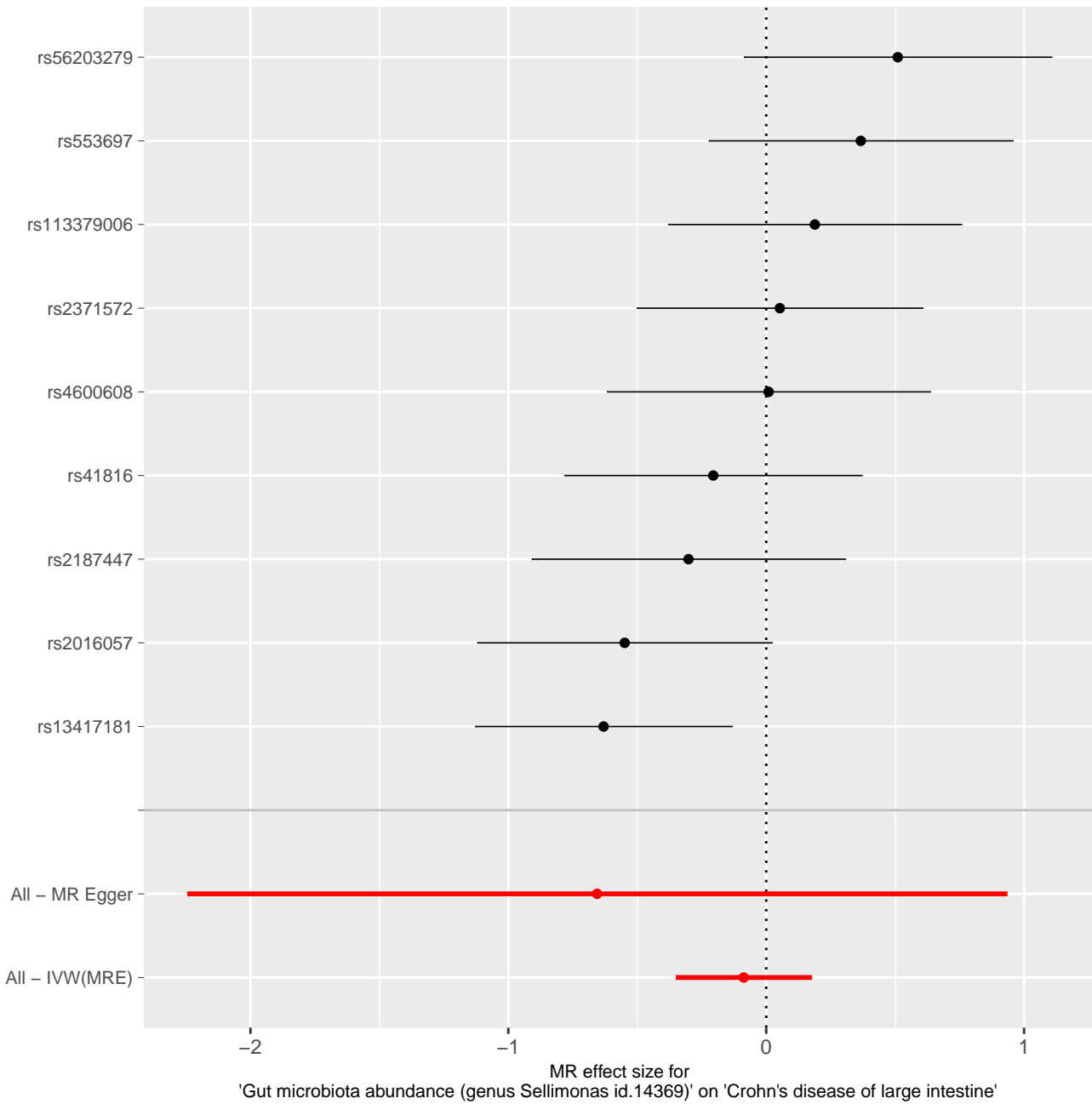


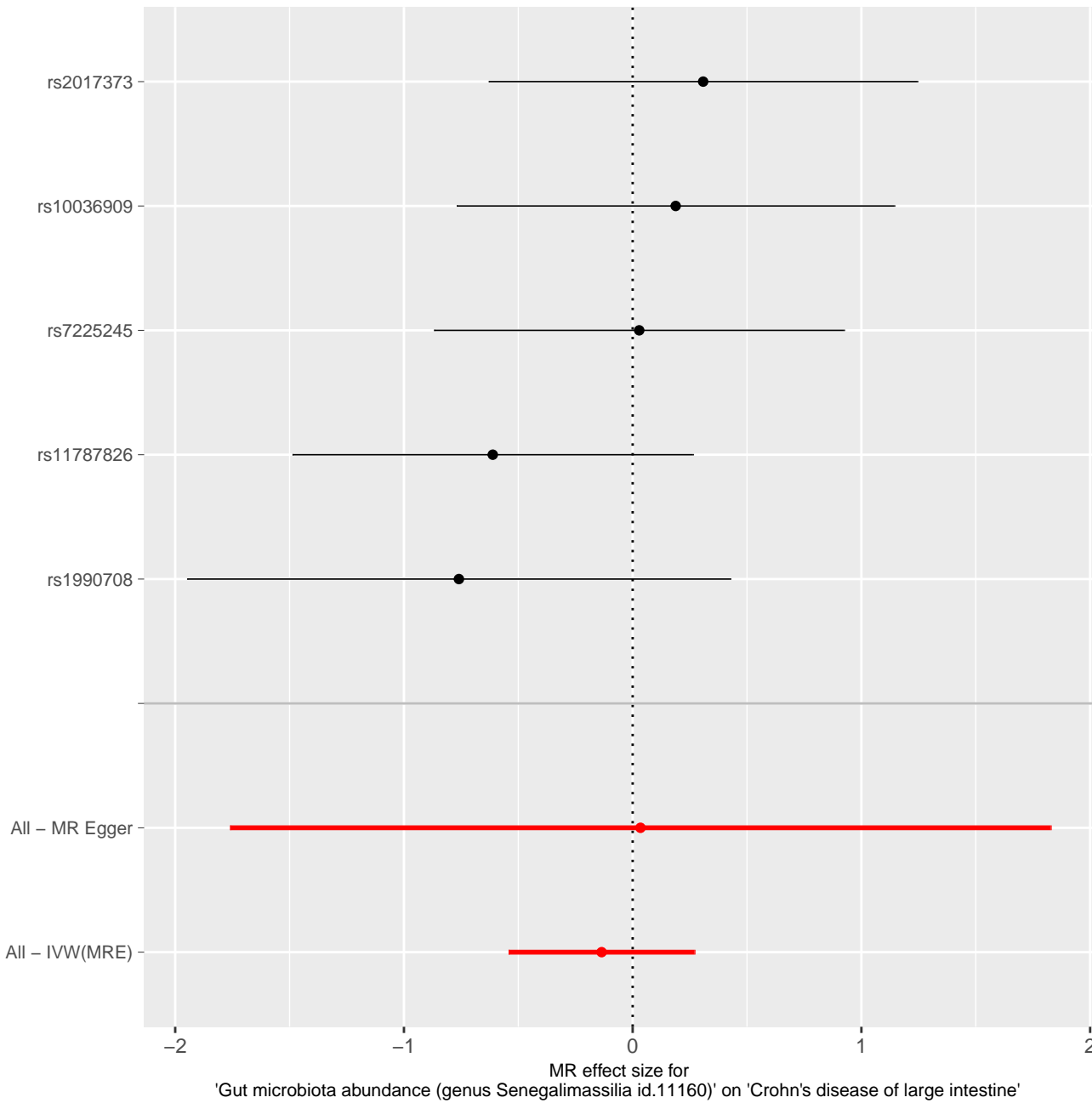


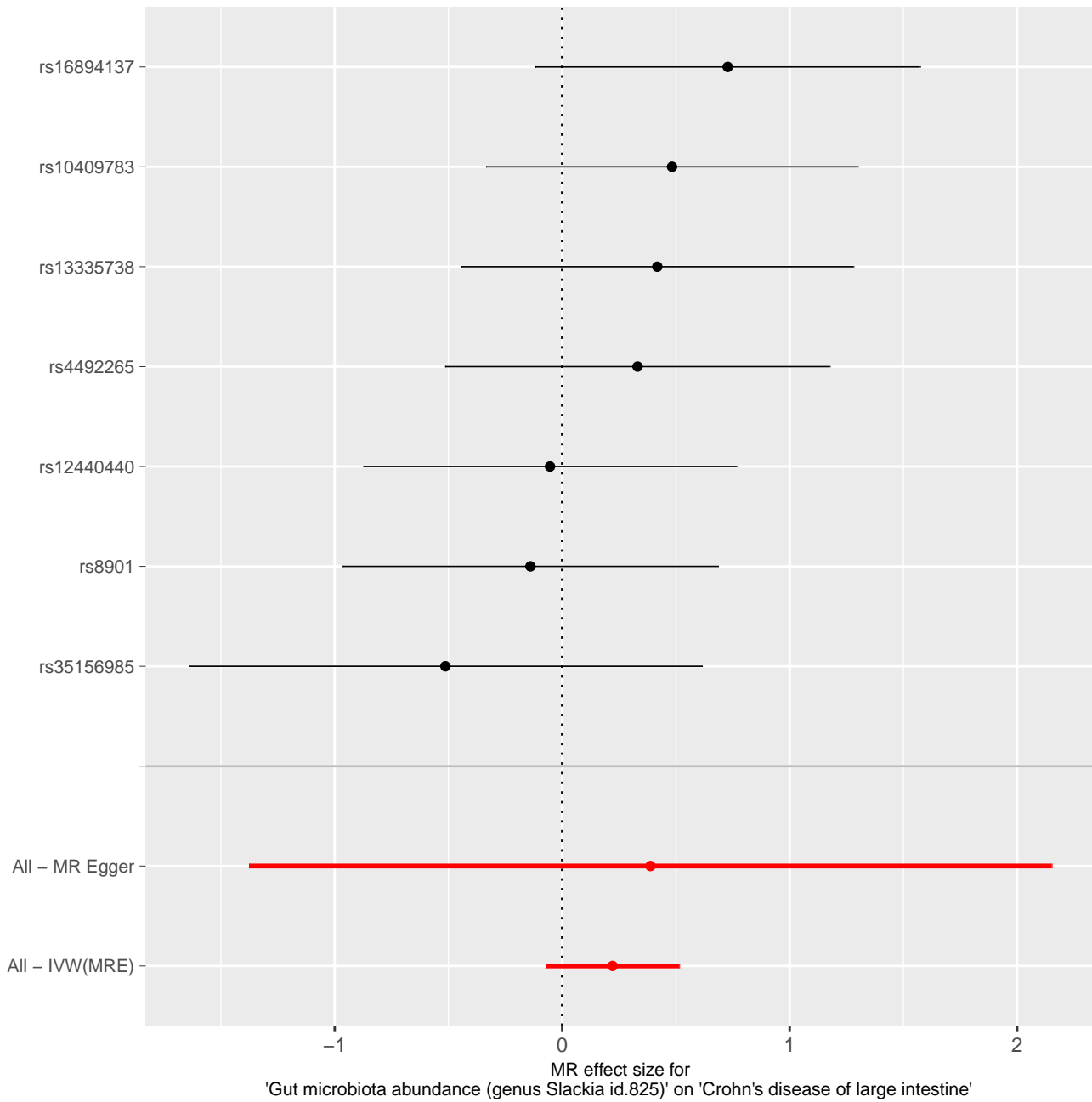
Batch 139 : Gut microbiota abundance (genus Ruminococcus gnavus group id.14376) on Crohn's disease of large intestine



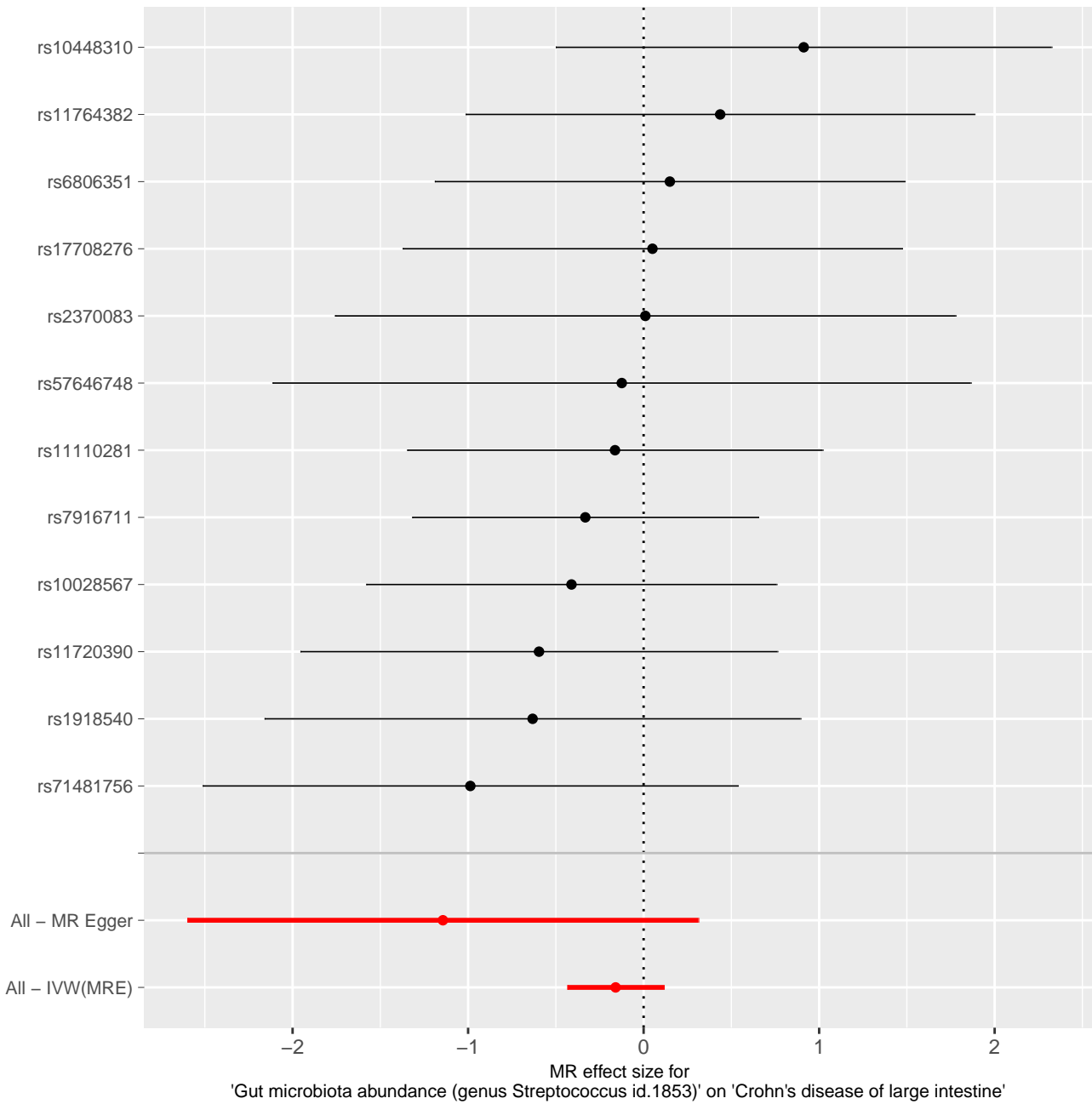


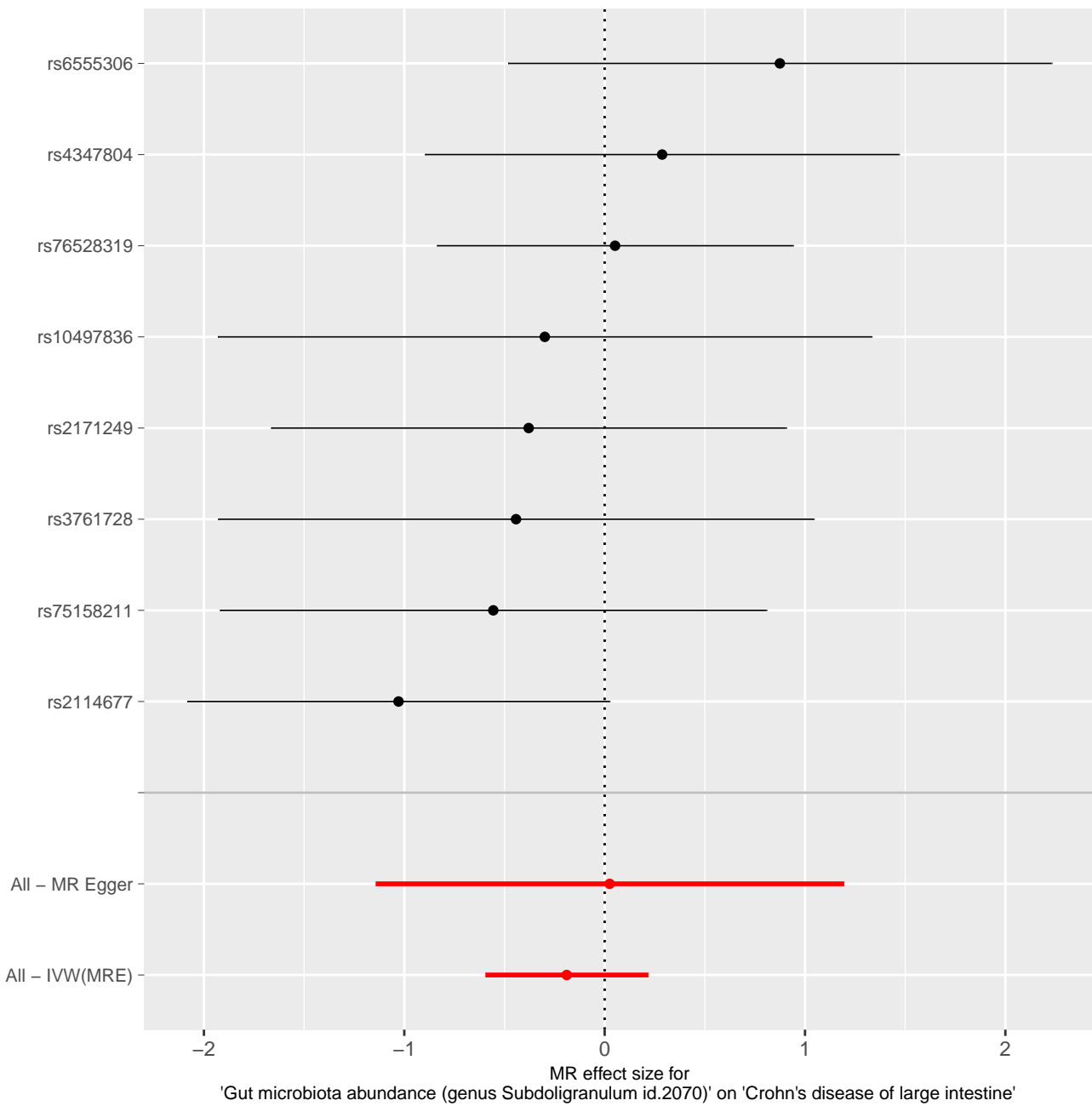




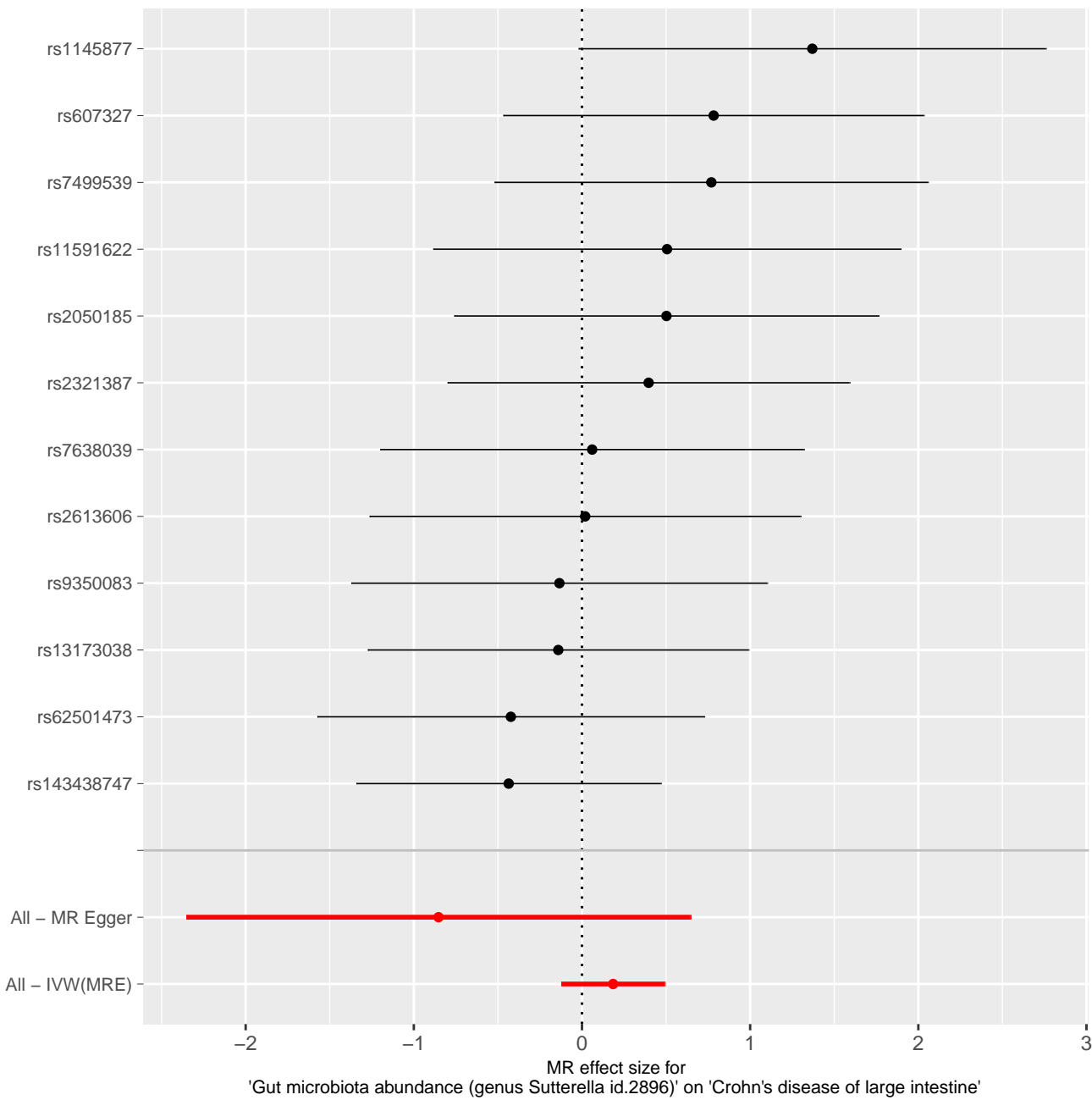


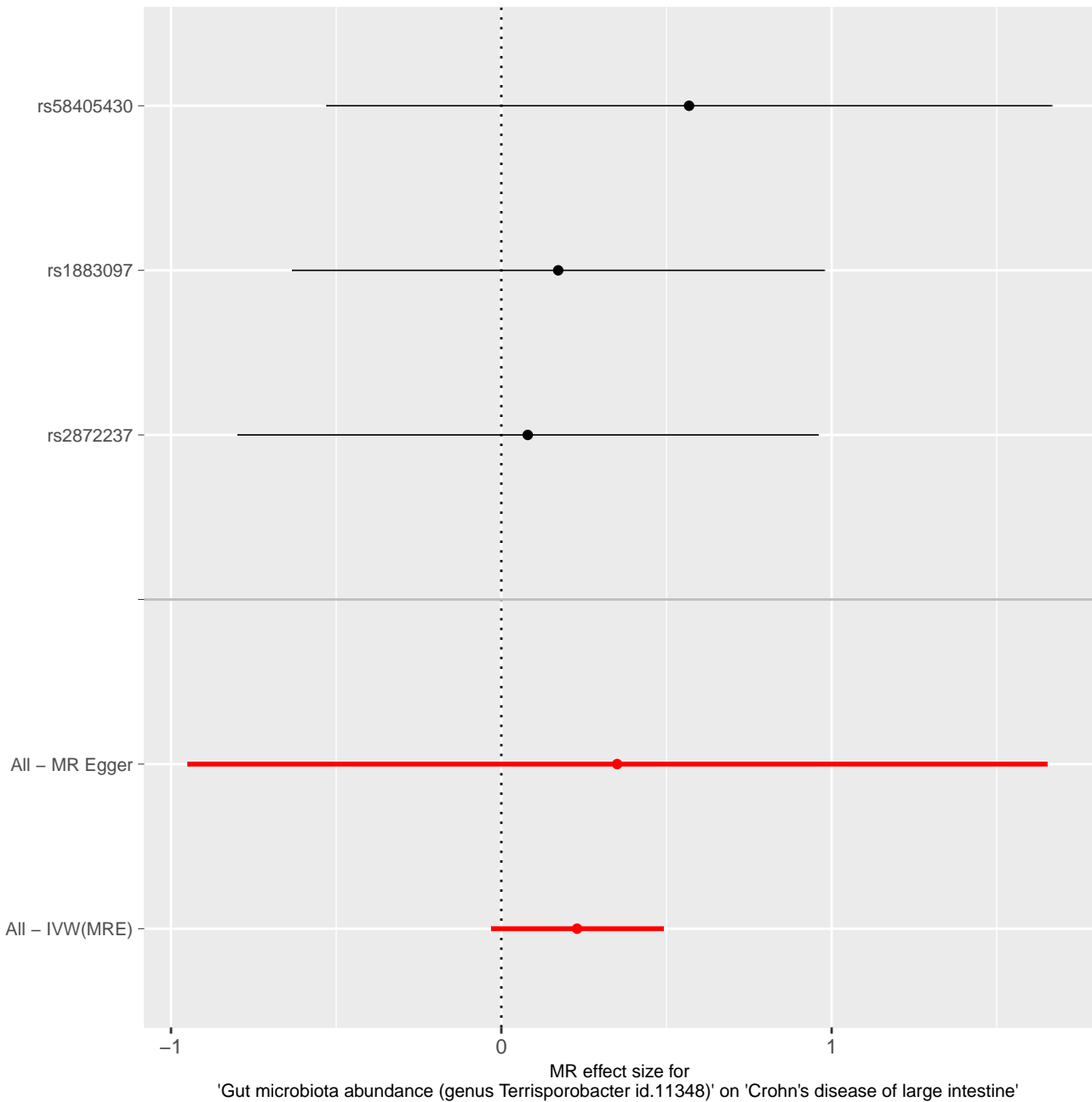
Batch 144 : Gut microbiota abundance (genus Streptococcus id.1853) on Crohn's disease of large intestine

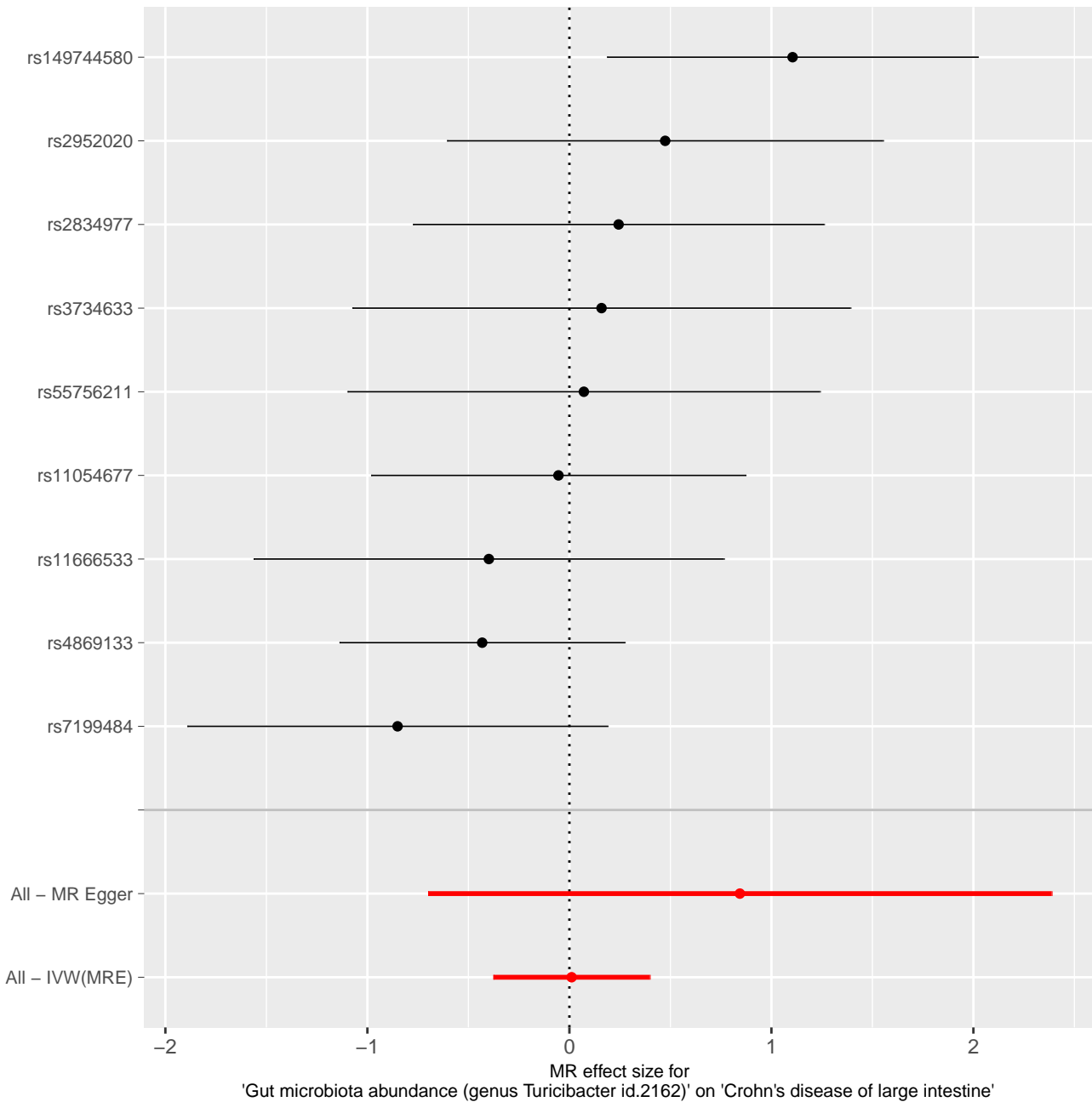


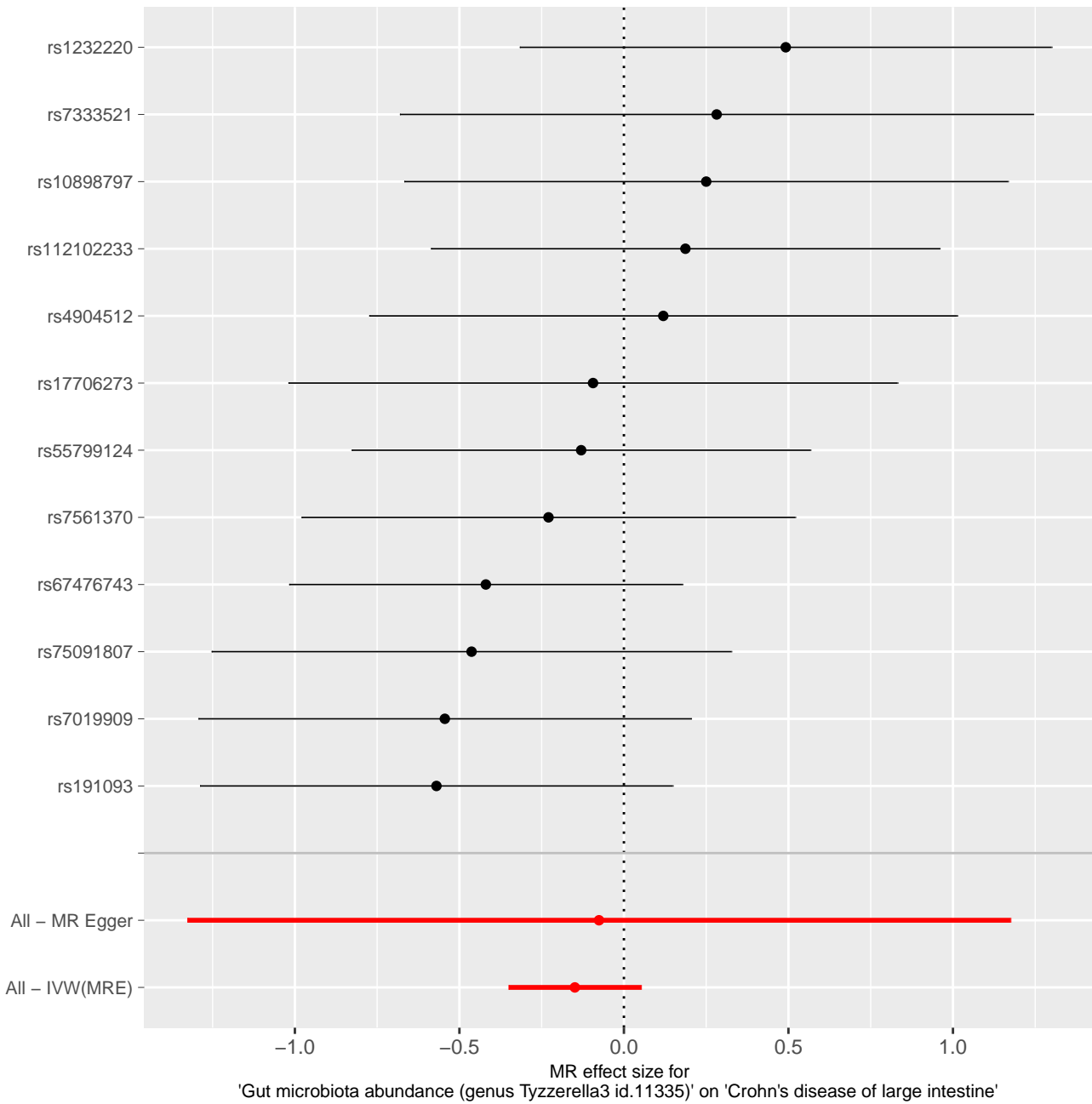


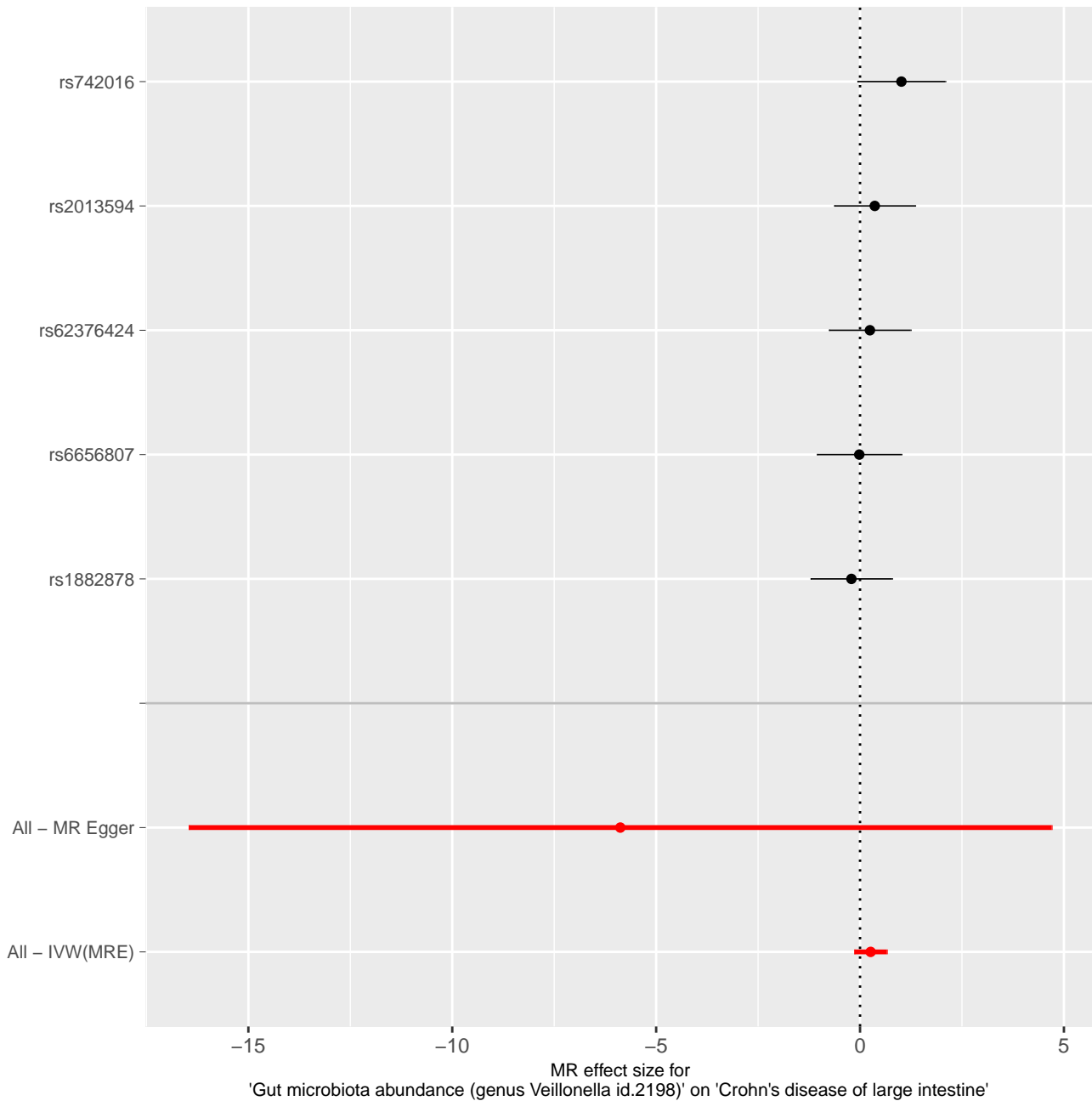
Batch 146 : Gut microbiota abundance (genus Sutterella id.2896) on Crohn's disease of large intestine

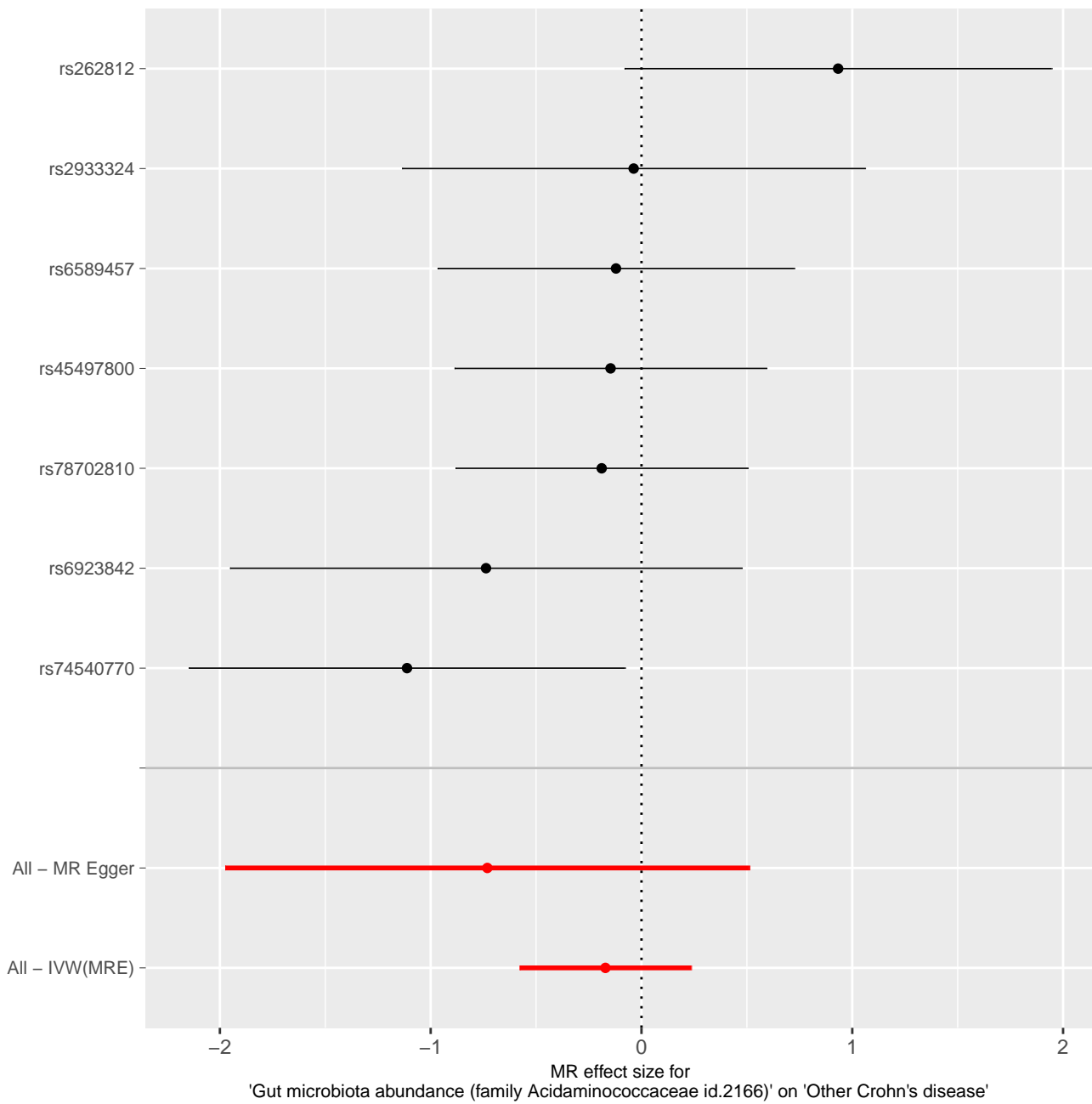


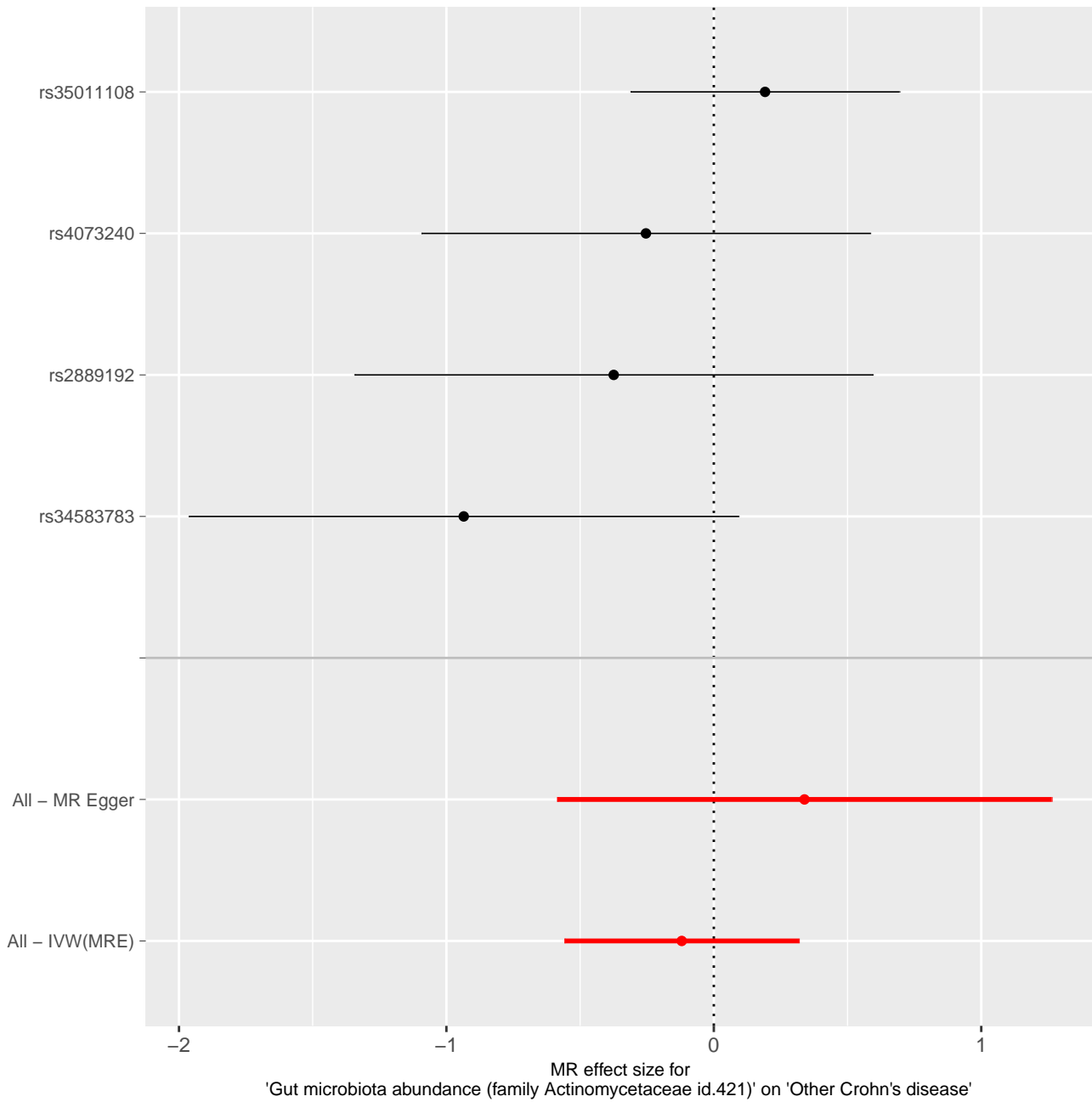


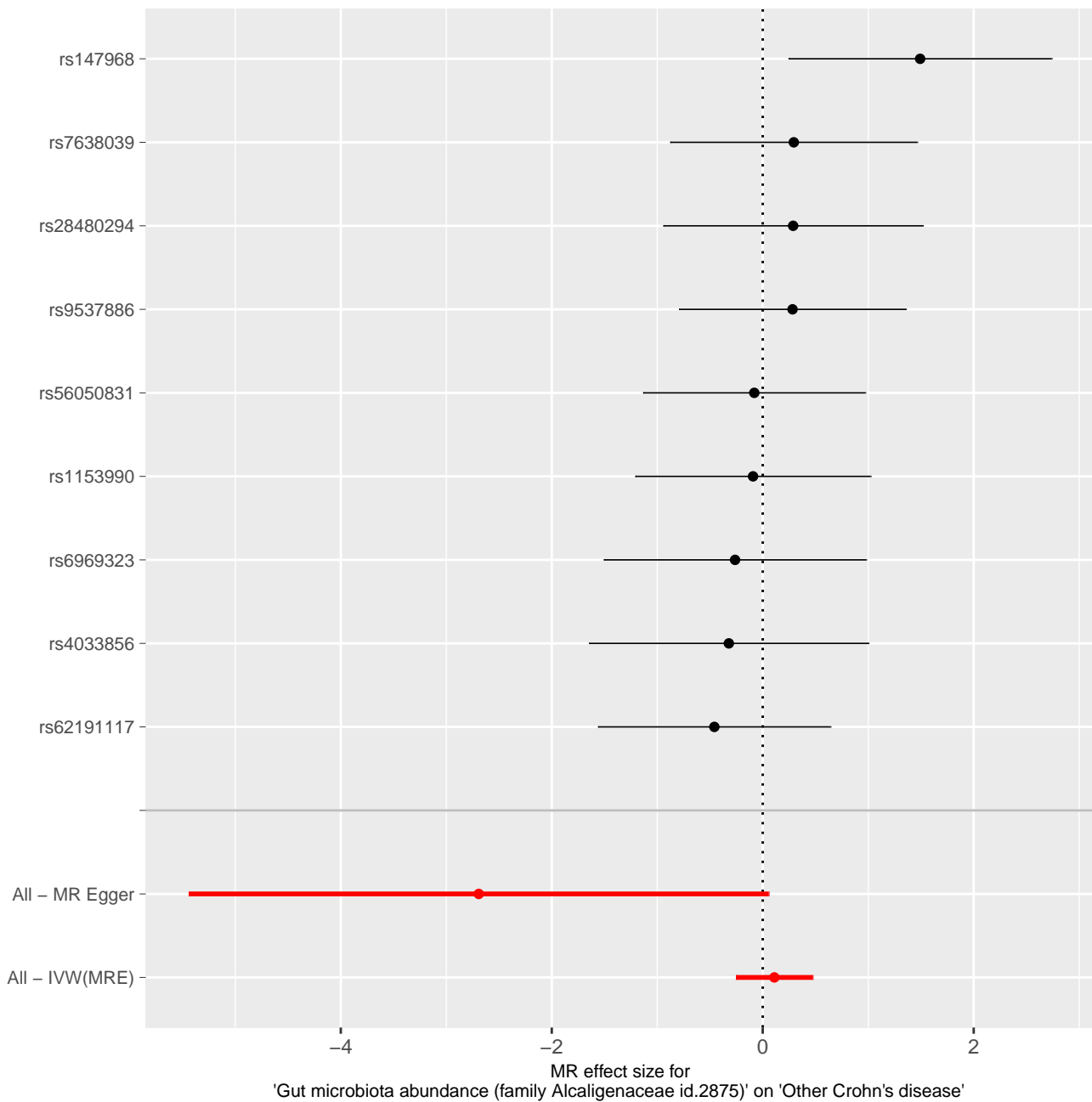


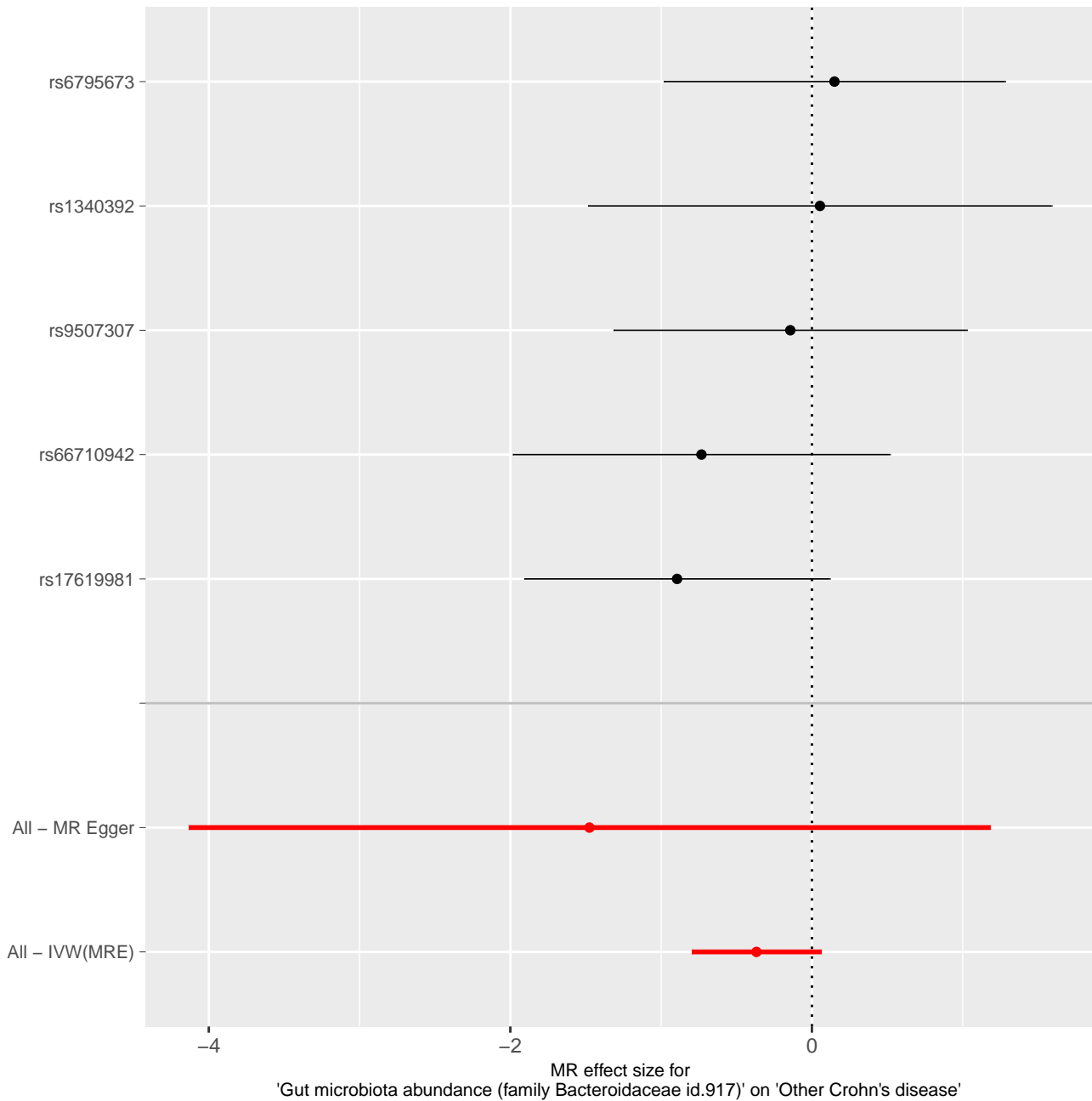


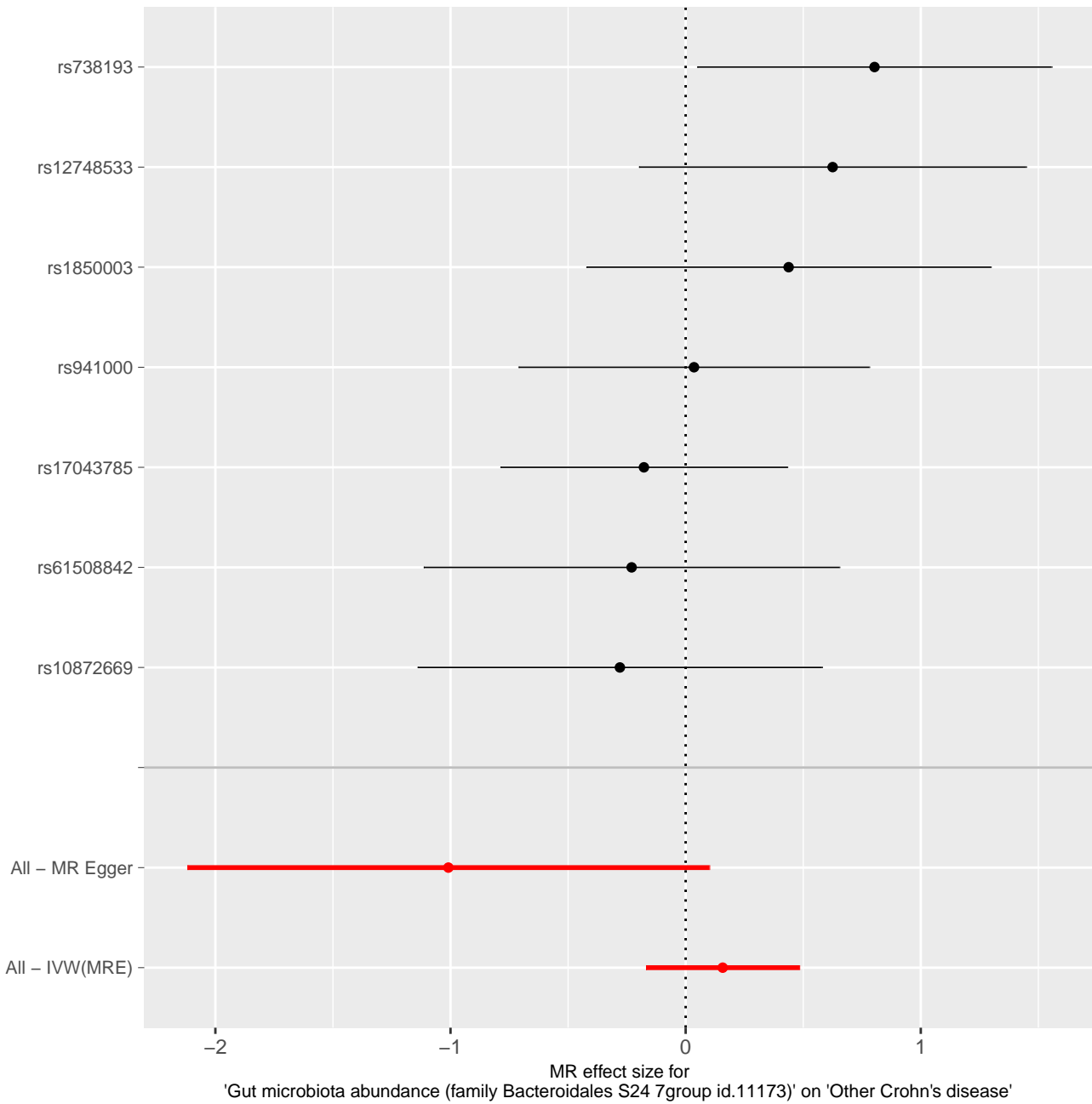


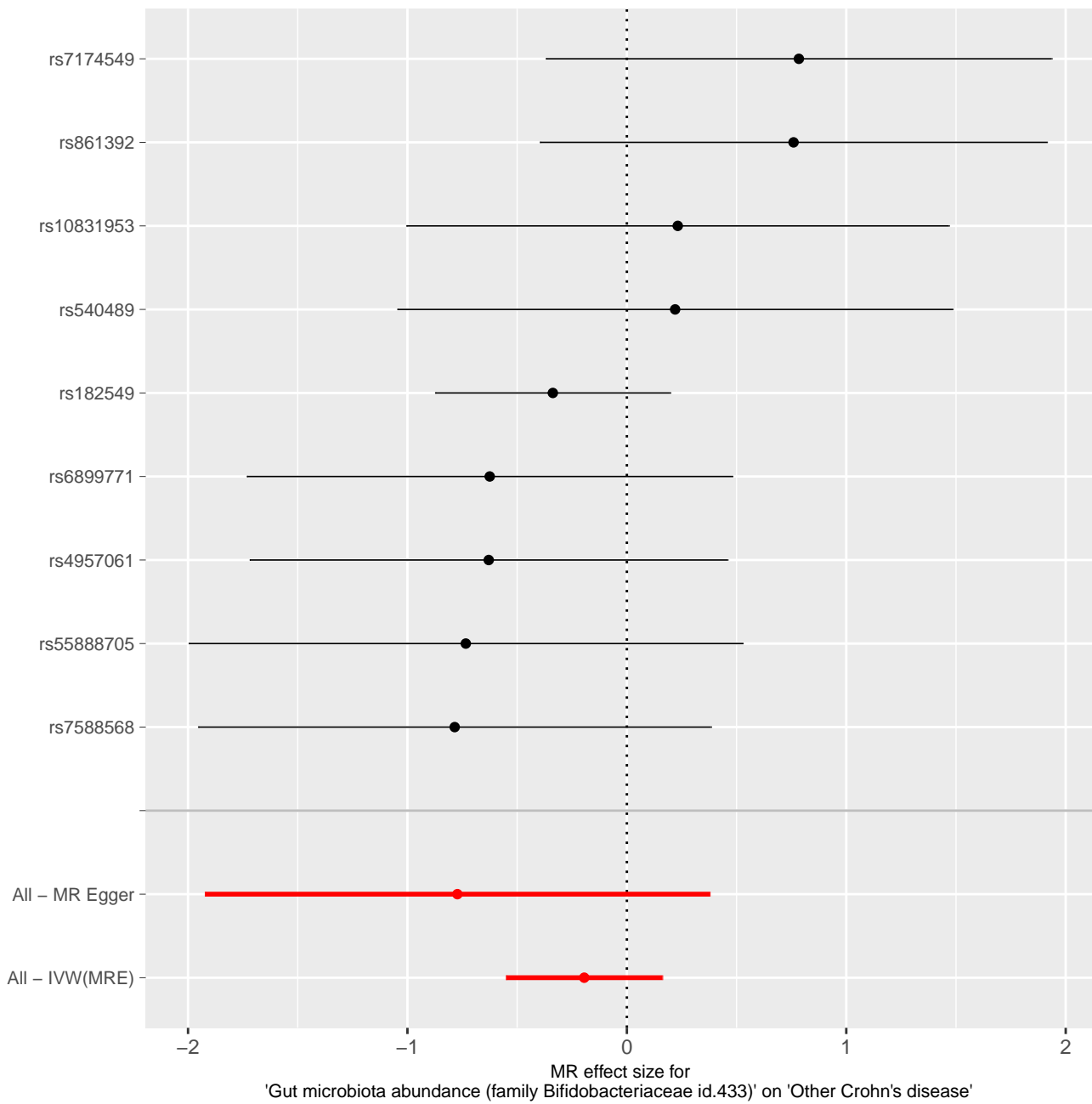


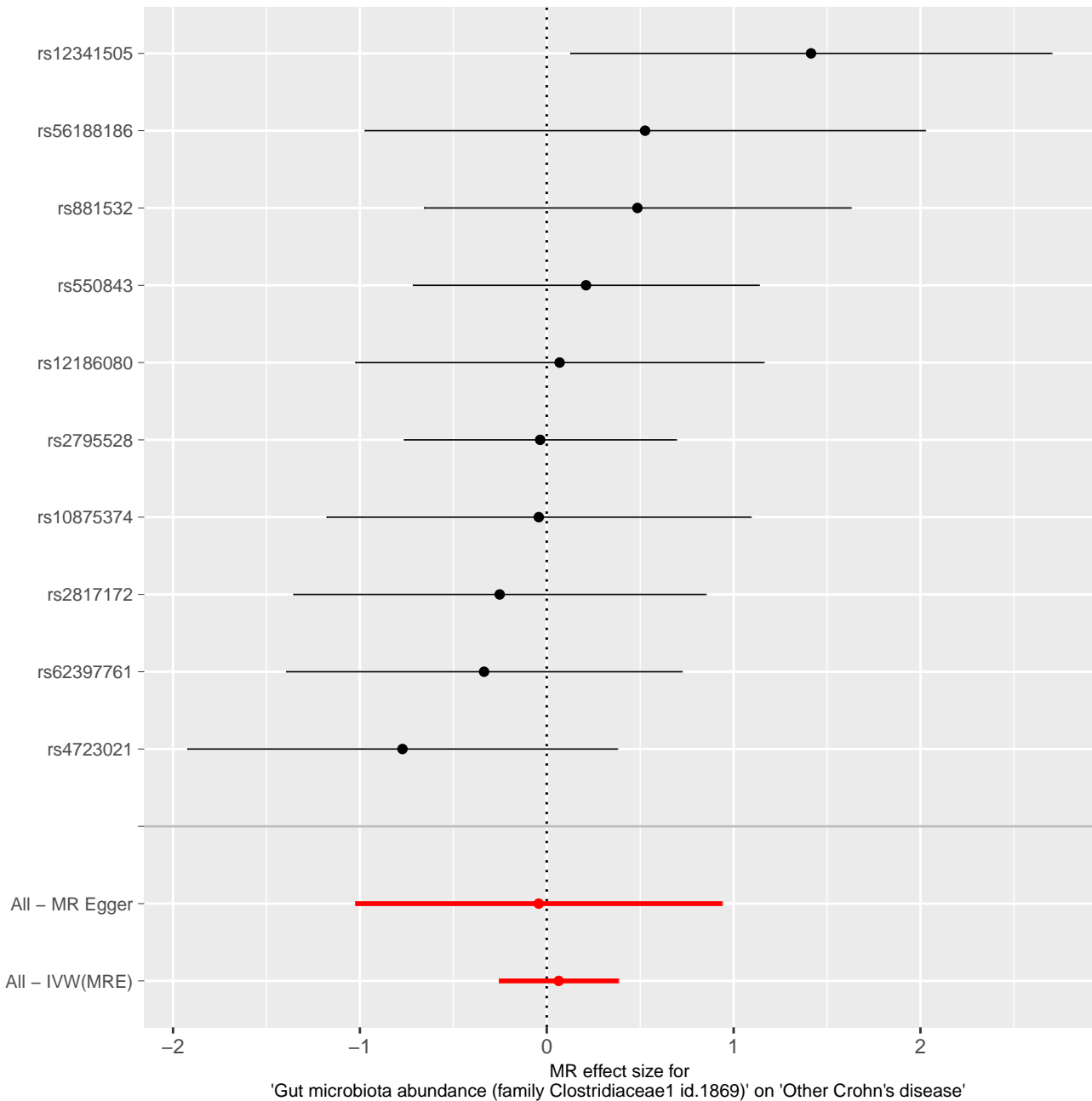


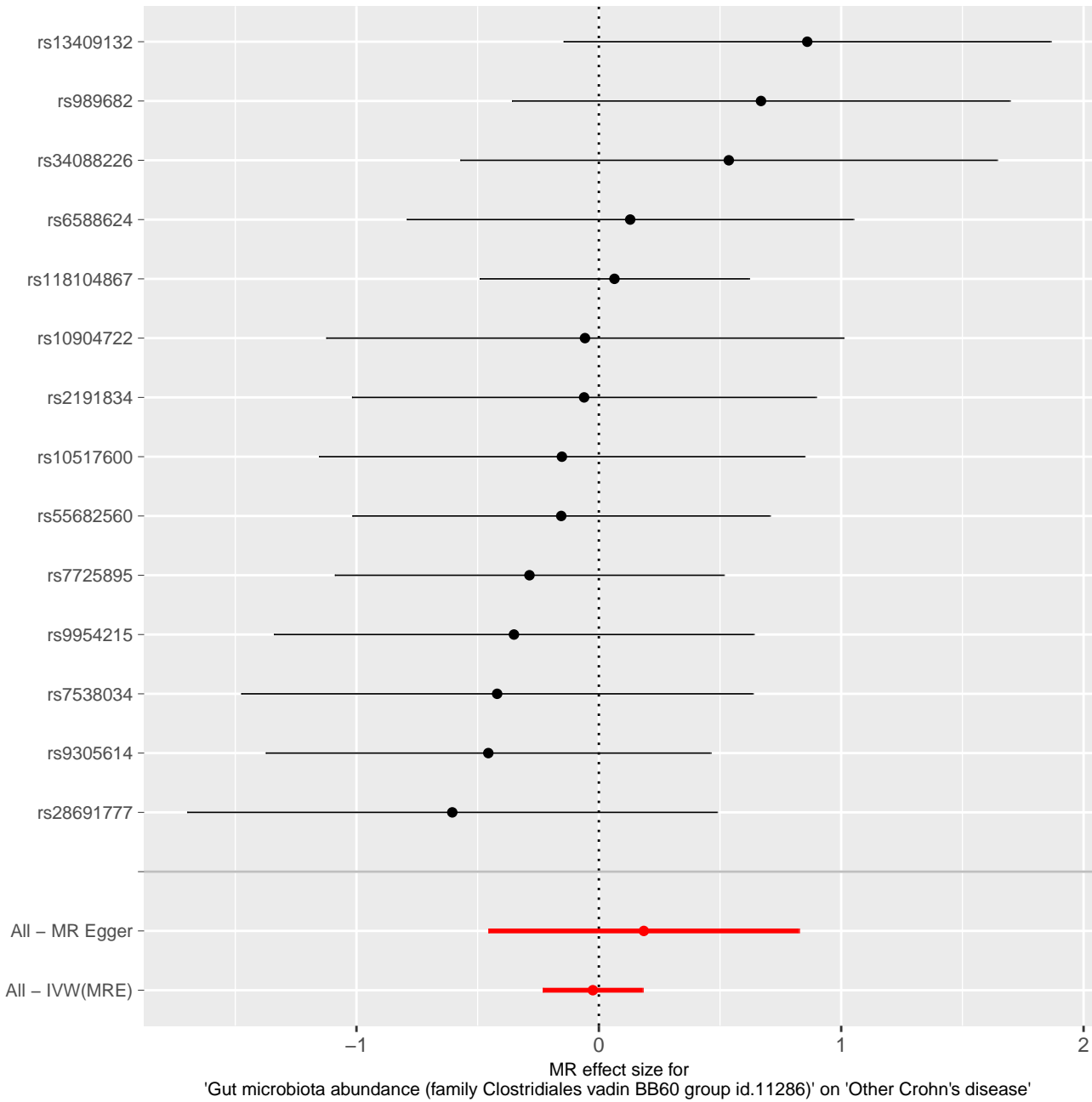




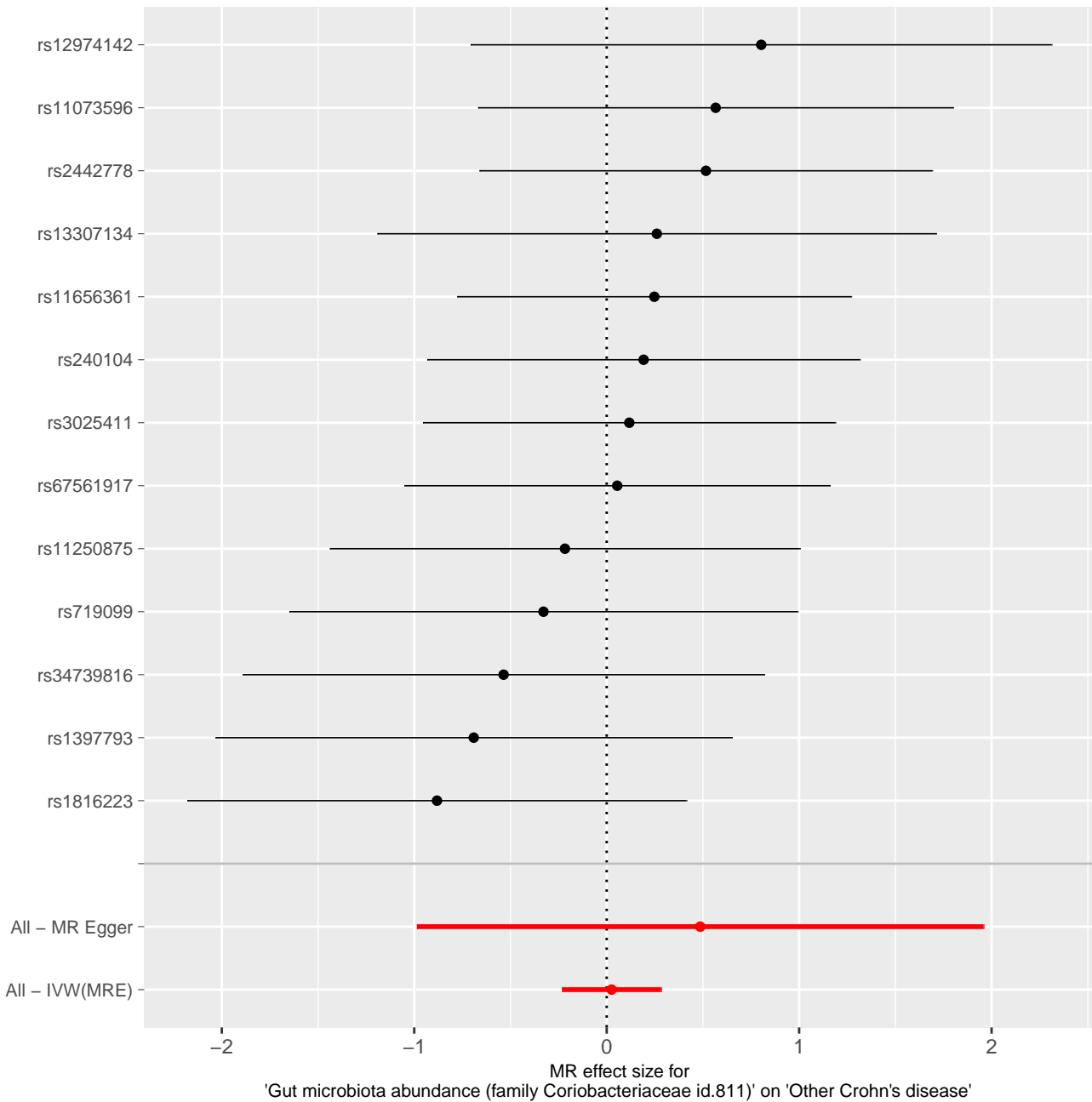


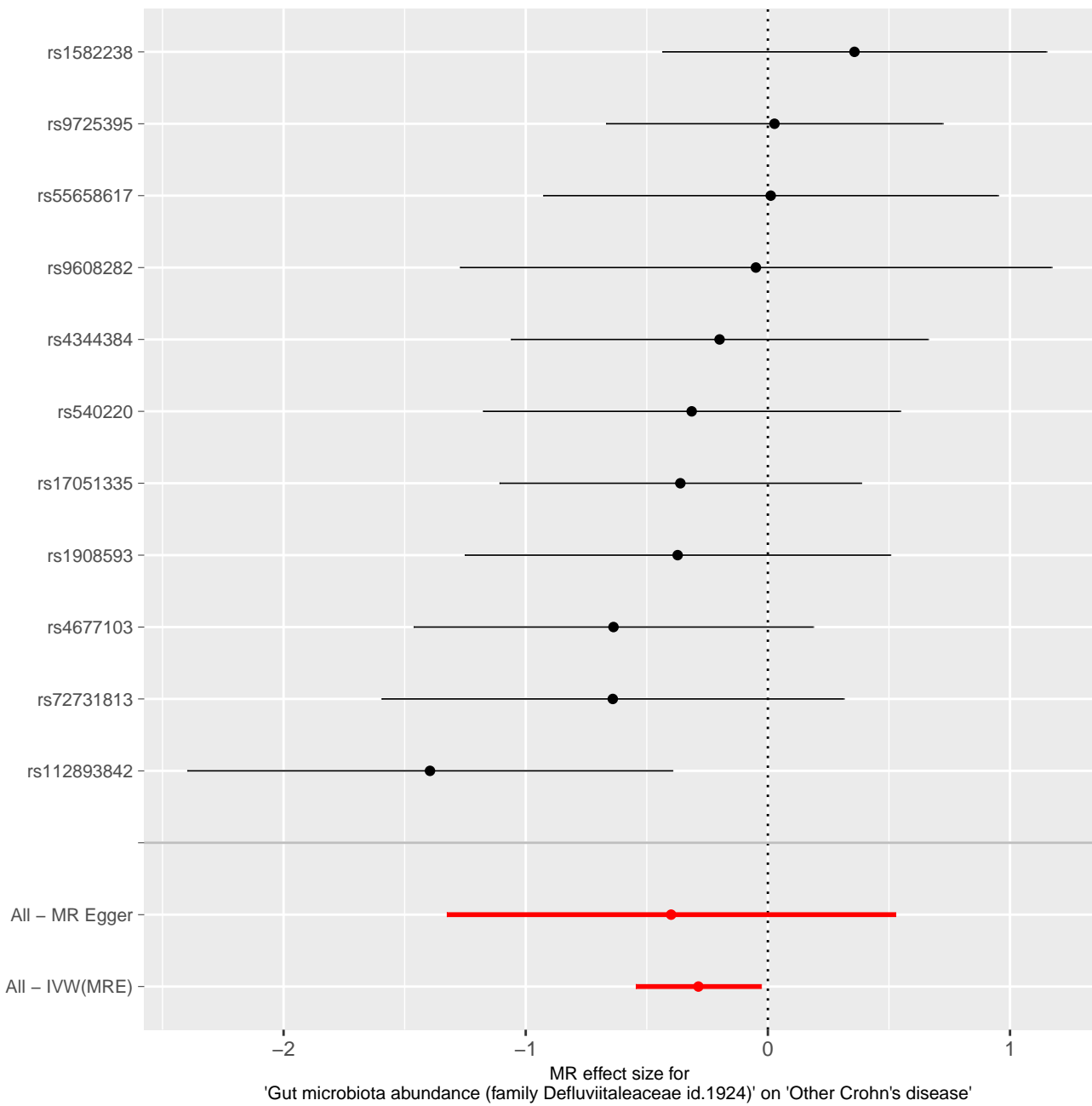


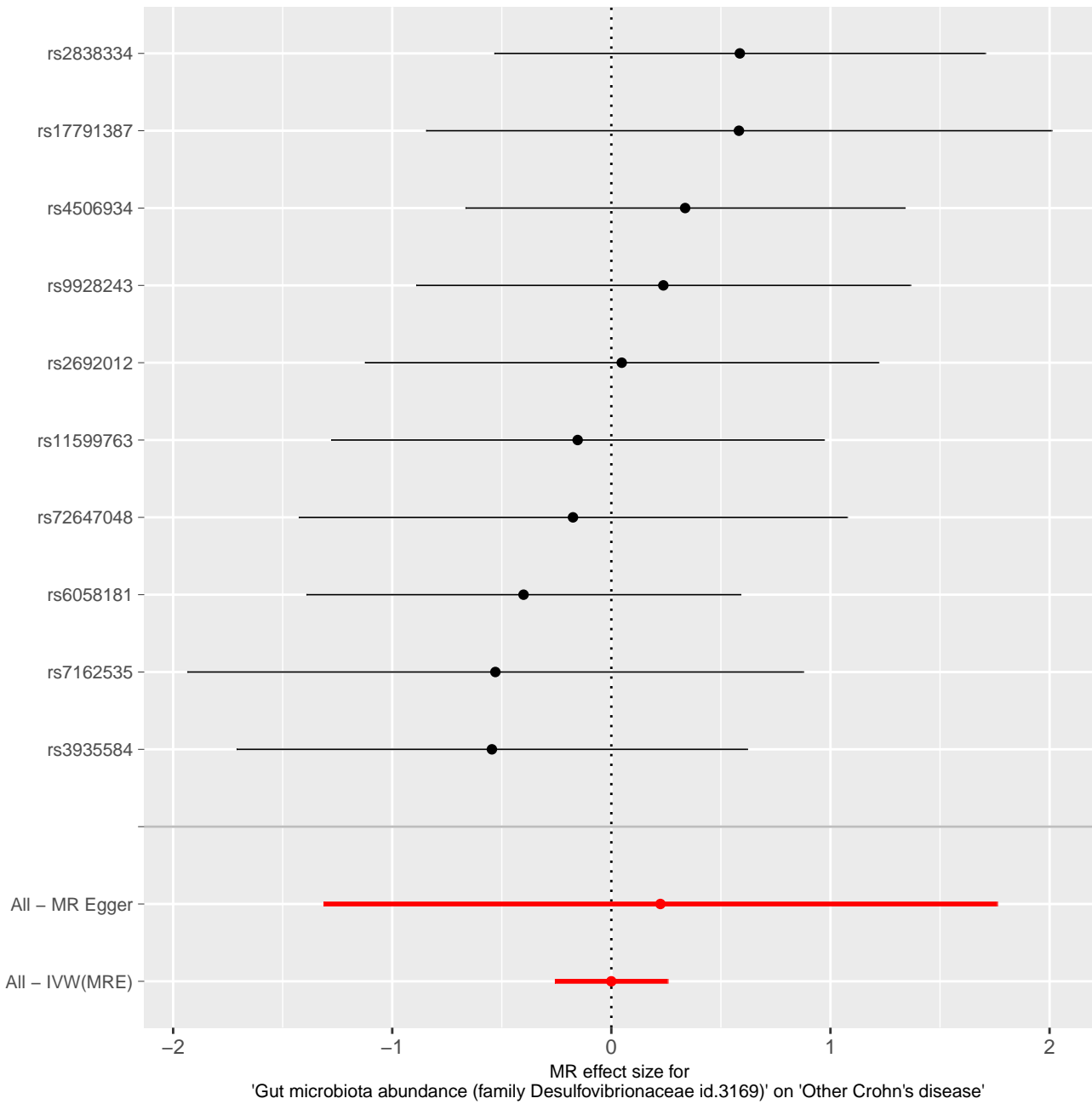


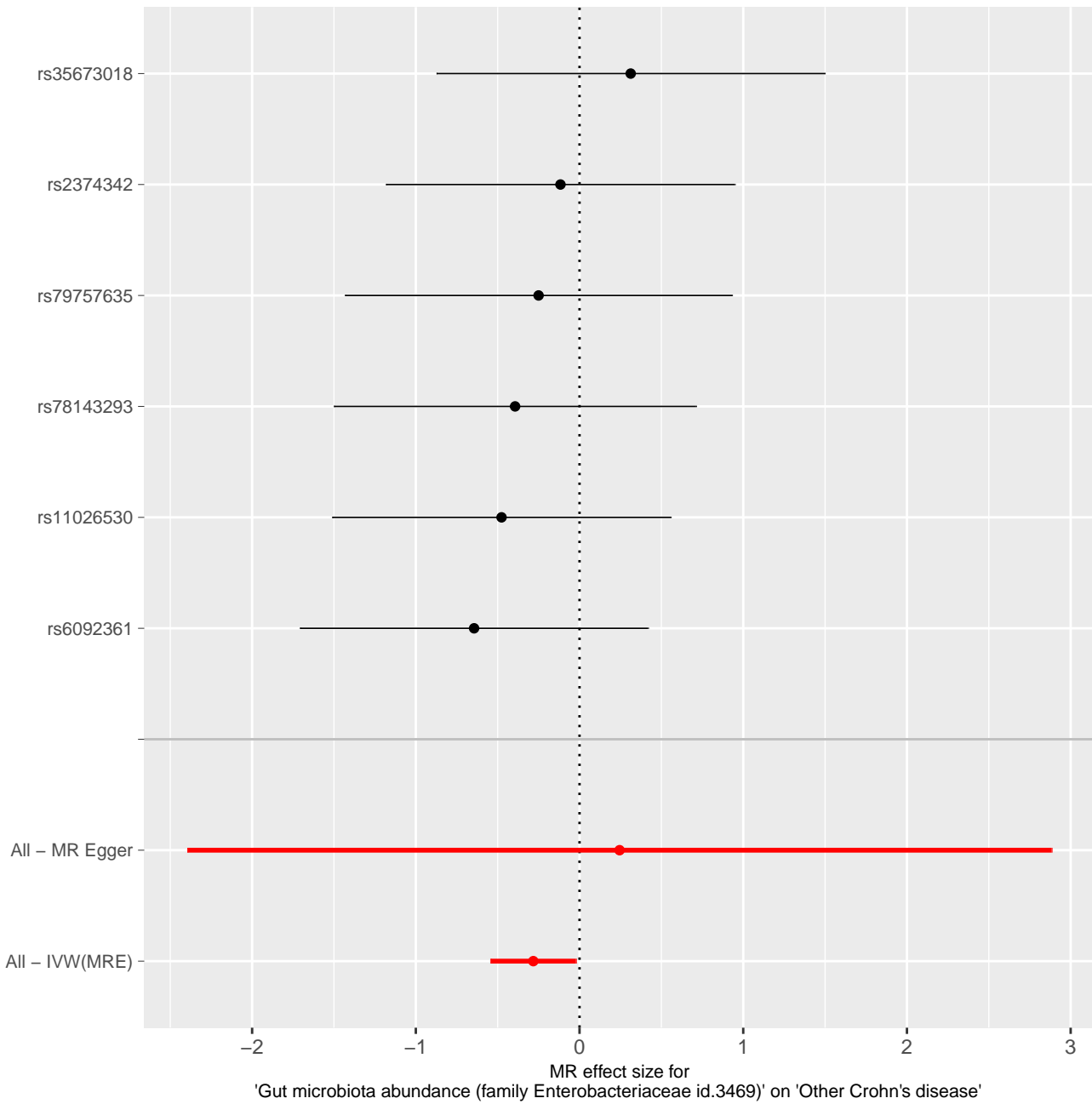


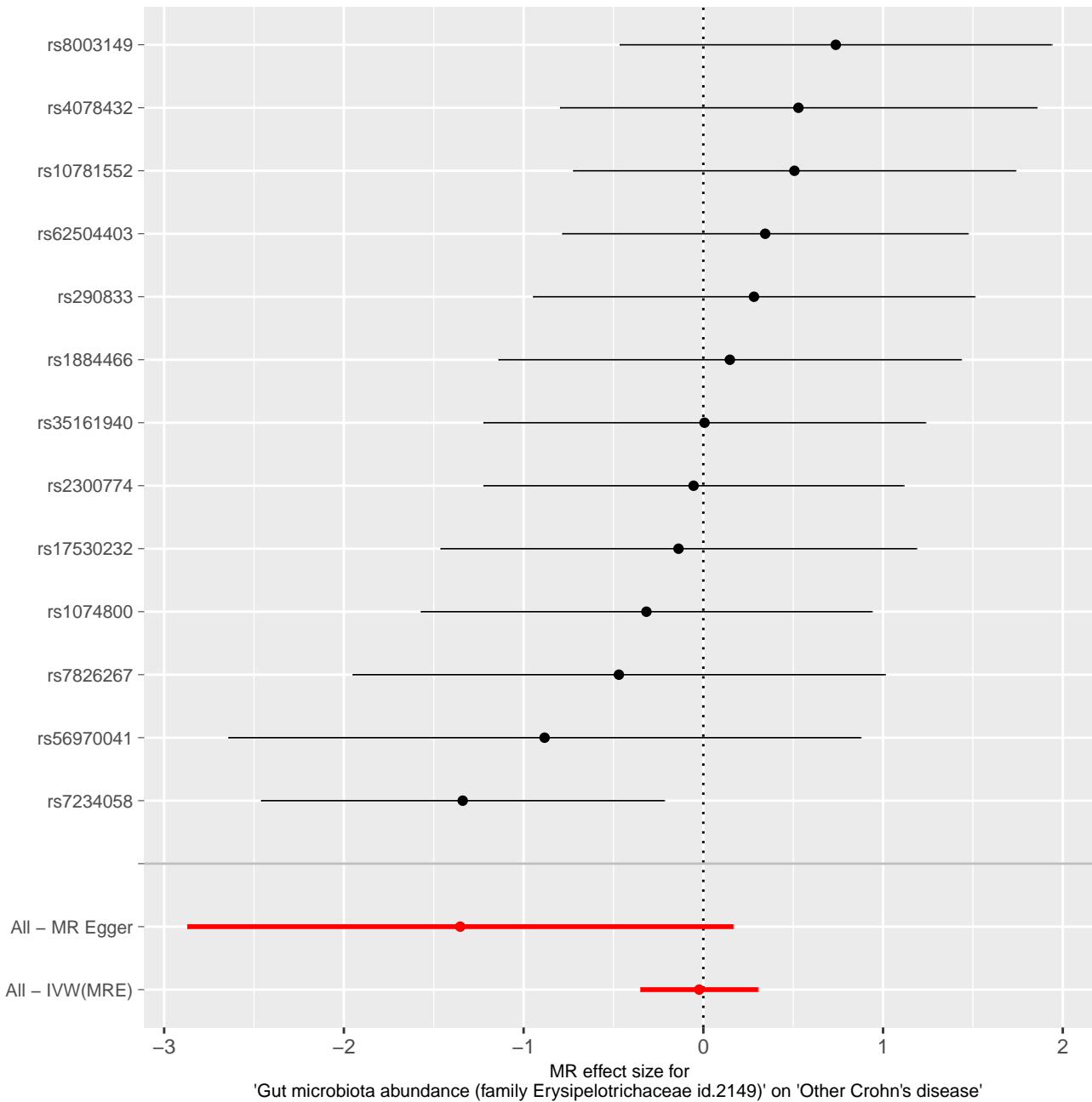
Batch 160 : Gut microbiota abundance (family Coriobacteriaceae id.811) on Other Crohn's disease

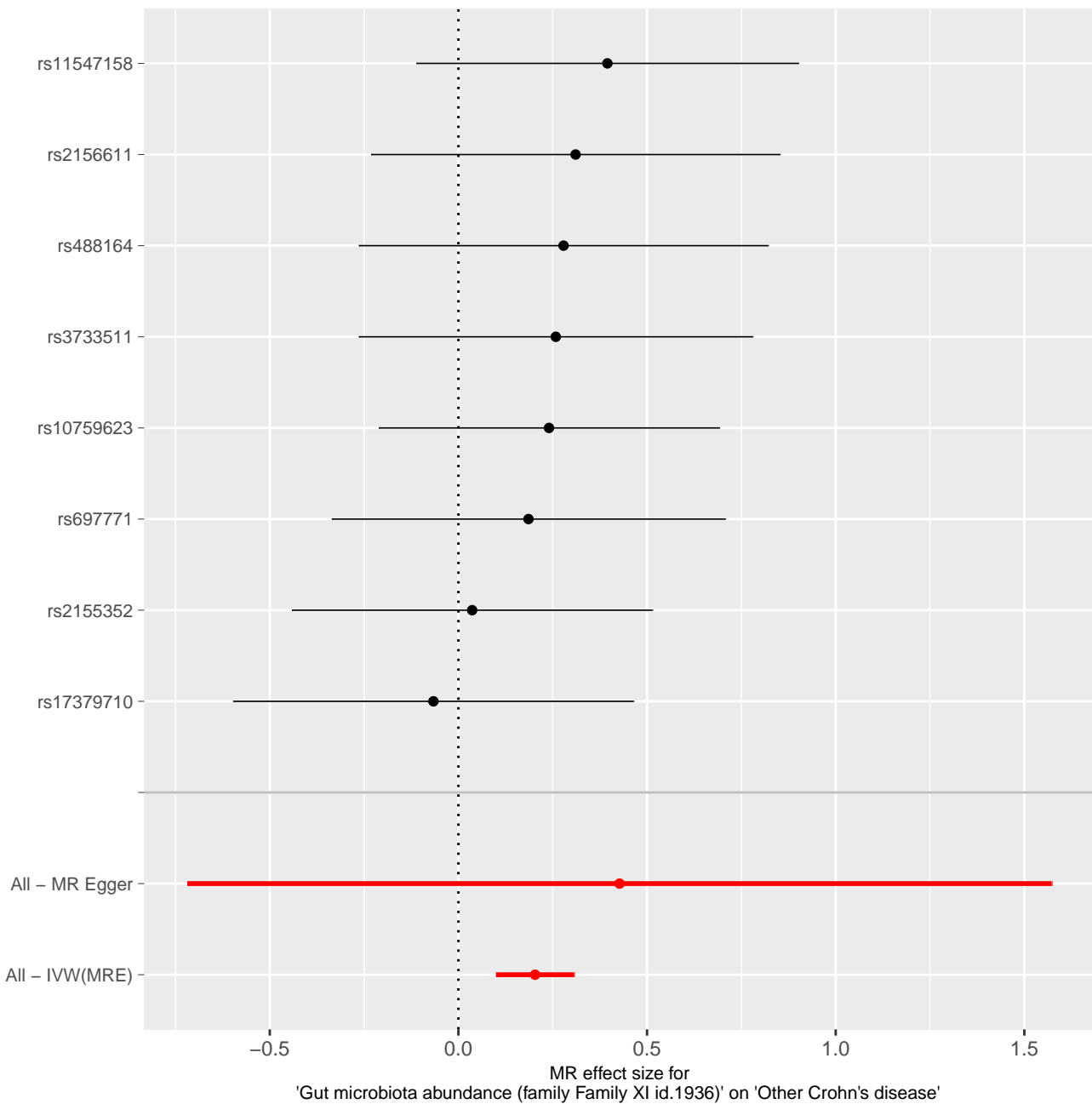


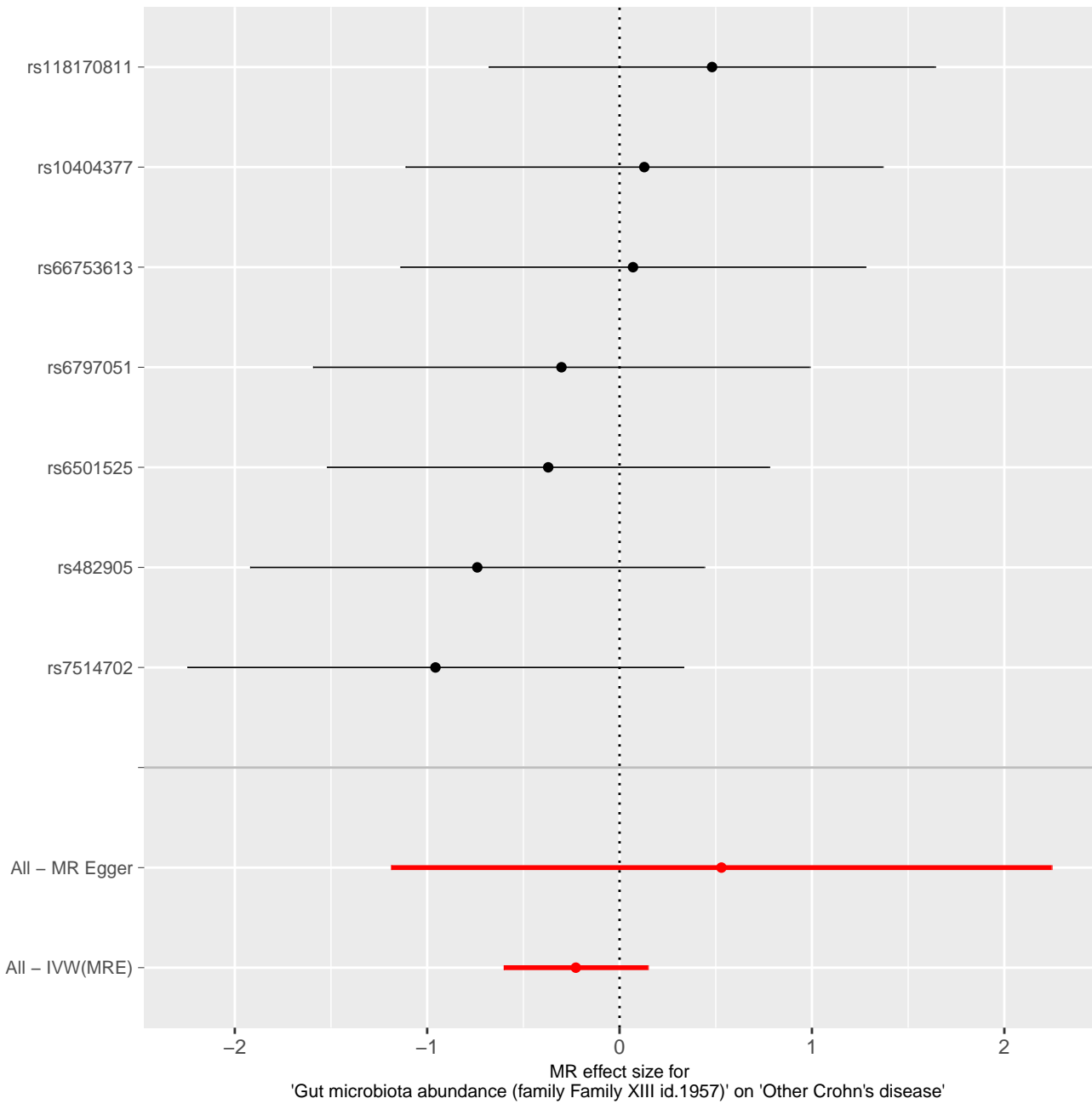




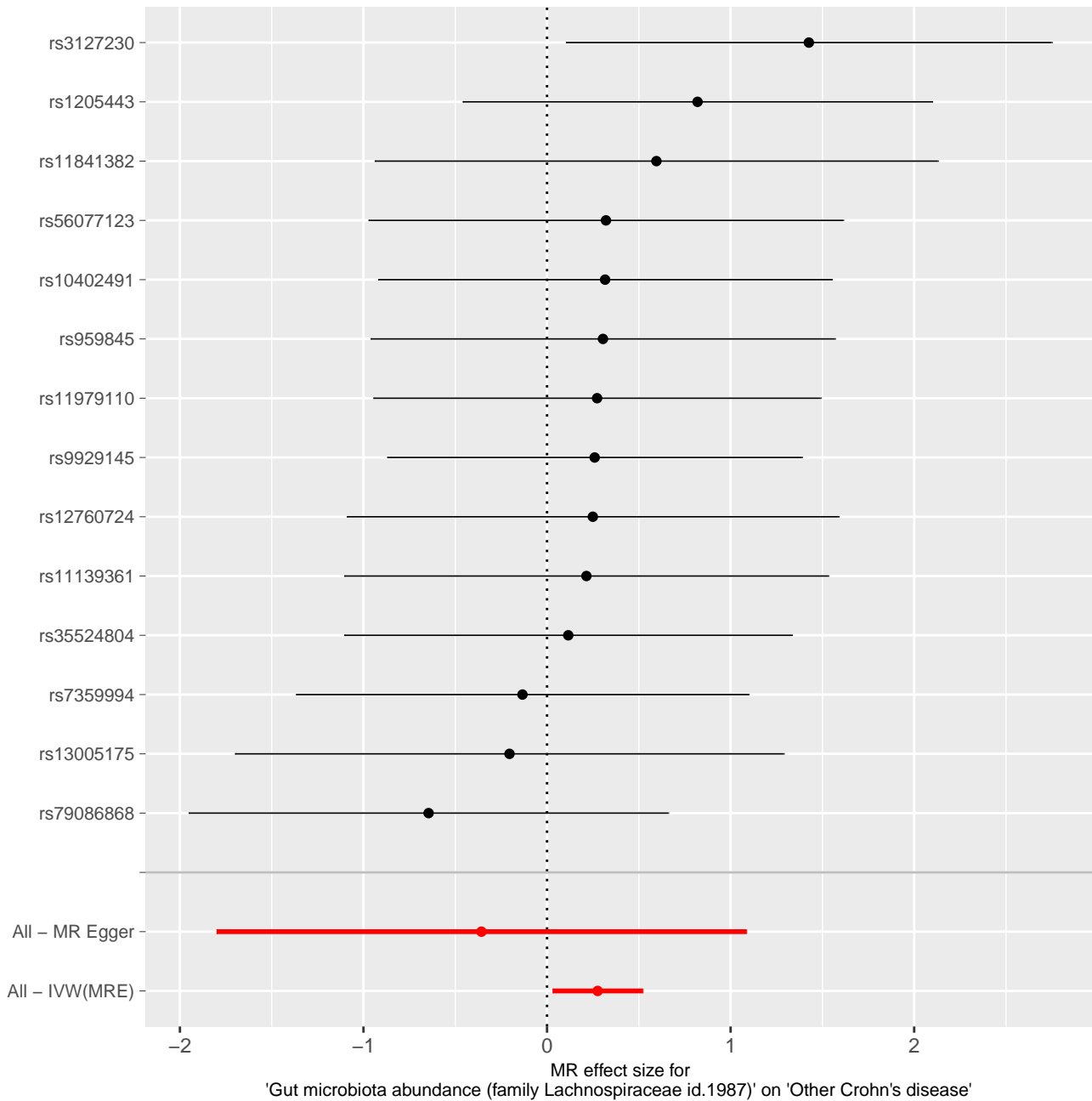


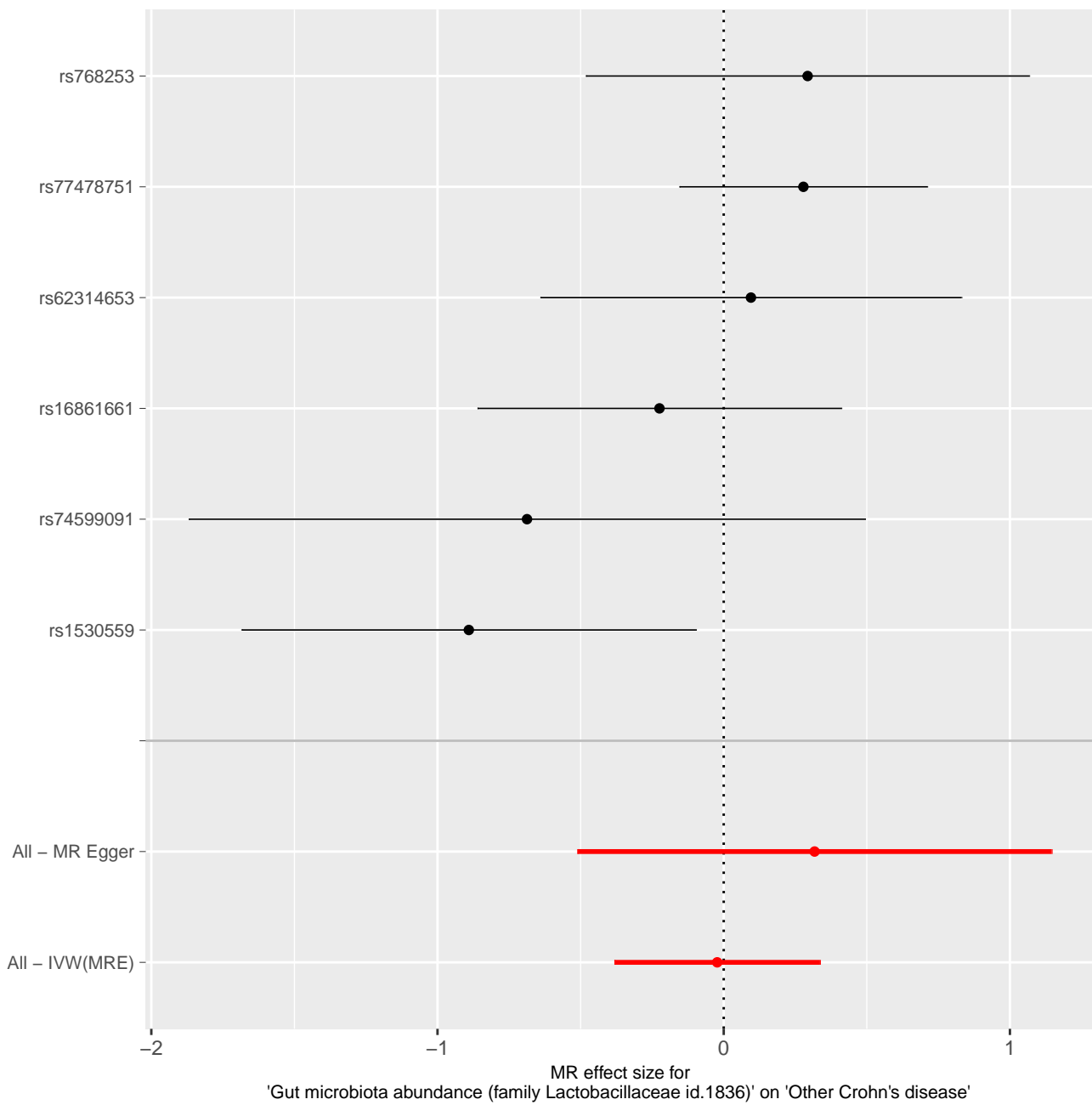


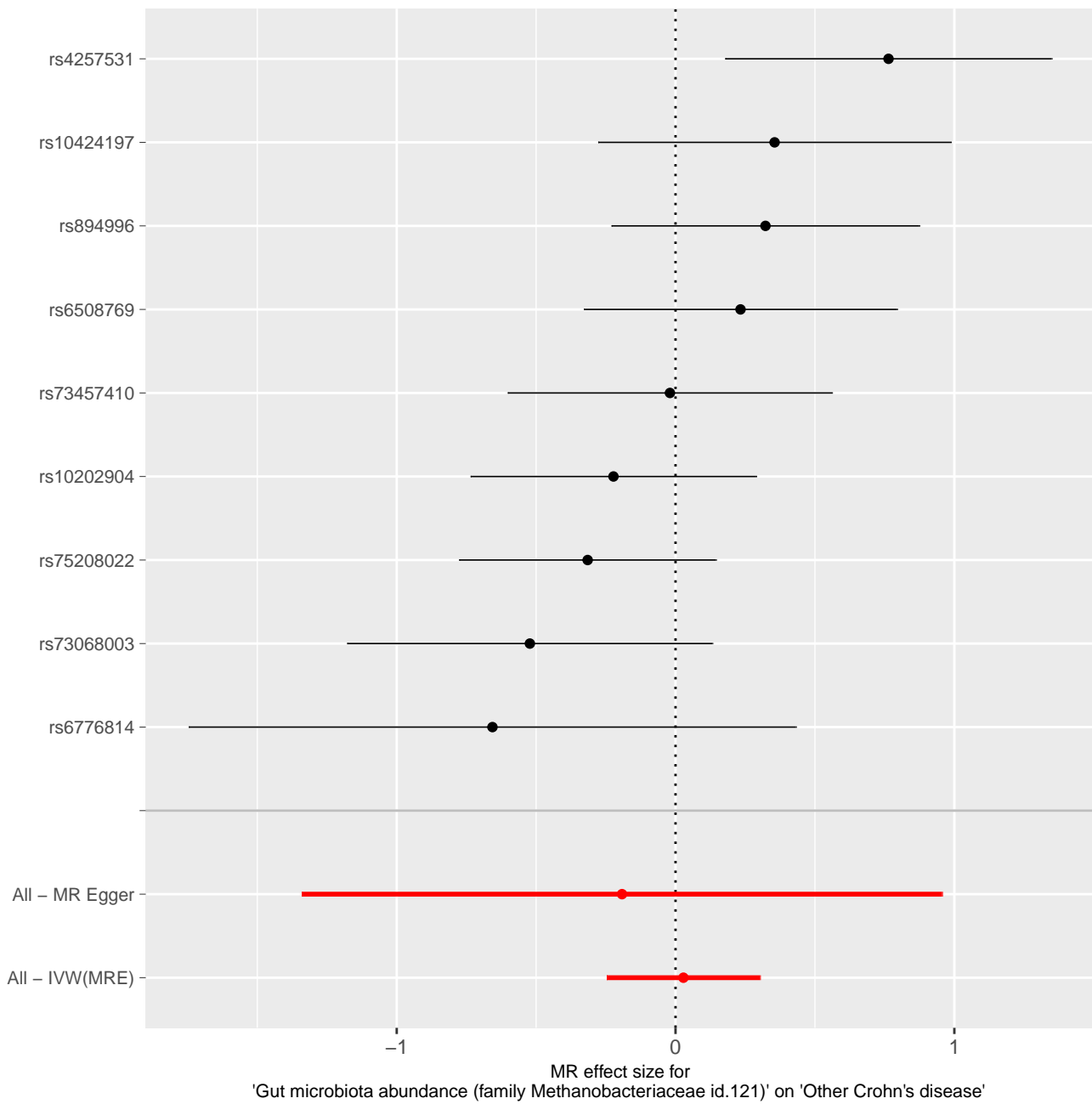


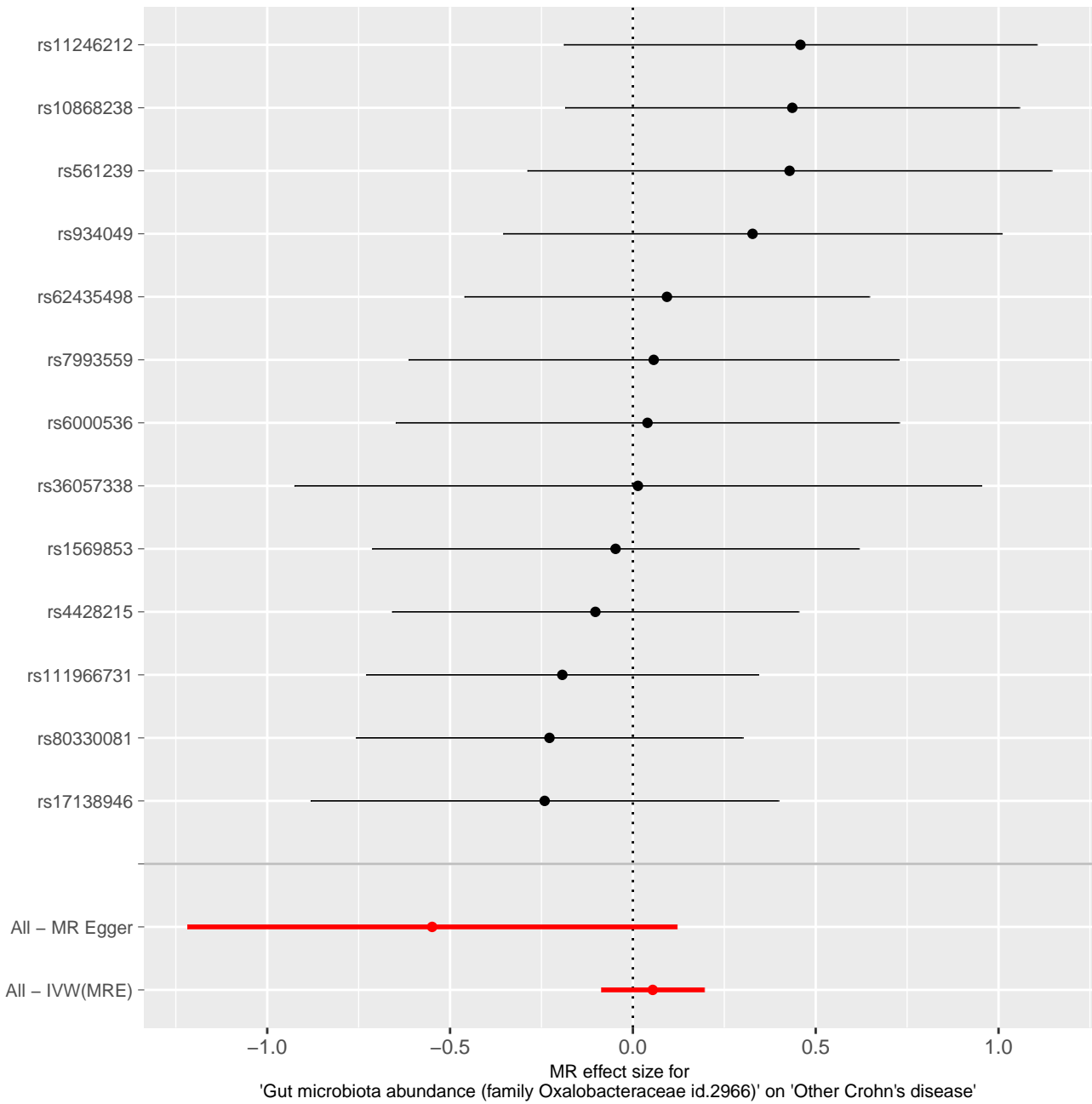


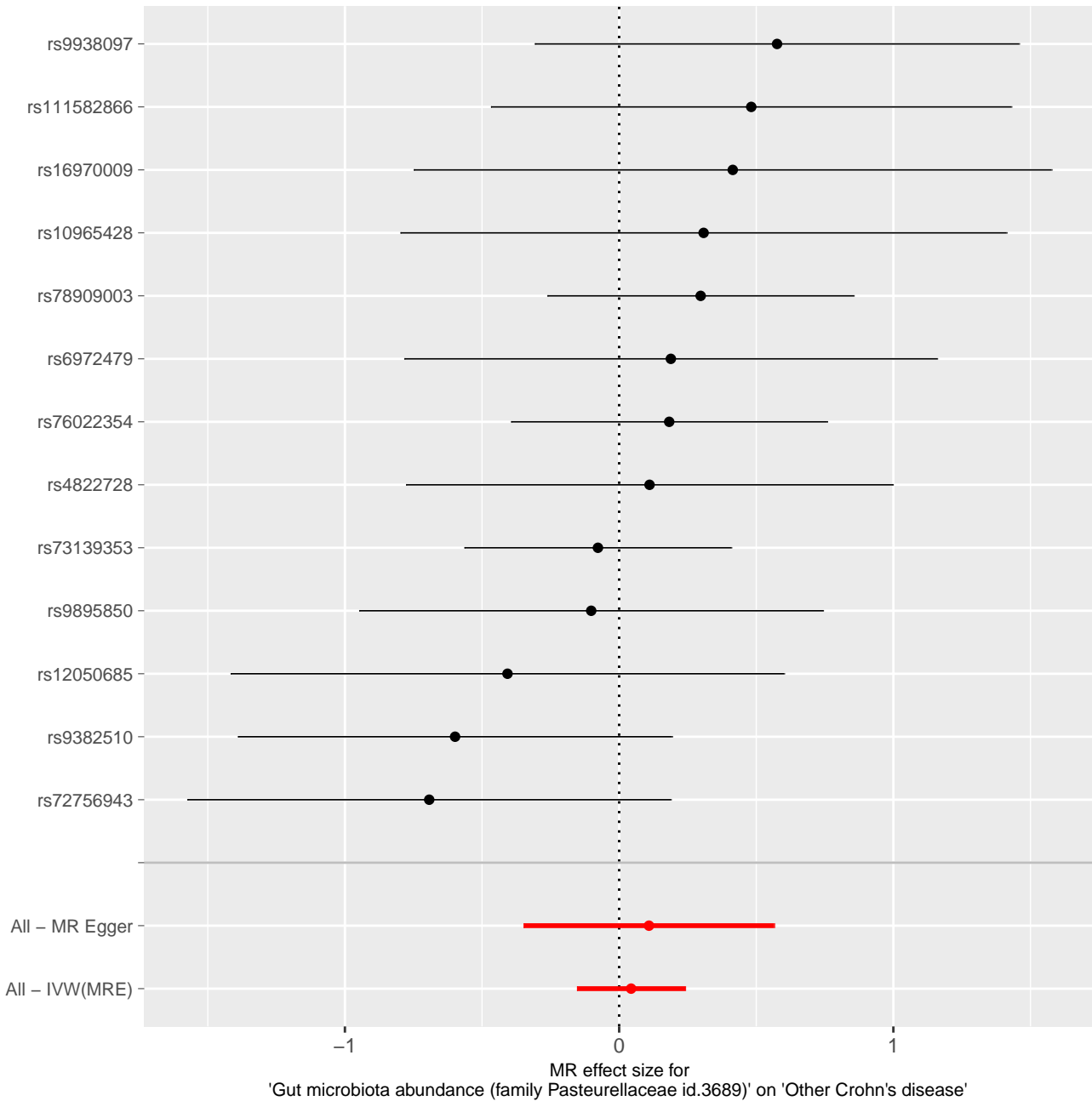
Batch 167 : Gut microbiota abundance (family Lachnospiraceae id.1987) on Other Crohn's disease

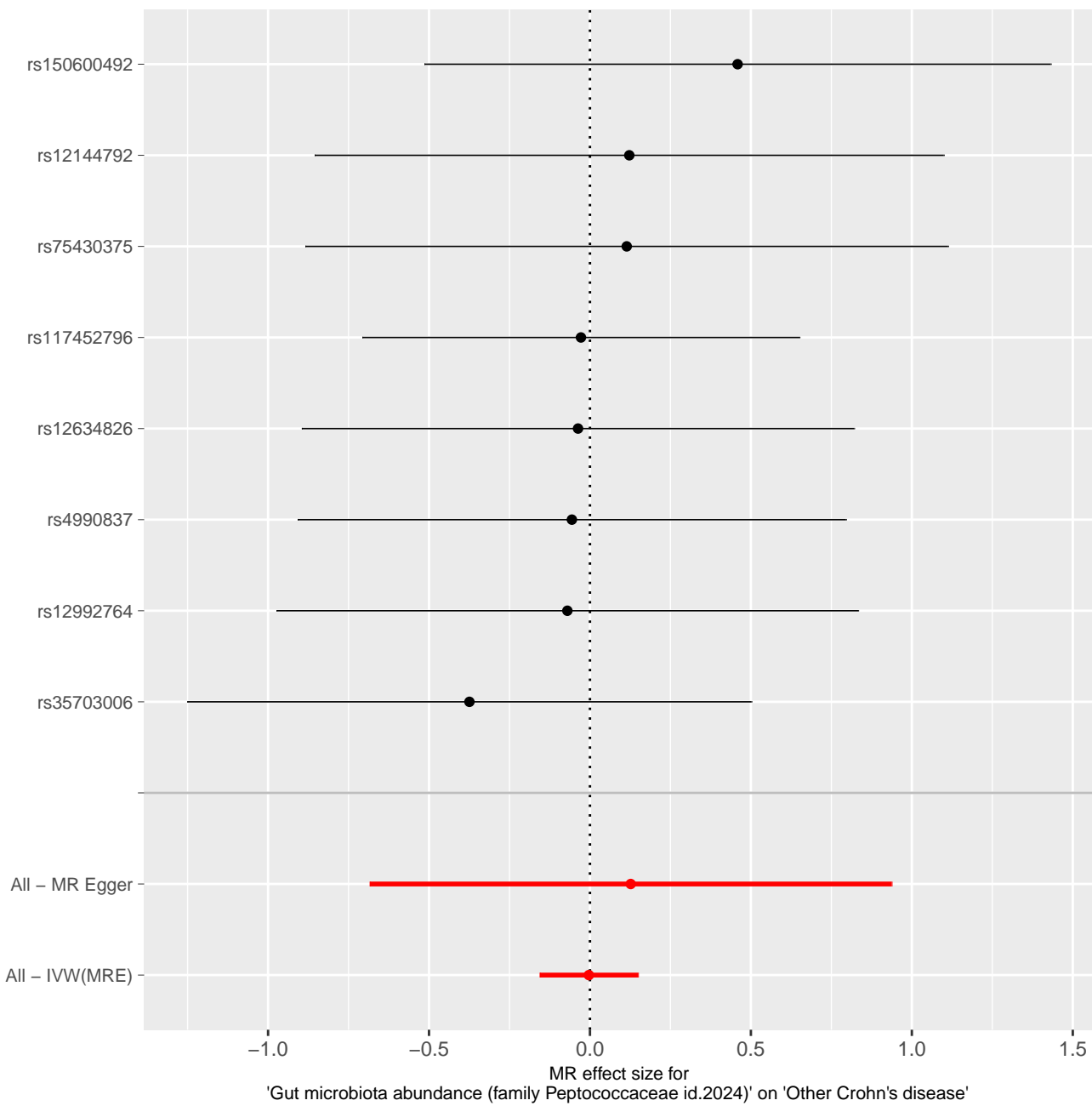




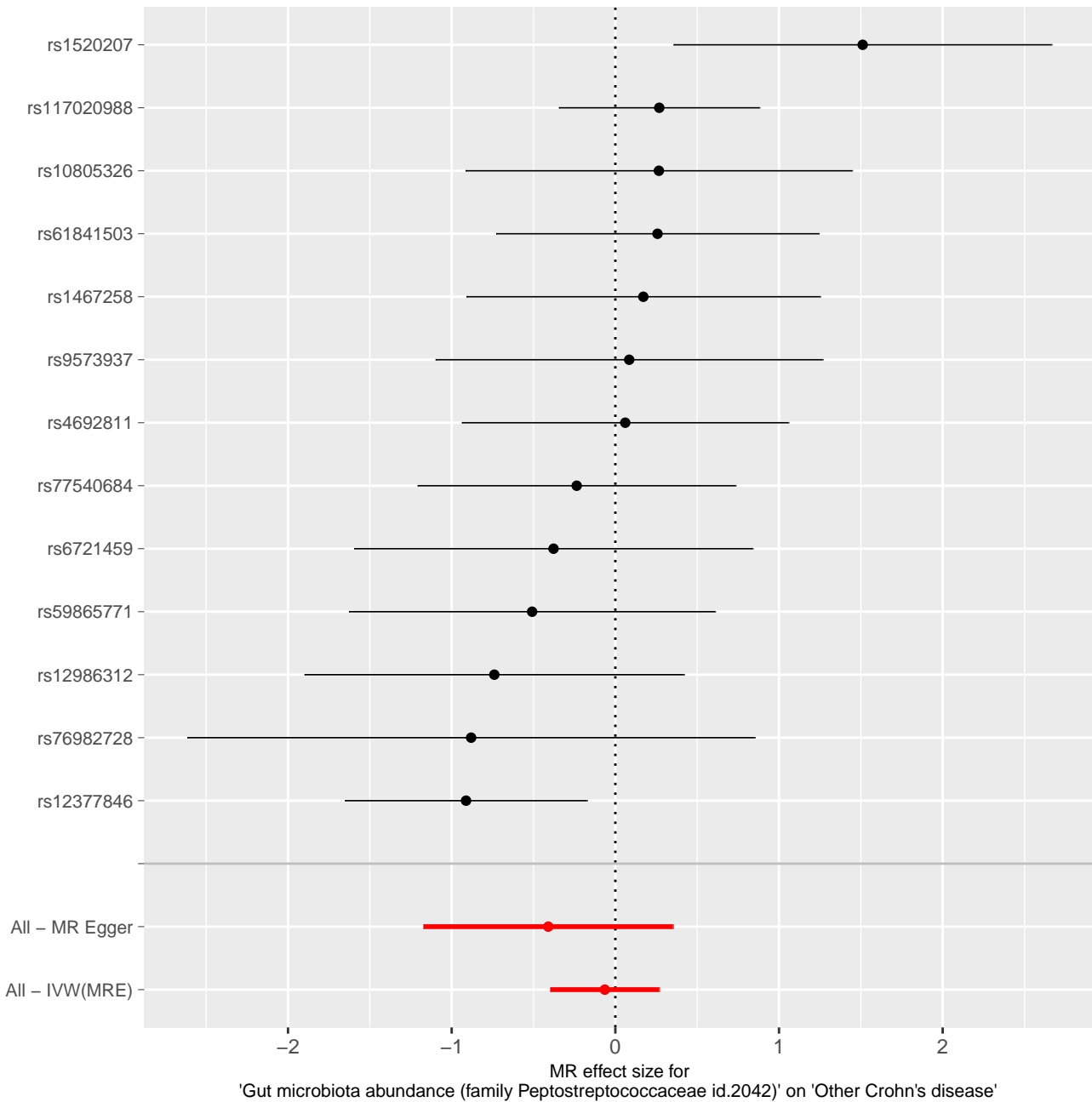




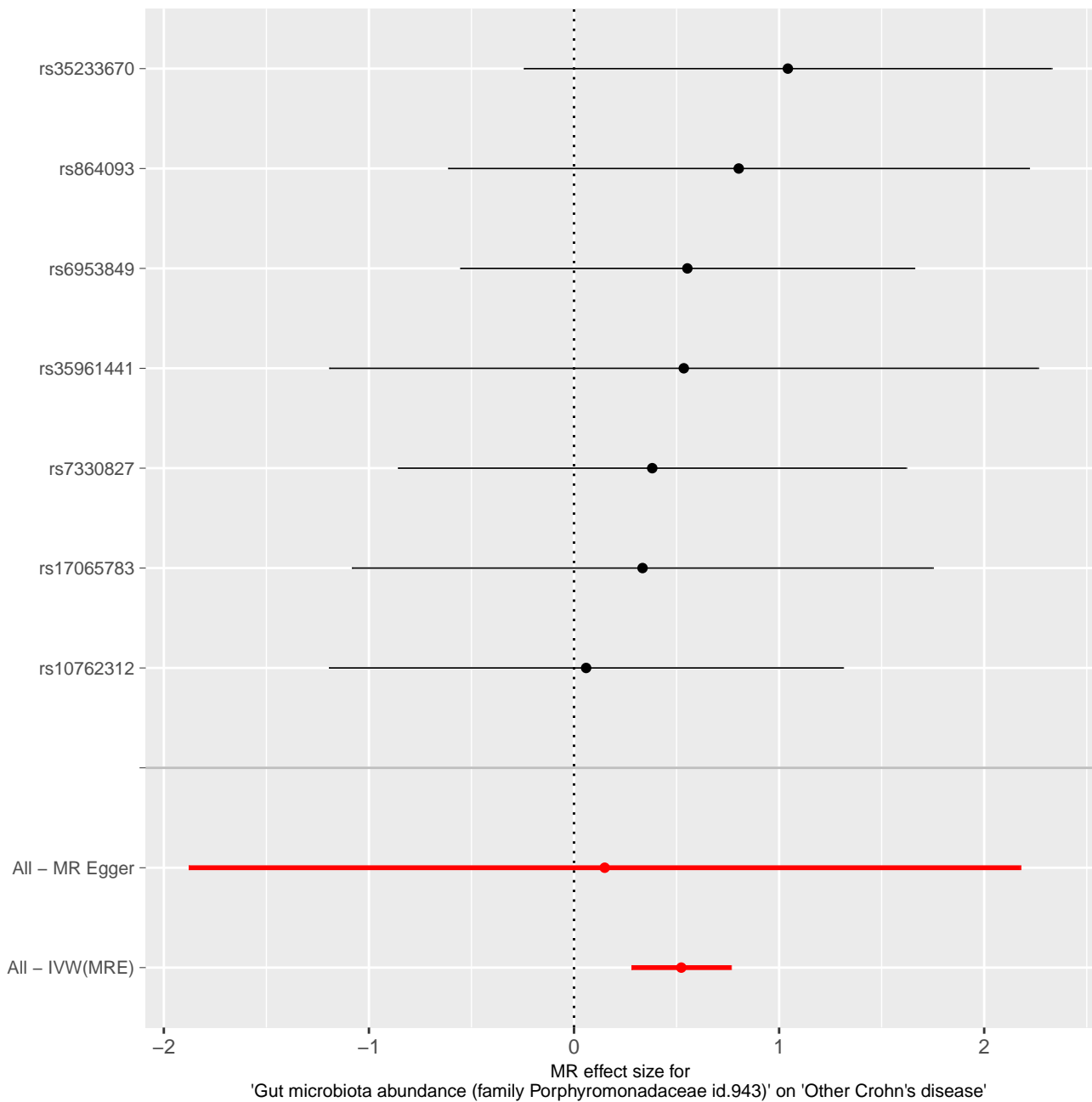




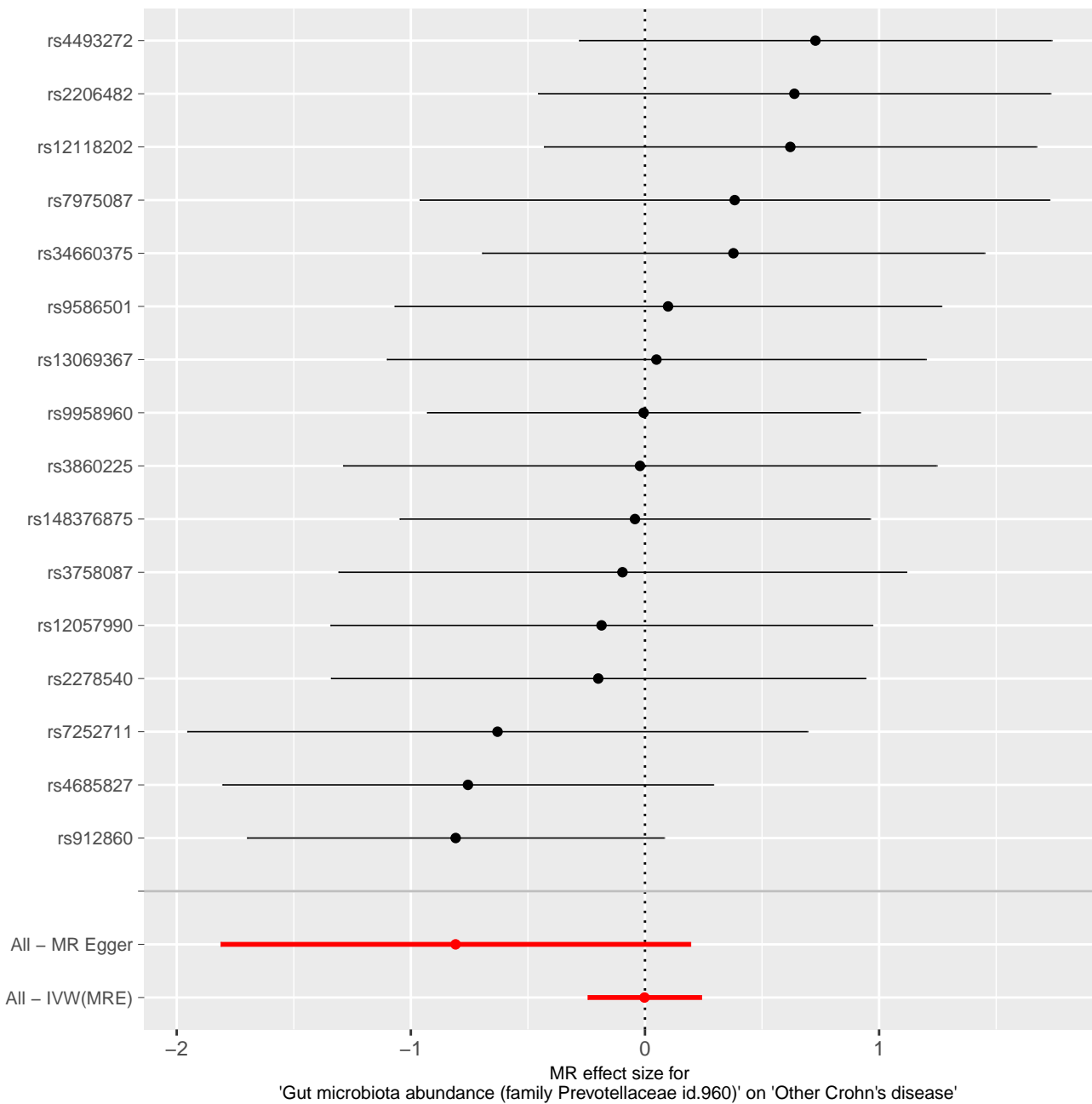
Batch 173 : Gut microbiota abundance (family Peptostreptococcaceae id.2042) on Other Crohn's disease

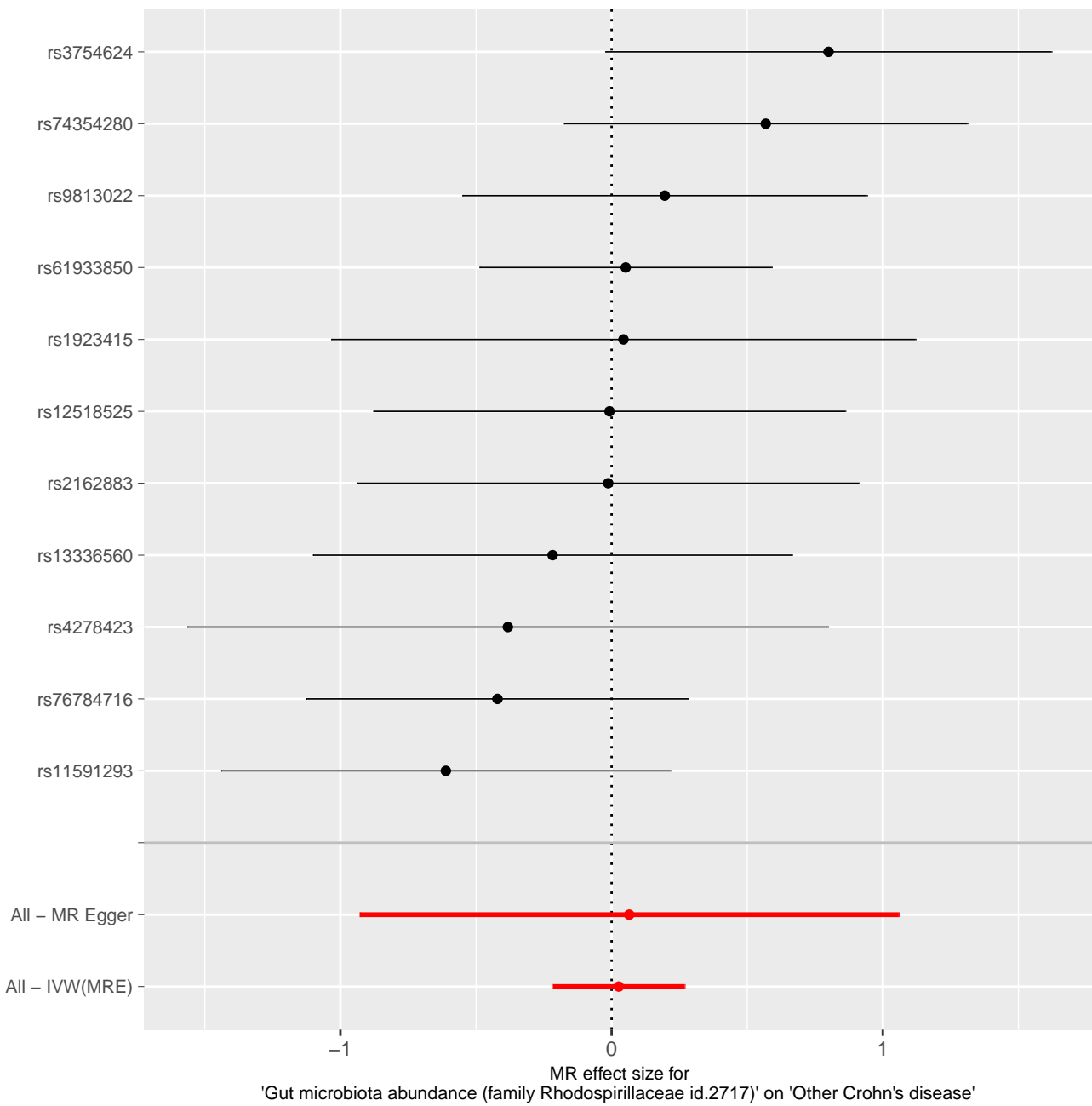


Batch 174 : Gut microbiota abundance (family Porphyromonadaceae id.943) on Other Crohn's disease

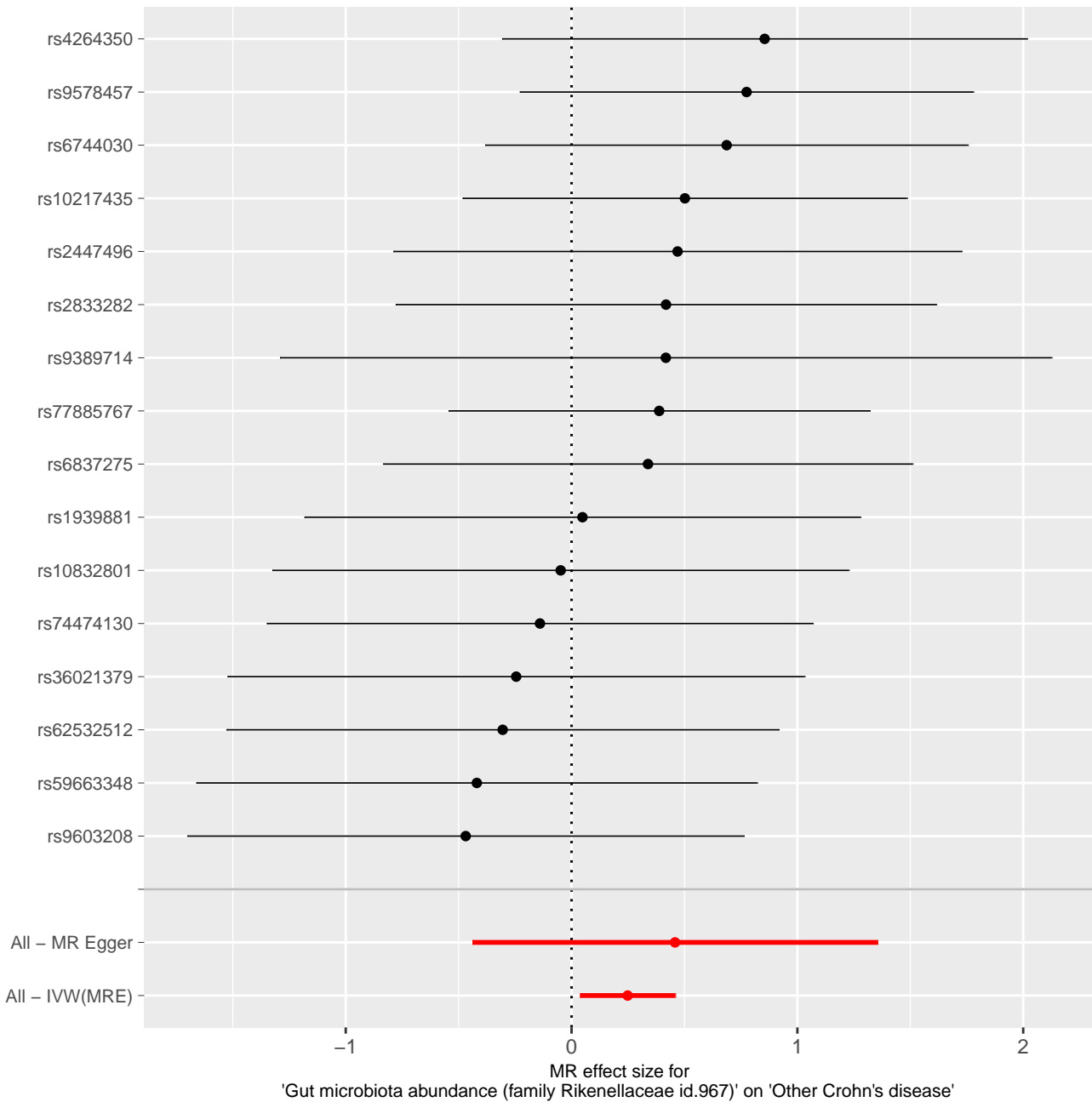


Batch 175 : Gut microbiota abundance (family Prevotellaceae id.960) on Other Crohn's disease

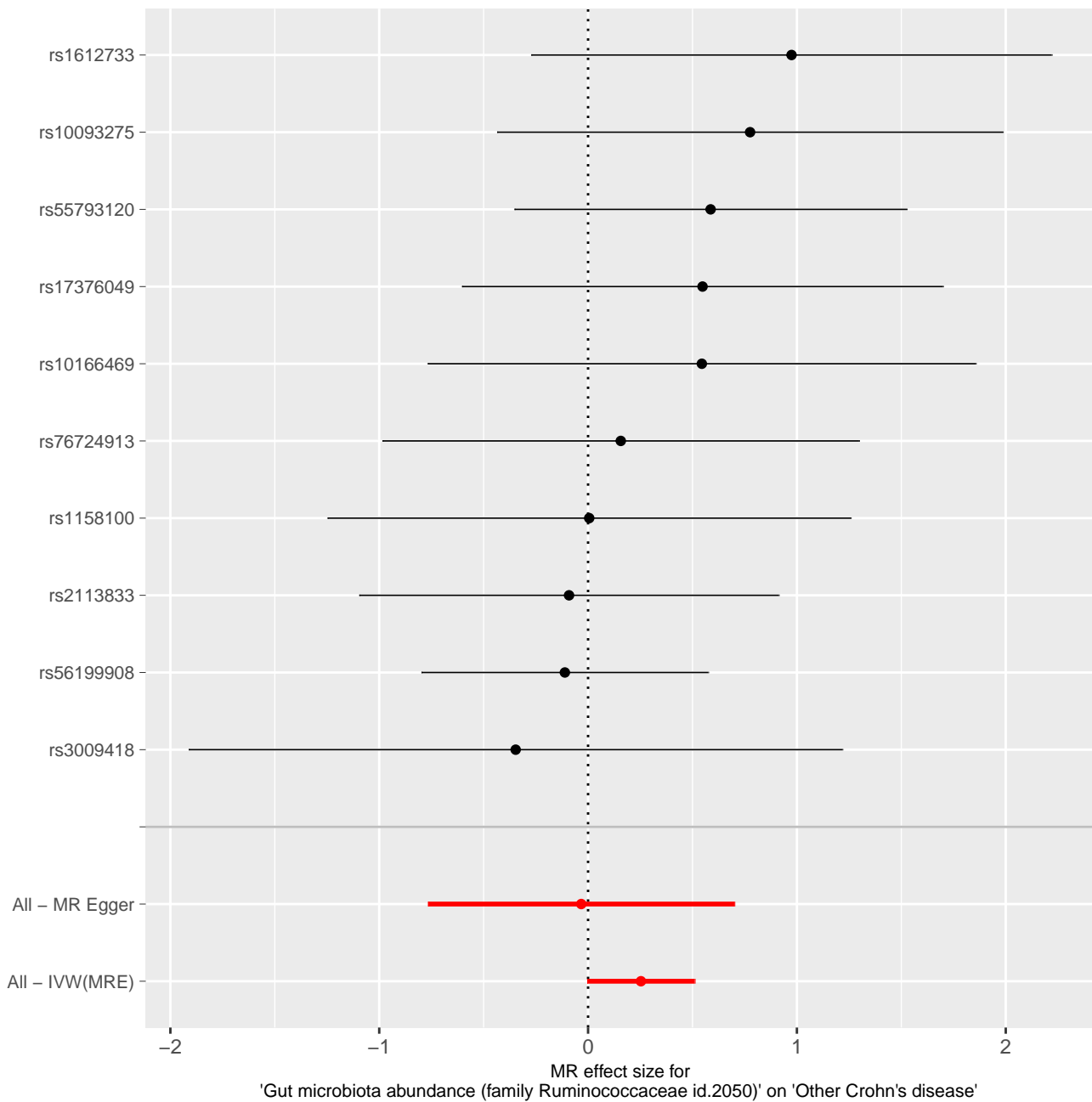


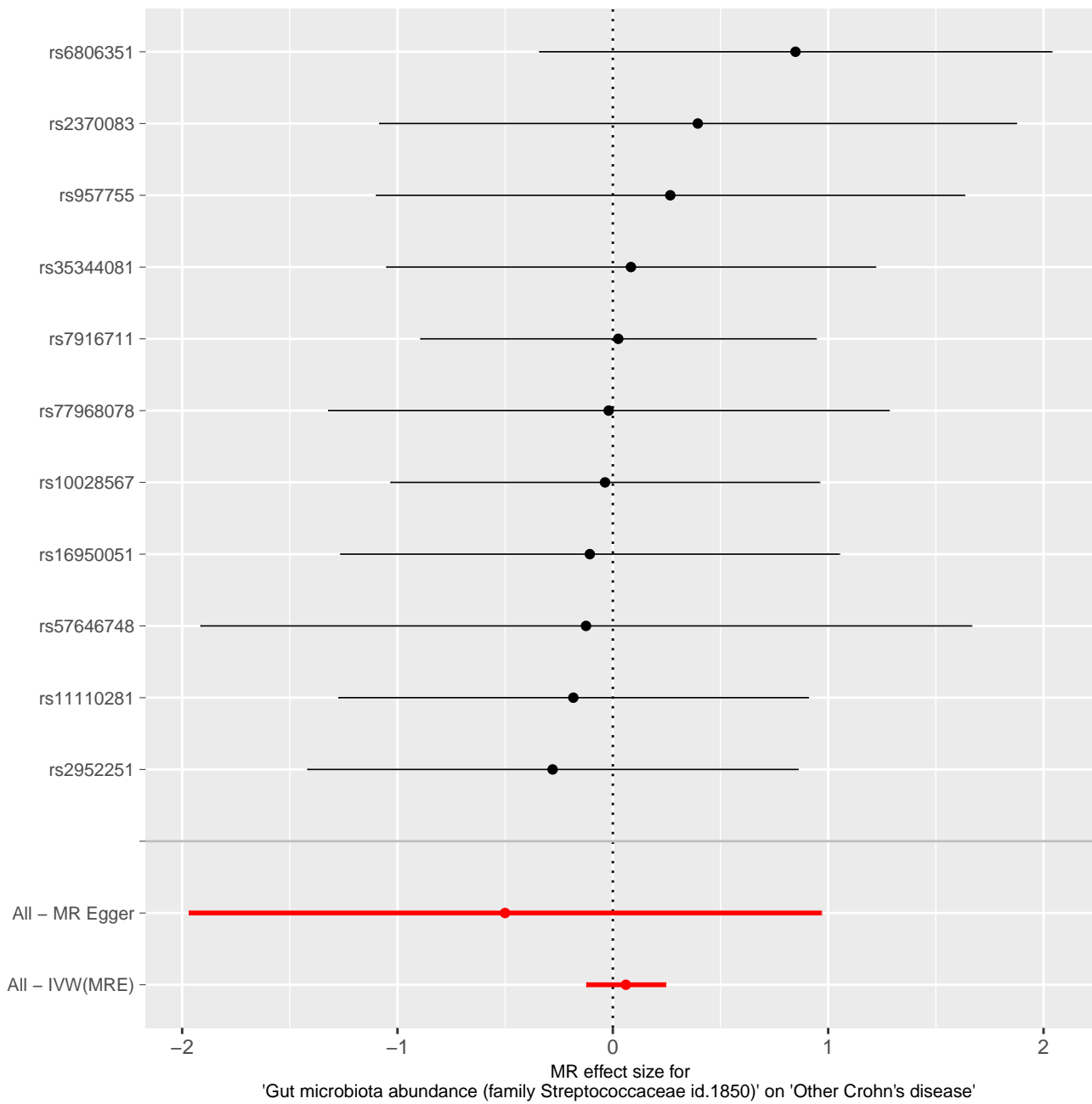


Batch 177 : Gut microbiota abundance (family Rikenellaceae id.967) on Other Crohn's disease

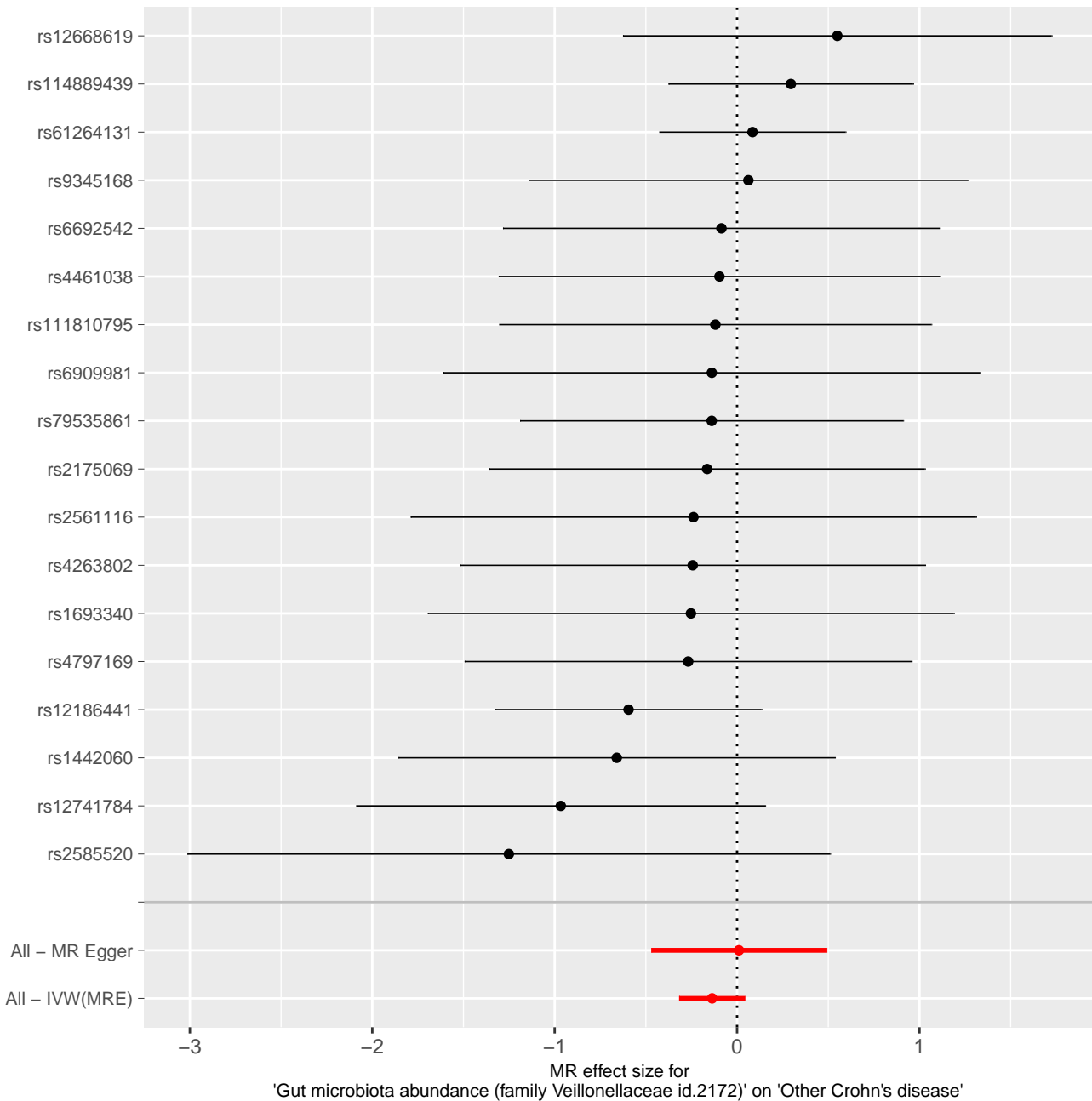


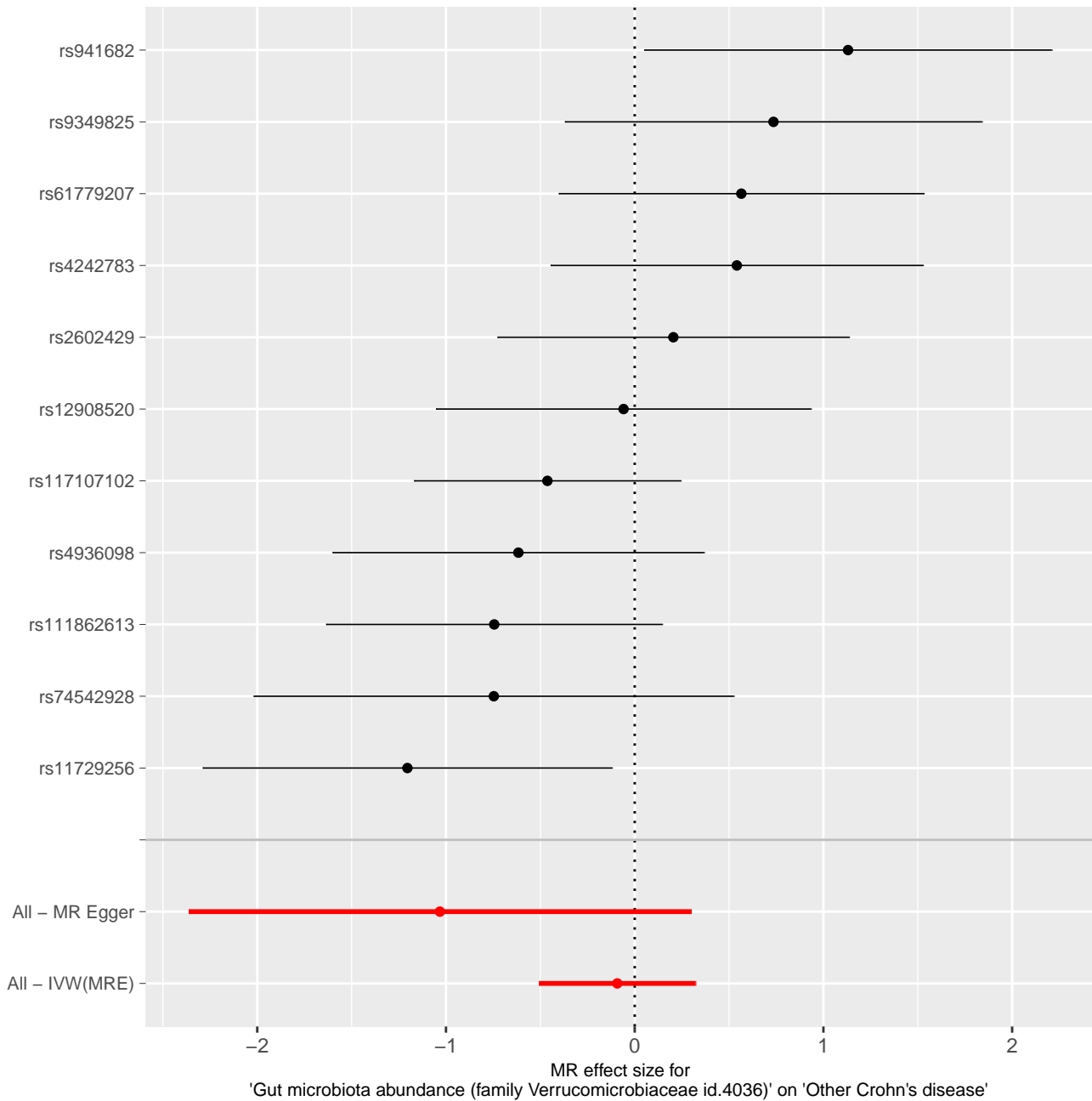
Batch 178 : Gut microbiota abundance (family Ruminococcaceae id.2050) on Other Crohn's disease



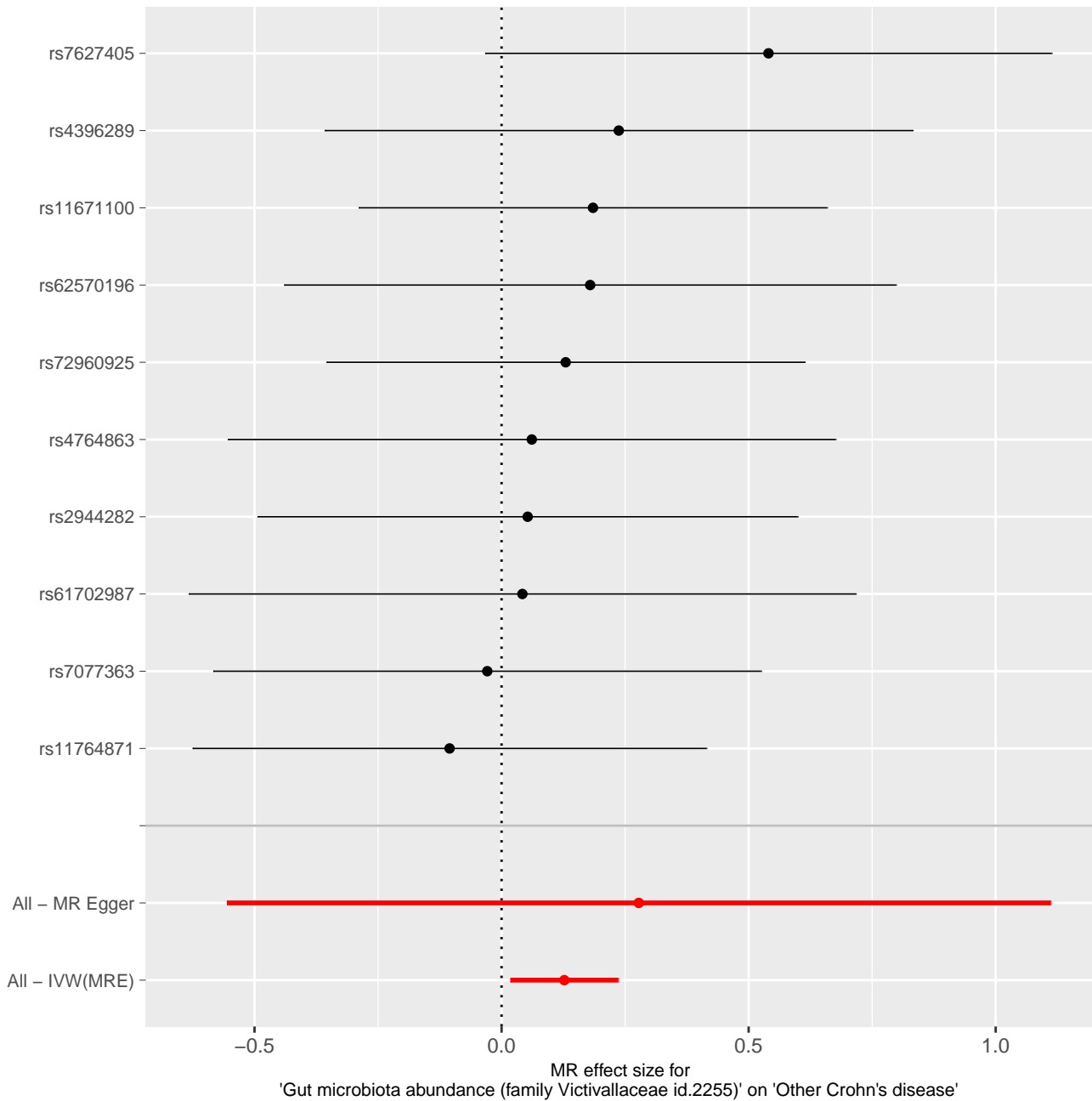


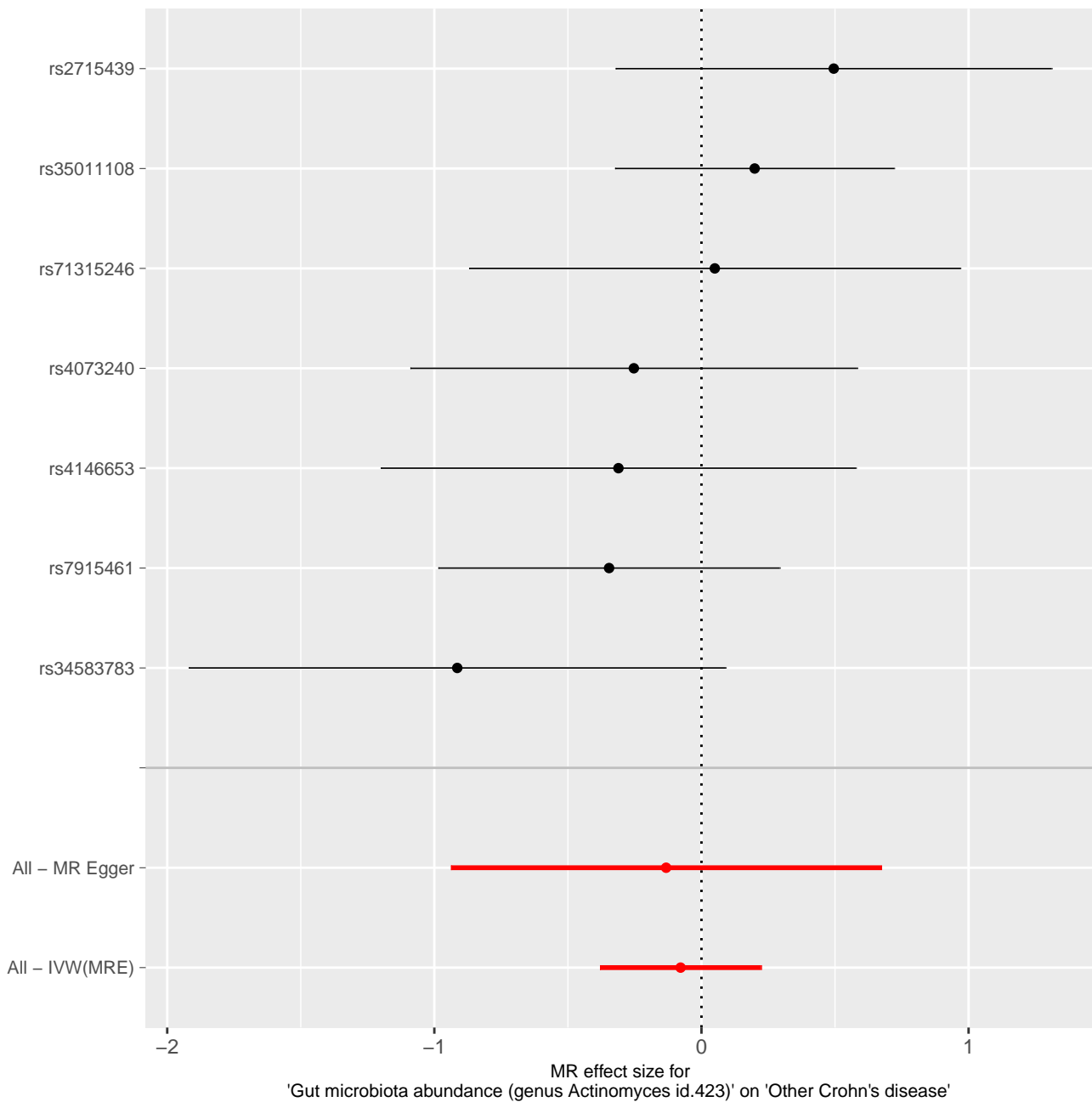
Batch 180 : Gut microbiota abundance (family Veillonellaceae id.2172) on Other Crohn's disease

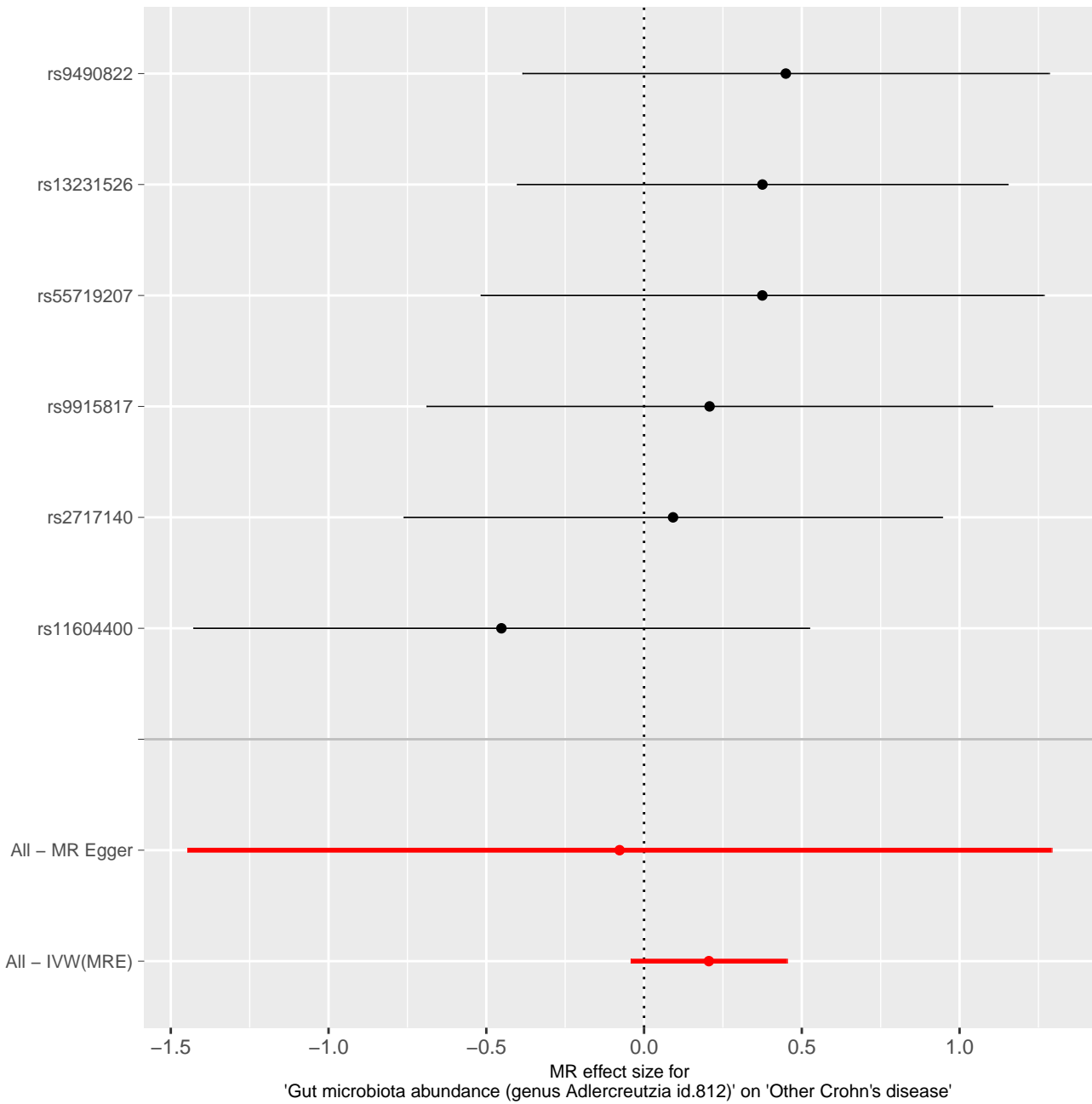




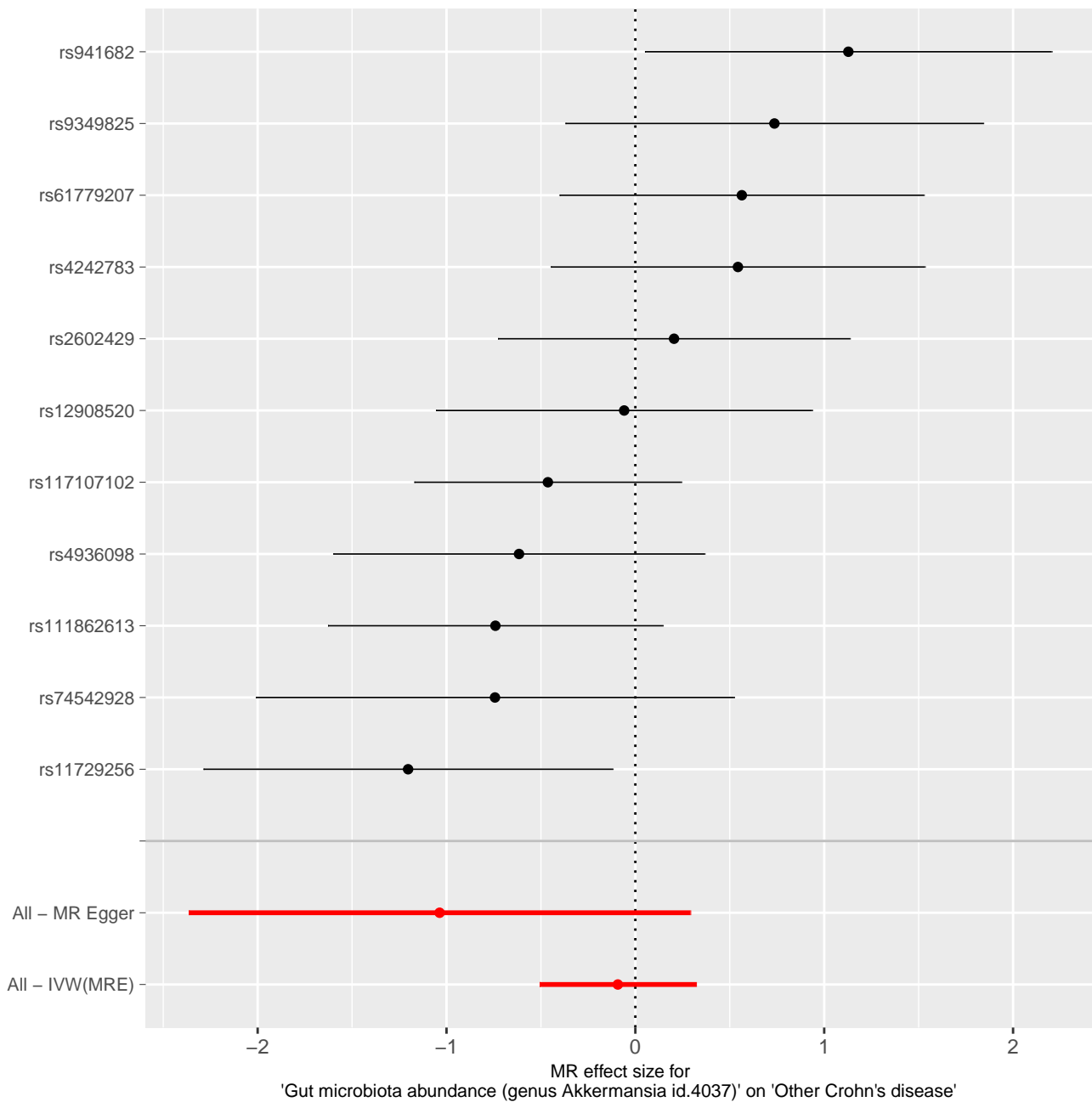
Batch 182 : Gut microbiota abundance (family Victivallaceae id.2255) on Other Crohn's disease



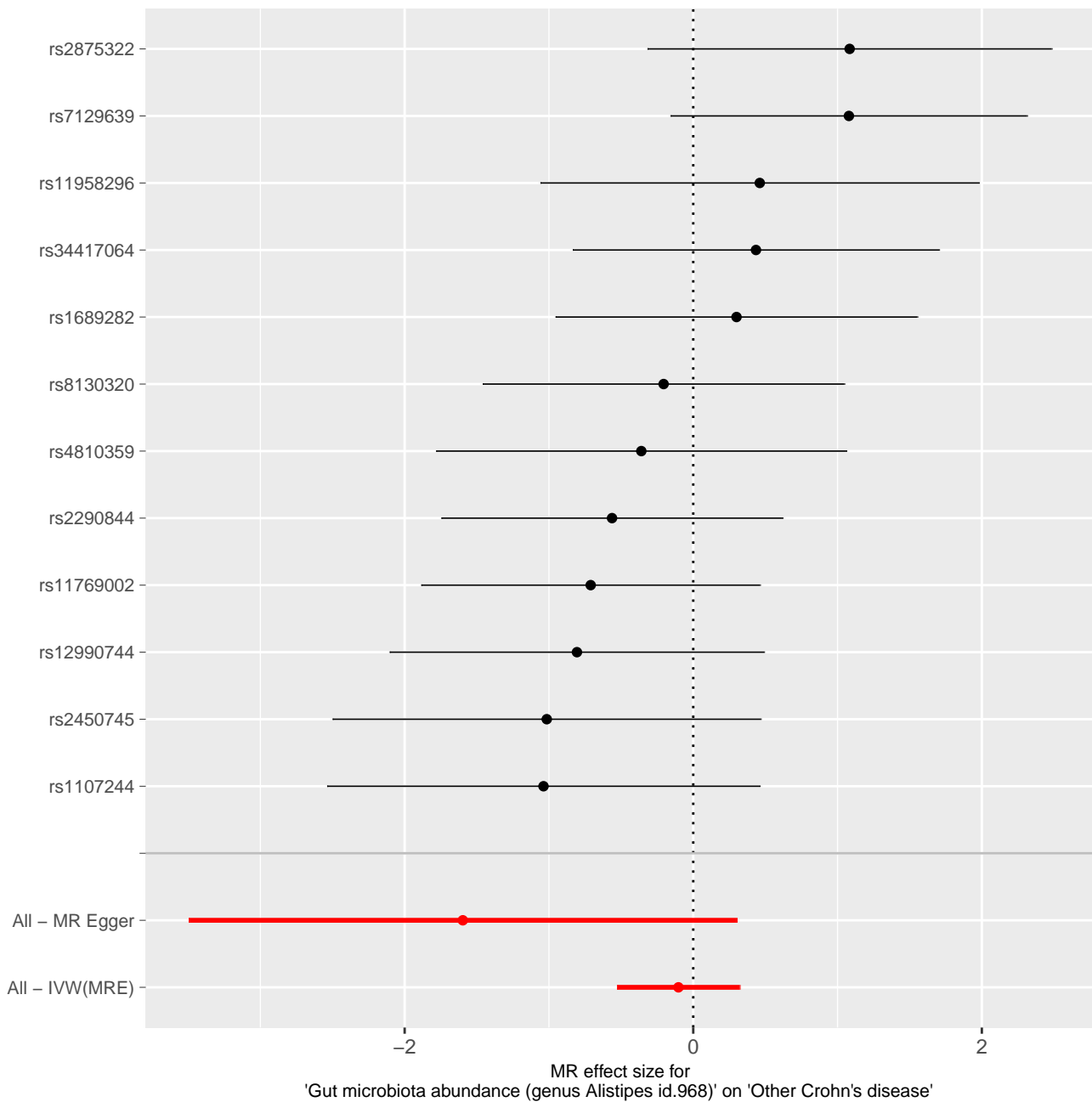




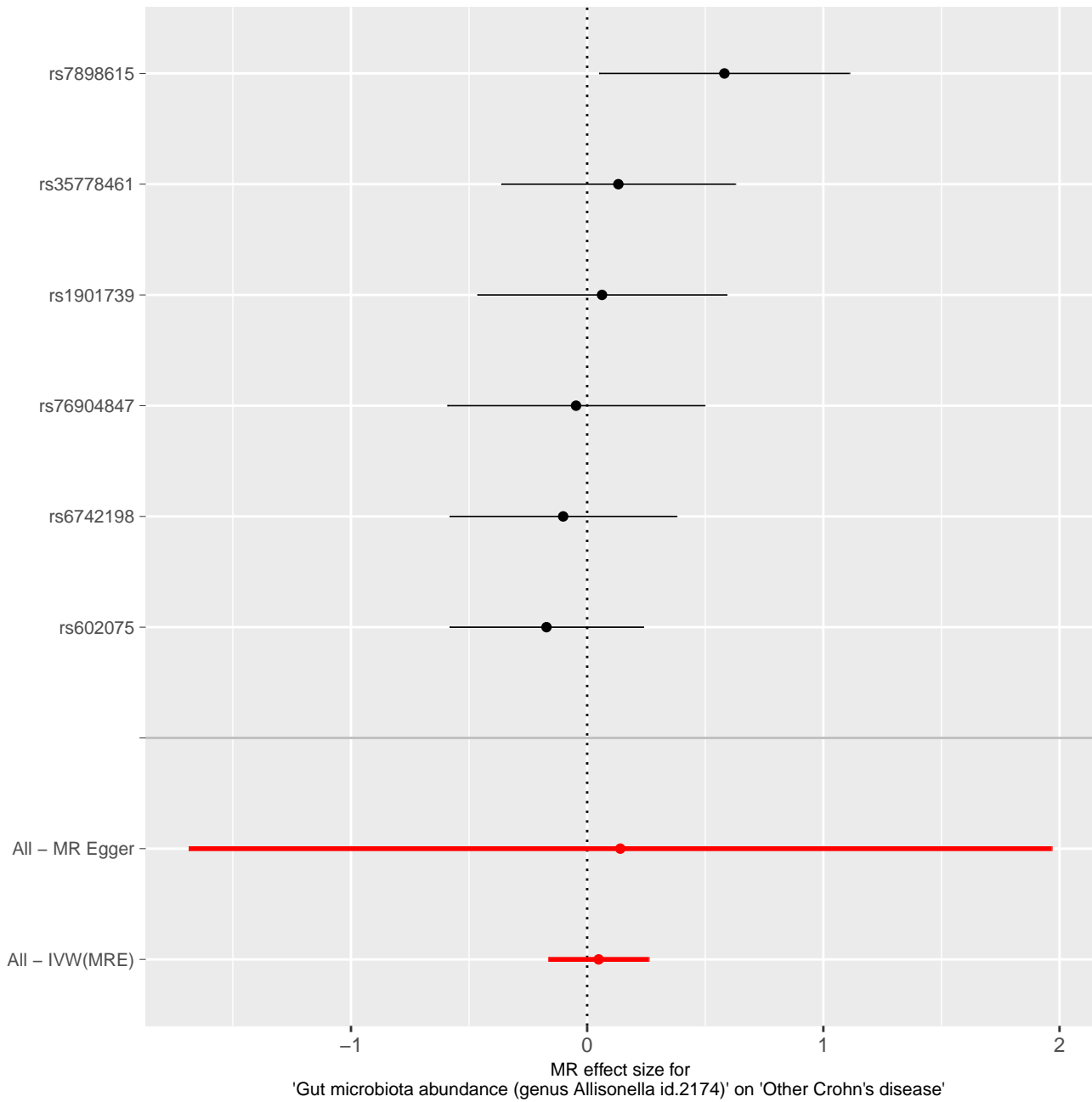
Batch 185 : Gut microbiota abundance (genus Akkermansia id.4037) on Other Crohn's disease

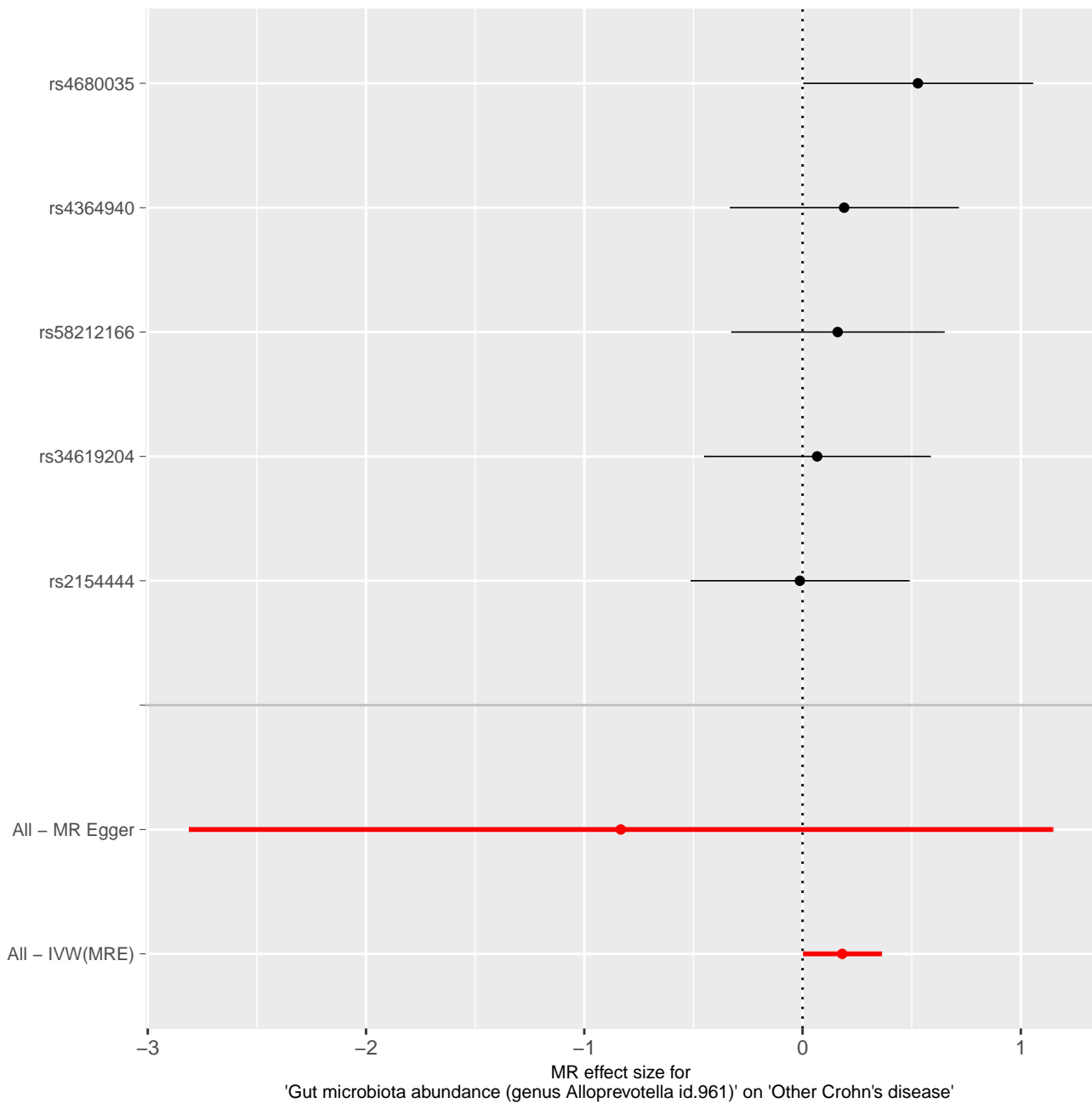


Batch 186 : Gut microbiota abundance (genus Alistipes id.968) on Other Crohn's disease

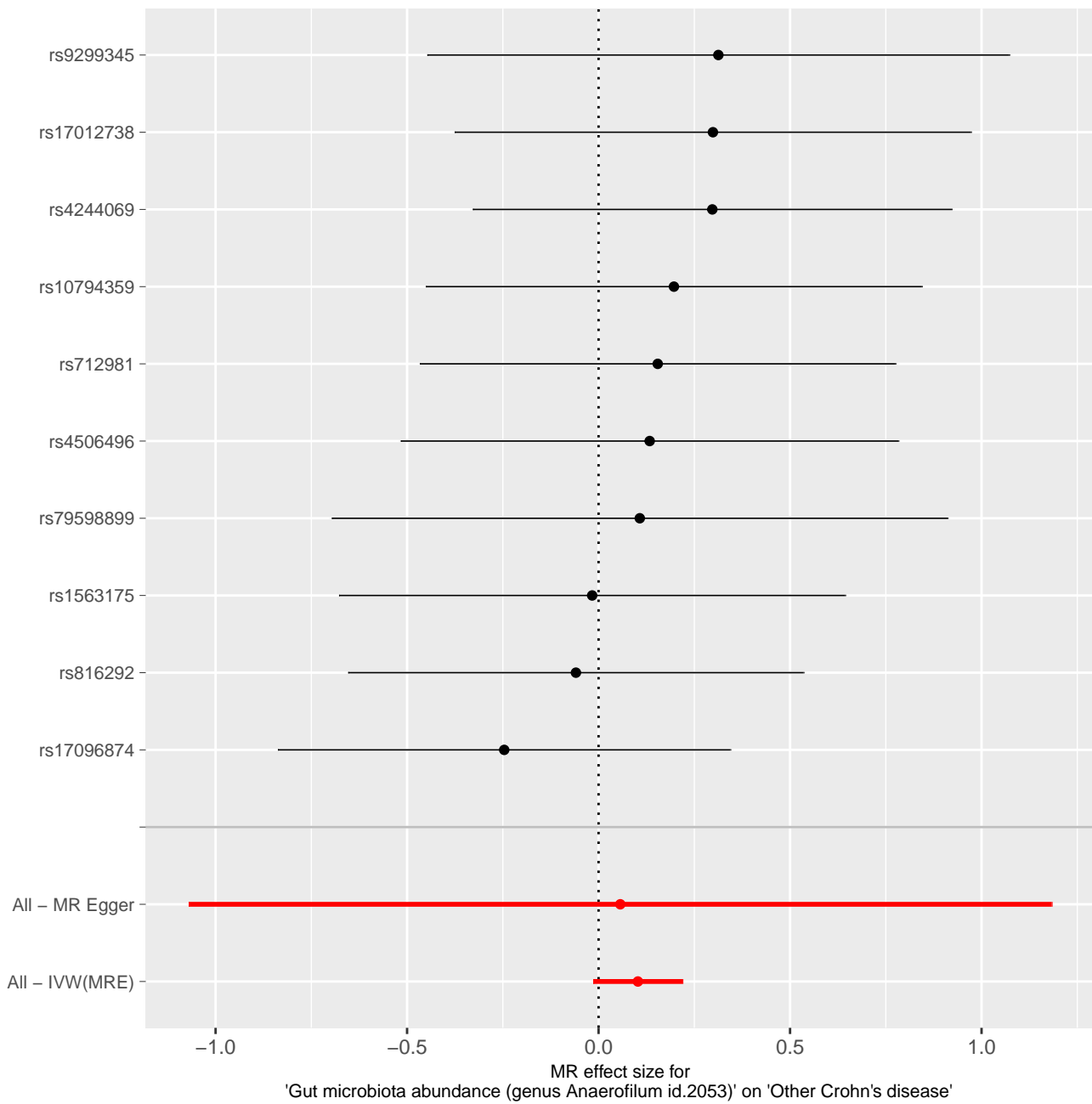


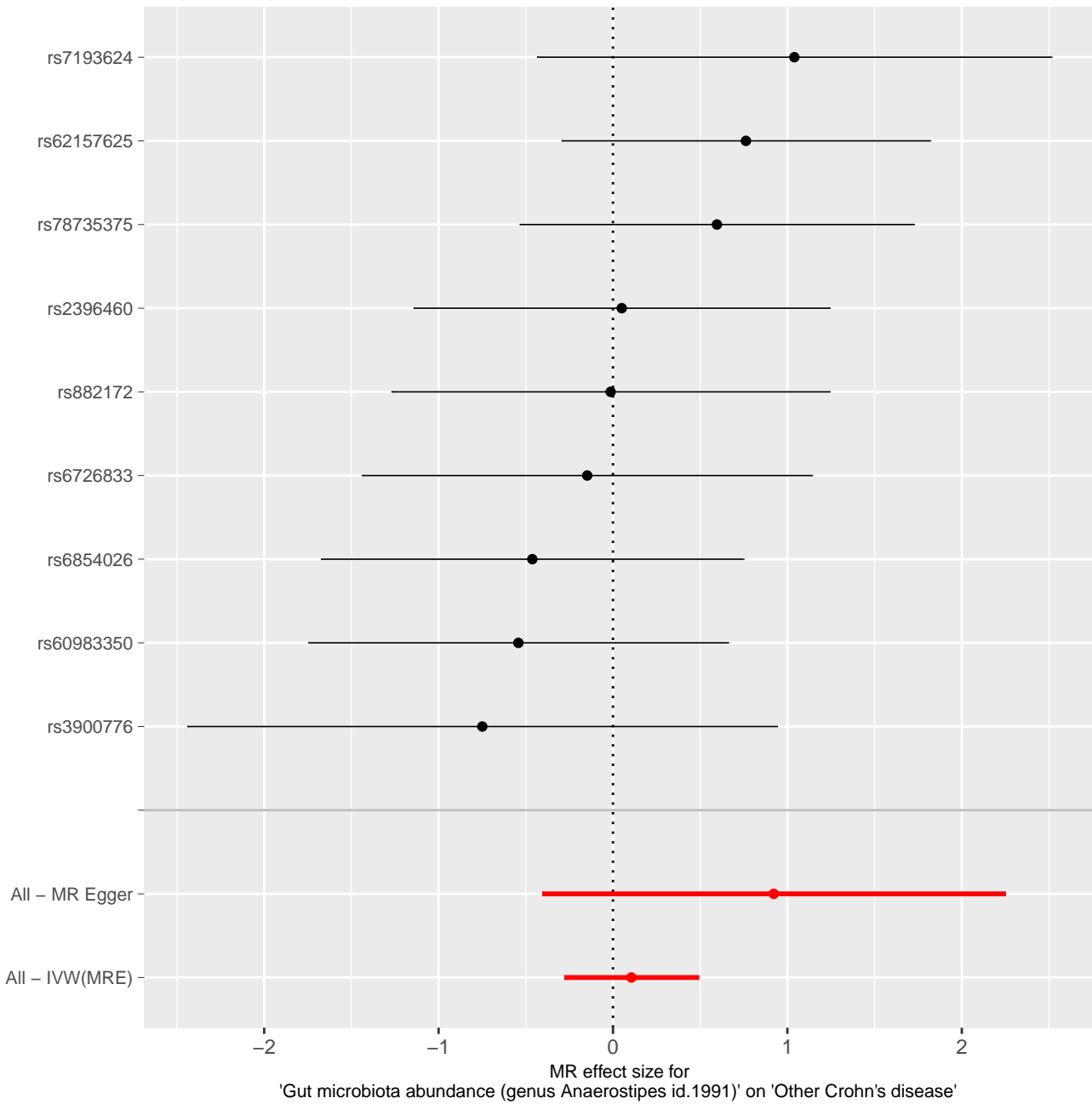
Batch 187 : Gut microbiota abundance (genus Allisonella id.2174) on Other Crohn's disease



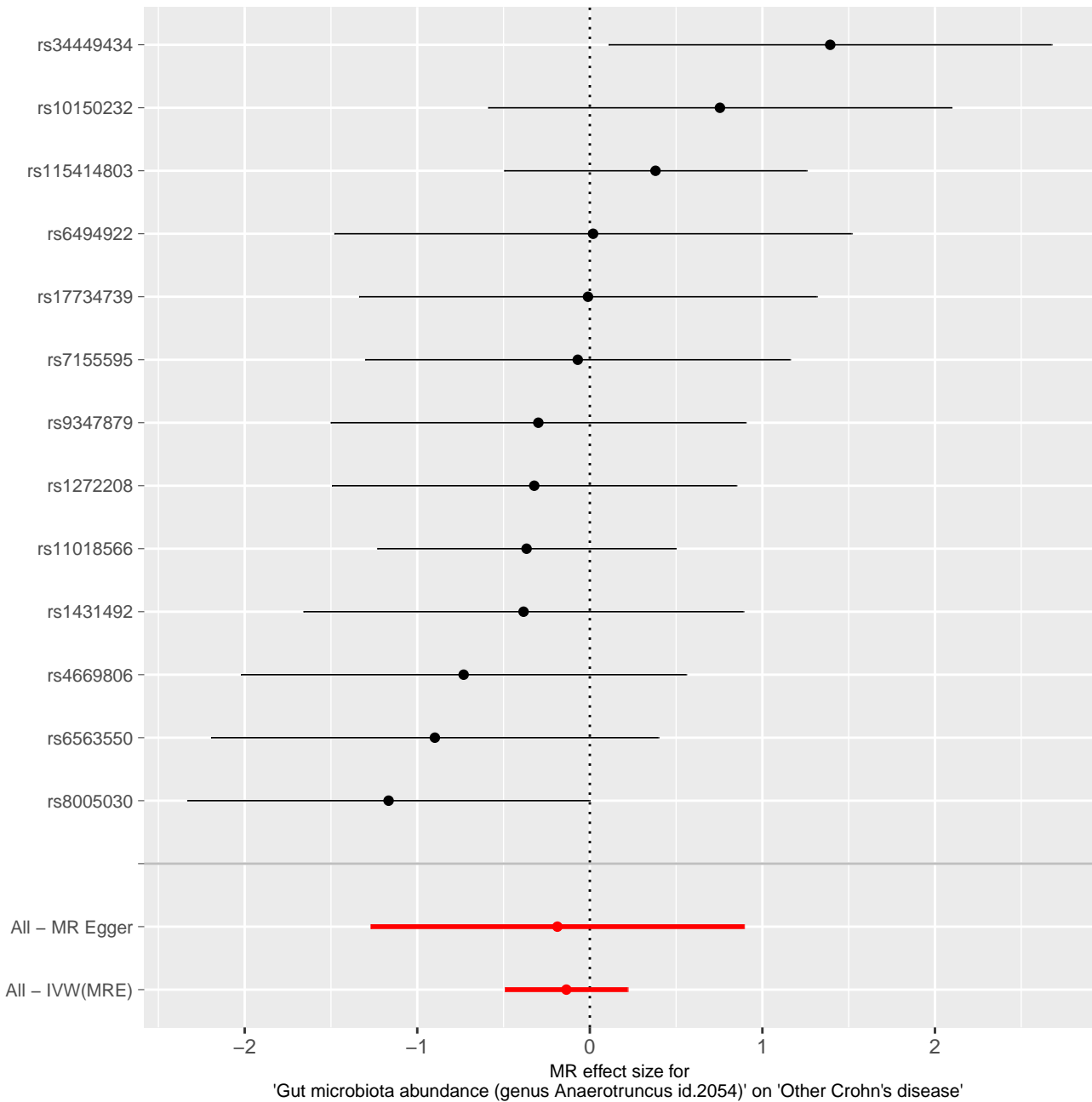


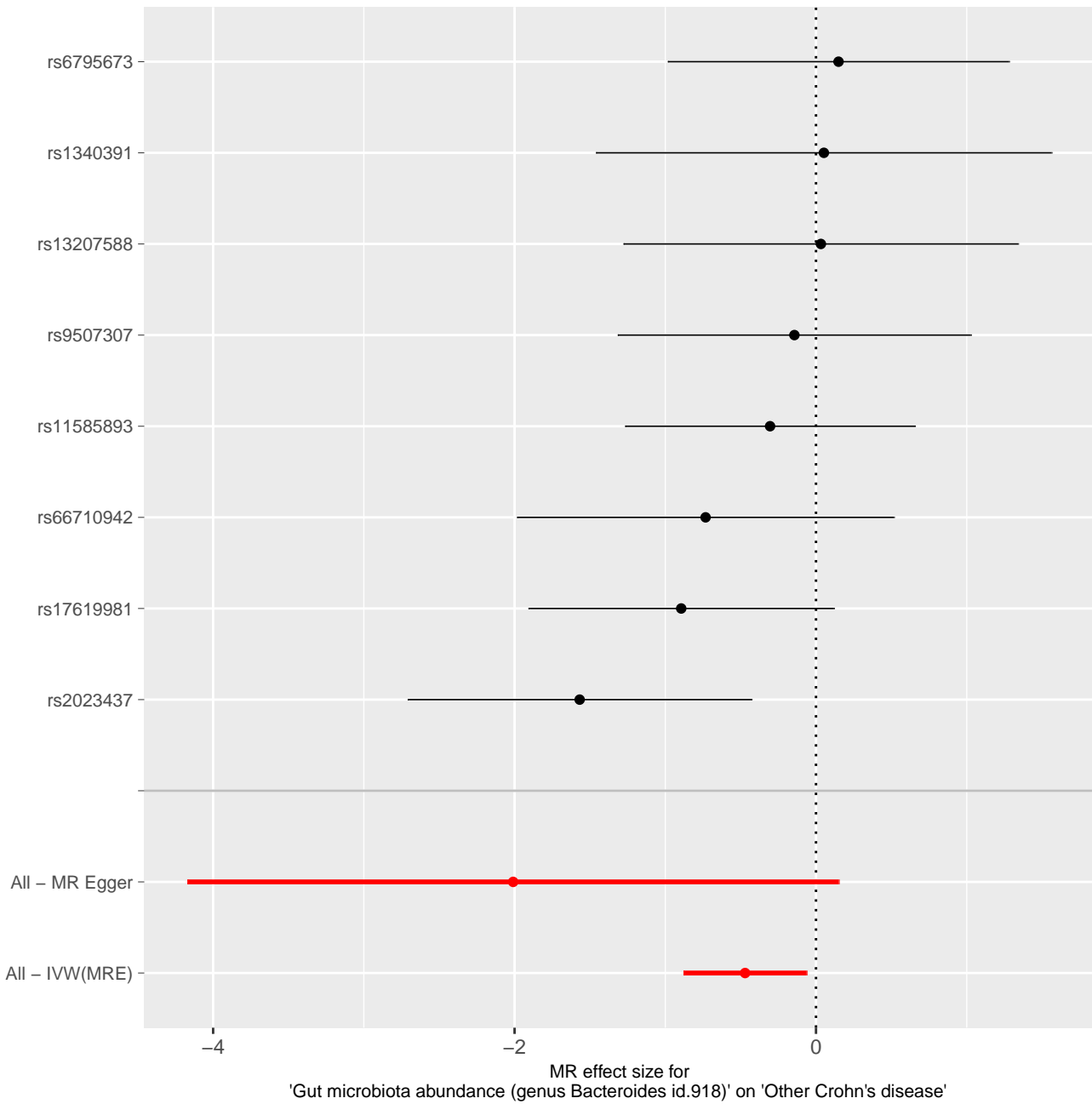
Batch 189 : Gut microbiota abundance (genus Anaerofilum id.2053) on Other Crohn's disease



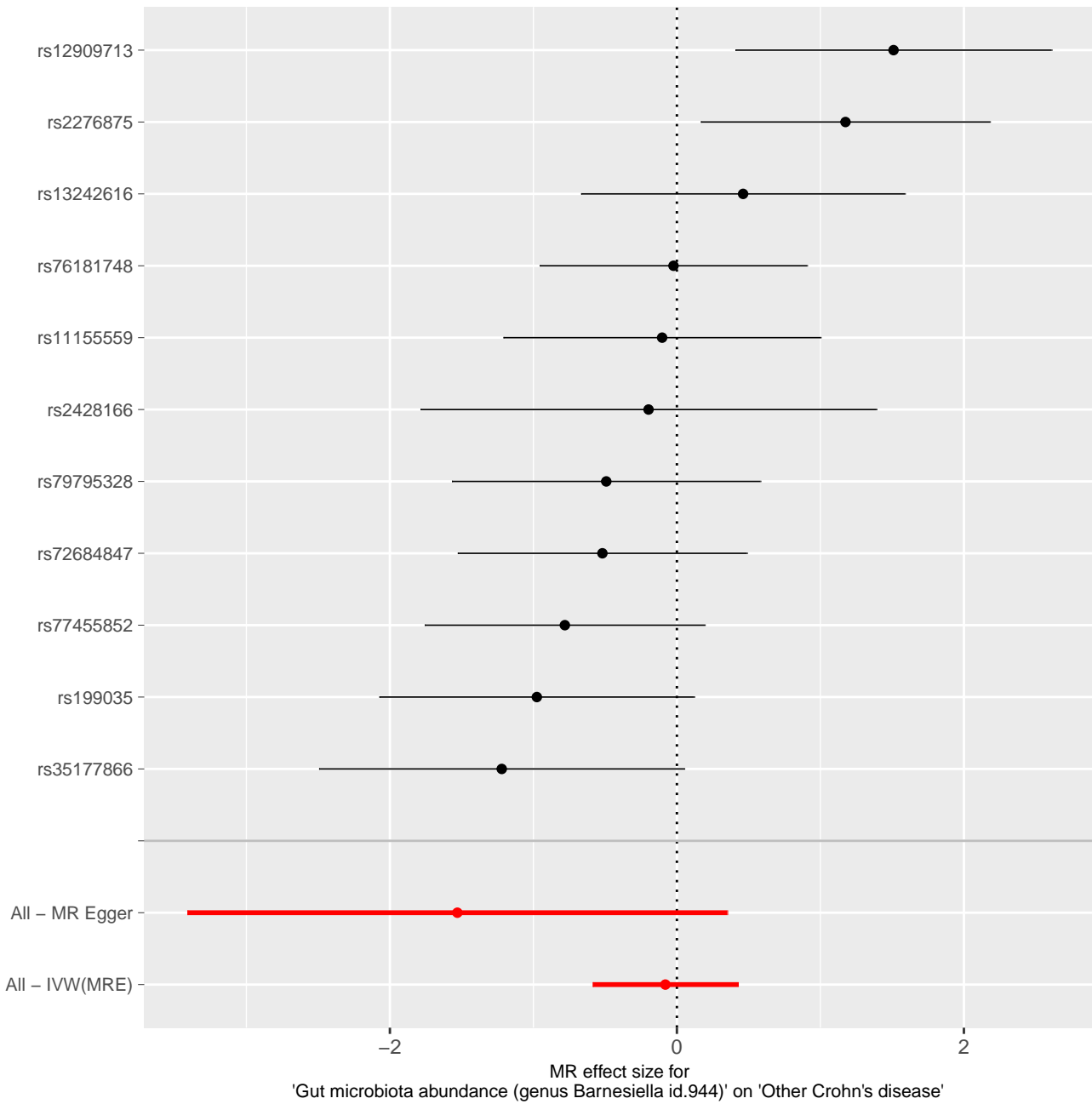


Batch 191 : Gut microbiota abundance (genus Anaerotruncus id.2054) on Other Crohn's disease

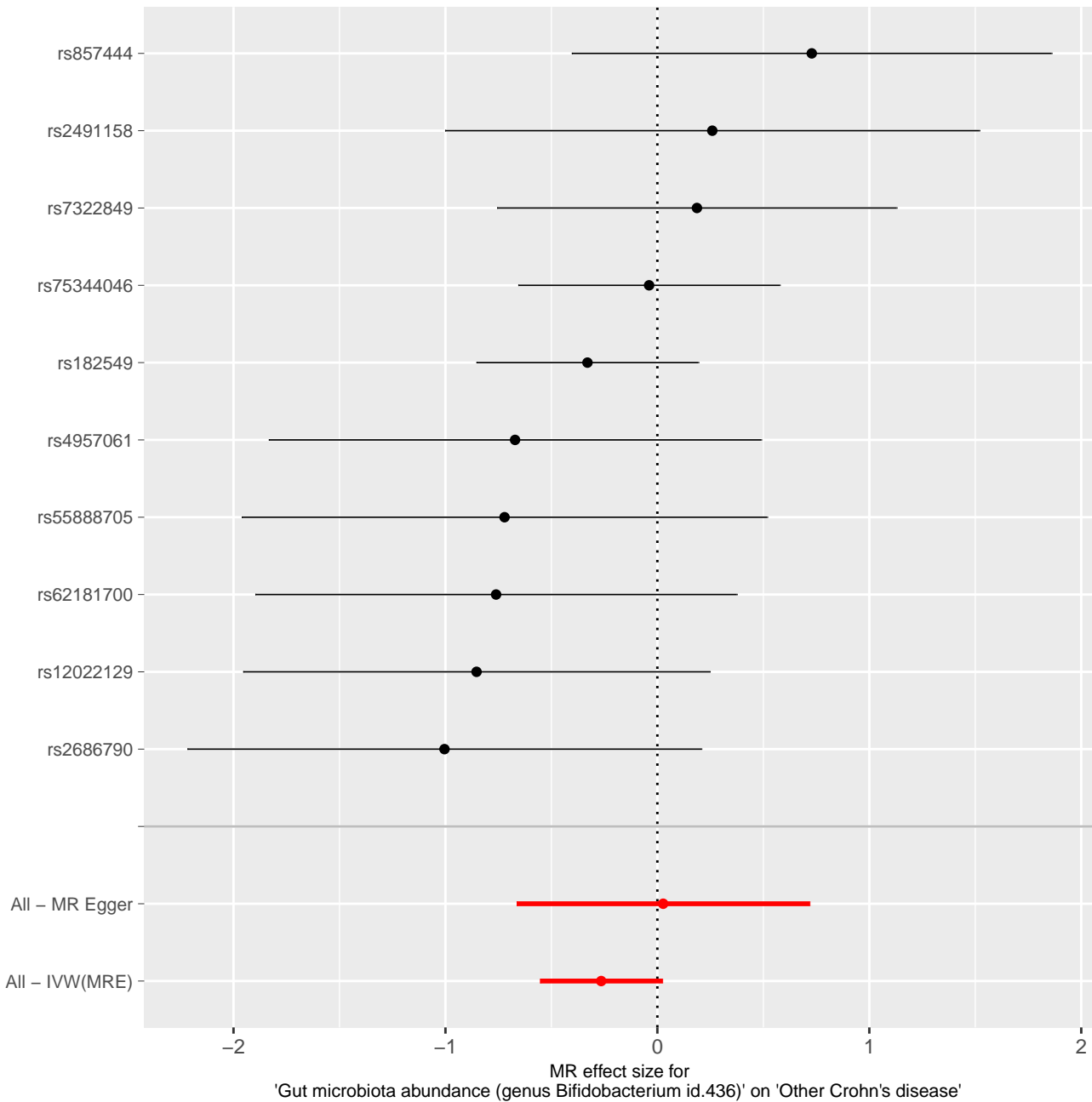




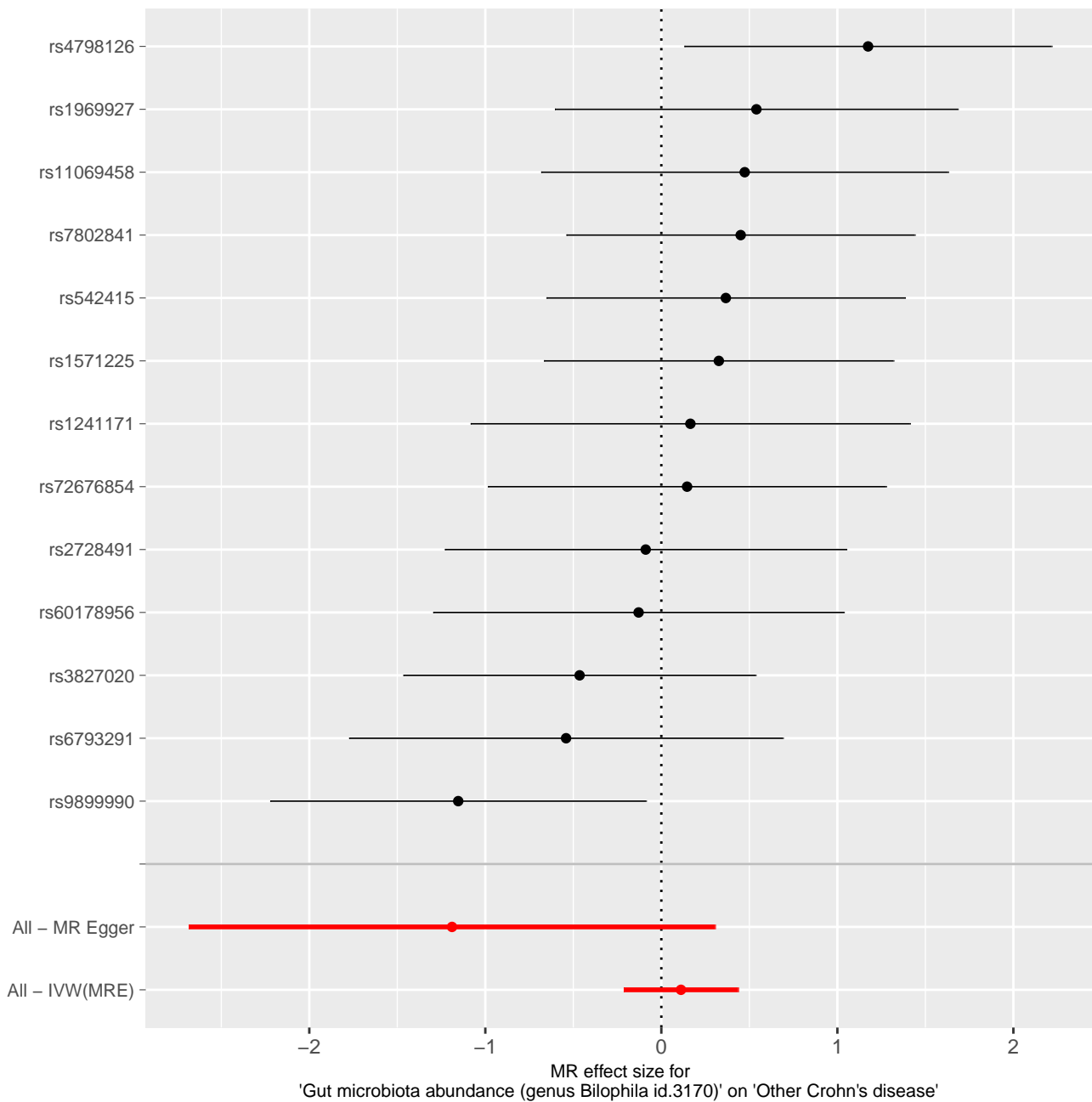
Batch 193 : Gut microbiota abundance (genus *Barnesiella* id.944) on Other Crohn's disease

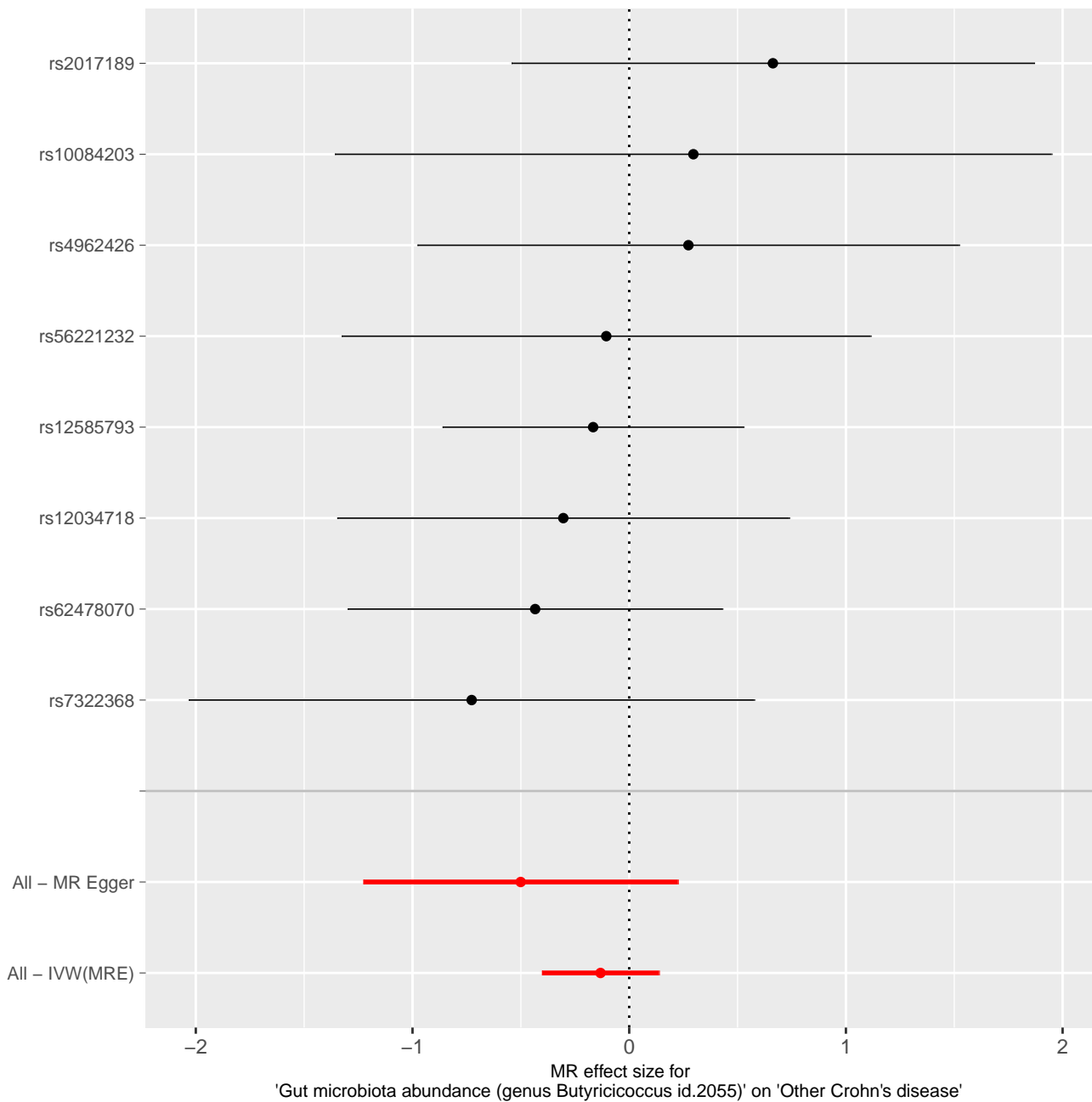


Batch 194 : Gut microbiota abundance (genus Bifidobacterium id.436) on Other Crohn's disease

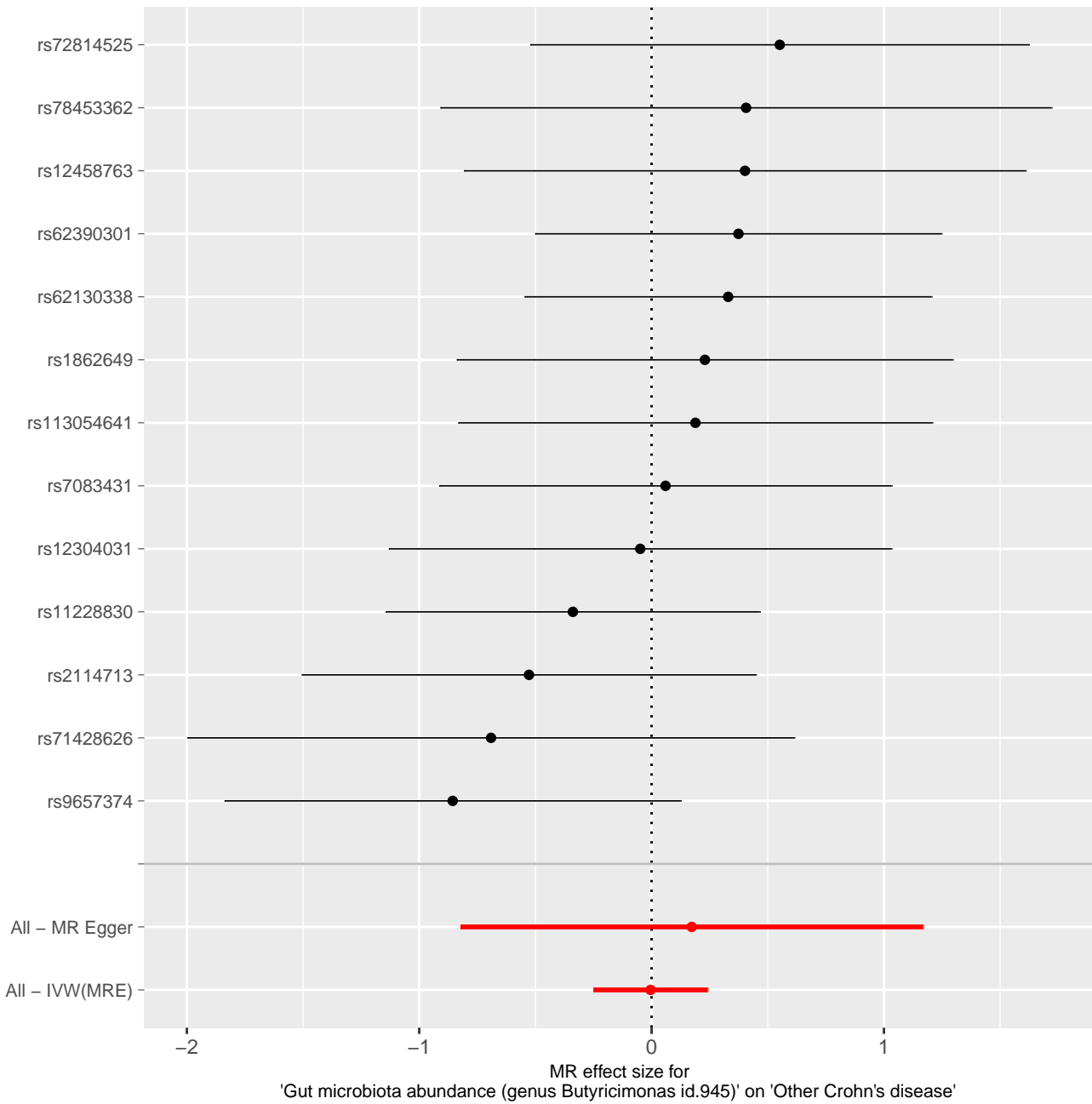


Batch 195 : Gut microbiota abundance (genus Bilophila id.3170) on Other Crohn's disease

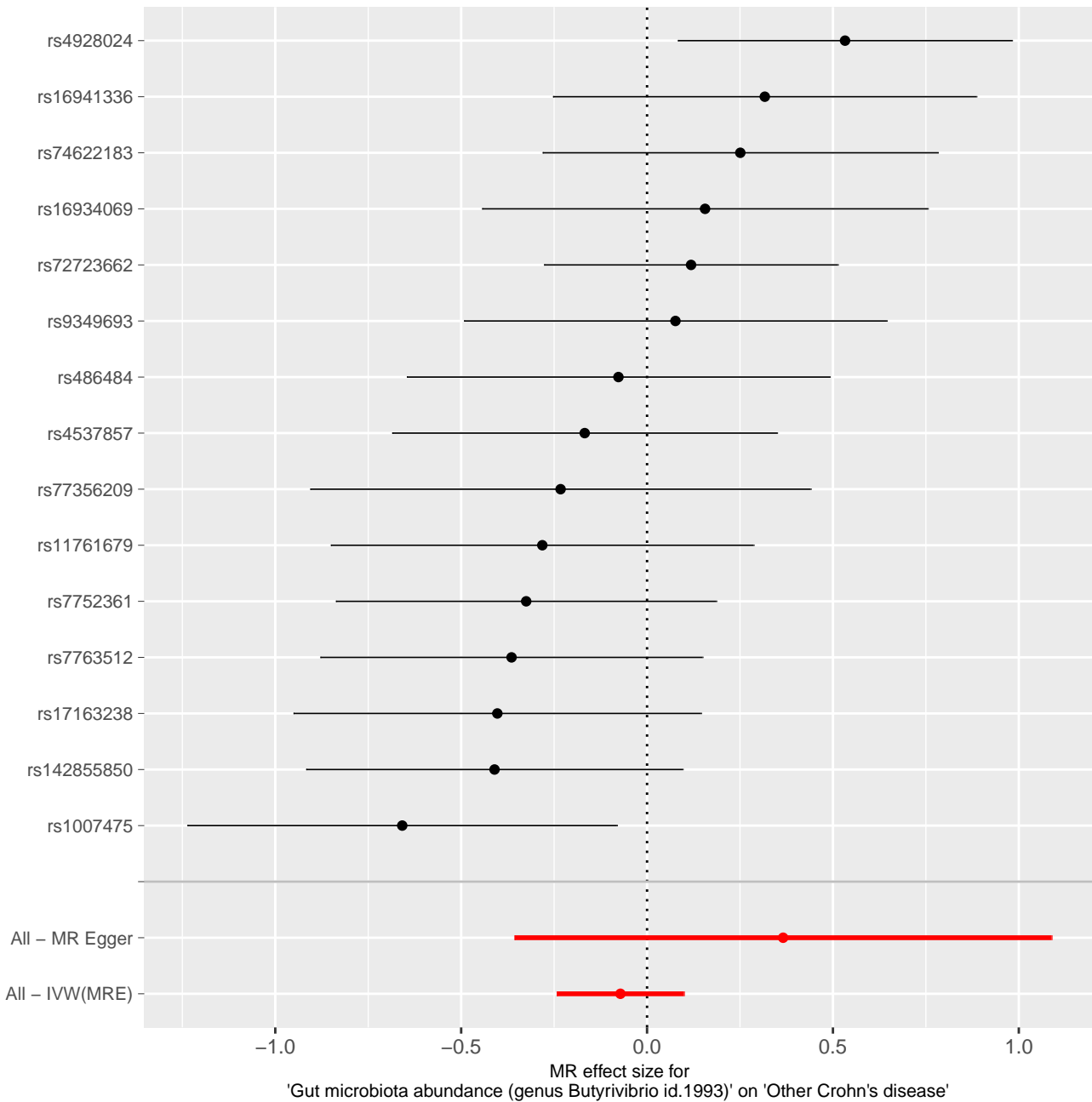


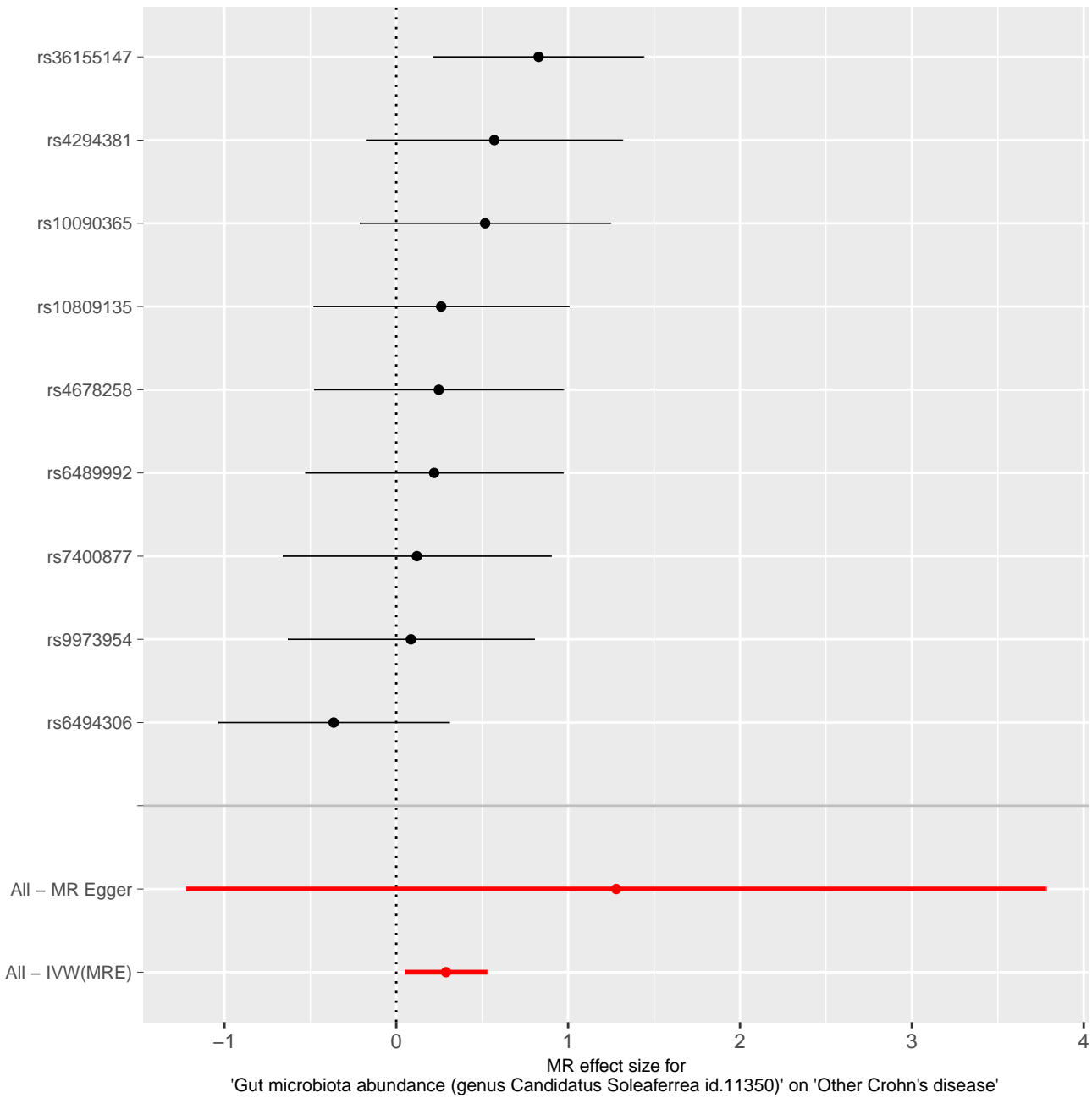


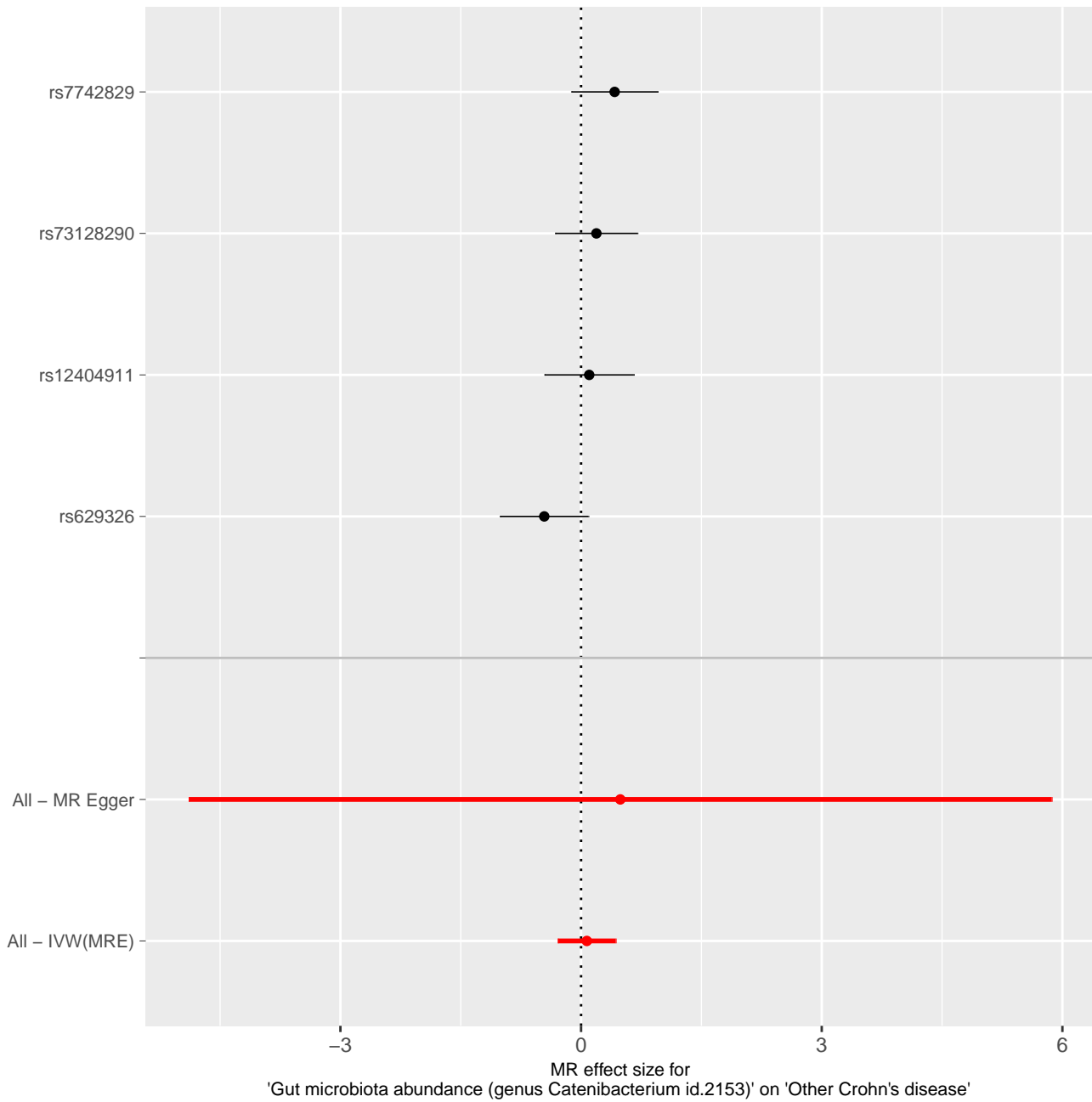
Batch 198 : Gut microbiota abundance (genus Butyricimonas id.945) on Other Crohn's disease

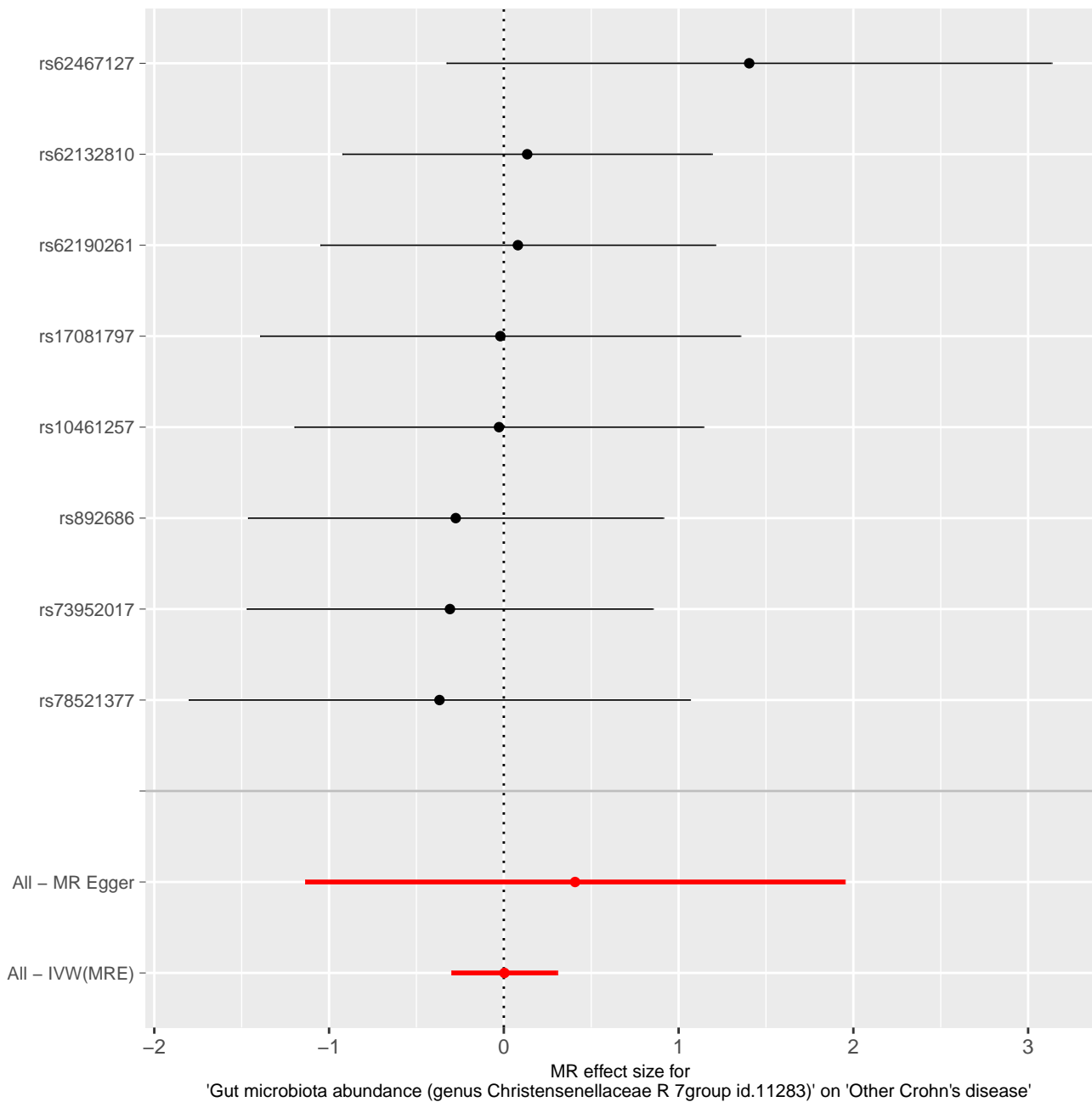


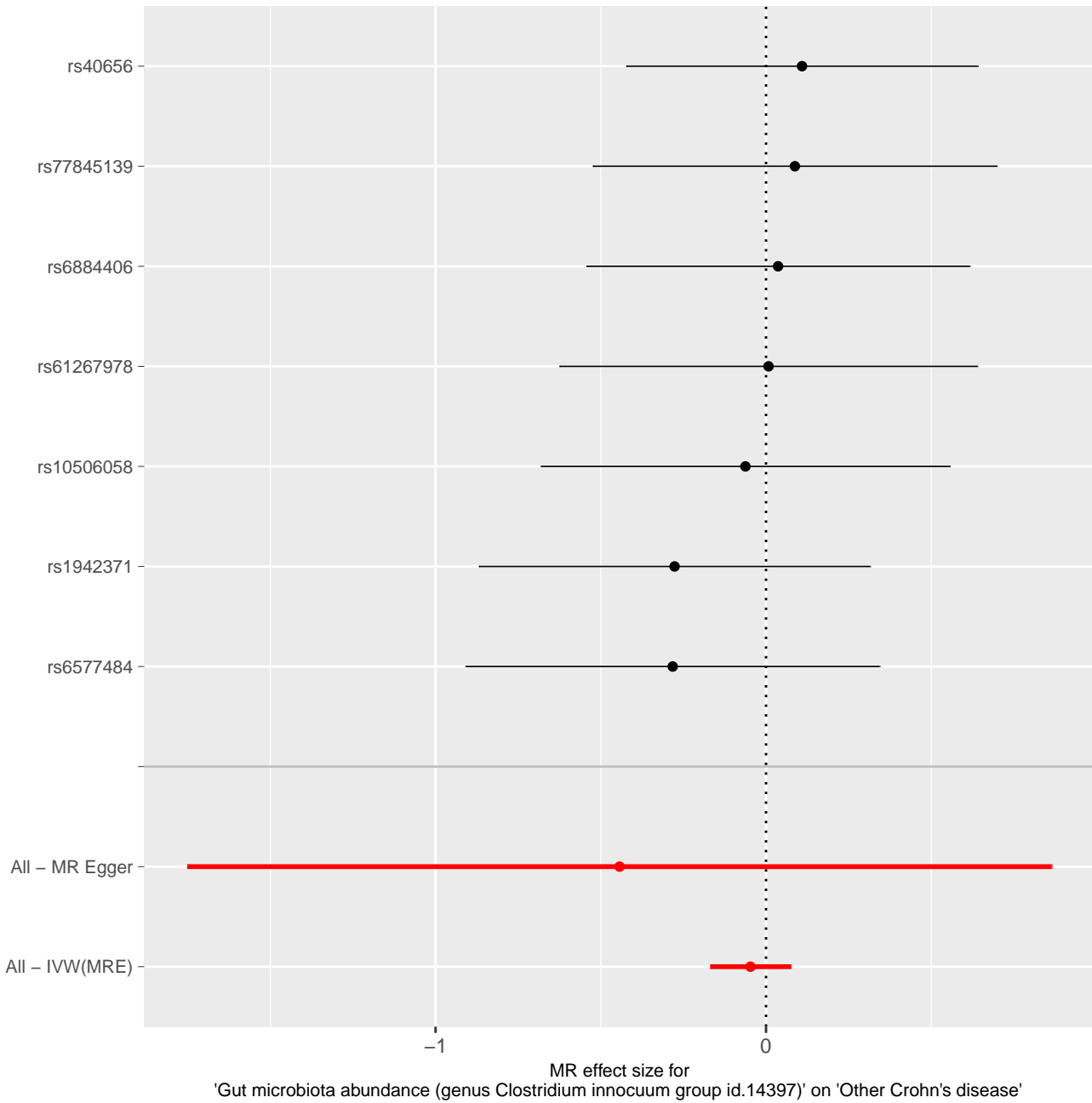
Batch 199 : Gut microbiota abundance (genus Butyrivibrio id.1993) on Other Crohn's disease

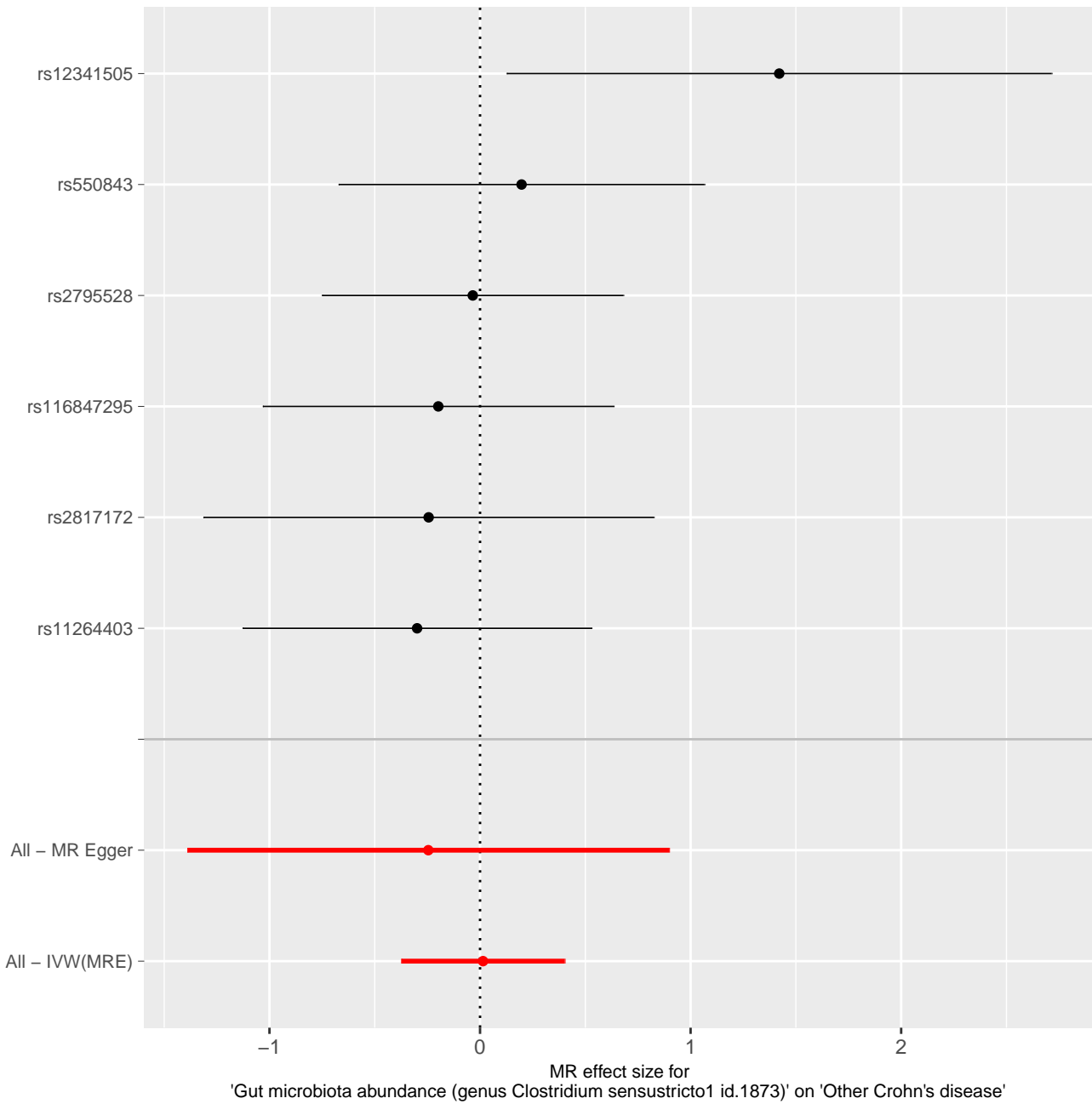


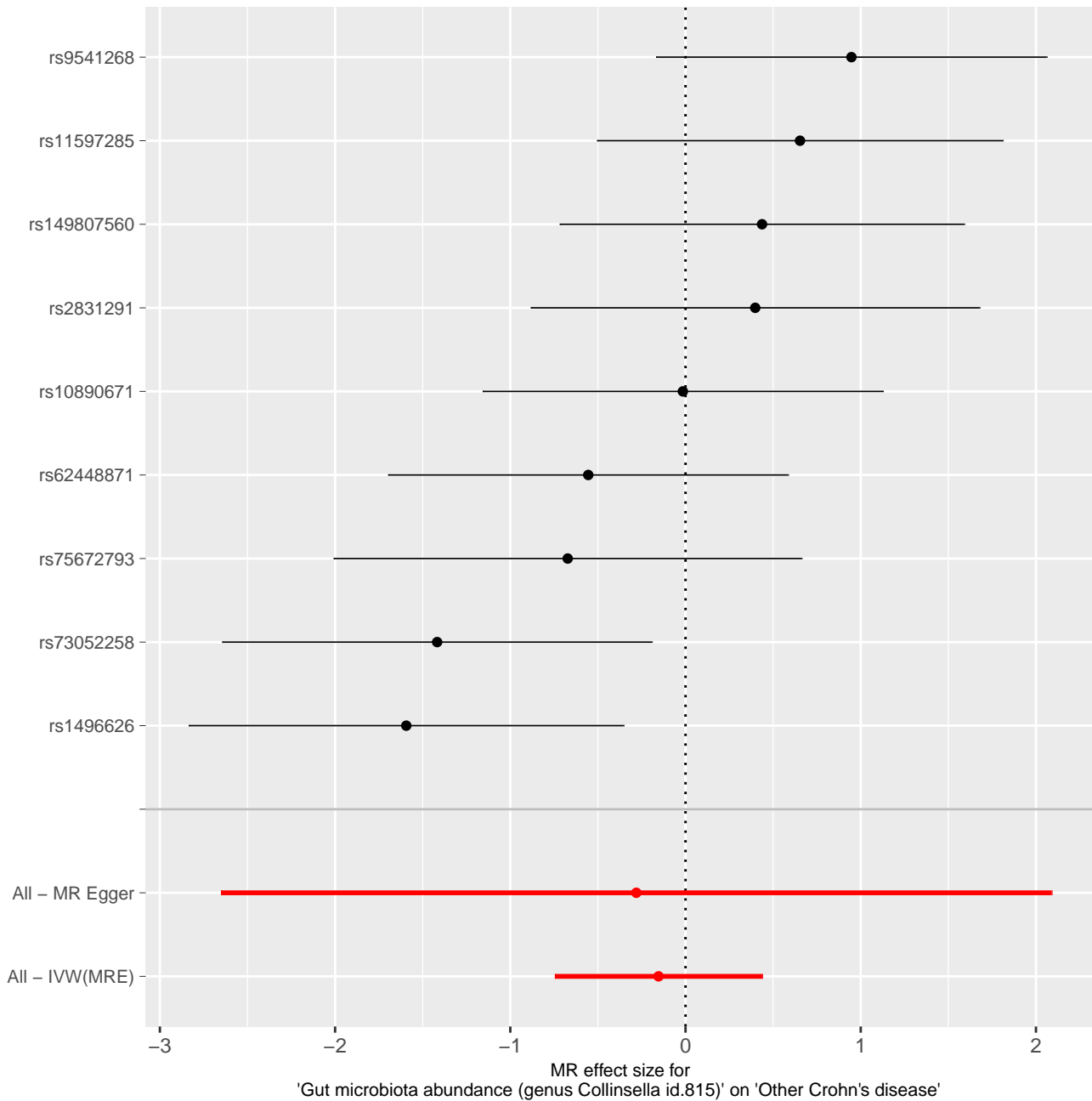


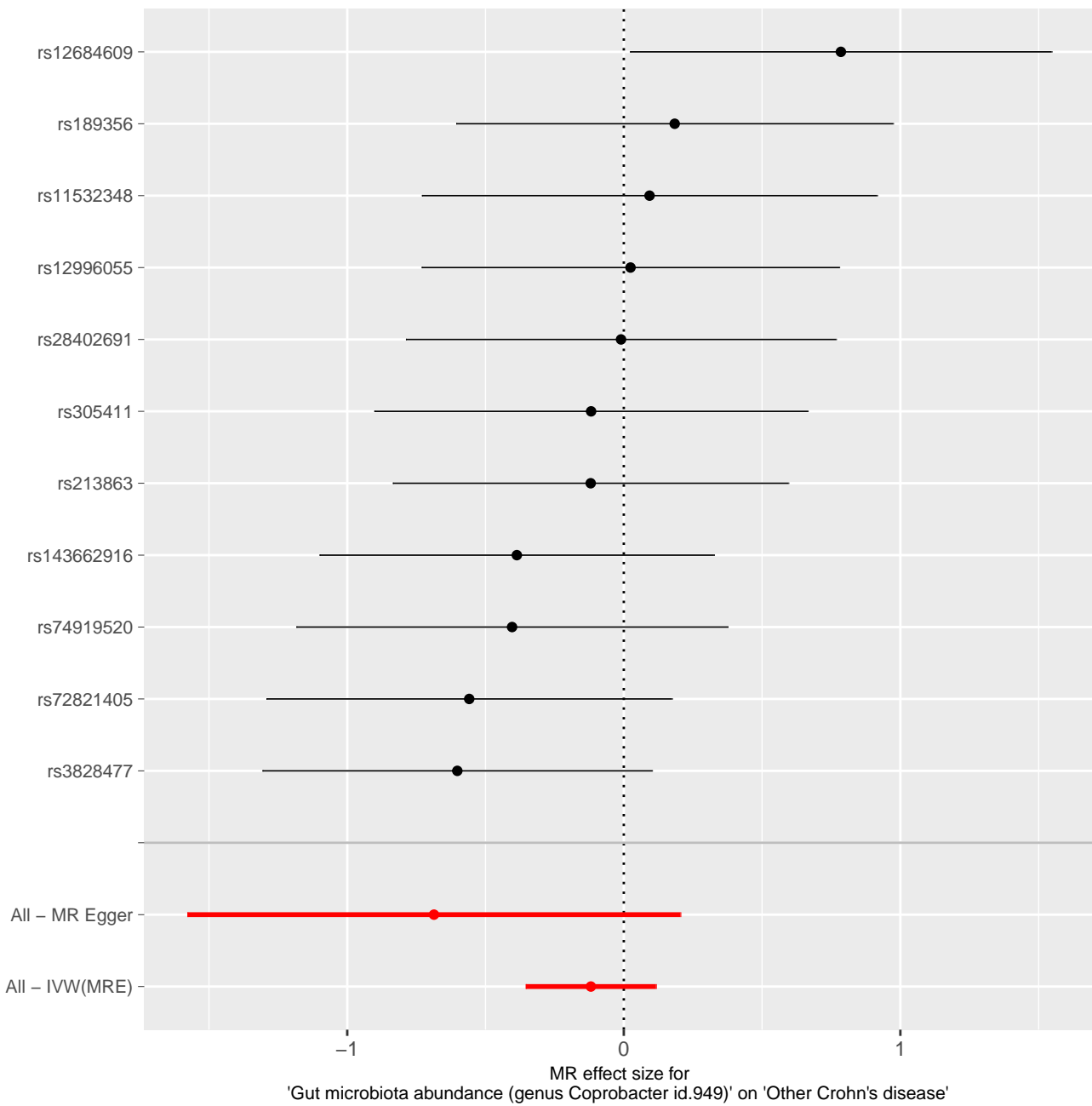


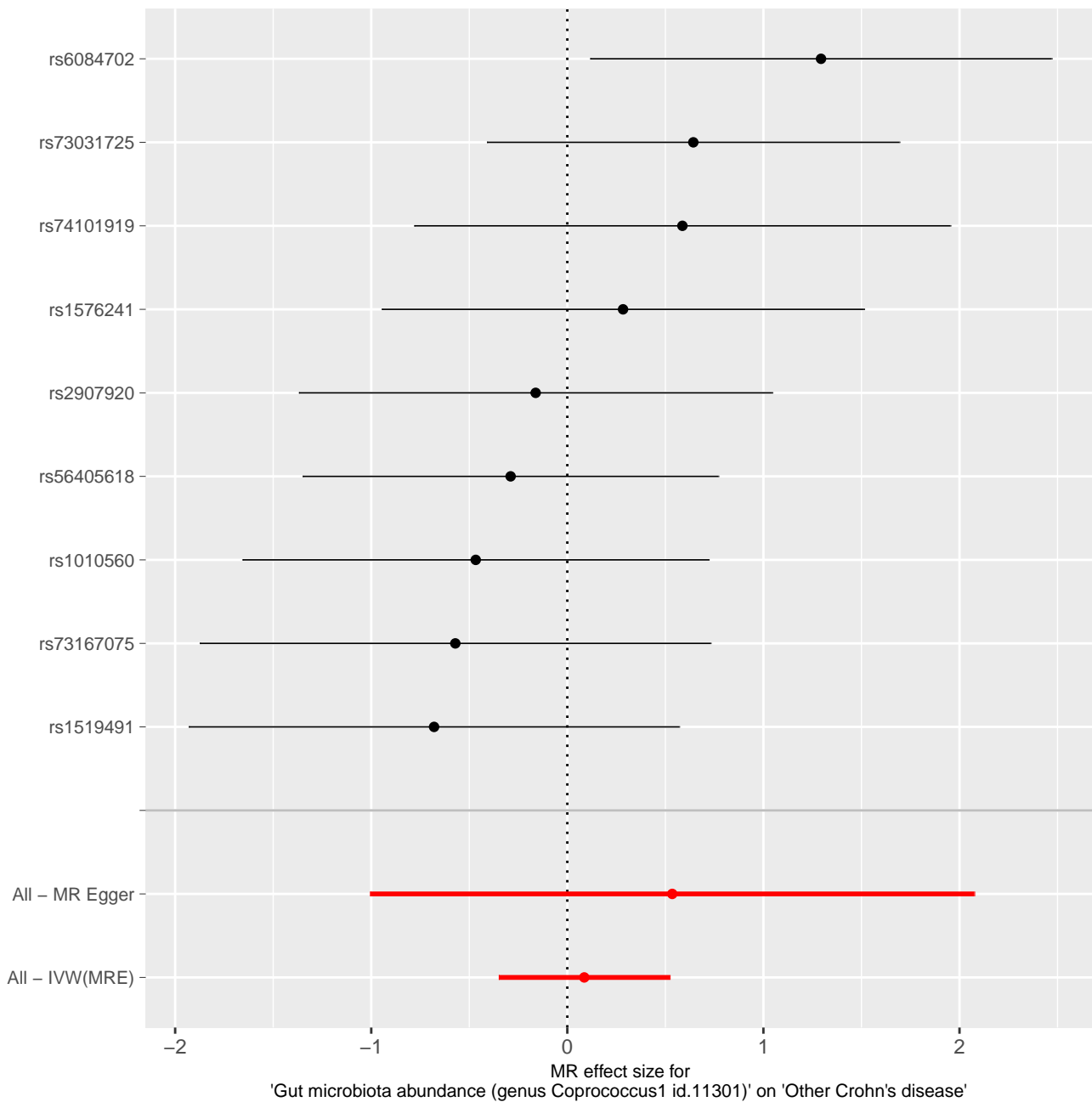


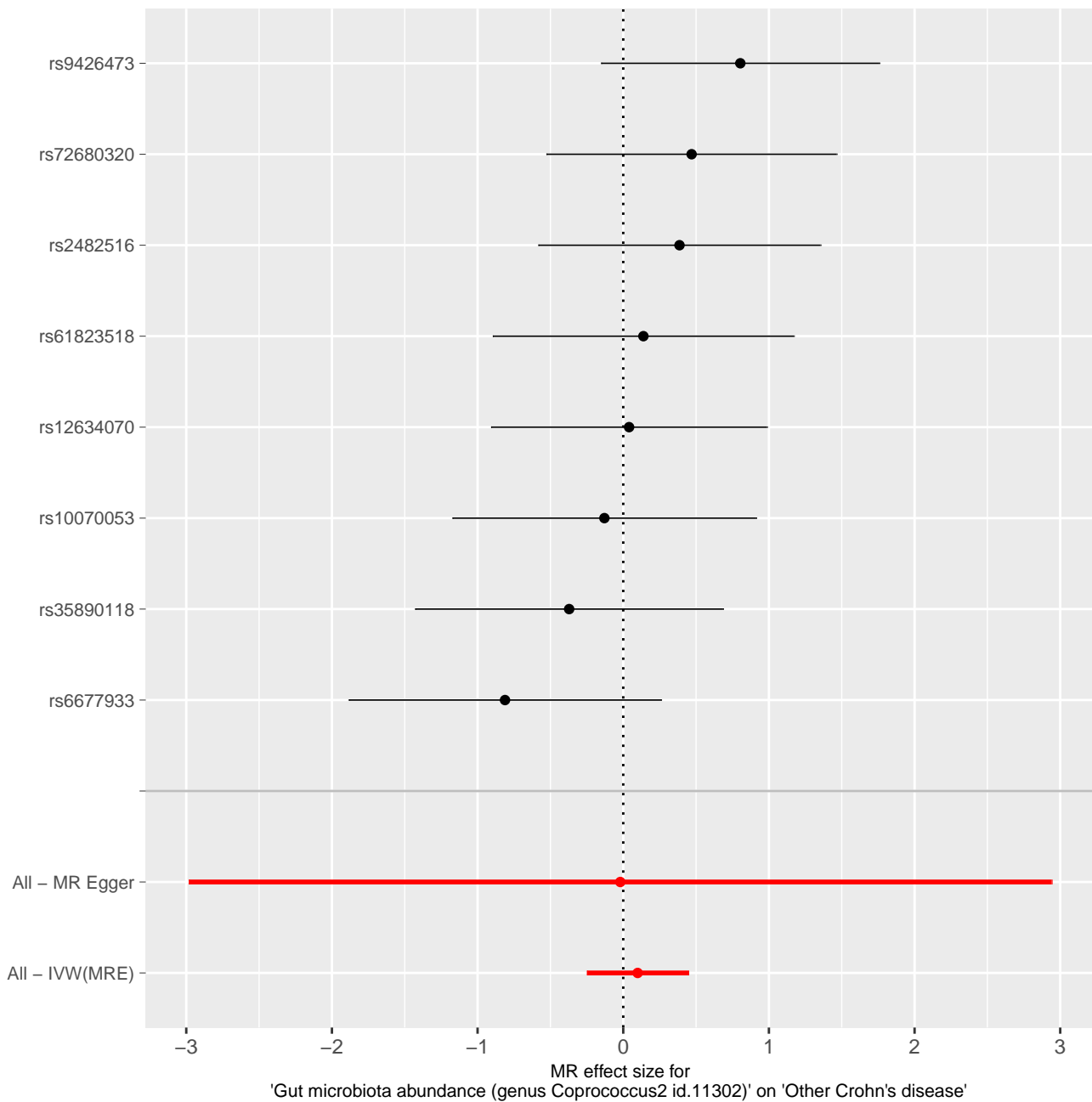


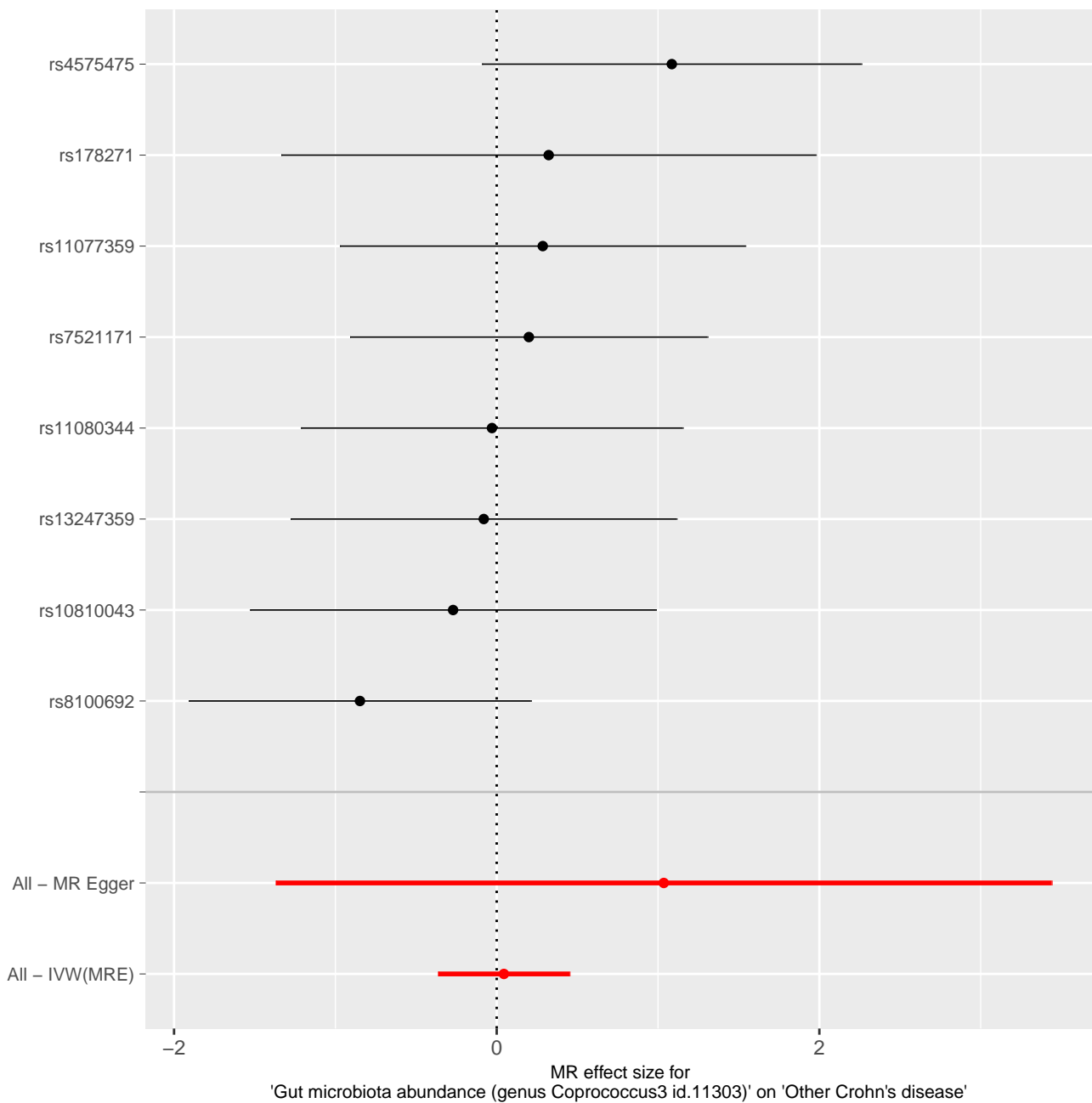


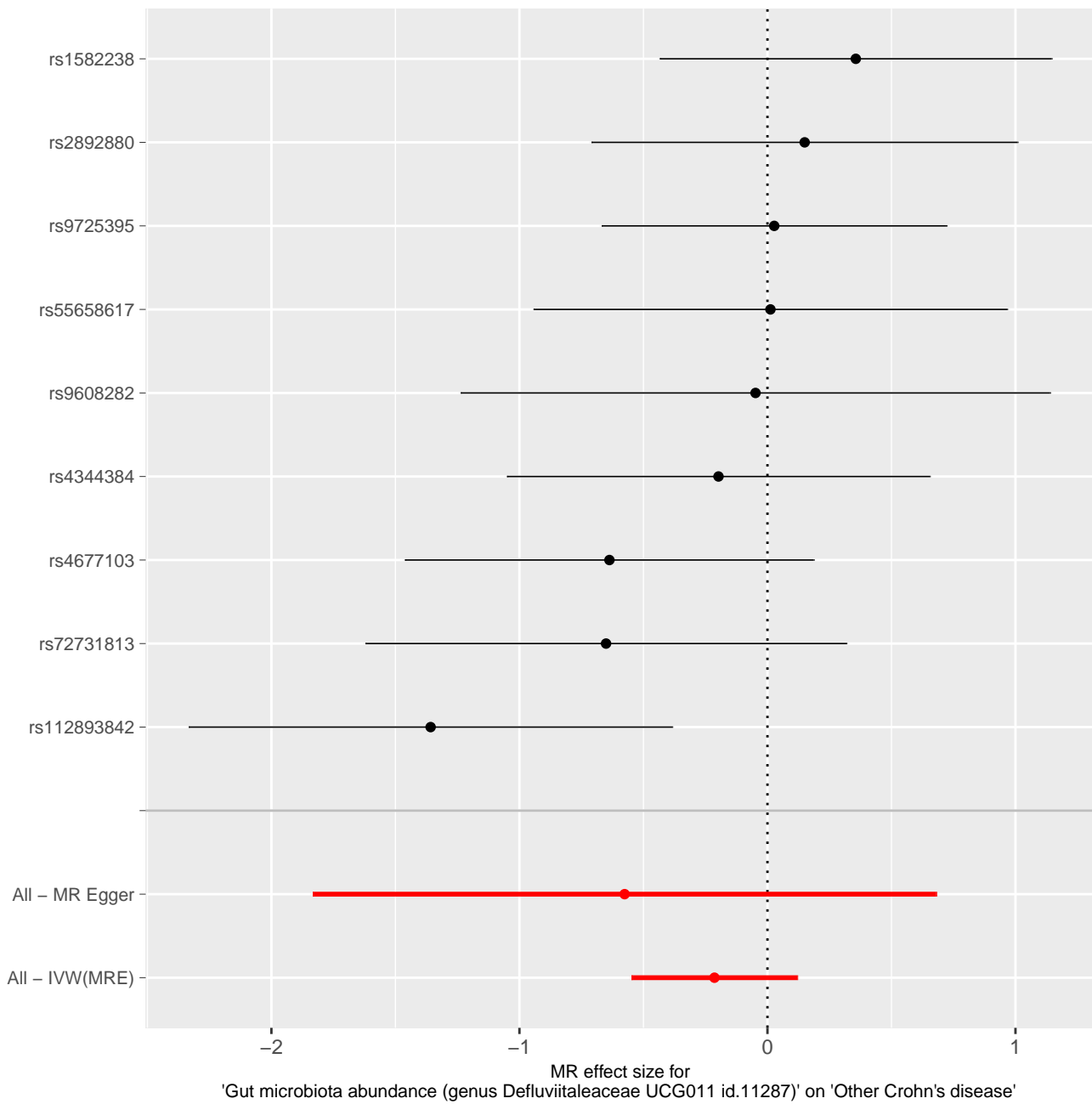


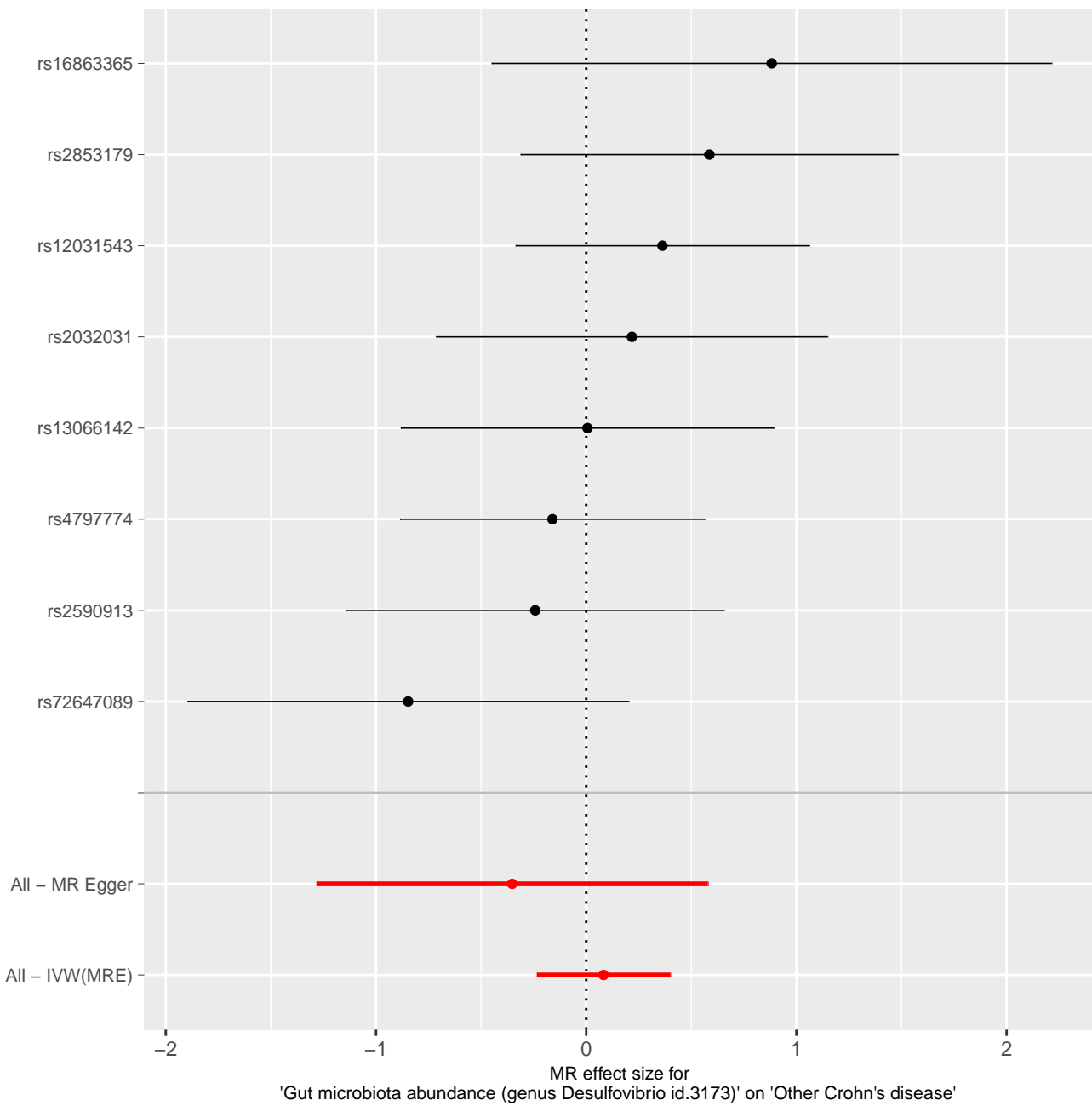




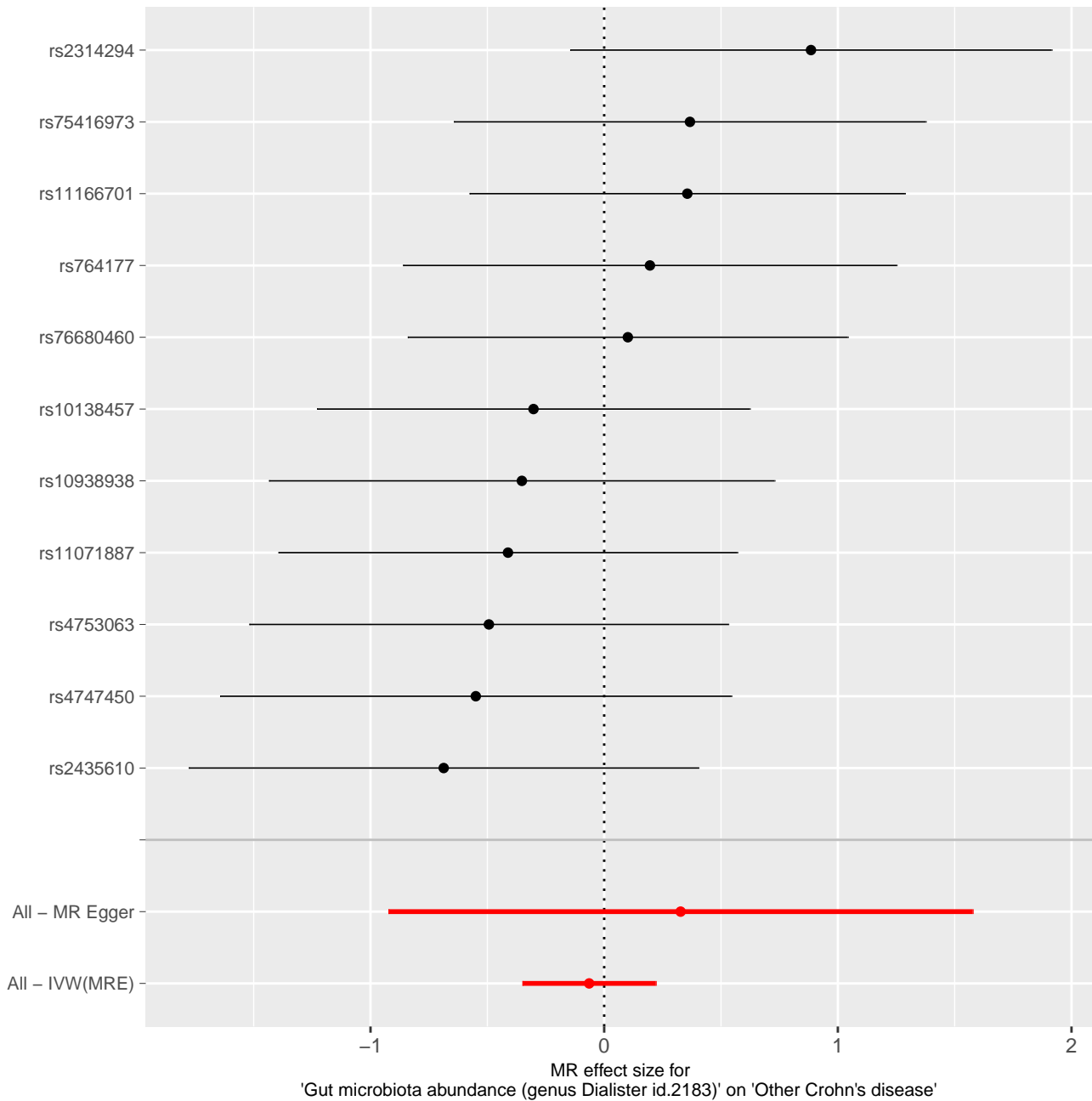




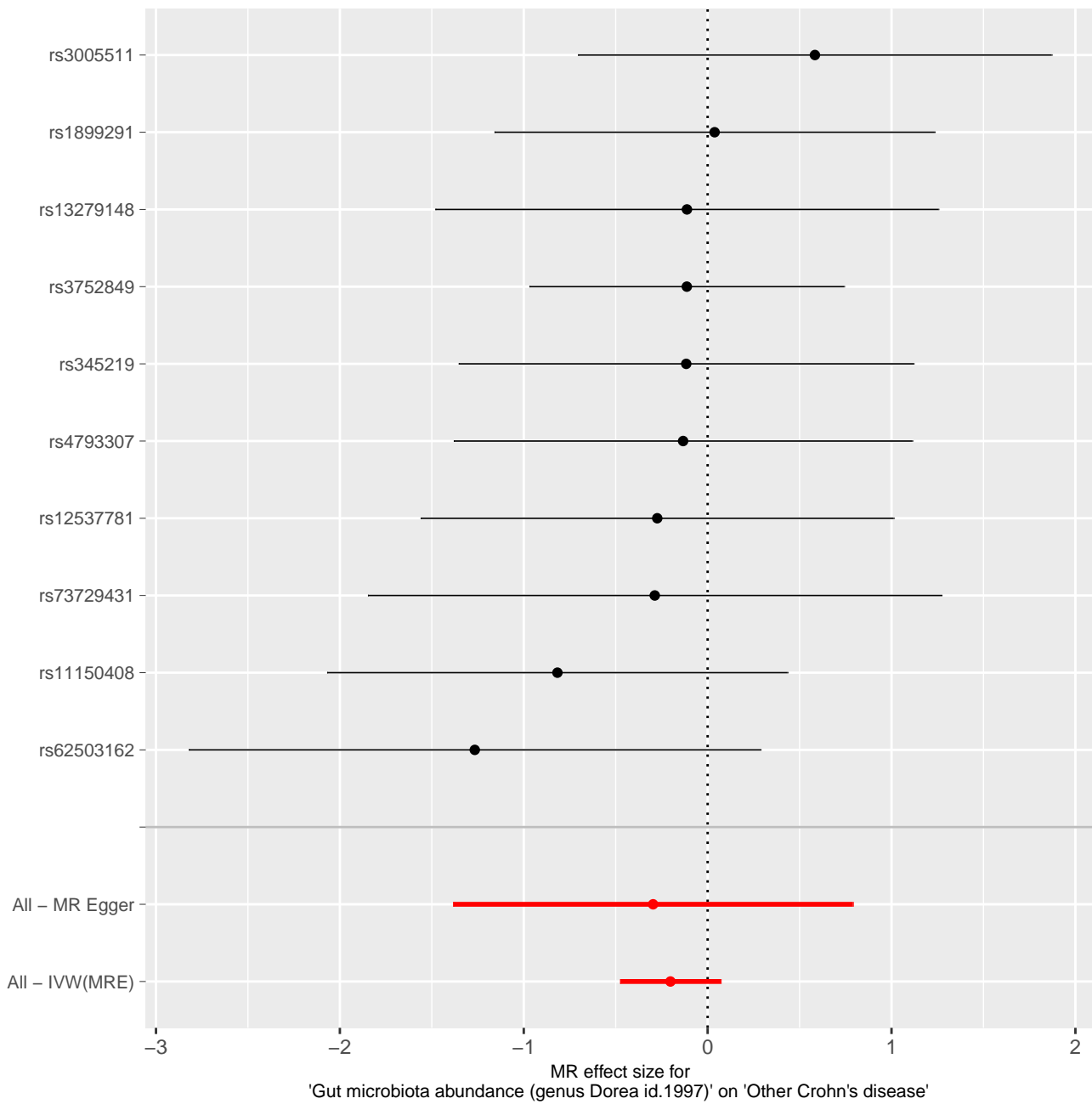


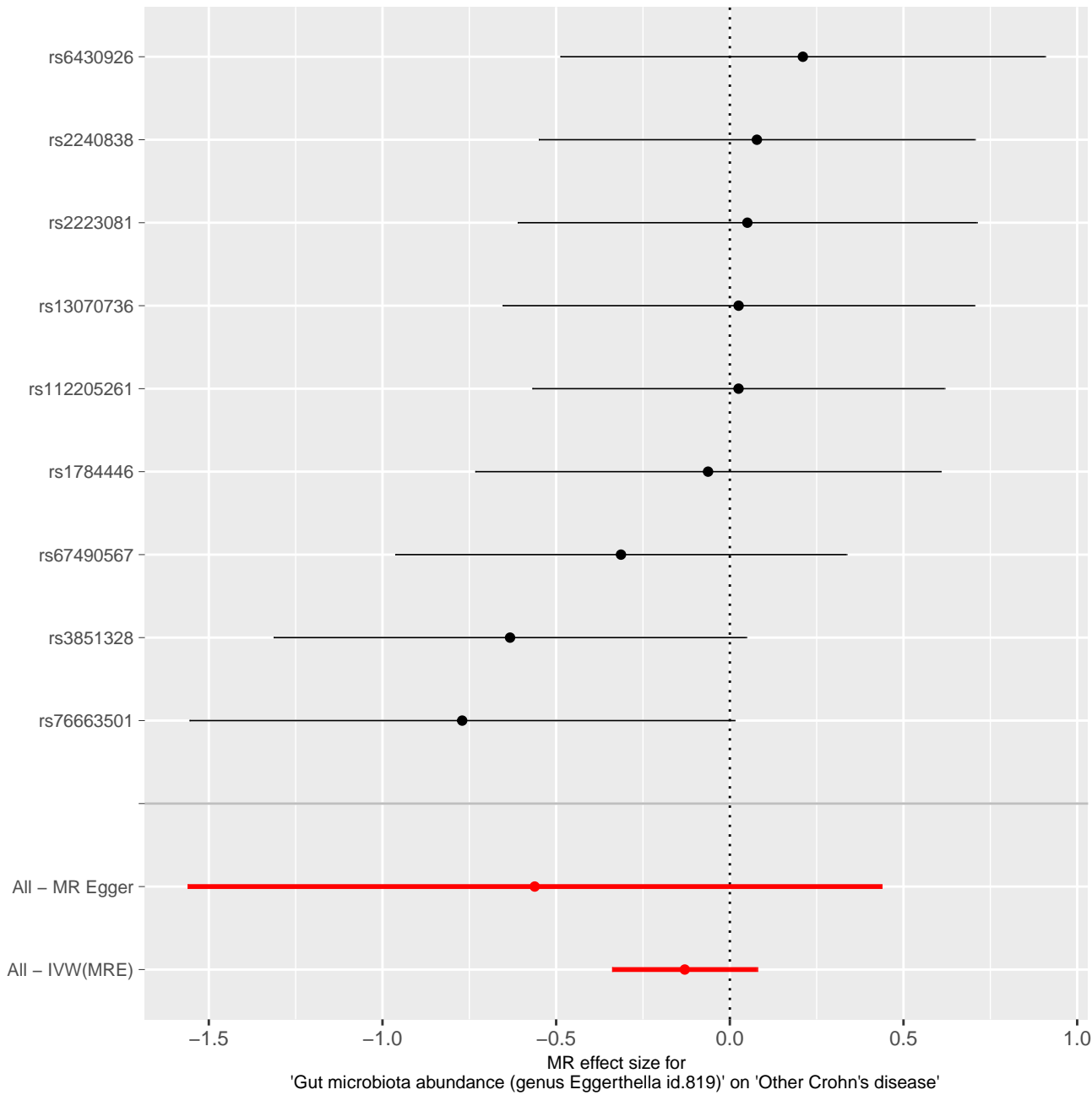


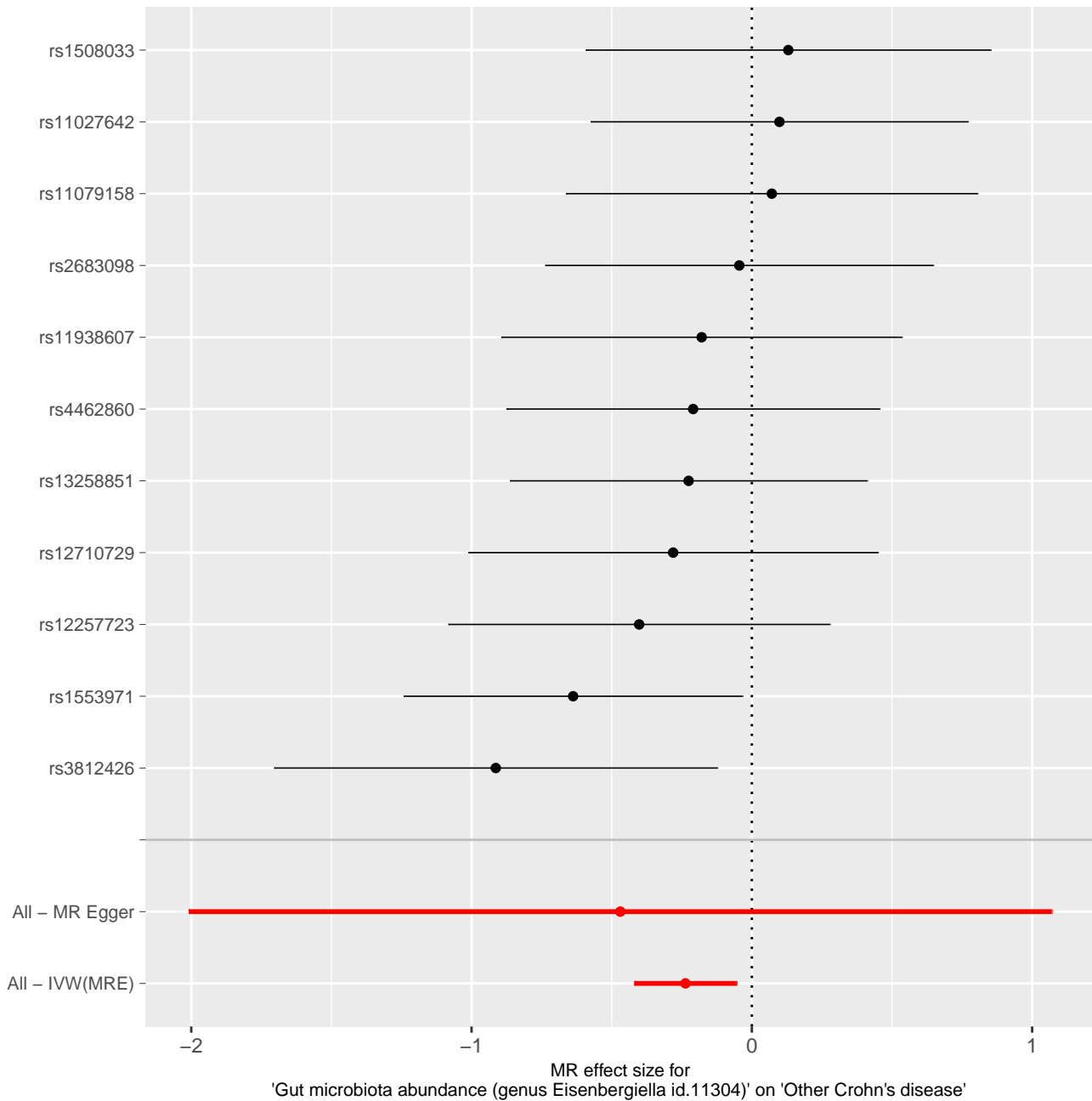
Batch 212 : Gut microbiota abundance (genus Dialister id.2183) on Other Crohn's disease

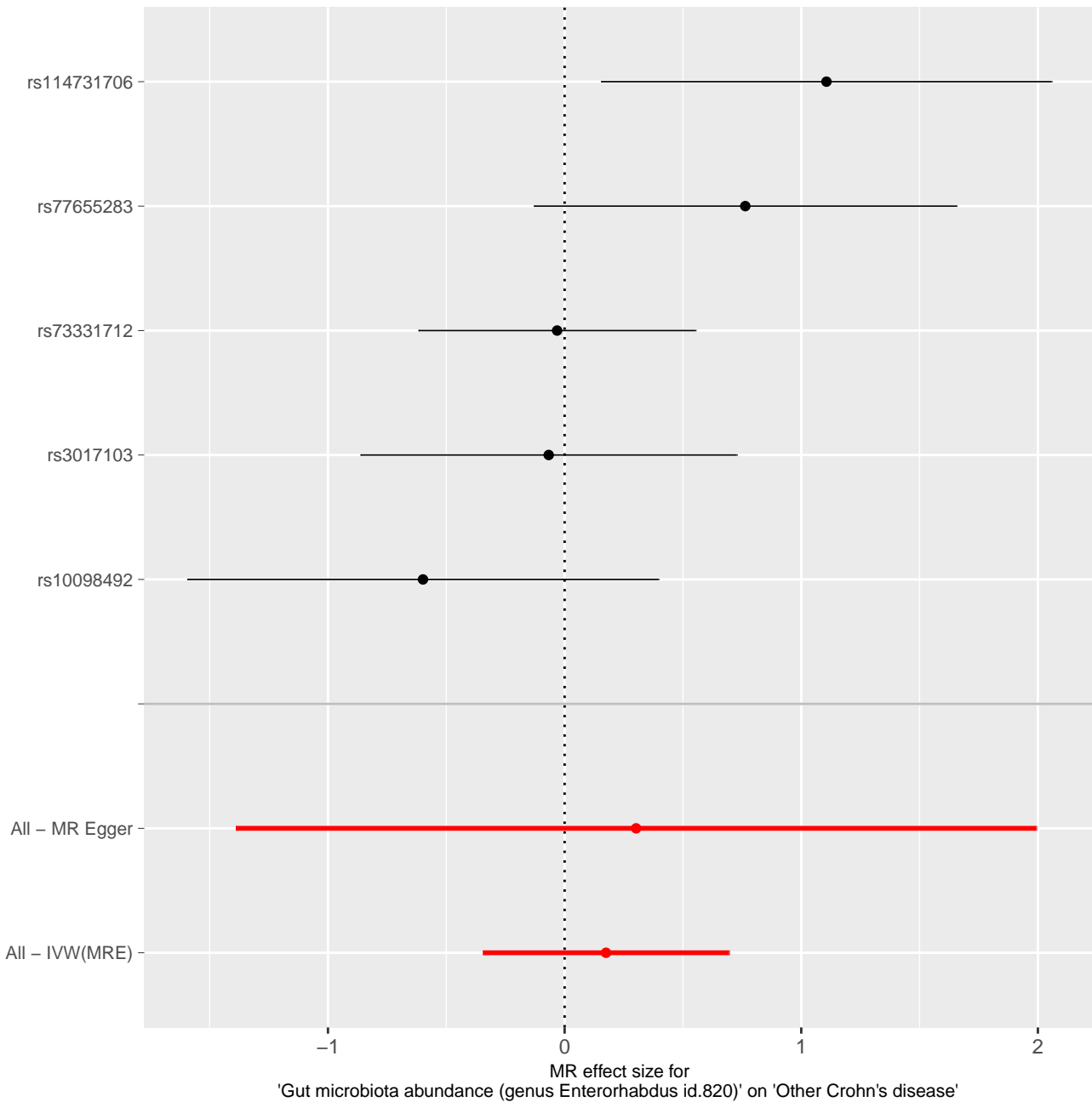


Batch 213 : Gut microbiota abundance (genus Dorea id.1997) on Other Crohn's disease

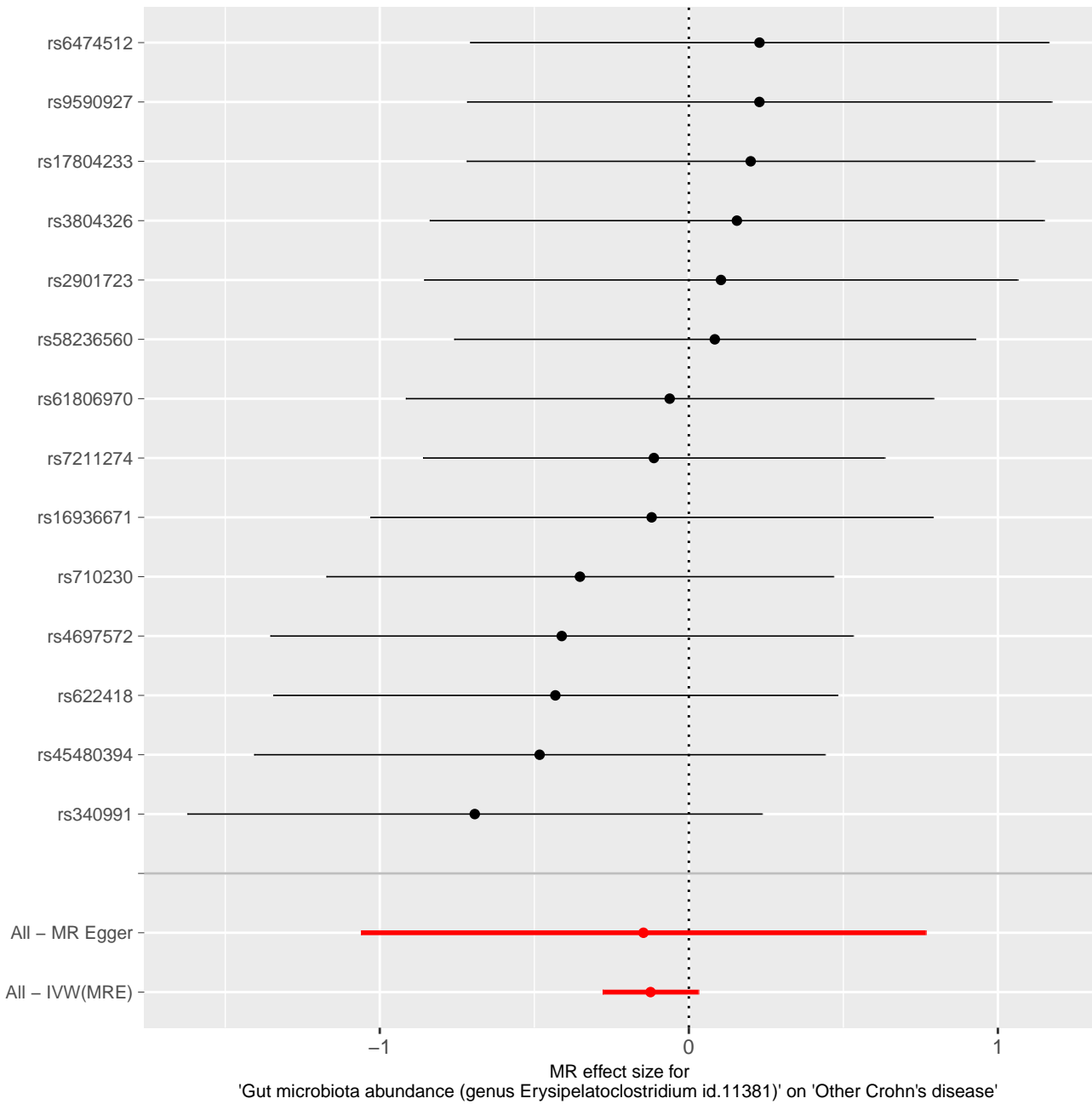


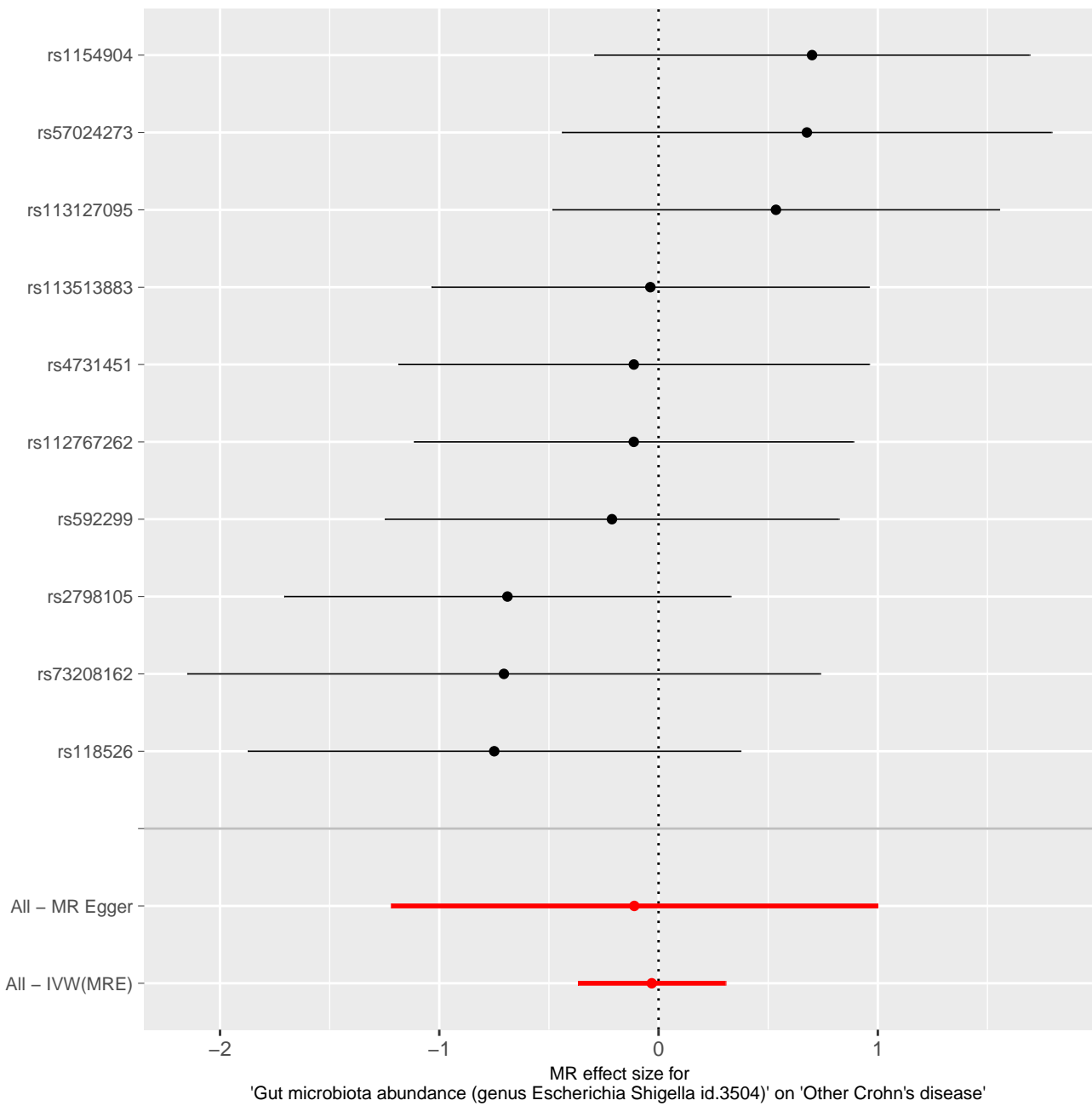


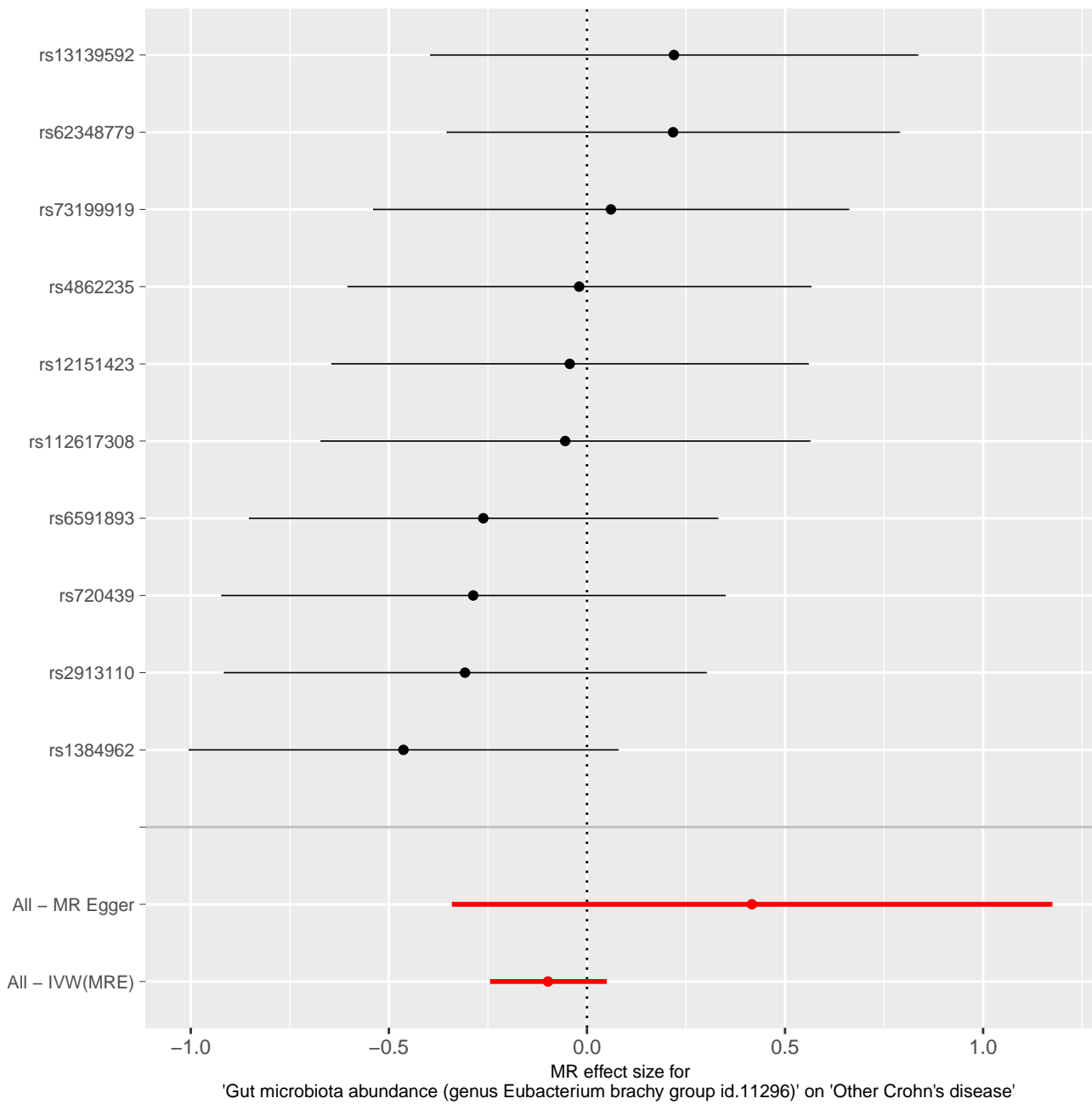


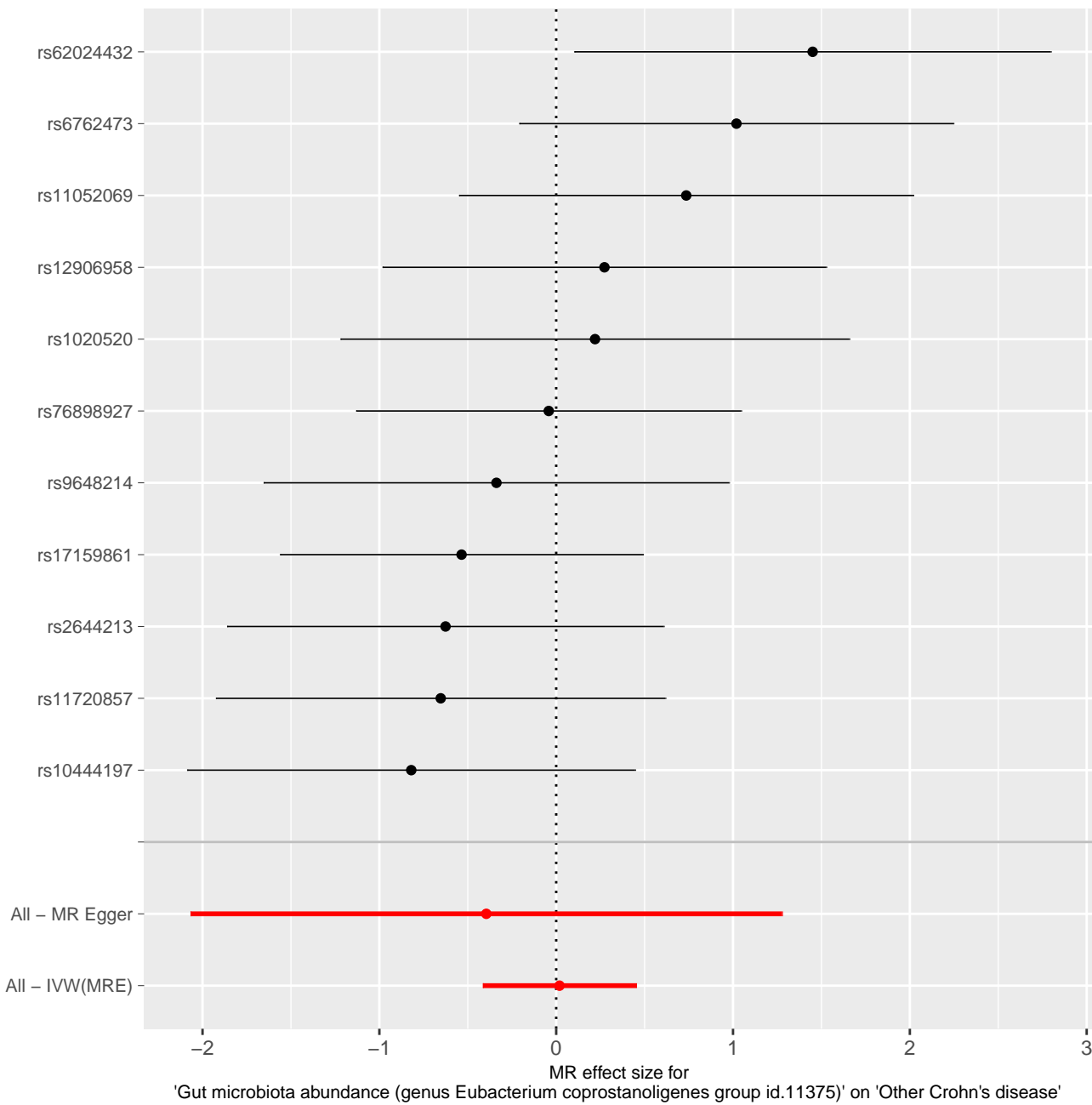


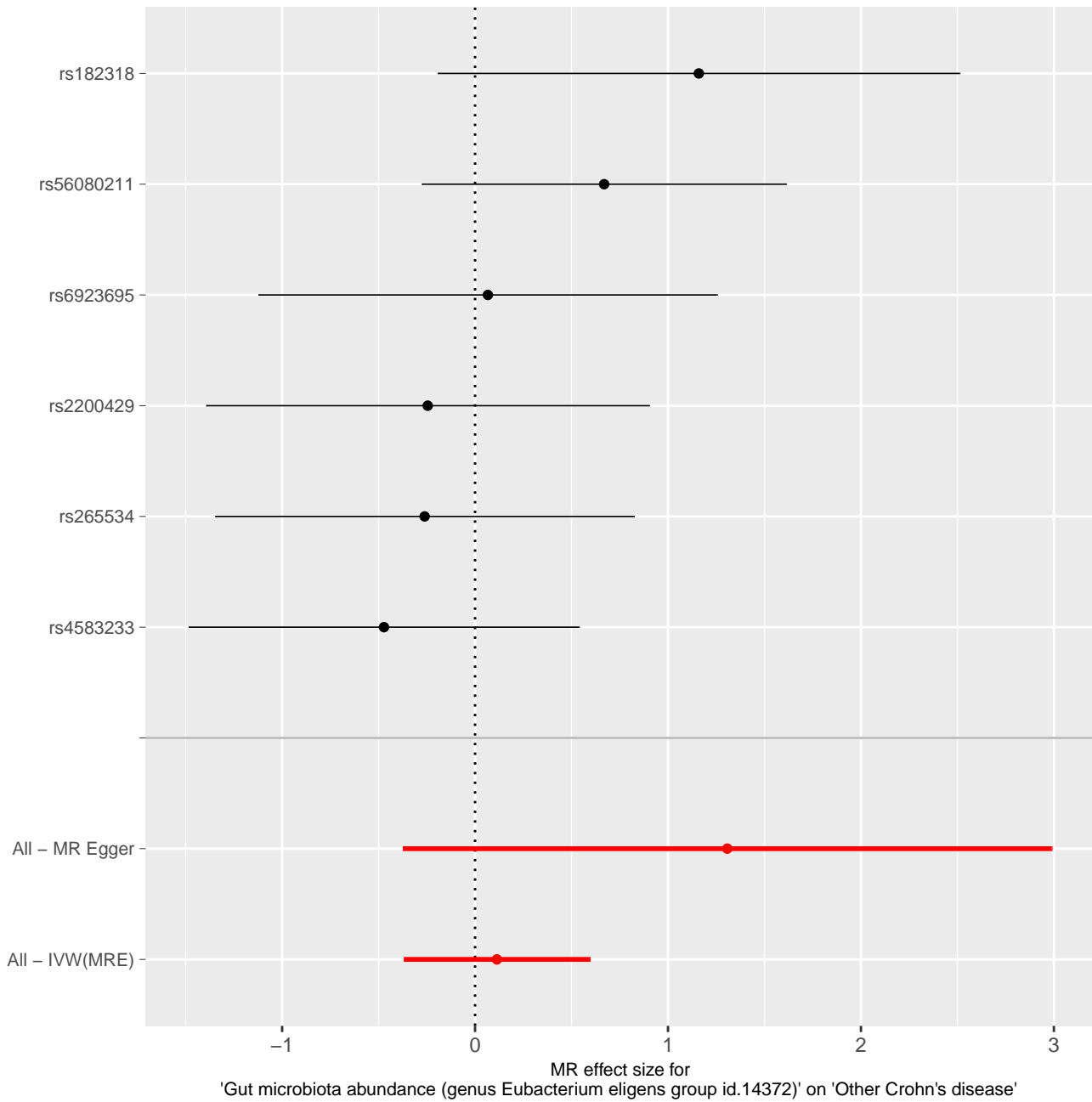
Batch 217 : Gut microbiota abundance (genus Erysipelatoclostridium id.11381) on Other Crohn's disease

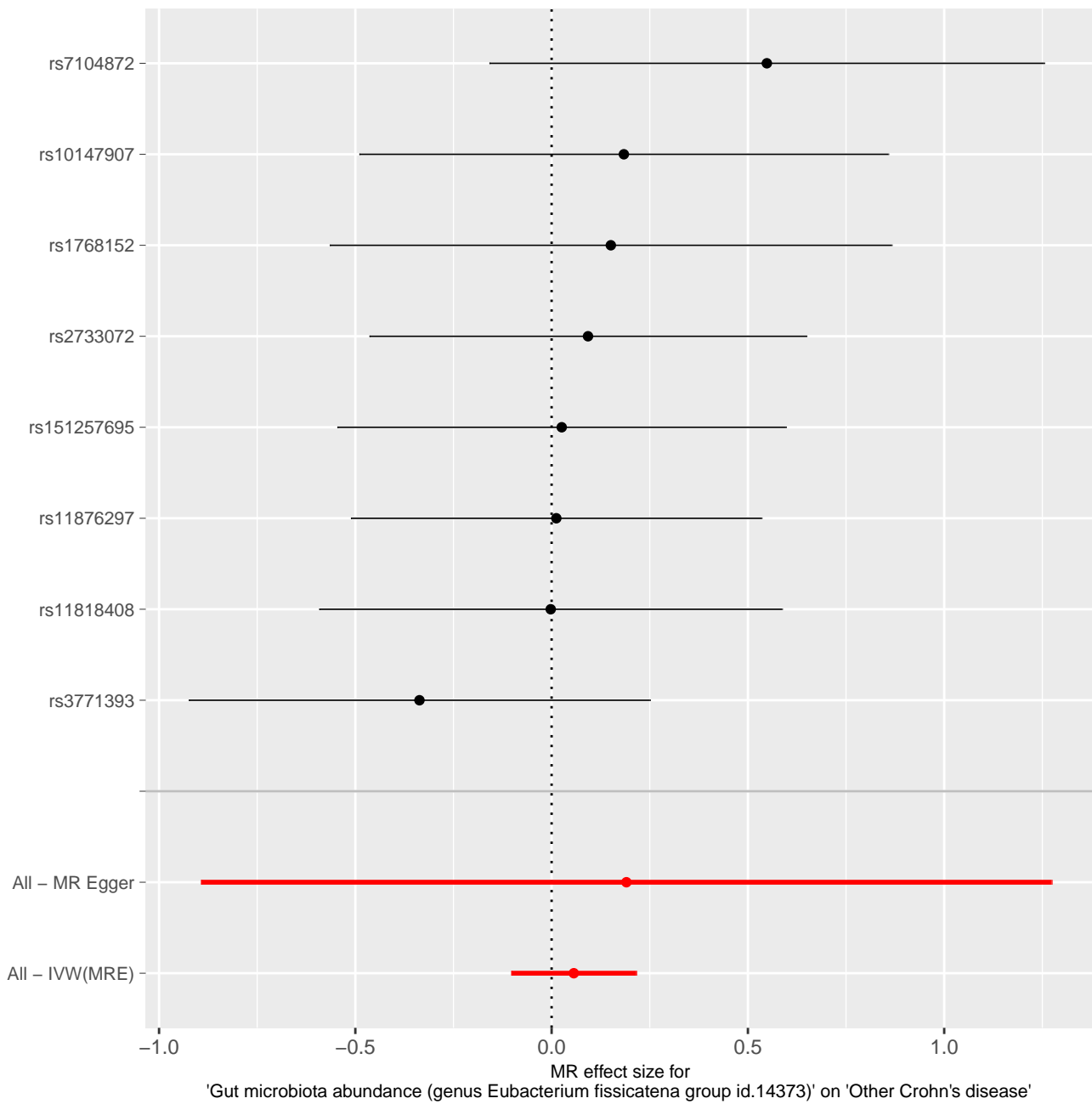


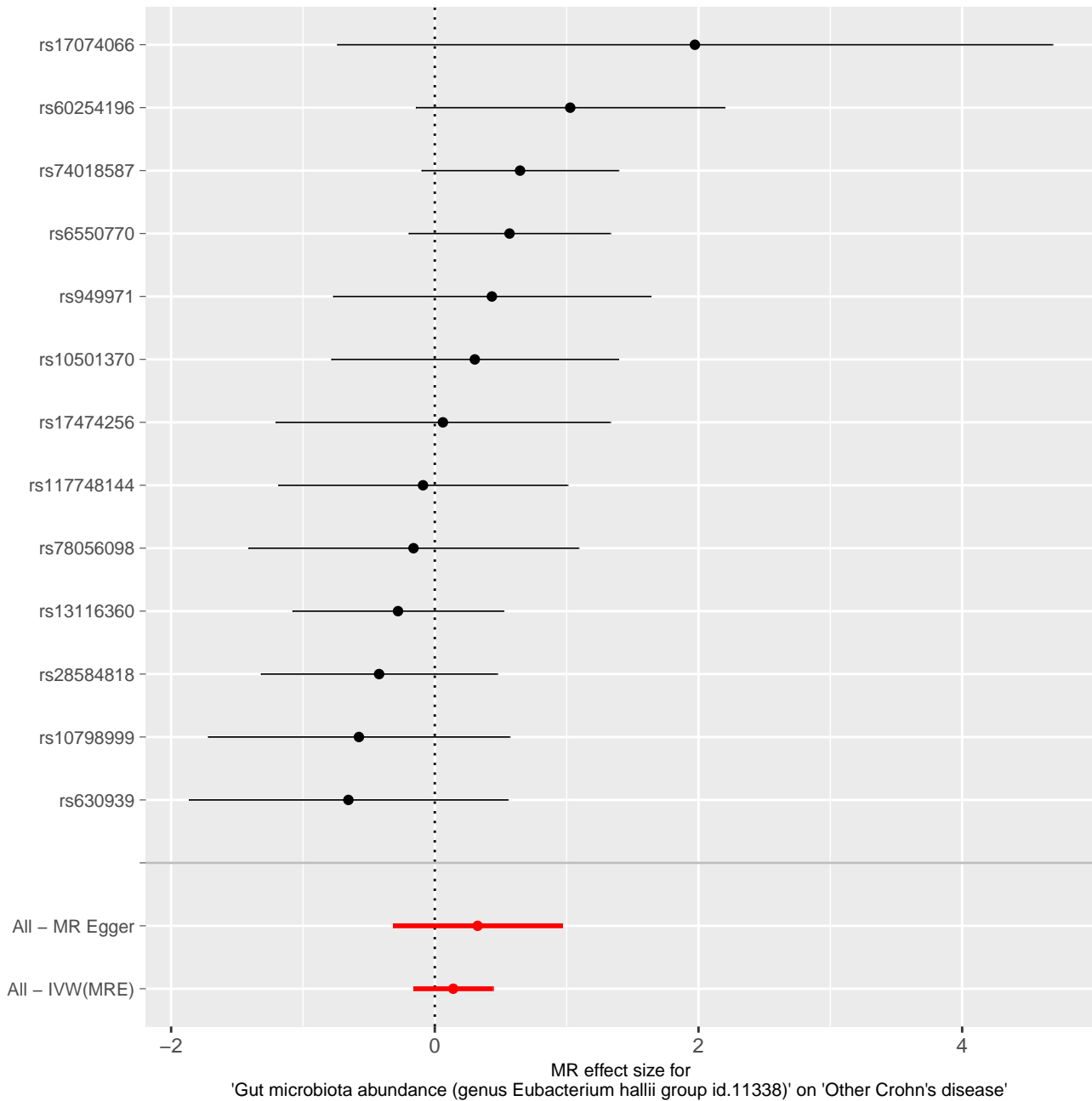




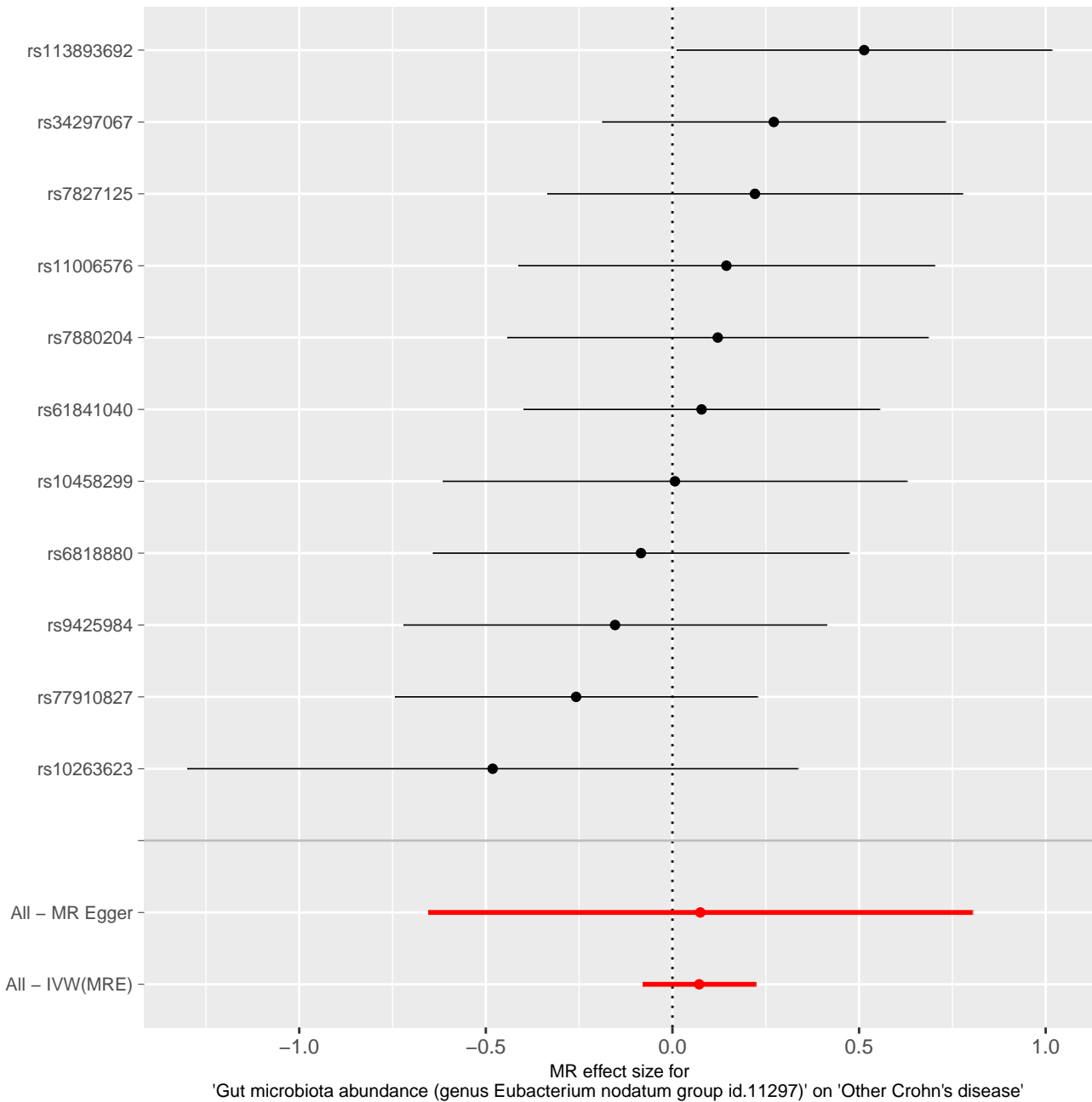


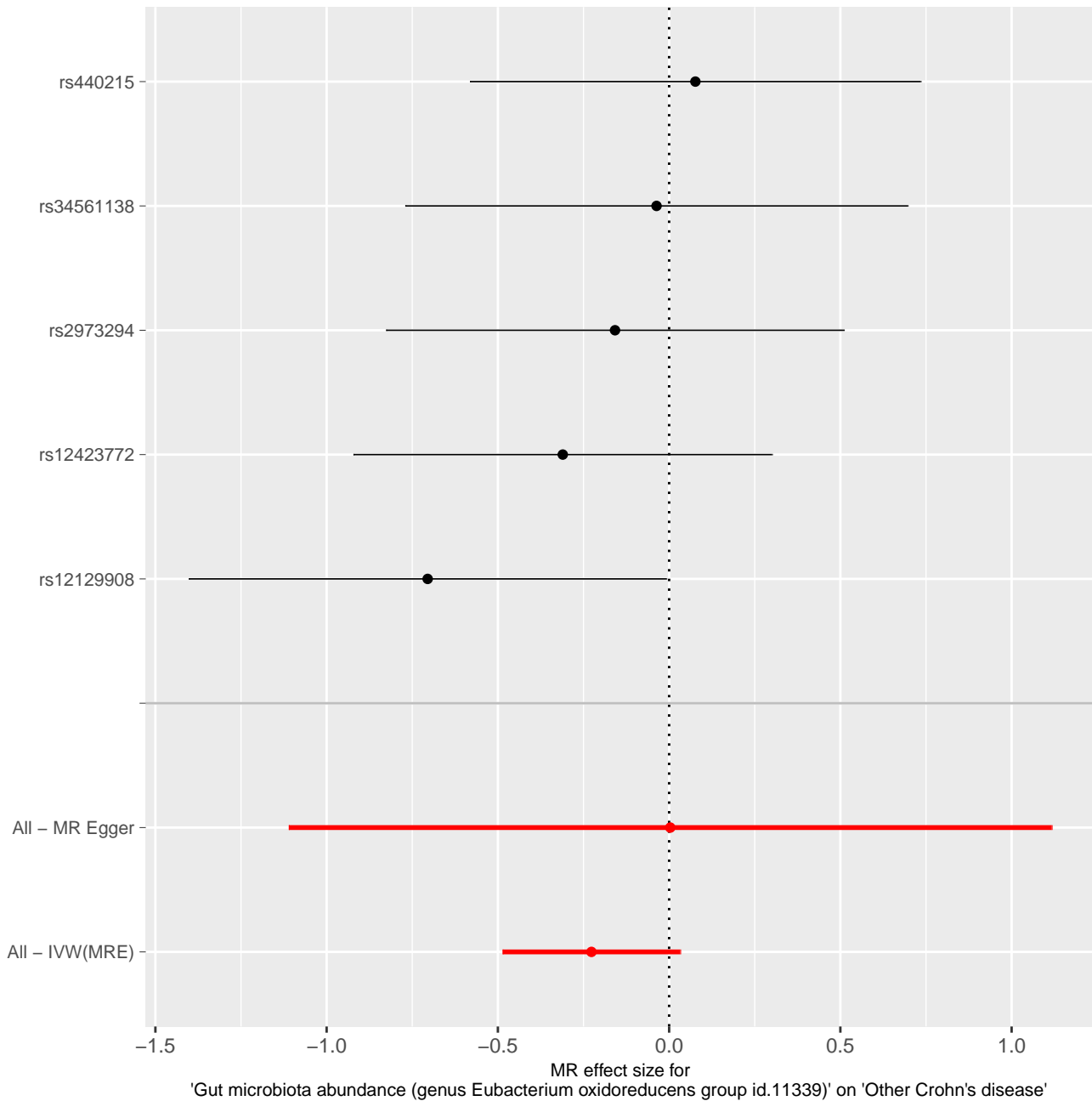


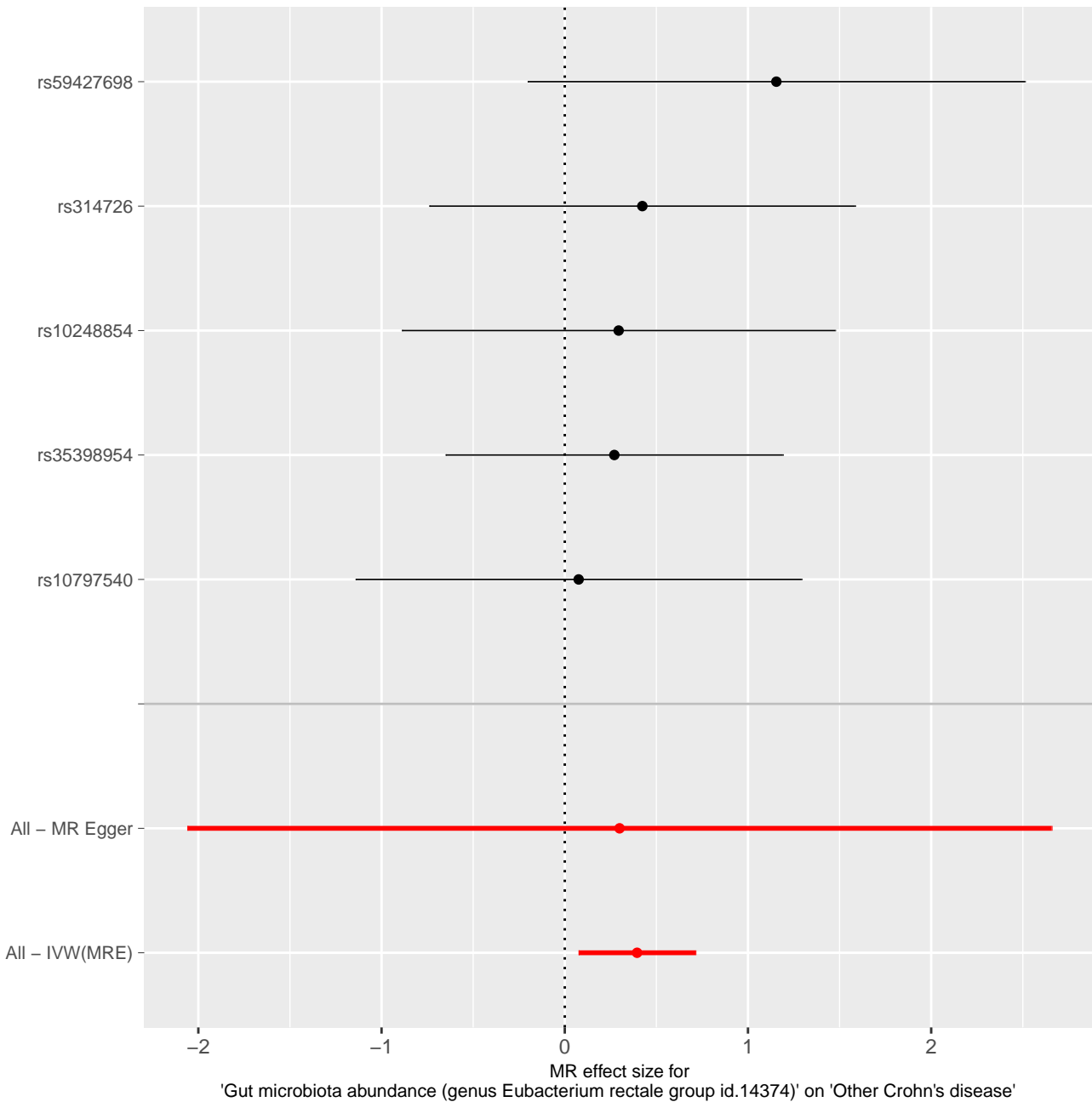




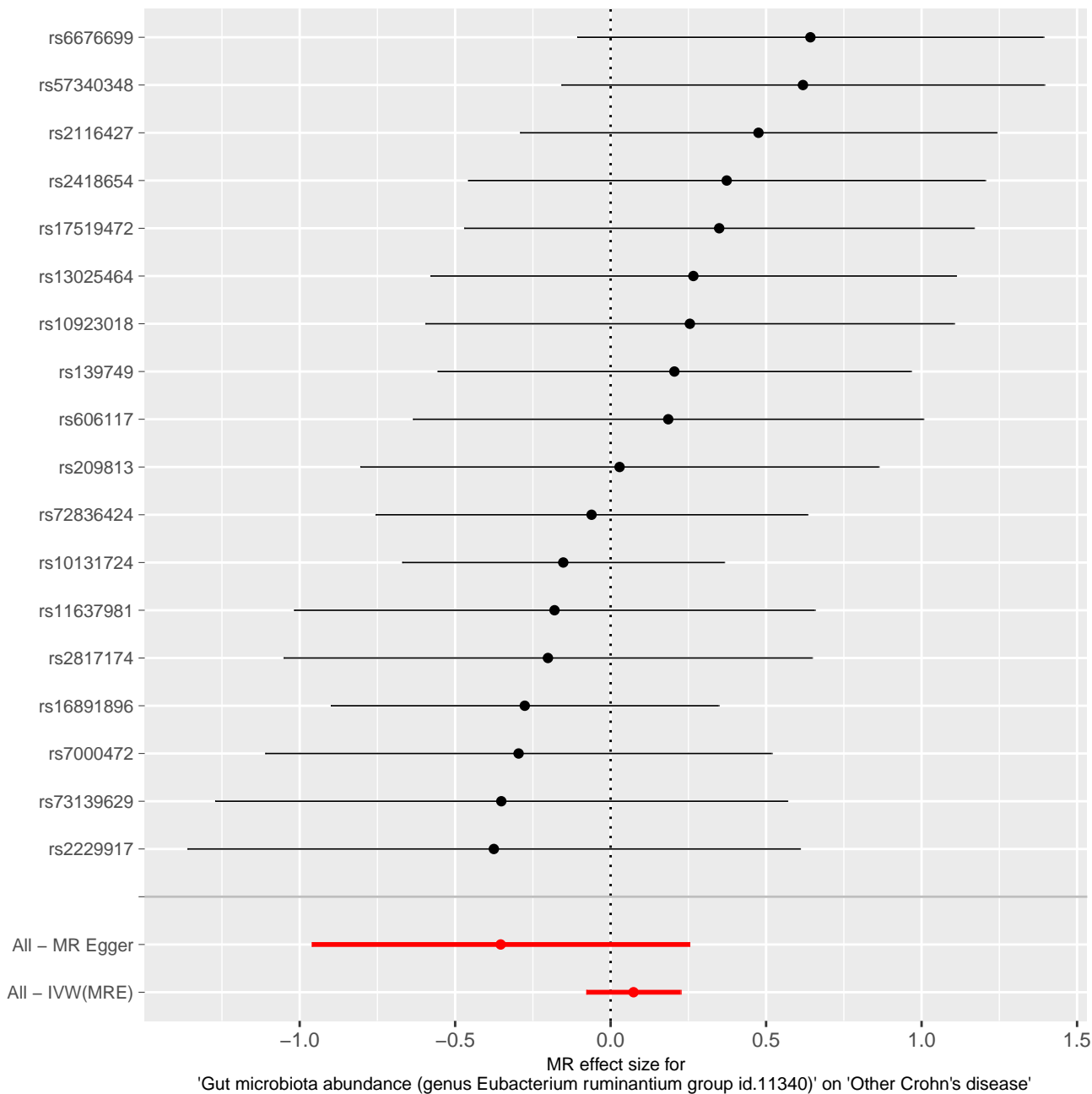
Batch 225 : Gut microbiota abundance (genus Eubacterium nodatum group id.11297) on Other Crohn's disease

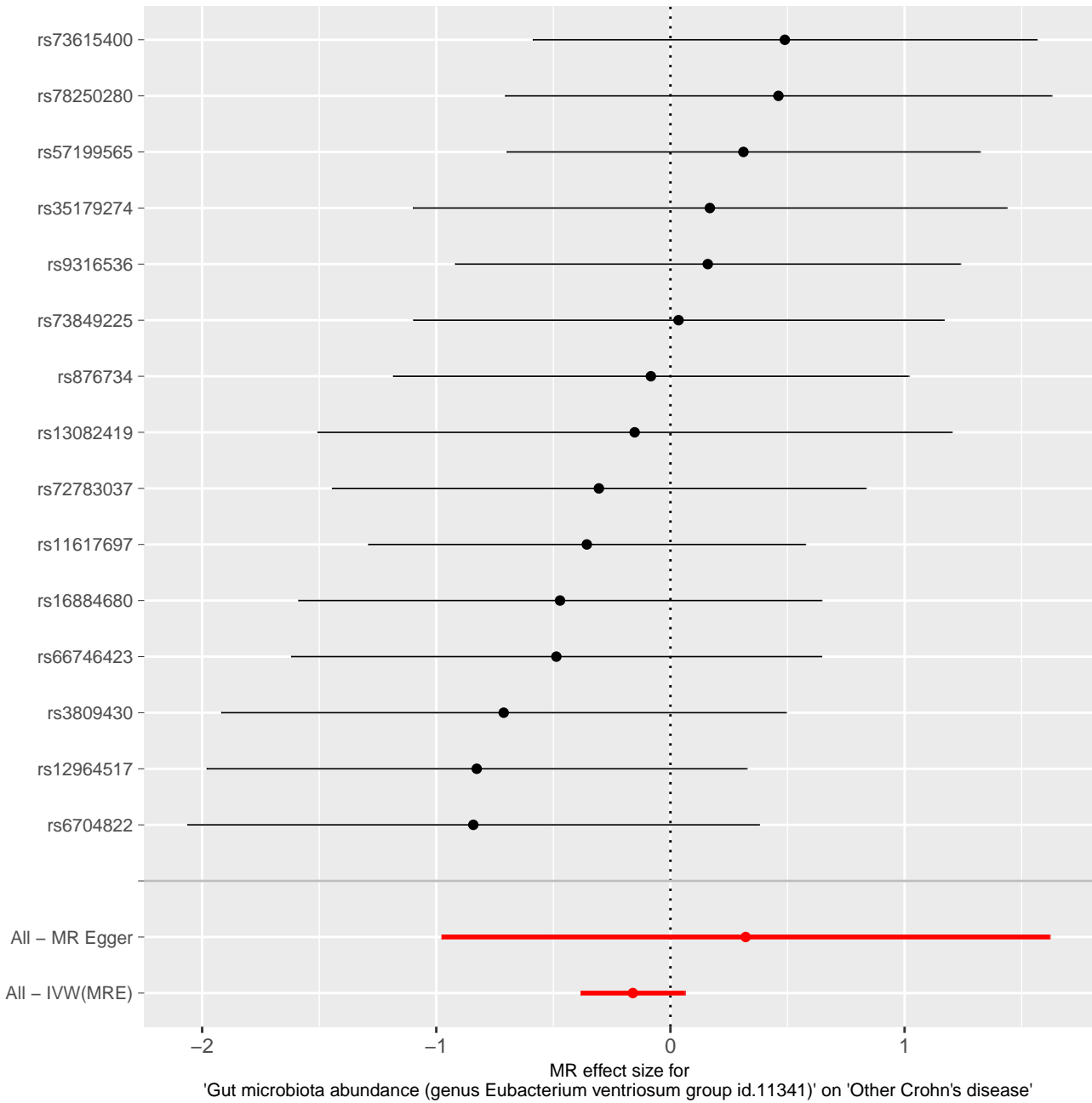


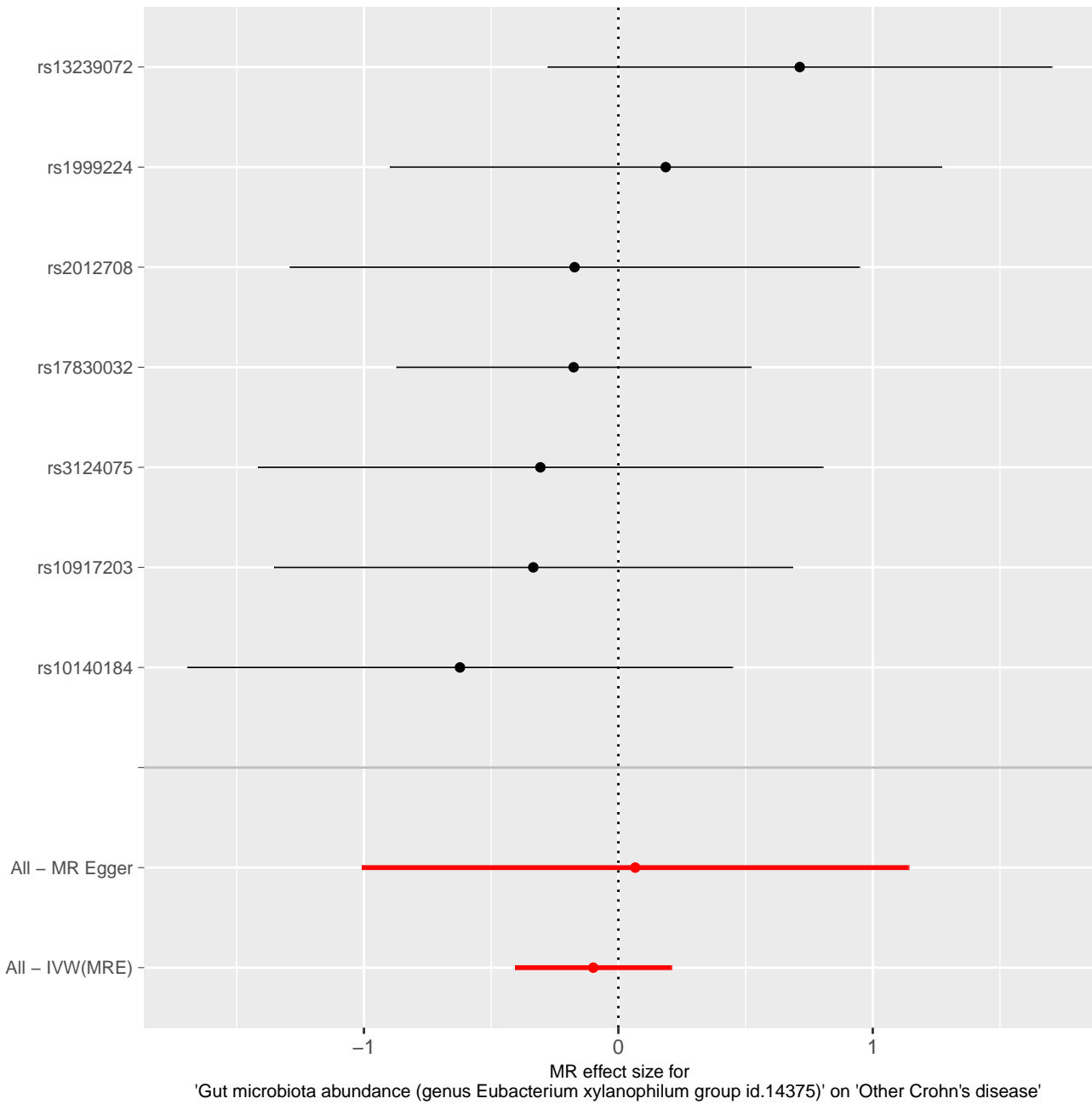




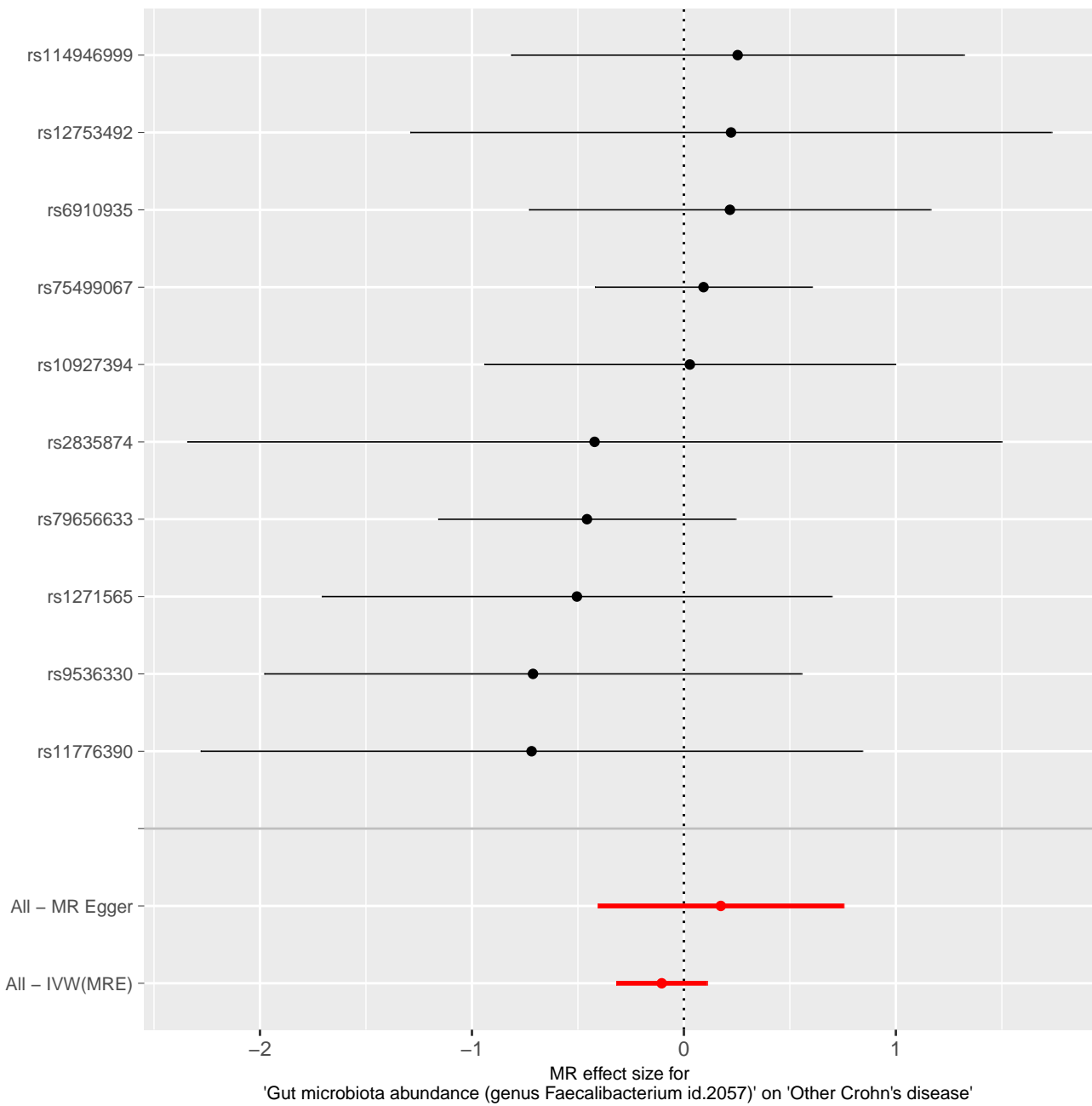
Batch 228 : Gut microbiota abundance (genus Eubacterium ruminantium group id.11340) on Other Crohn's disease

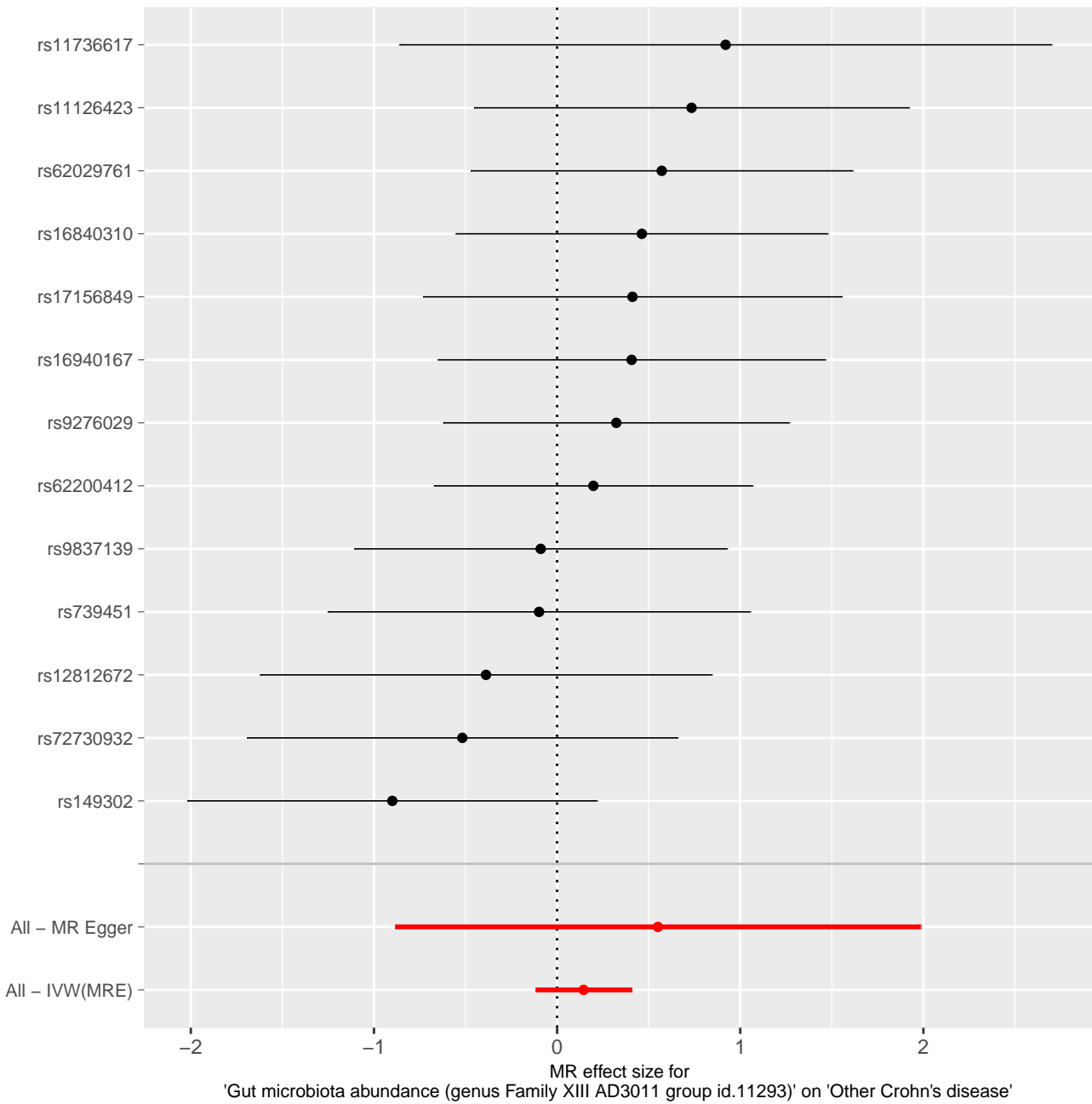


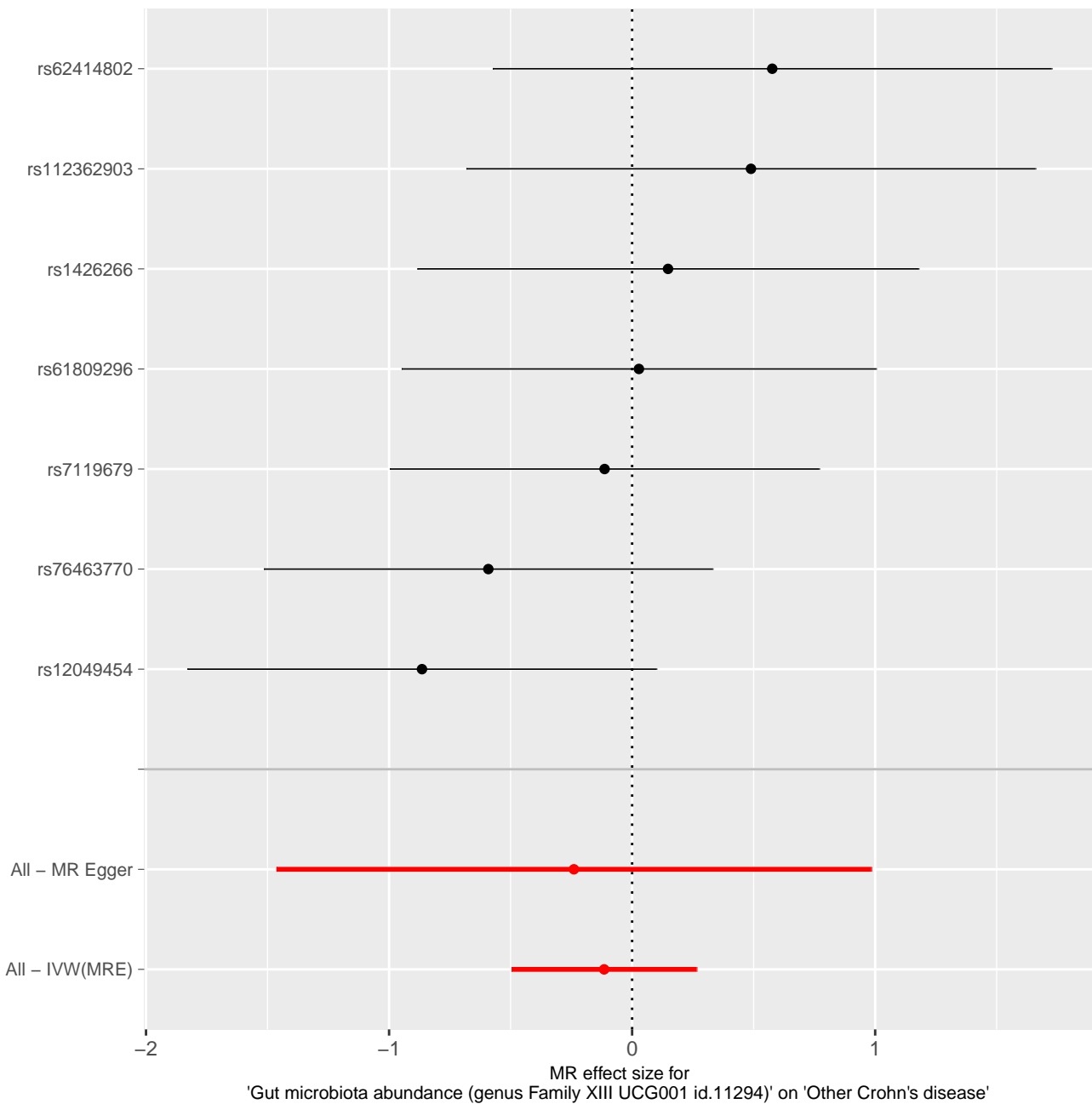


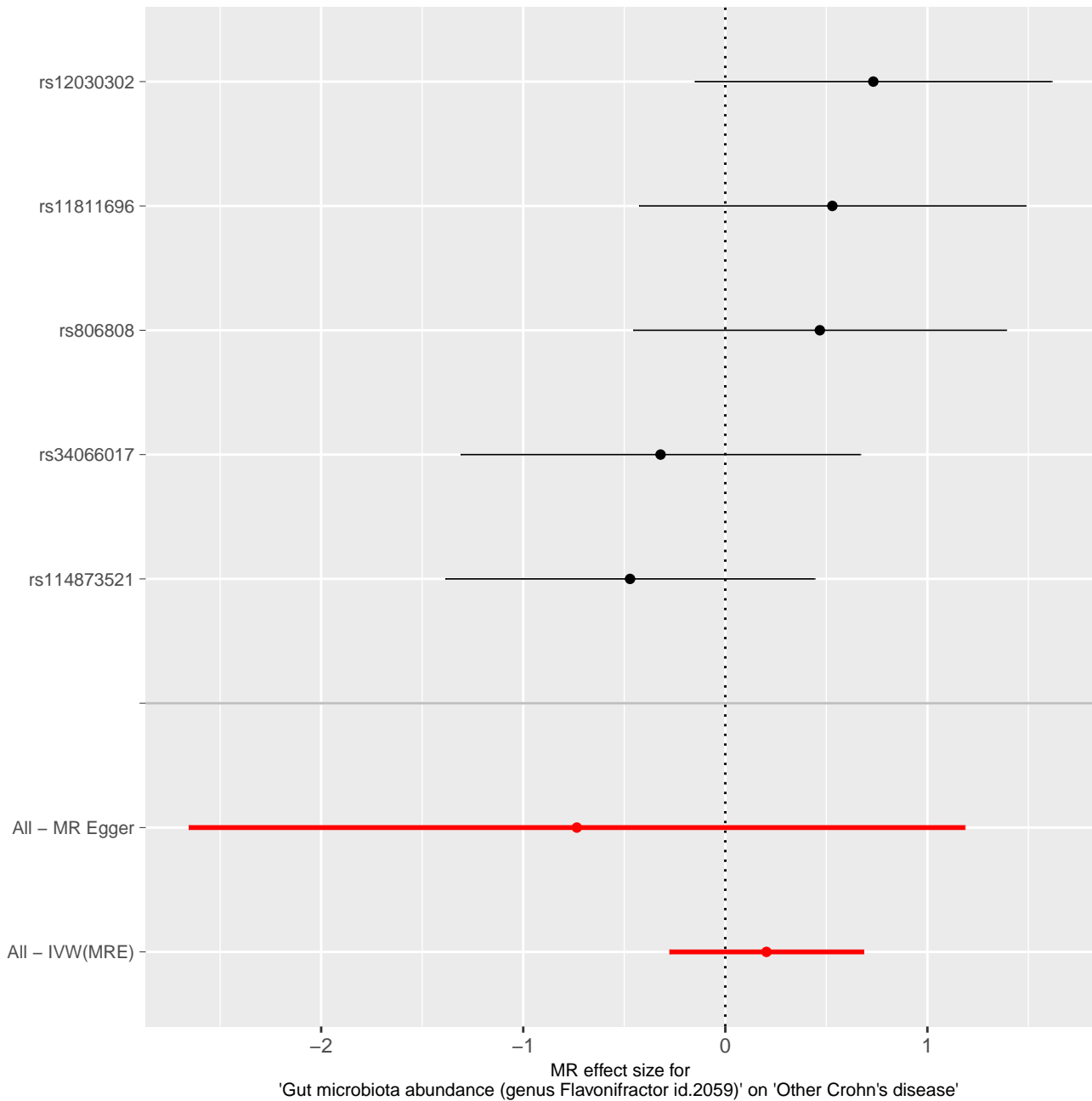


MR effect size for
'Gut microbiota abundance (genus Eubacterium xylanophilum group id.14375)' on 'Other Crohn's disease'

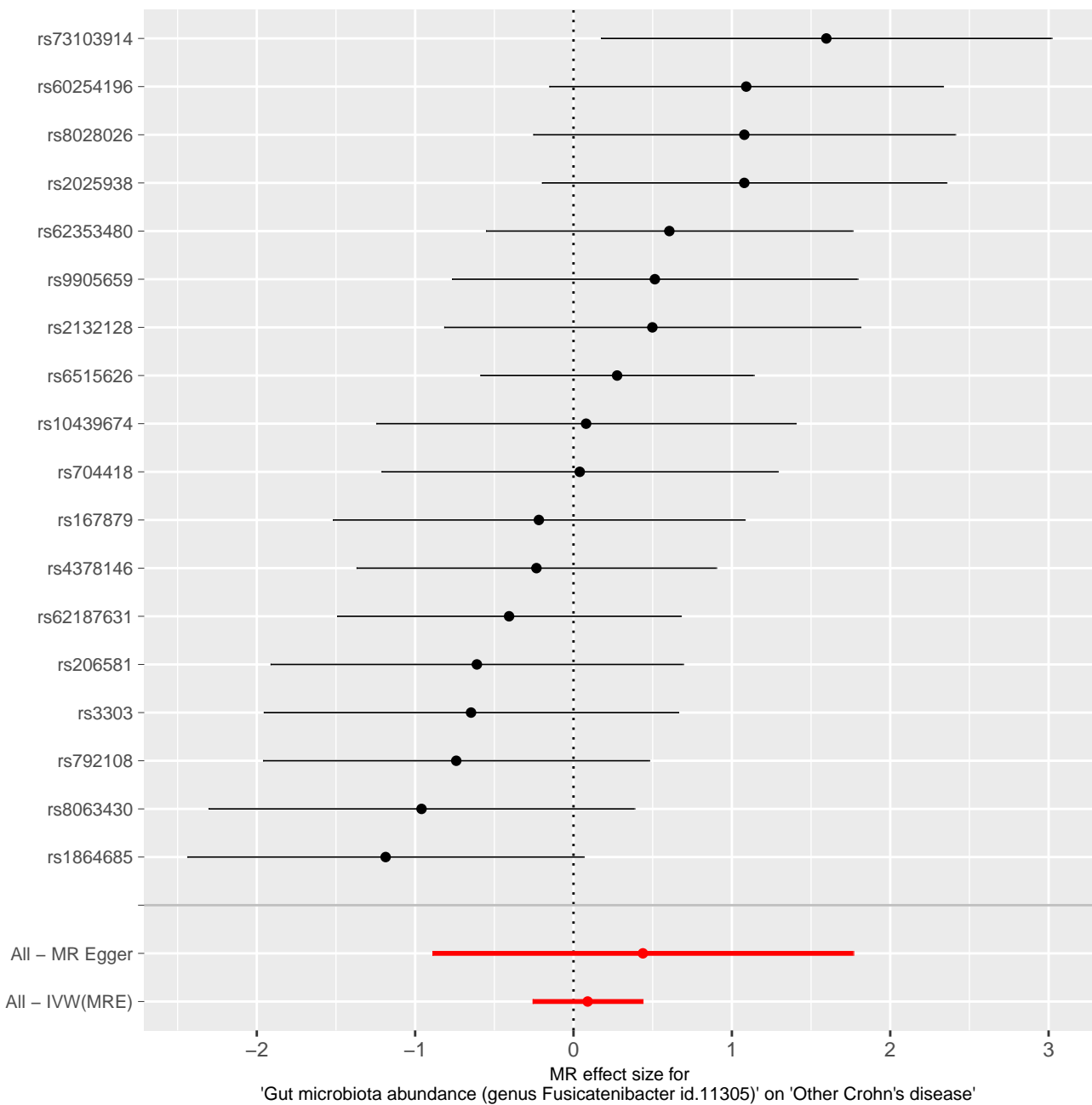


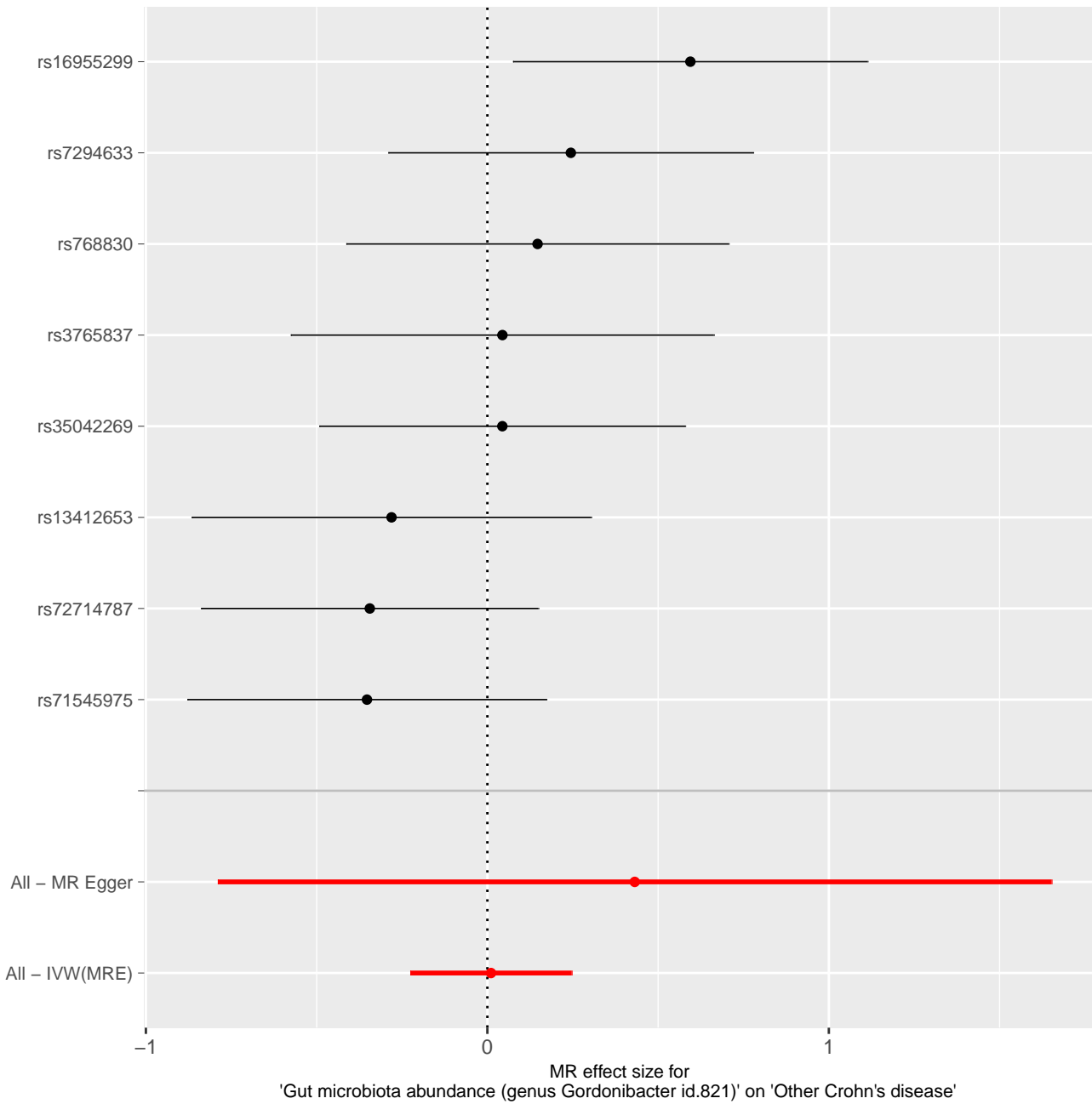


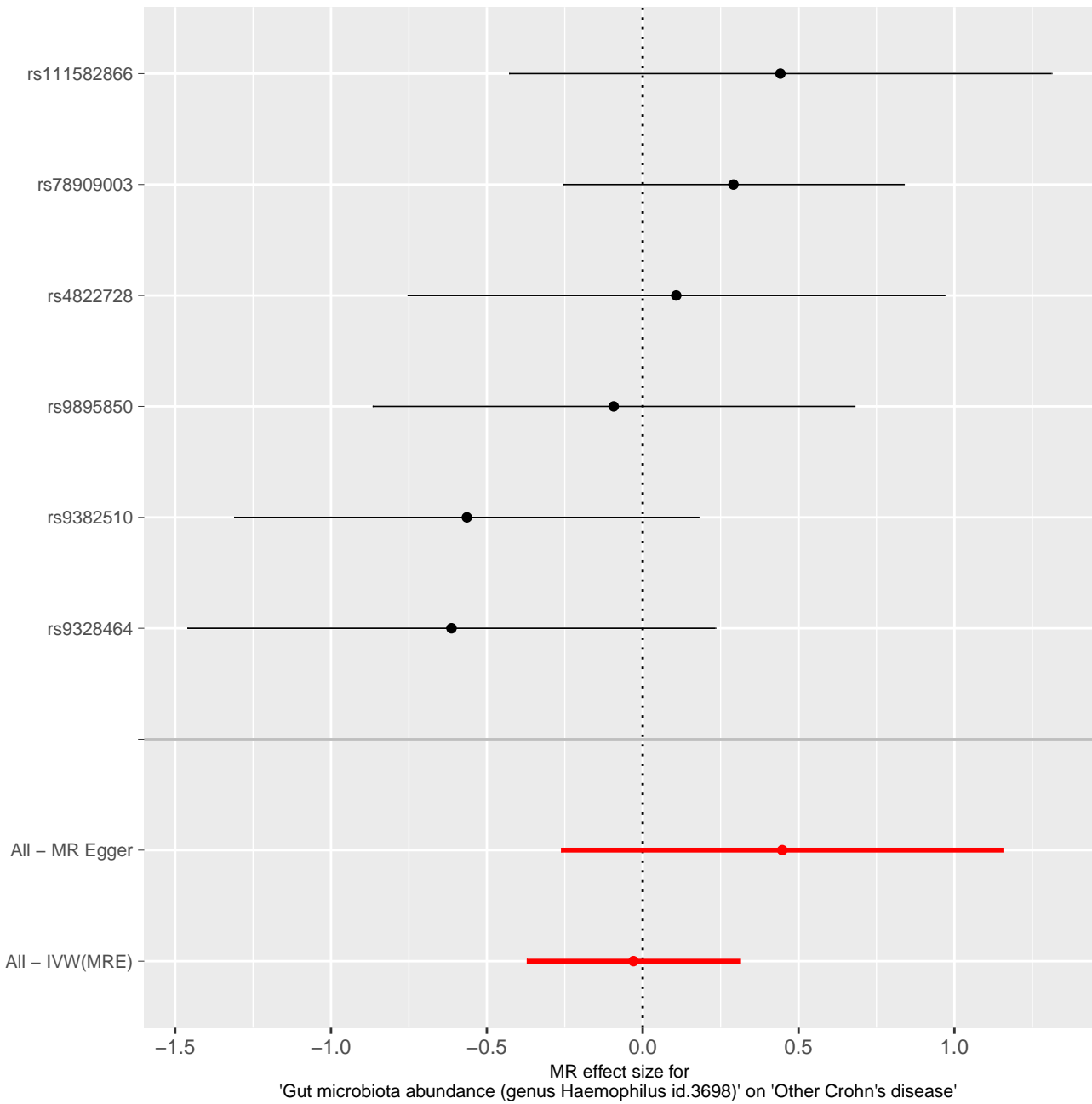




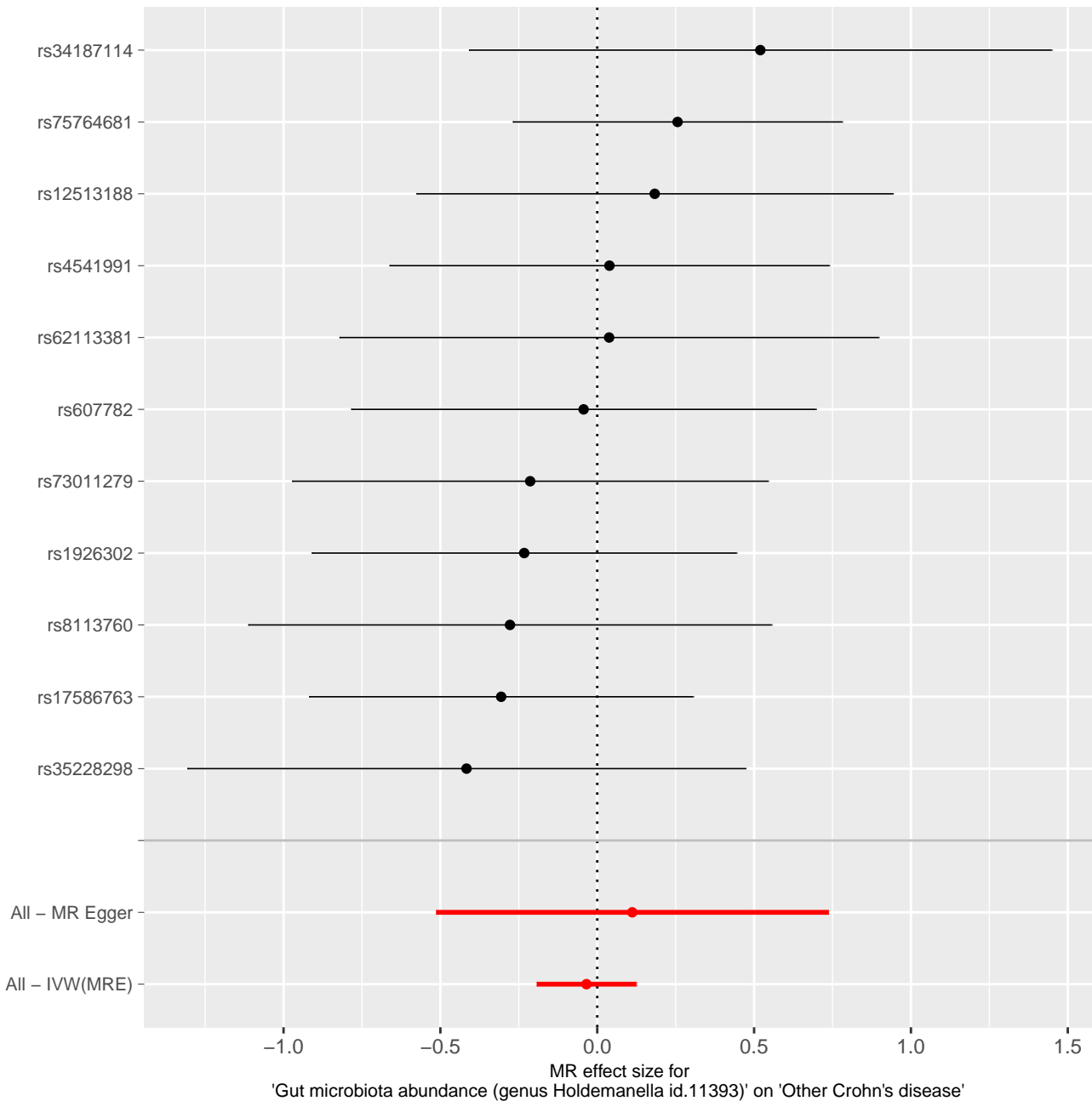
Batch 235 : Gut microbiota abundance (genus Fusicatenibacter id.11305) on Other Crohn's disease

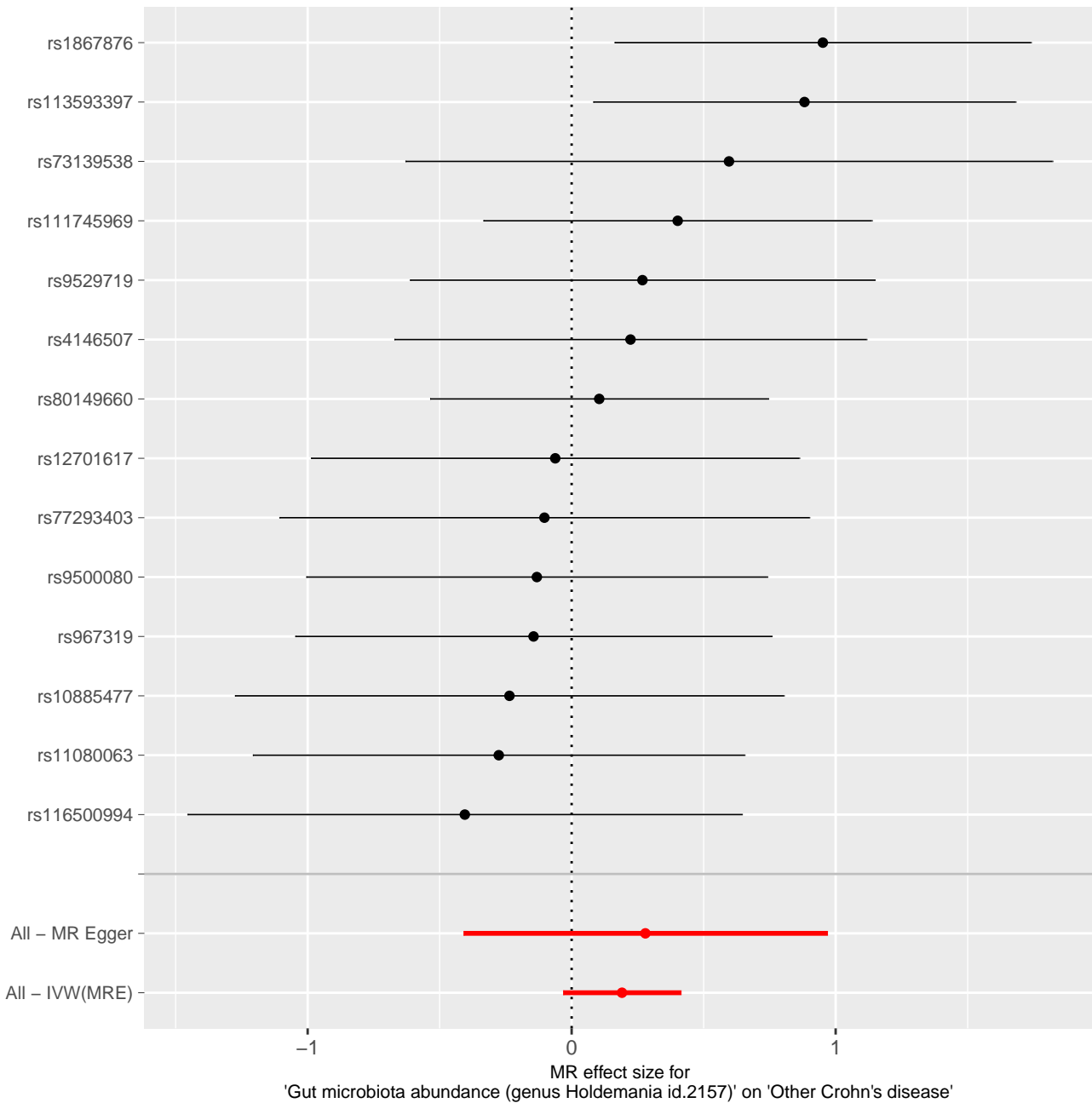


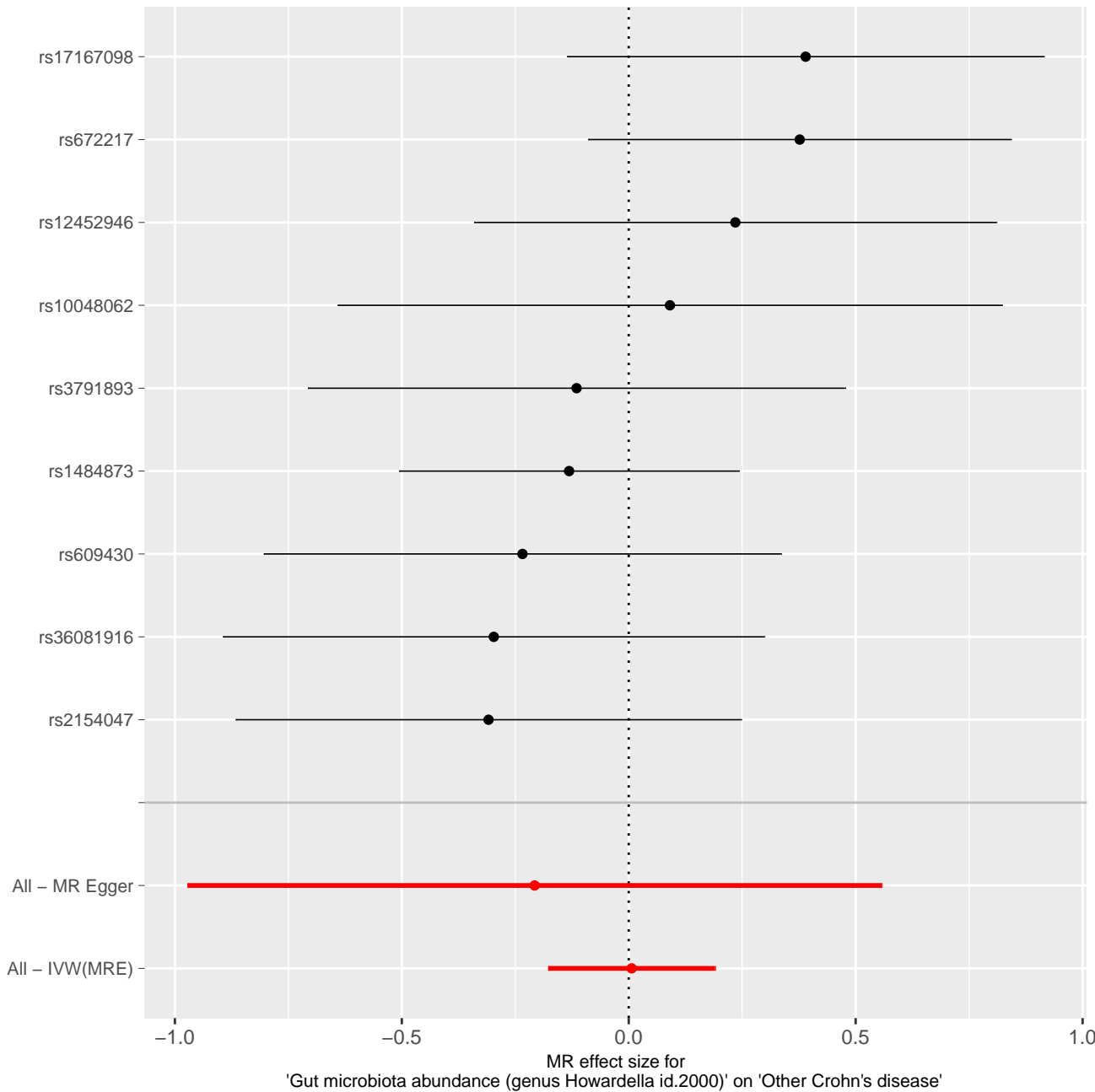




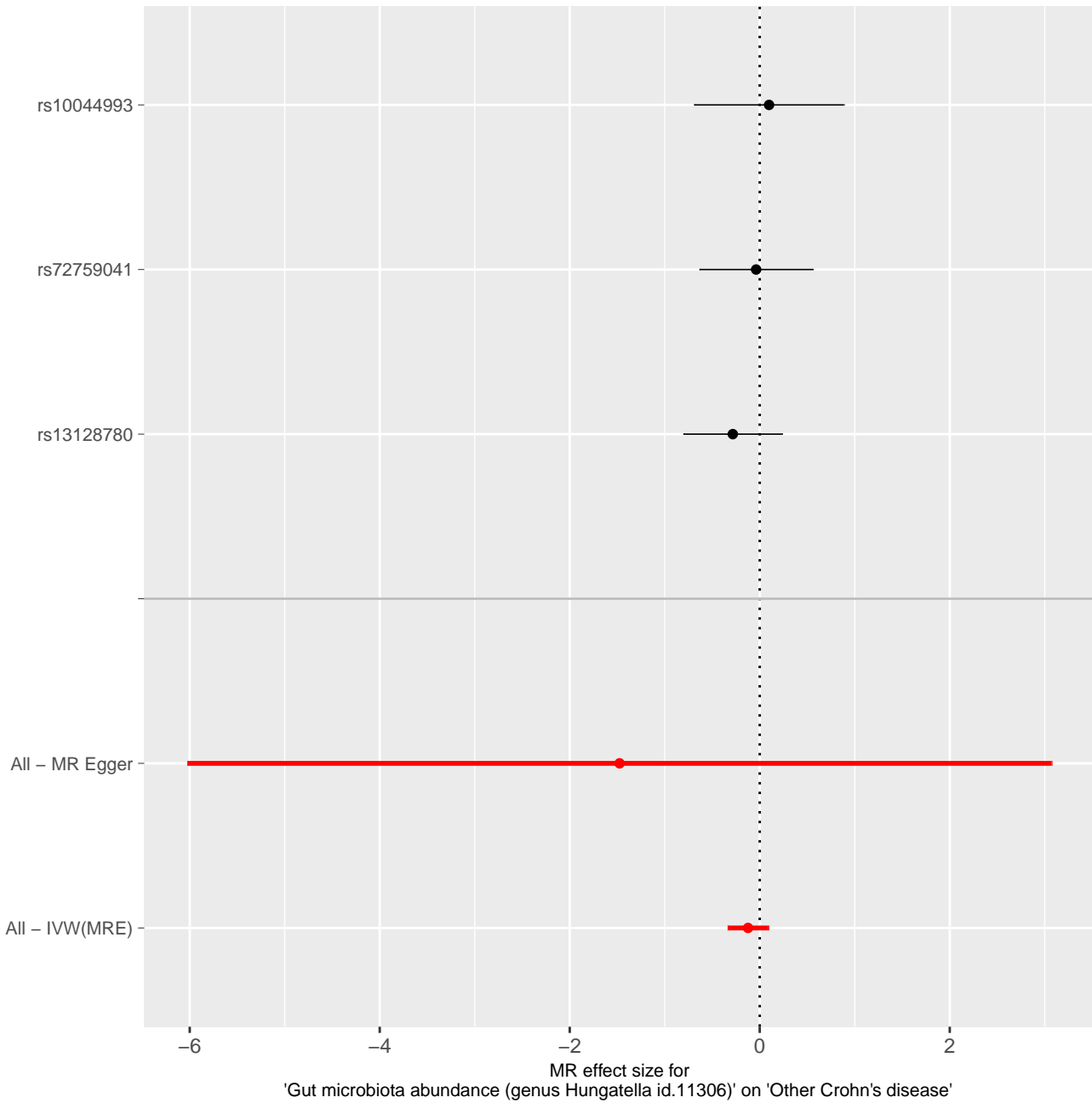
Batch 238 : Gut microbiota abundance (genus Holdemanella id.11393) on Other Crohn's disease



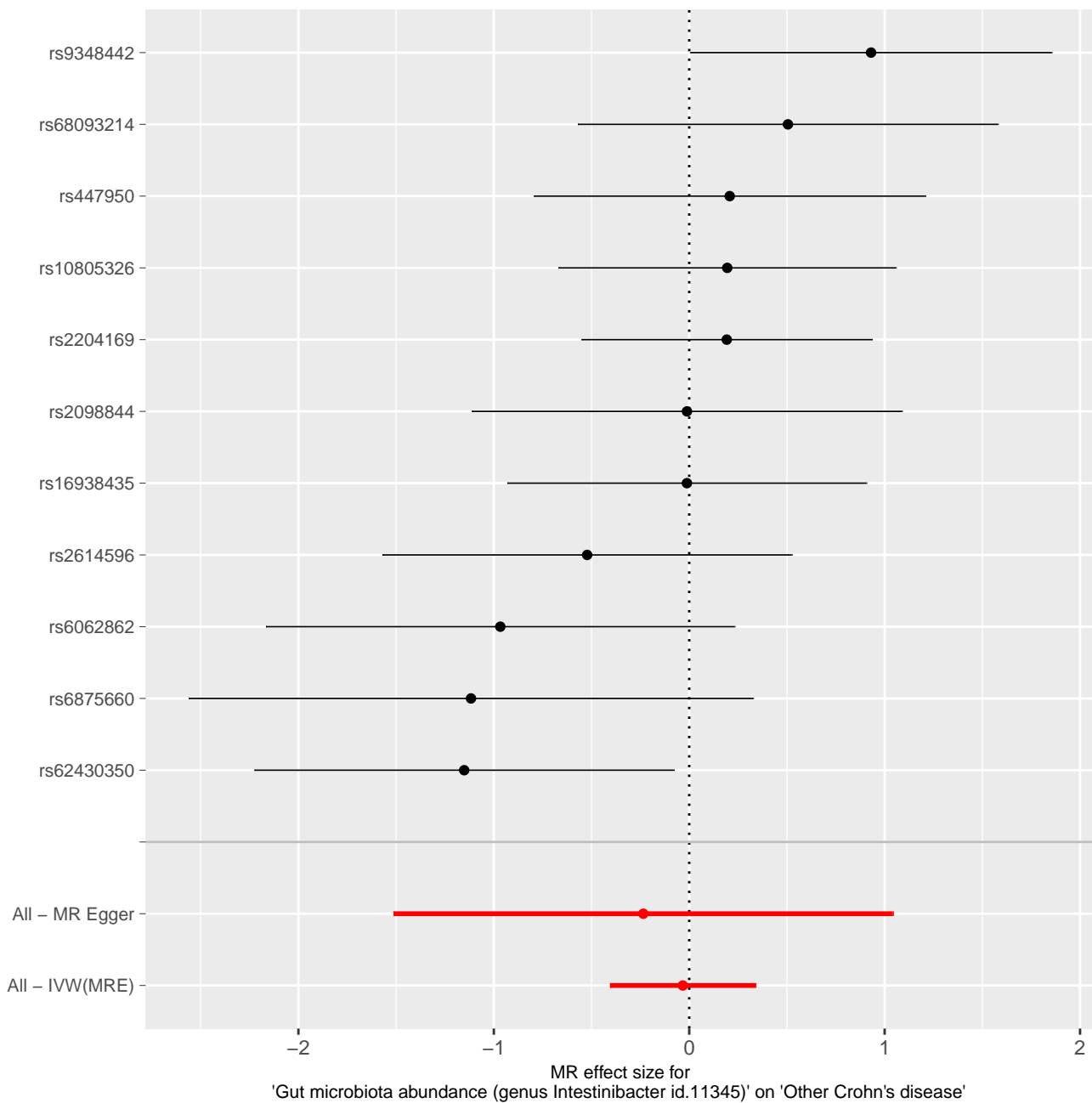


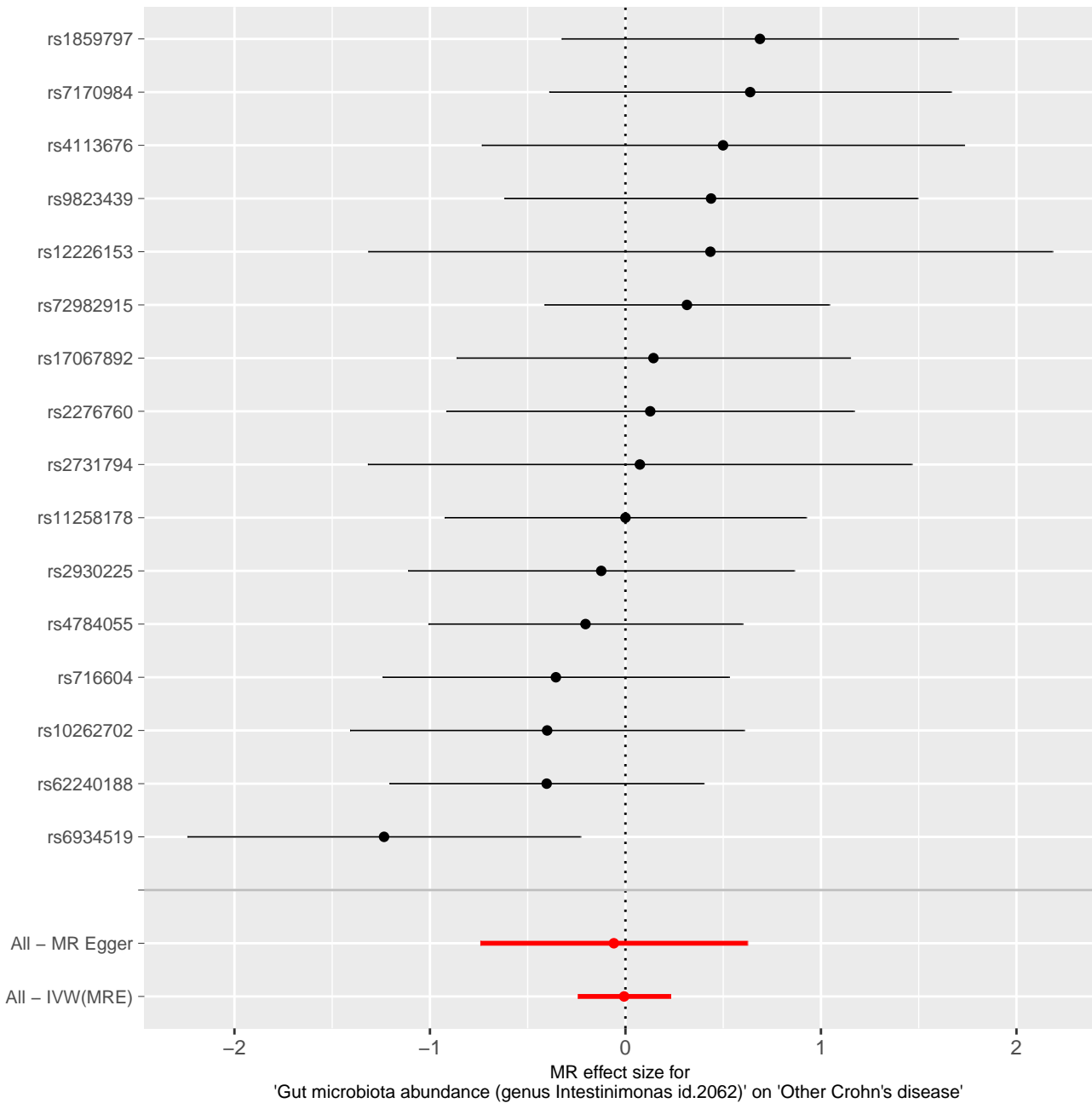


MR effect size for
'Gut microbiota abundance (genus Howardella id.2000)' on 'Other Crohn's disease'

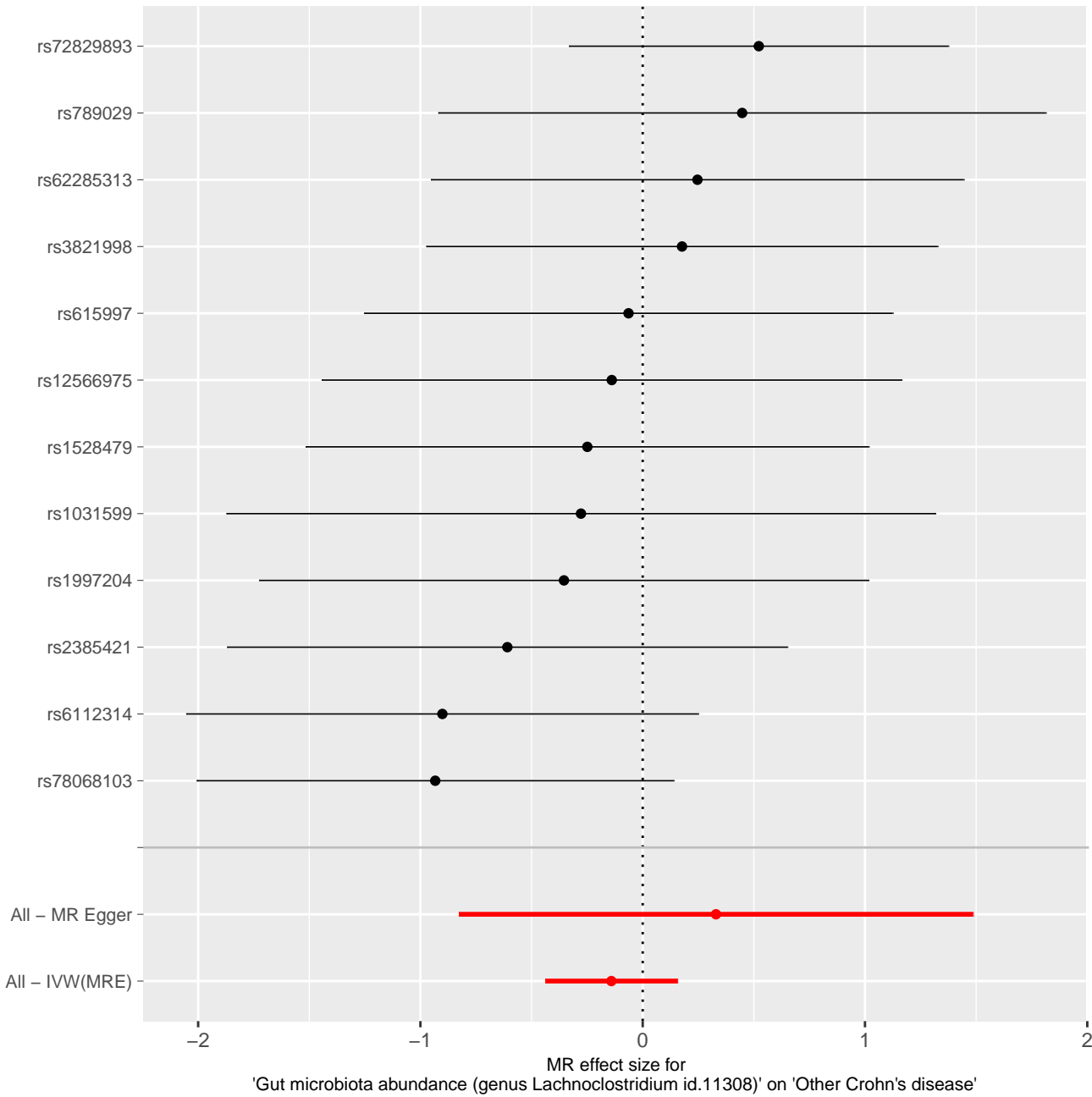


Batch 242 : Gut microbiota abundance (genus Intestinibacter id.11345) on Other Crohn's disease

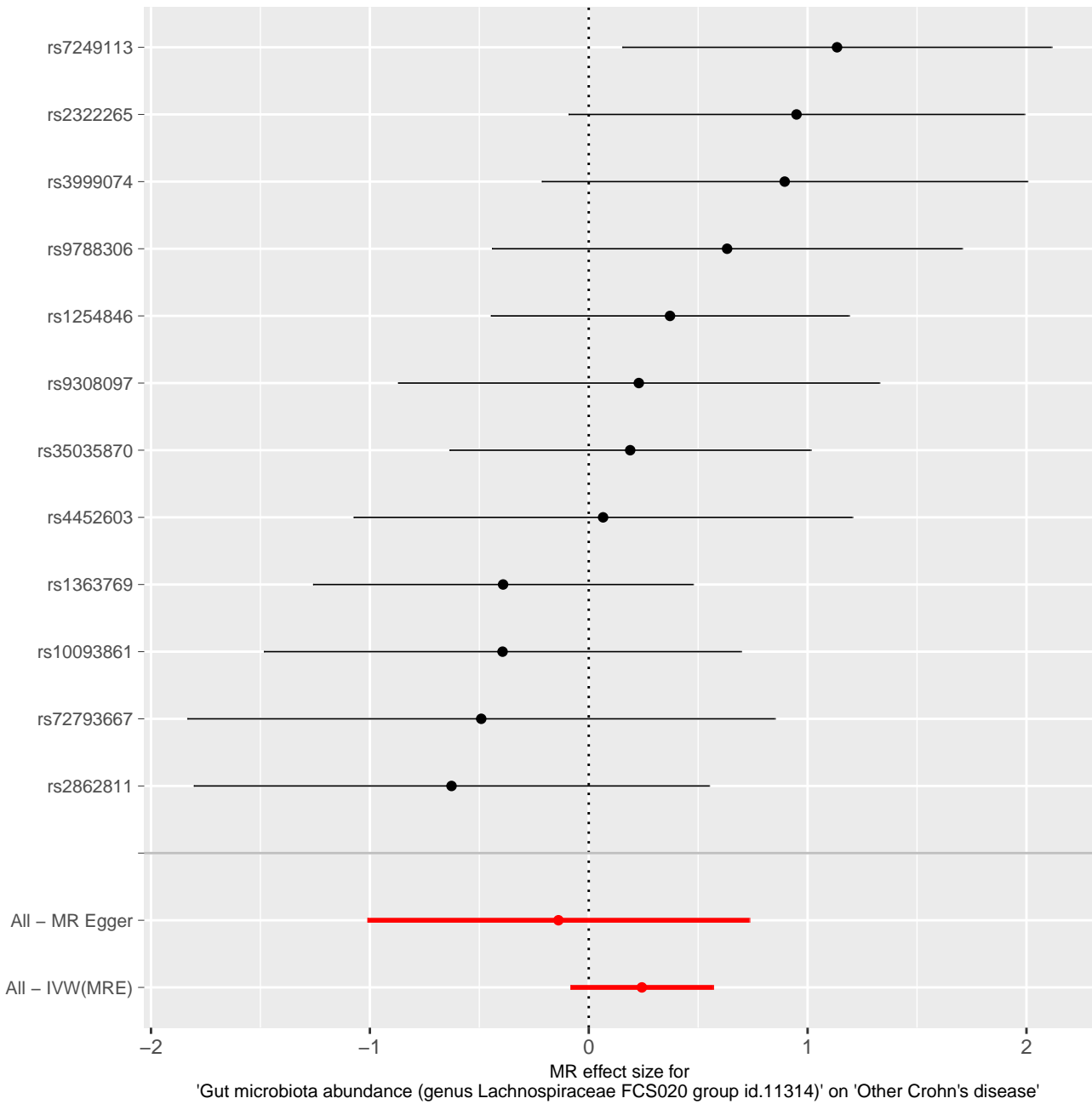


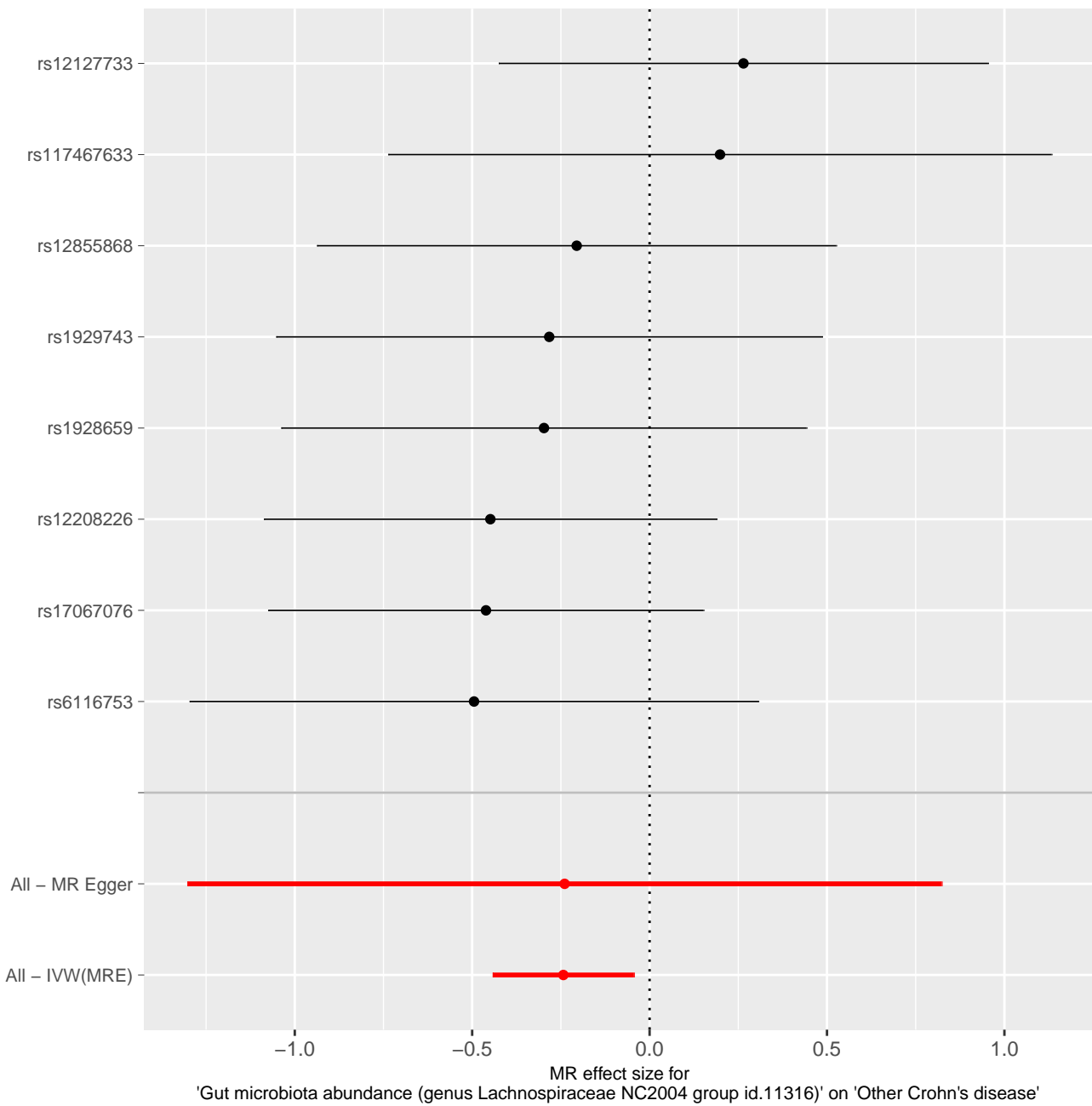


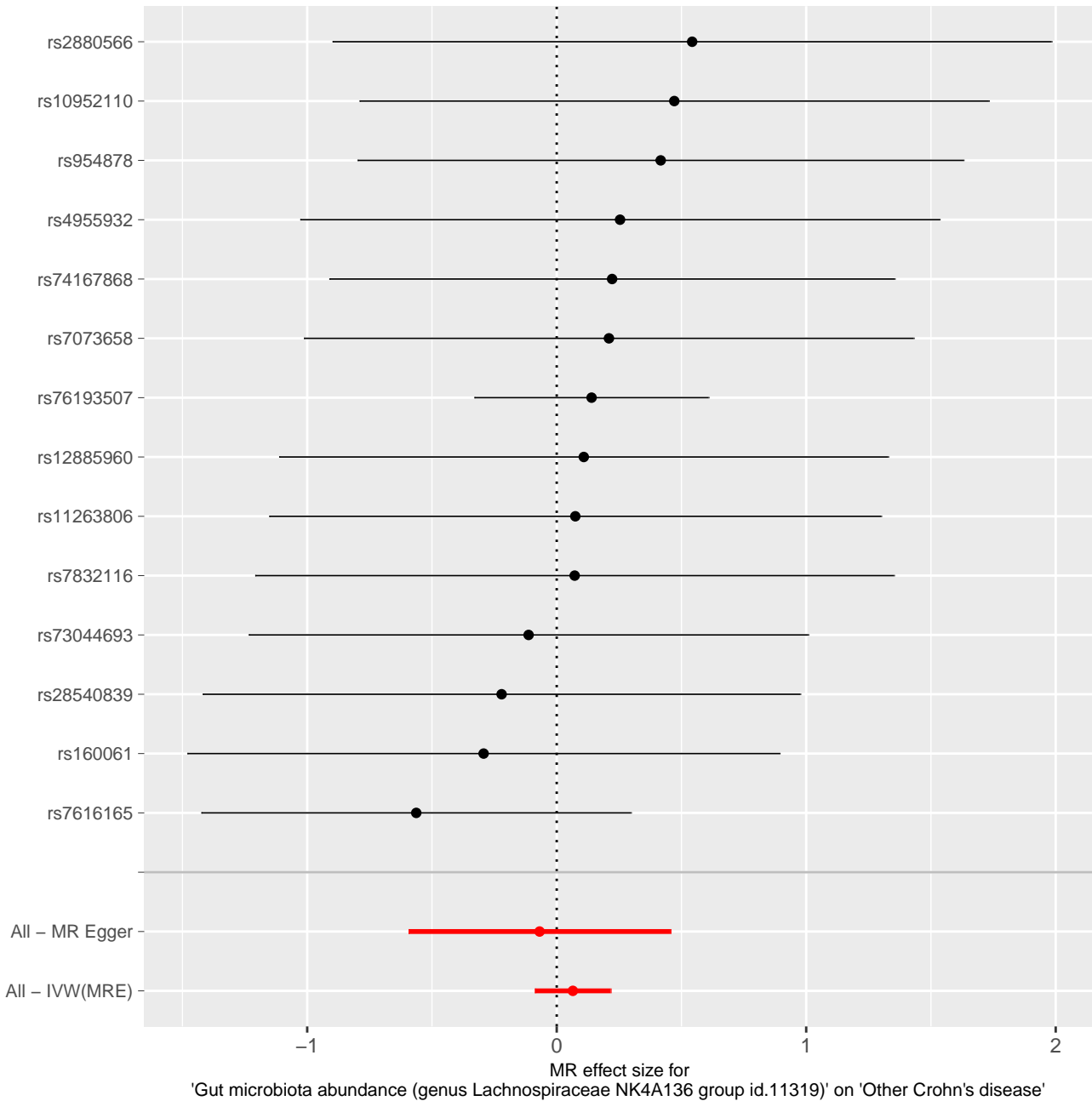
Batch 244 : Gut microbiota abundance (genus Lachnoclostridium id.11308) on Other Crohn's disease

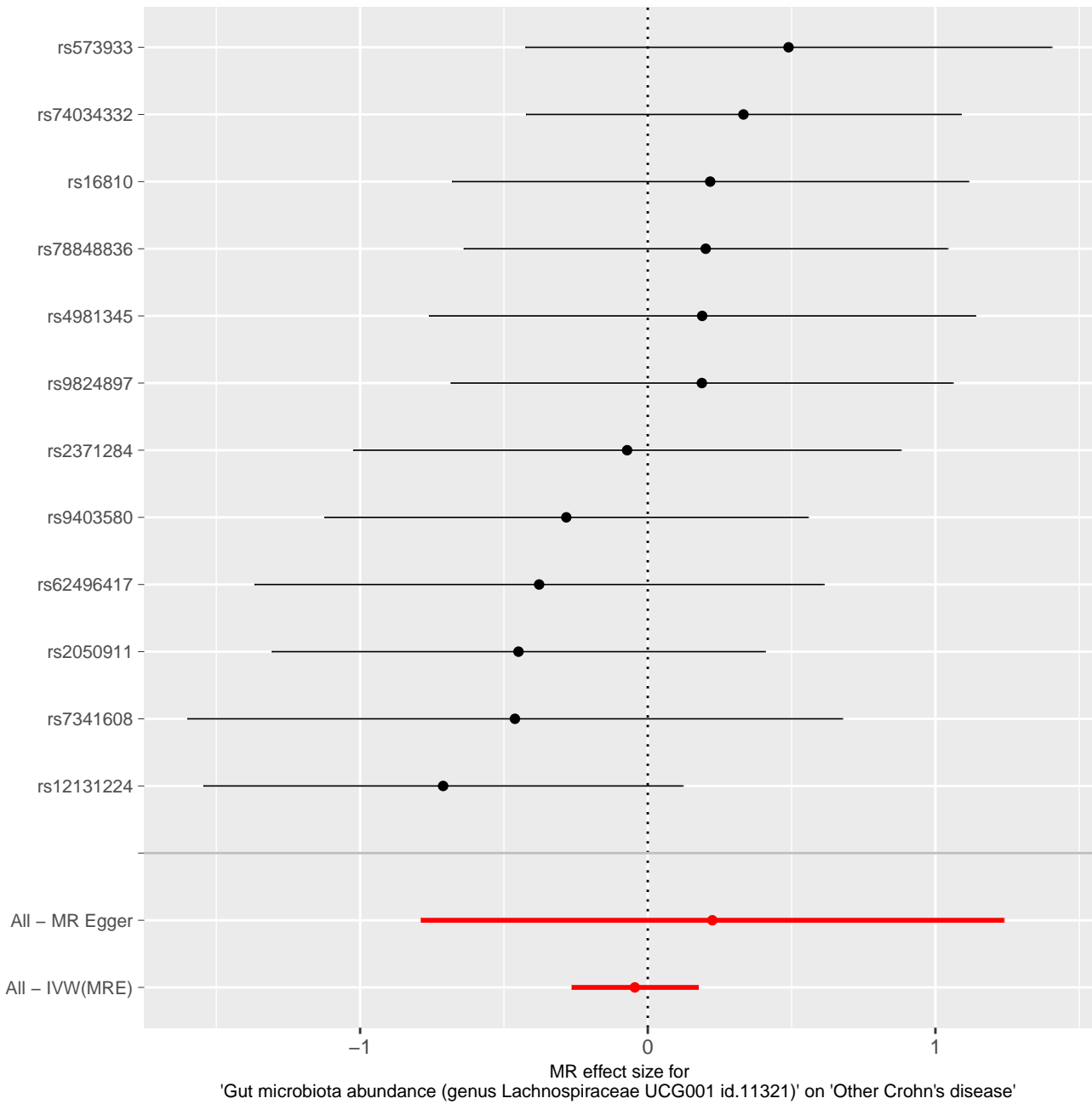


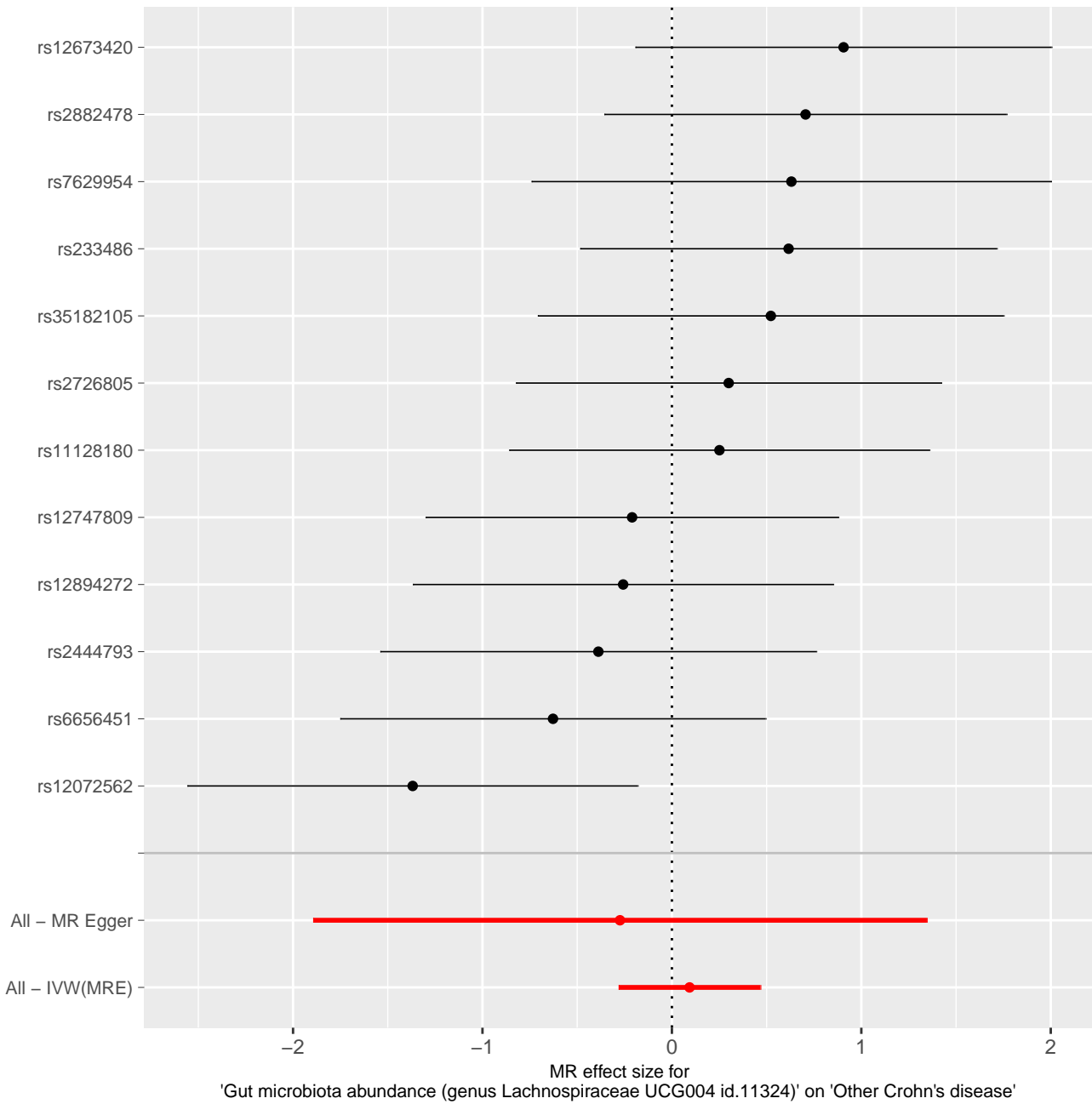
Batch 245 : Gut microbiota abundance (genus Lachnospiraceae FCS020 group id.11314) on Other Crohn's disease

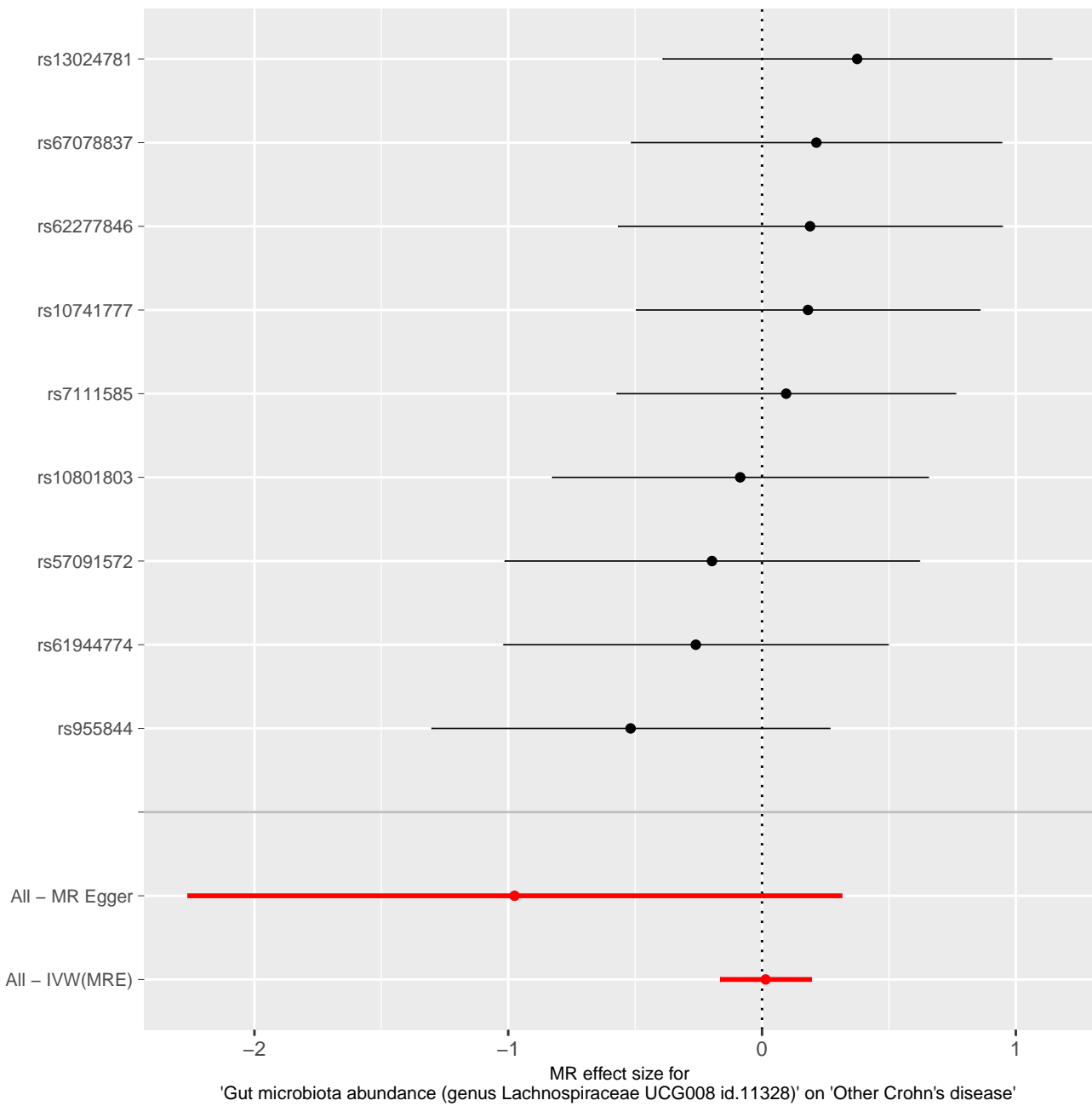


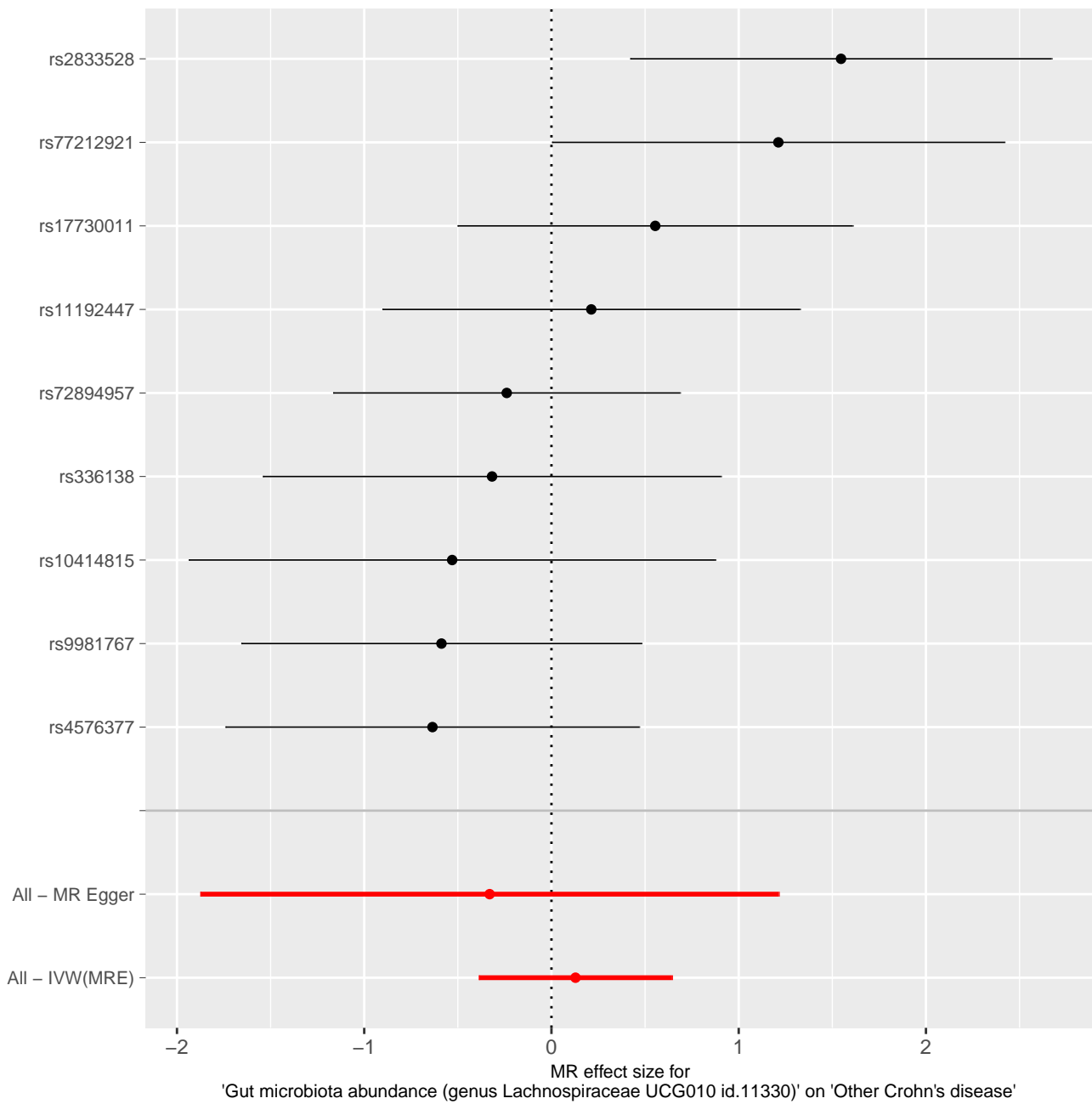


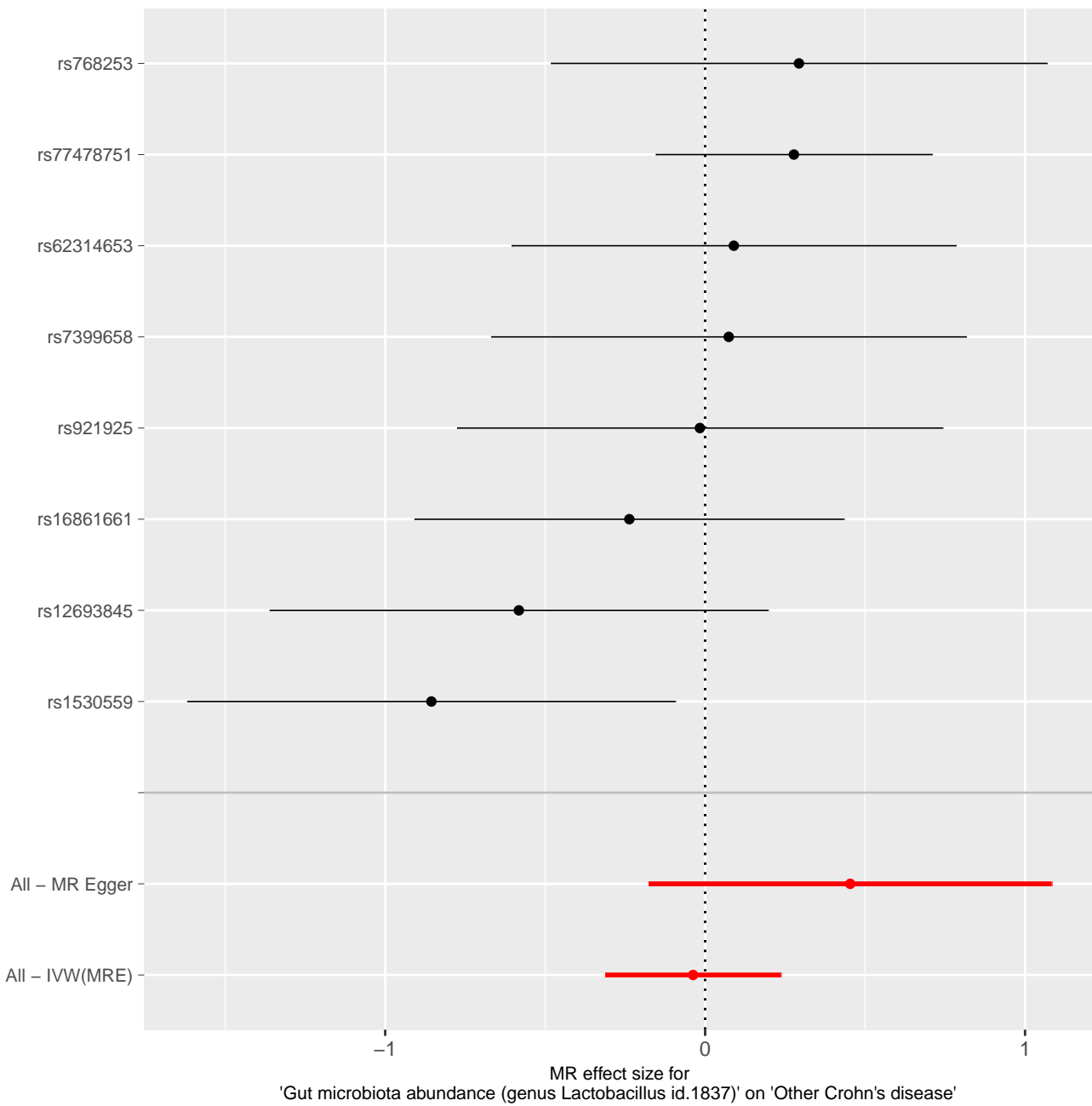


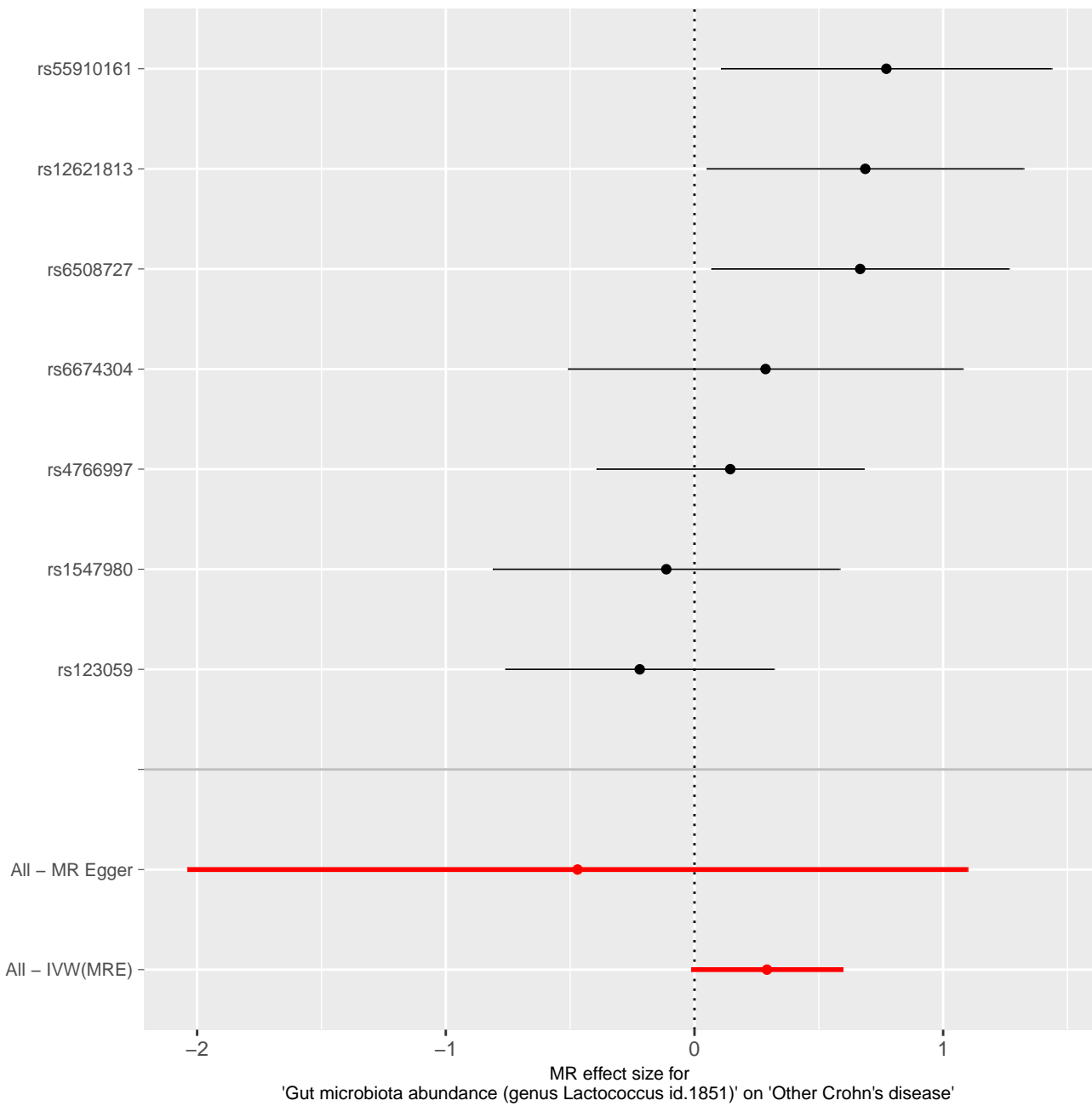


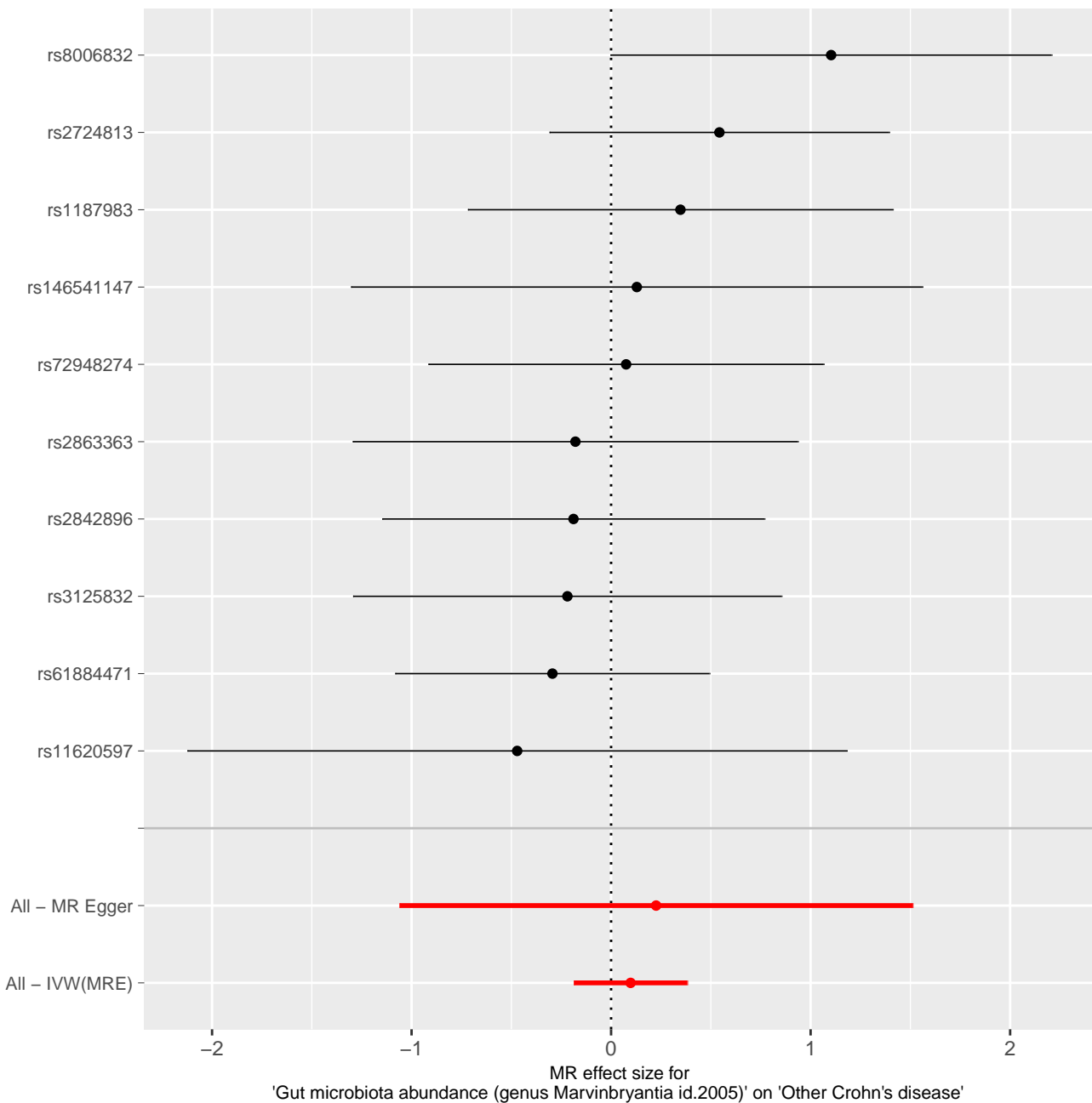


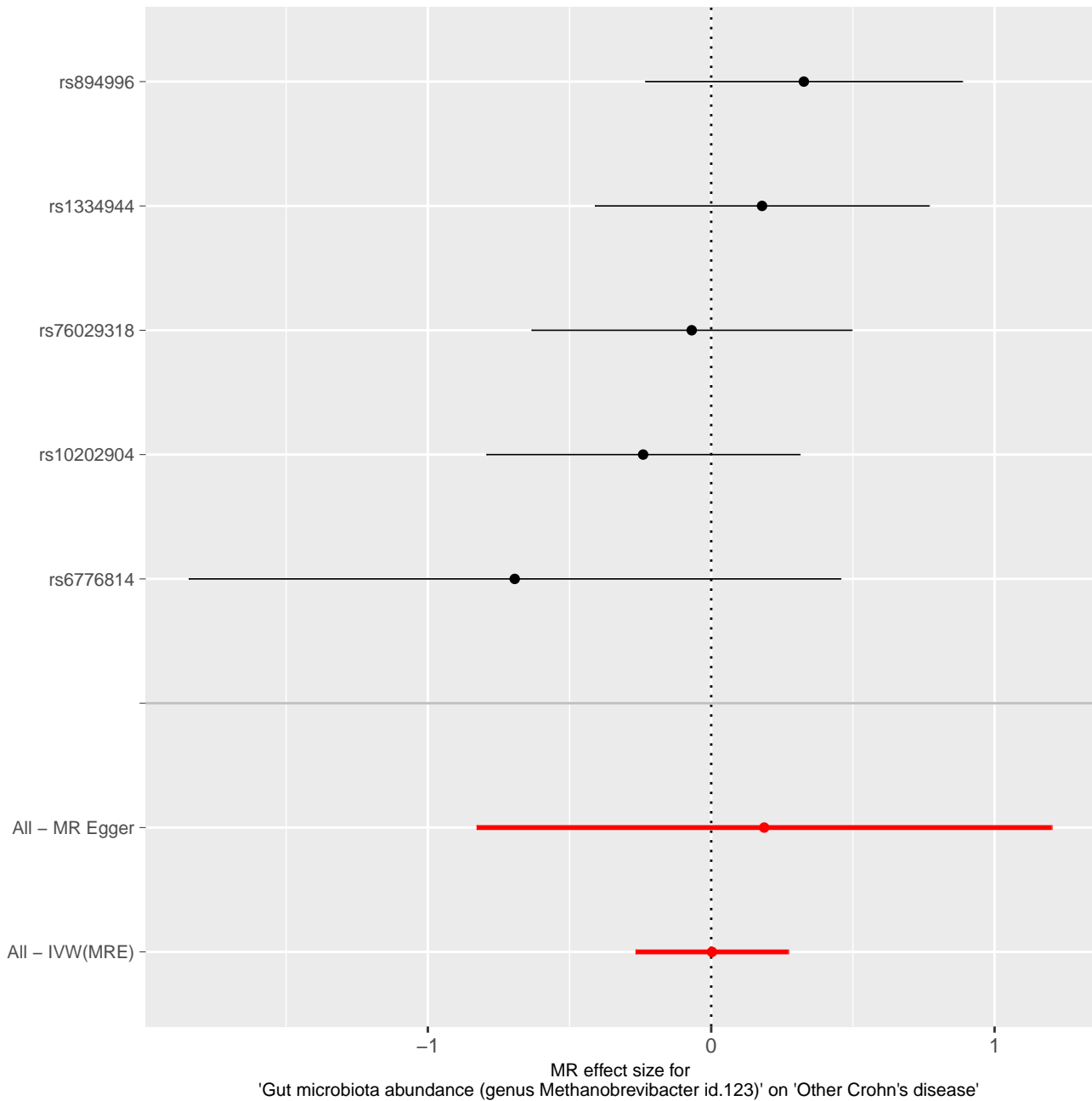


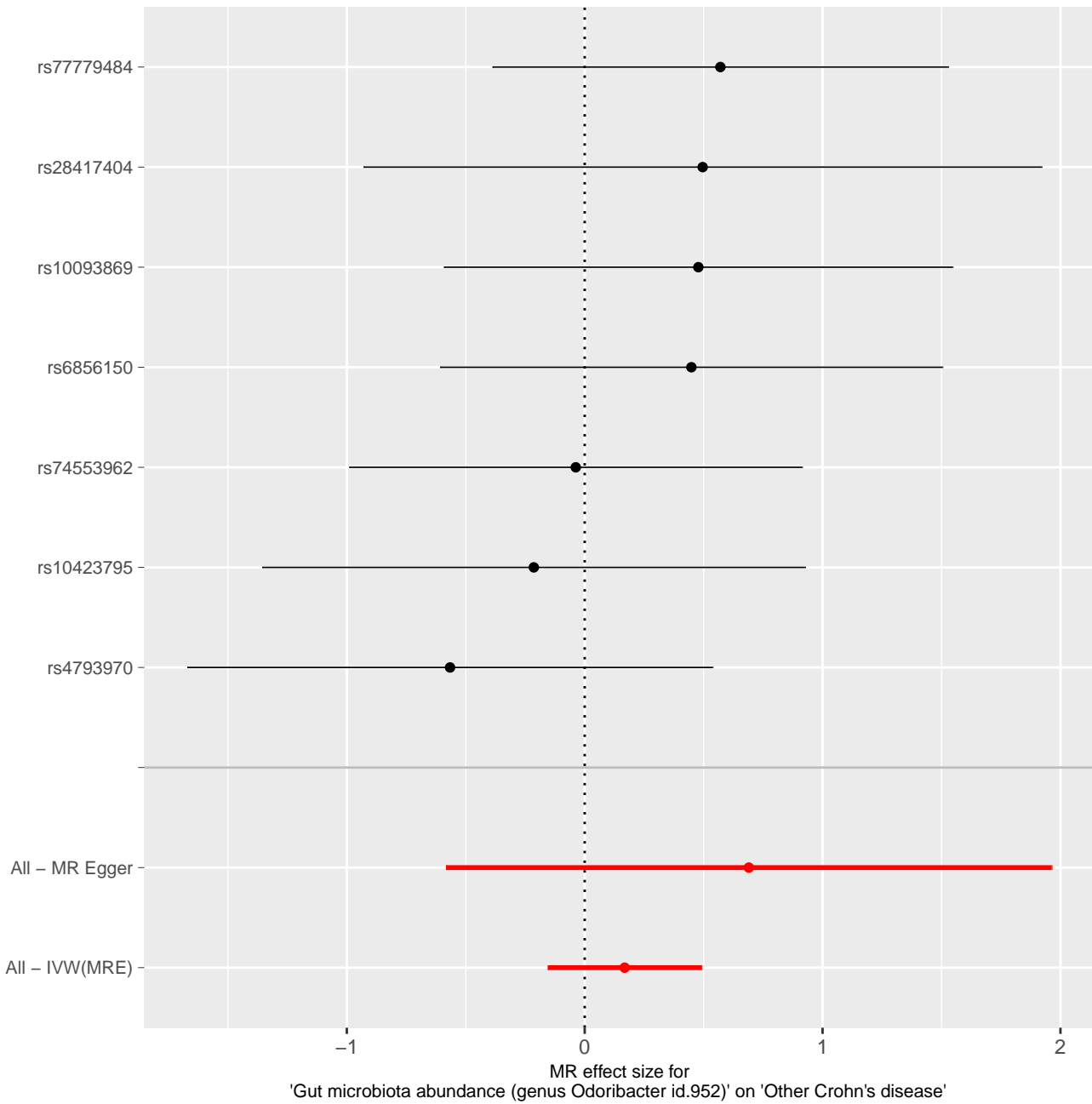




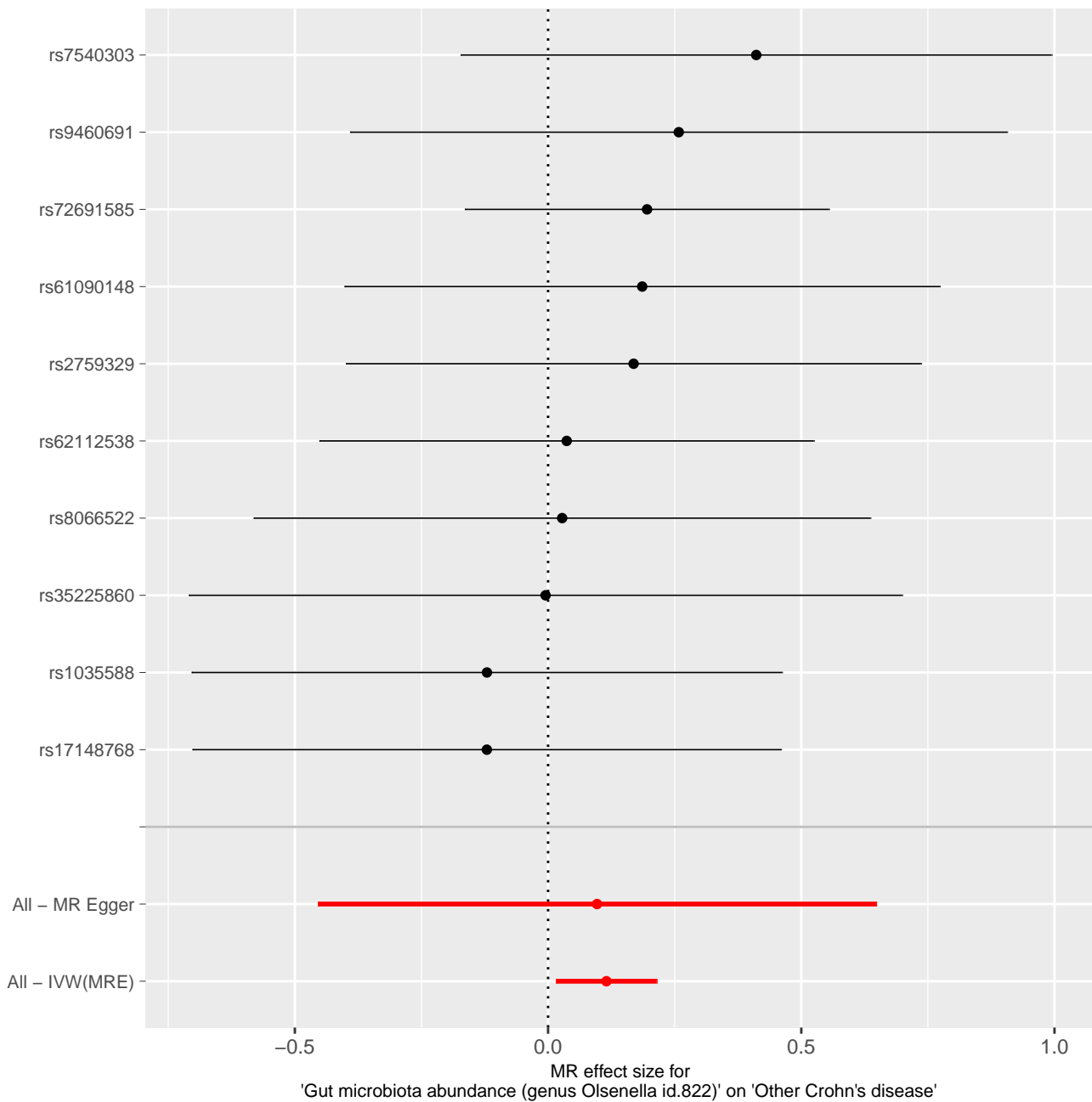




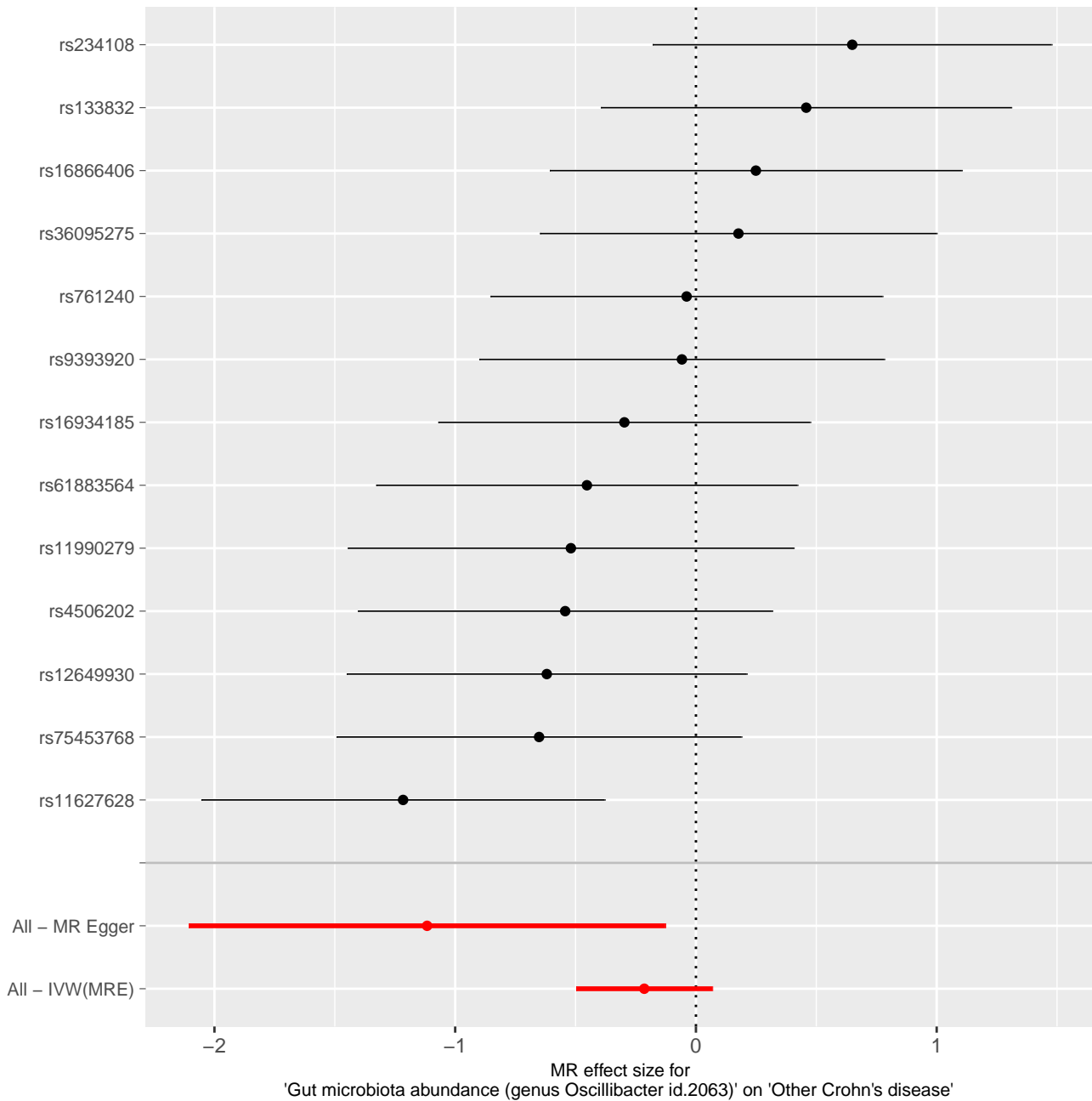


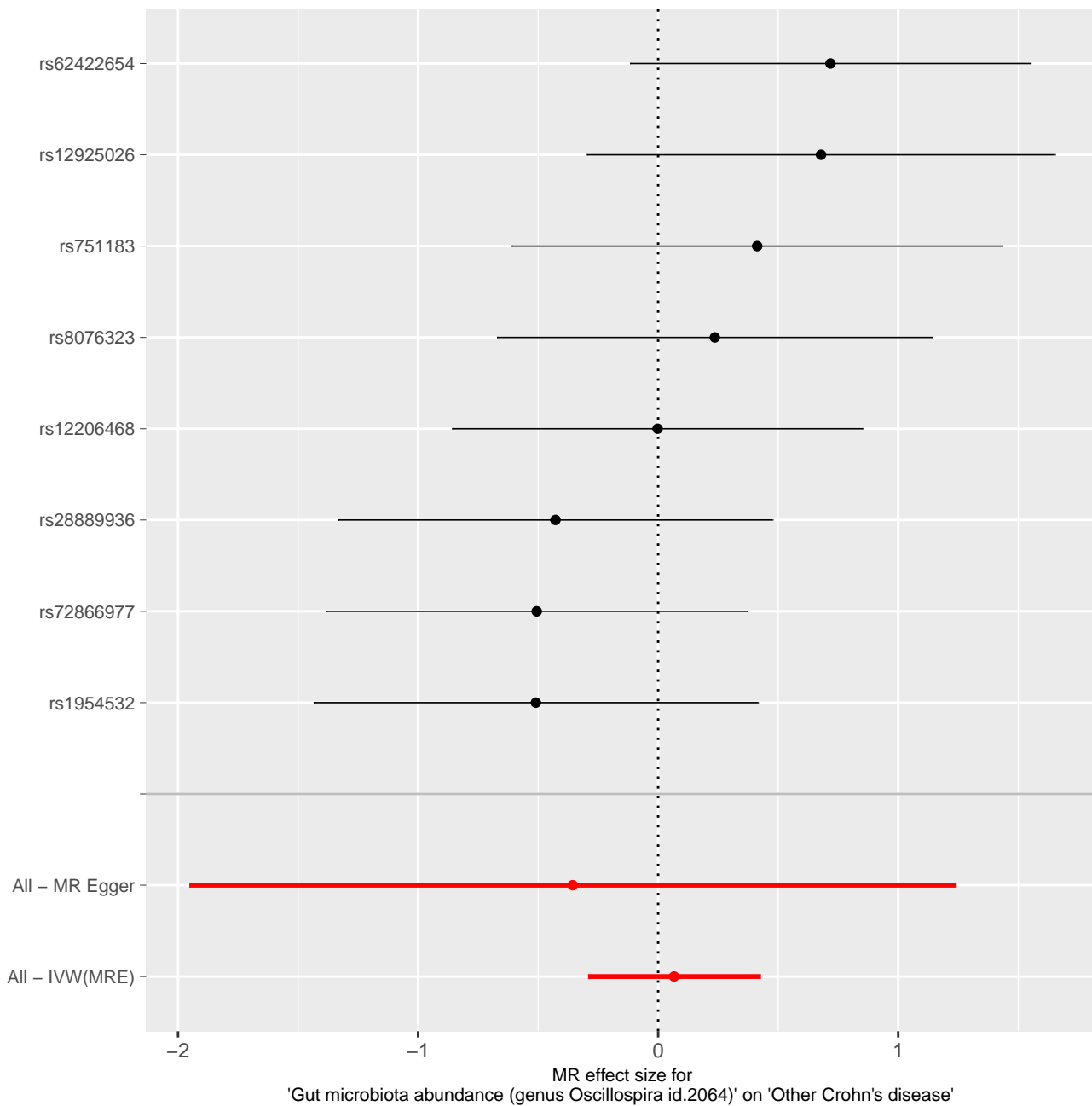


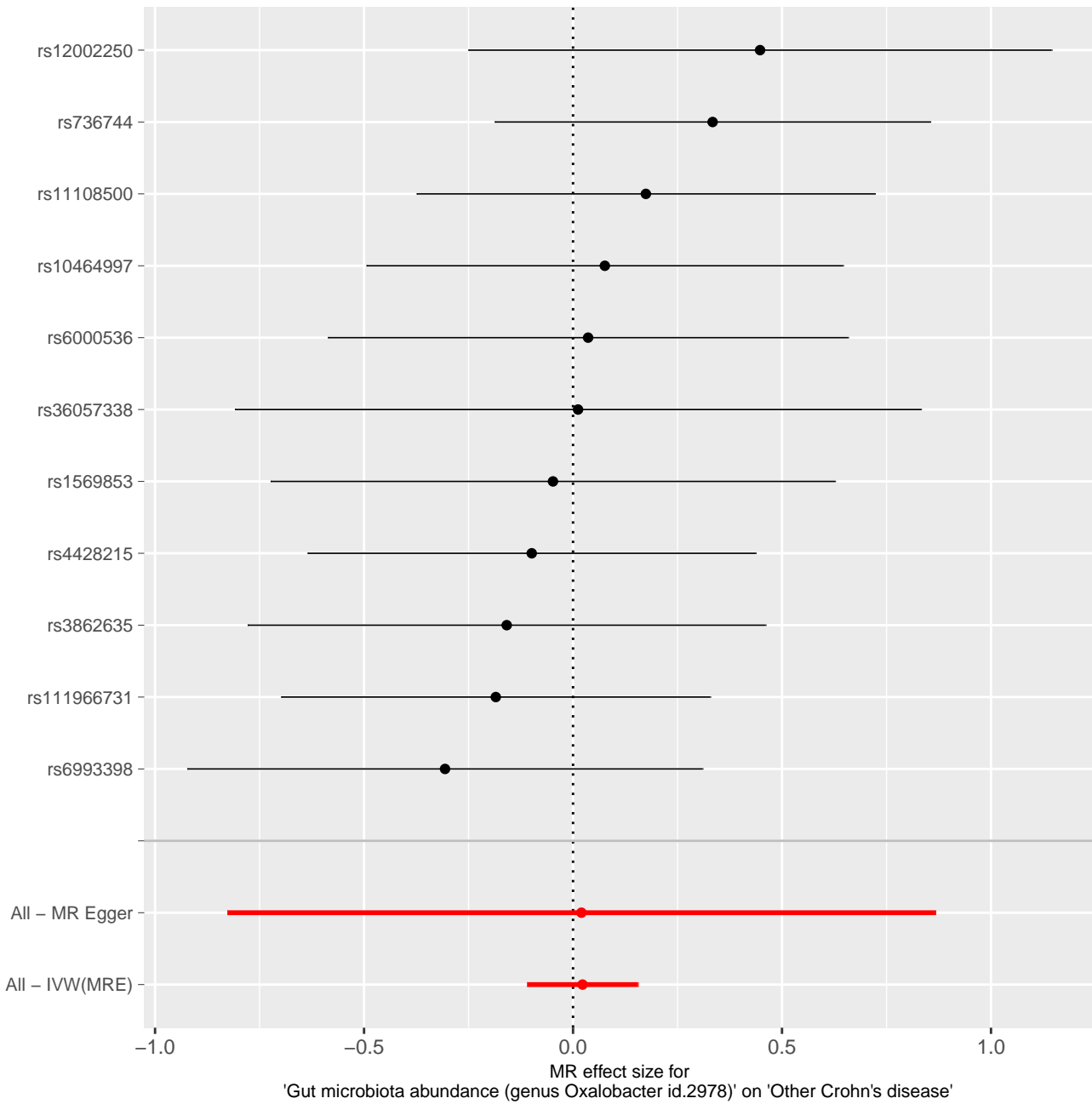
Batch 259 : Gut microbiota abundance (genus Olsenella id.822) on Other Crohn's disease

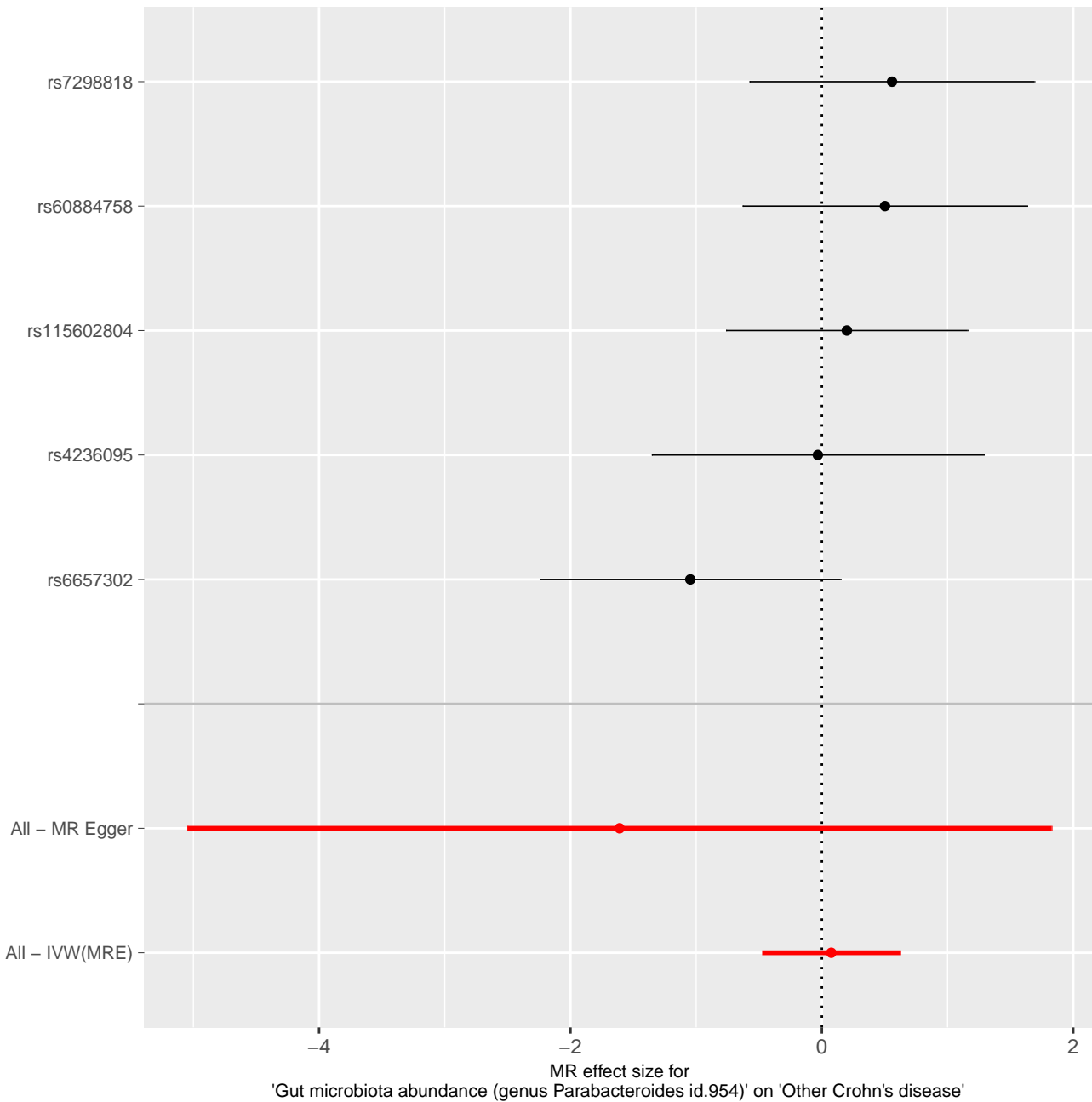


Batch 260 : Gut microbiota abundance (genus Oscillibacter id.2063) on Other Crohn's disease

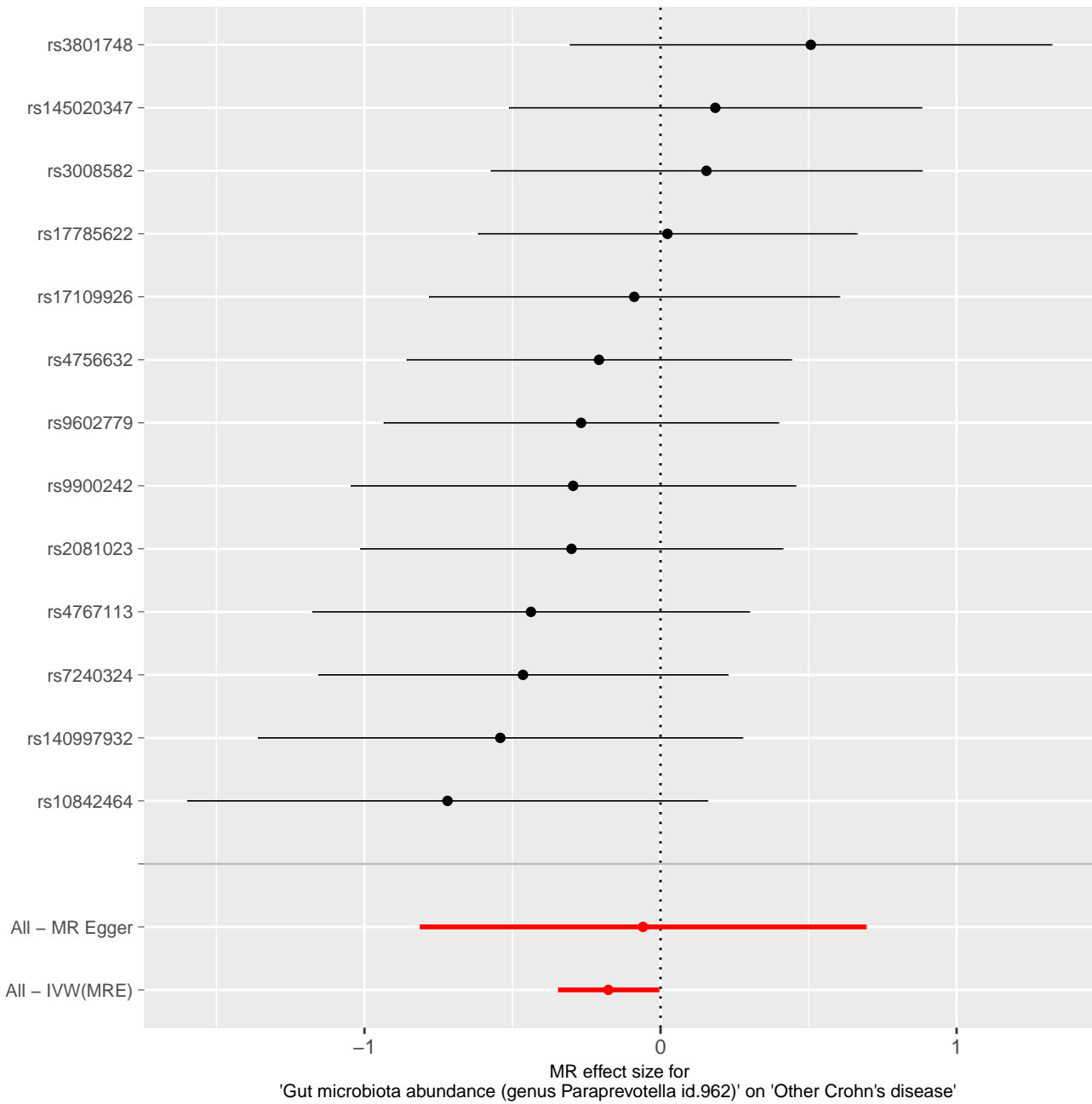




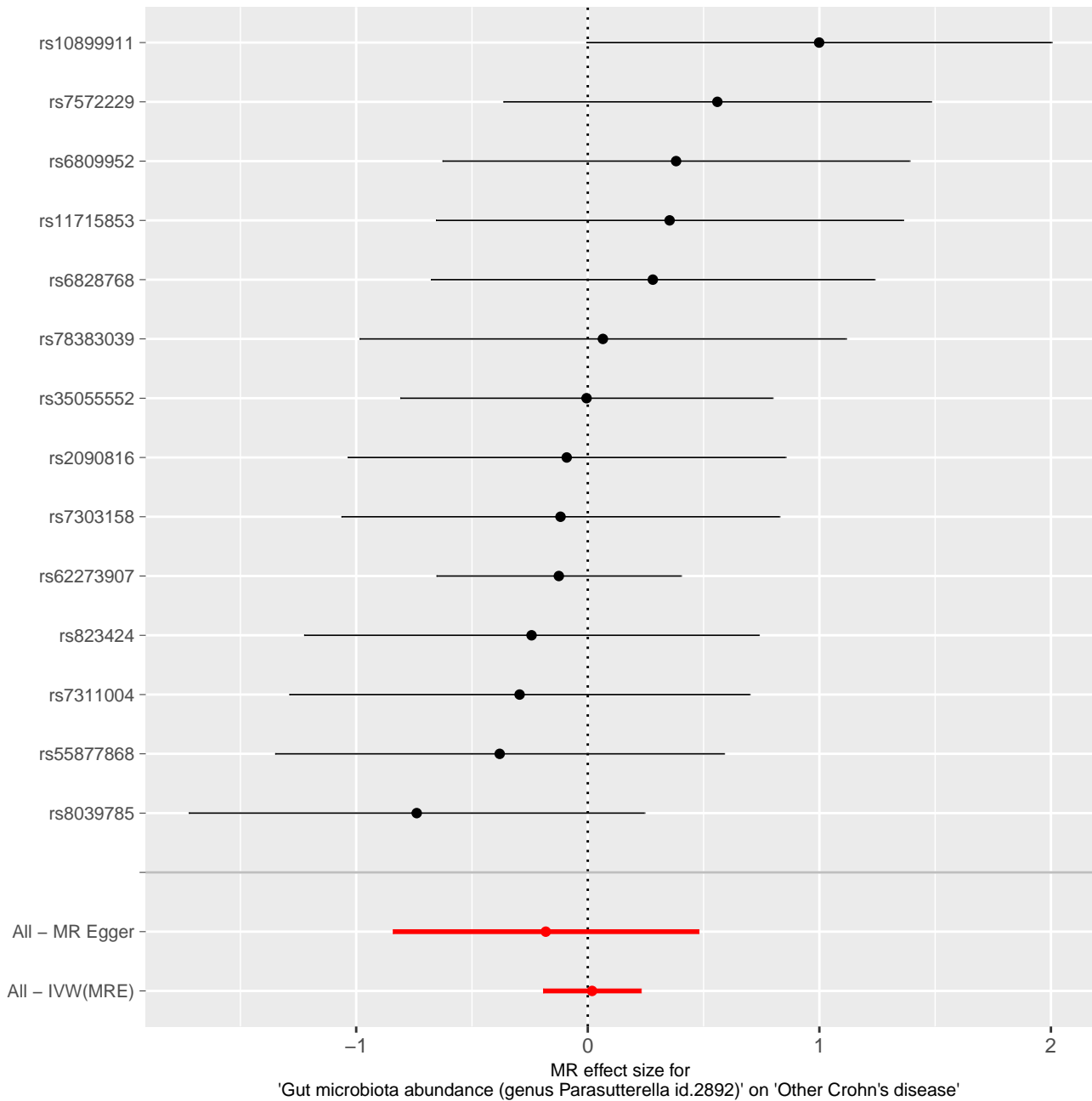


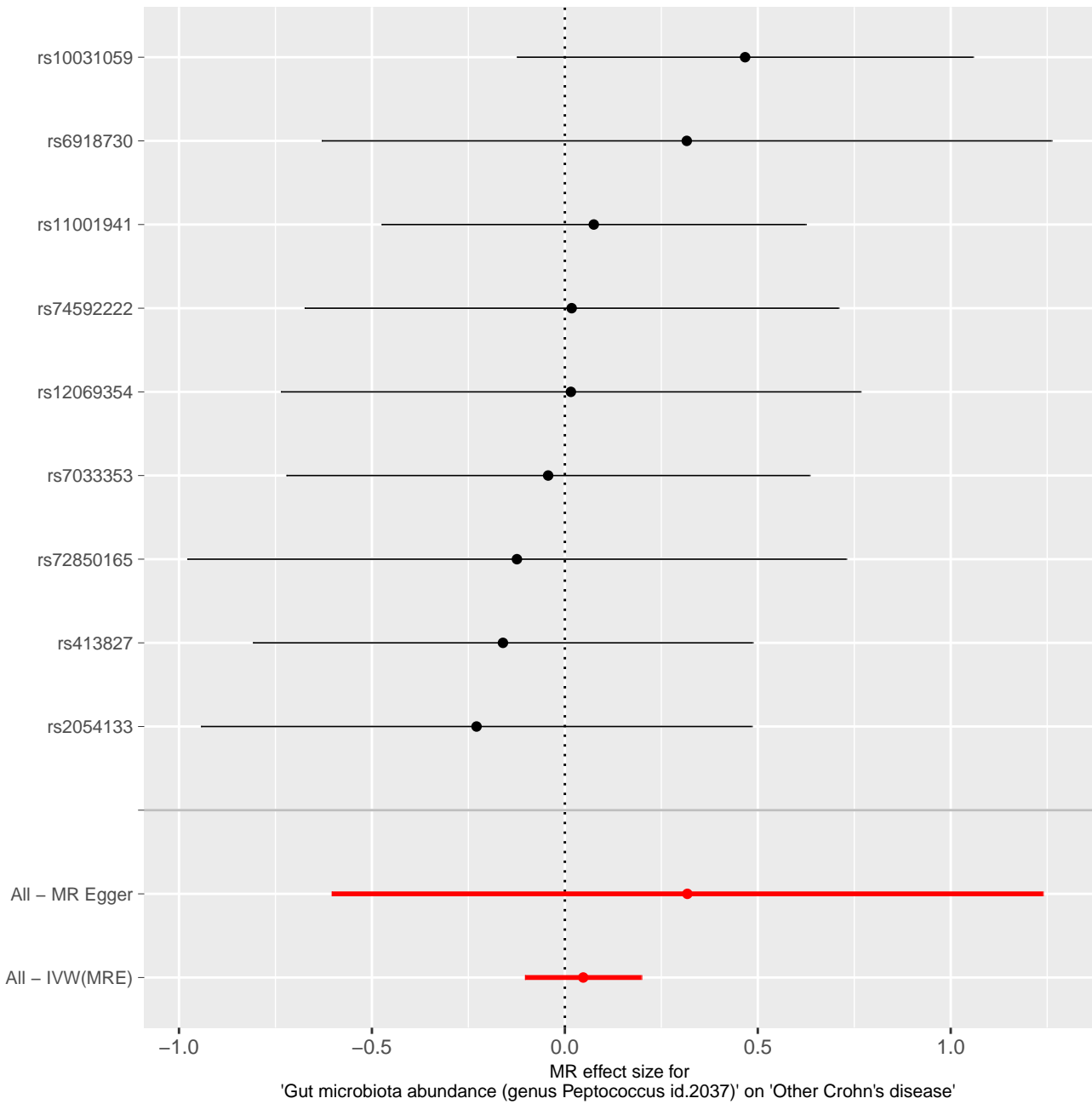


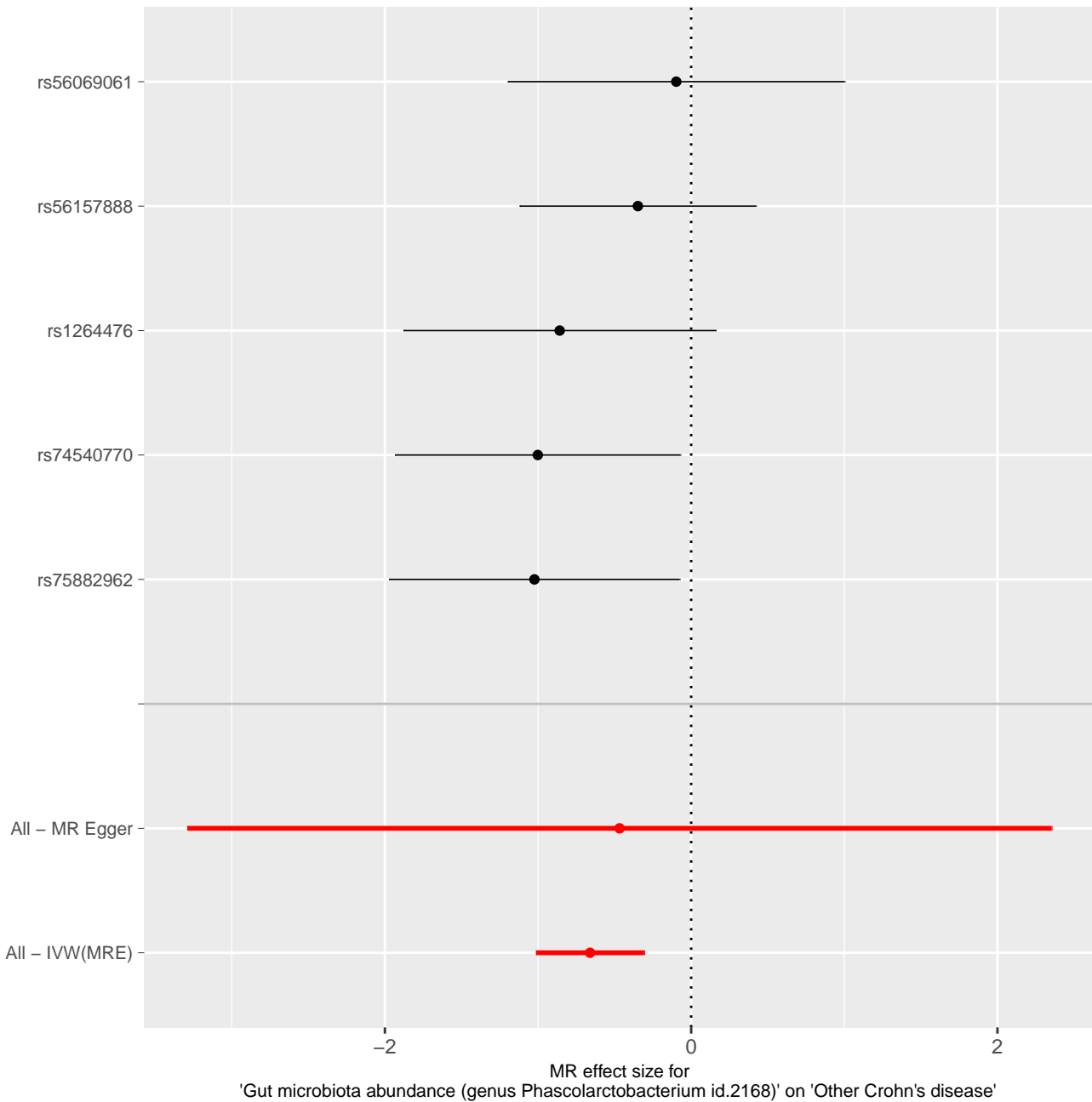
Batch 264 : Gut microbiota abundance (genus Paraprevotella id.962) on Other Crohn's disease



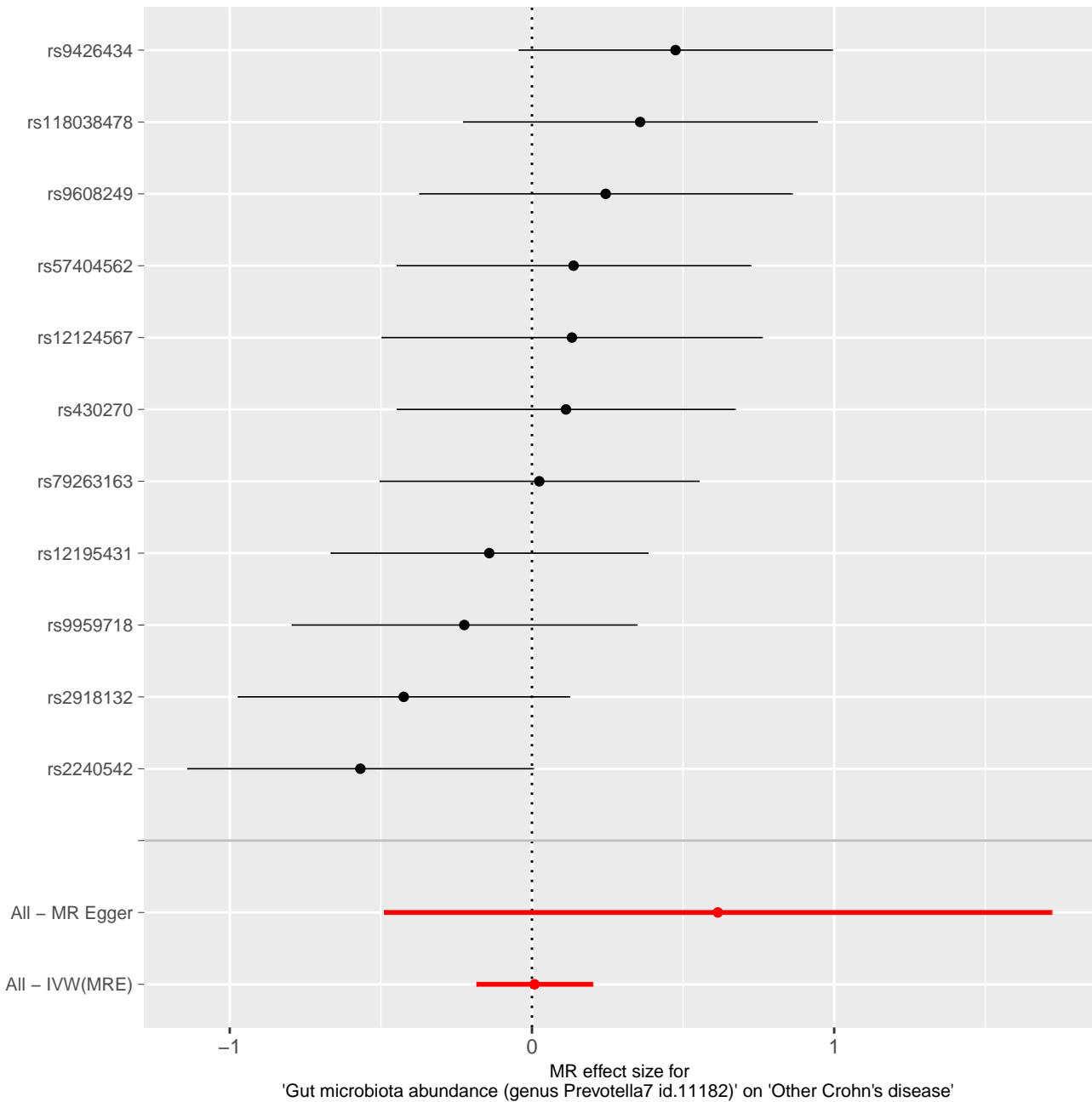
Batch 265 : Gut microbiota abundance (genus Parasutterella id.2892) on Other Crohn's disease

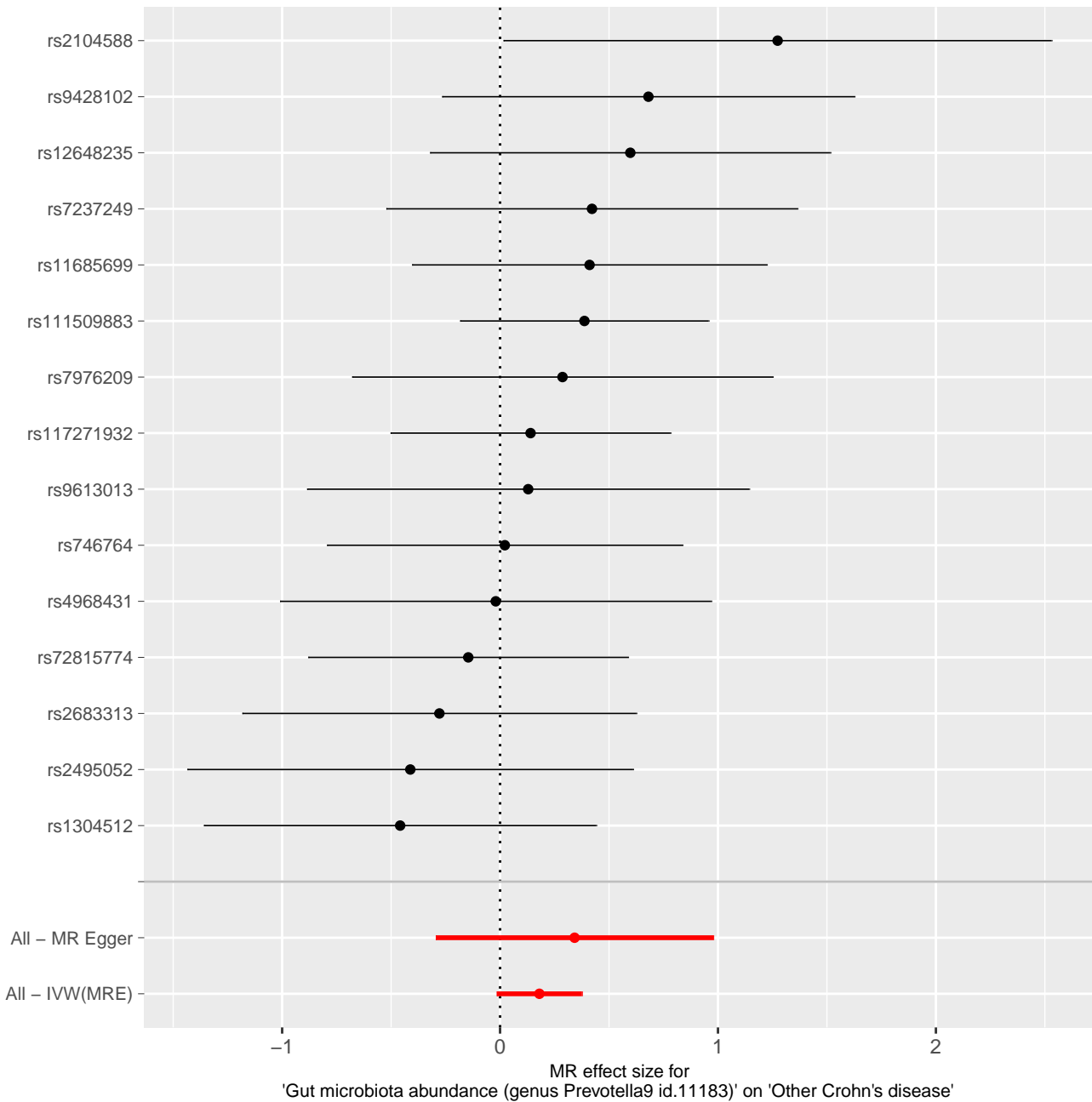


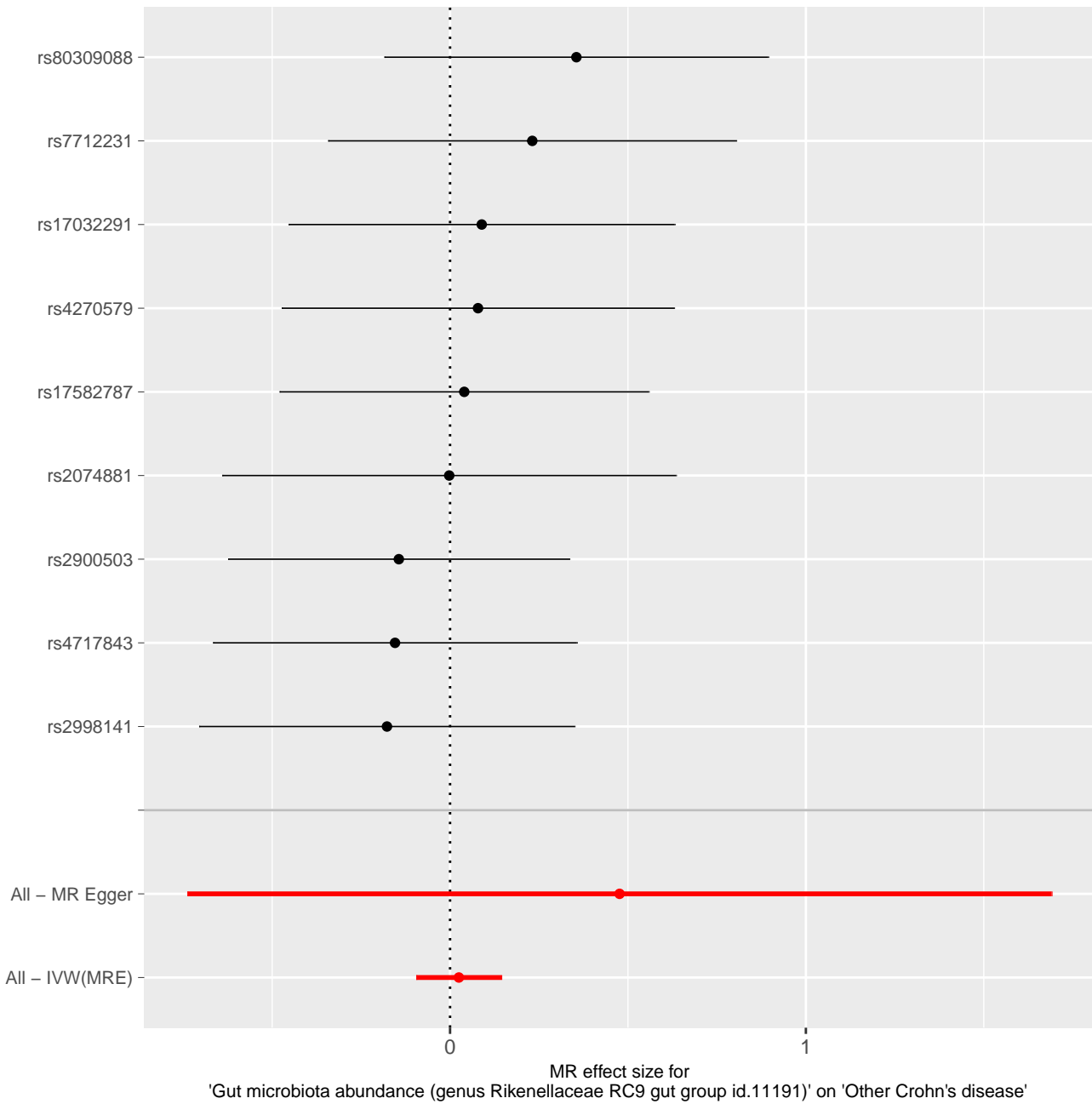




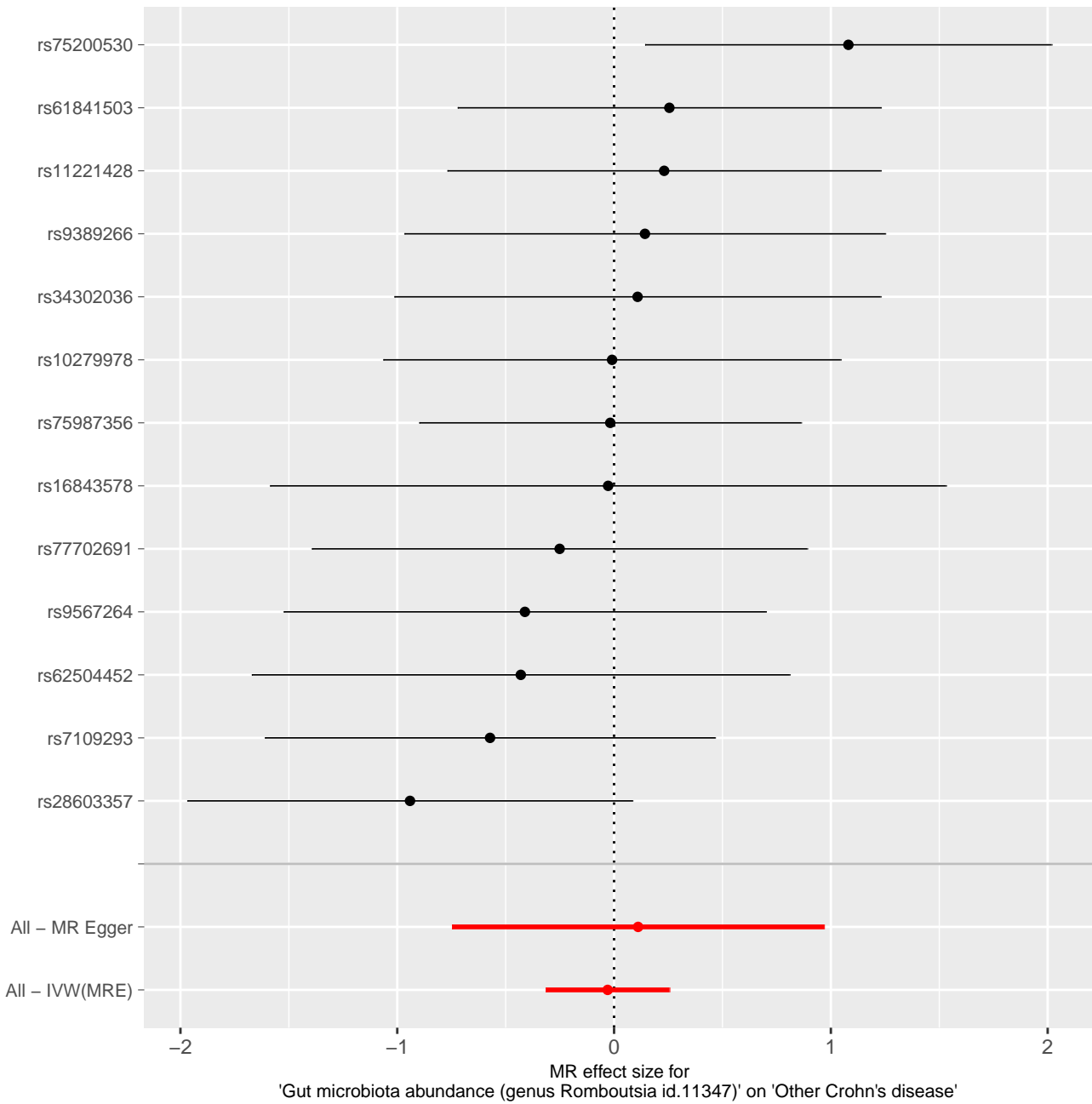
Batch 268 : Gut microbiota abundance (genus Prevotella7 id.11182) on Other Crohn's disease



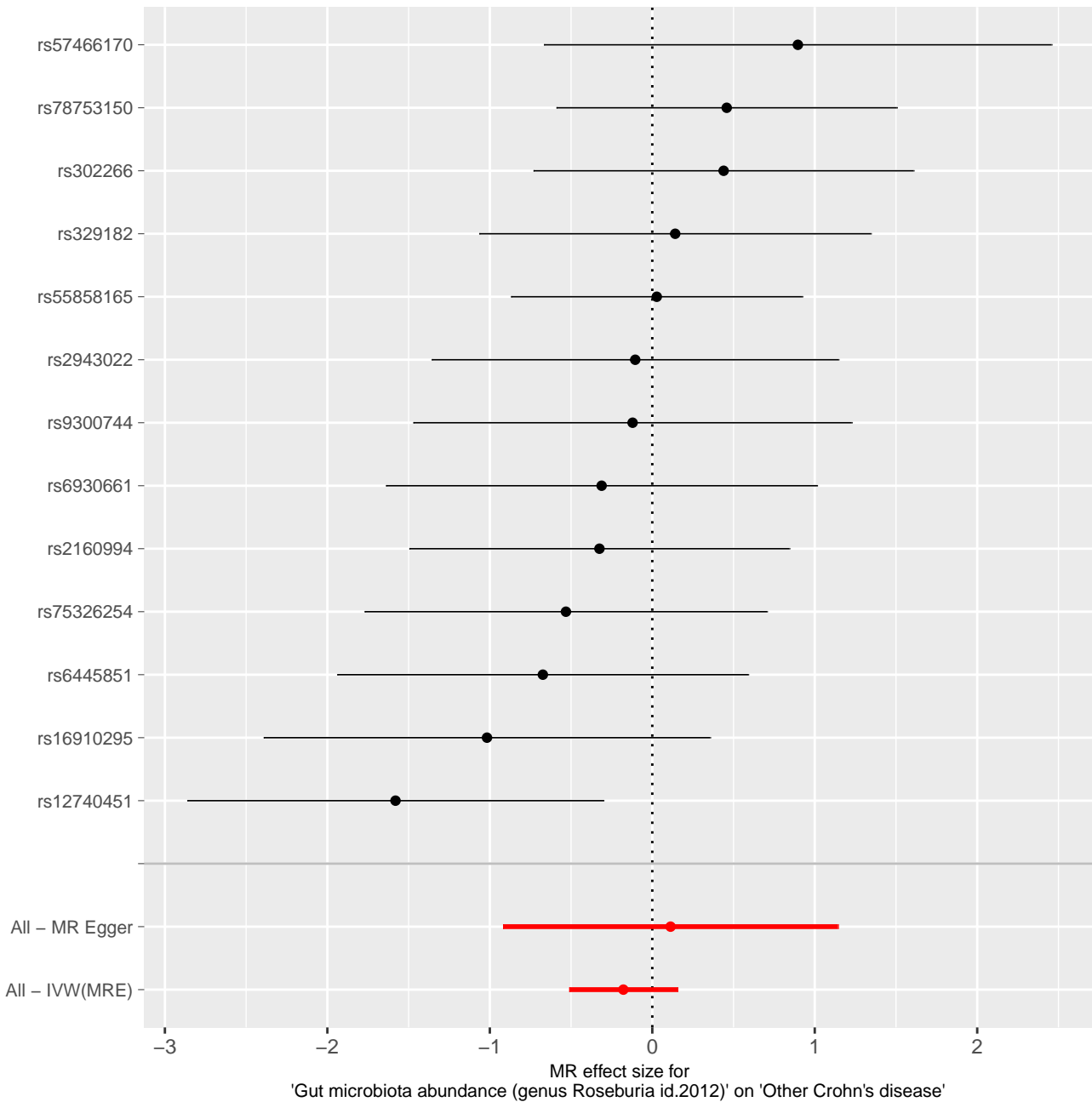




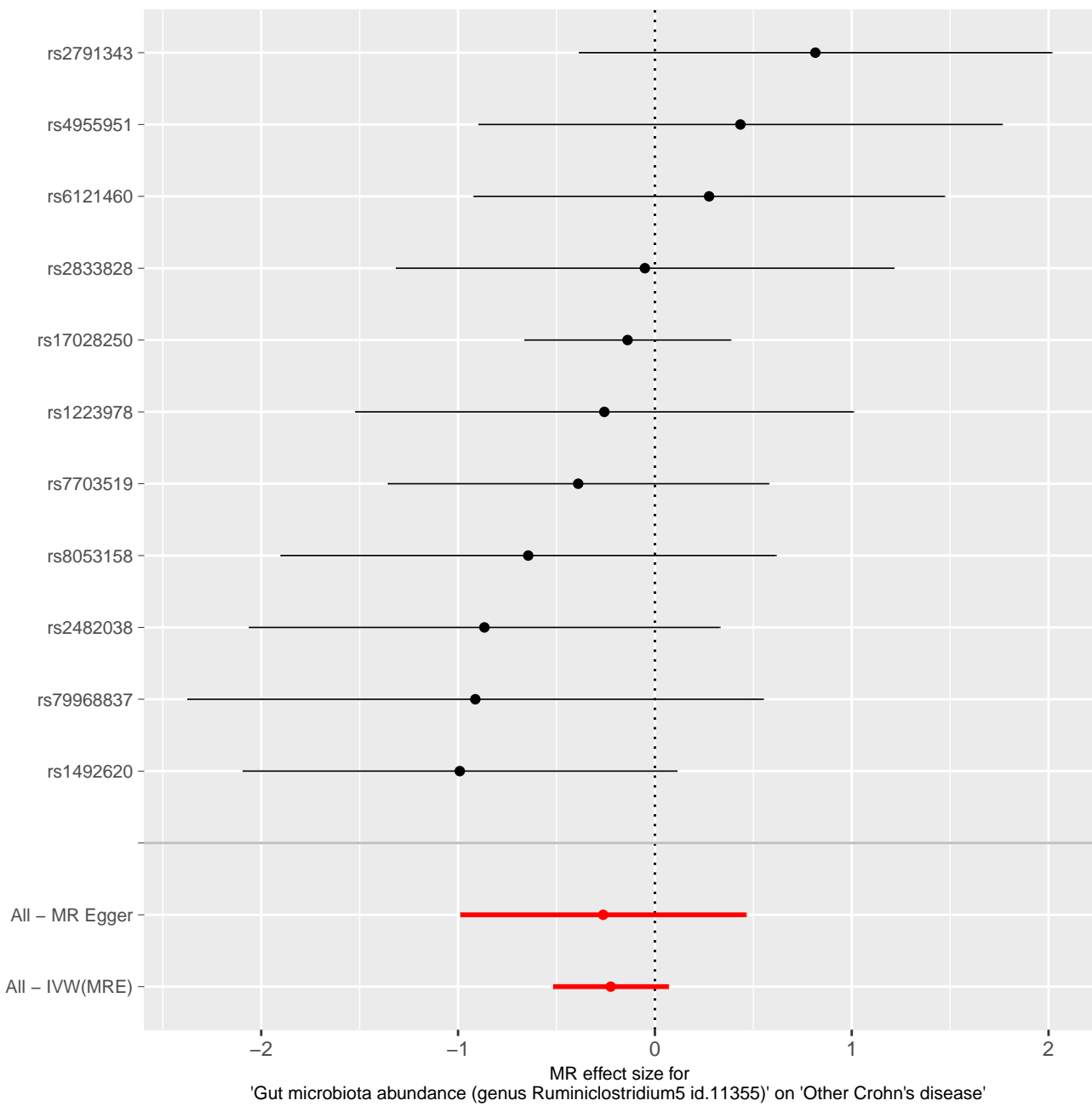
Batch 271 : Gut microbiota abundance (genus Romboutsia id.11347) on Other Crohn's disease



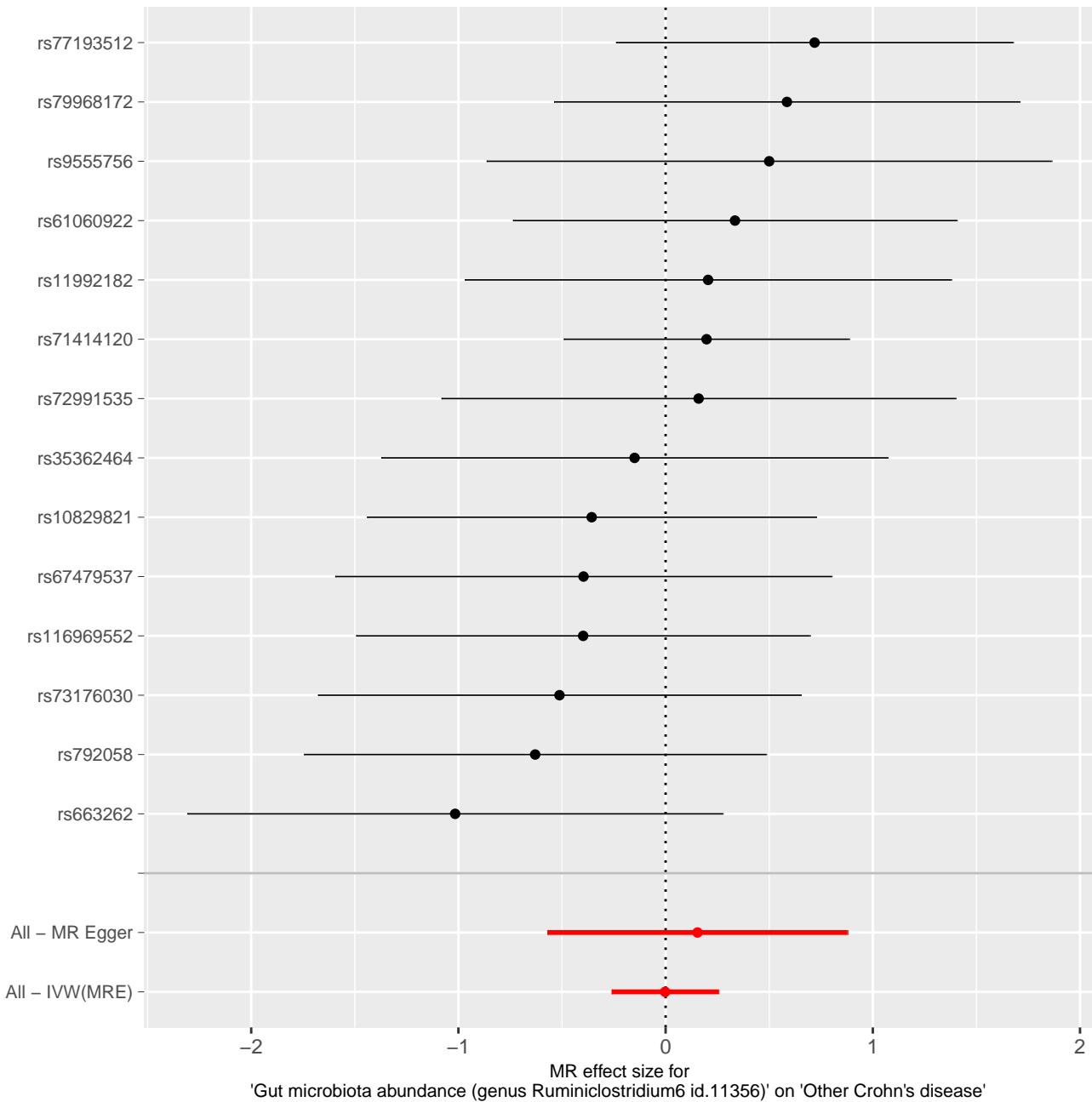
Batch 272 : Gut microbiota abundance (genus Roseburia id.2012) on Other Crohn's disease



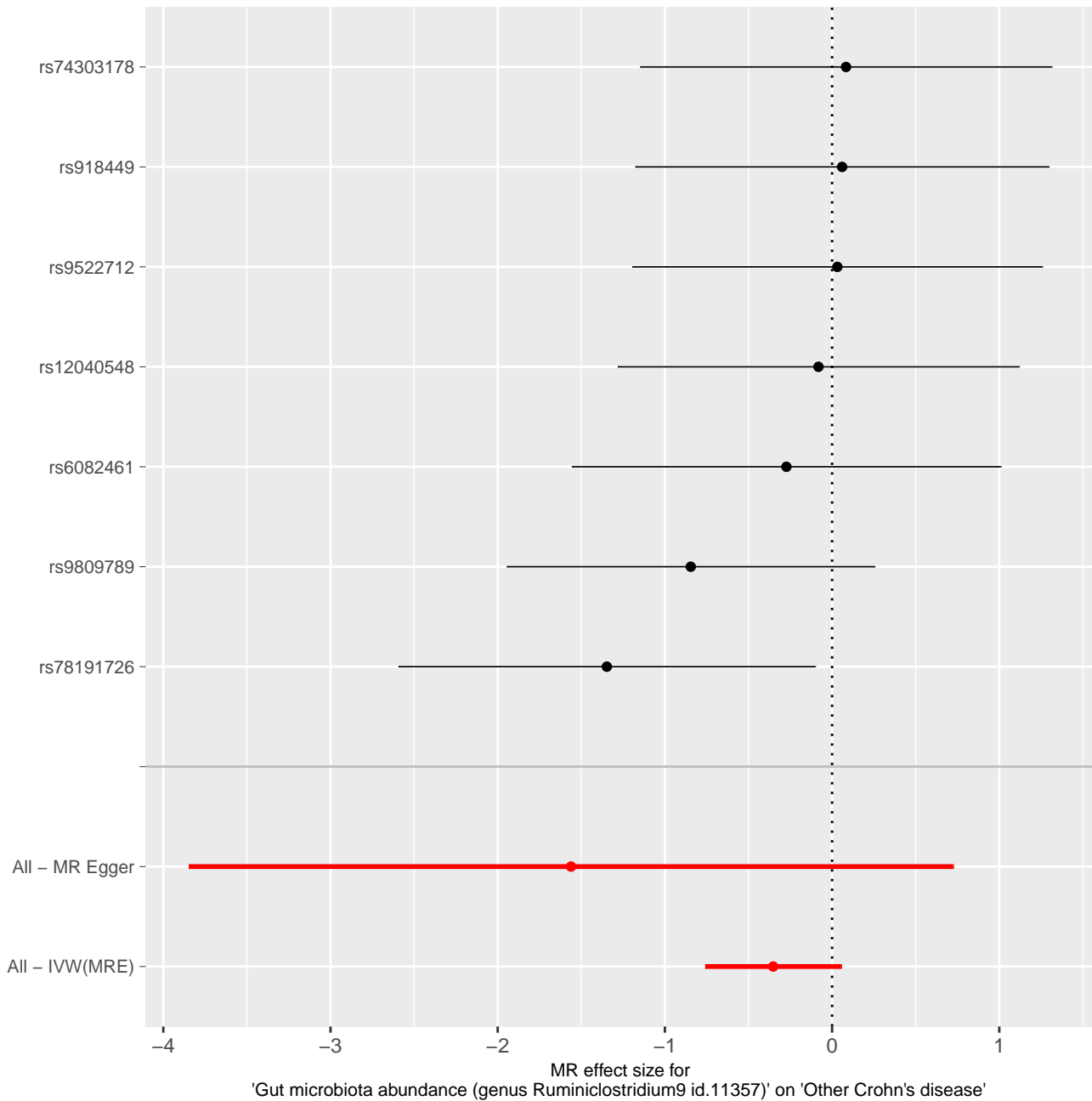
Batch 273 : Gut microbiota abundance (genus Ruminiclostridium5 id.11355) on Other Crohn's disease

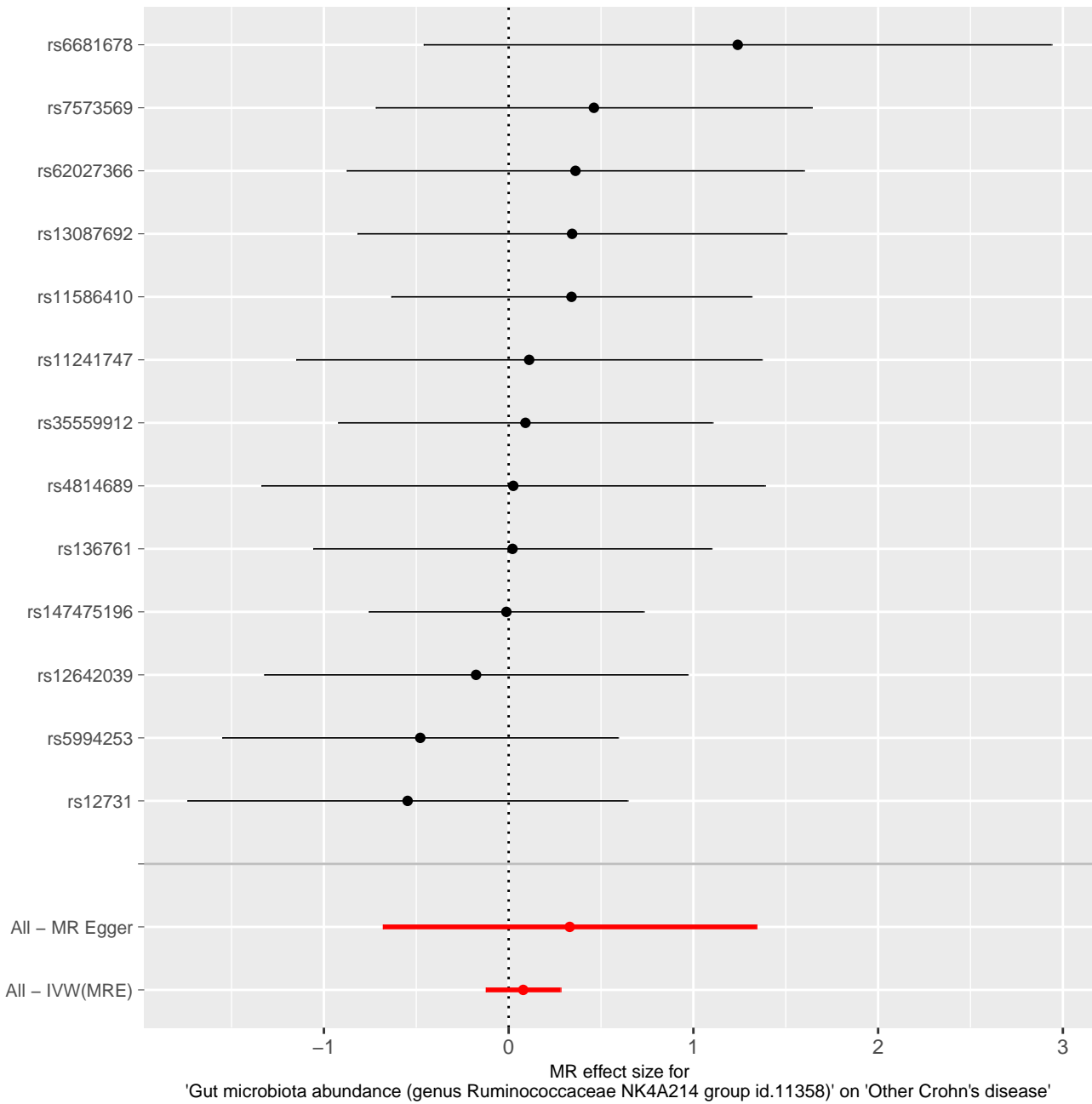


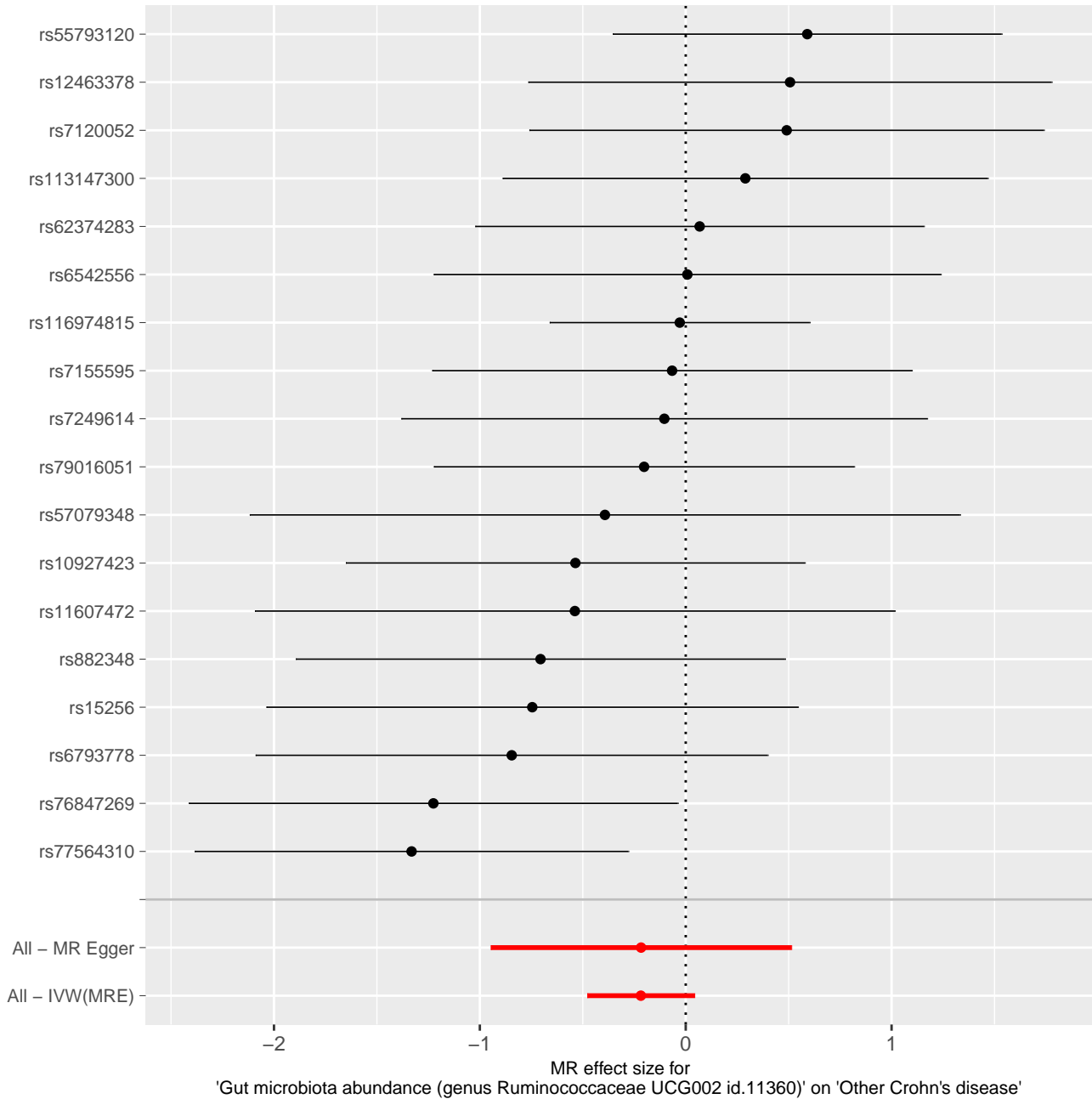
Batch 274 : Gut microbiota abundance (genus Ruminiclostridium6 id.11356) on Other Crohn's disease

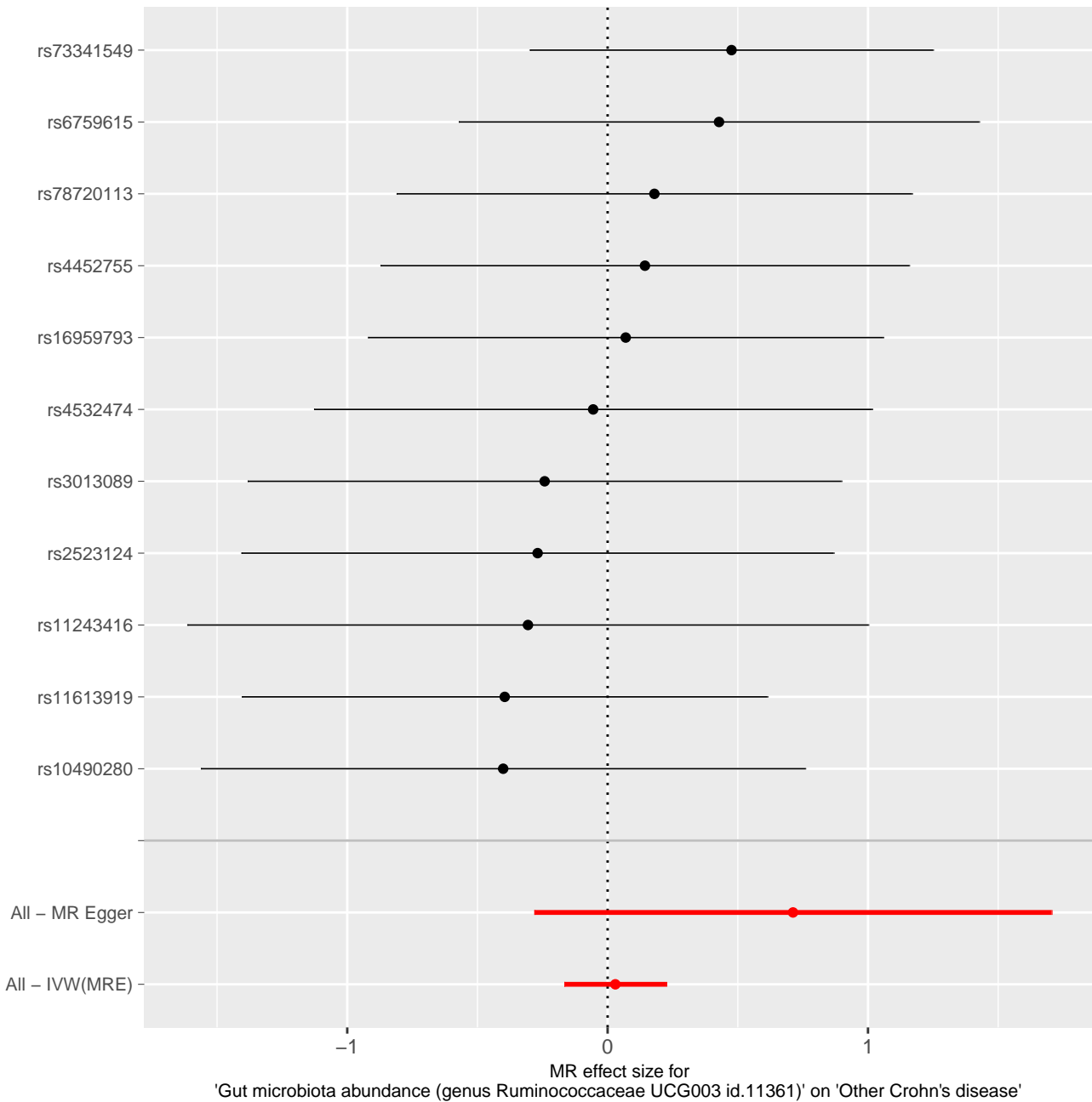


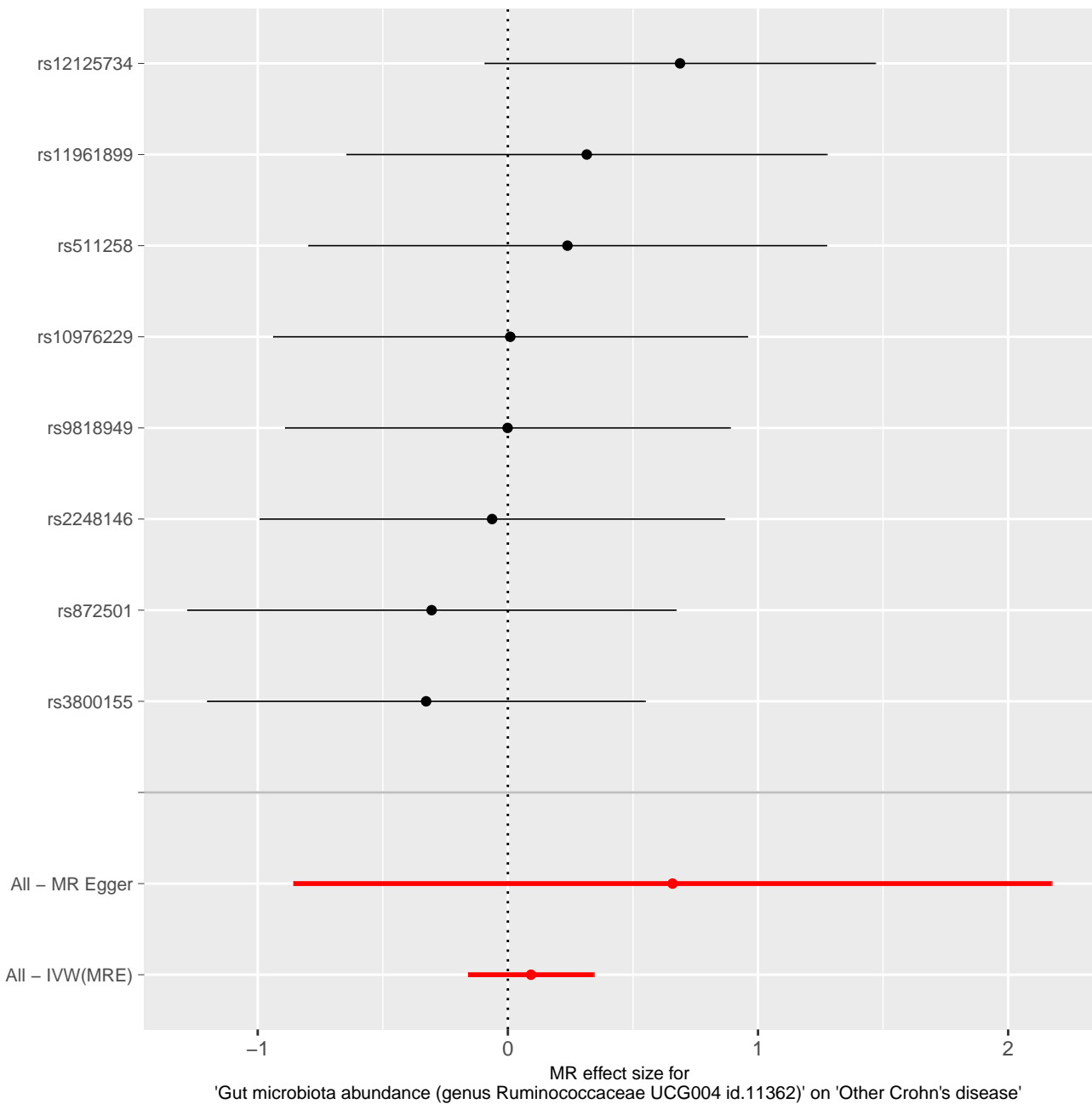
Batch 275 : Gut microbiota abundance (genus Ruminiclostridium9 id.11357) on Other Crohn's disease

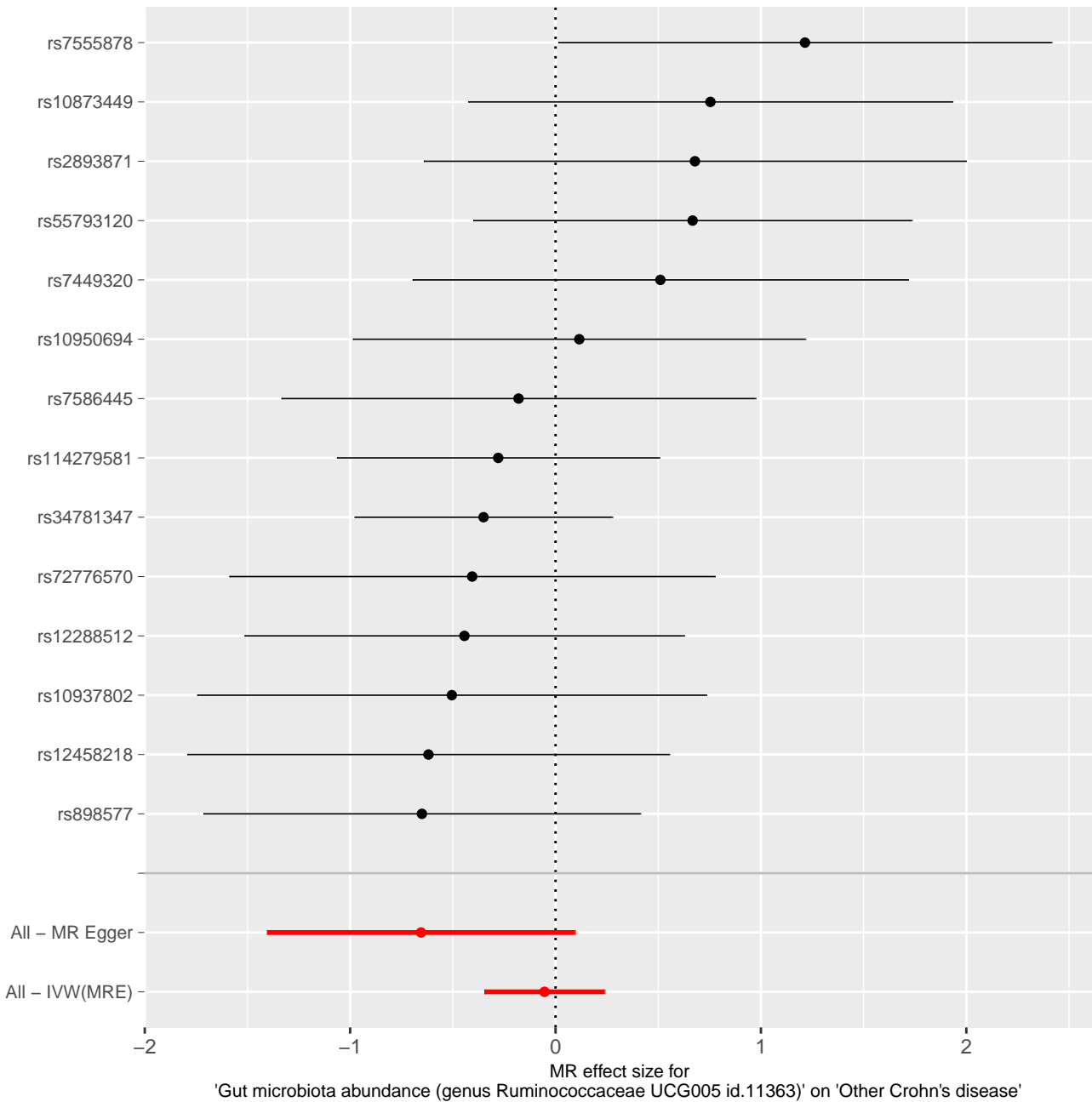


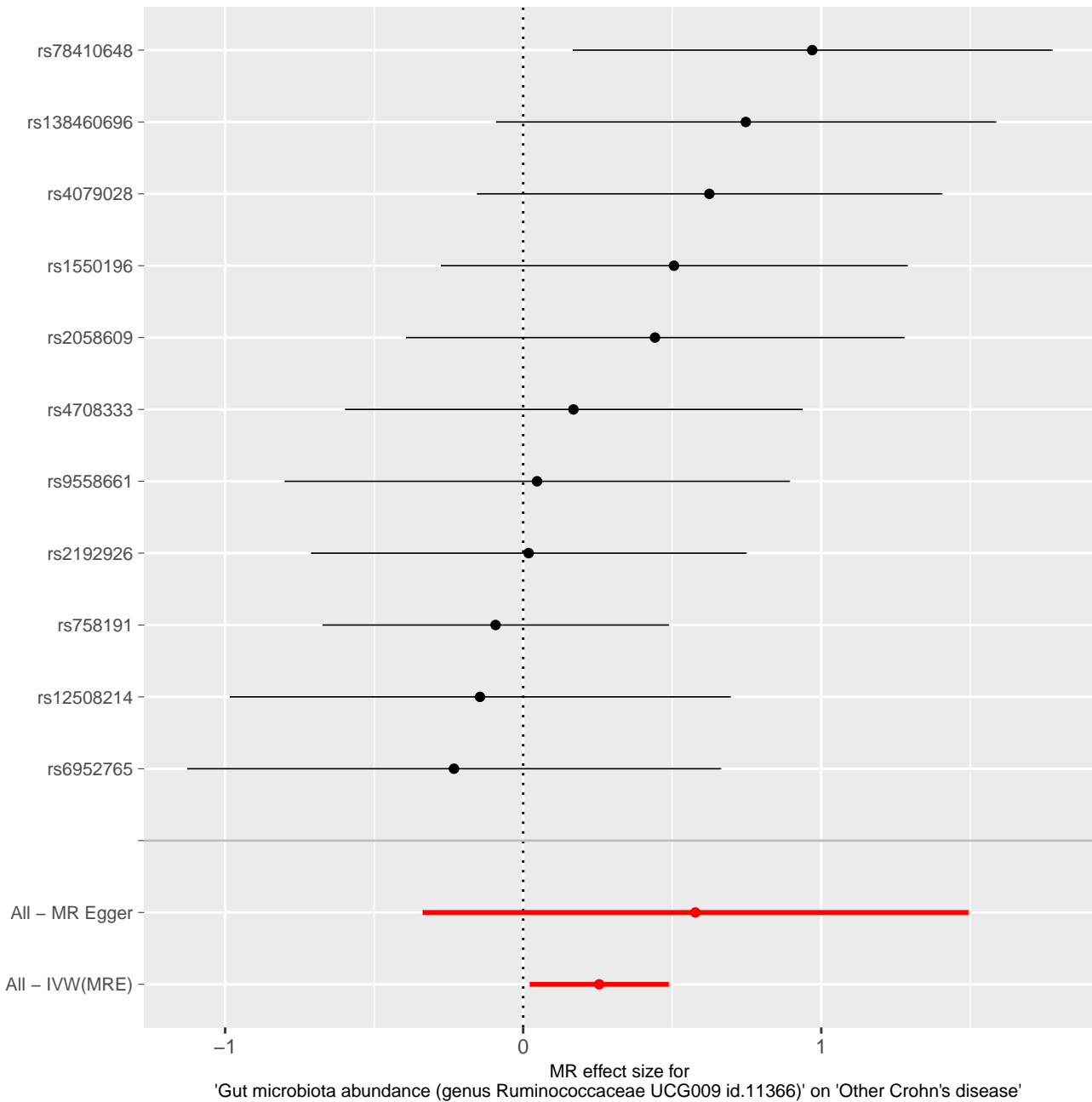


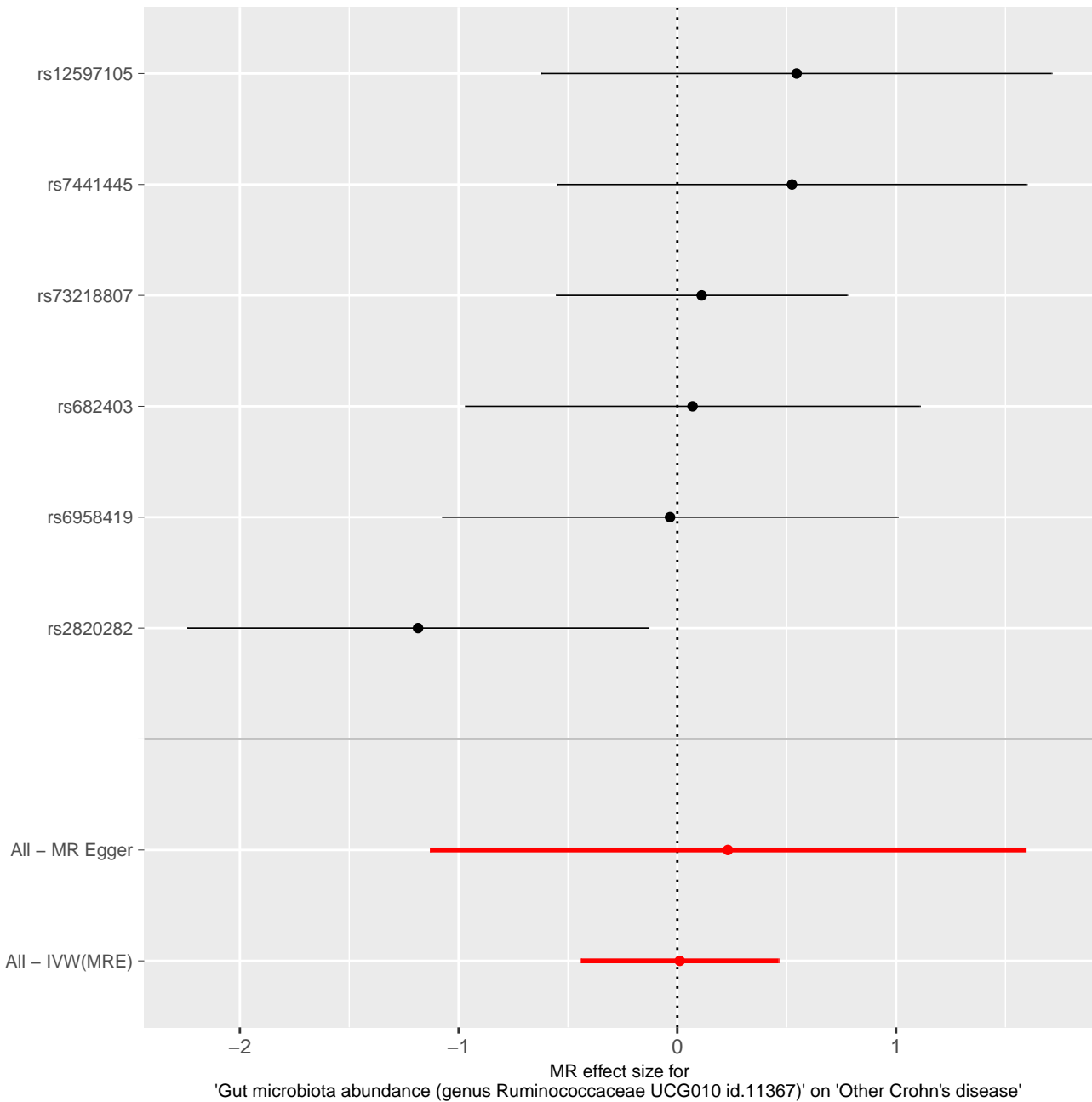


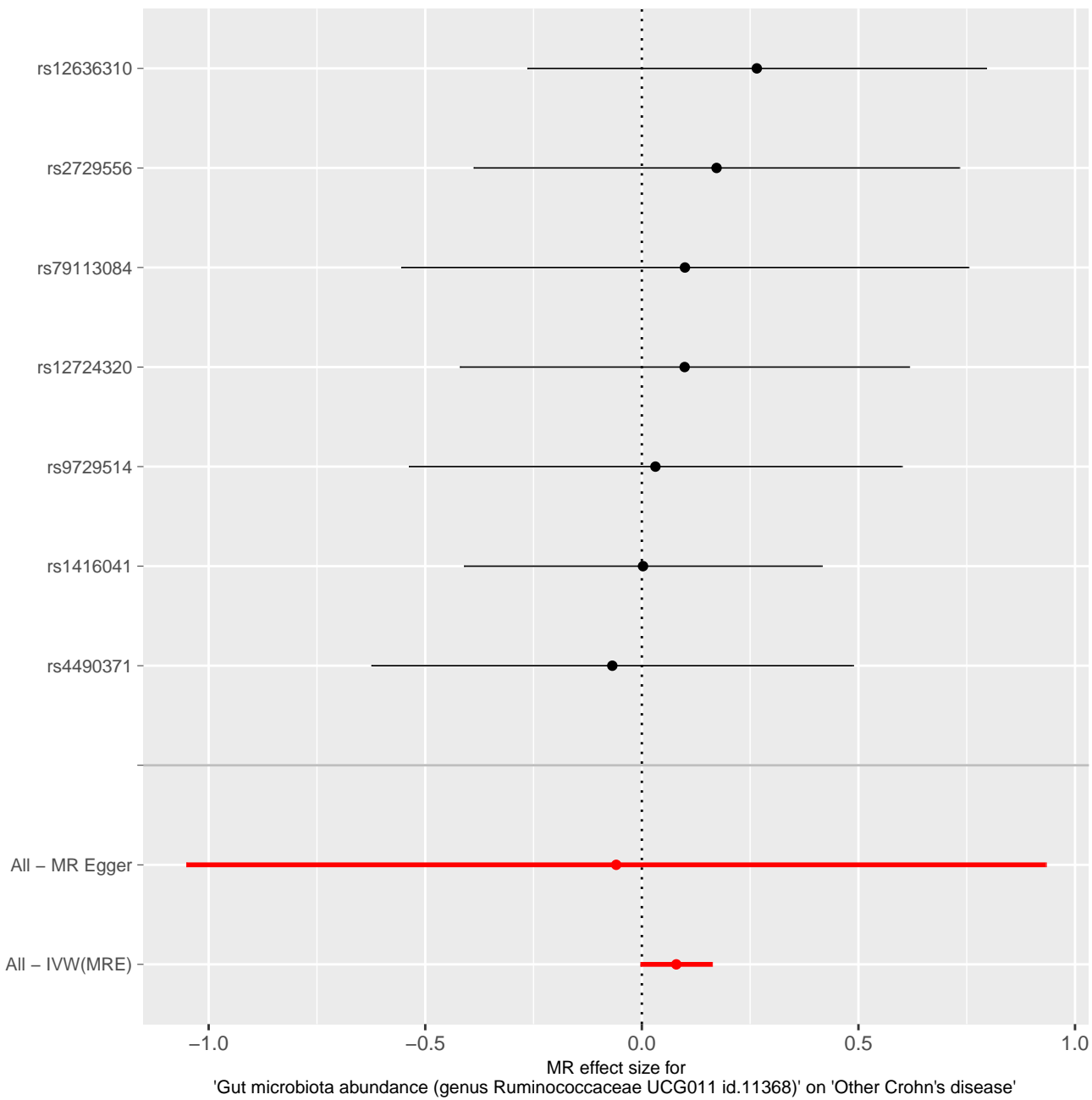




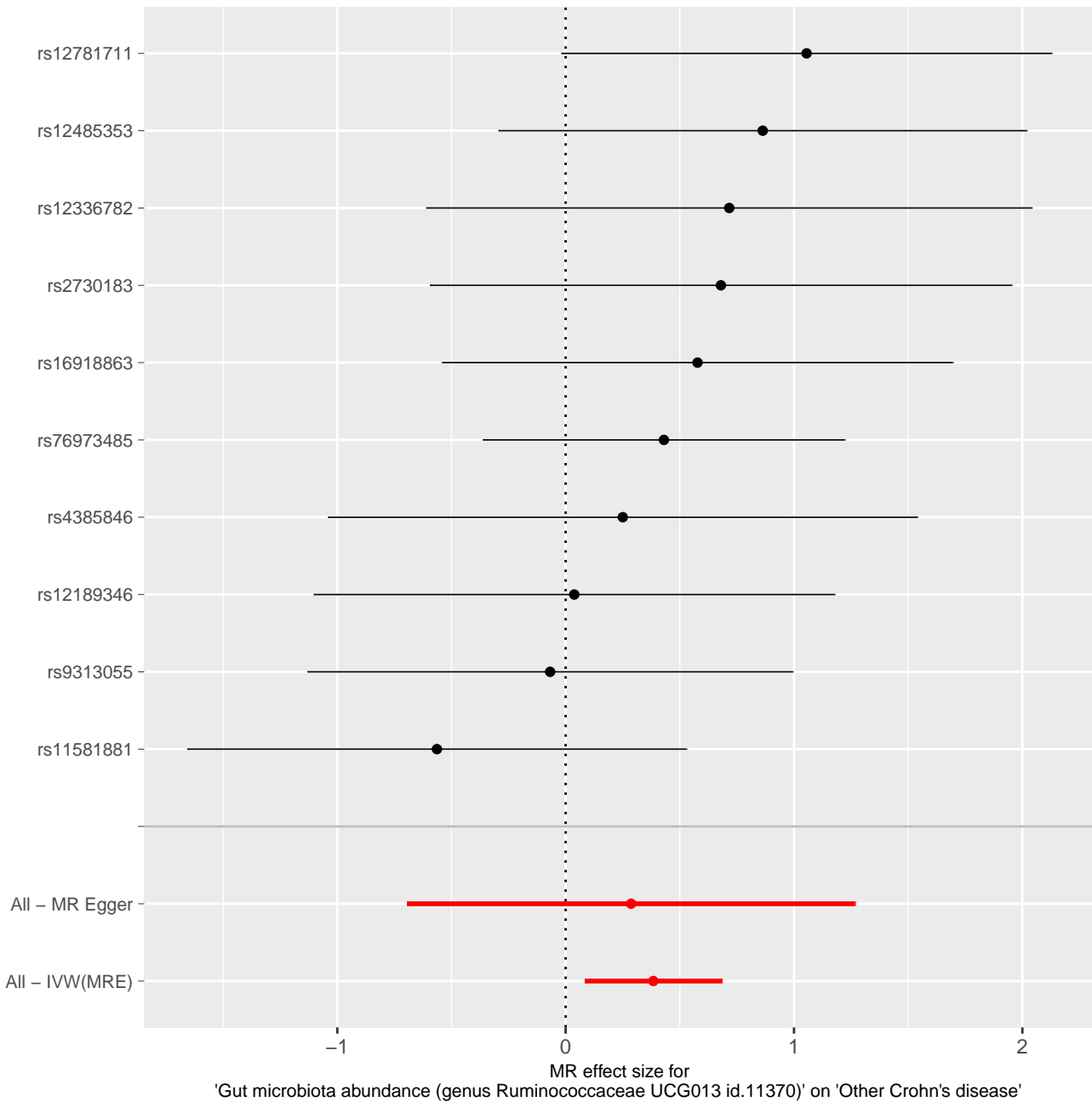


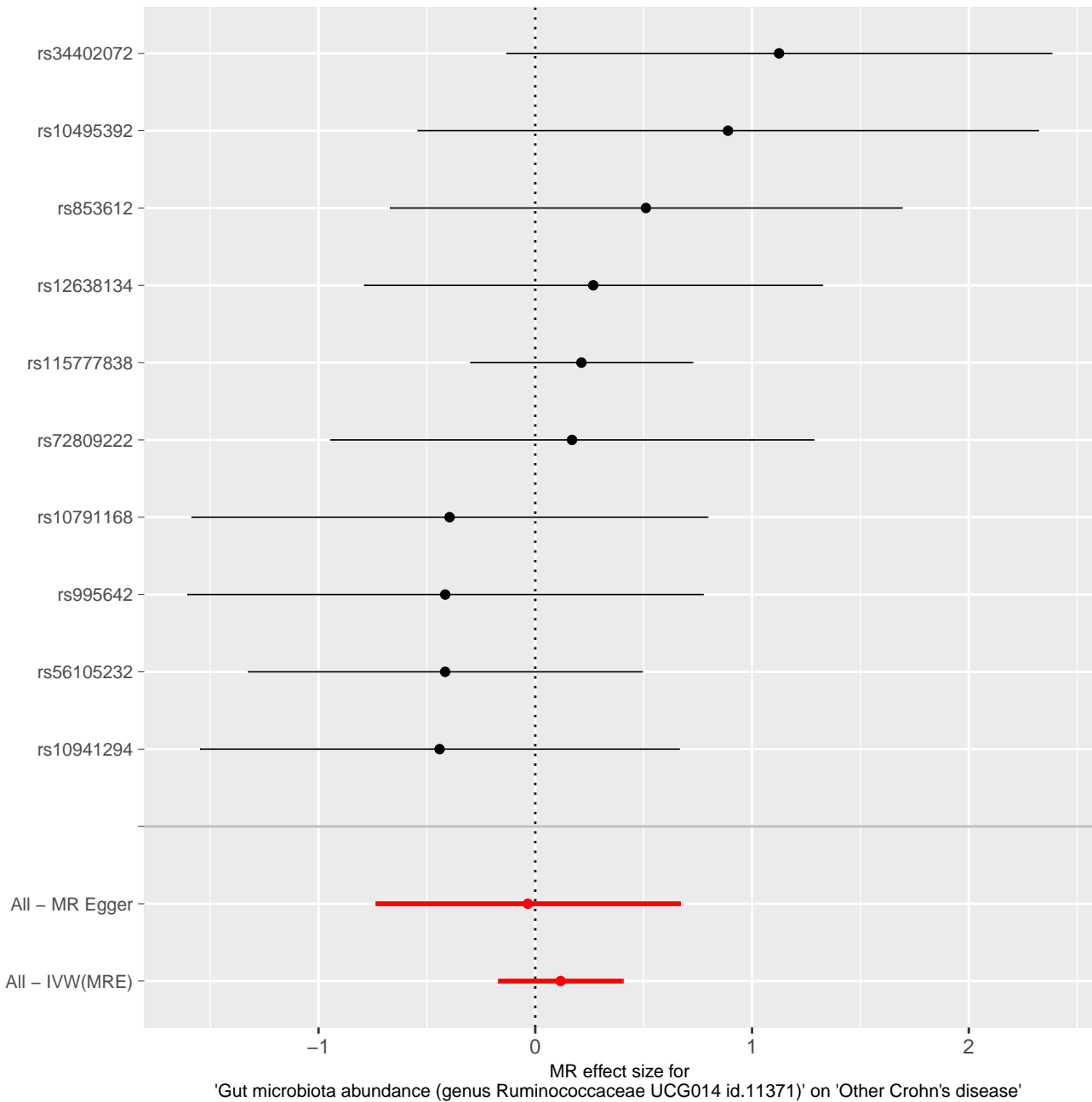


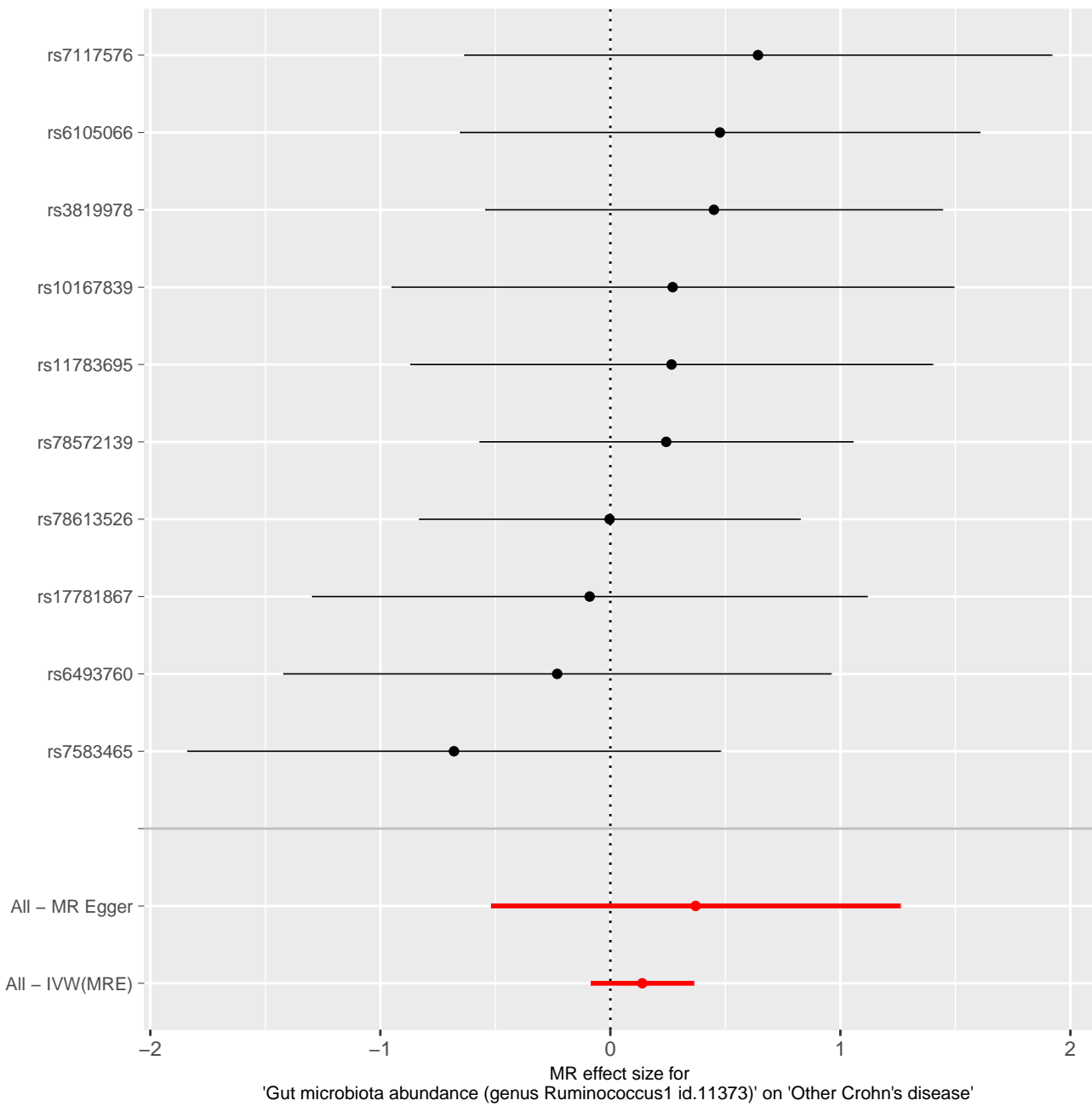




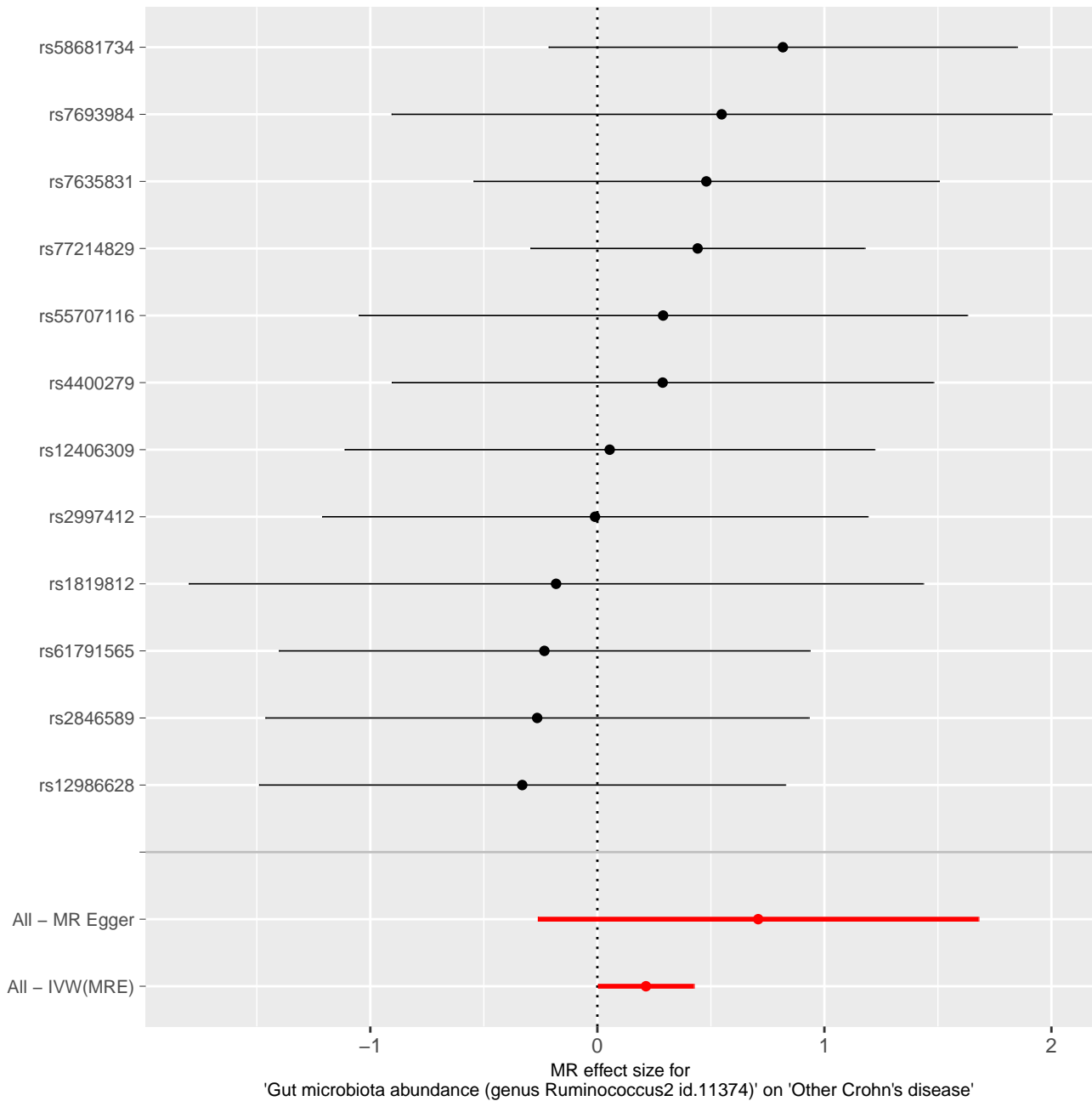
Batch 284 : Gut microbiota abundance (genus Ruminococcaceae UCG013 id.11370) on Other Crohn's disease

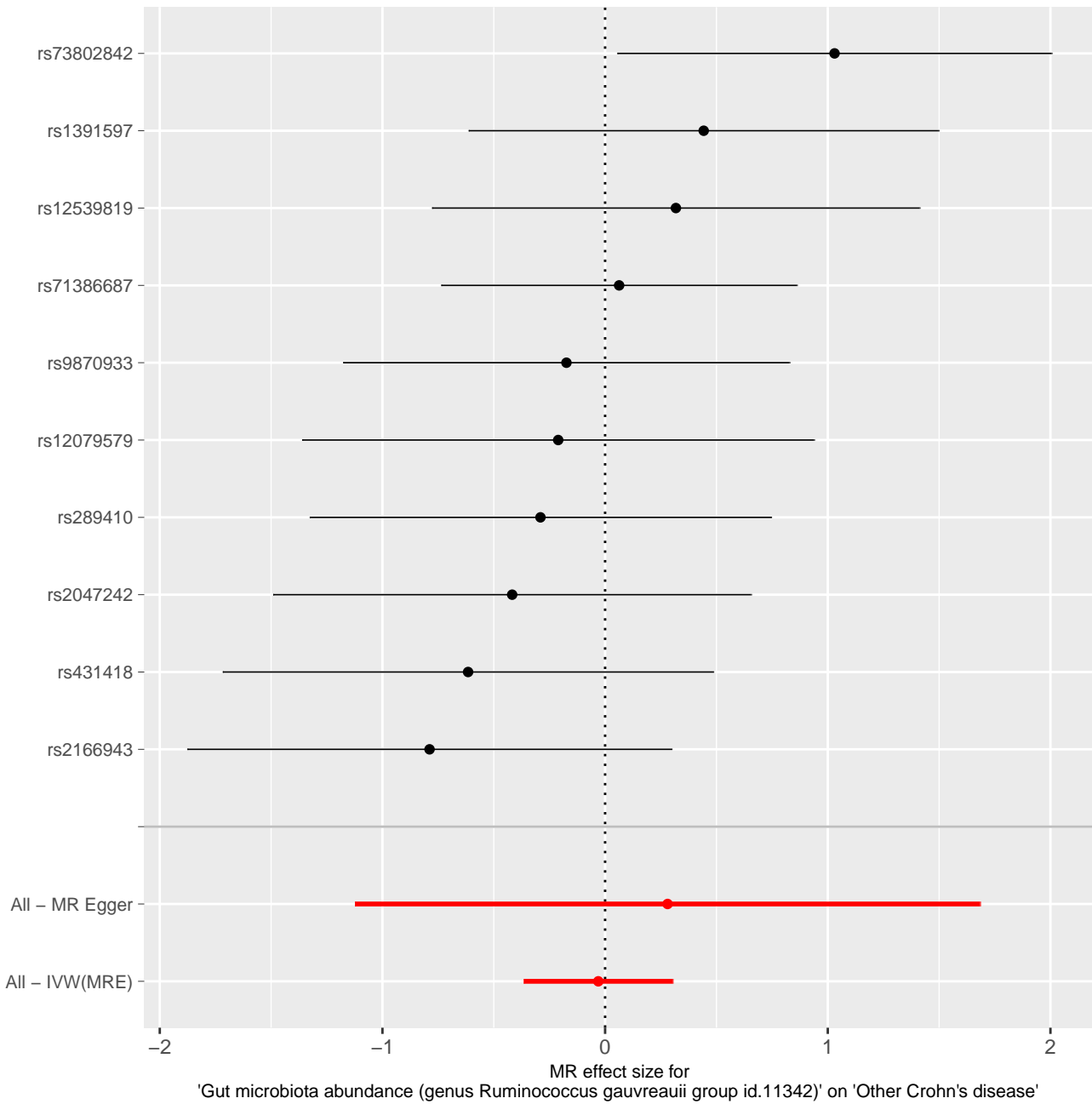


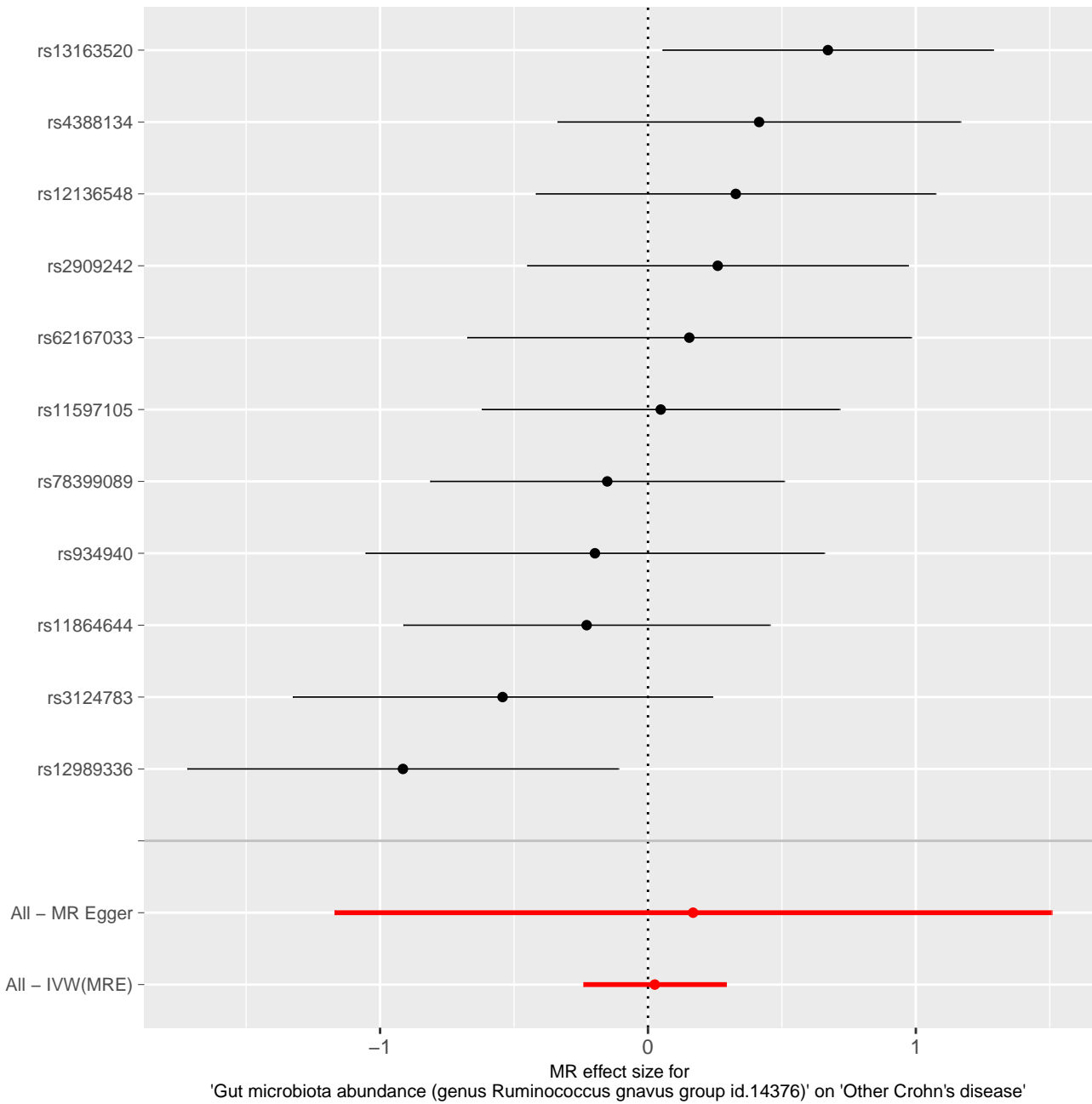


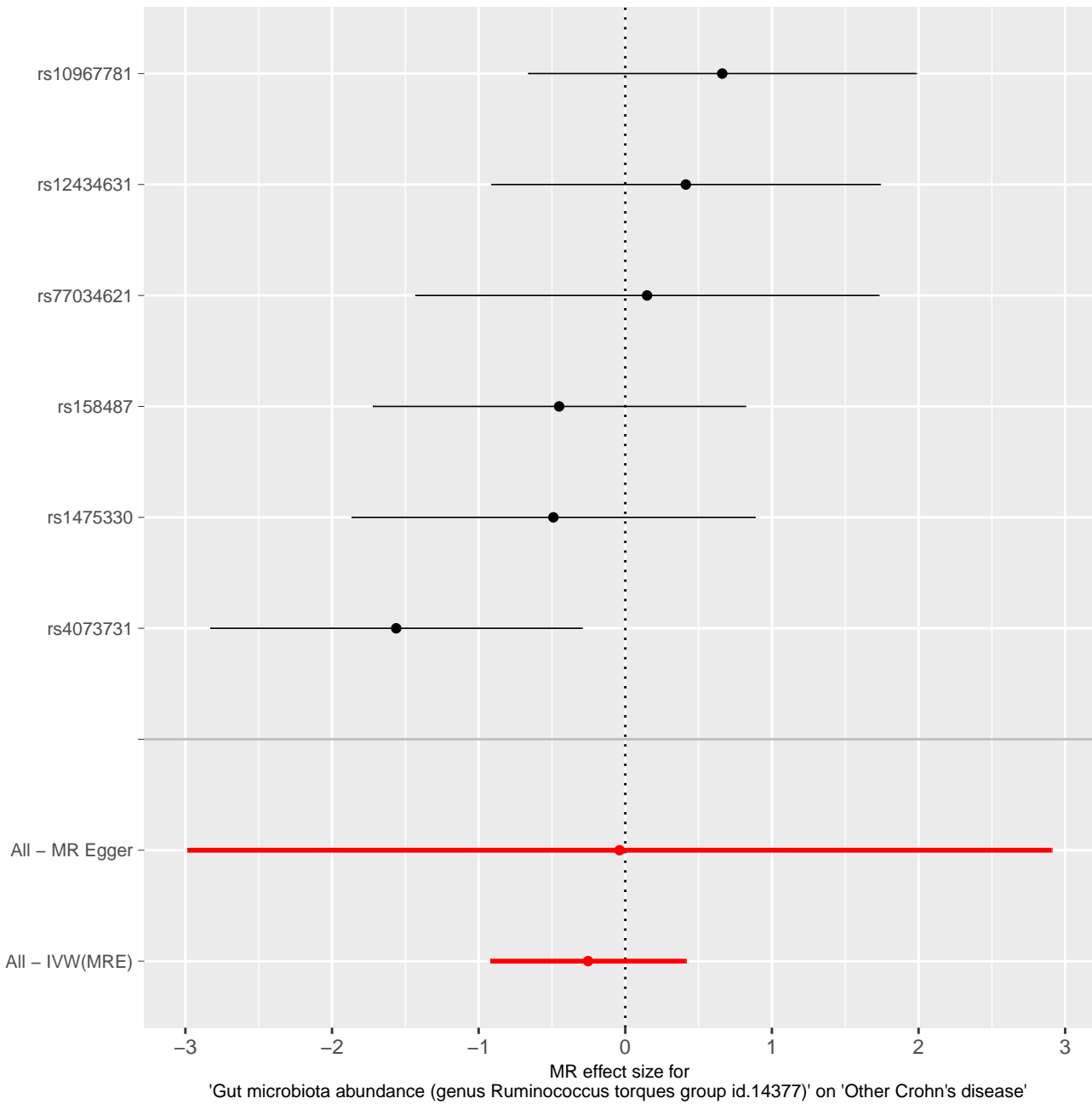


Batch 287 : Gut microbiota abundance (genus Ruminococcus2 id.11374) on Other Crohn's disease

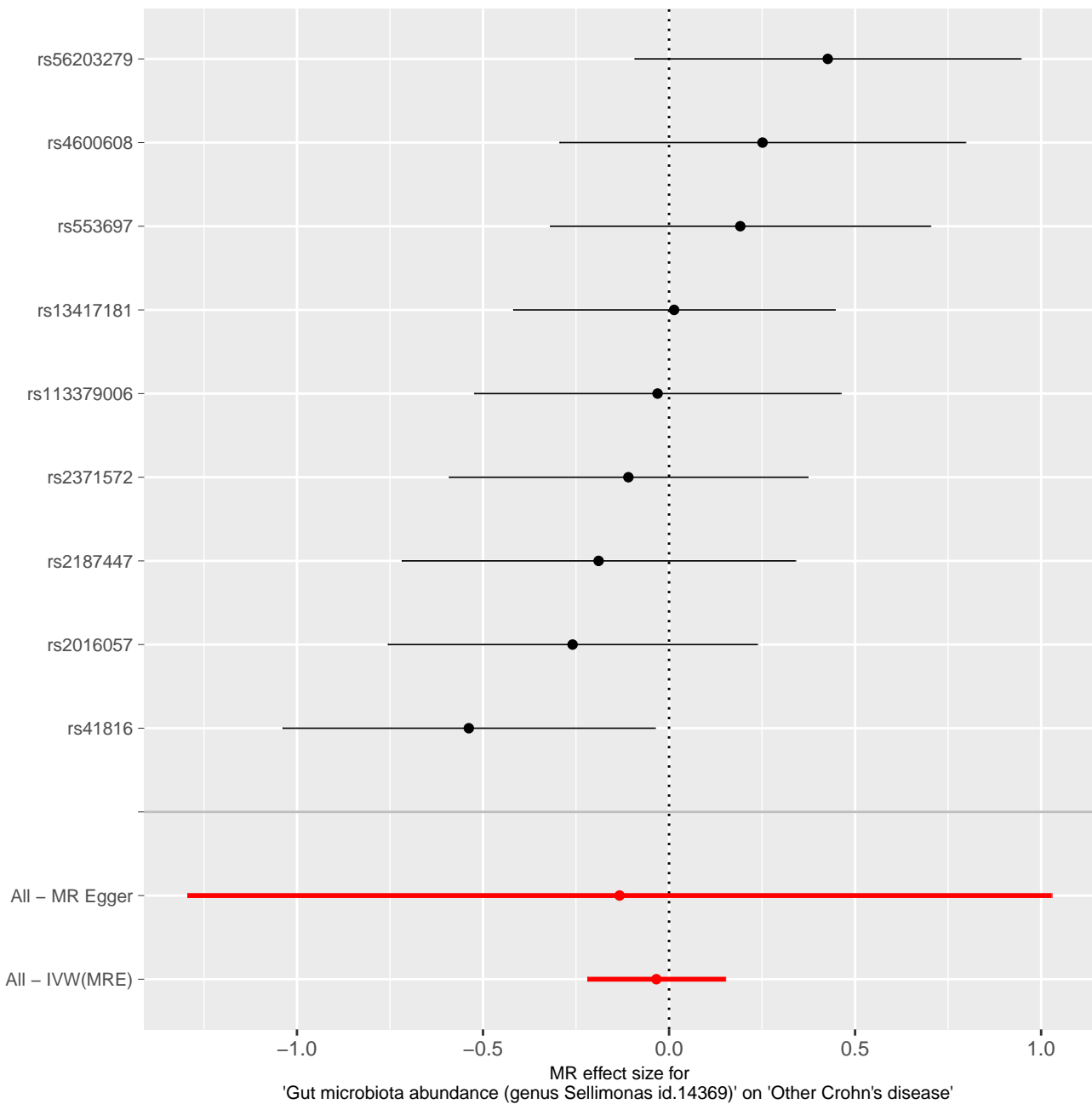


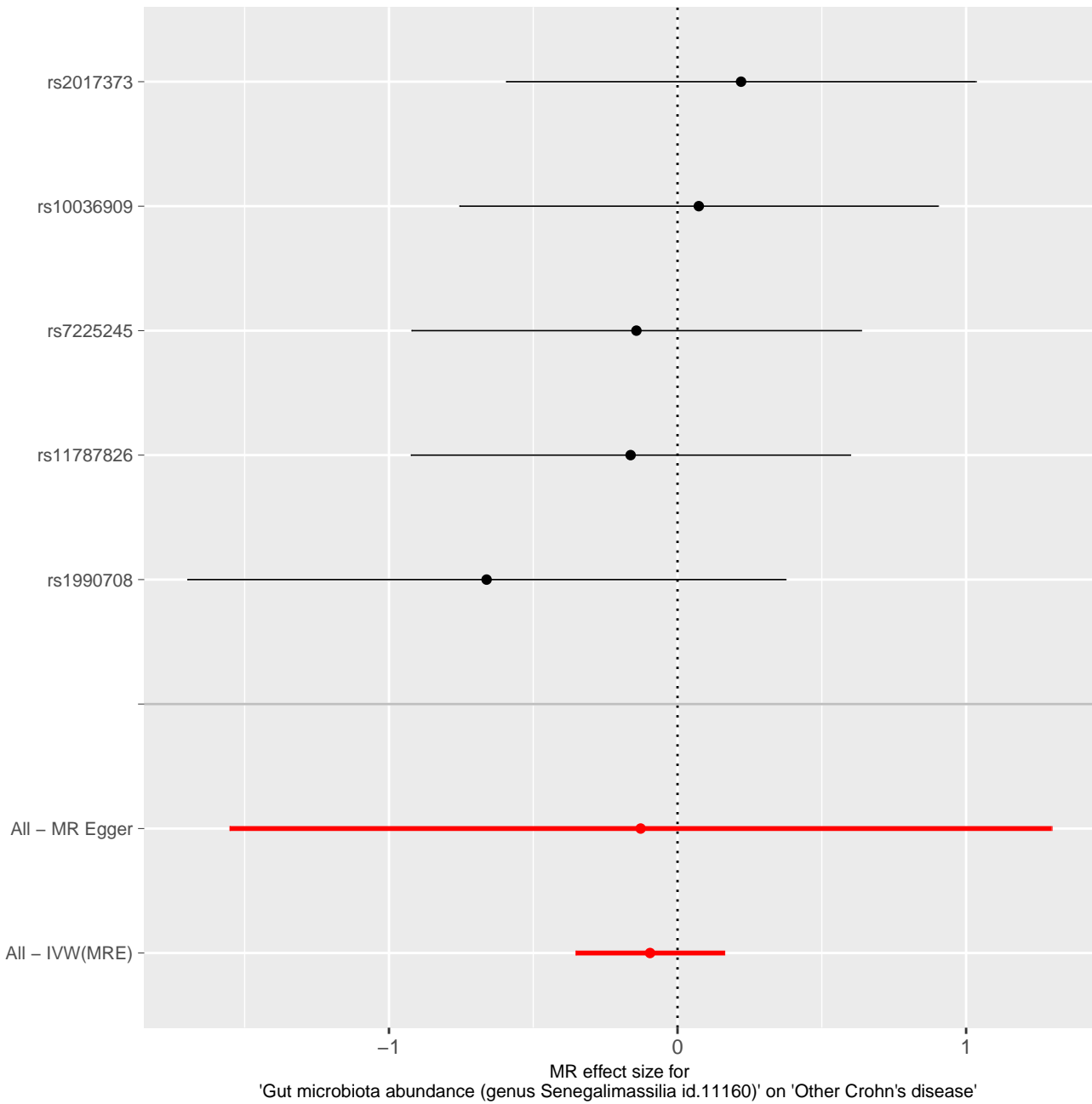




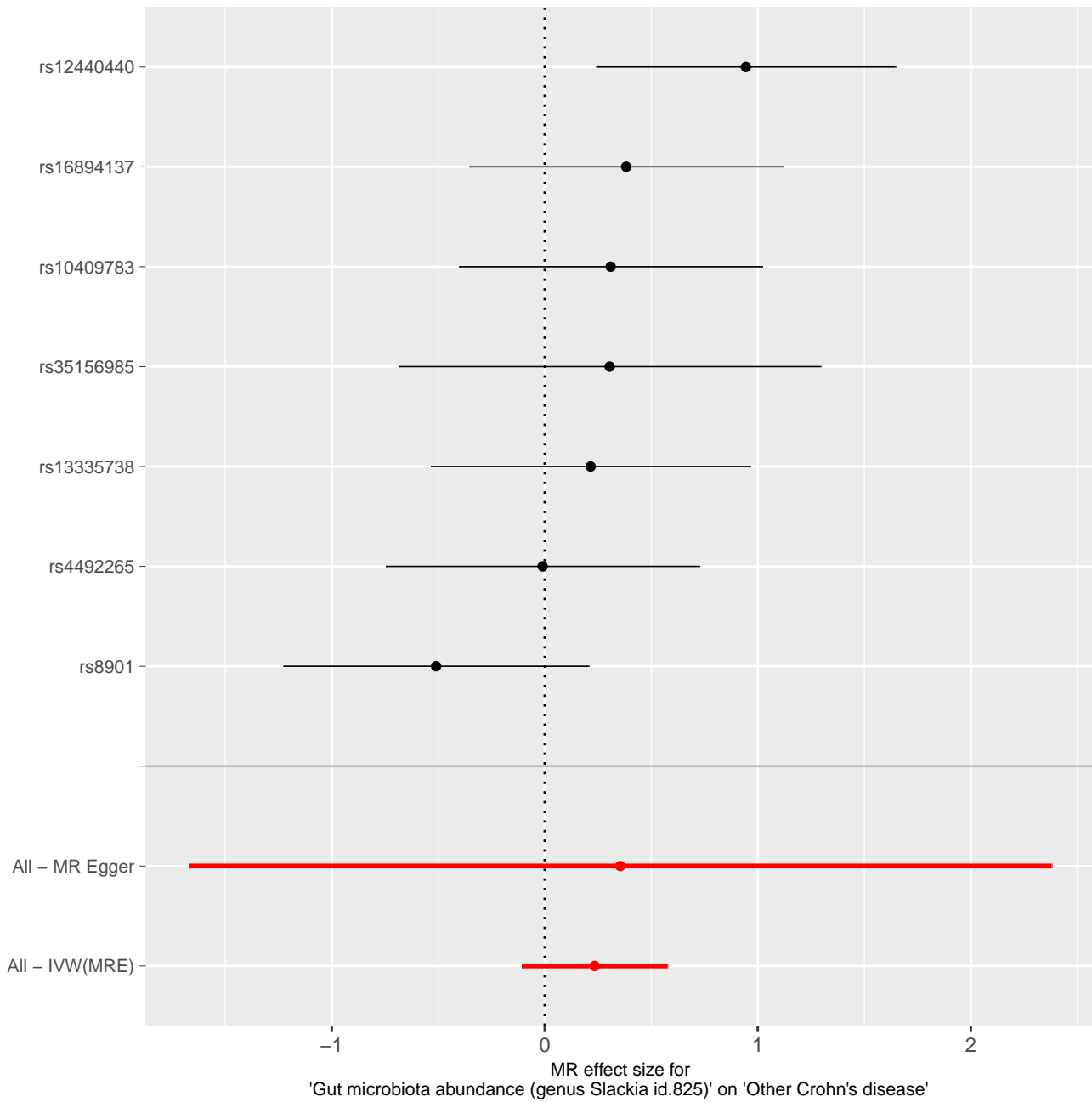


Batch 291 : Gut microbiota abundance (genus Sellimonas id.14369) on Other Crohn's disease

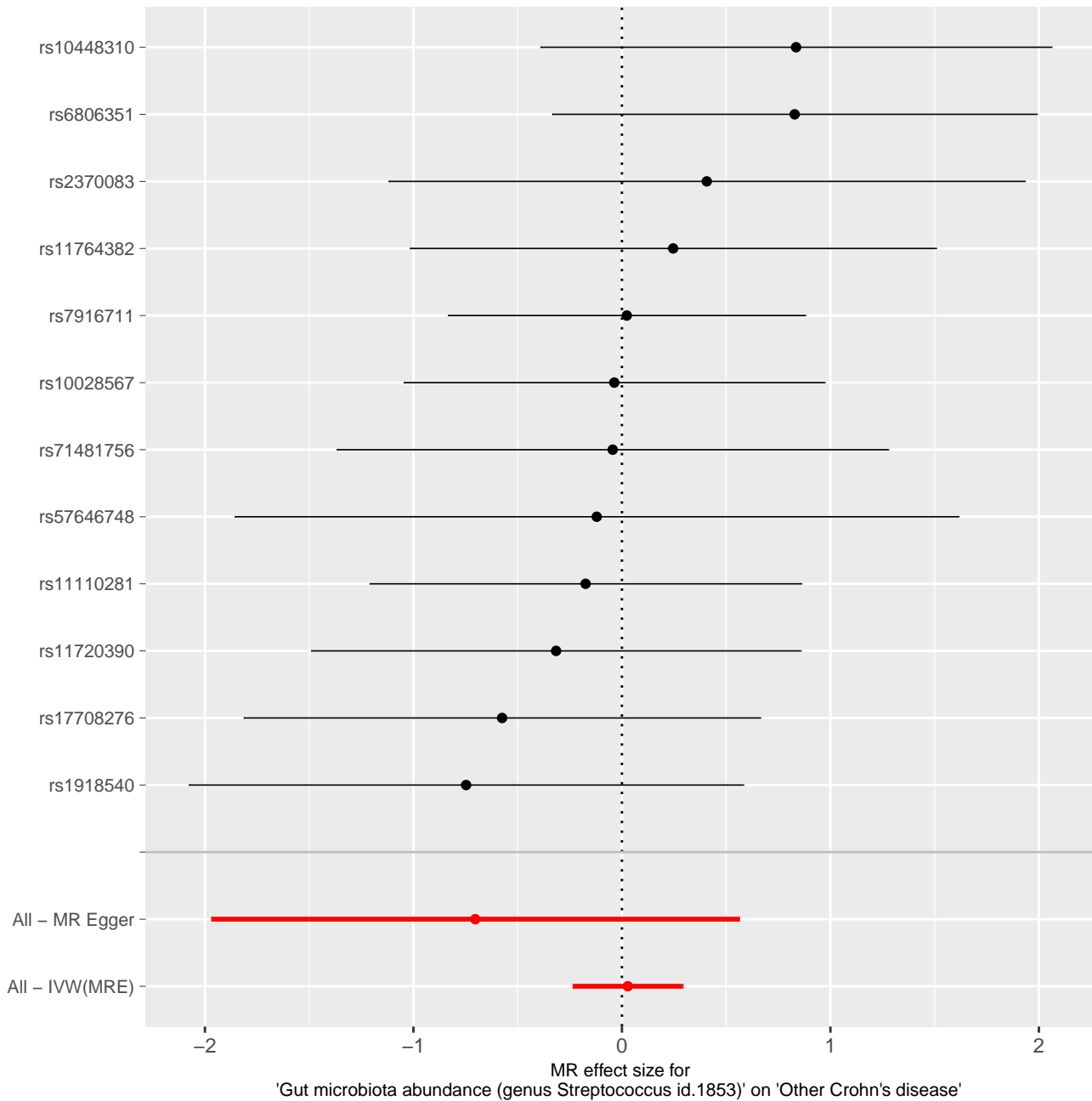


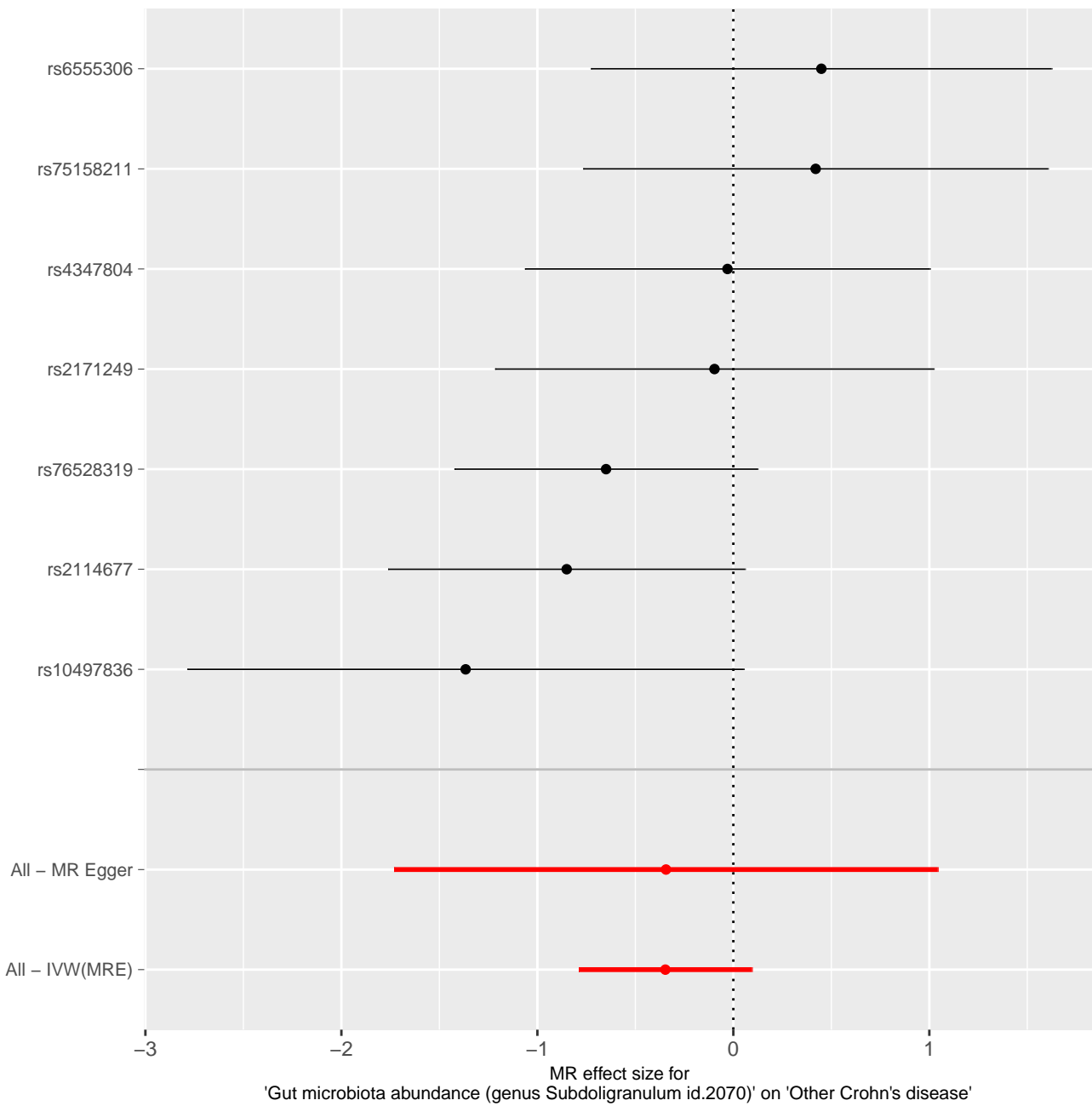


Batch 293 : Gut microbiota abundance (genus Slackia id.825) on Other Crohn's disease

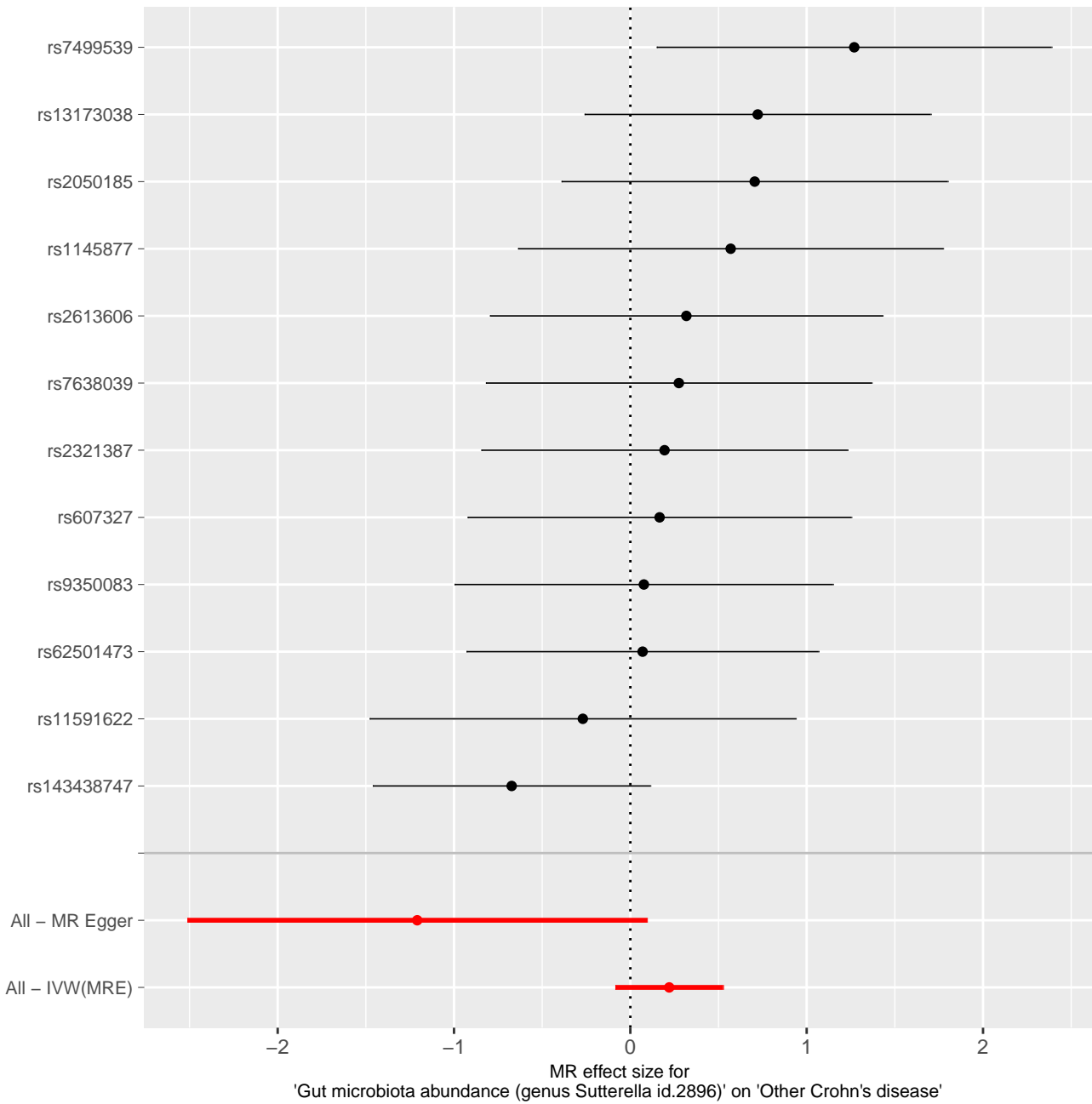


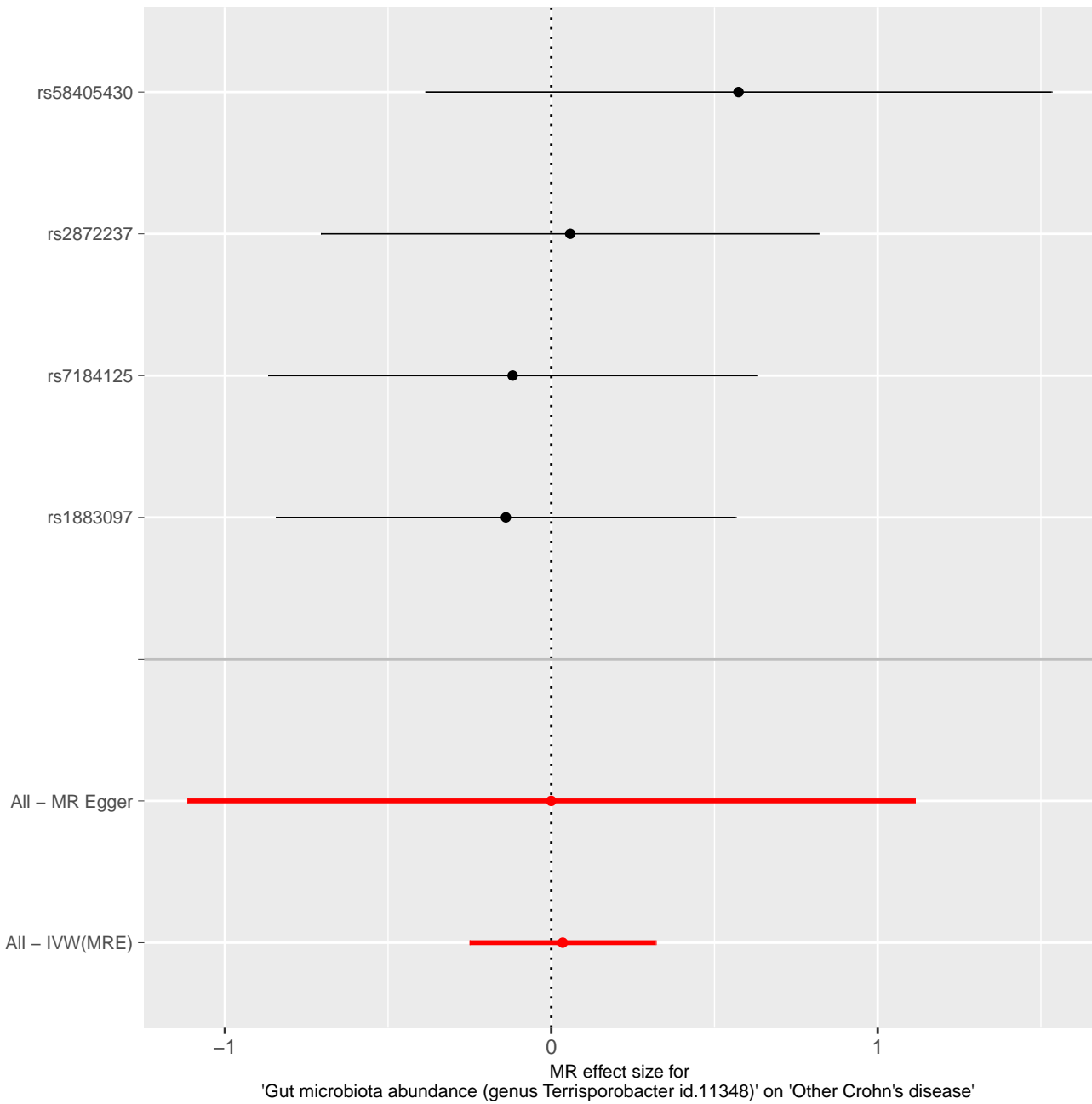
Batch 294 : Gut microbiota abundance (genus Streptococcus id.1853) on Other Crohn's disease



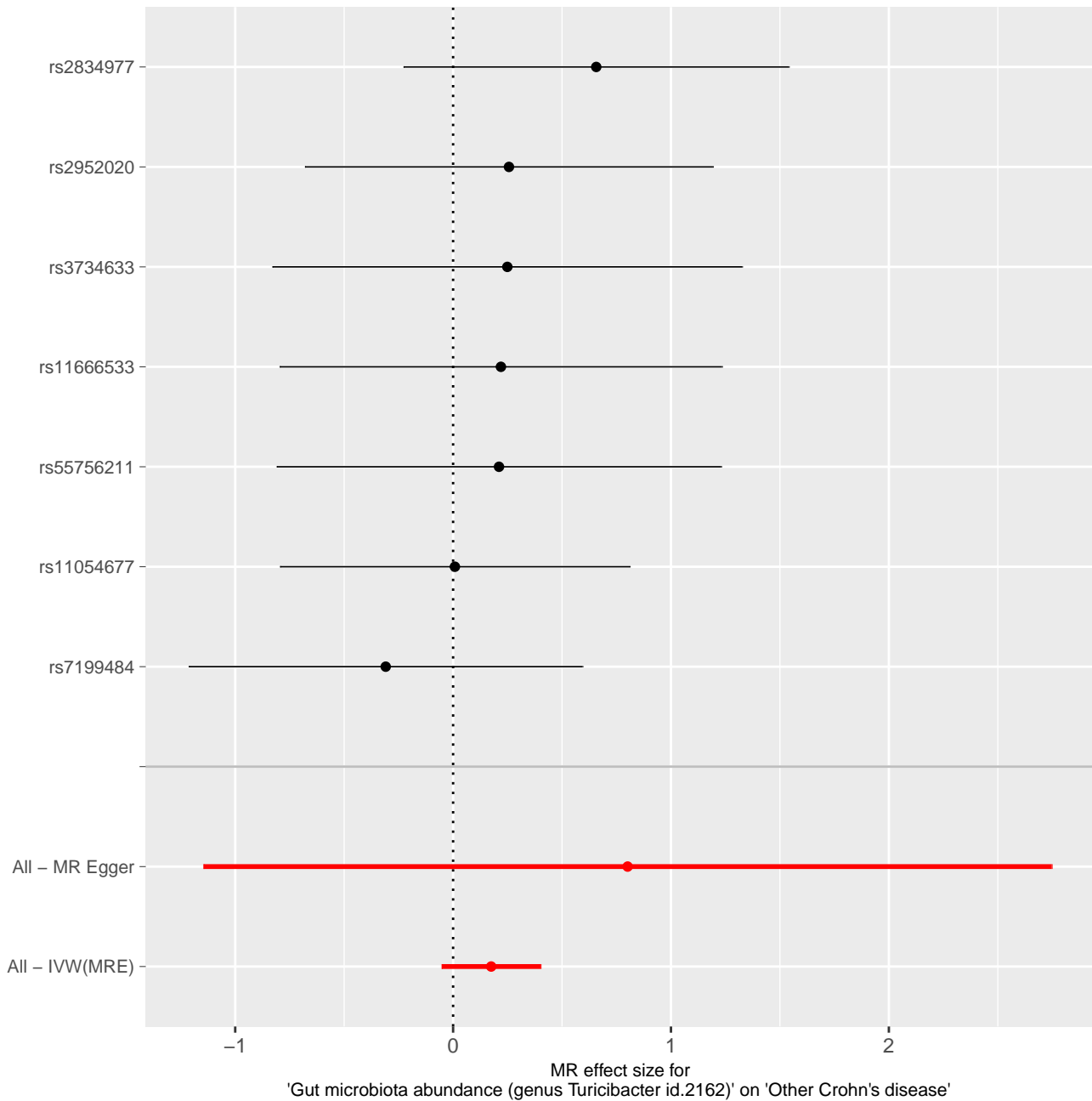


Batch 296 : Gut microbiota abundance (genus Sutterella id.2896) on Other Crohn's disease

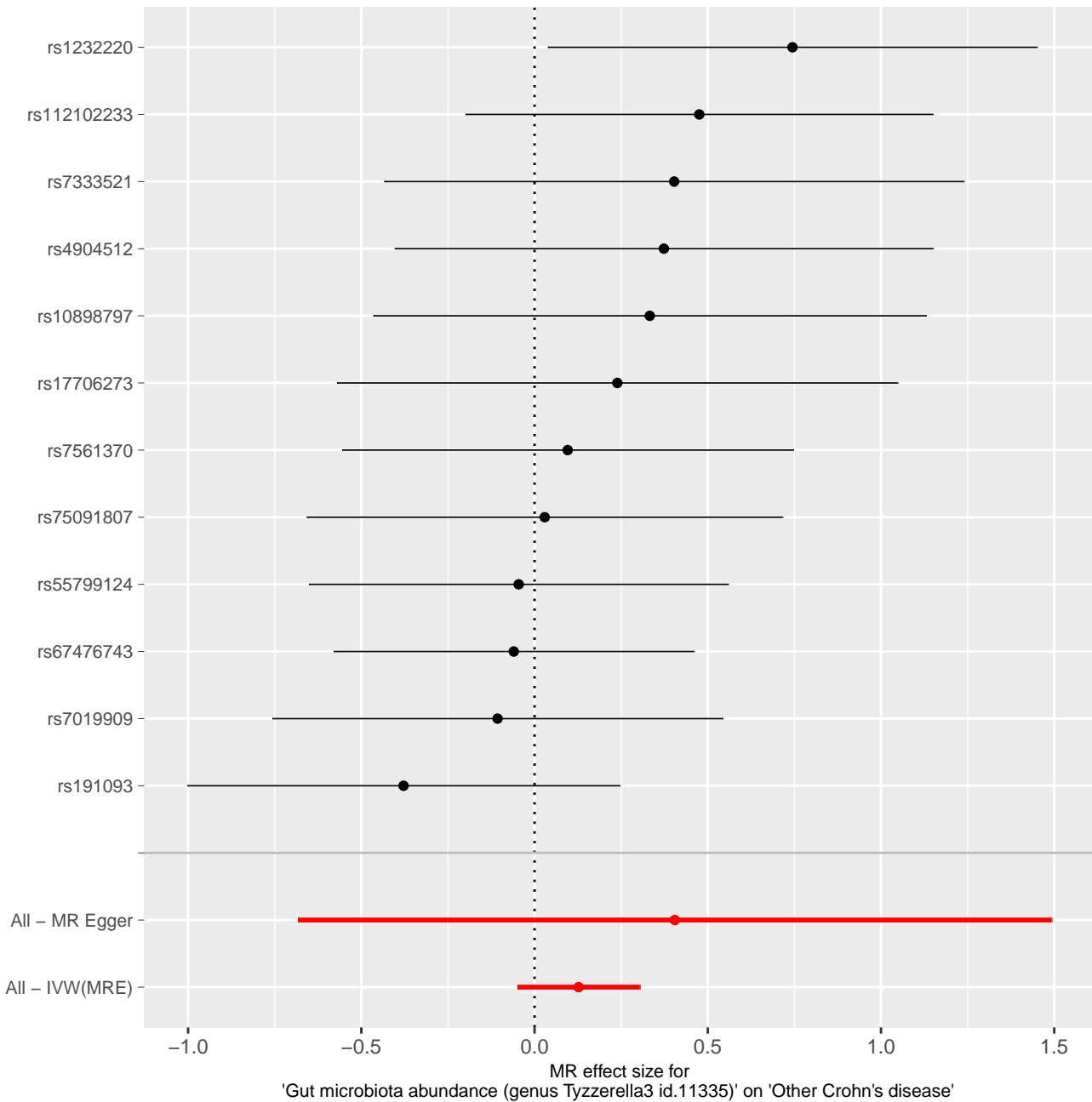


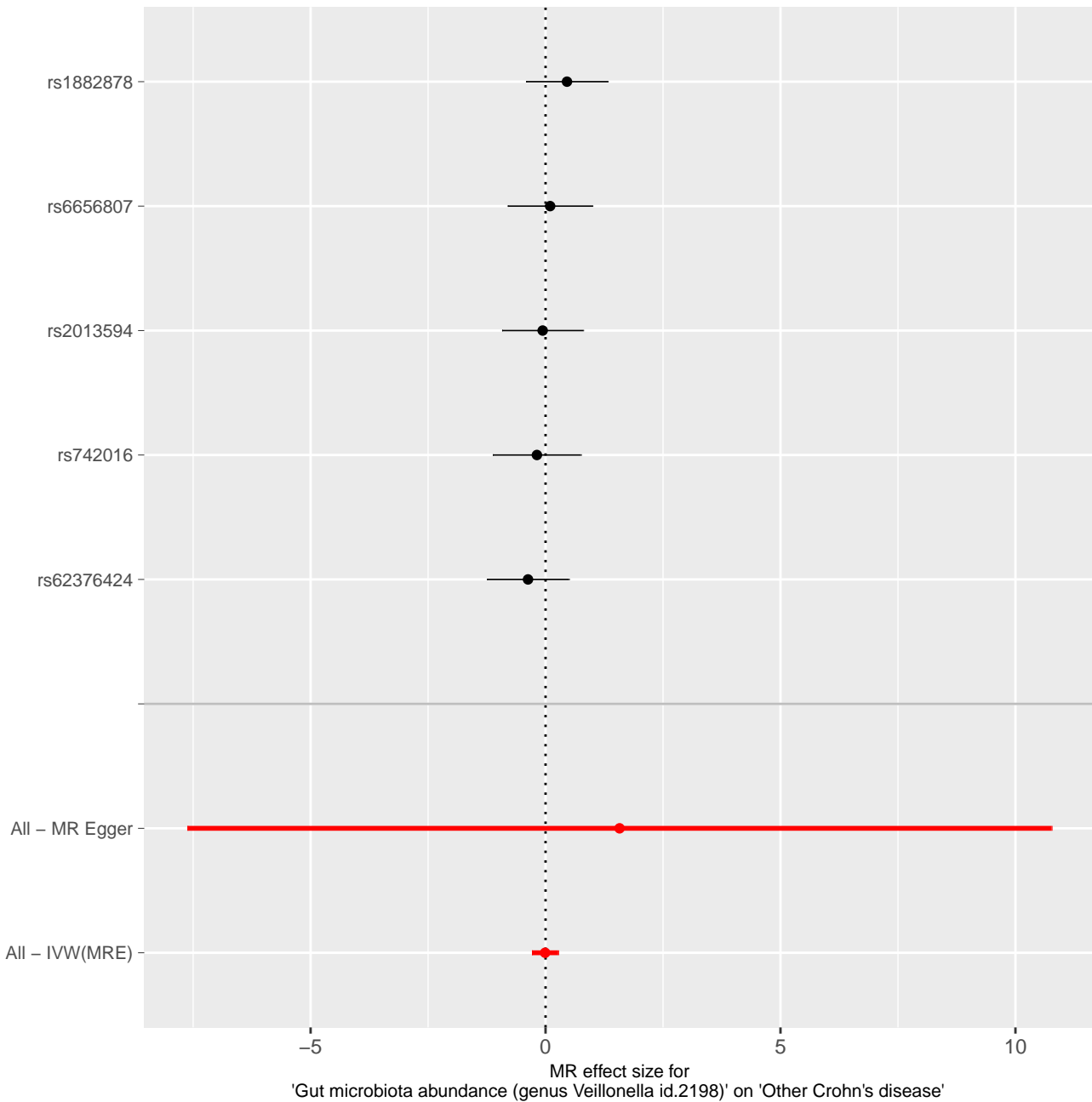


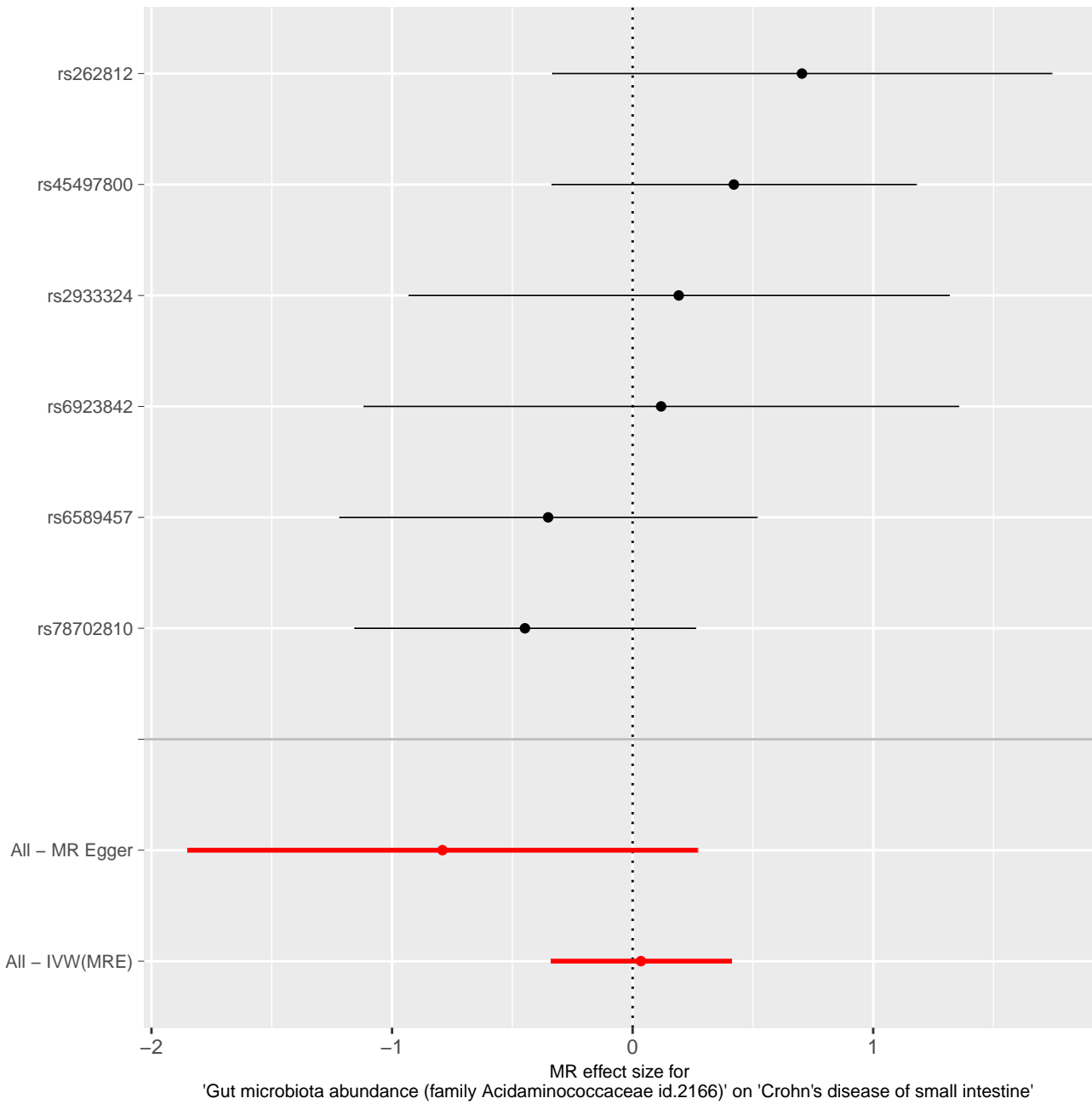
Batch 298 : Gut microbiota abundance (genus Turicibacter id.2162) on Other Crohn's disease

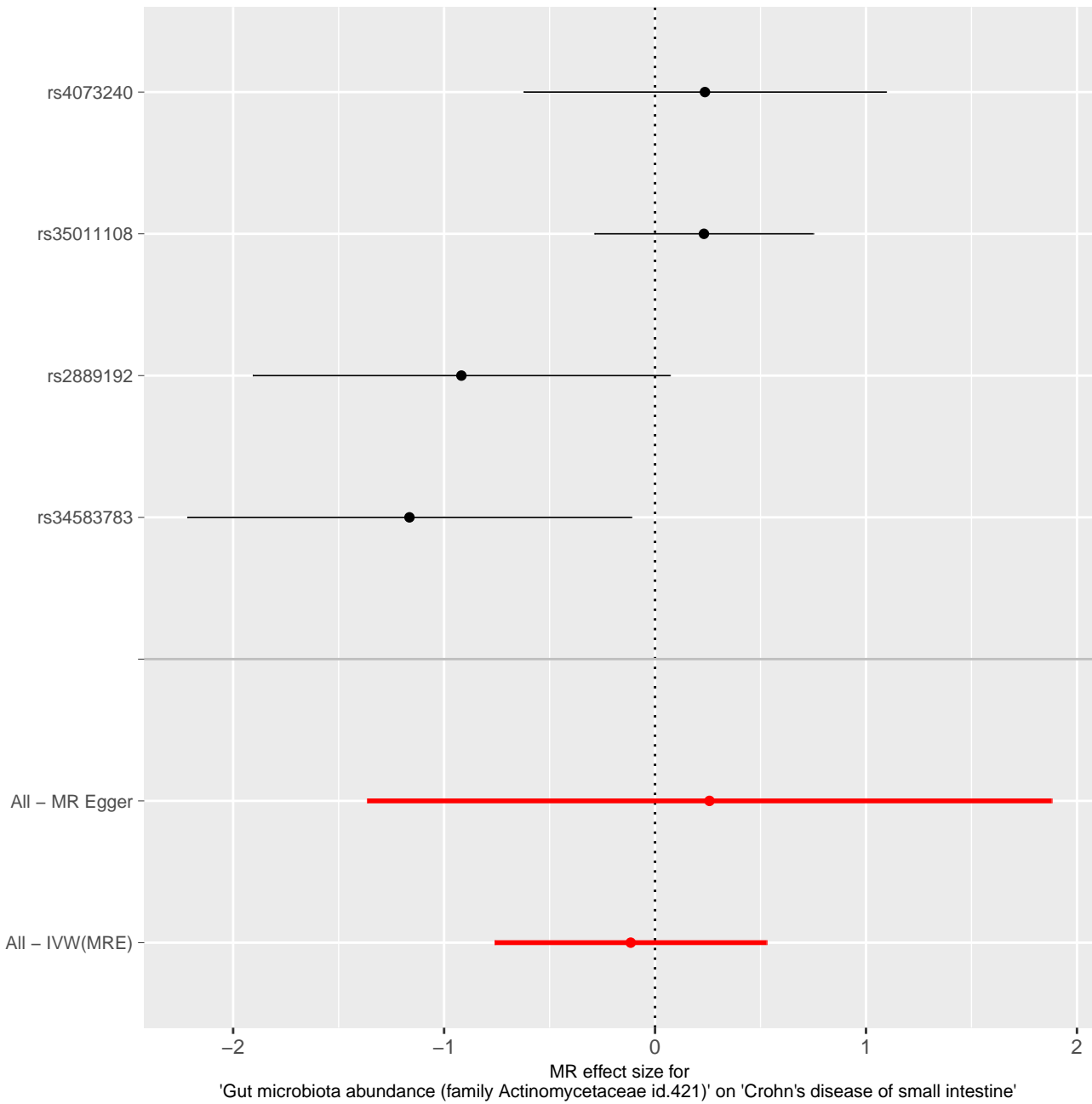


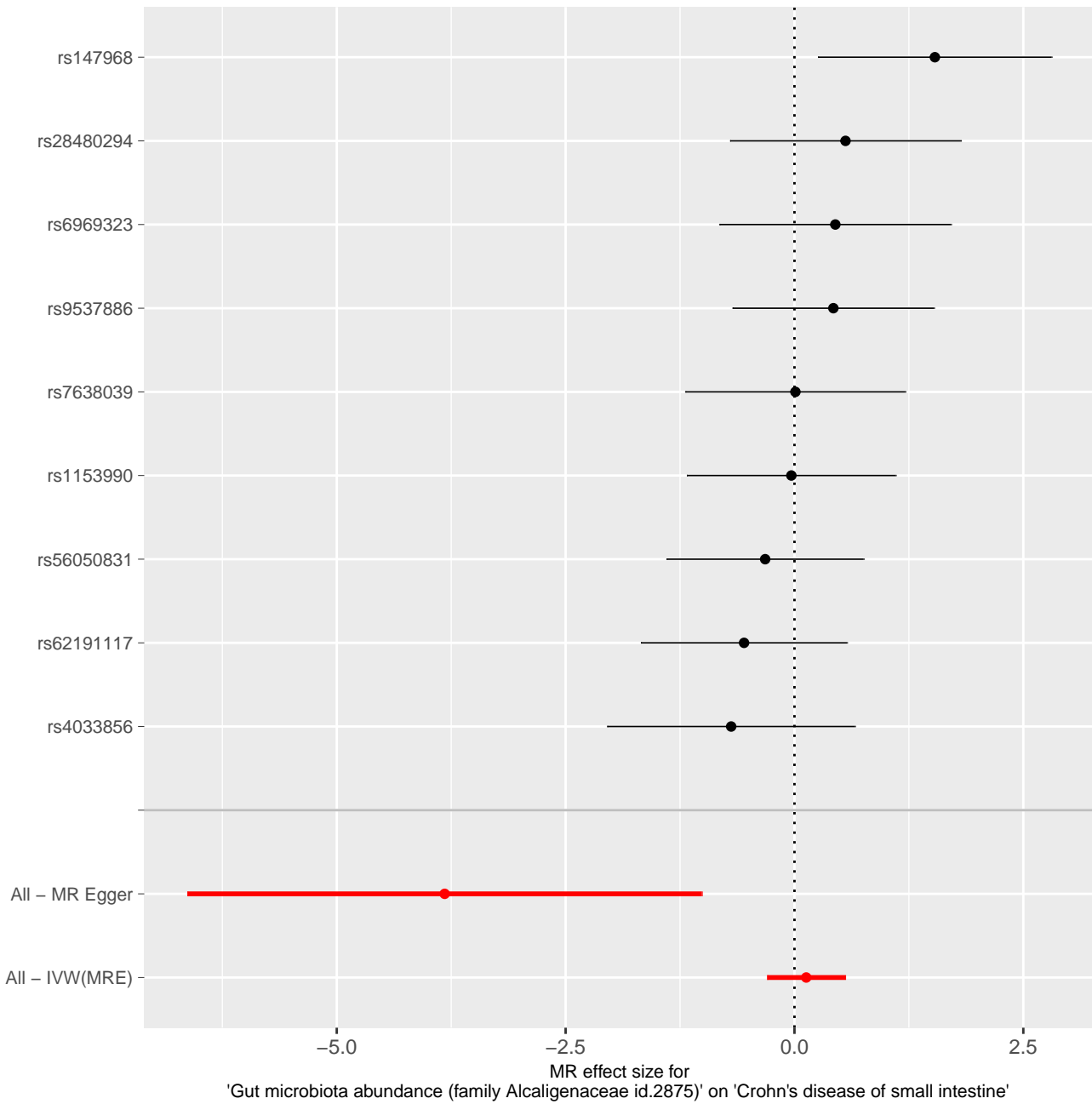
Batch 299 : Gut microbiota abundance (genus Tyzzerella3 id.11335) on Other Crohn's disease



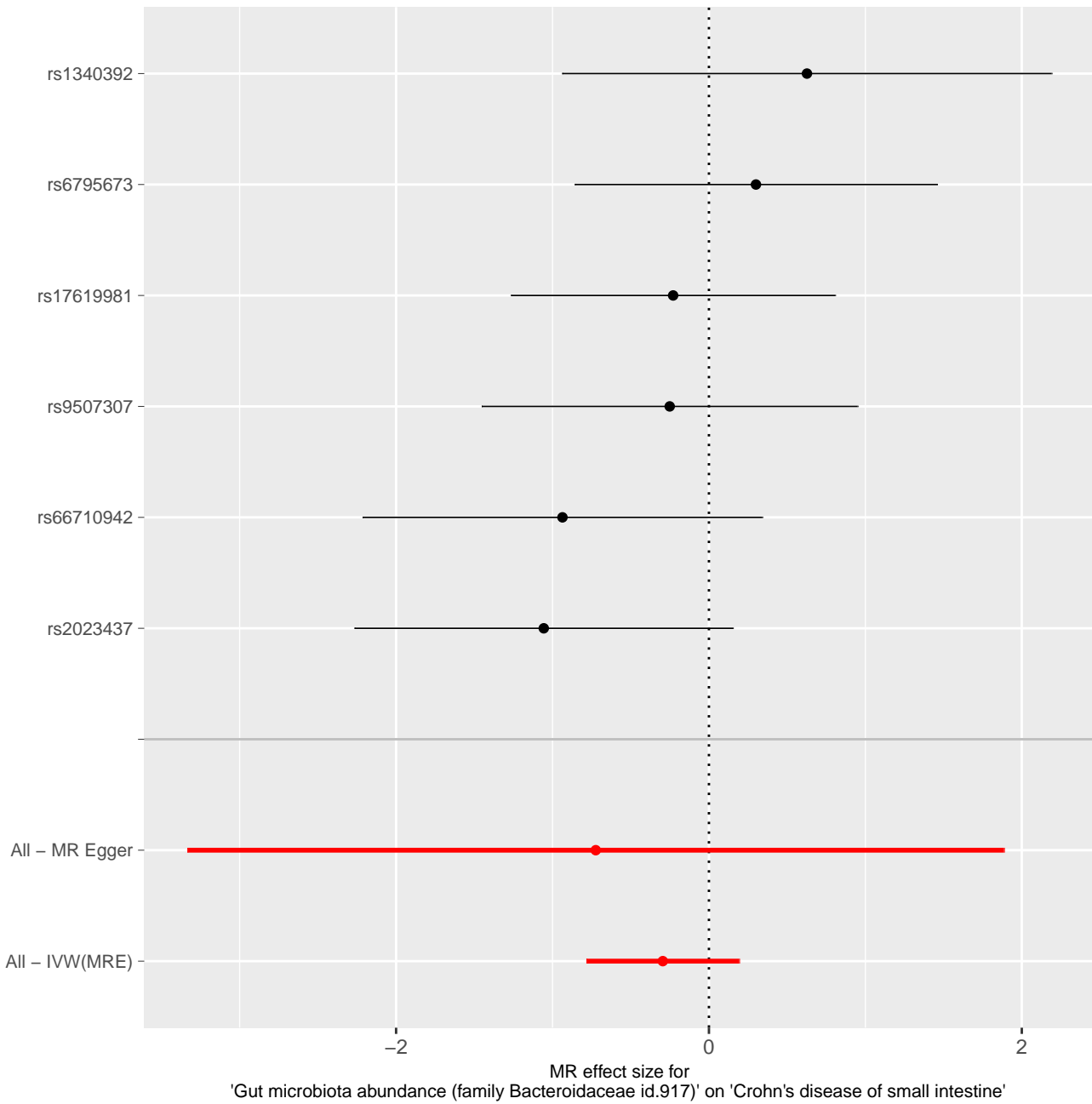


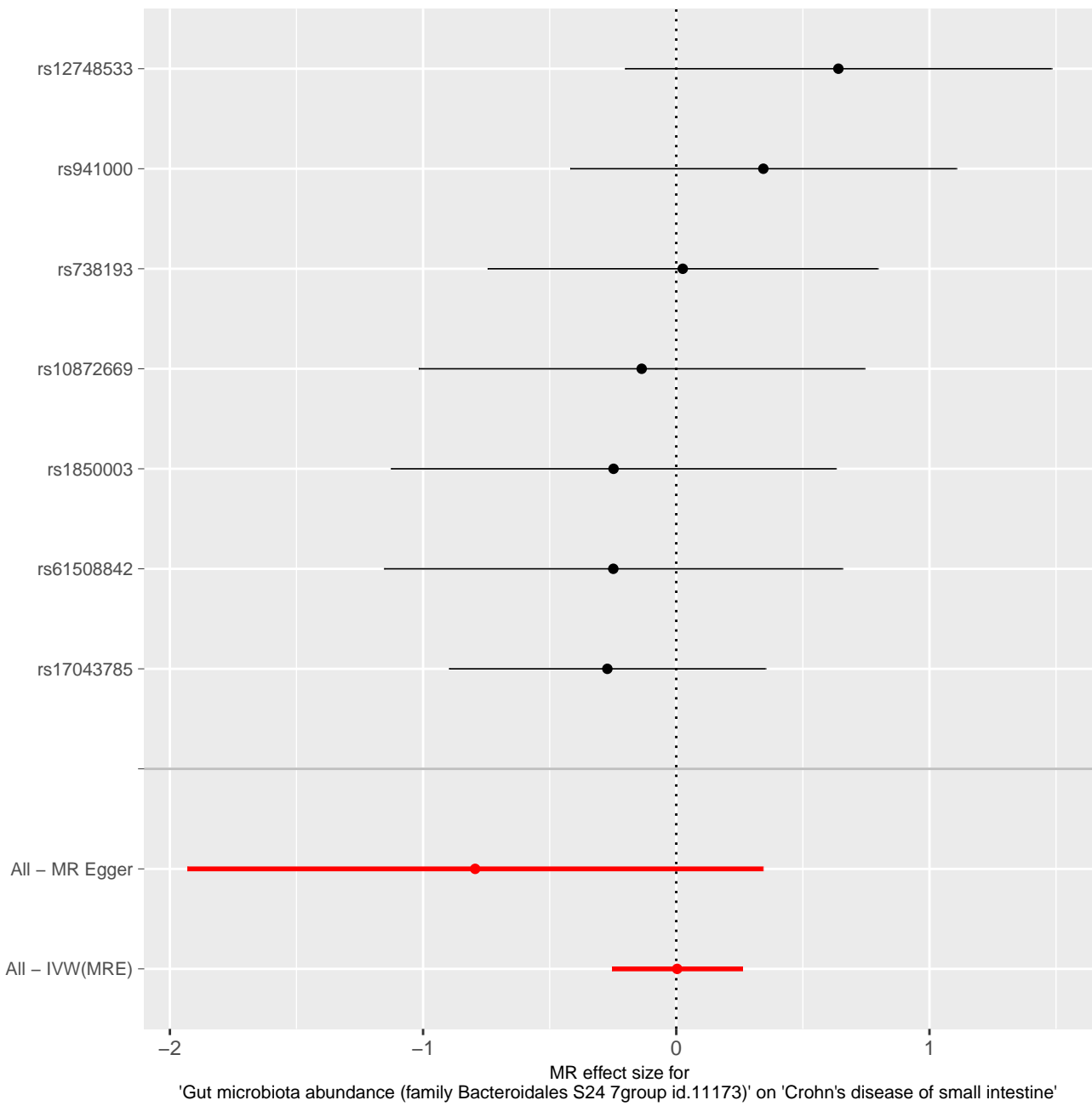


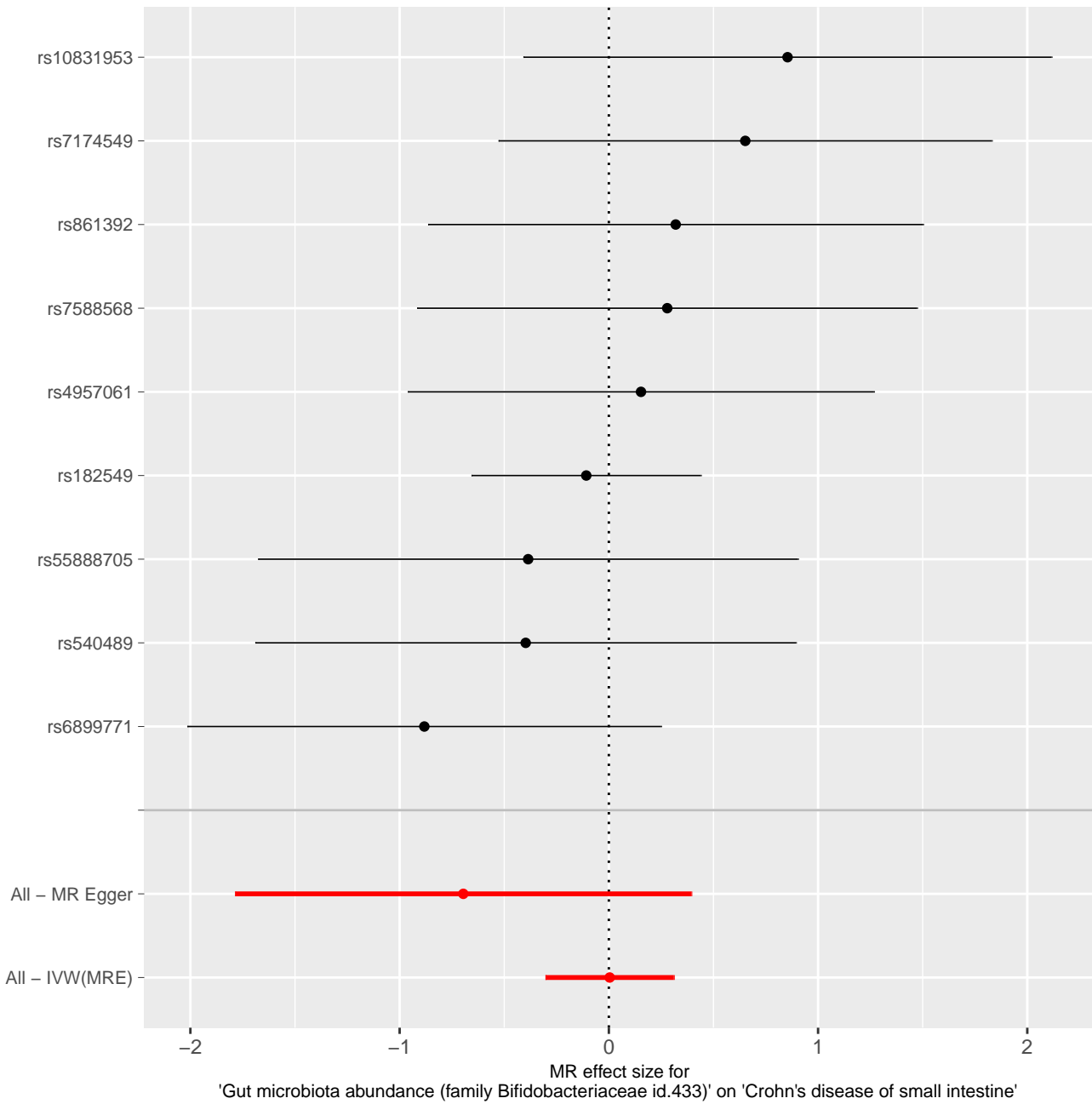


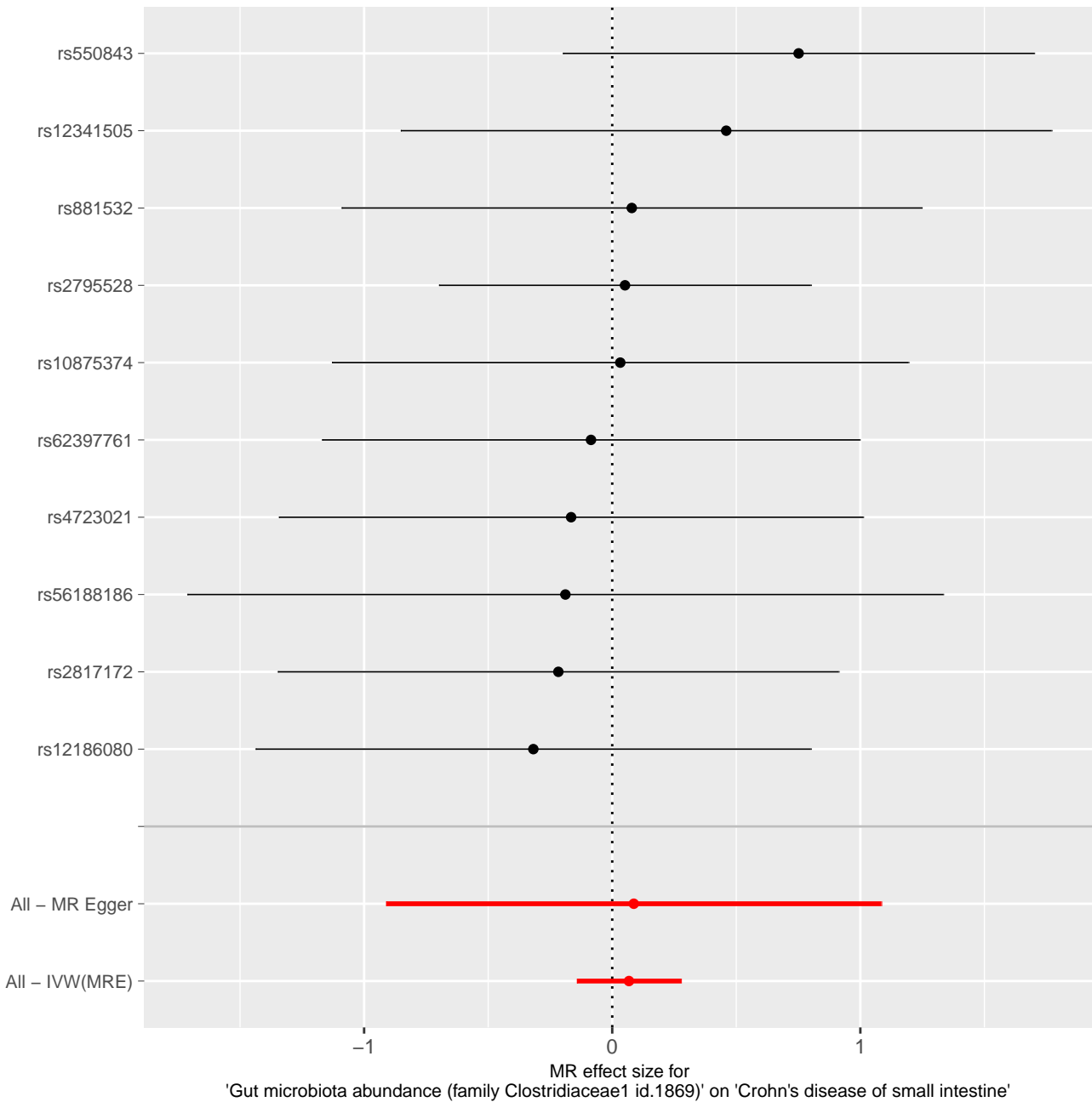


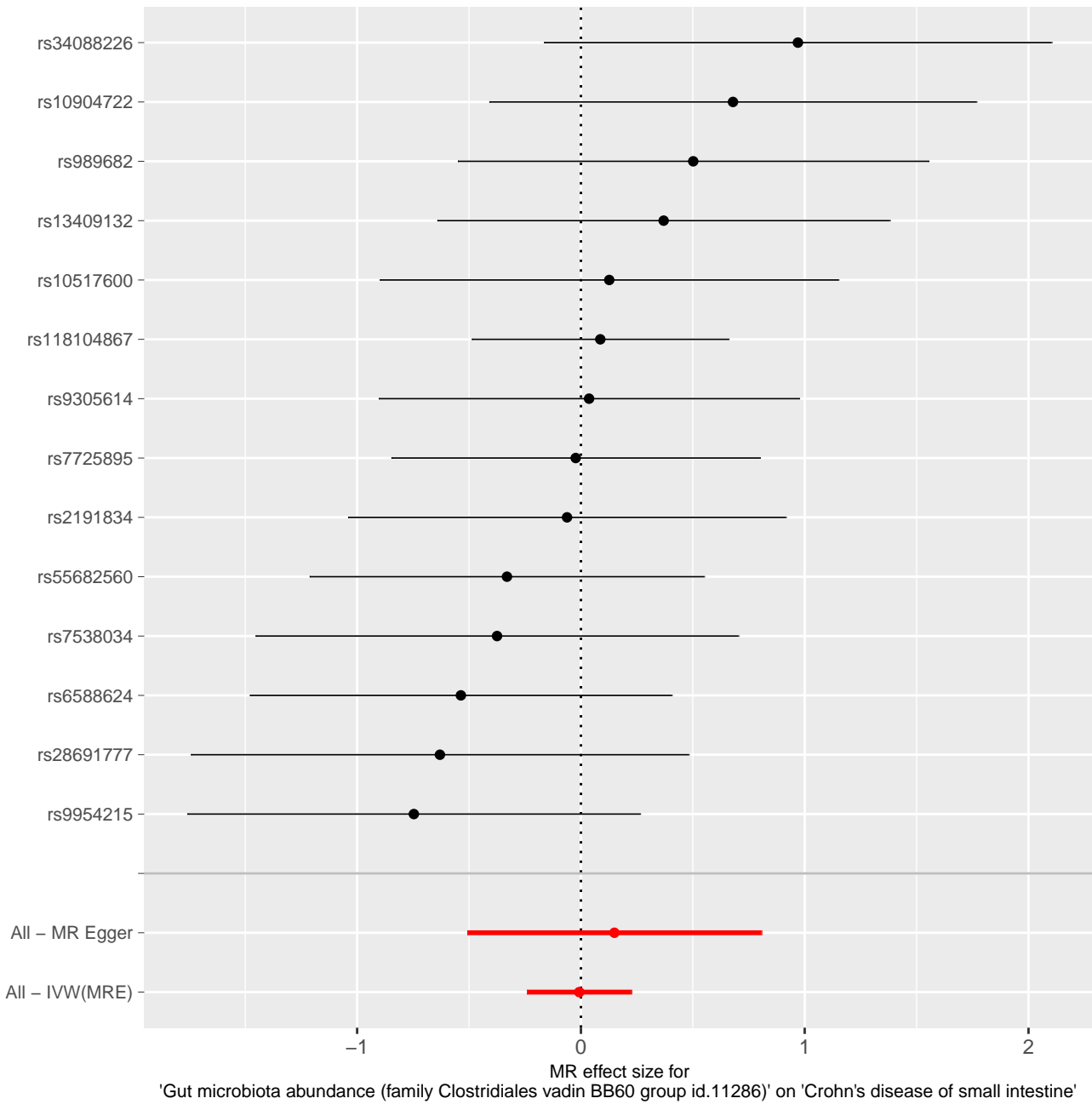
Batch 304 : Gut microbiota abundance (family Bacteroidaceae id.917) on Crohn's disease of small intestine



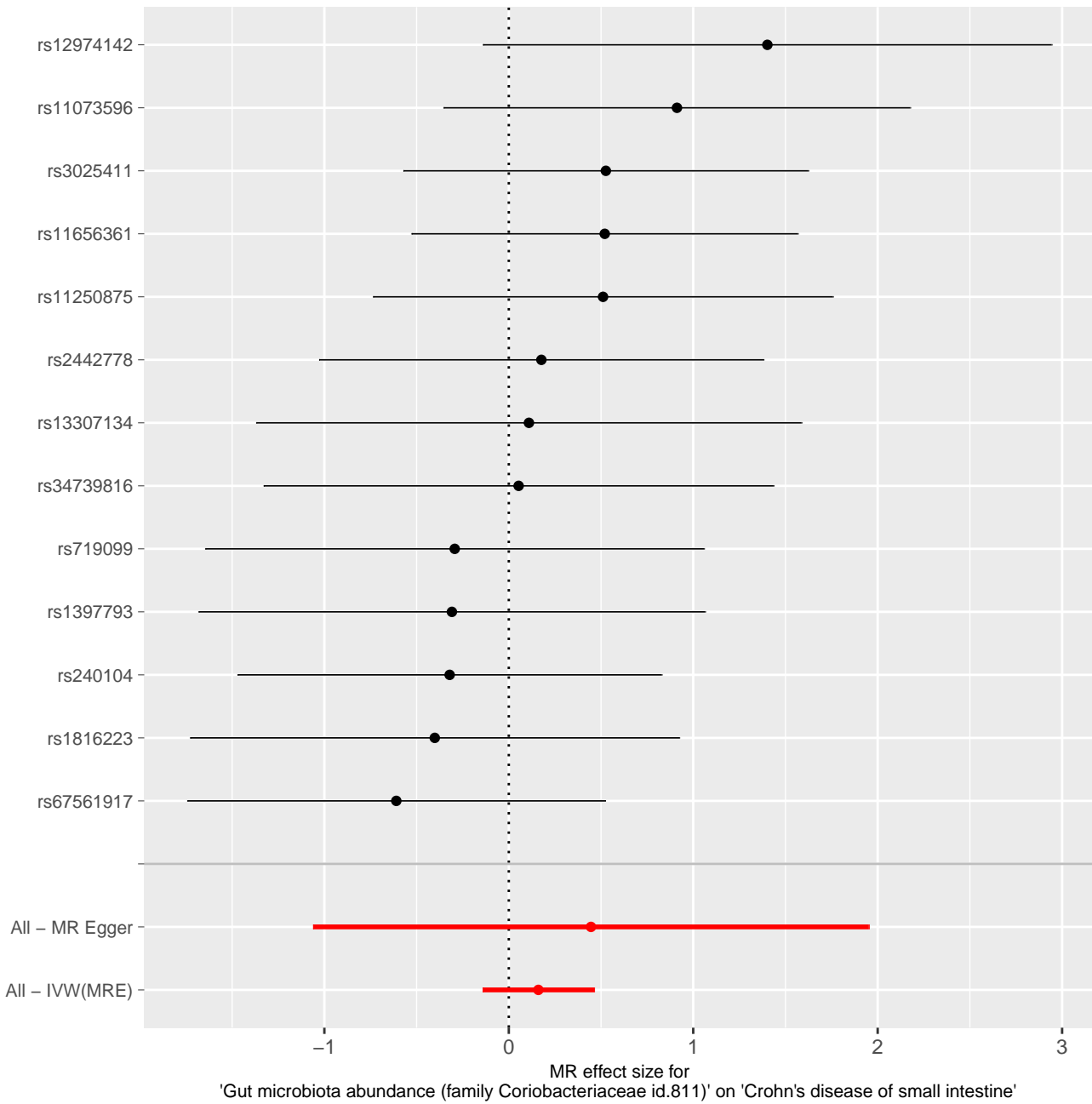


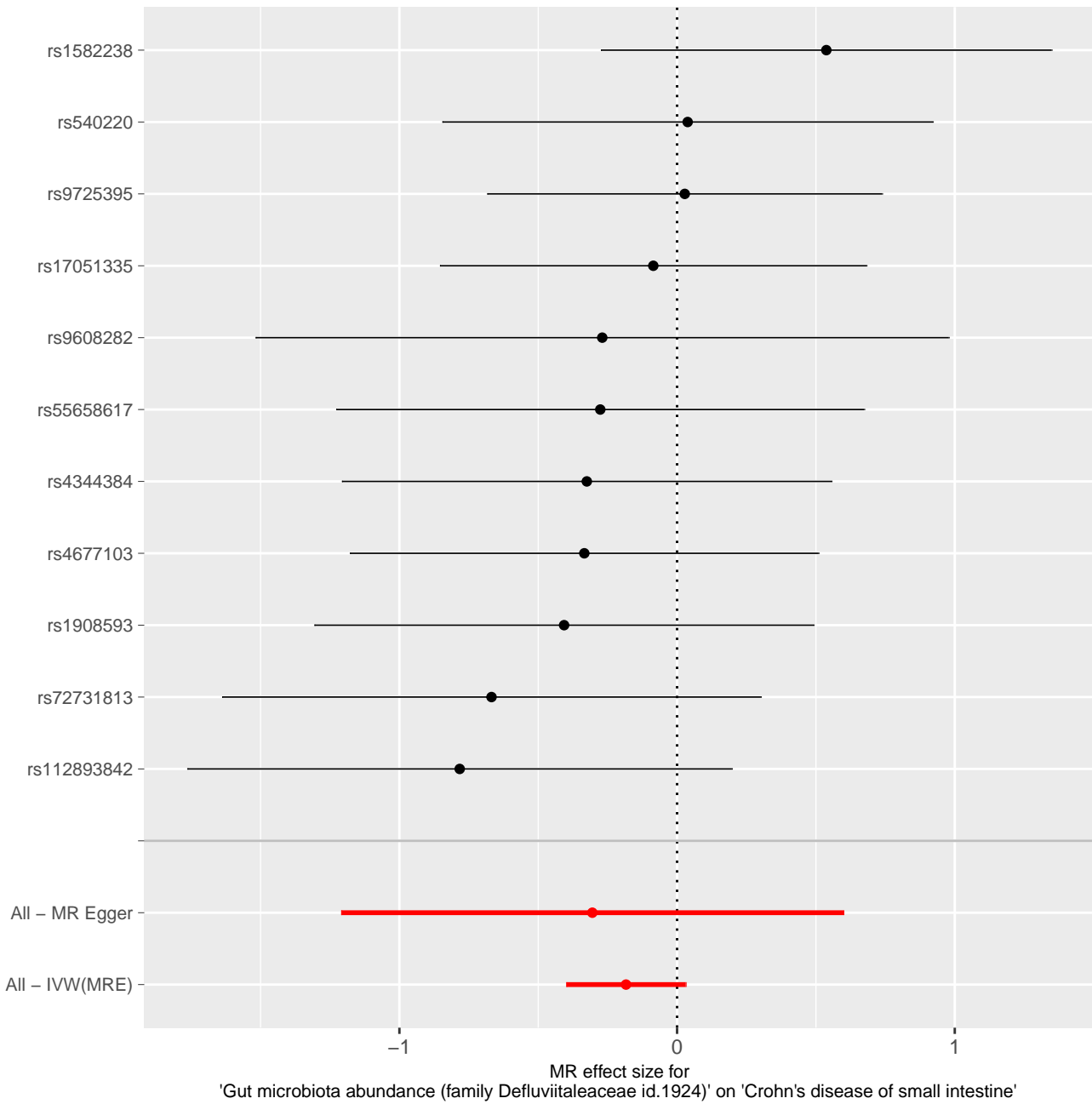




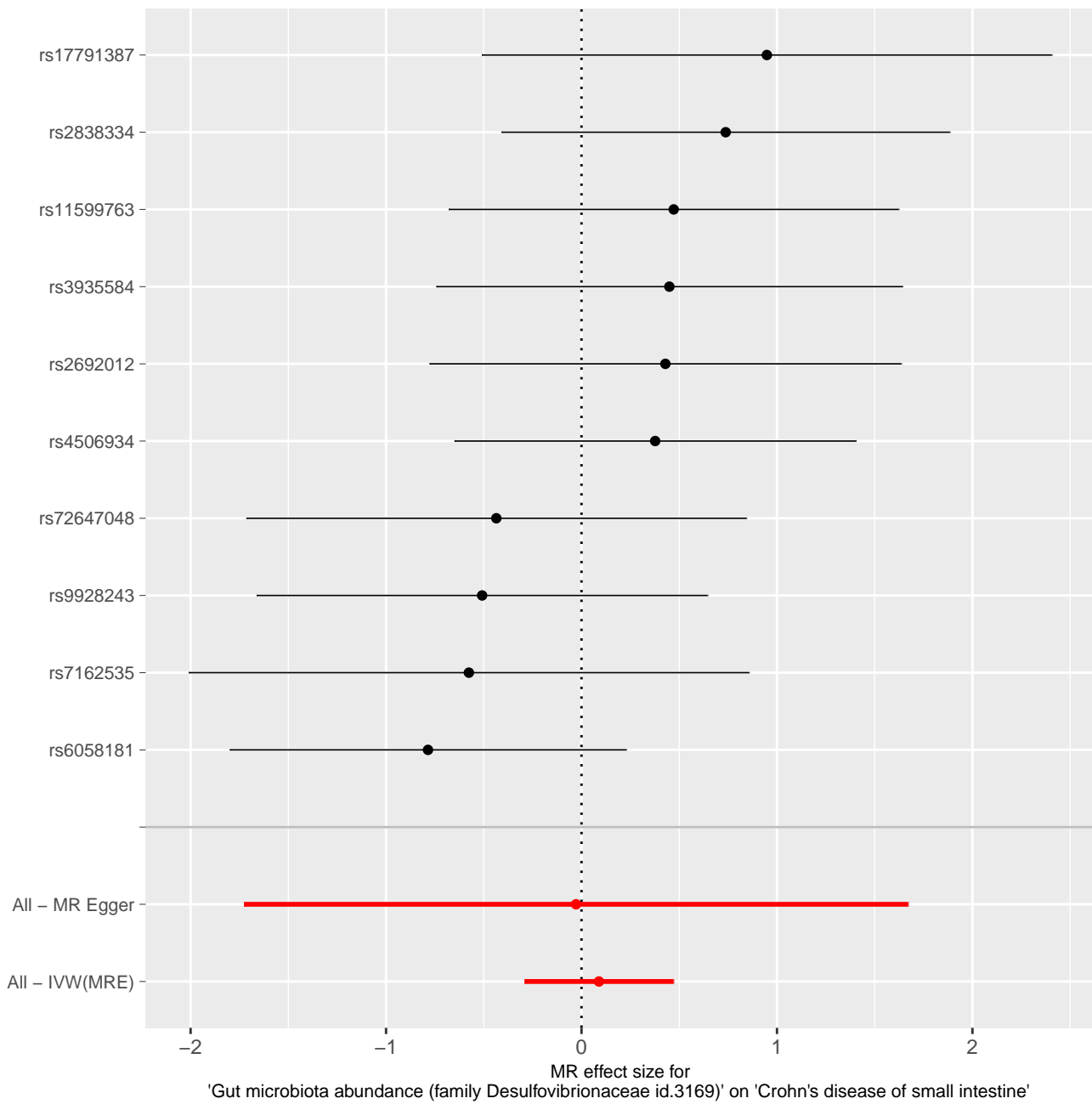


Batch 310 : Gut microbiota abundance (family Coriobacteriaceae id.811) on Crohn's disease of small intestine

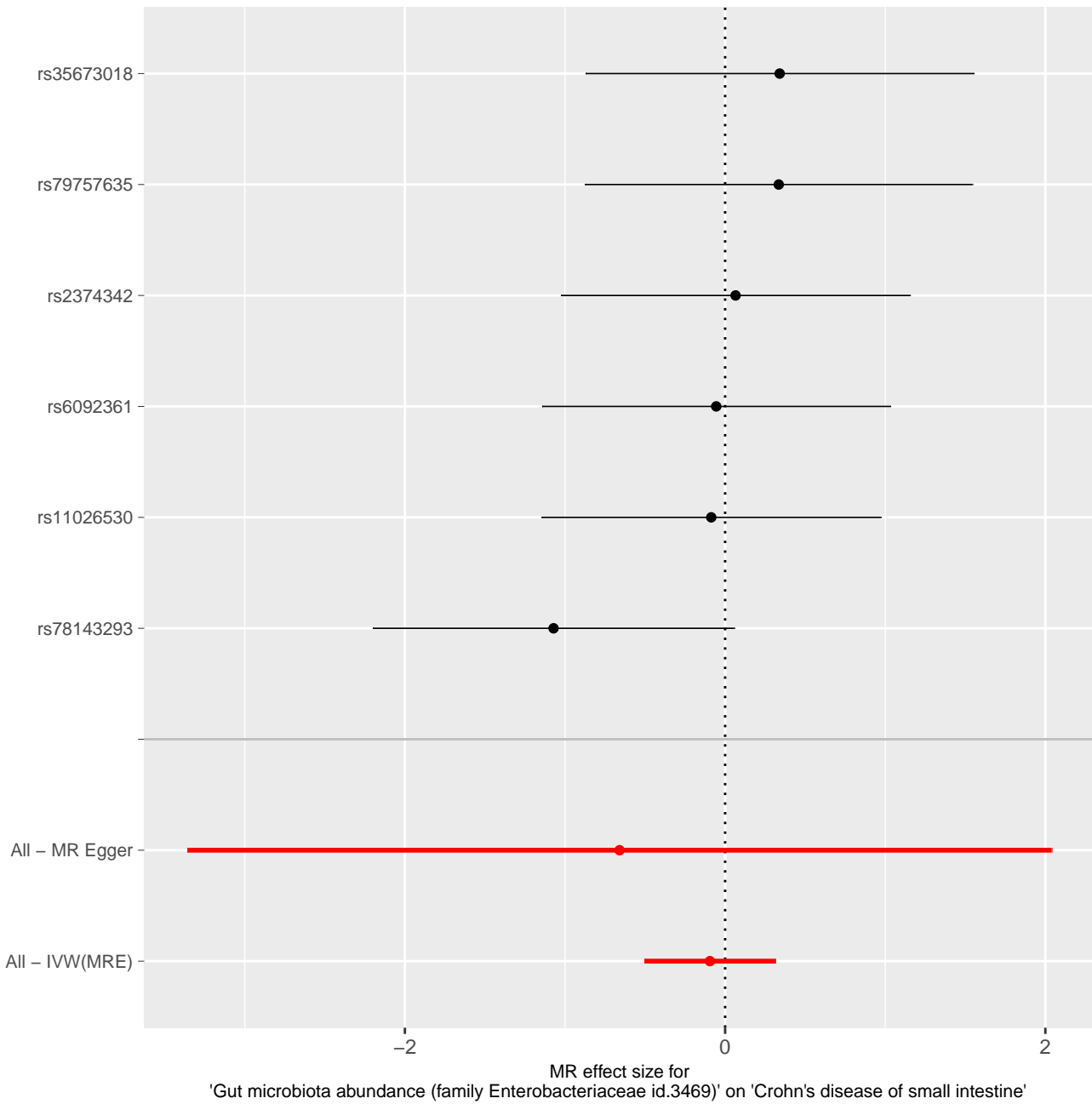


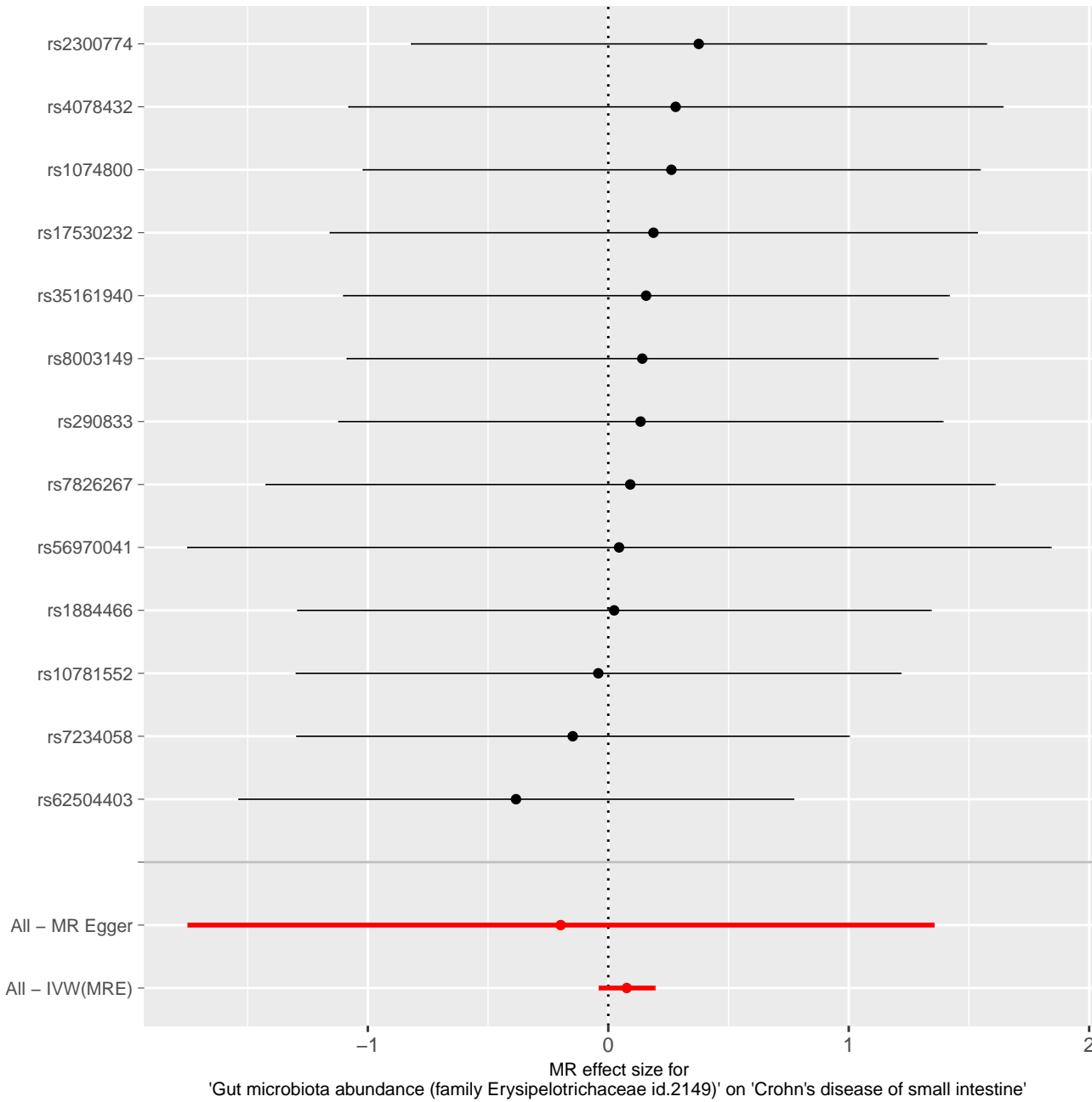


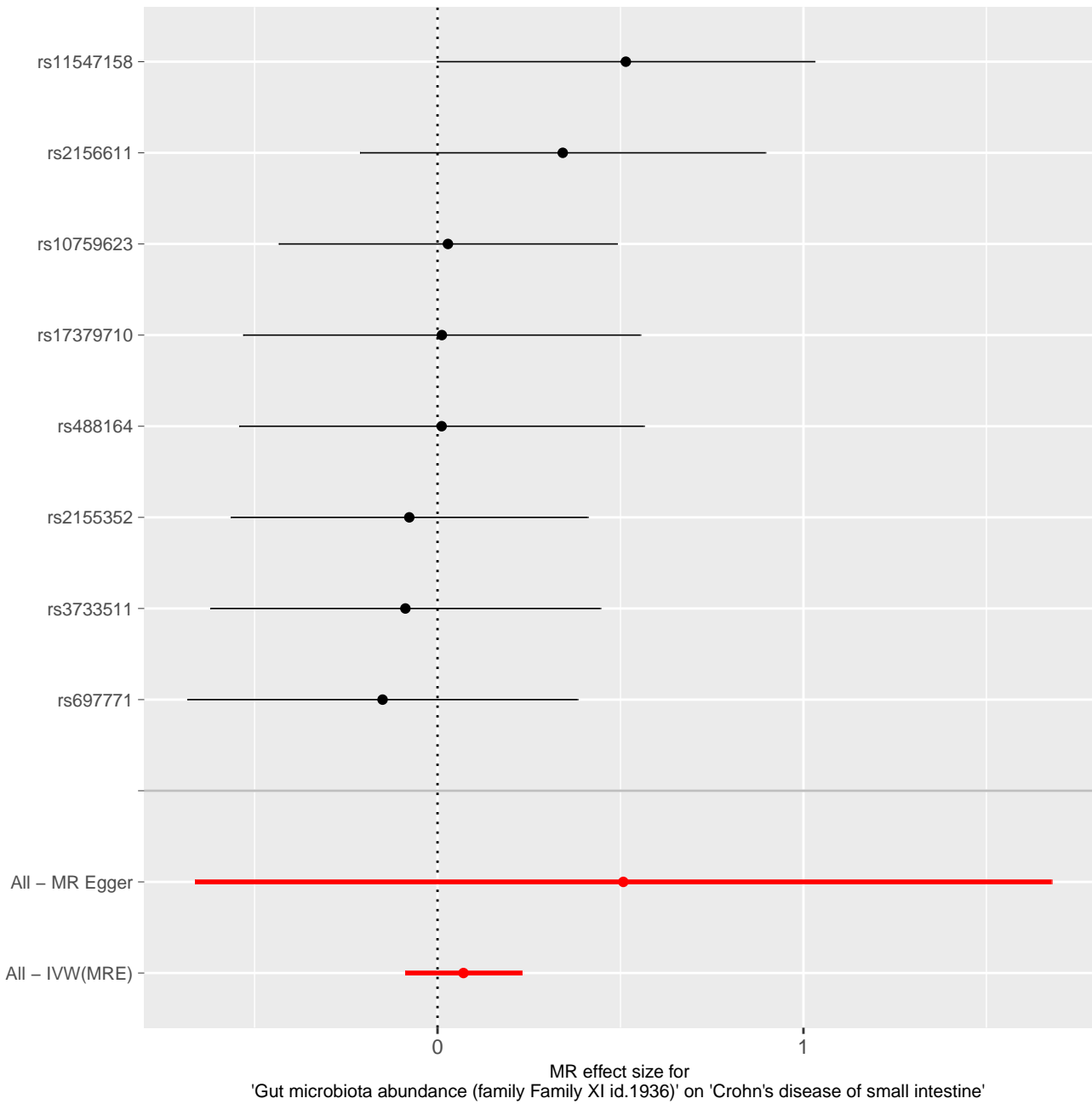
Batch 312 : Gut microbiota abundance (family Desulfovibrionaceae id.3169) on Crohn's disease of small intestine

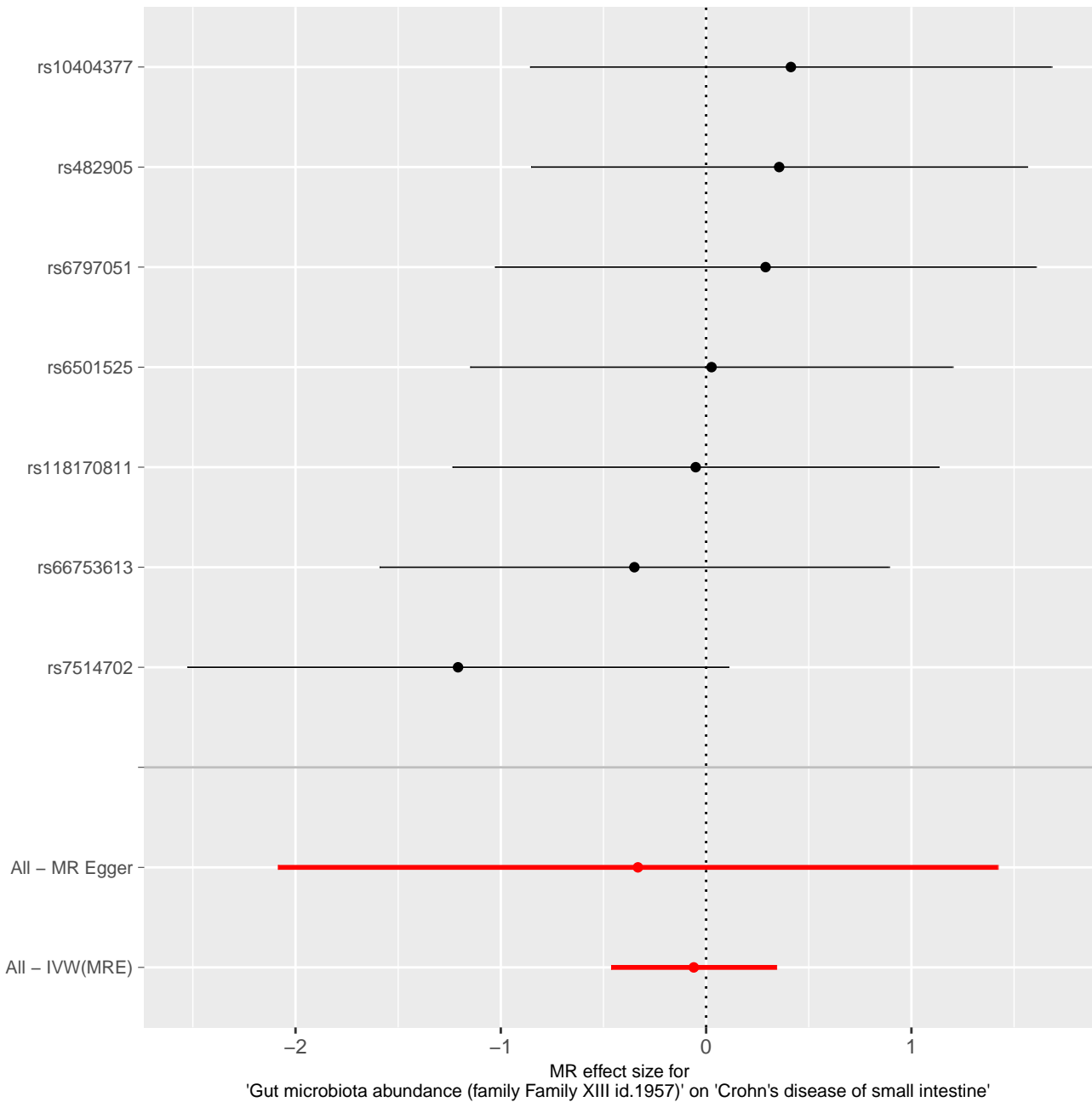


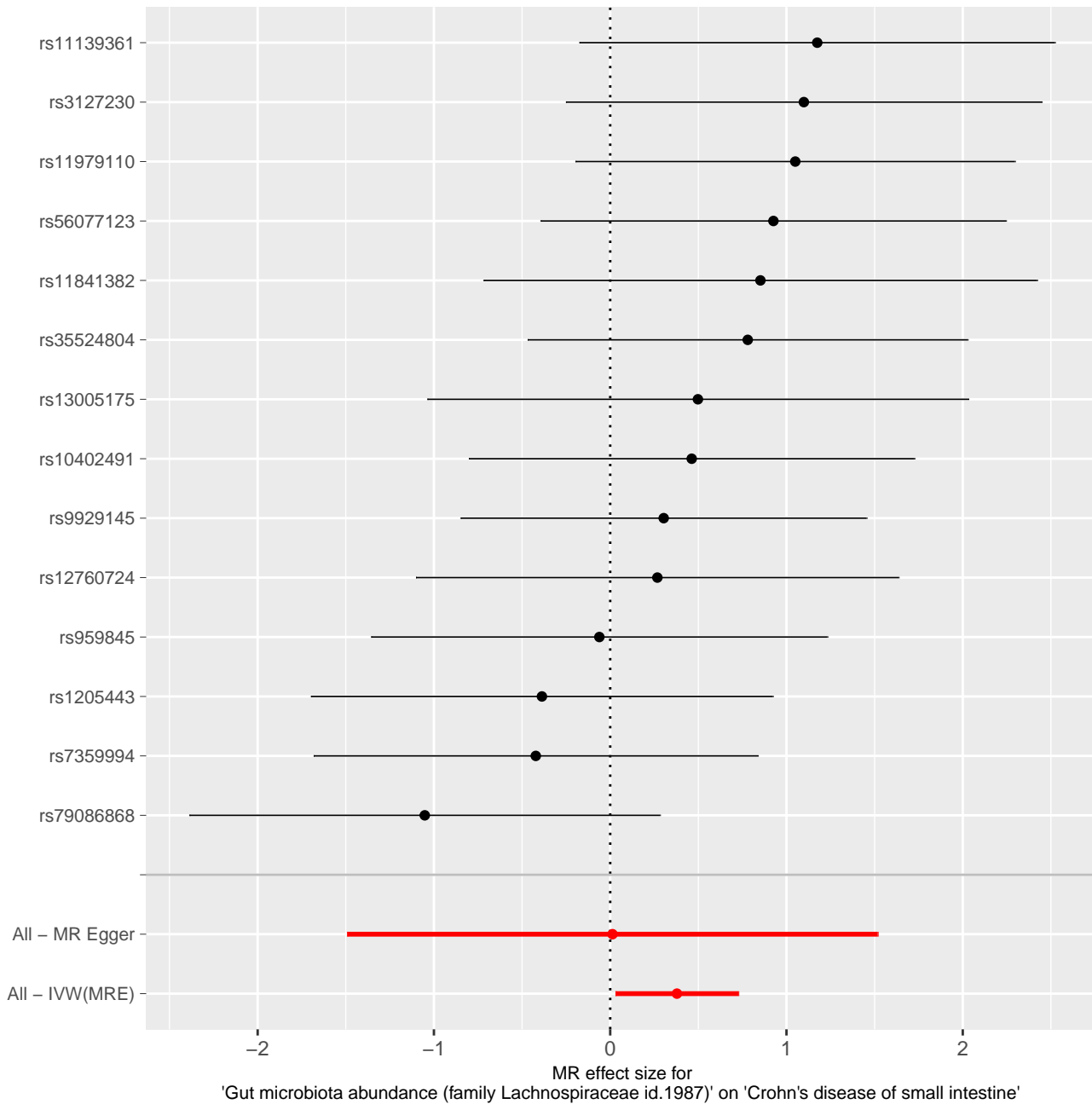
Batch 313 : Gut microbiota abundance (family Enterobacteriaceae id.3469) on Crohn's disease of small intestine

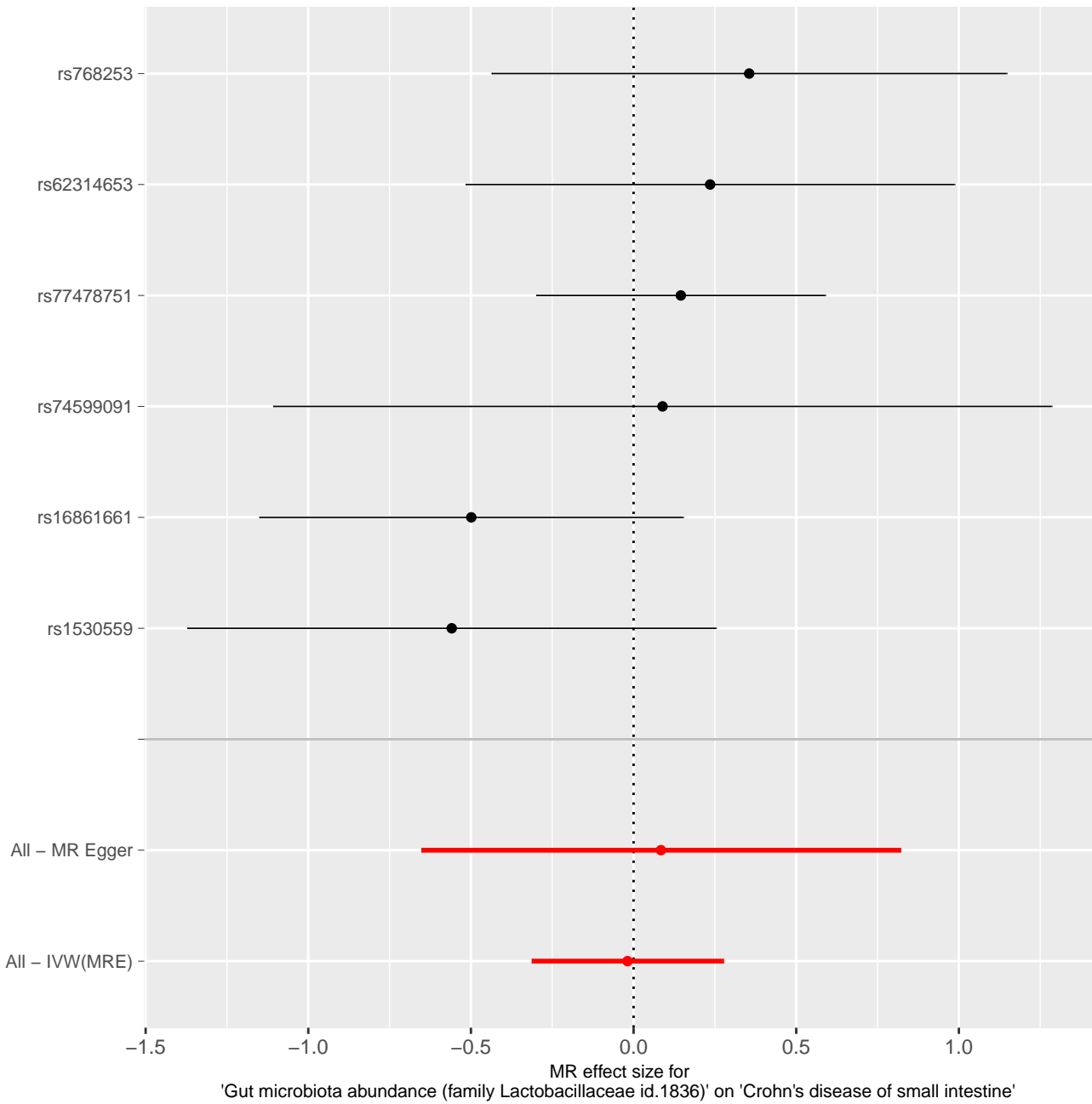


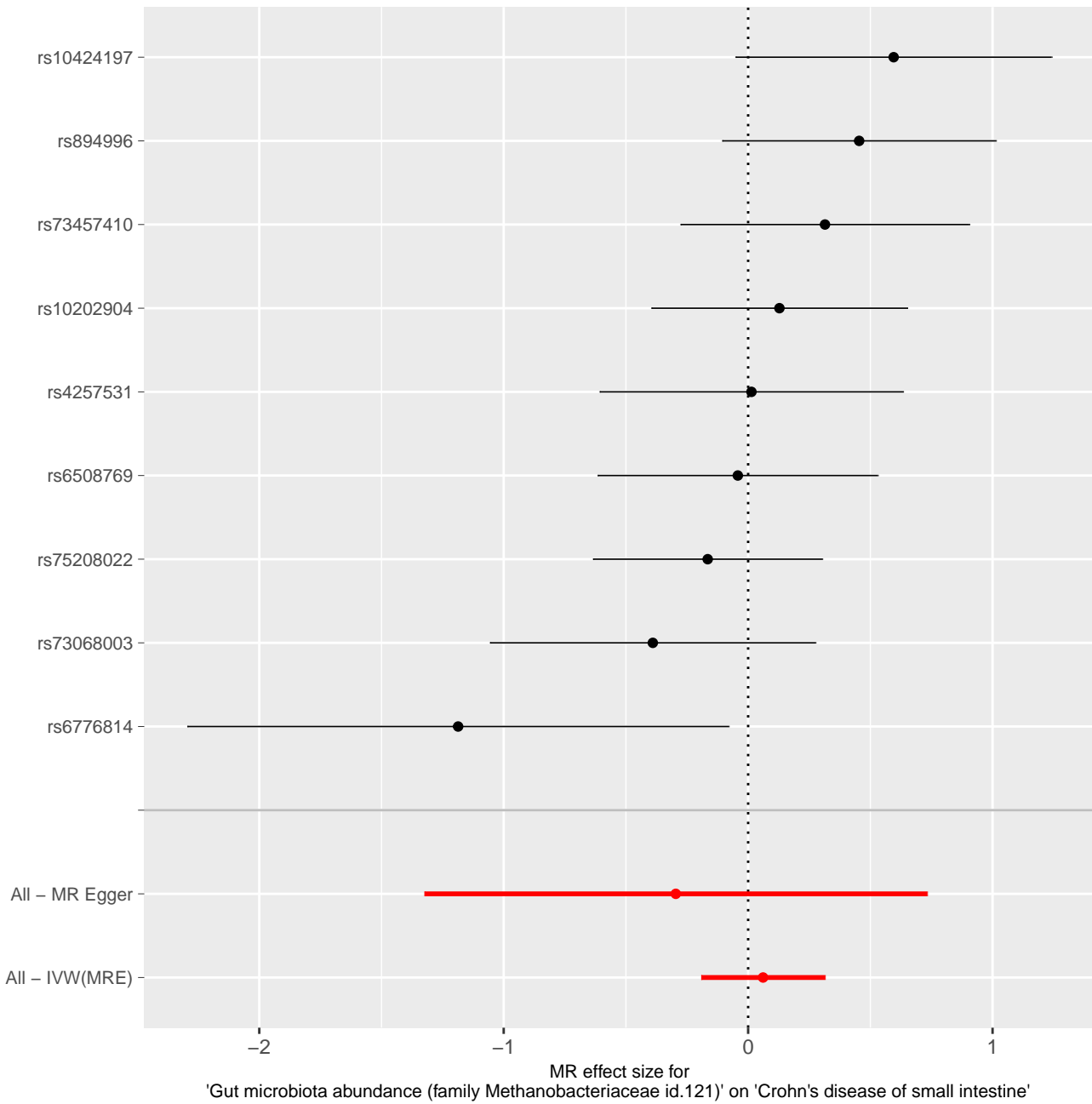


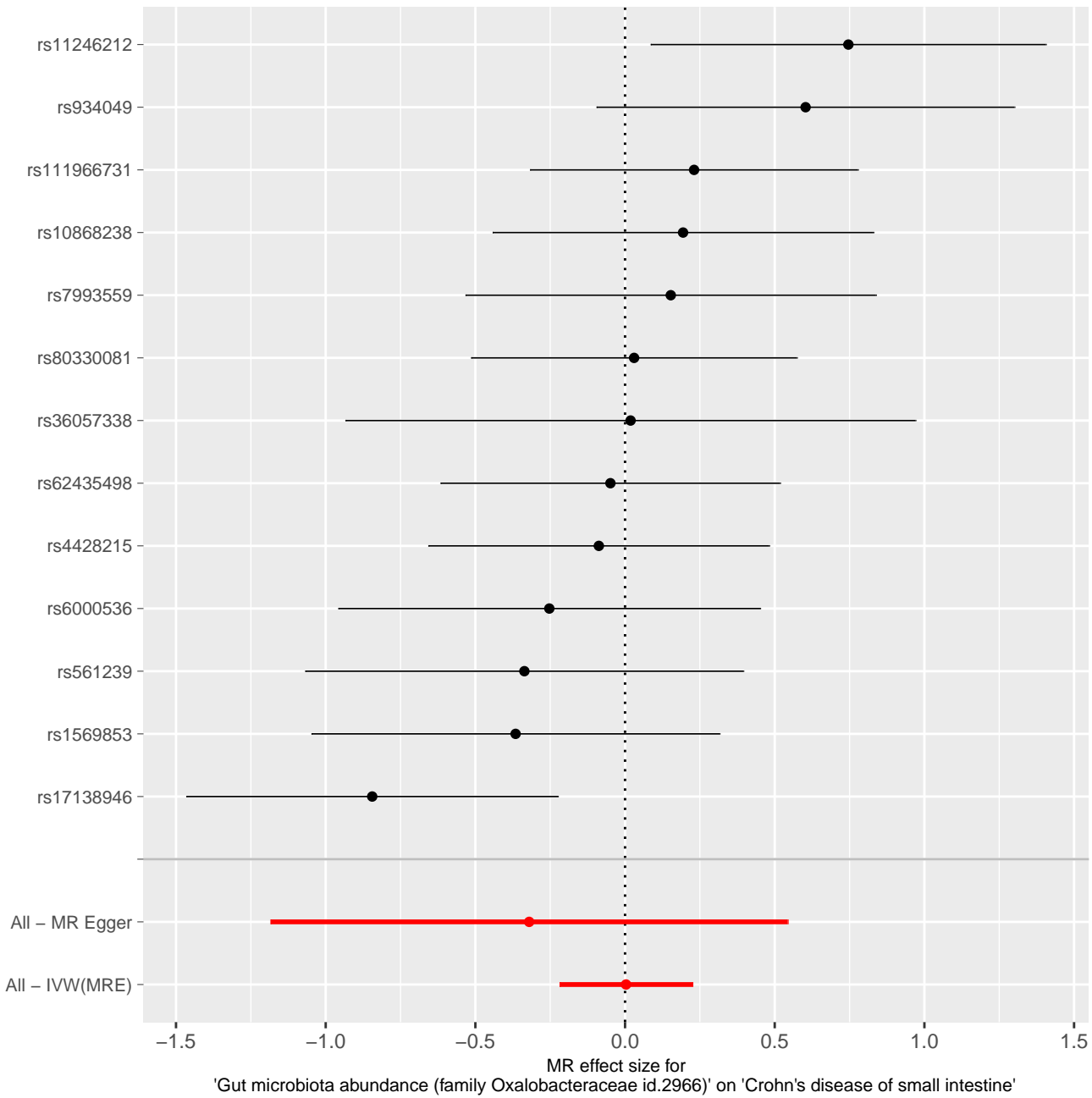


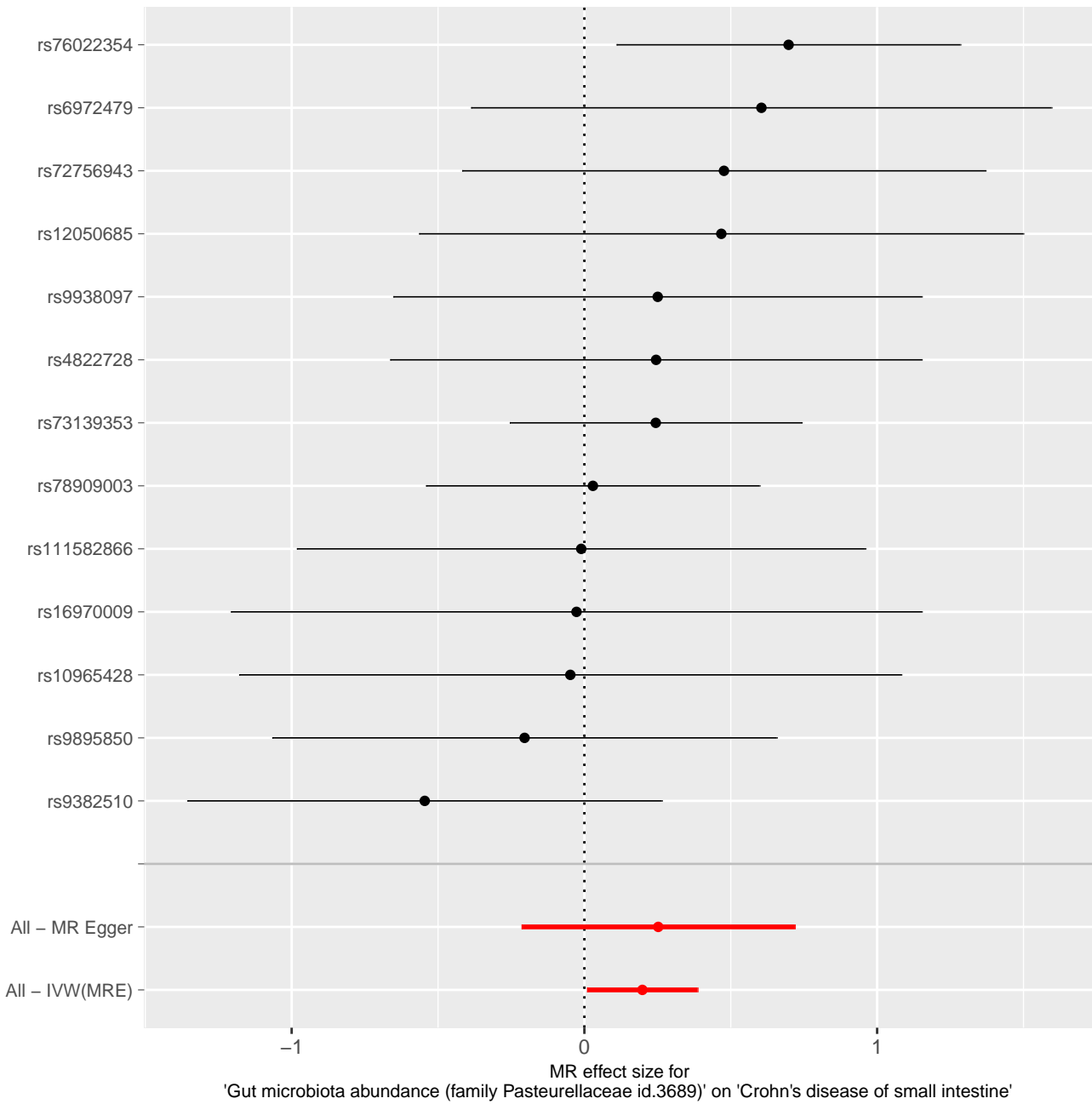




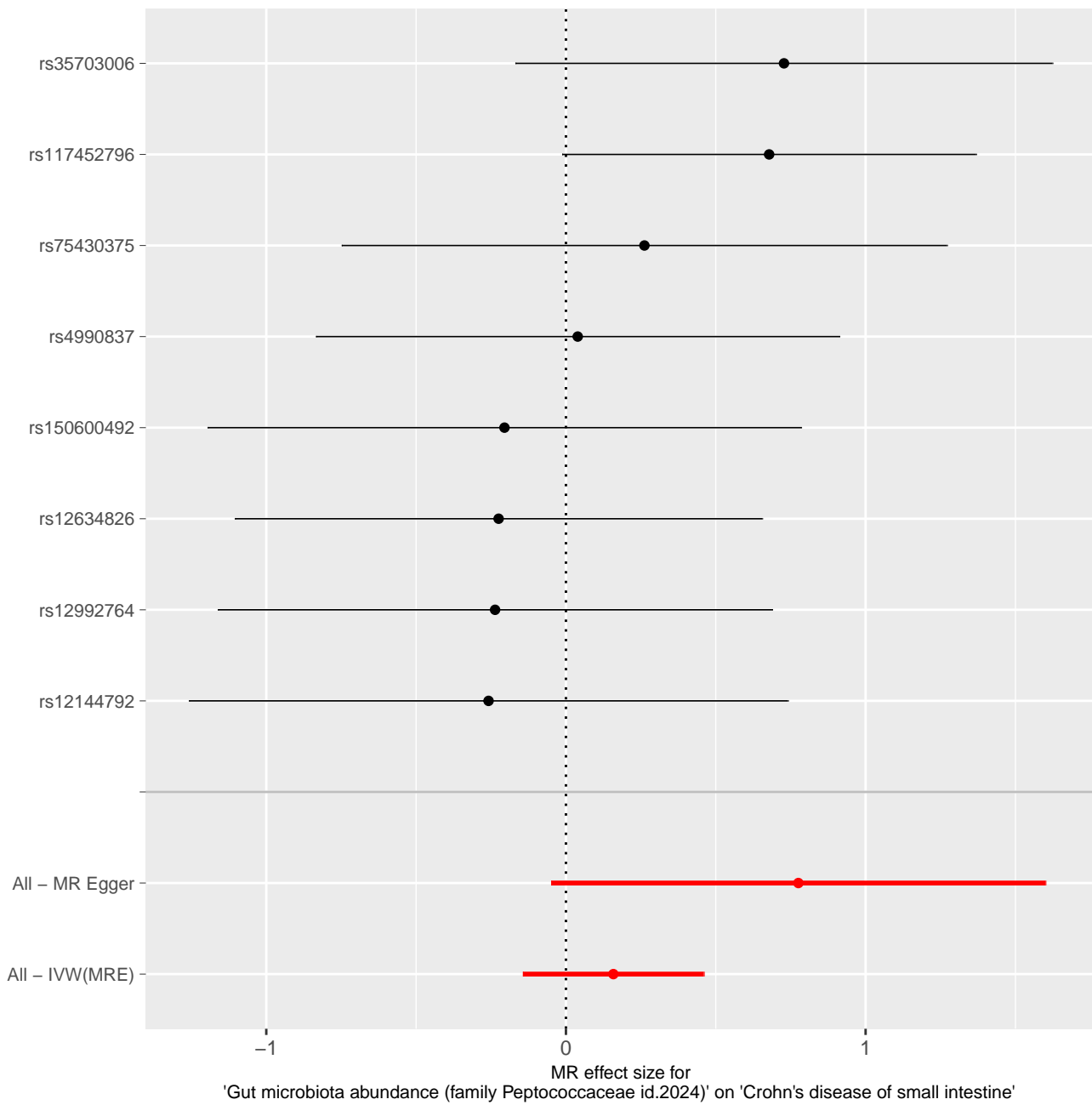


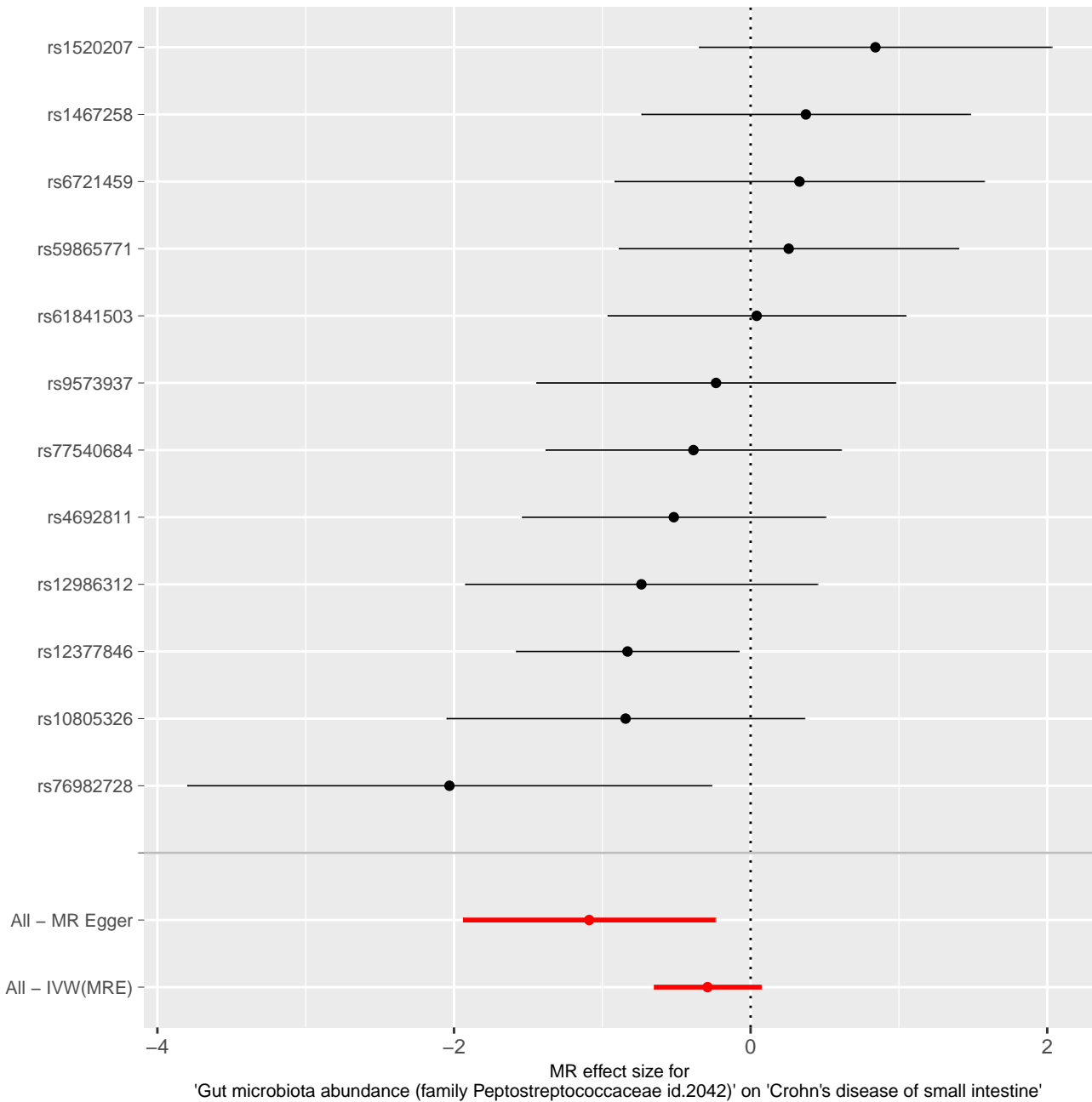


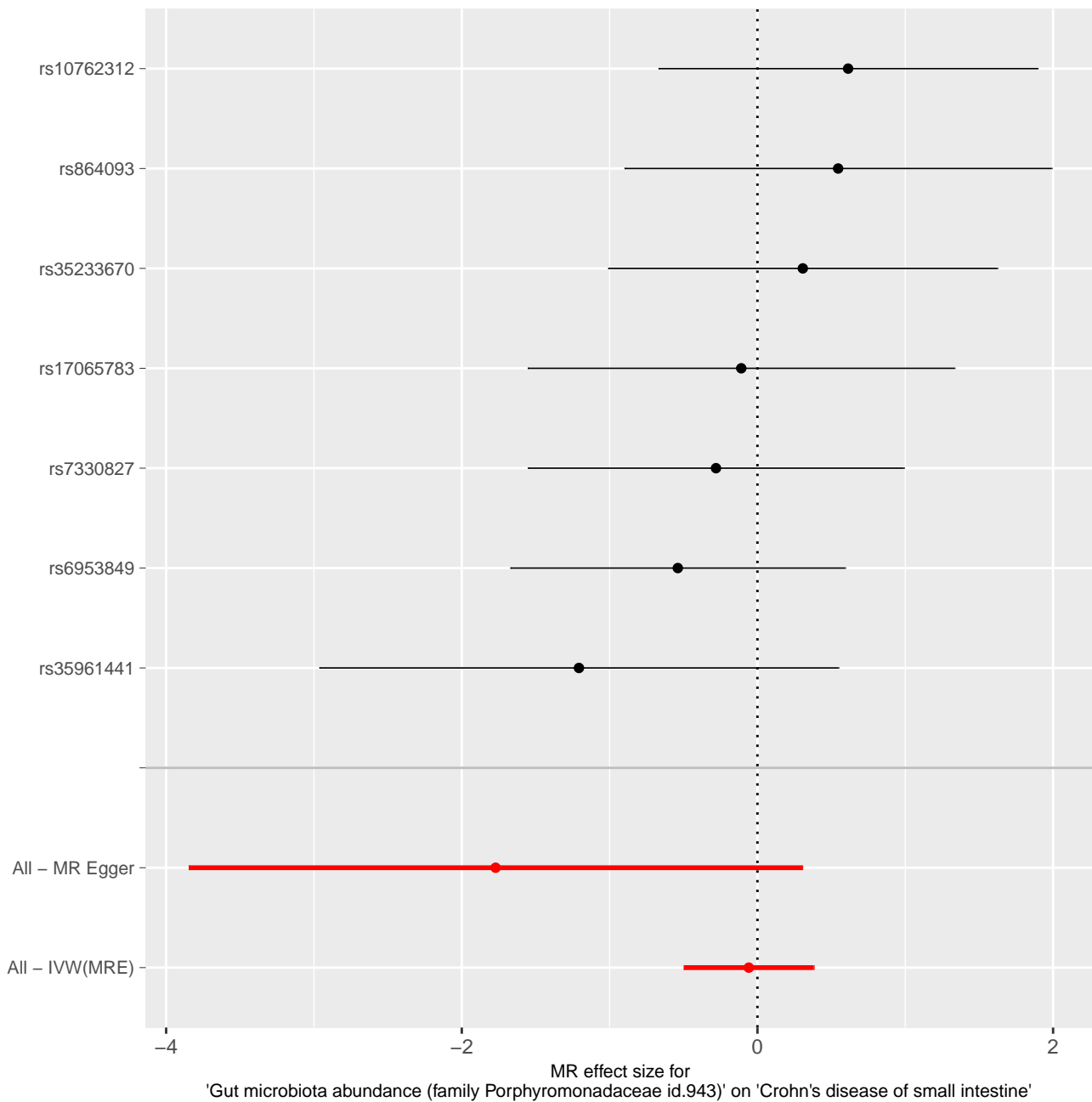




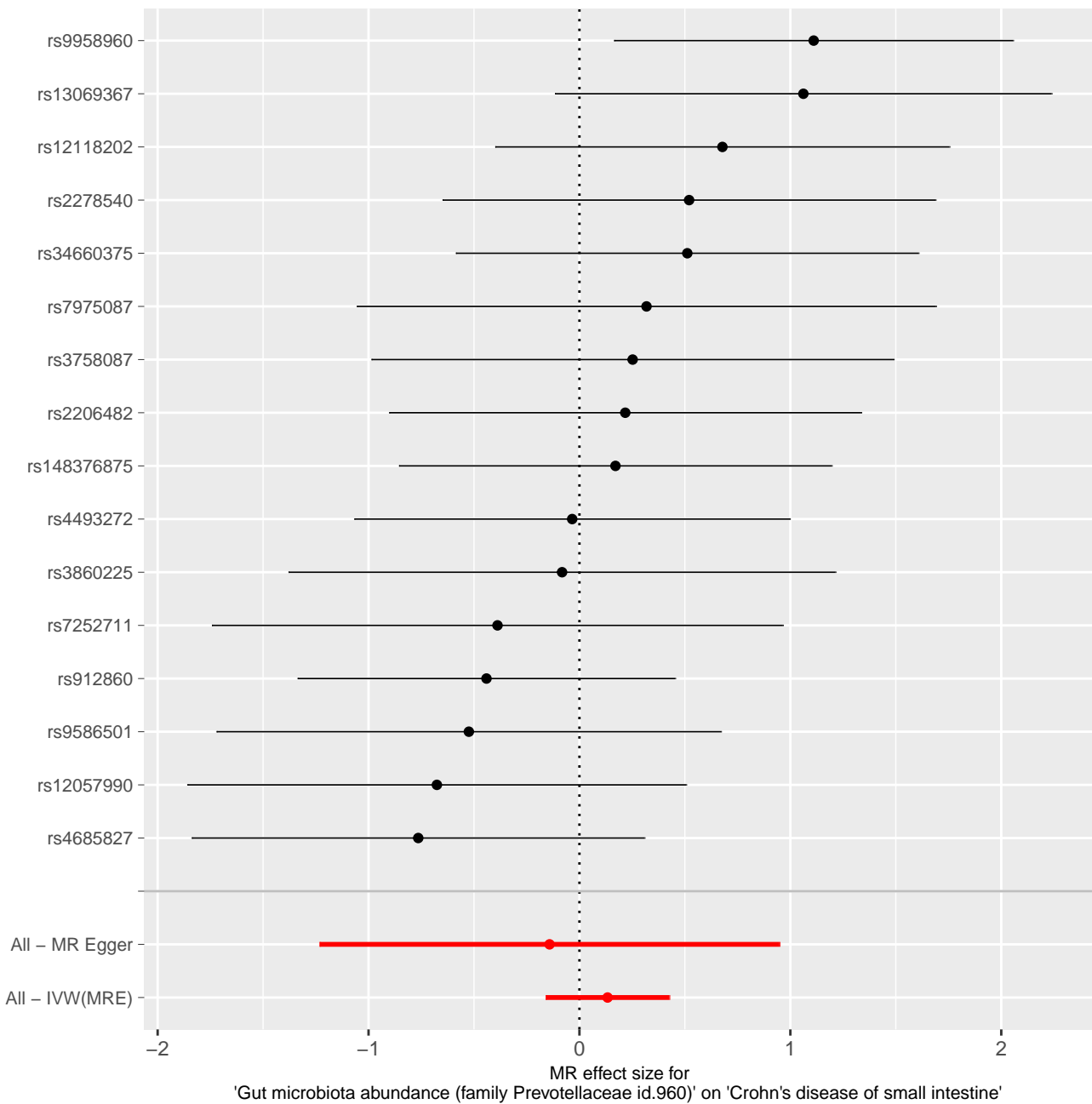
Batch 322 : Gut microbiota abundance (family Peptococcaceae id.2024) on Crohn's disease of small intestine

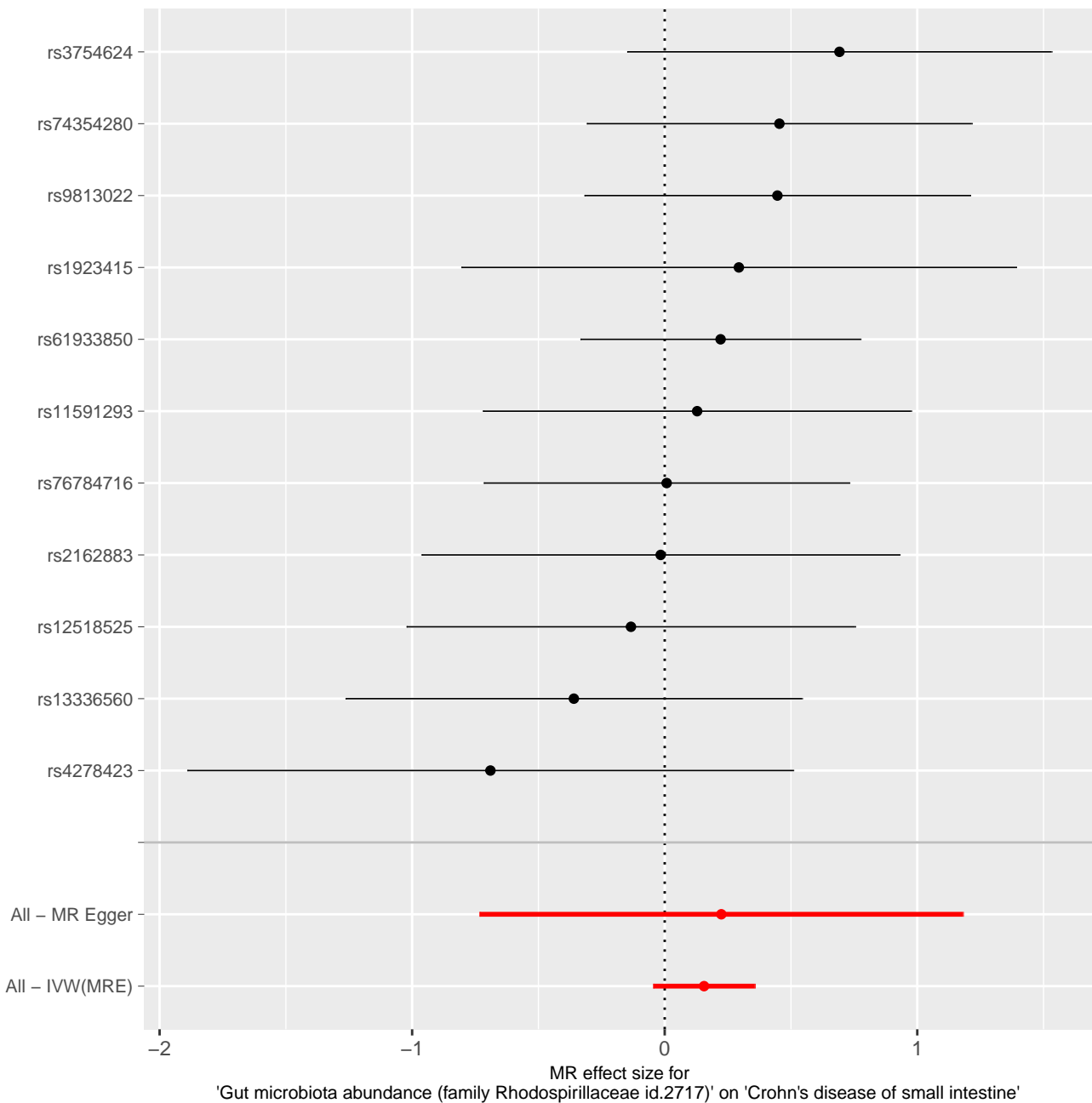




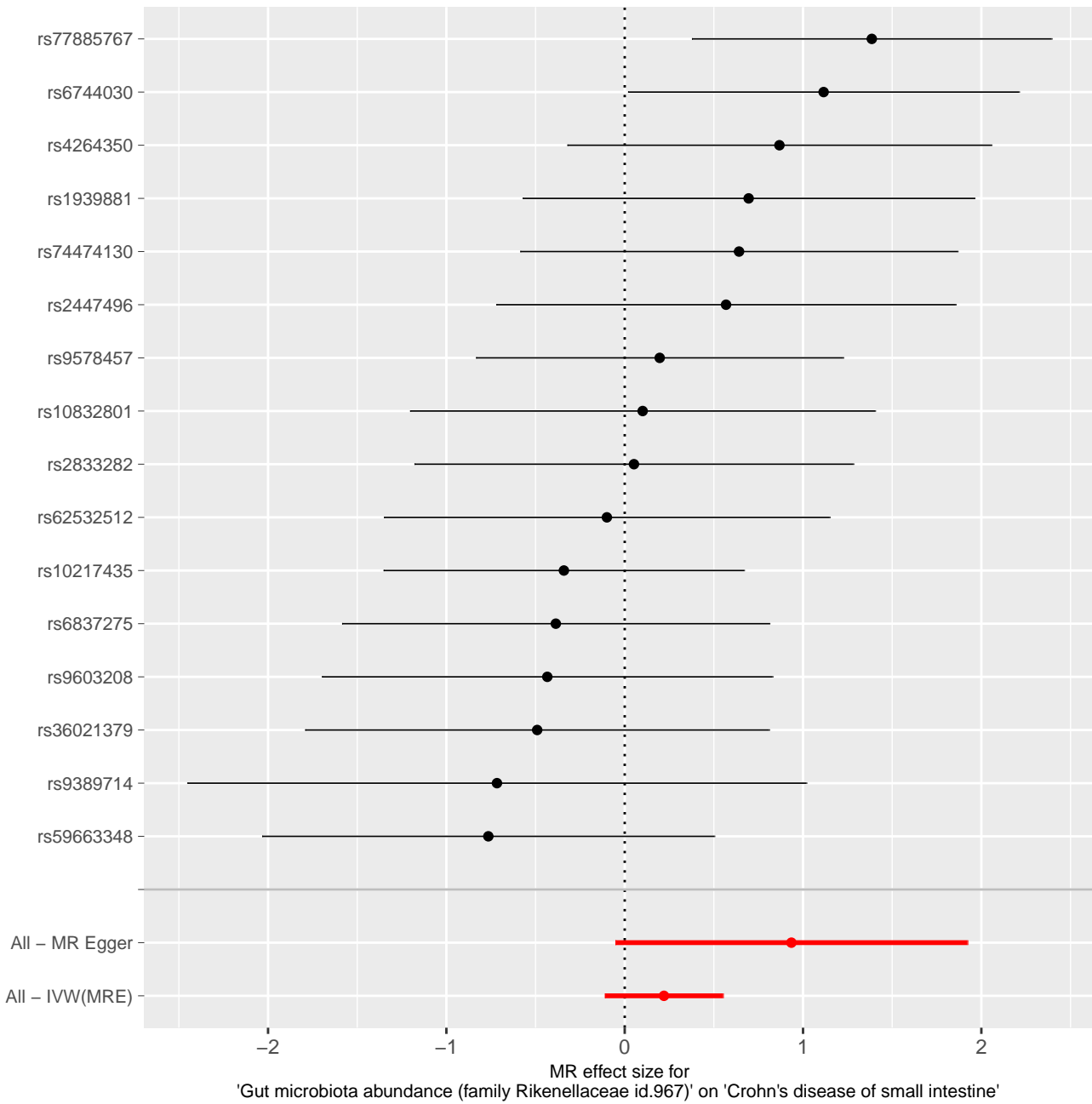


Batch 325 : Gut microbiota abundance (family Prevotellaceae id.960) on Crohn's disease of small intestine

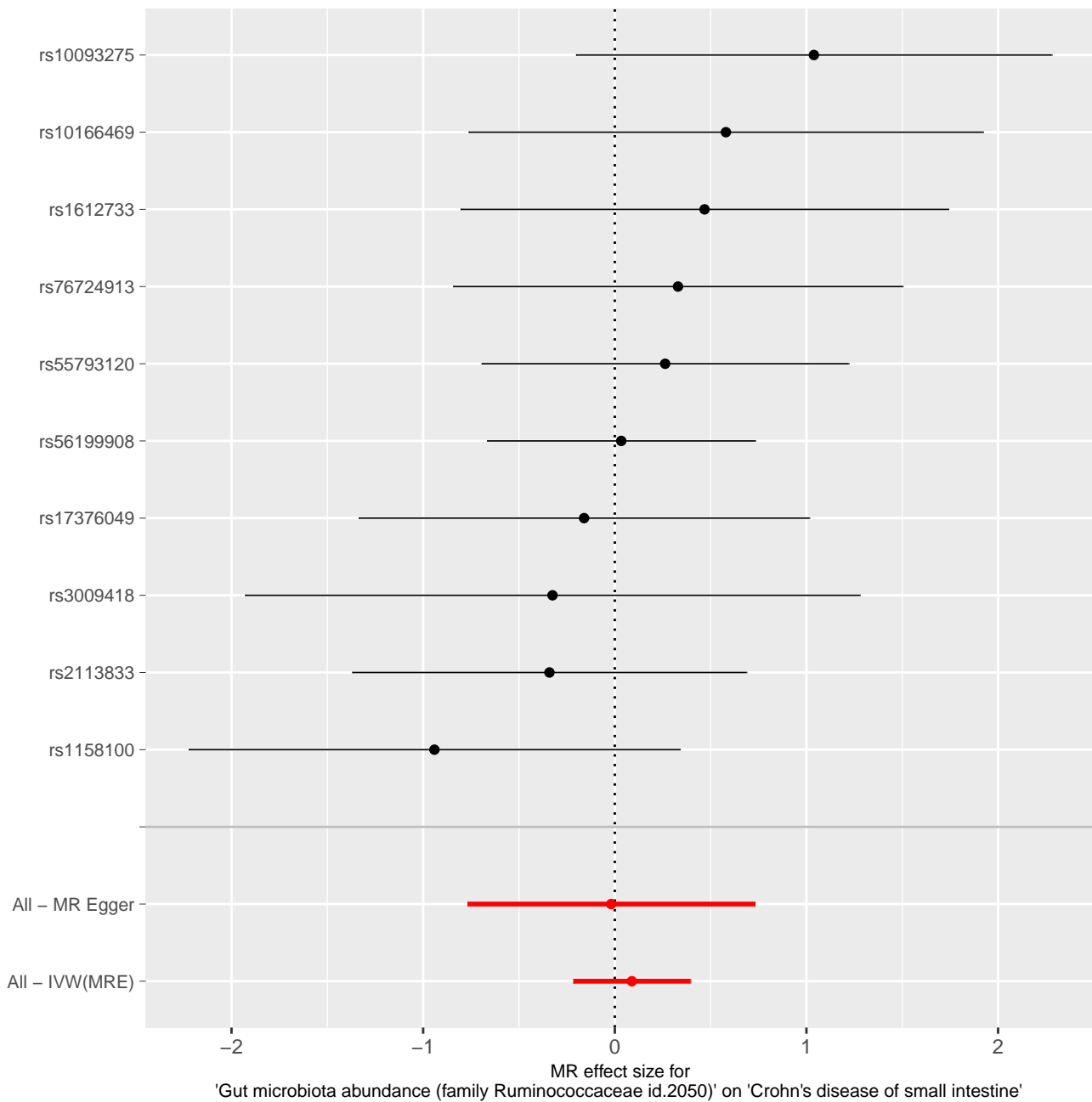




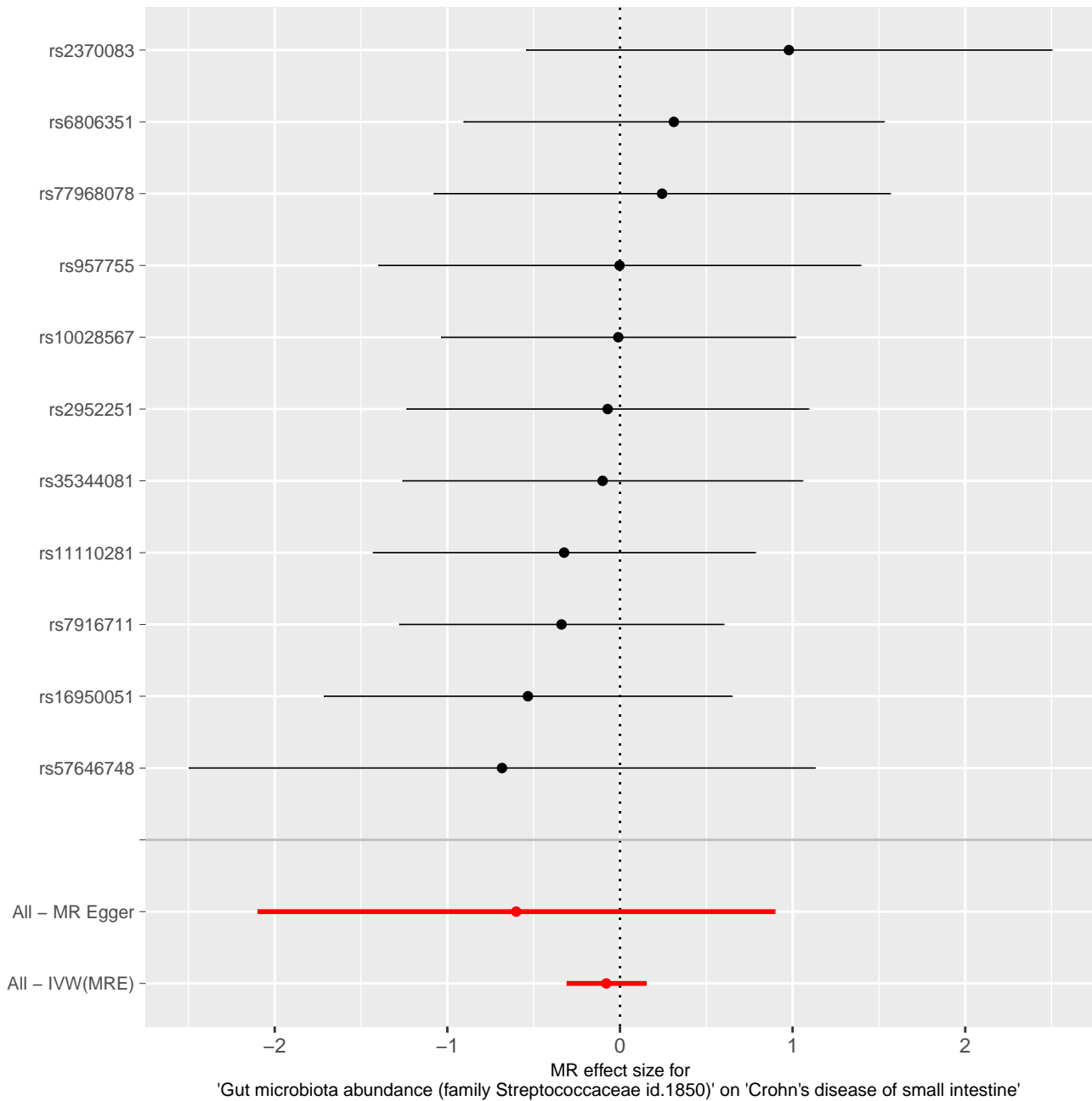
Batch 327 : Gut microbiota abundance (family Rikenellaceae id.967) on Crohn's disease of small intestine



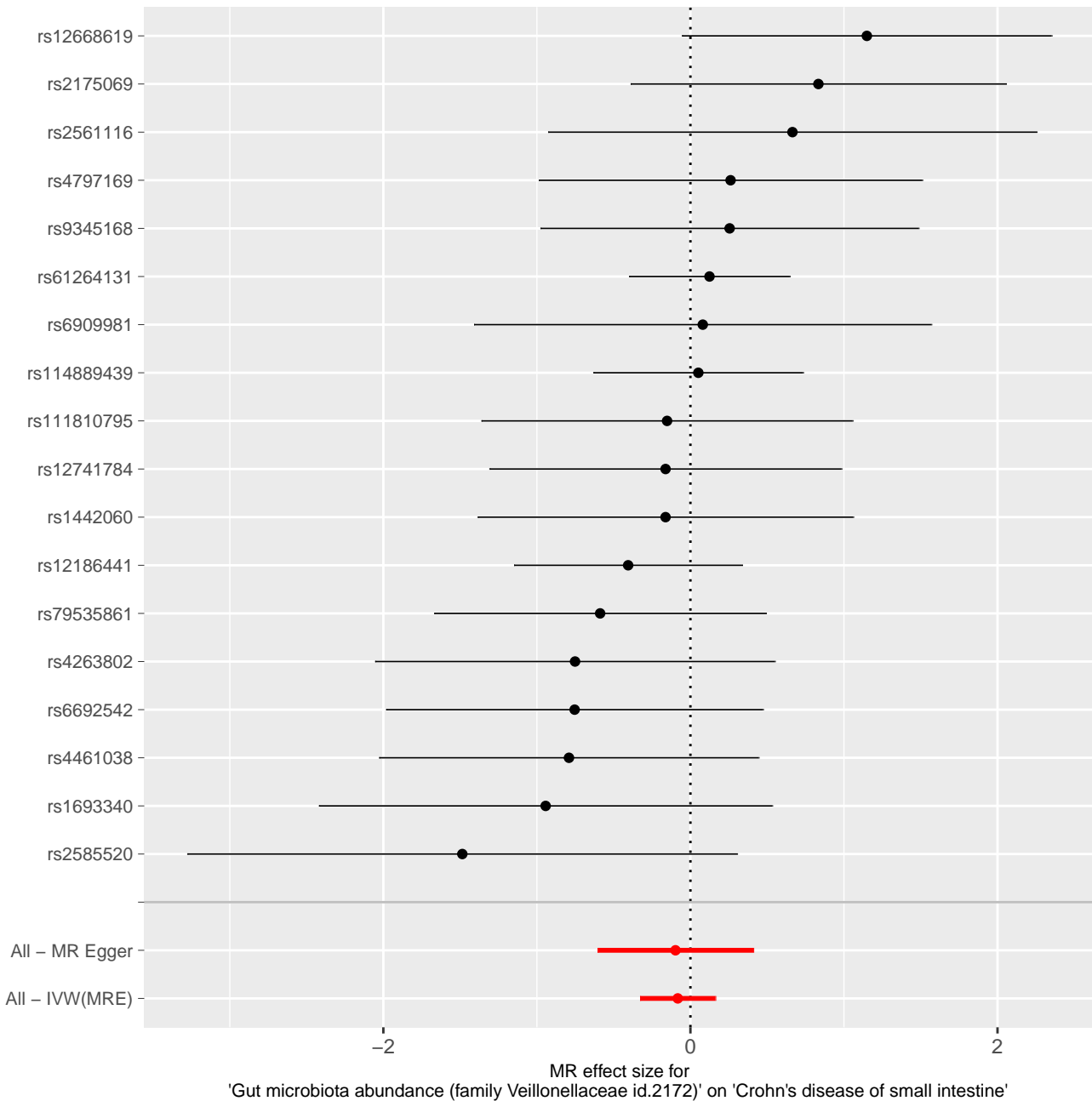
Batch 328 : Gut microbiota abundance (family Ruminococcaceae id.2050) on Crohn's disease of small intestine

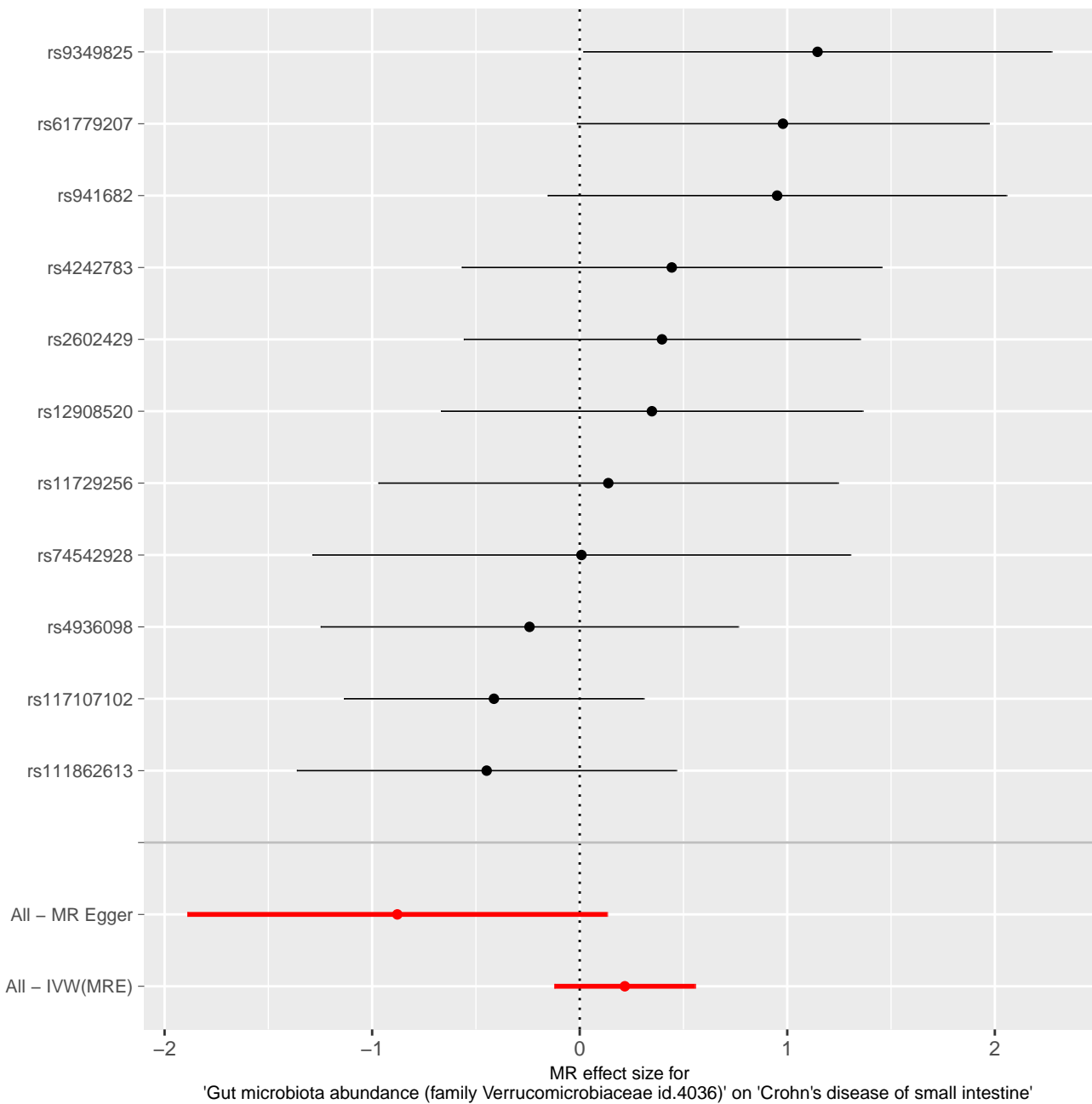


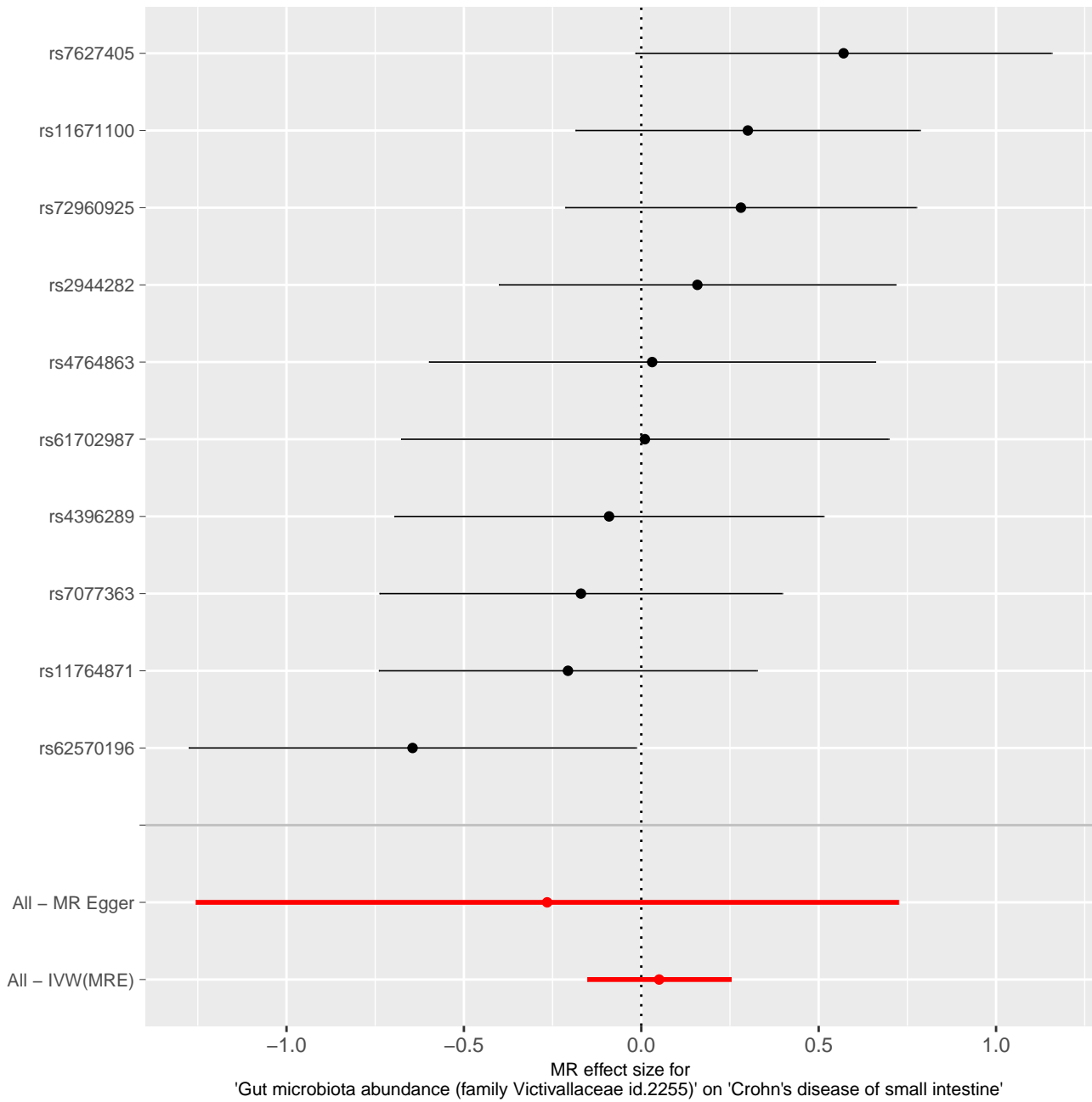
Batch 329 : Gut microbiota abundance (family Streptococcaceae id.1850) on Crohn's disease of small intestine

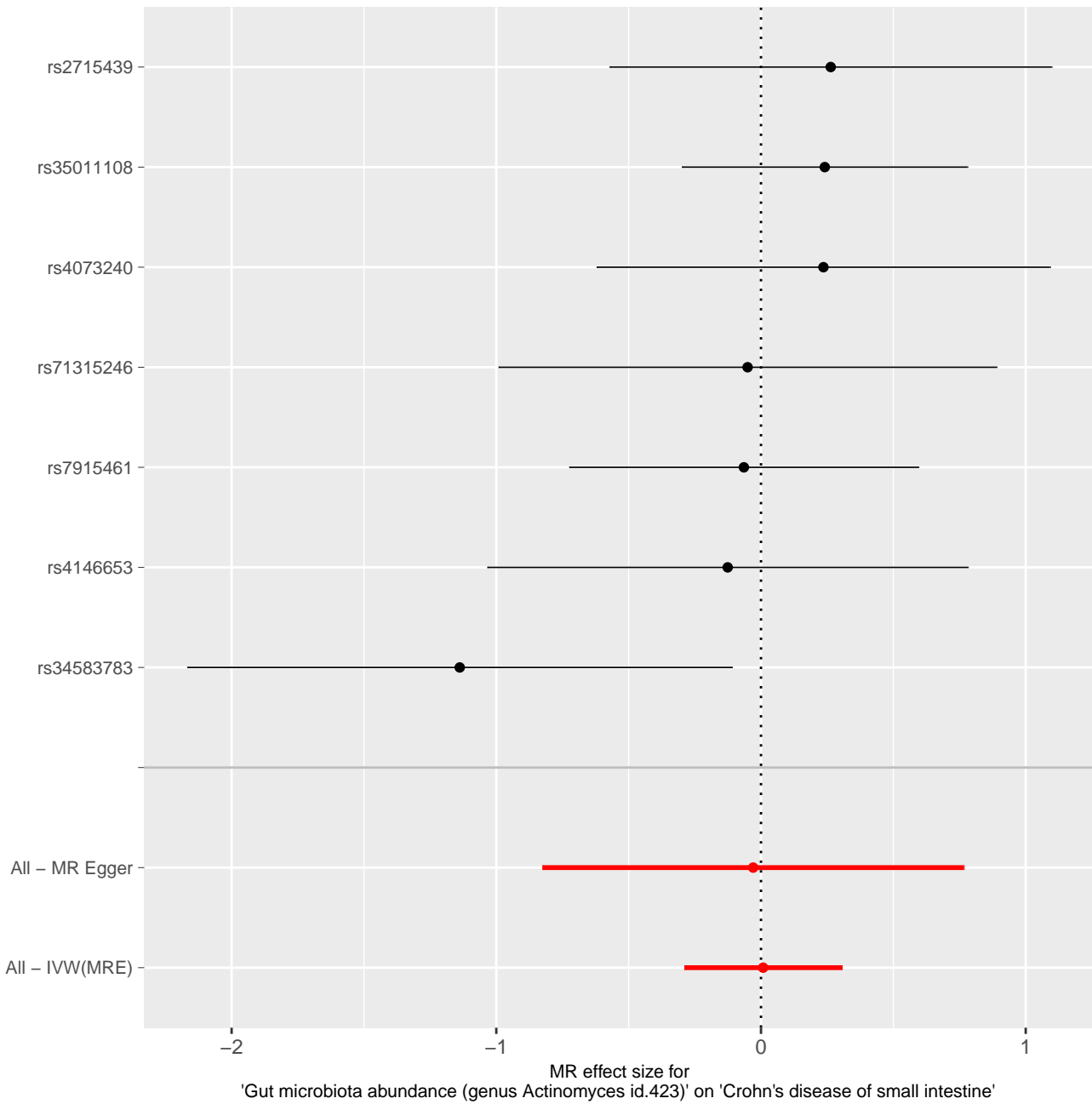


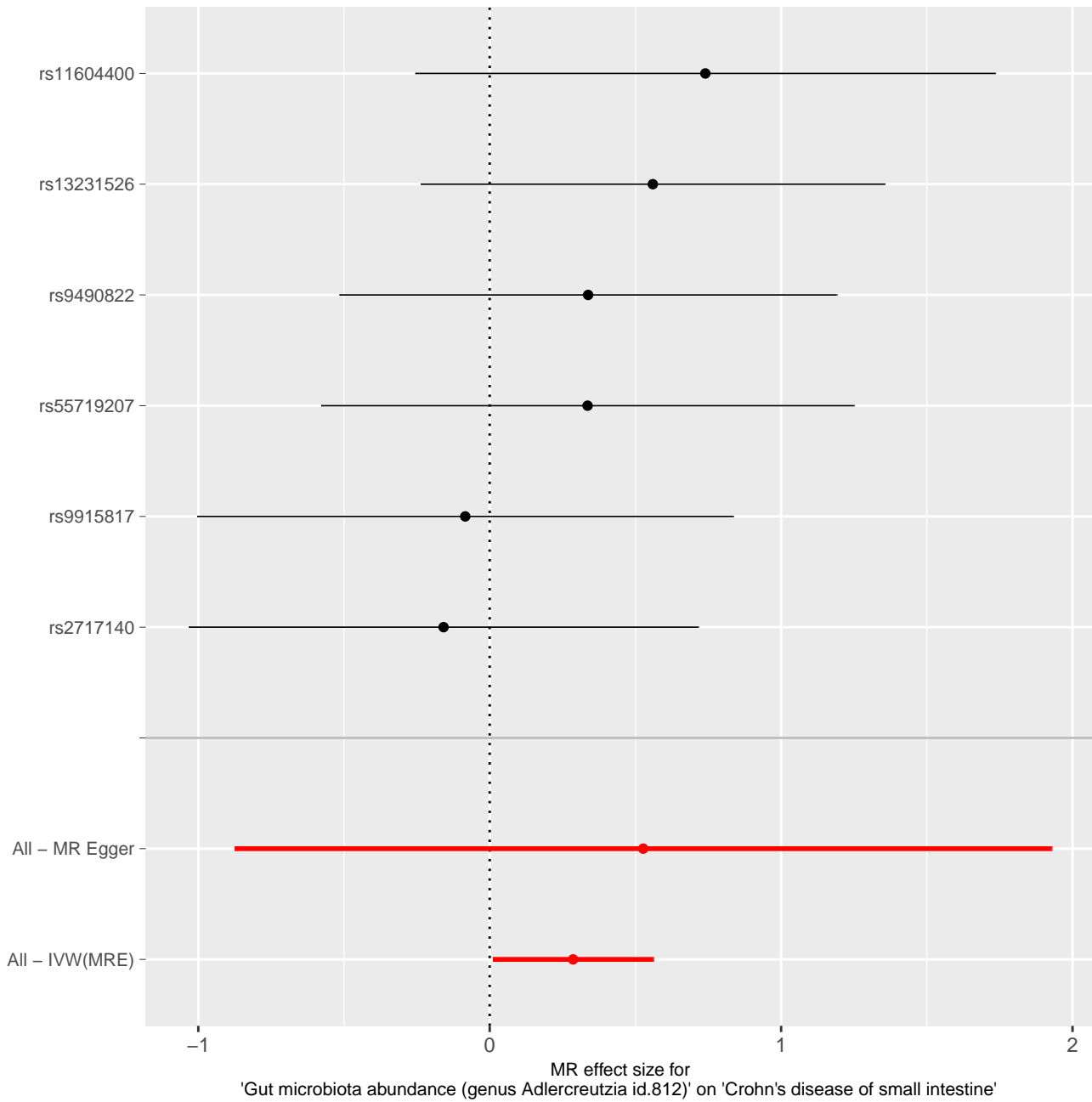
Batch 330 : Gut microbiota abundance (family Veillonellaceae id.2172) on Crohn's disease of small intestine

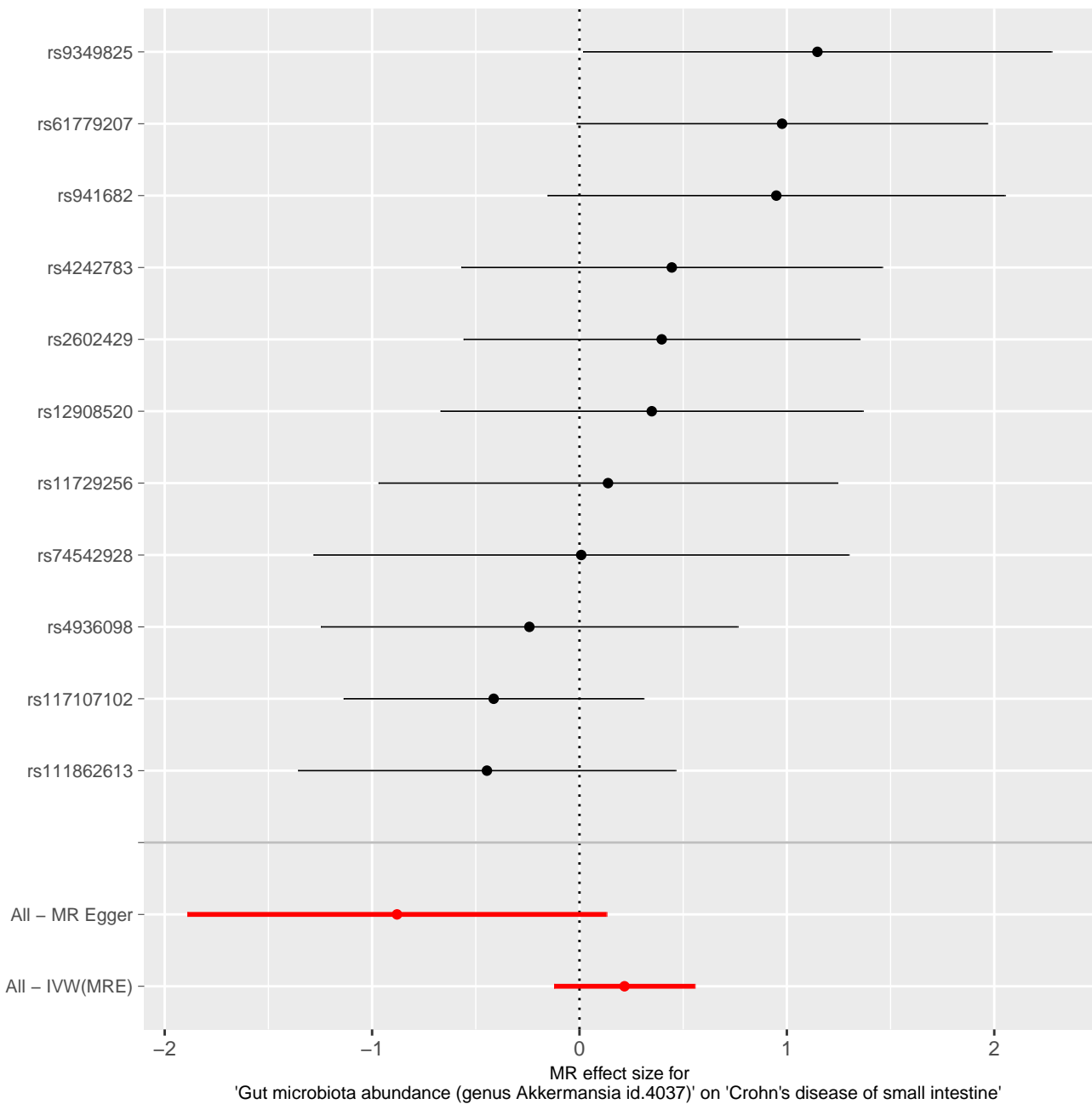




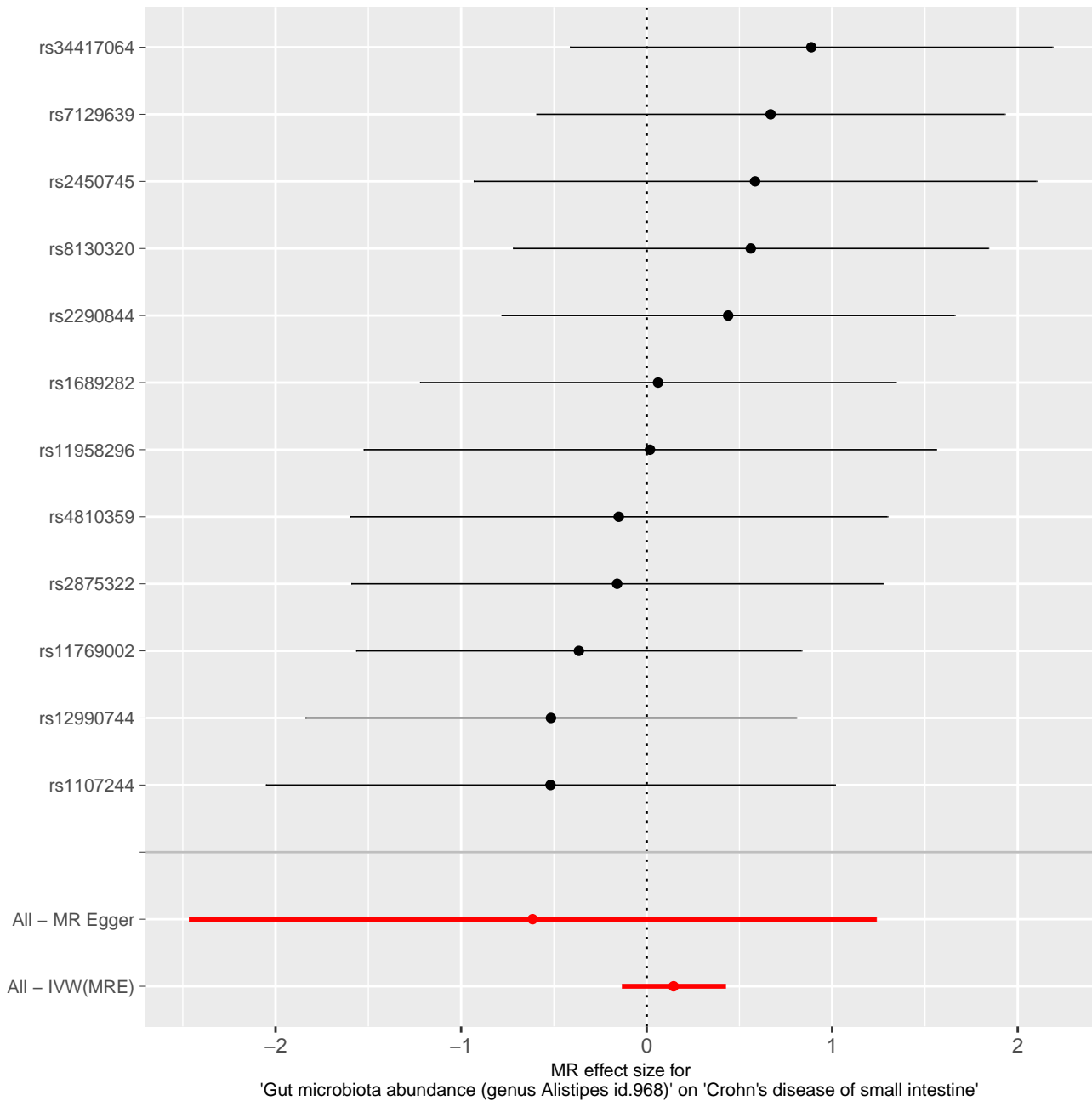


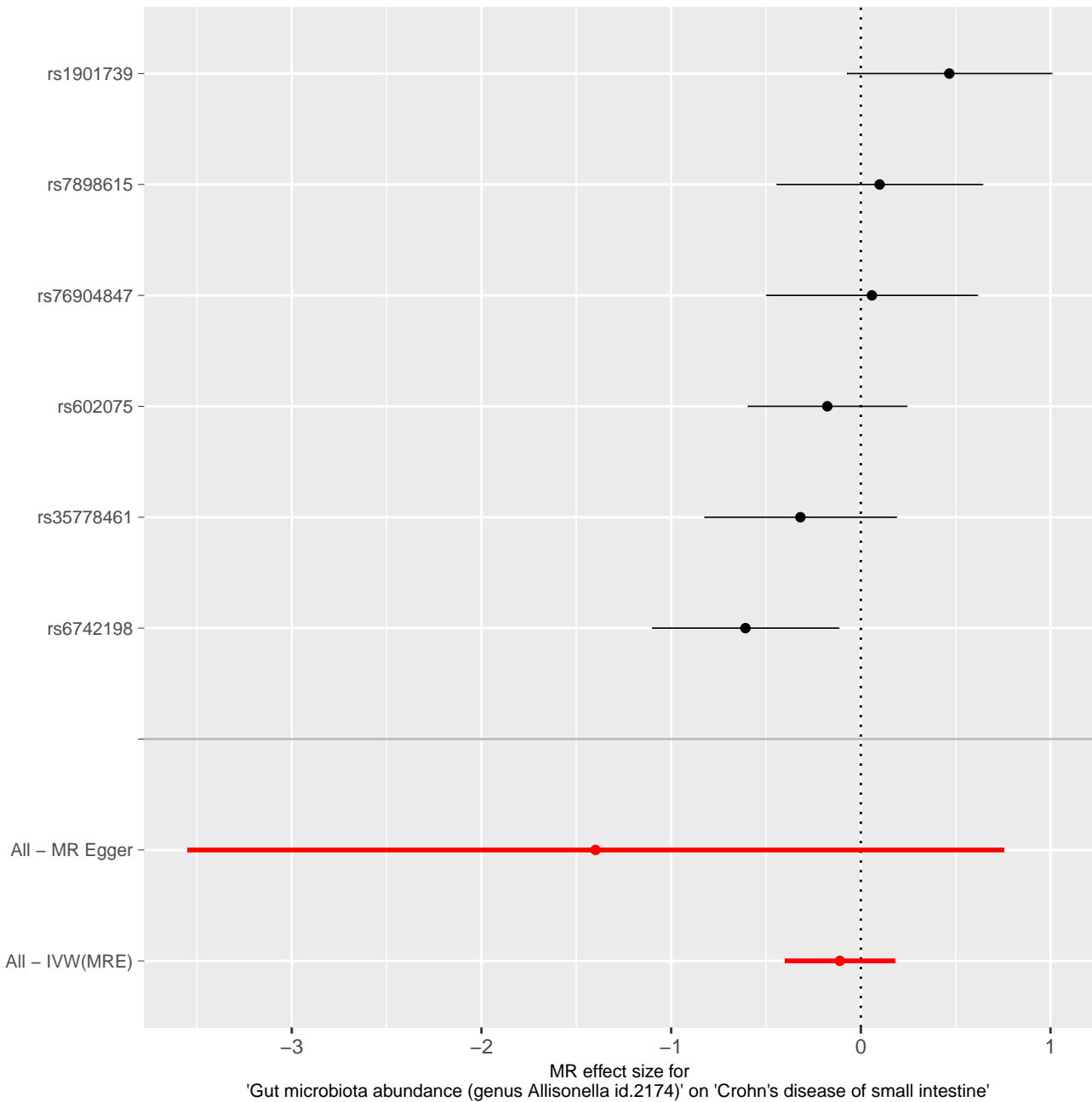


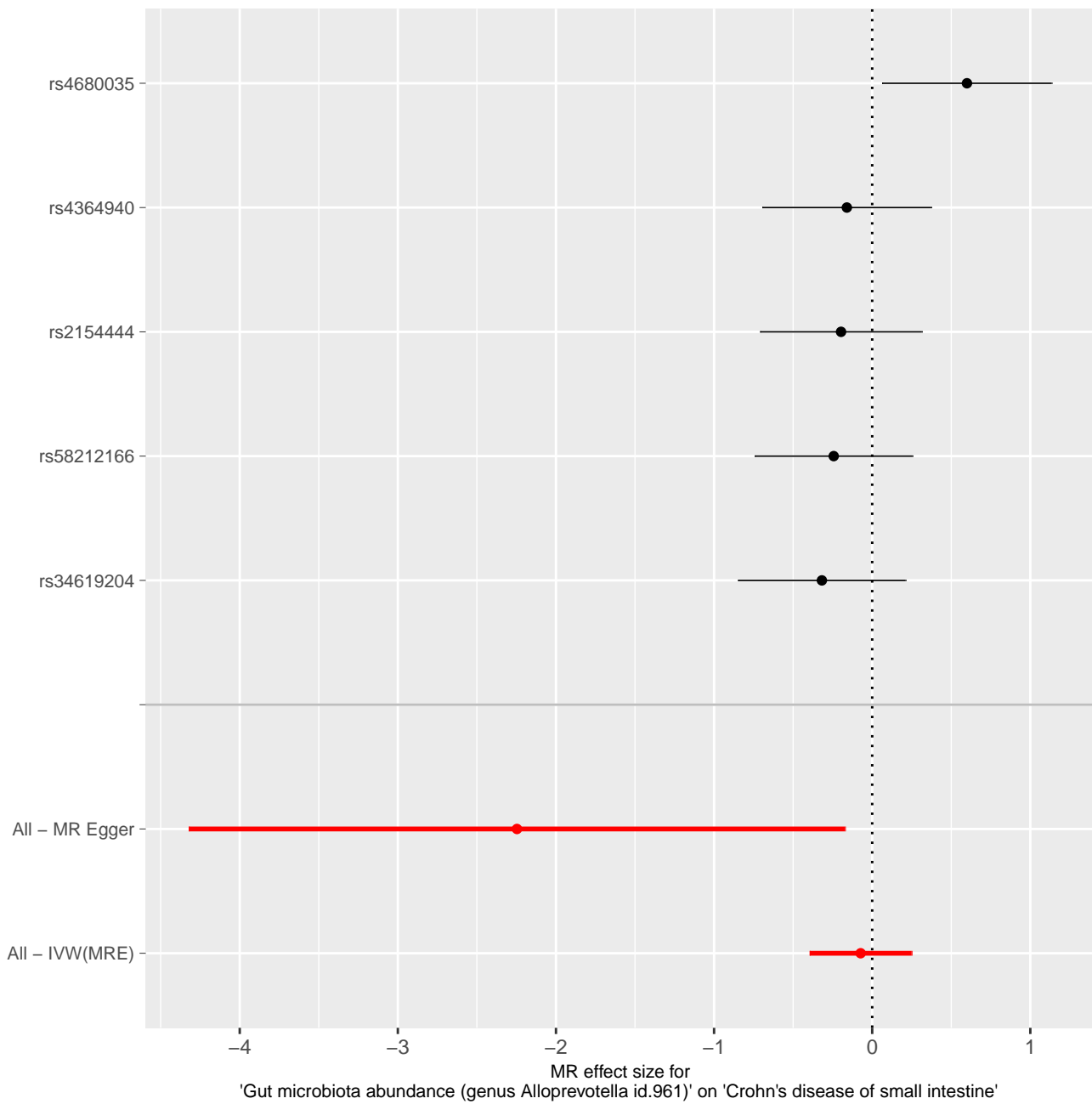




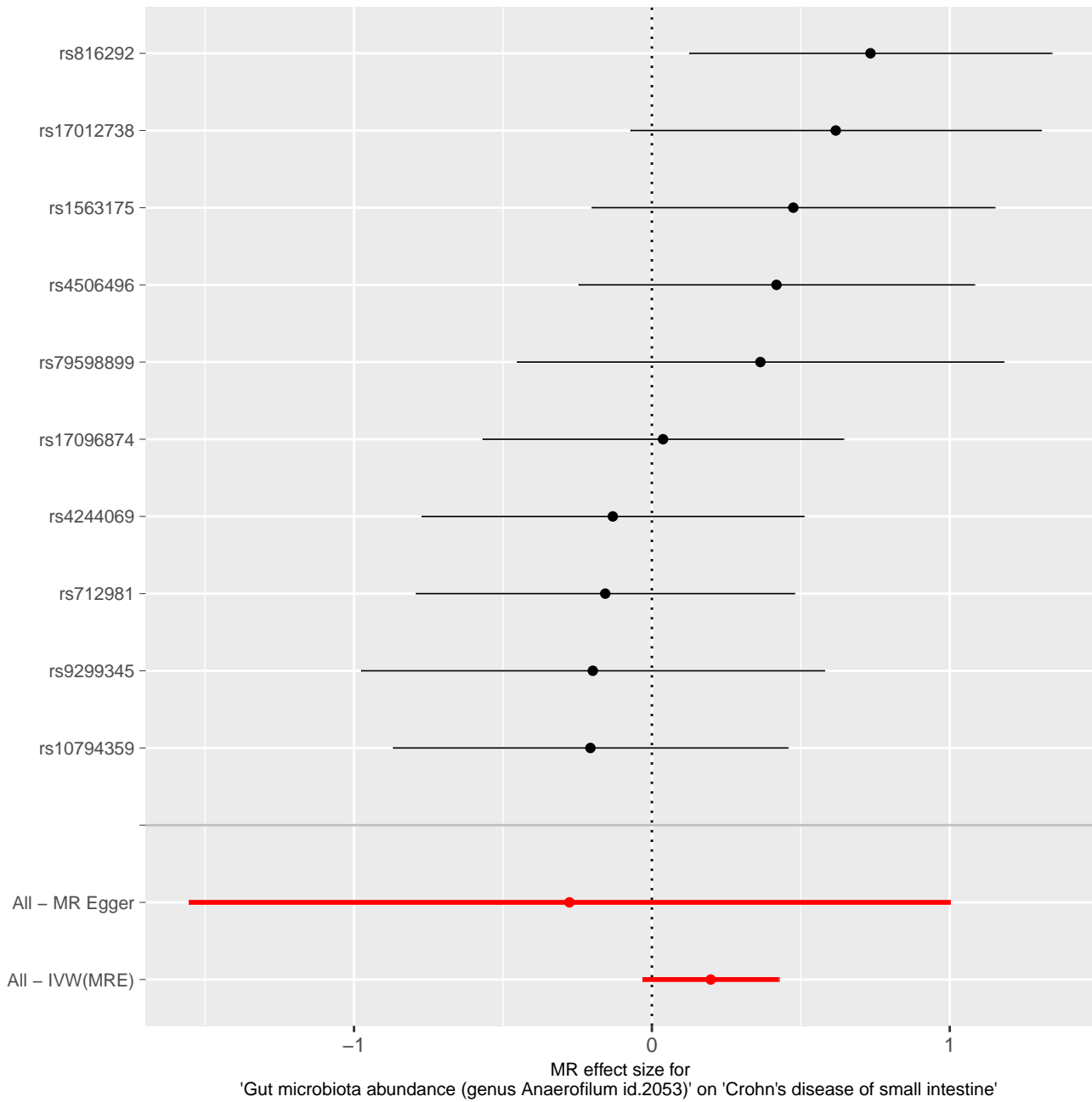
Batch 336 : Gut microbiota abundance (genus Alistipes id.968) on Crohn's disease of small intestine



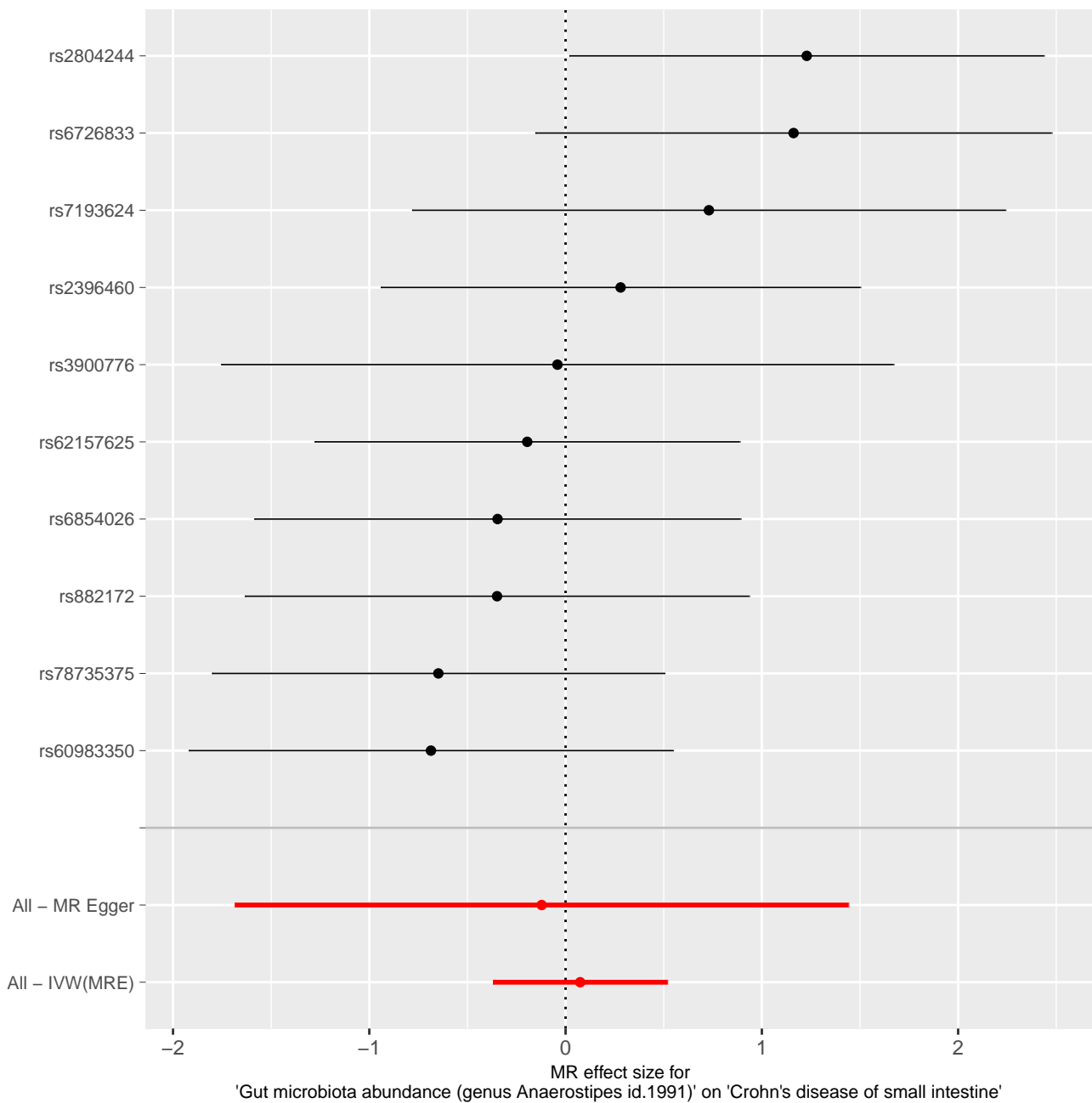




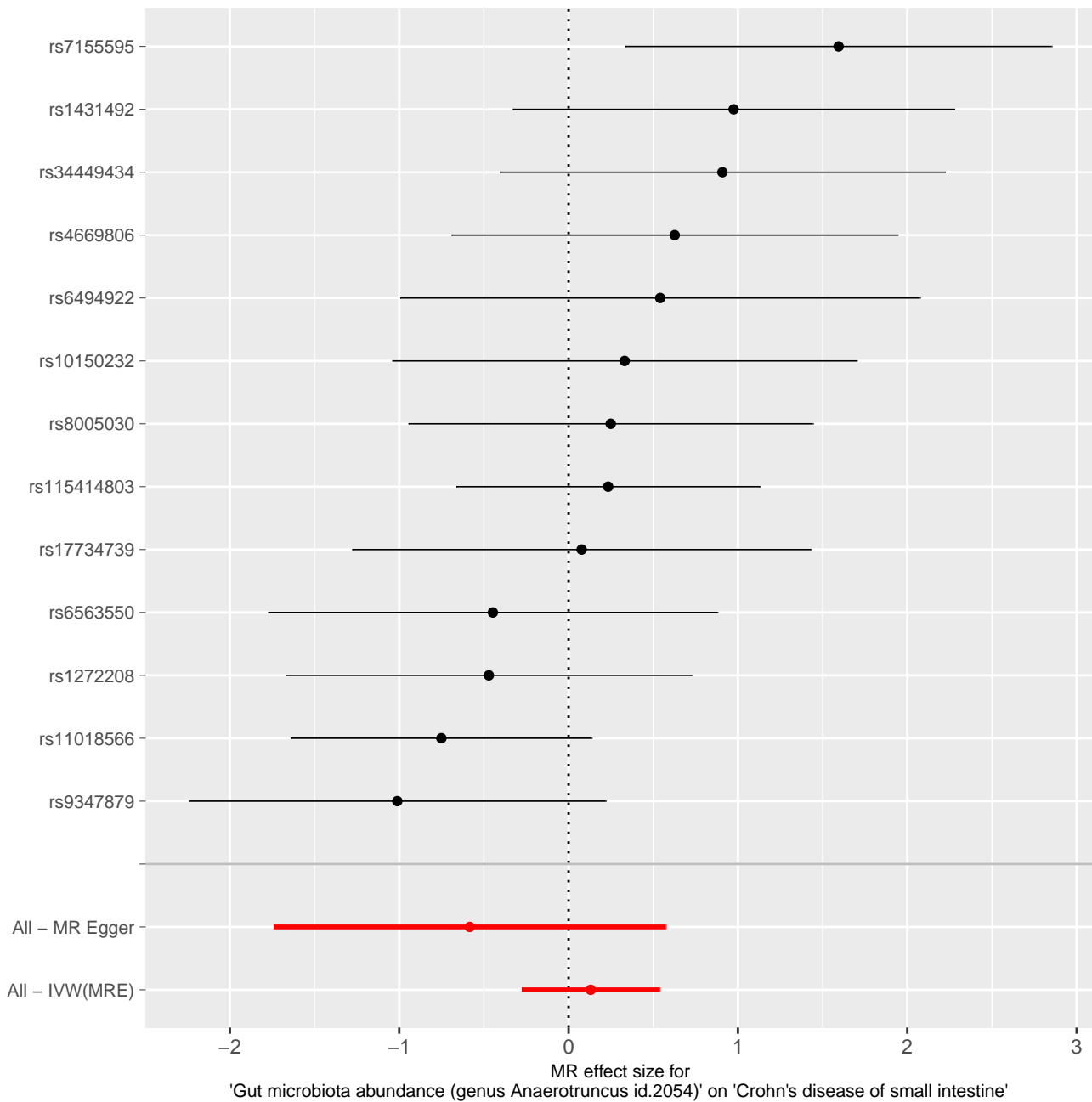
Batch 339 : Gut microbiota abundance (genus Anaerofilum id.2053) on Crohn's disease of small intestine



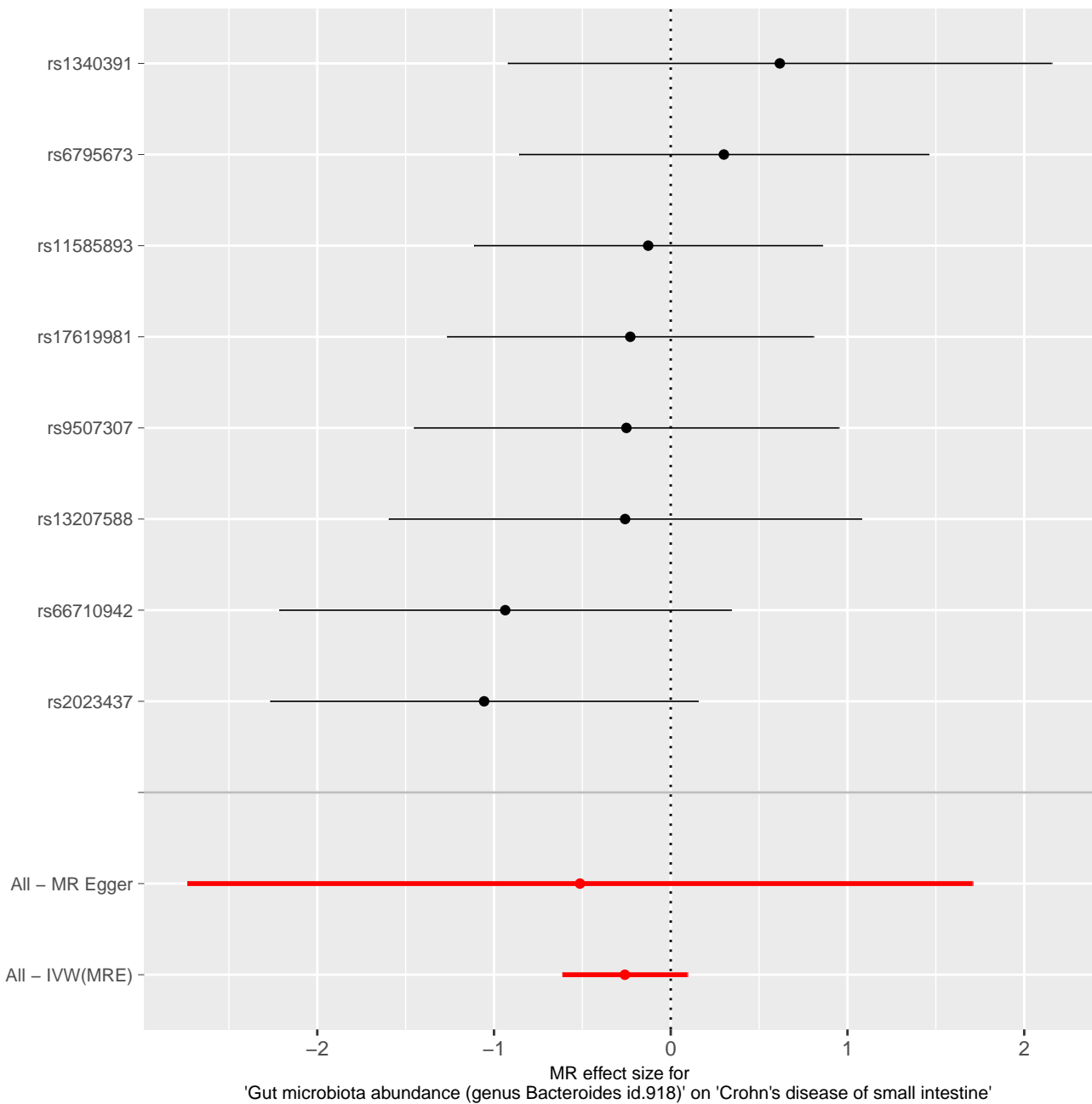
Batch 340 : Gut microbiota abundance (genus Anaerostipes id.1991) on Crohn's disease of small intestine



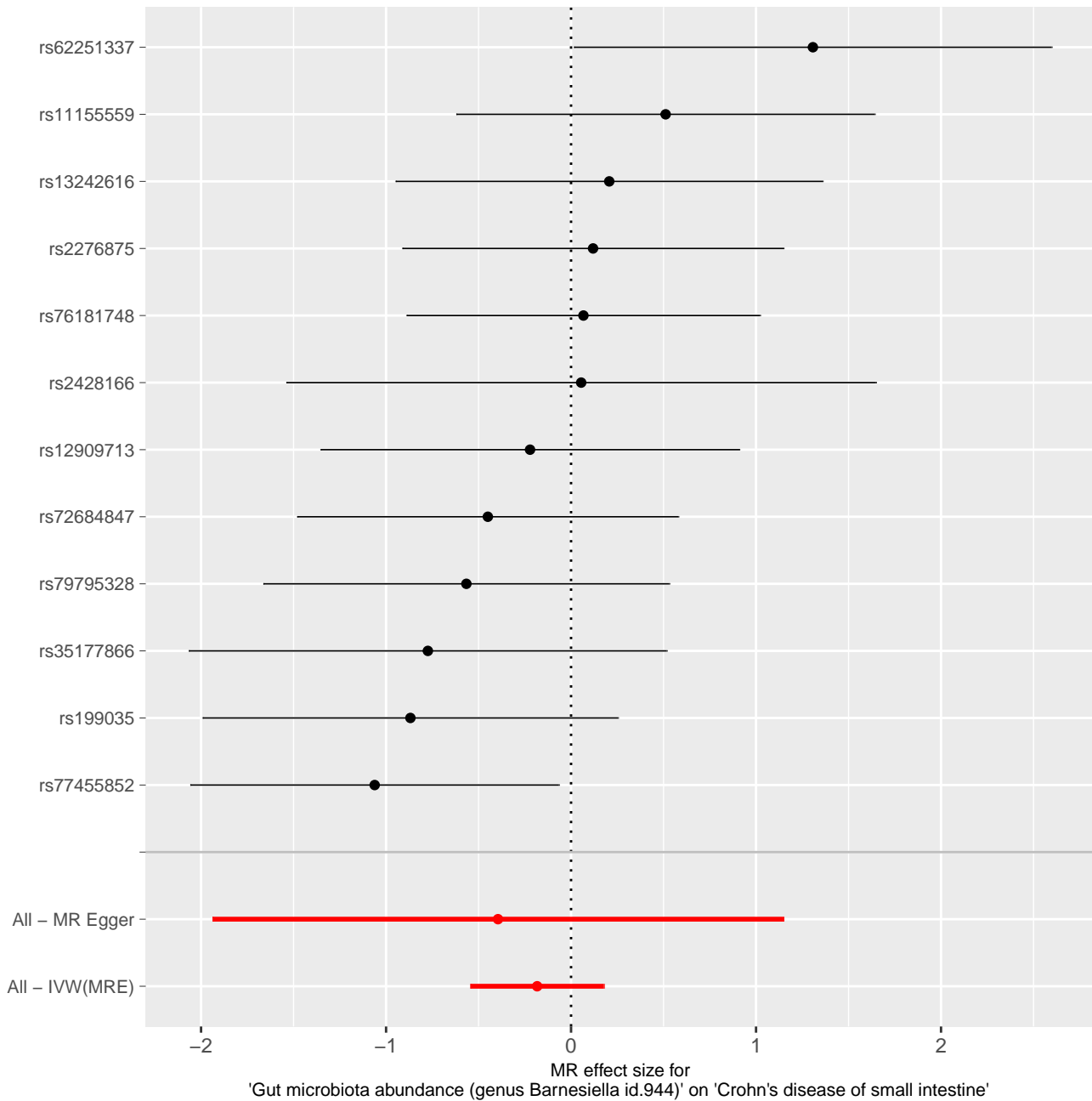
Batch 341 : Gut microbiota abundance (genus Anaerotruncus id.2054) on Crohn's disease of small intestine

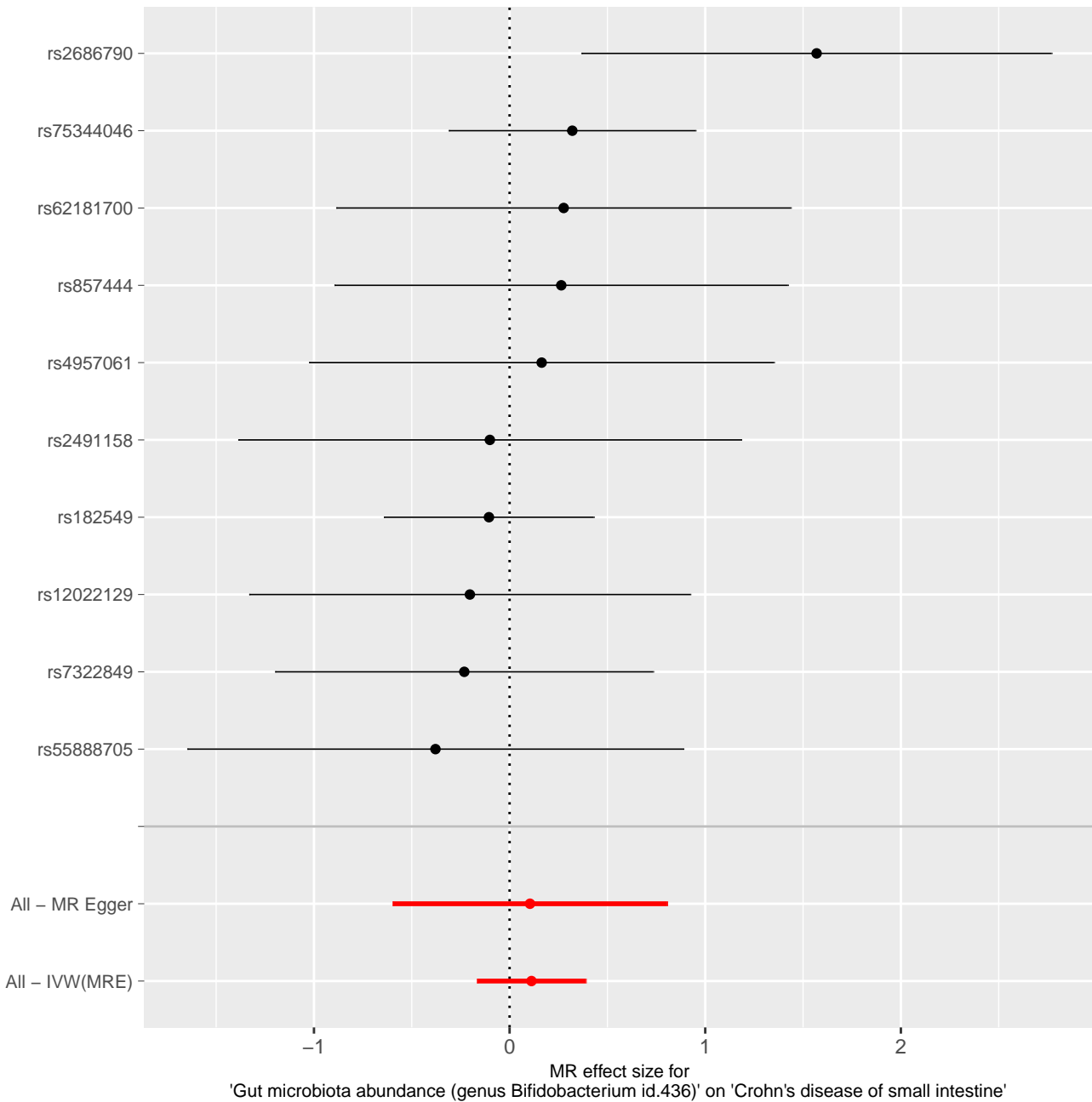


Batch 342 : Gut microbiota abundance (genus Bacteroides id.918) on Crohn's disease of small intestine

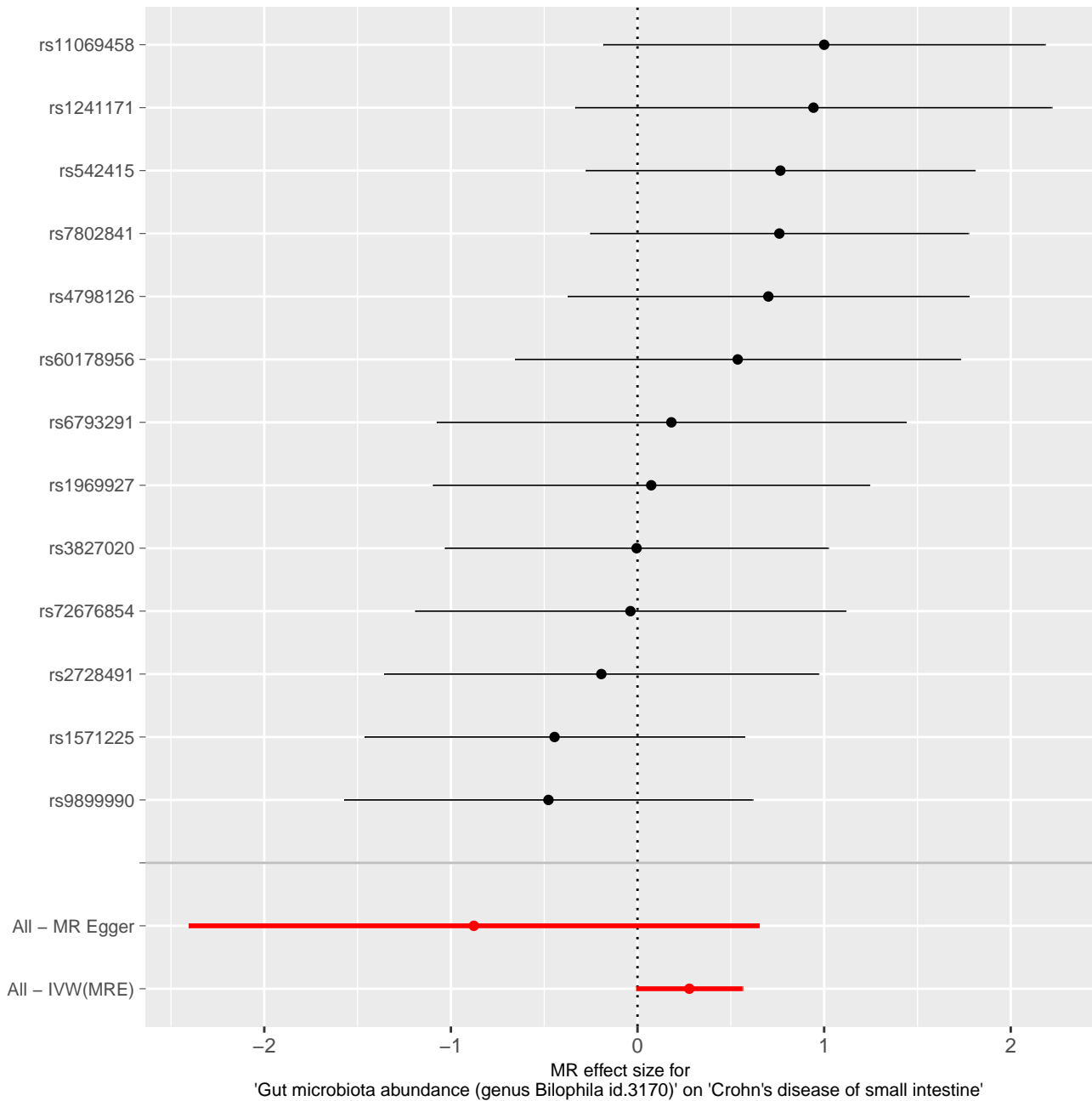


Batch 343 : Gut microbiota abundance (genus Barnesiella id.944) on Crohn's disease of small intestine

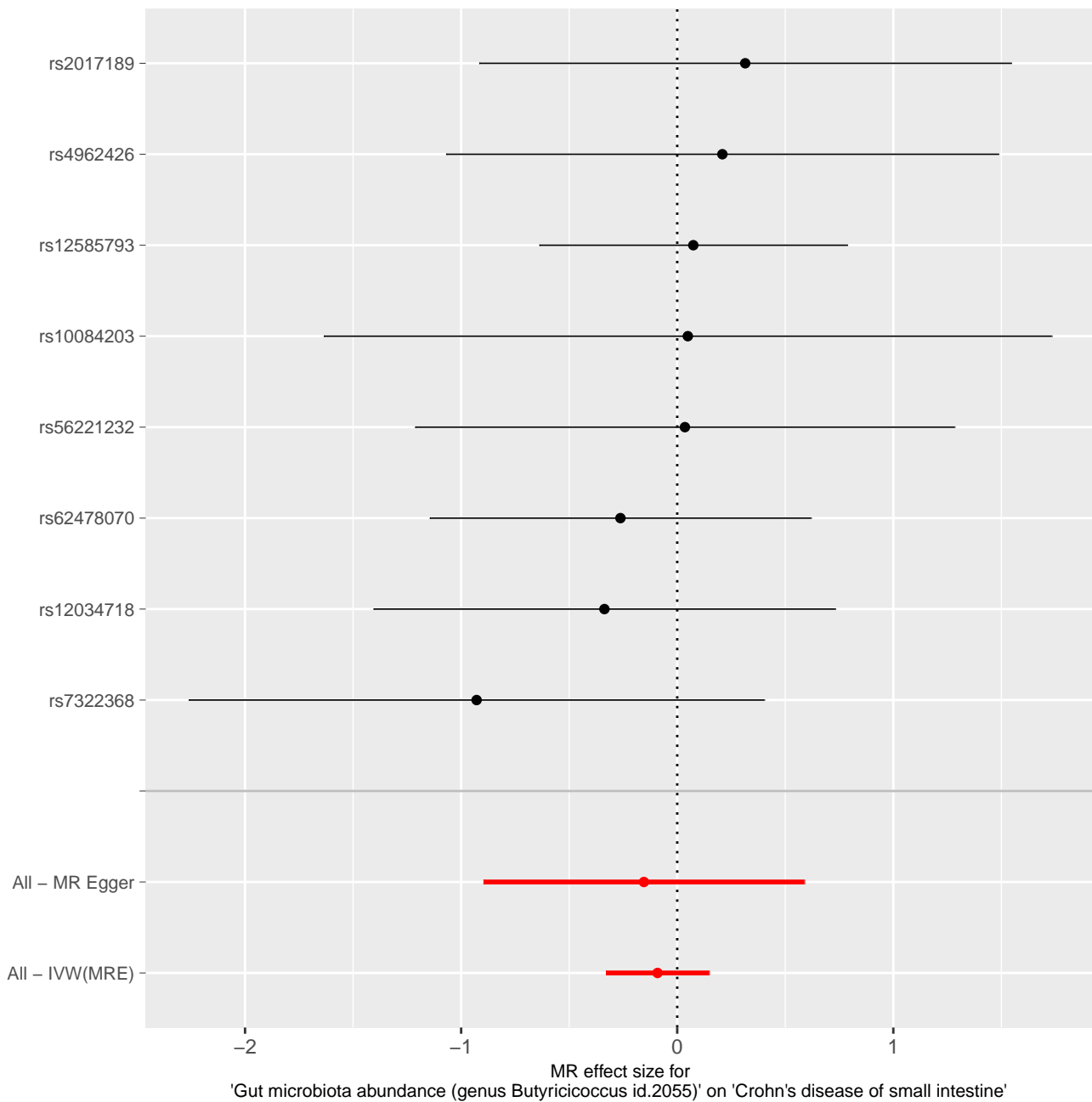


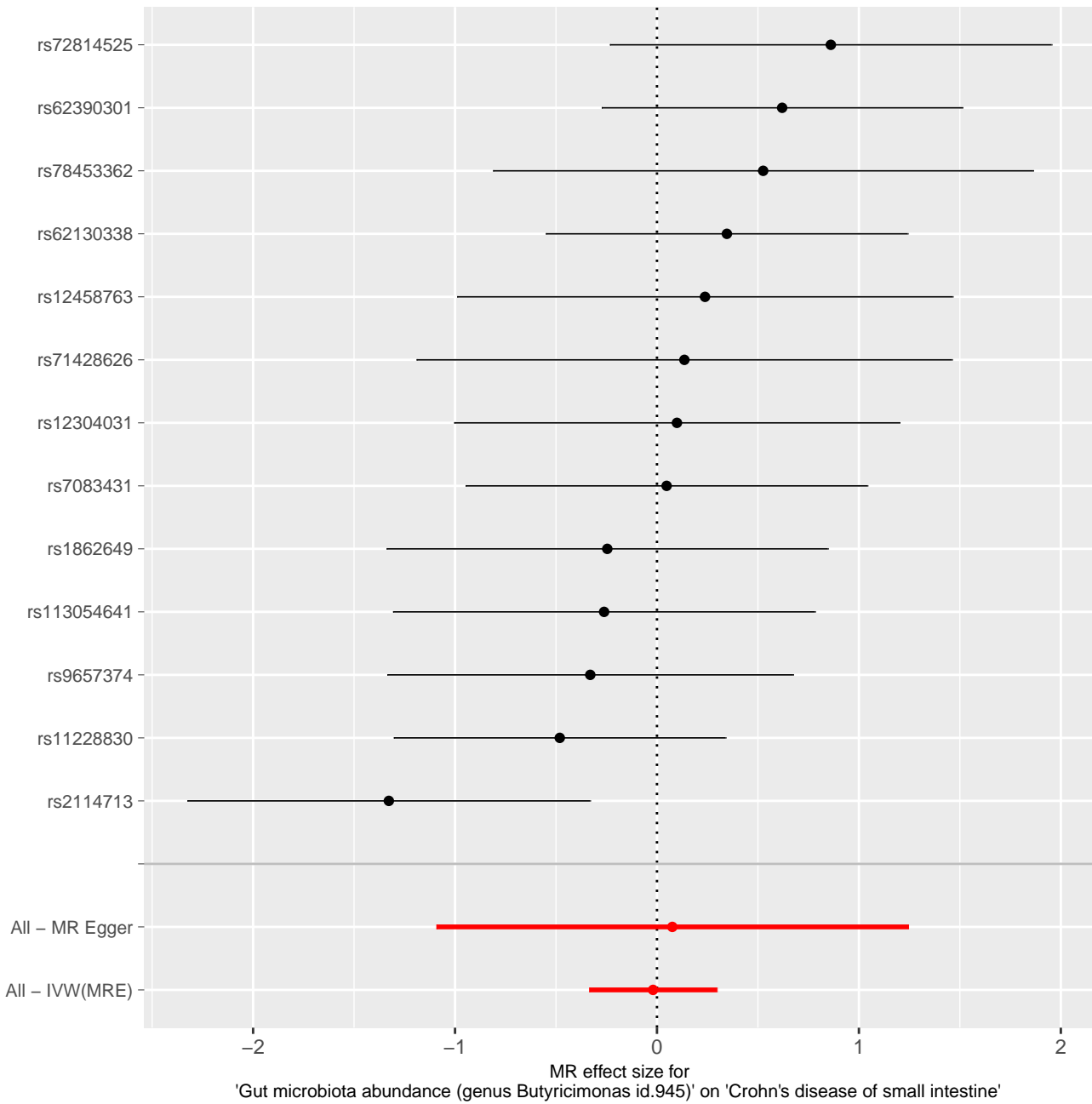


Batch 345 : Gut microbiota abundance (genus Bilophila id.3170) on Crohn's disease of small intestine

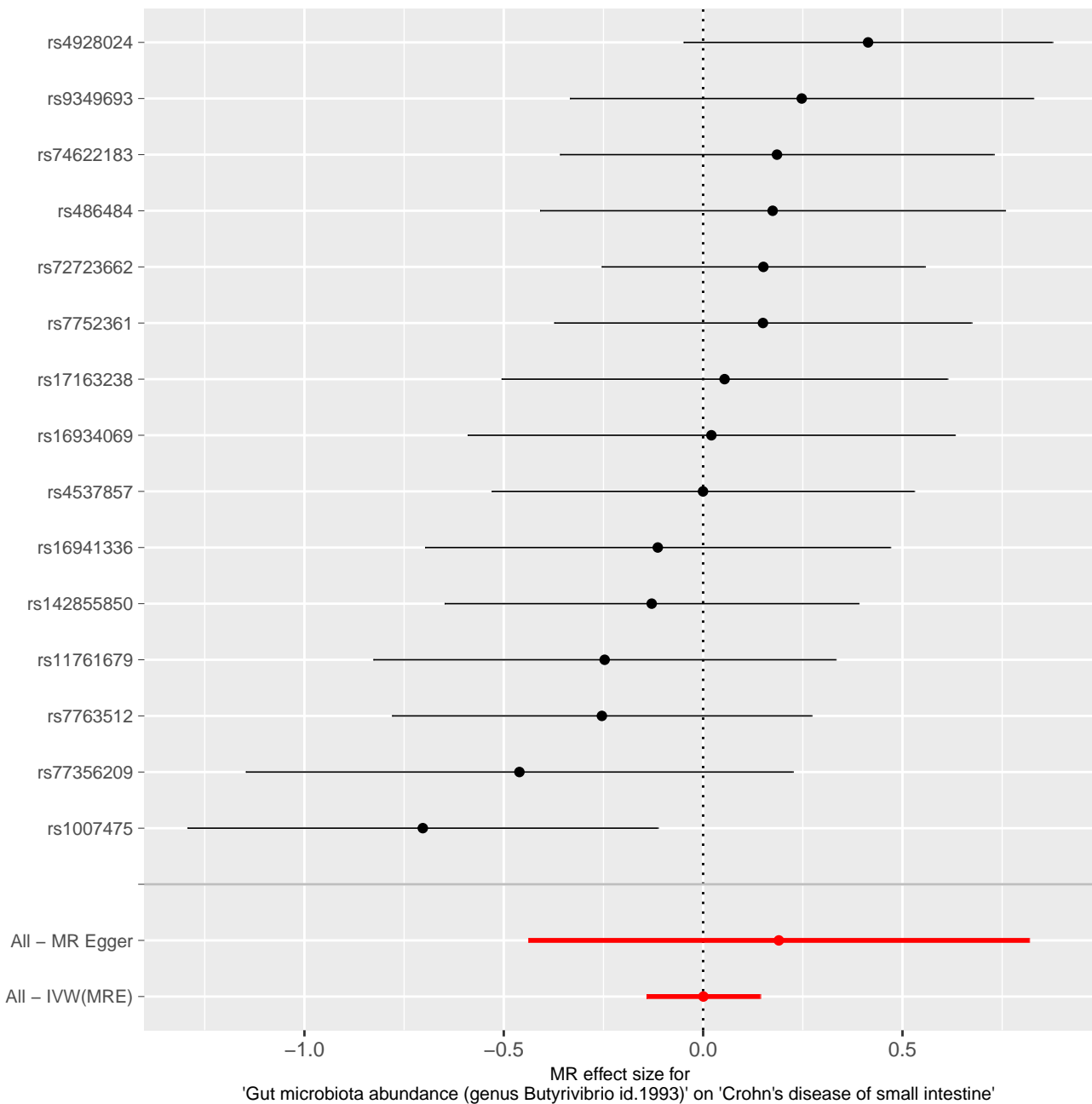


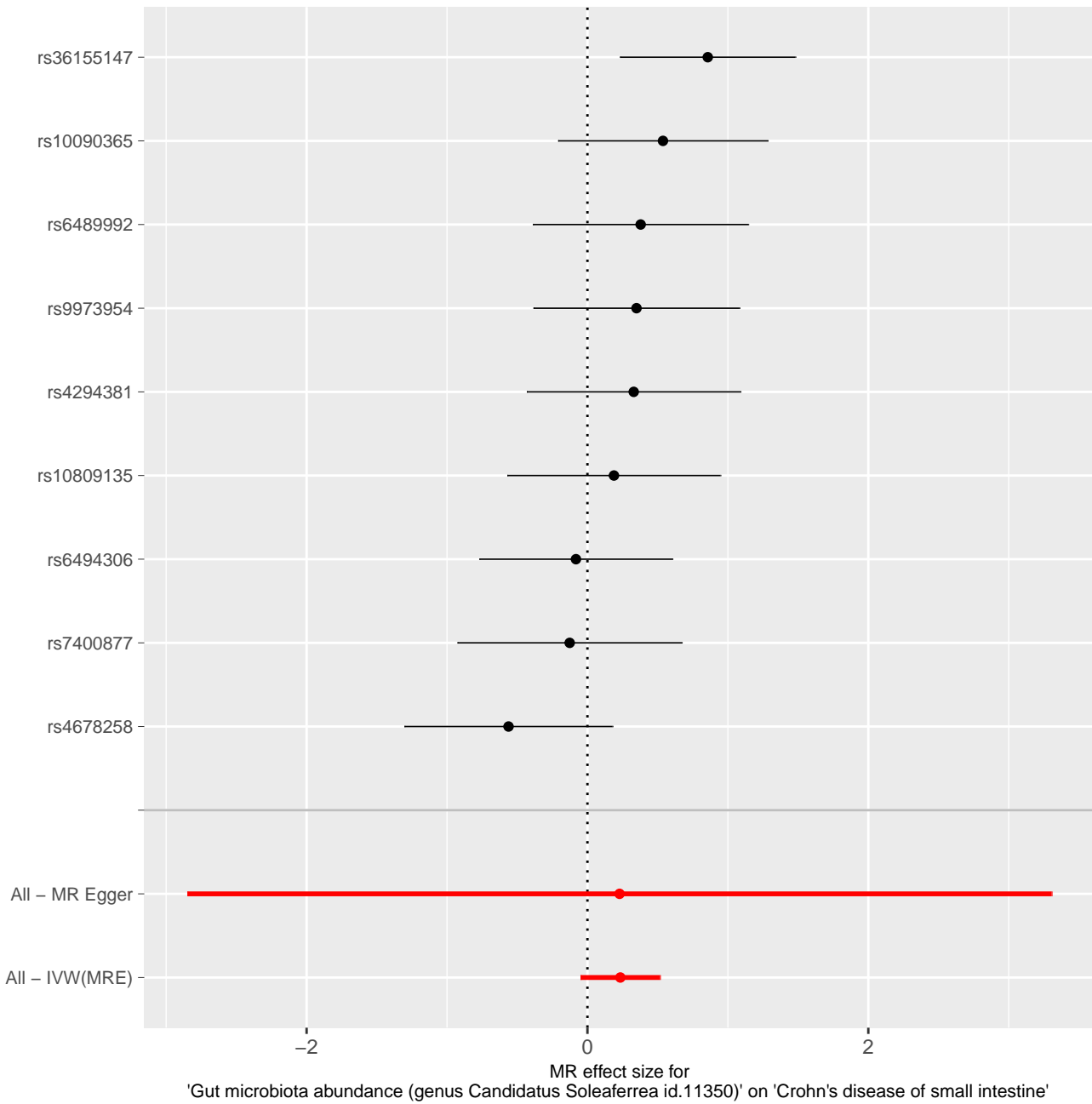
Batch 347 : Gut microbiota abundance (genus *Butyricoccus* id.2055) on Crohn's disease of small intestine

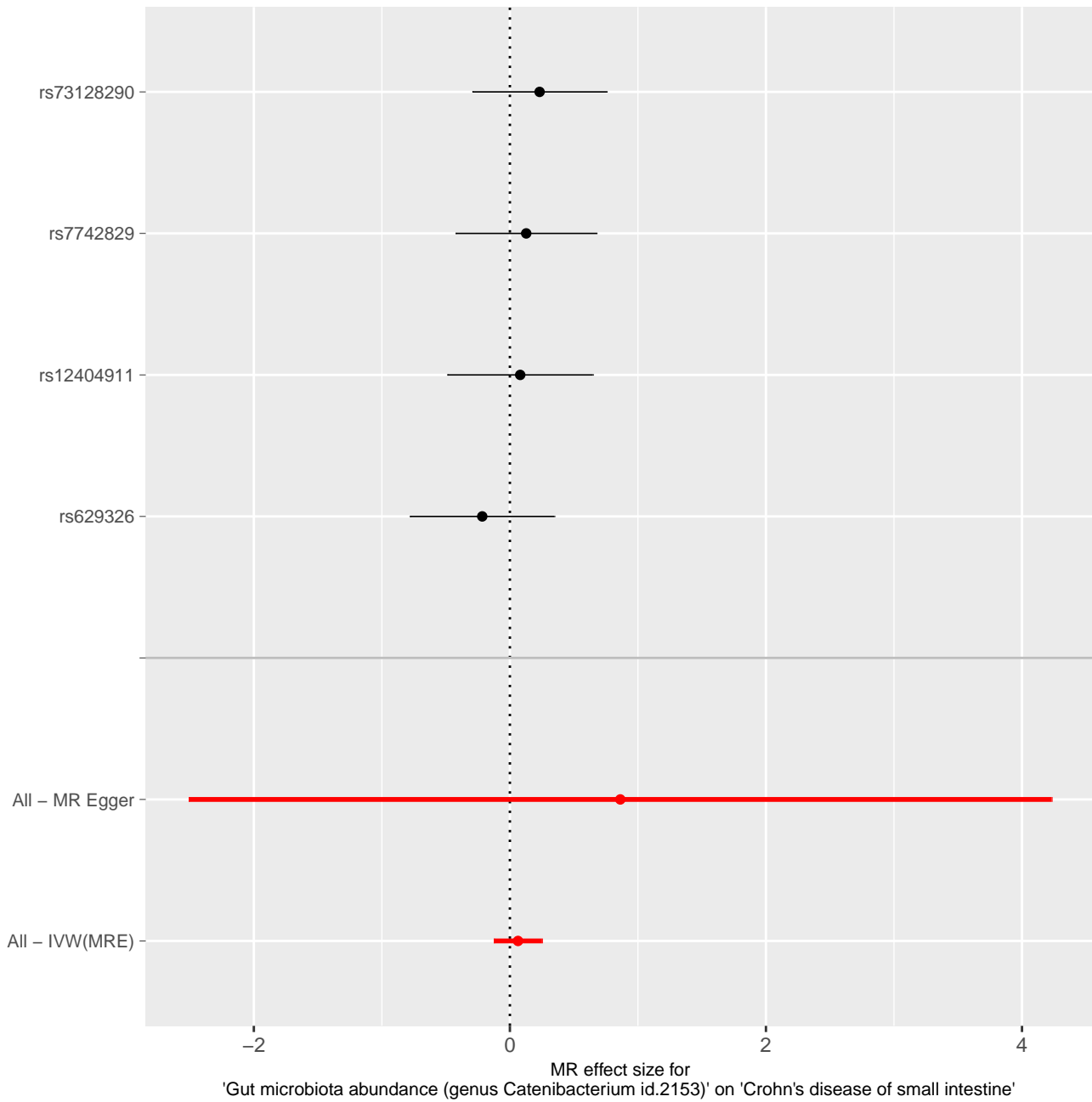


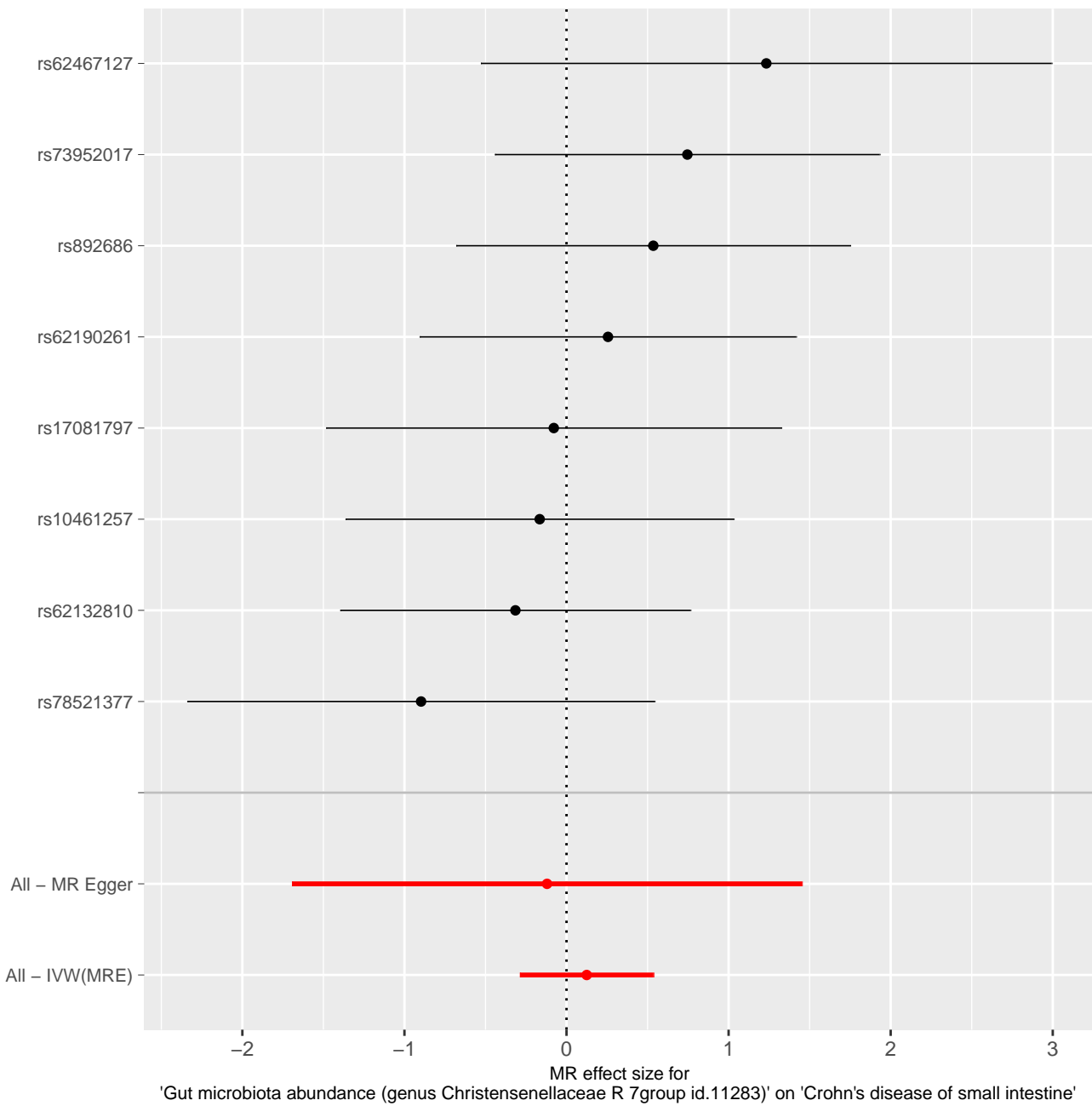


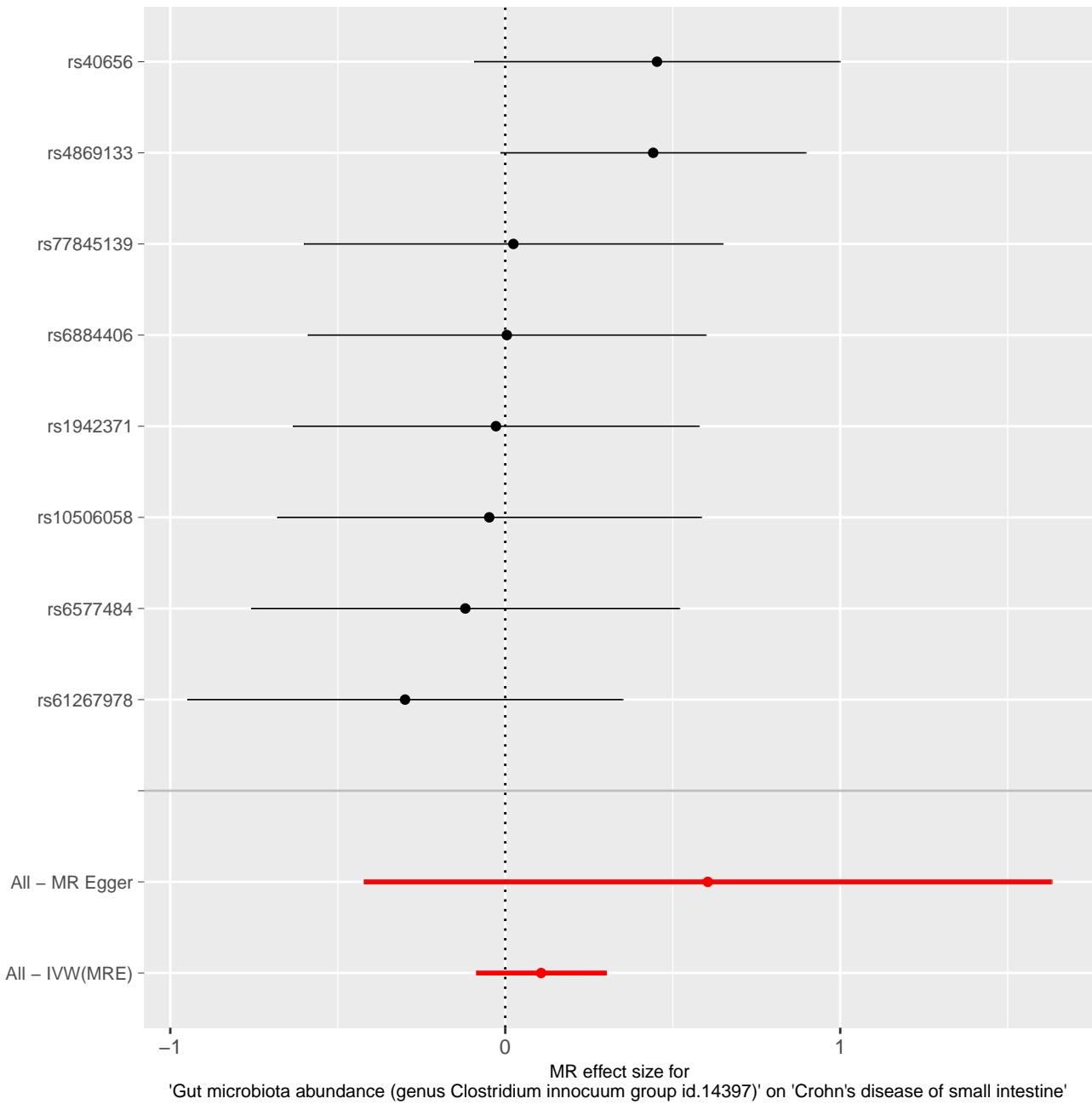
Batch 349 : Gut microbiota abundance (genus Butyrivibrio id.1993) on Crohn's disease of small intestine



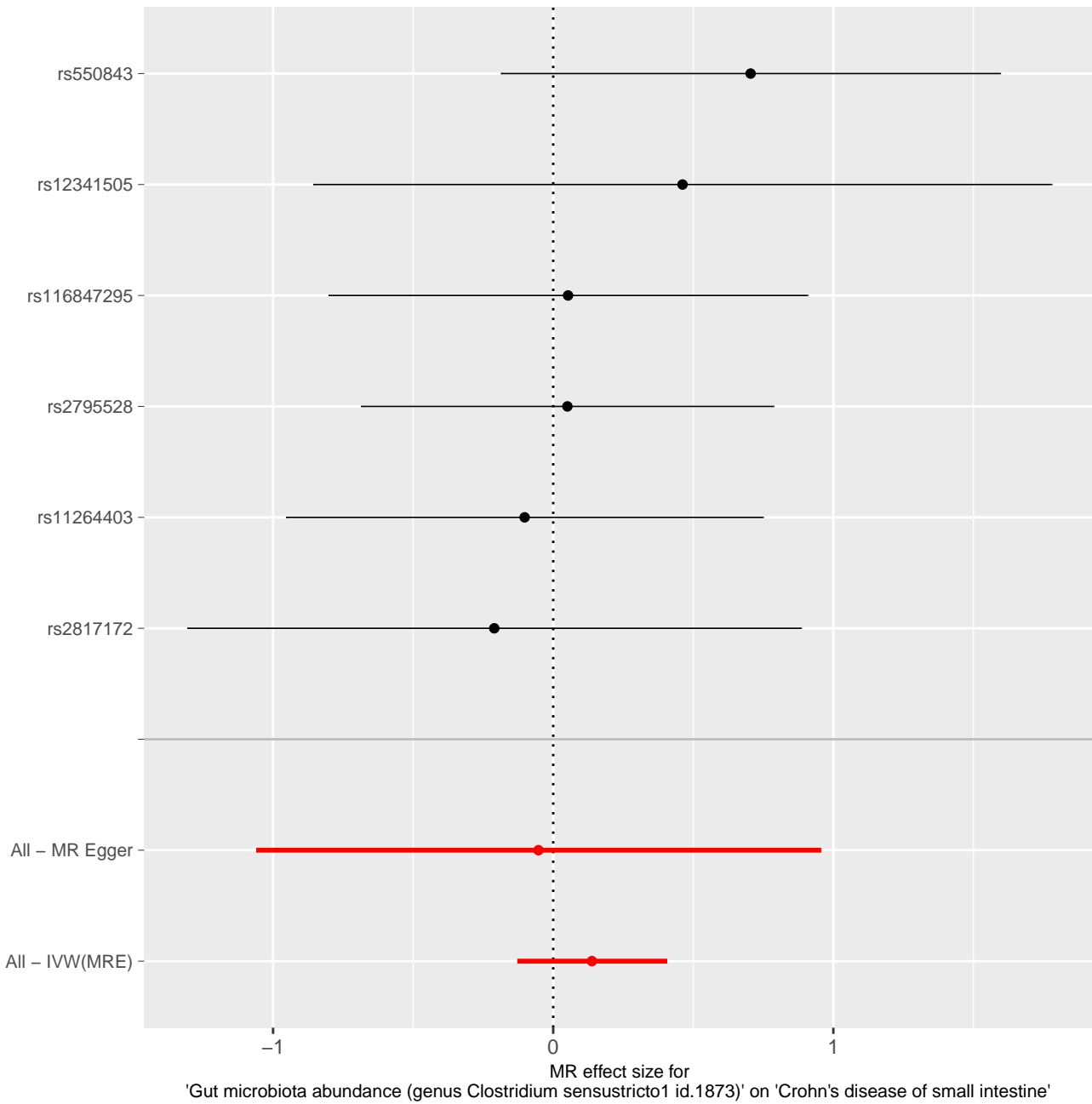




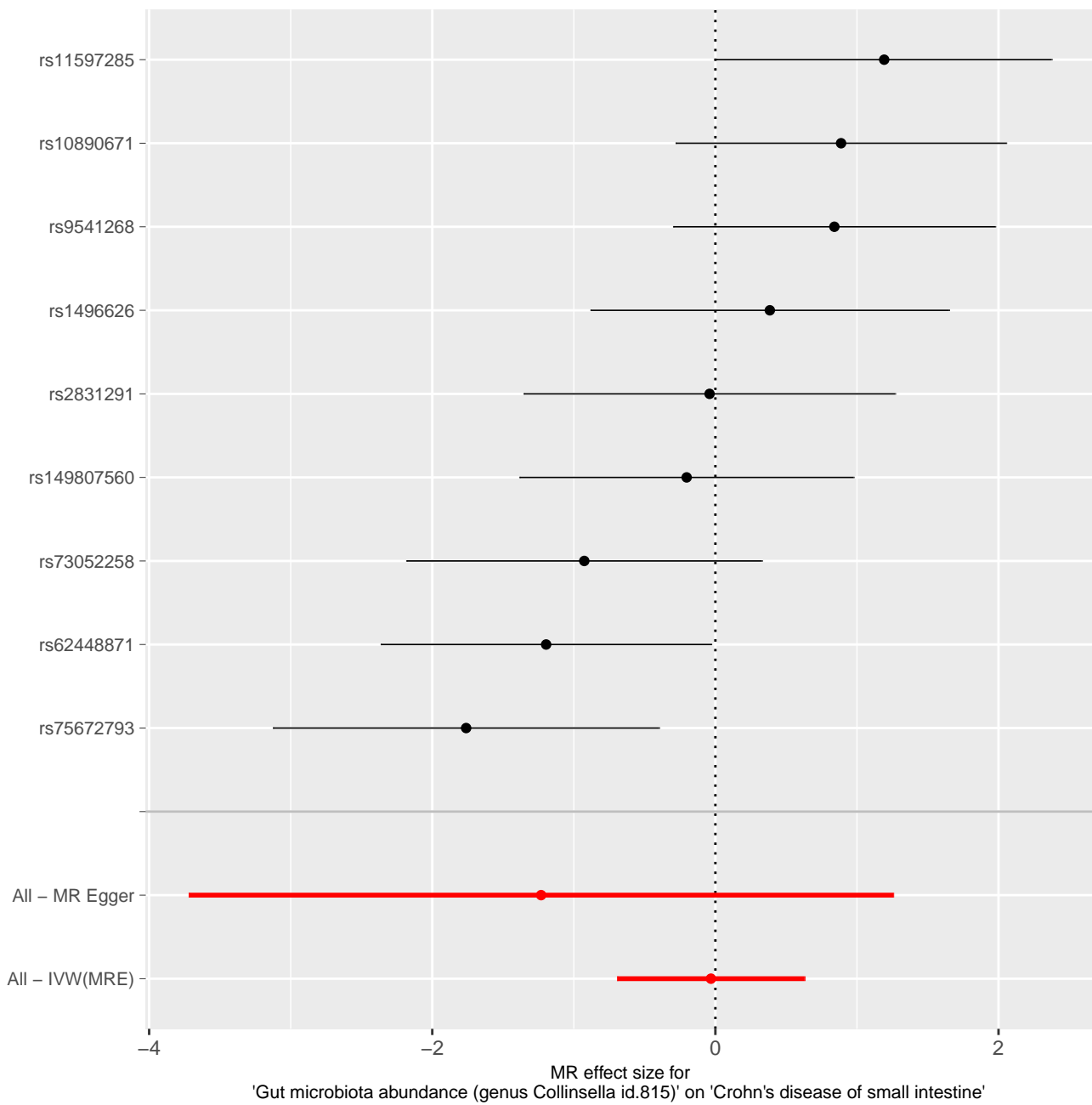




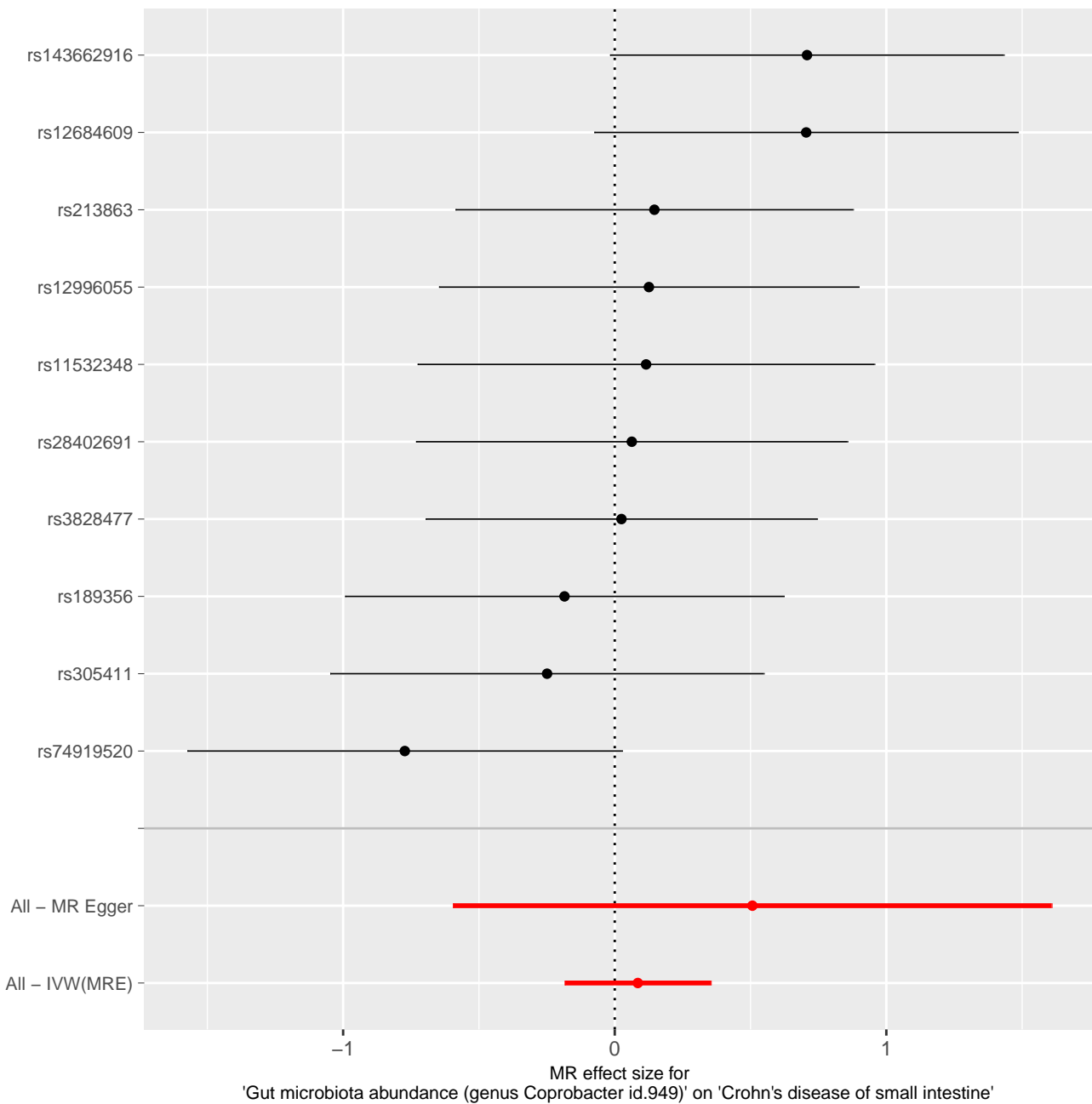
MR effect size for 'Gut microbiota abundance (genus Clostridium innocuum group id.14397)' on 'Crohn's disease of small intestine'



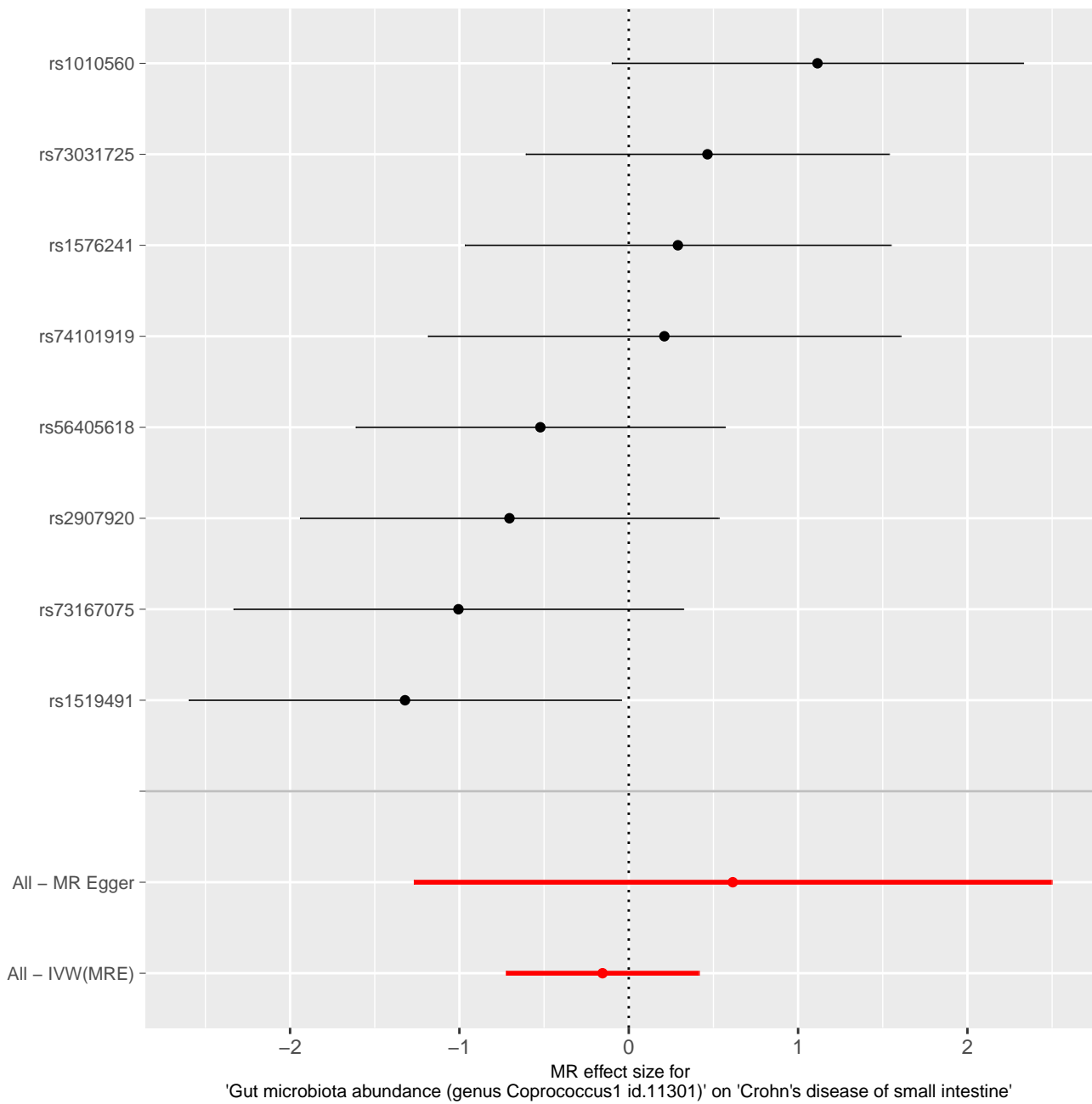
Batch 355 : Gut microbiota abundance (genus Collinsella id.815) on Crohn's disease of small intestine

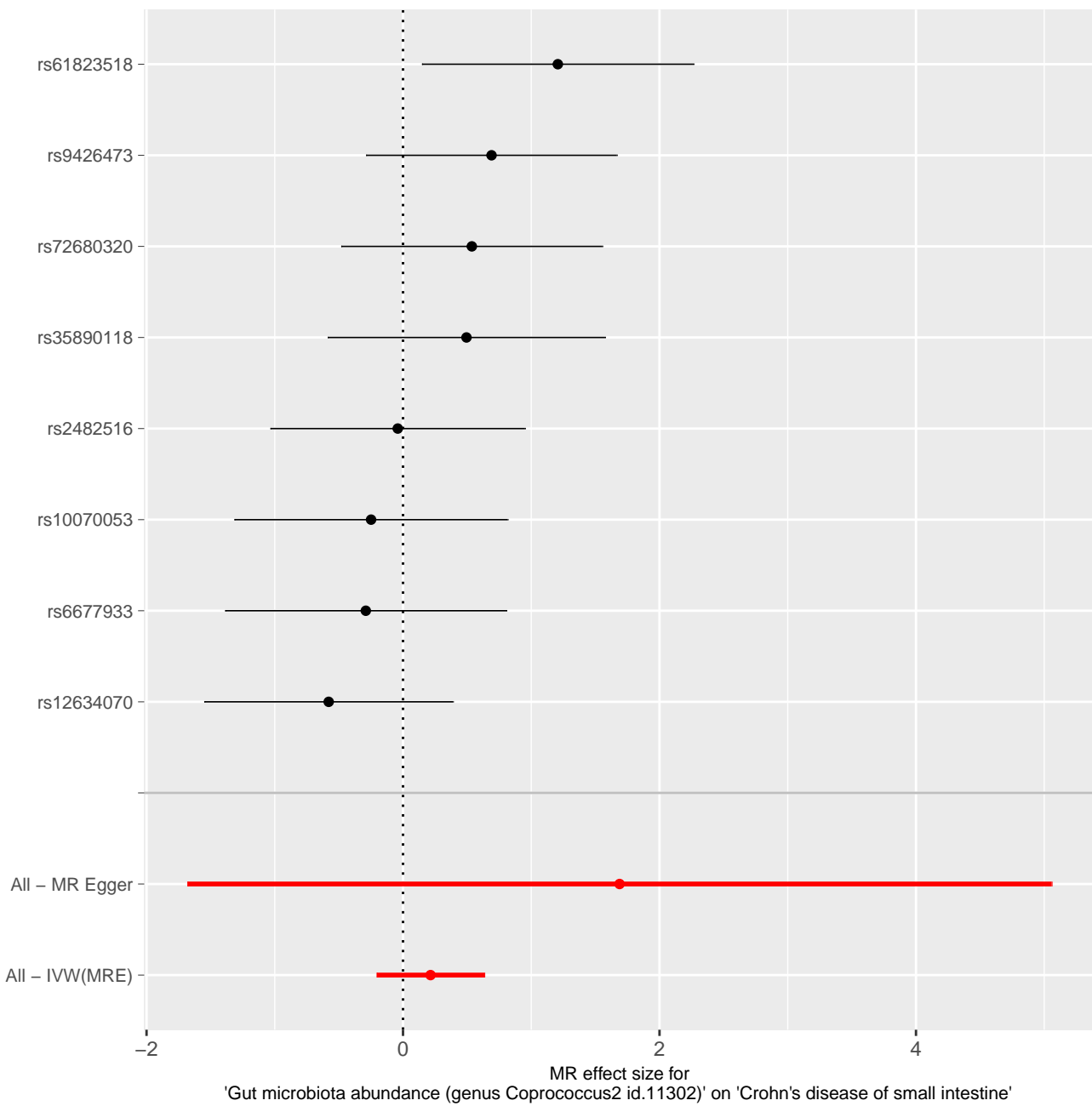


Batch 356 : Gut microbiota abundance (genus Coprobacter id.949) on Crohn's disease of small intestine

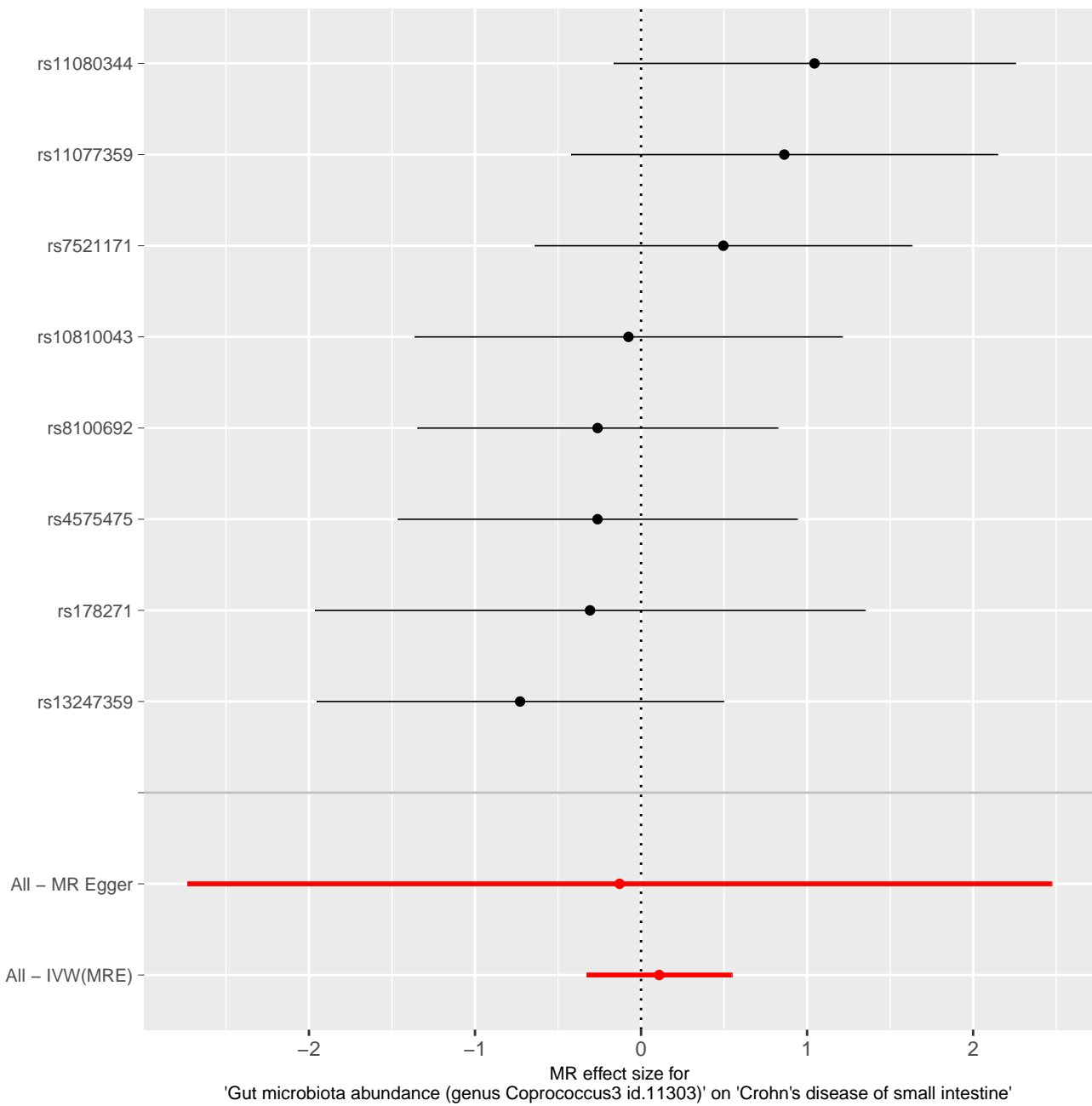


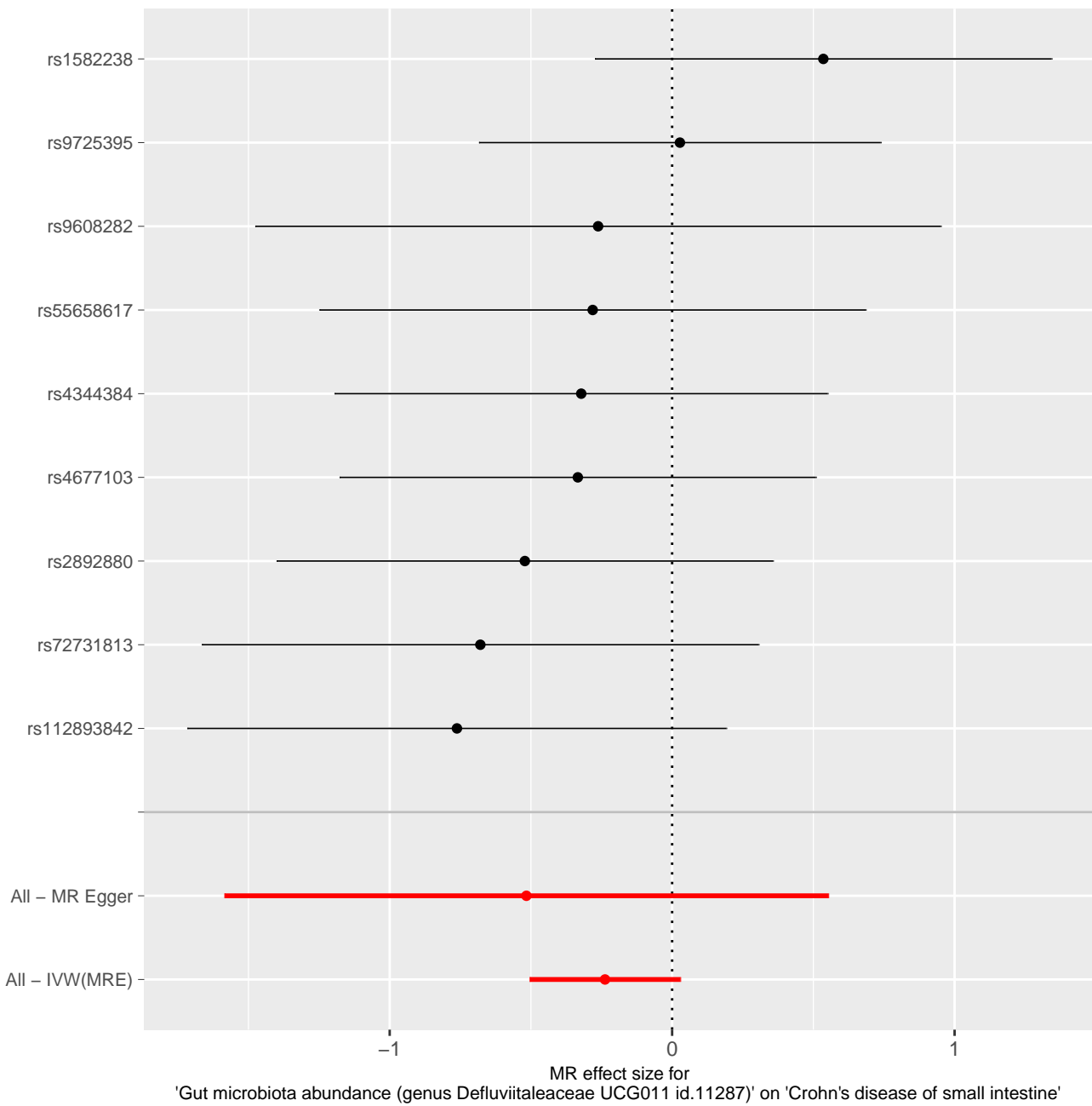
Batch 357 : Gut microbiota abundance (genus Coprococcus1 id.11301) on Crohn's disease of small intestine

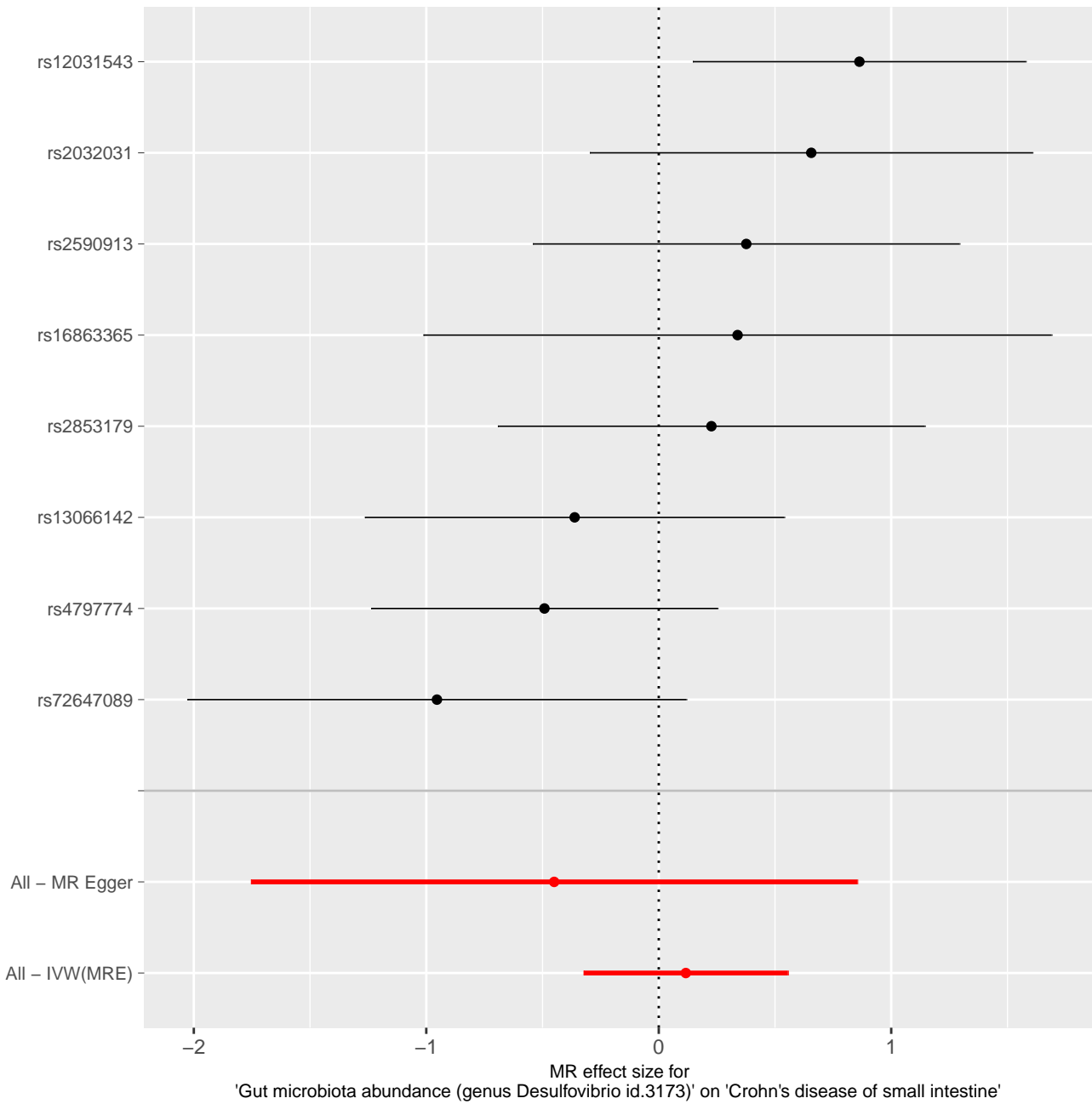


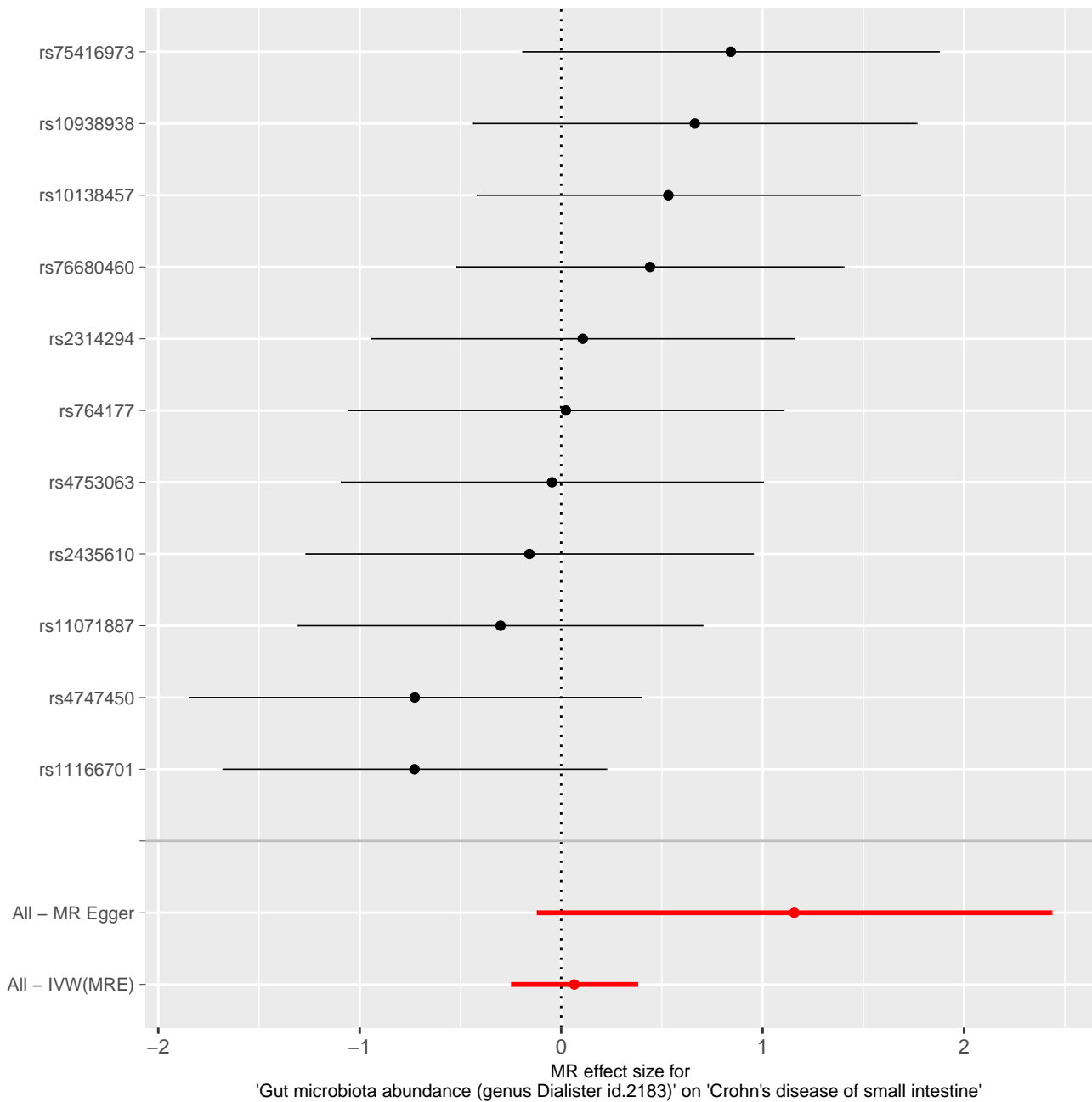


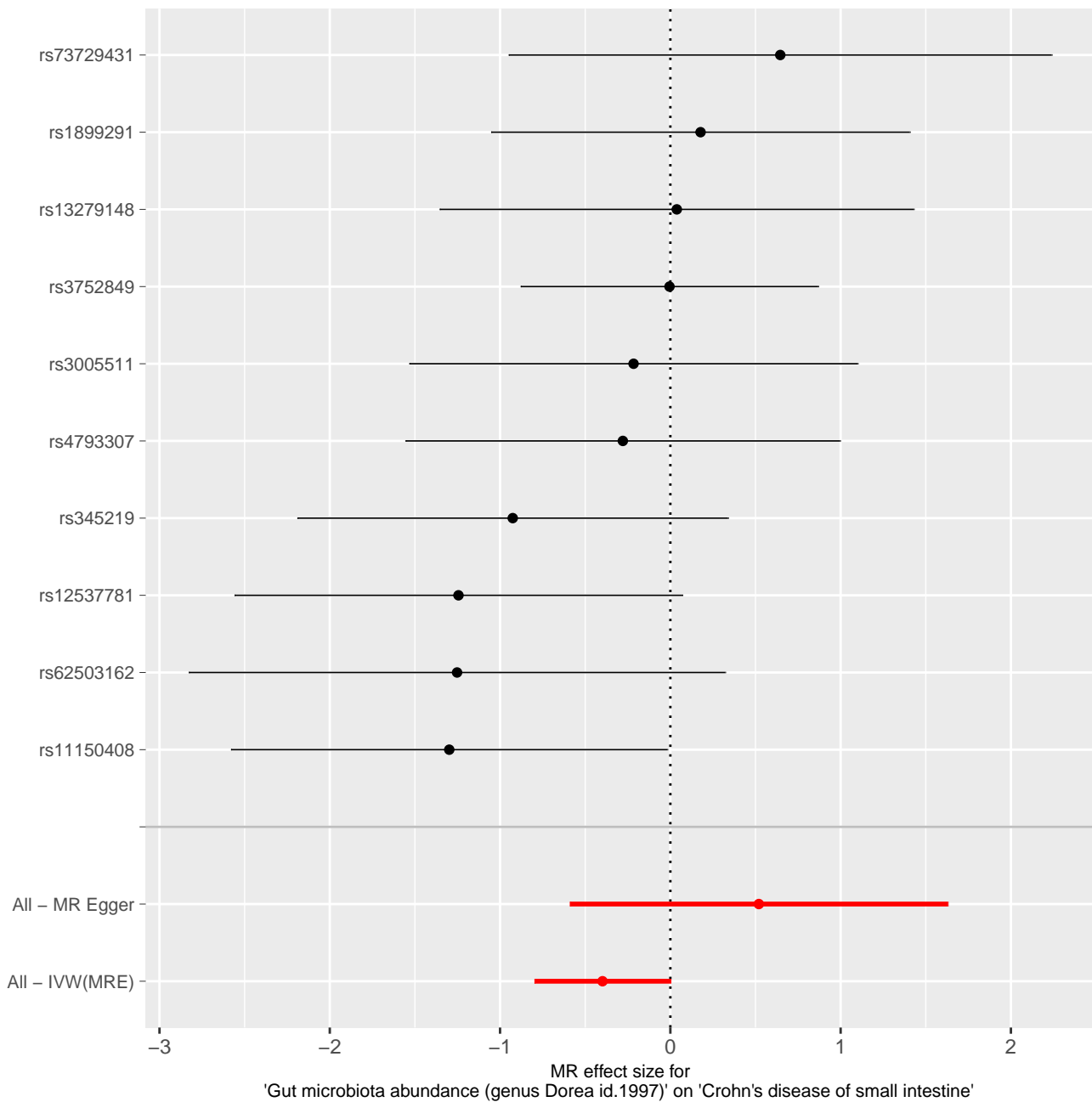
Batch 359 : Gut microbiota abundance (genus Coprococcus3 id.11303) on Crohn's disease of small intestine



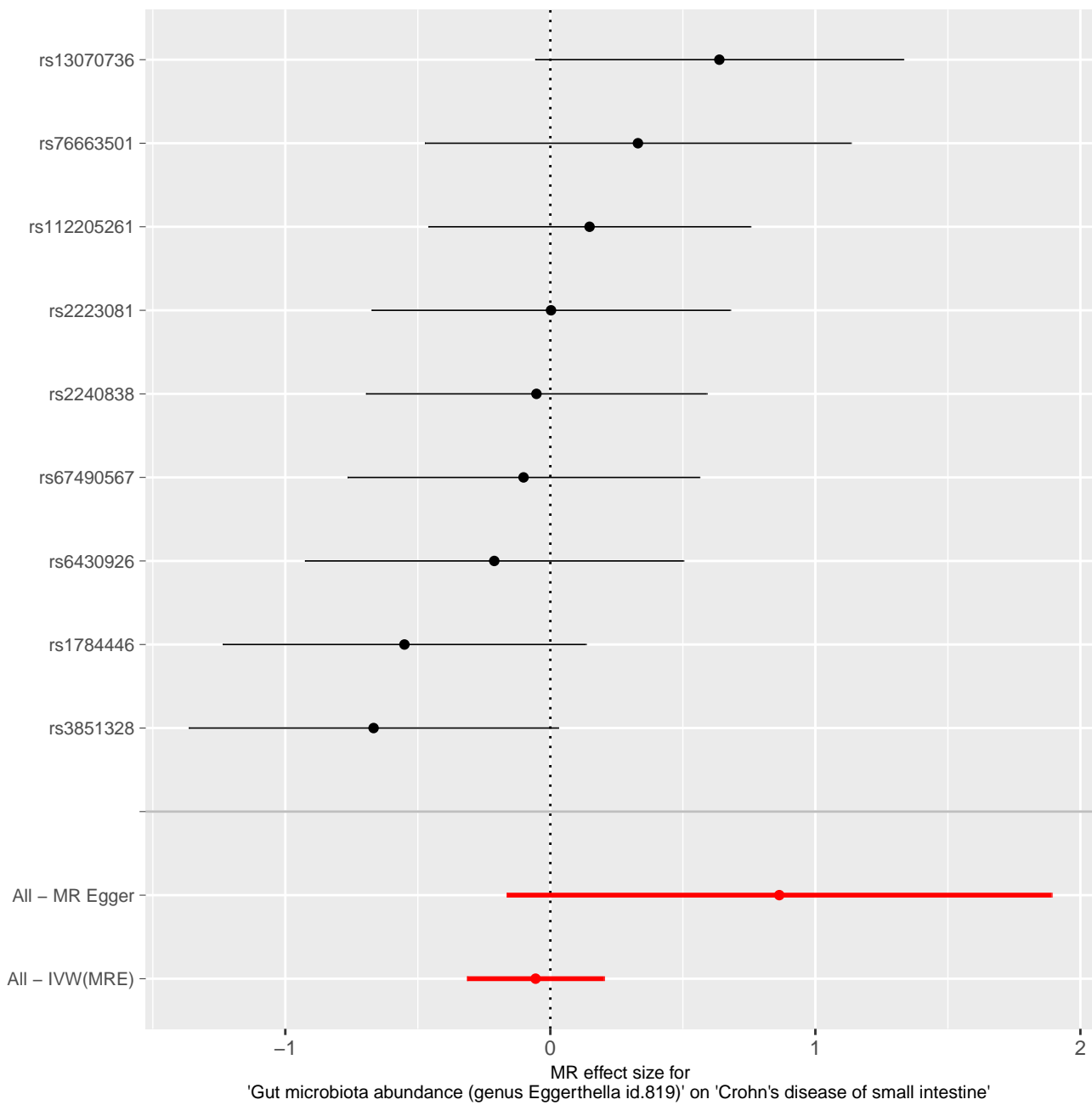


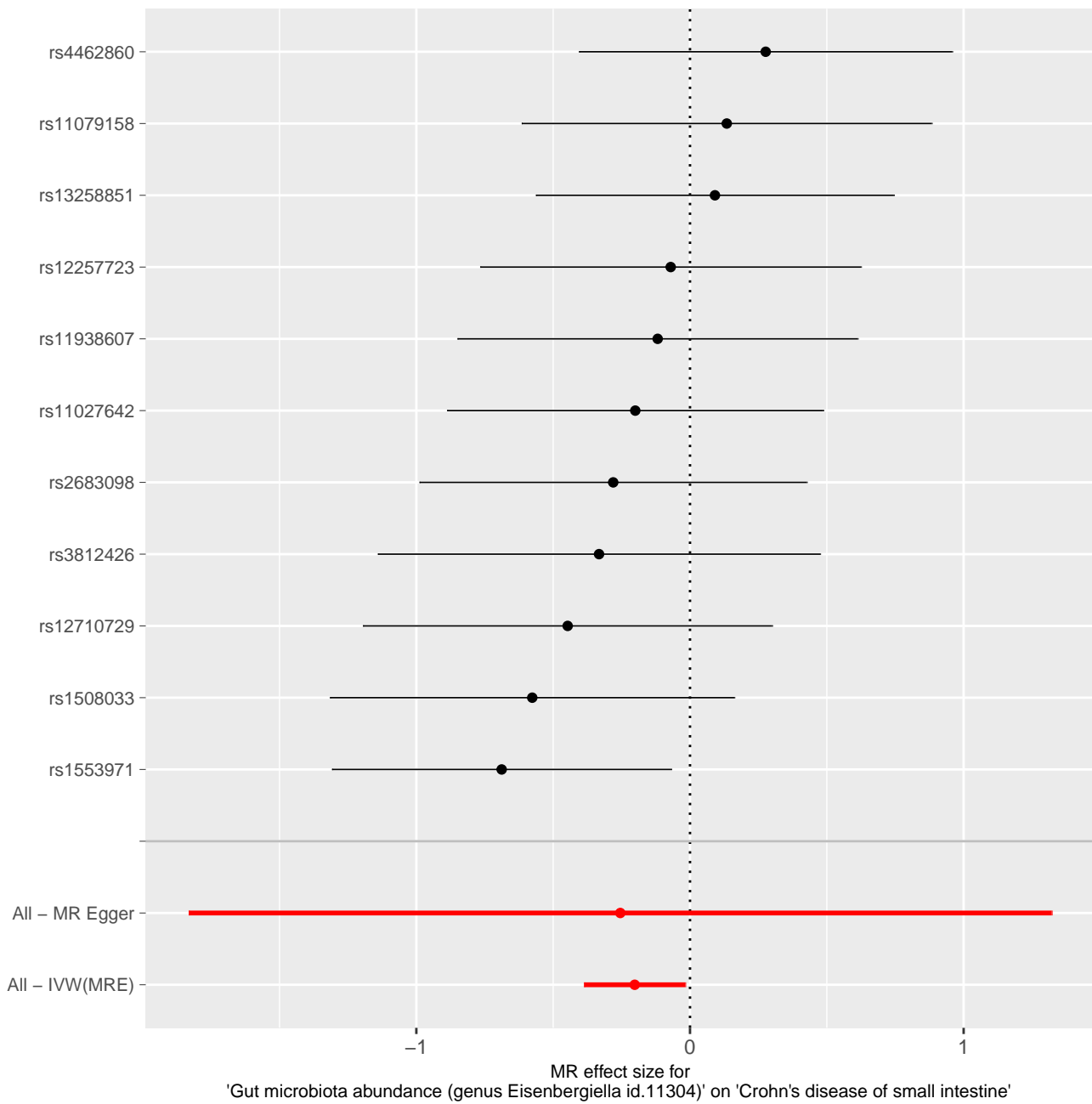


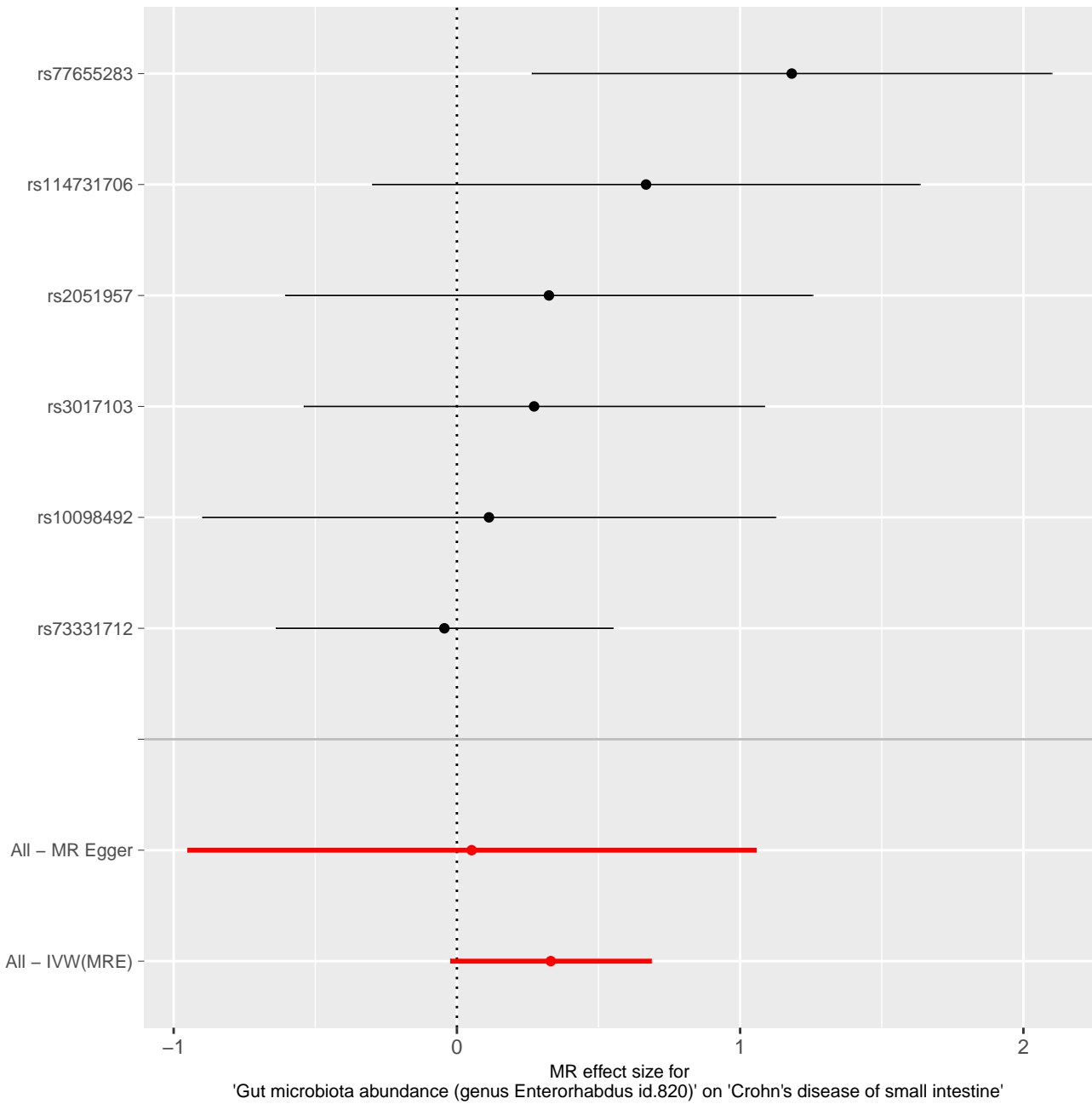


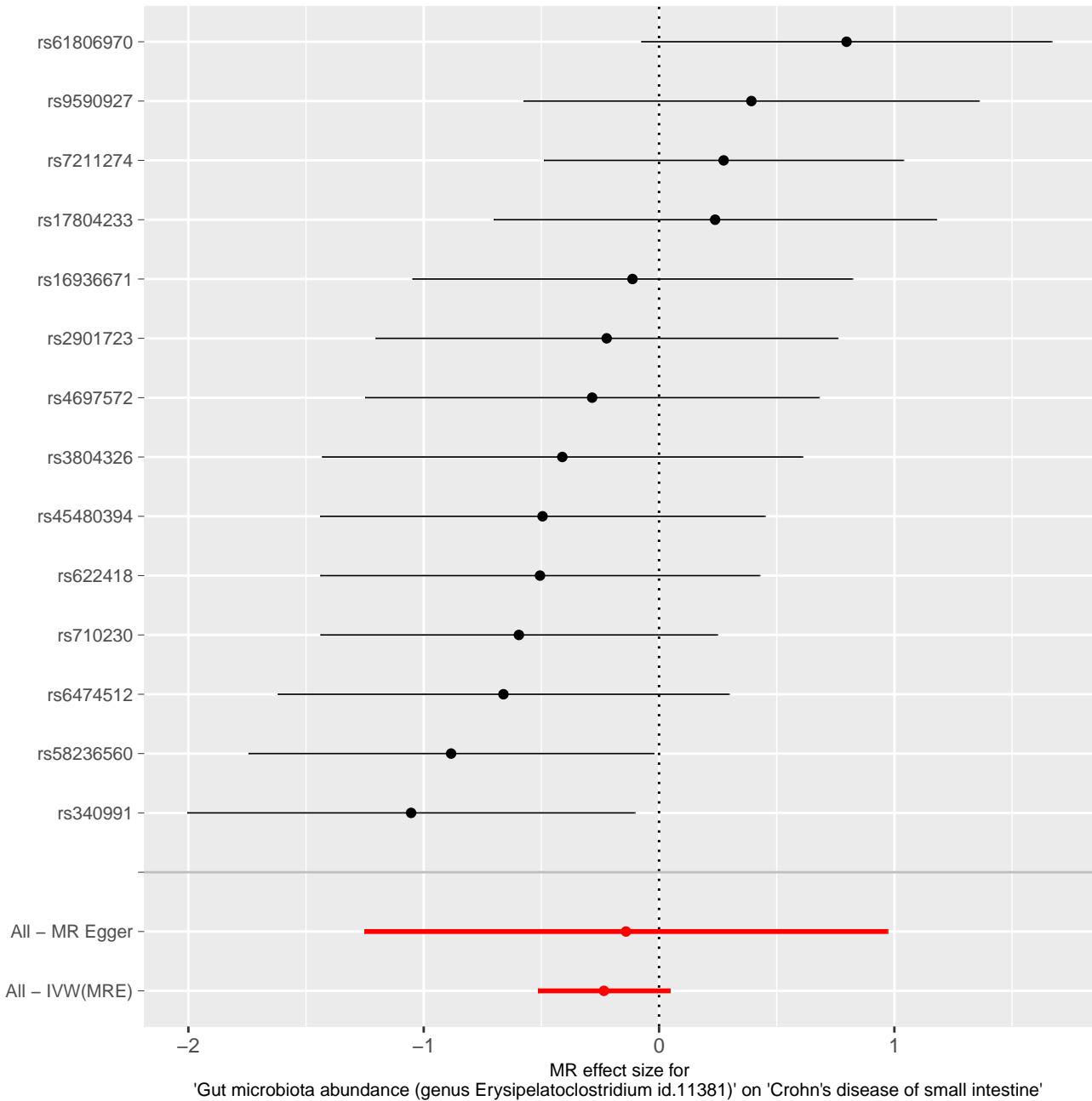


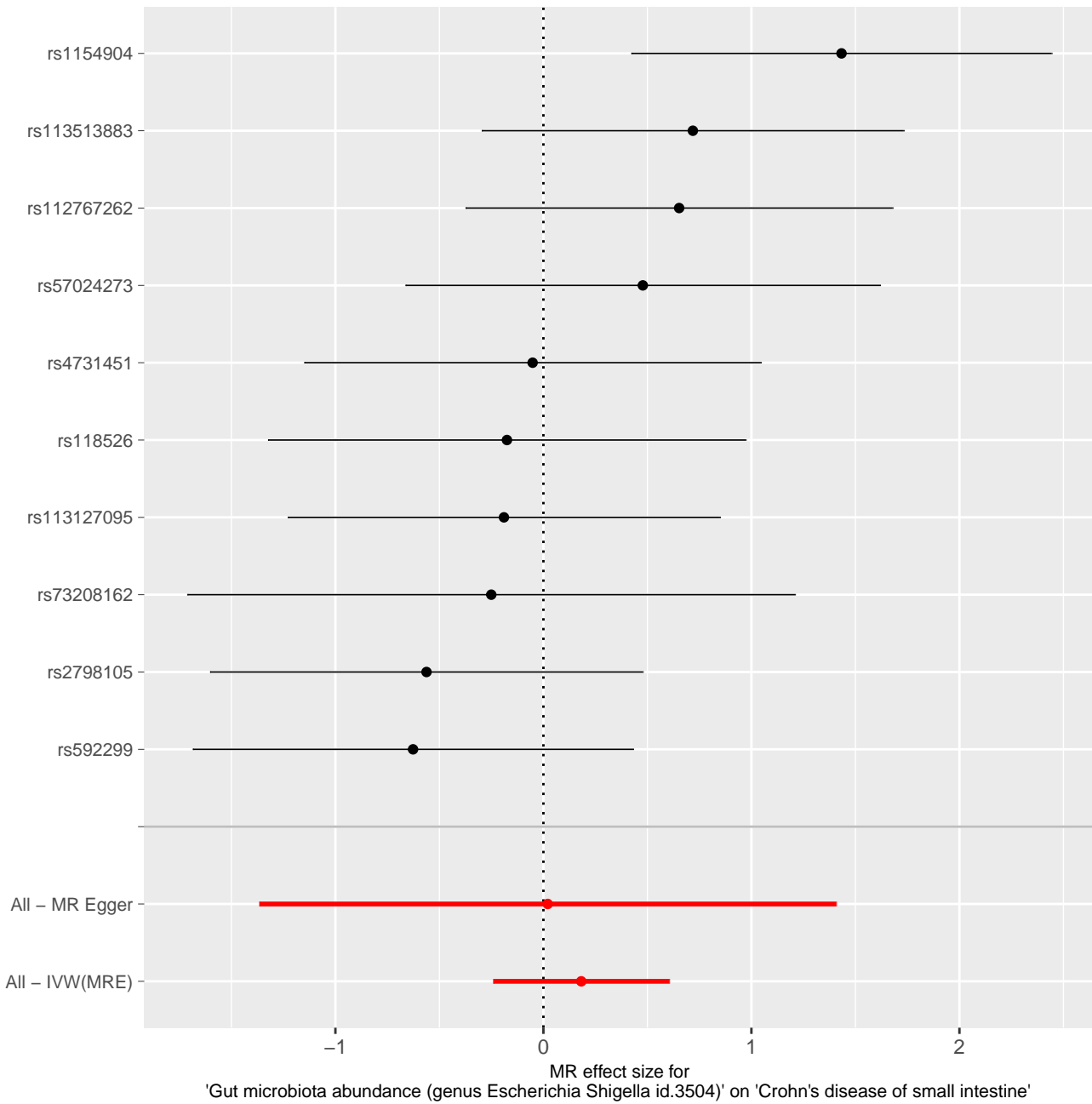
Batch 364 : Gut microbiota abundance (genus Eggerthella id.819) on Crohn's disease of small intestine



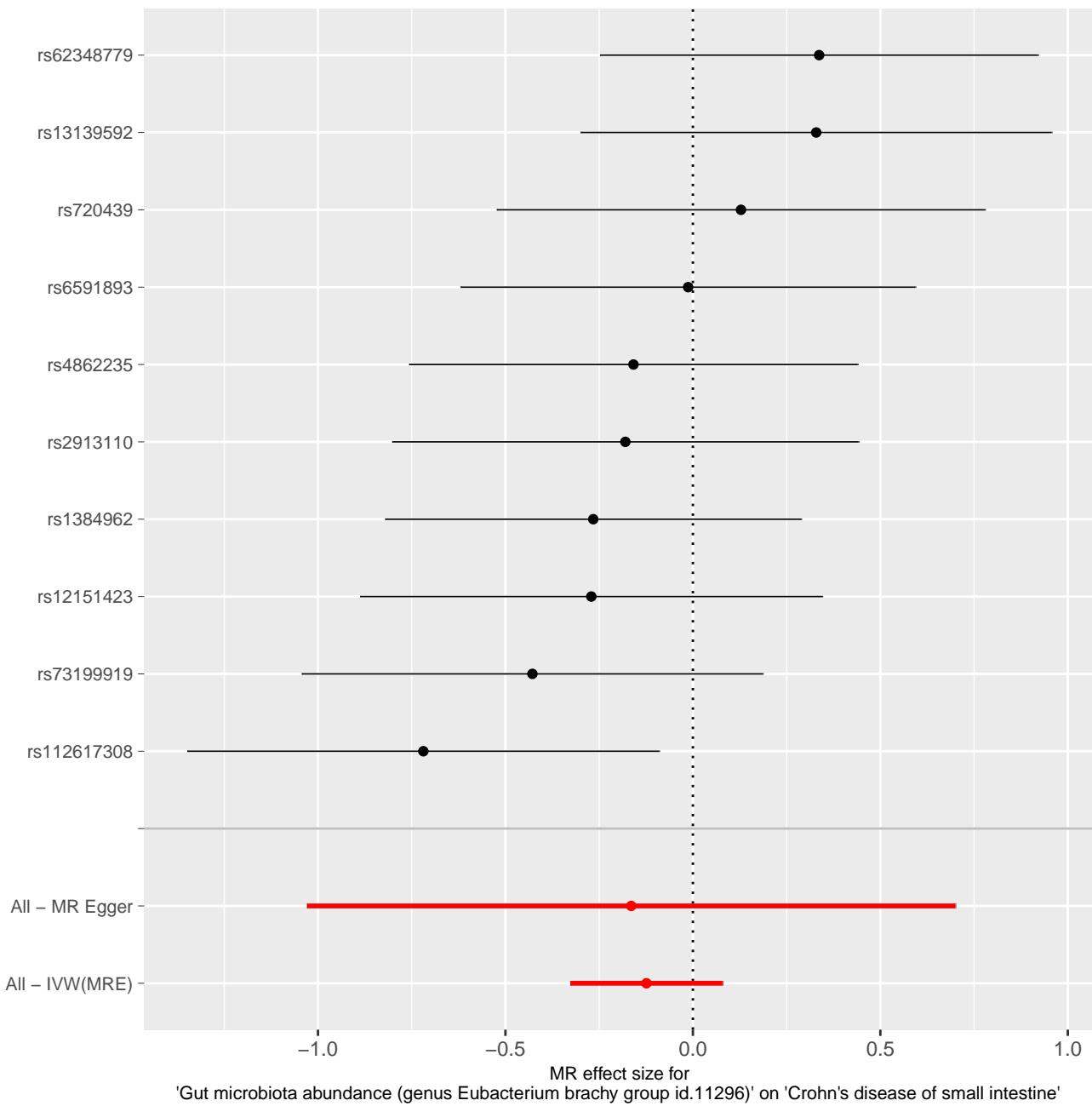


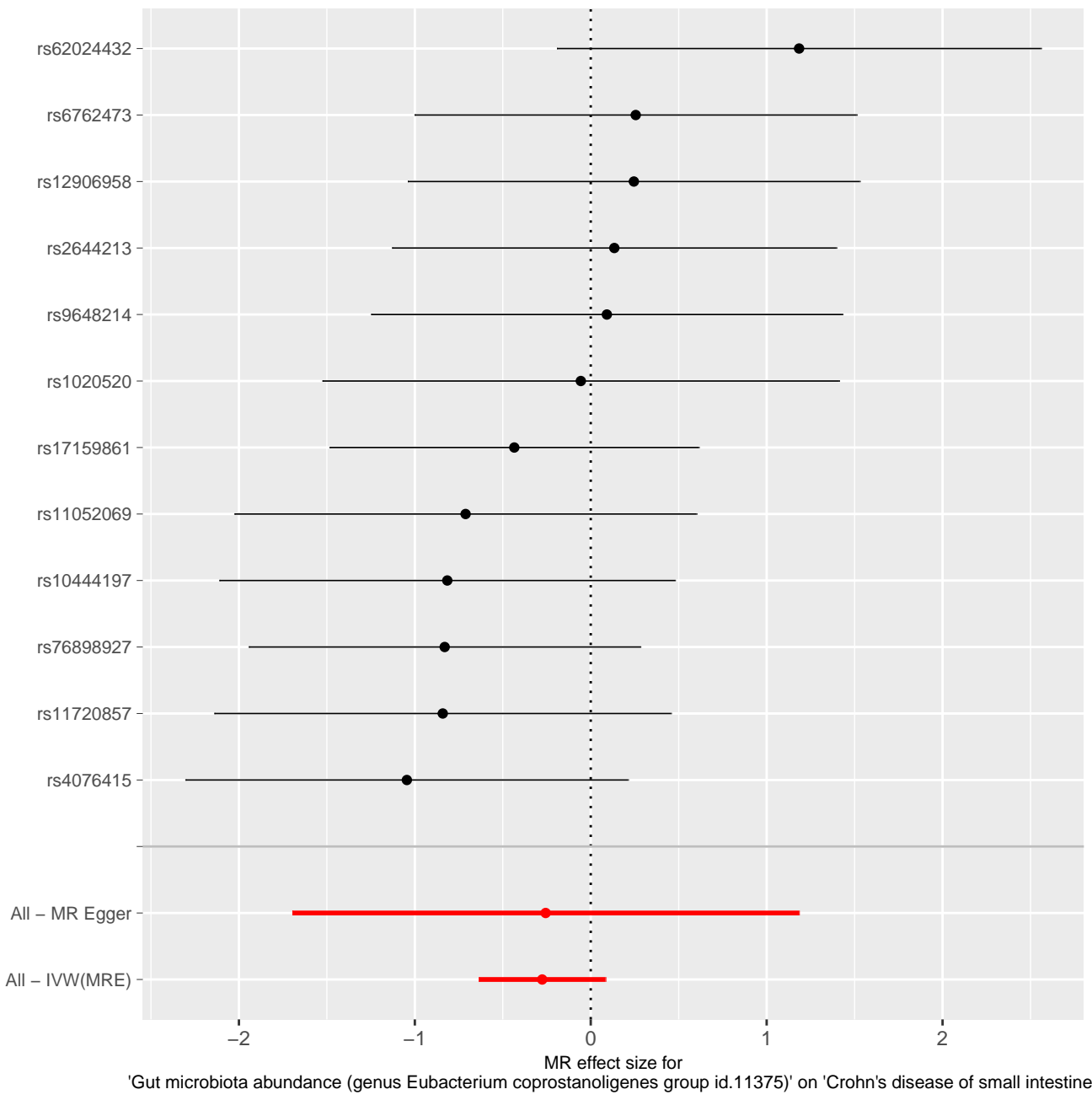


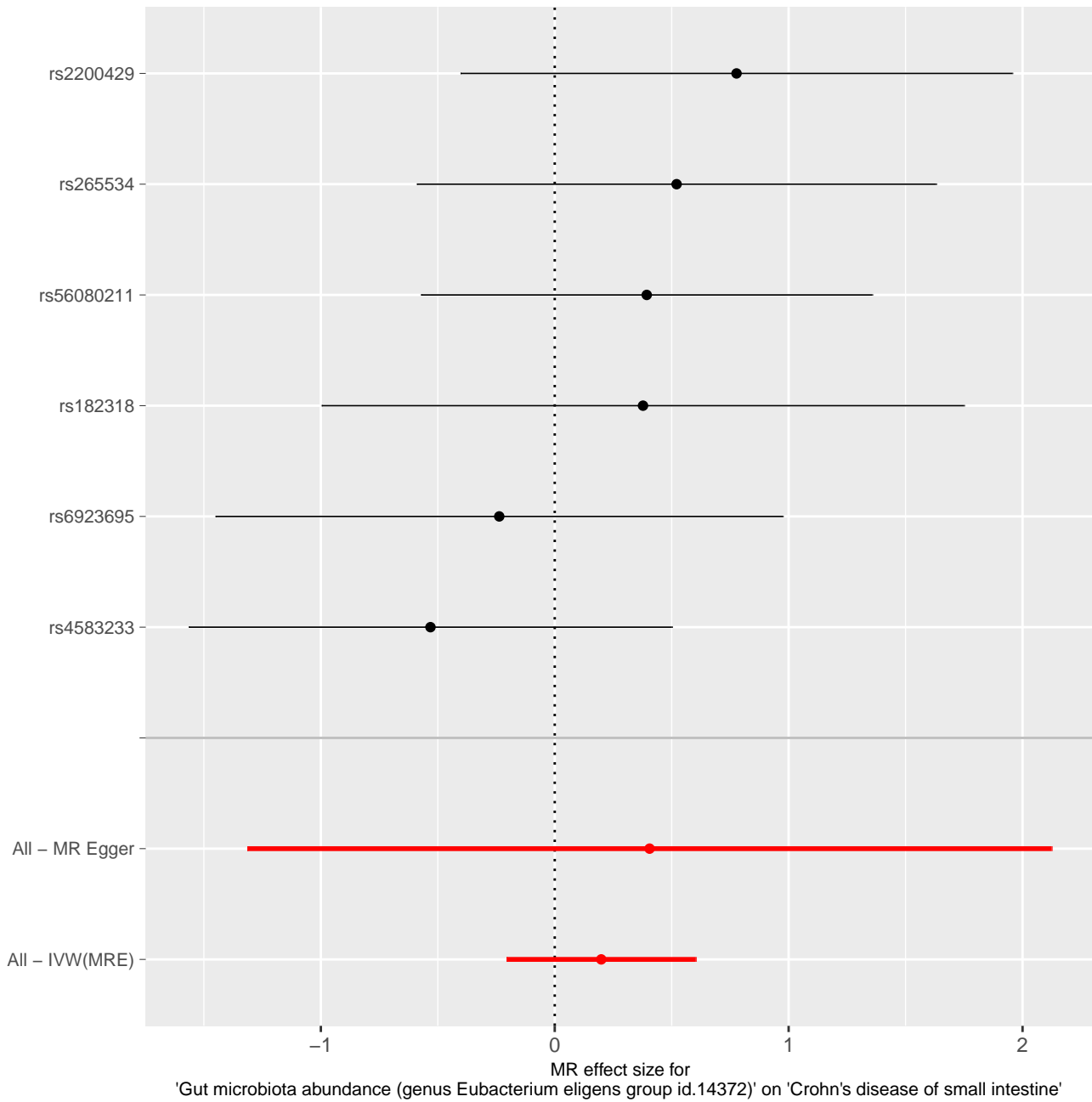


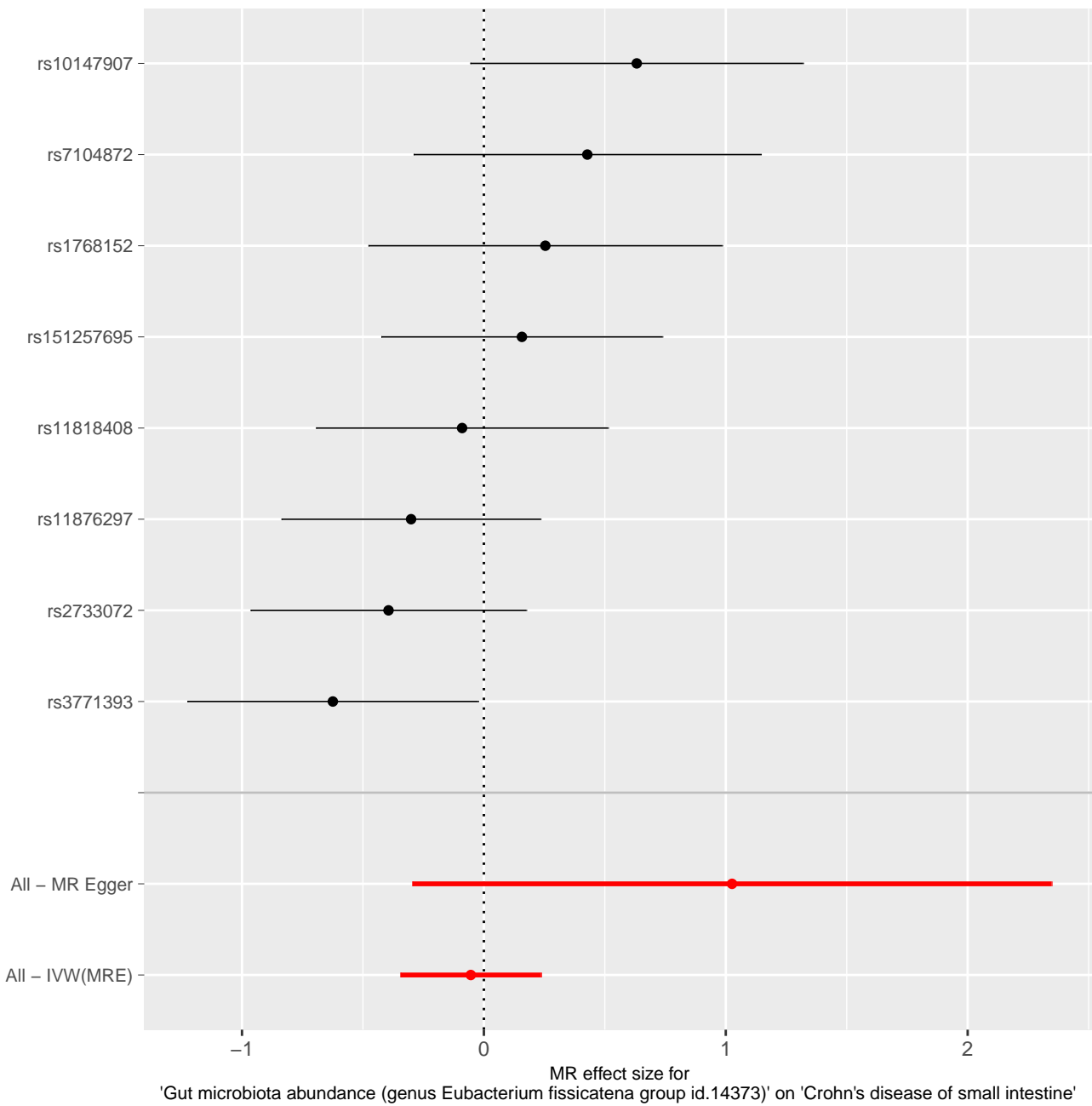


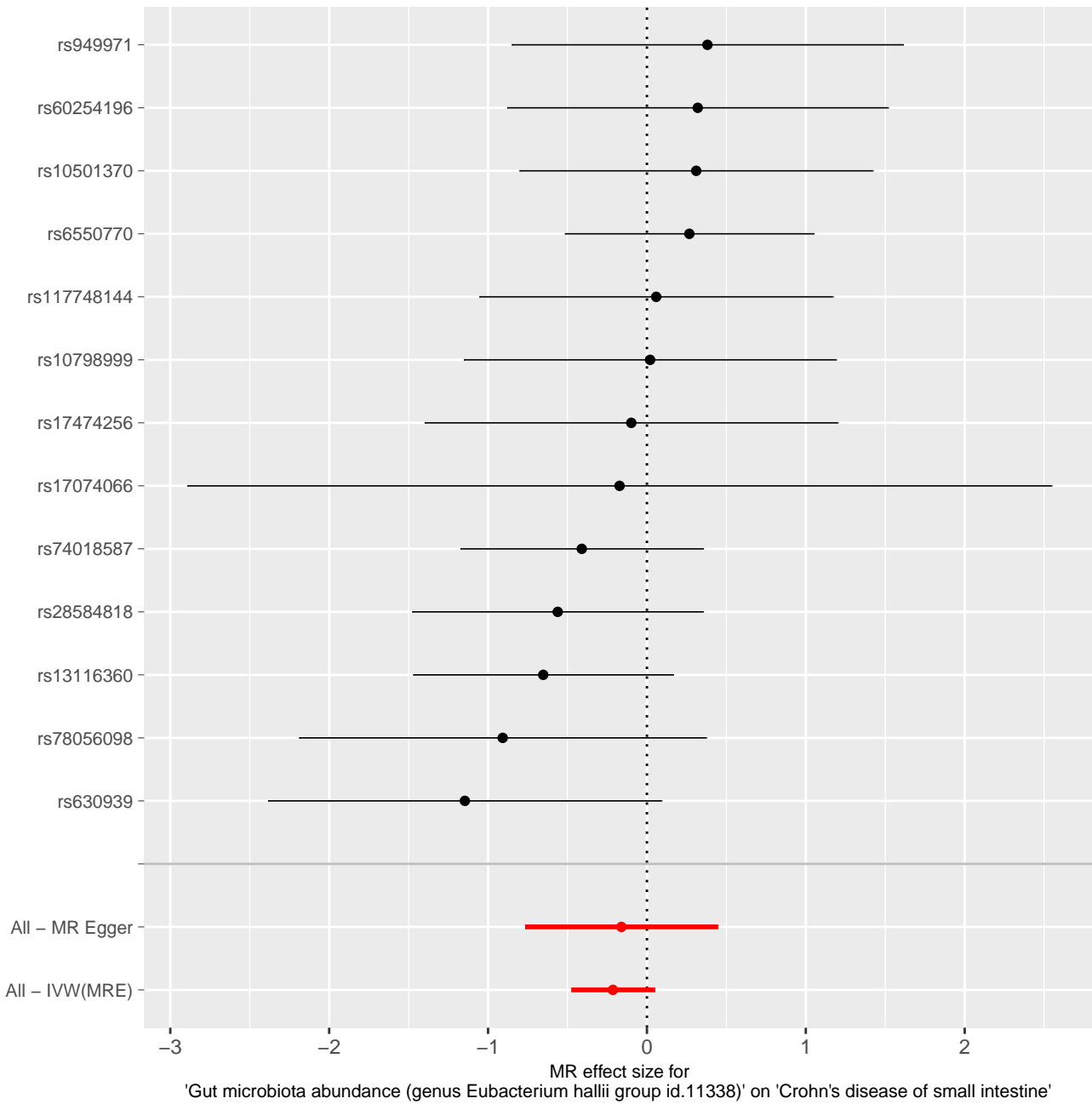
Batch 370 : Gut microbiota abundance (genus Eubacterium brachy group id.11296) on Crohn's disease of small intestine



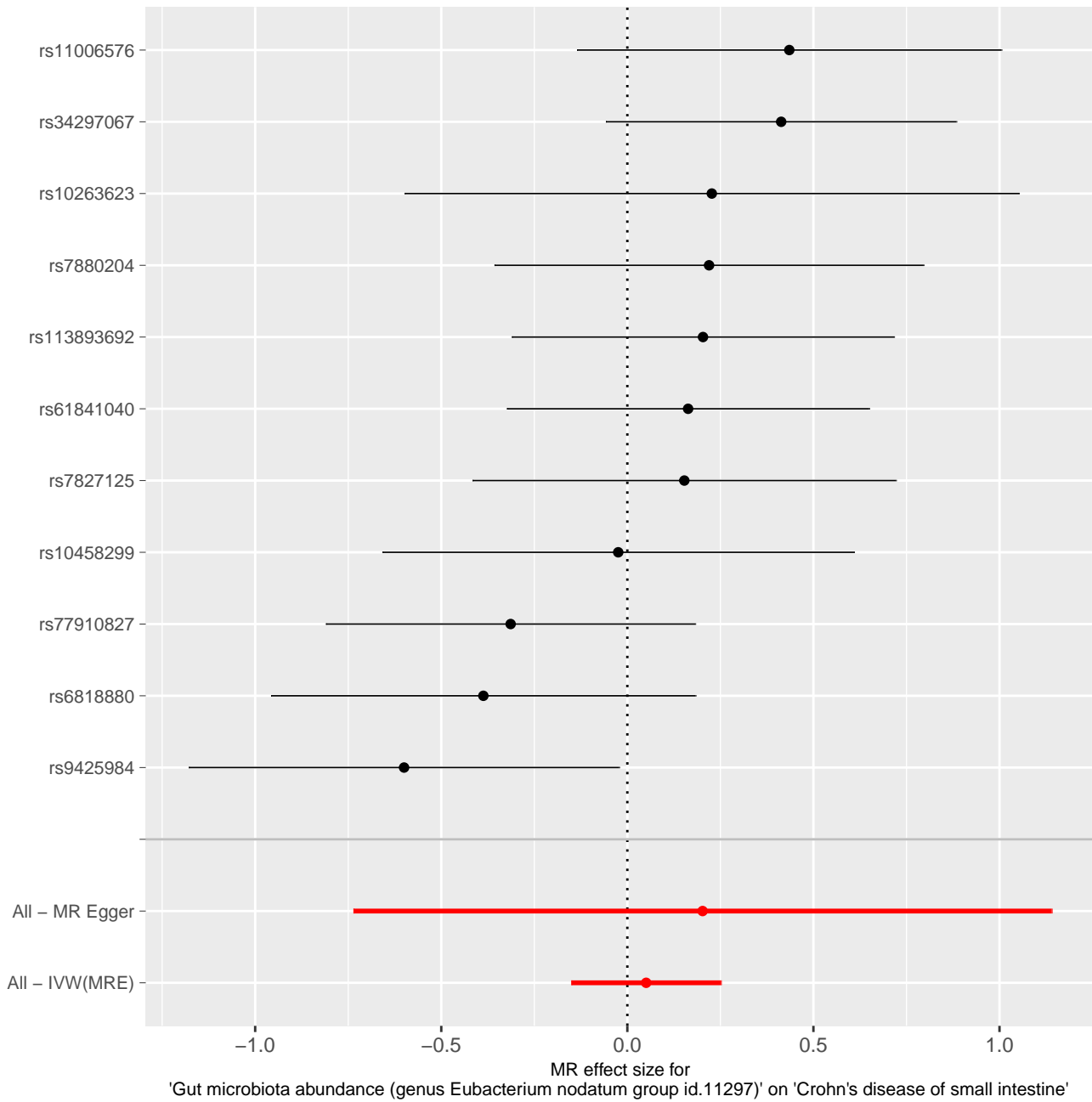


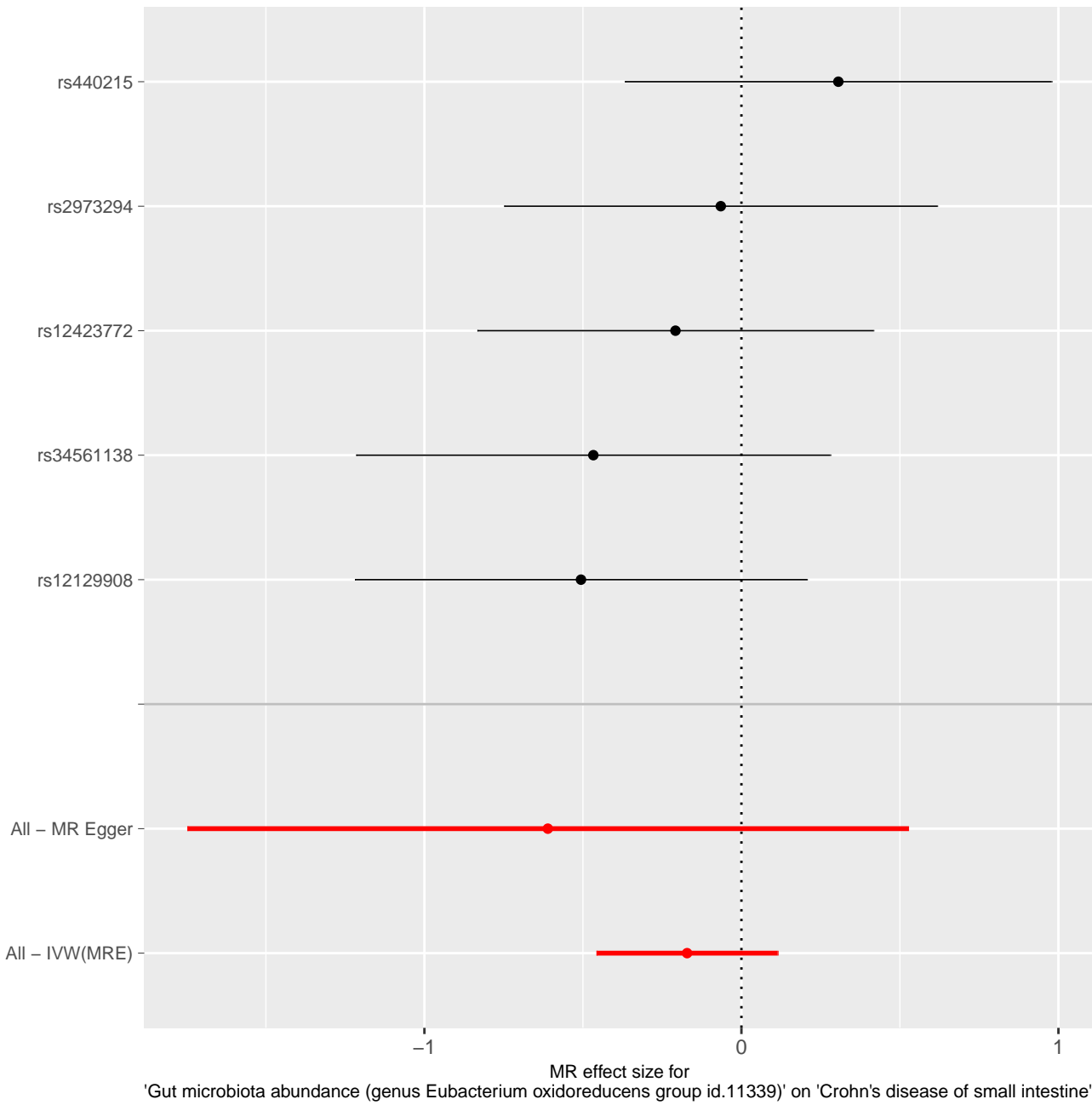


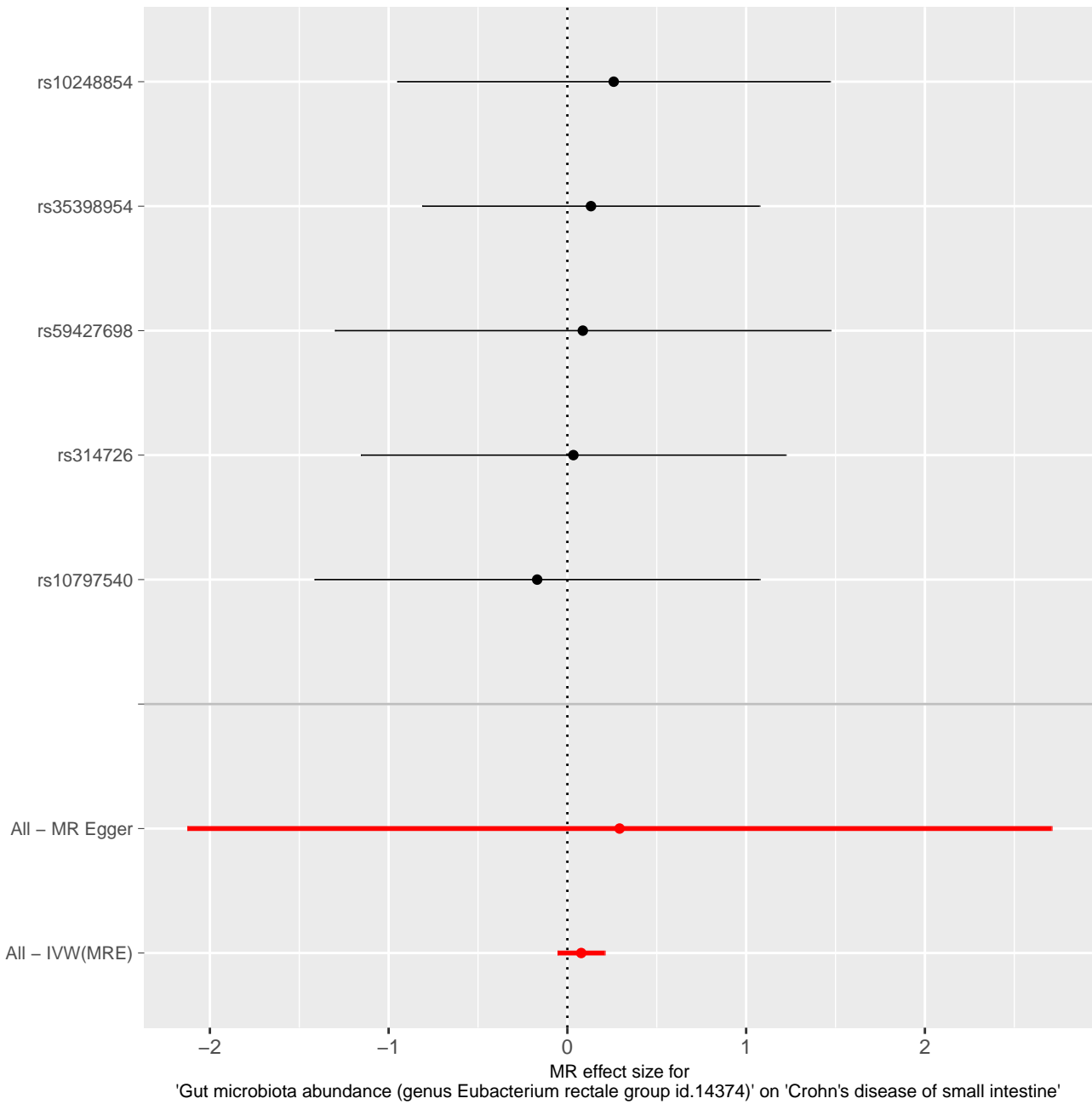


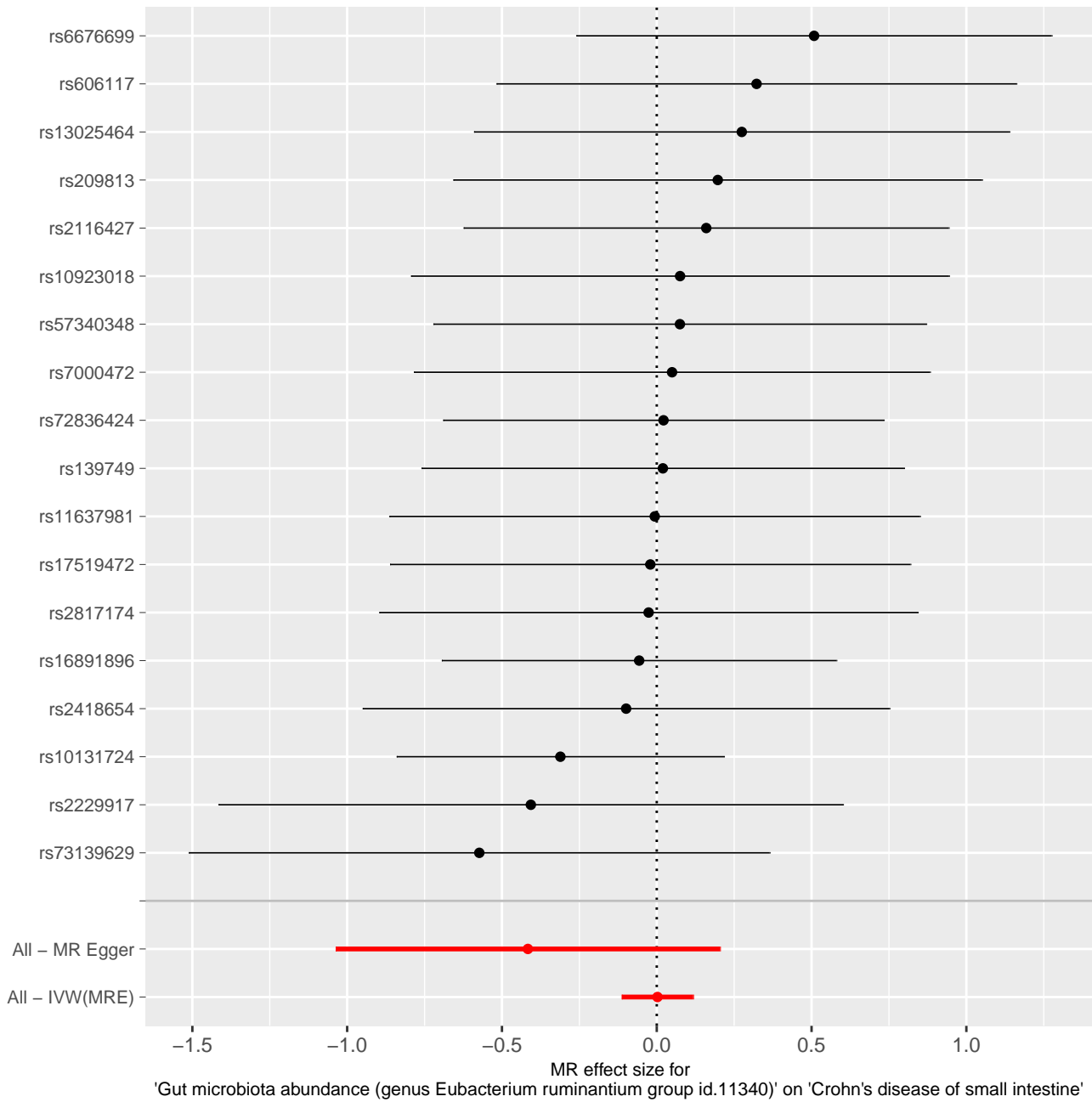


Batch 375 : Gut microbiota abundance (genus Eubacterium nodatum group id.11297) on Crohn's disease of small intestine

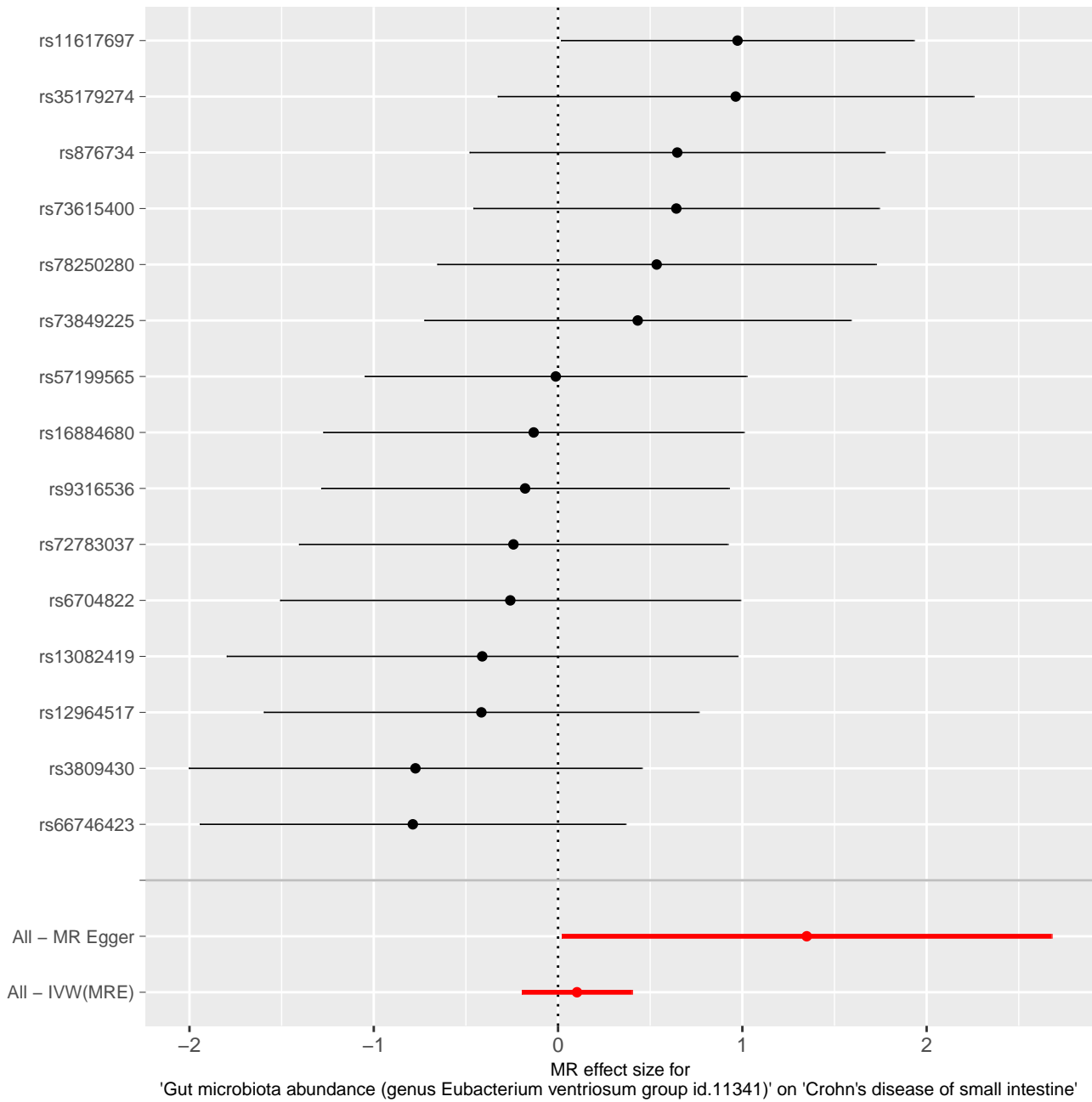


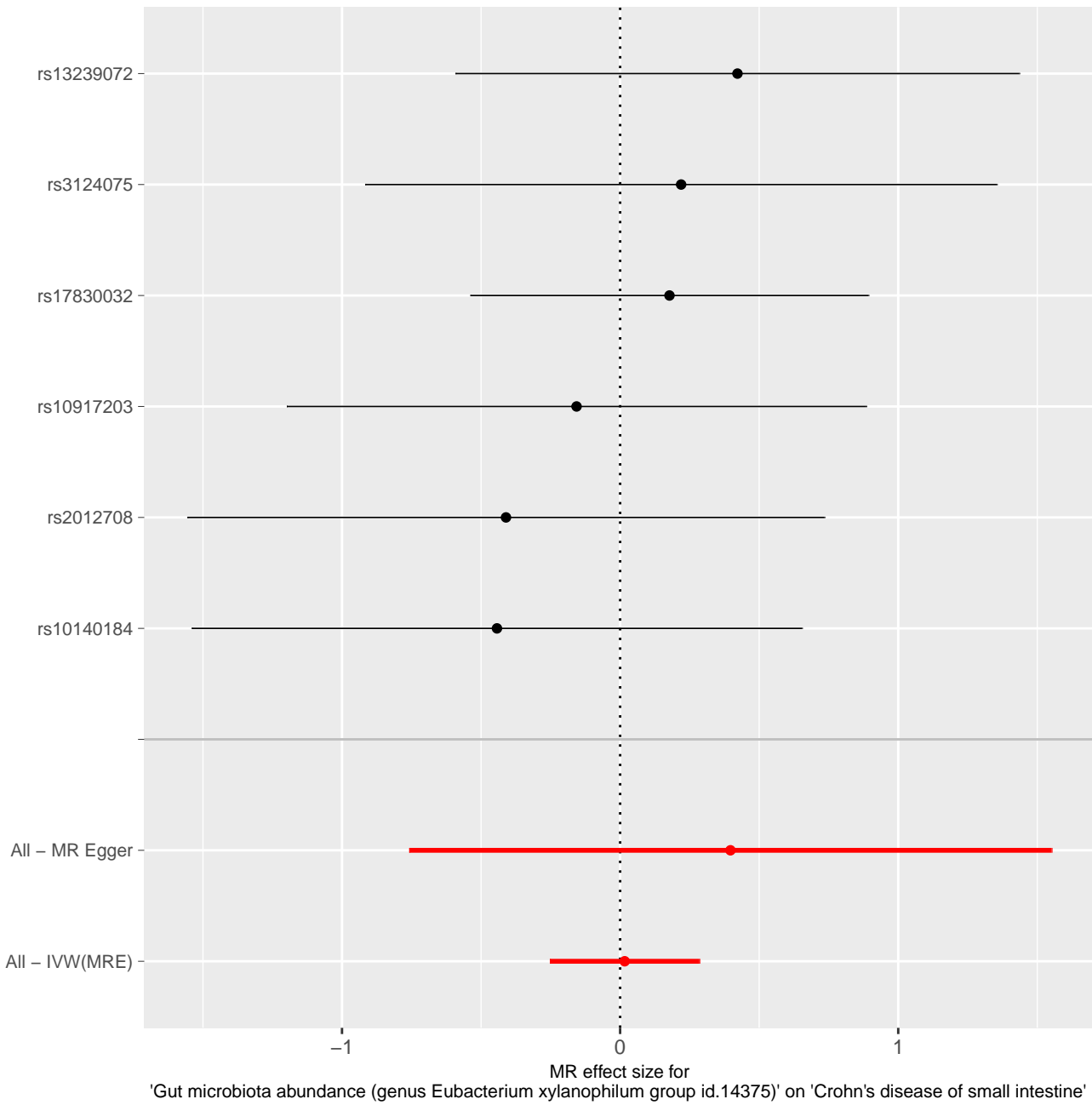




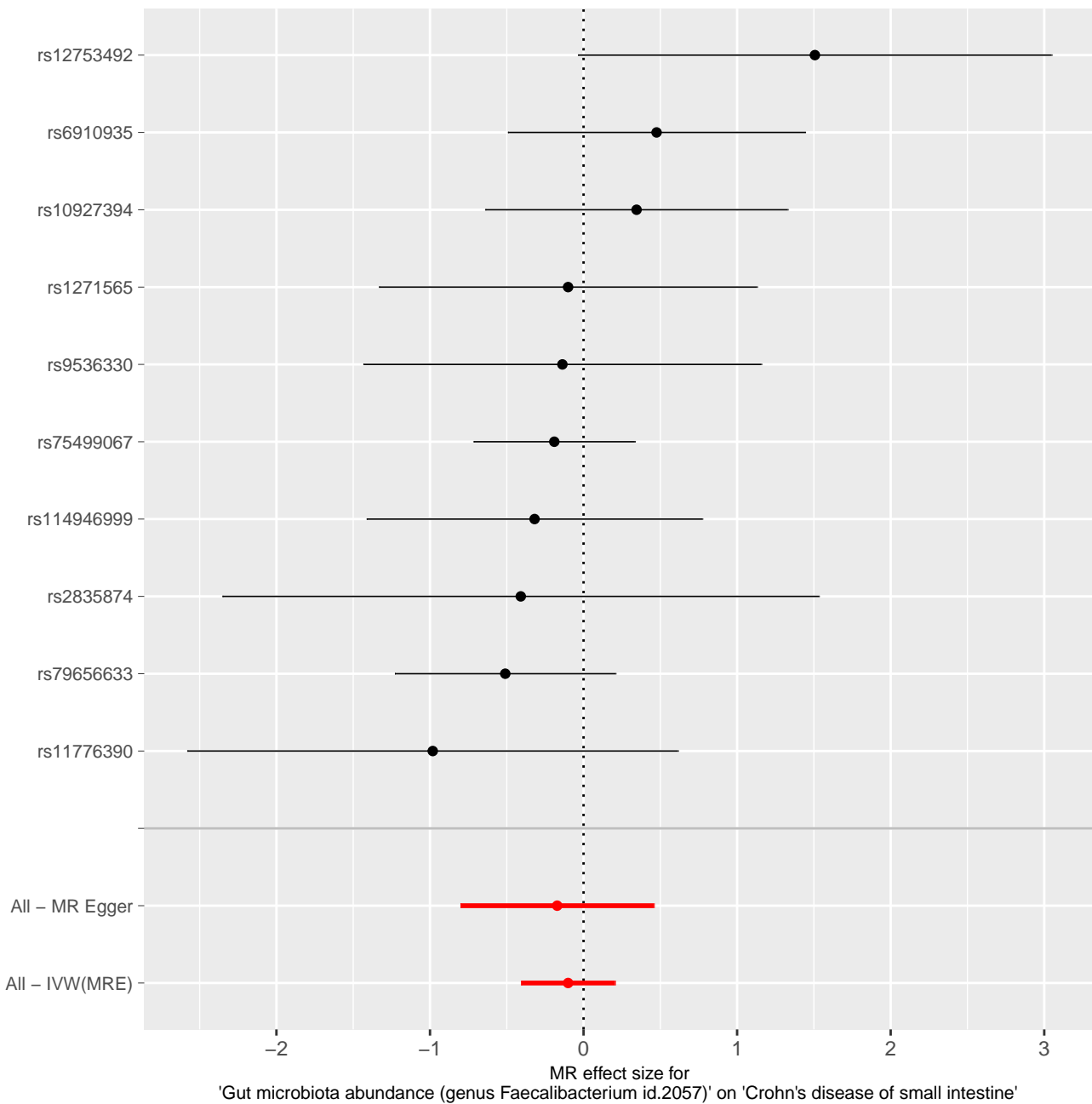


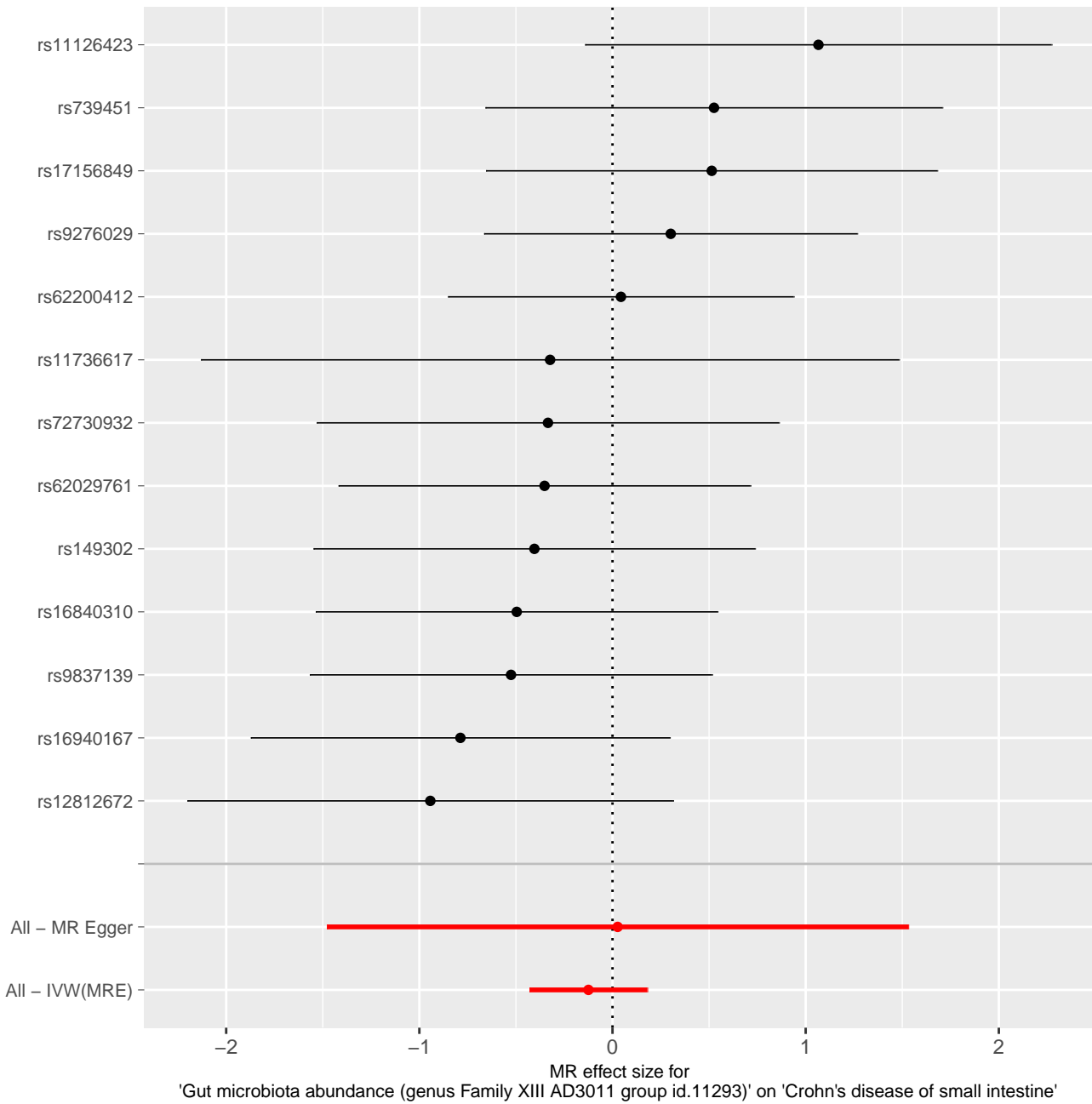
Batch 379 : Gut microbiota abundance (genus Eubacterium ventriosum group id.11341) on Crohn's disease of small intestine



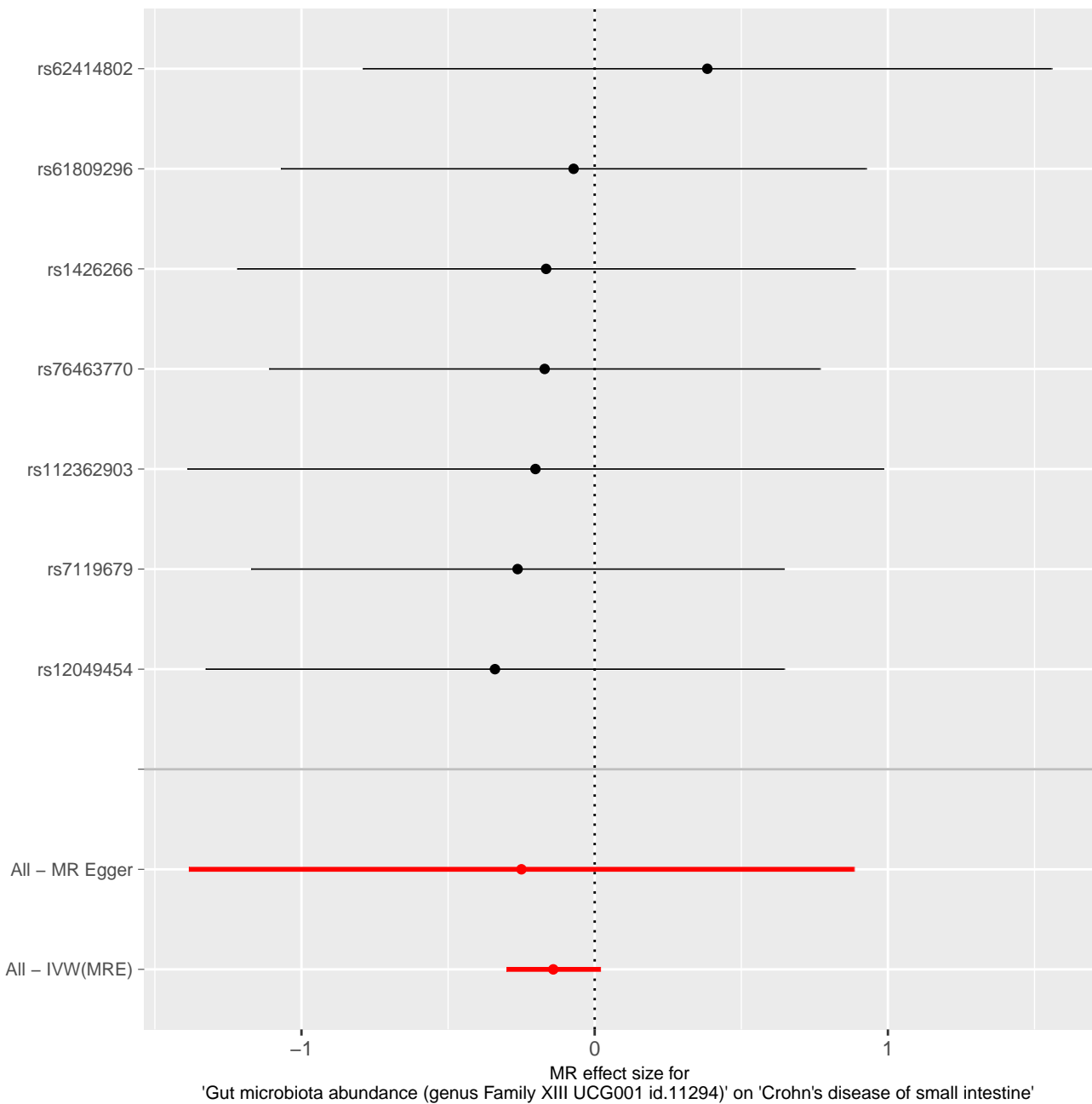


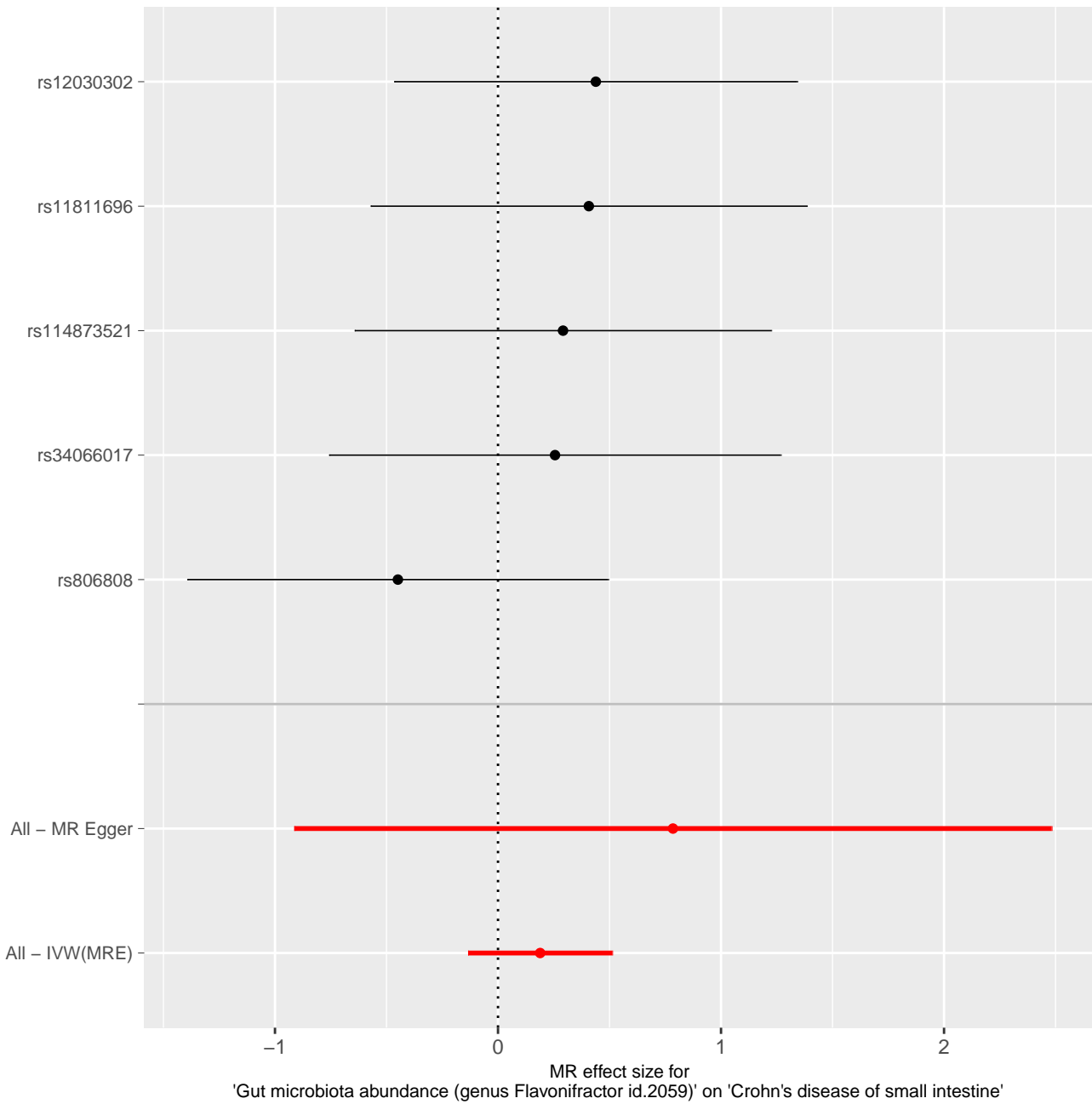
Batch 381 : Gut microbiota abundance (genus Faecalibacterium id.2057) on Crohn's disease of small intestine



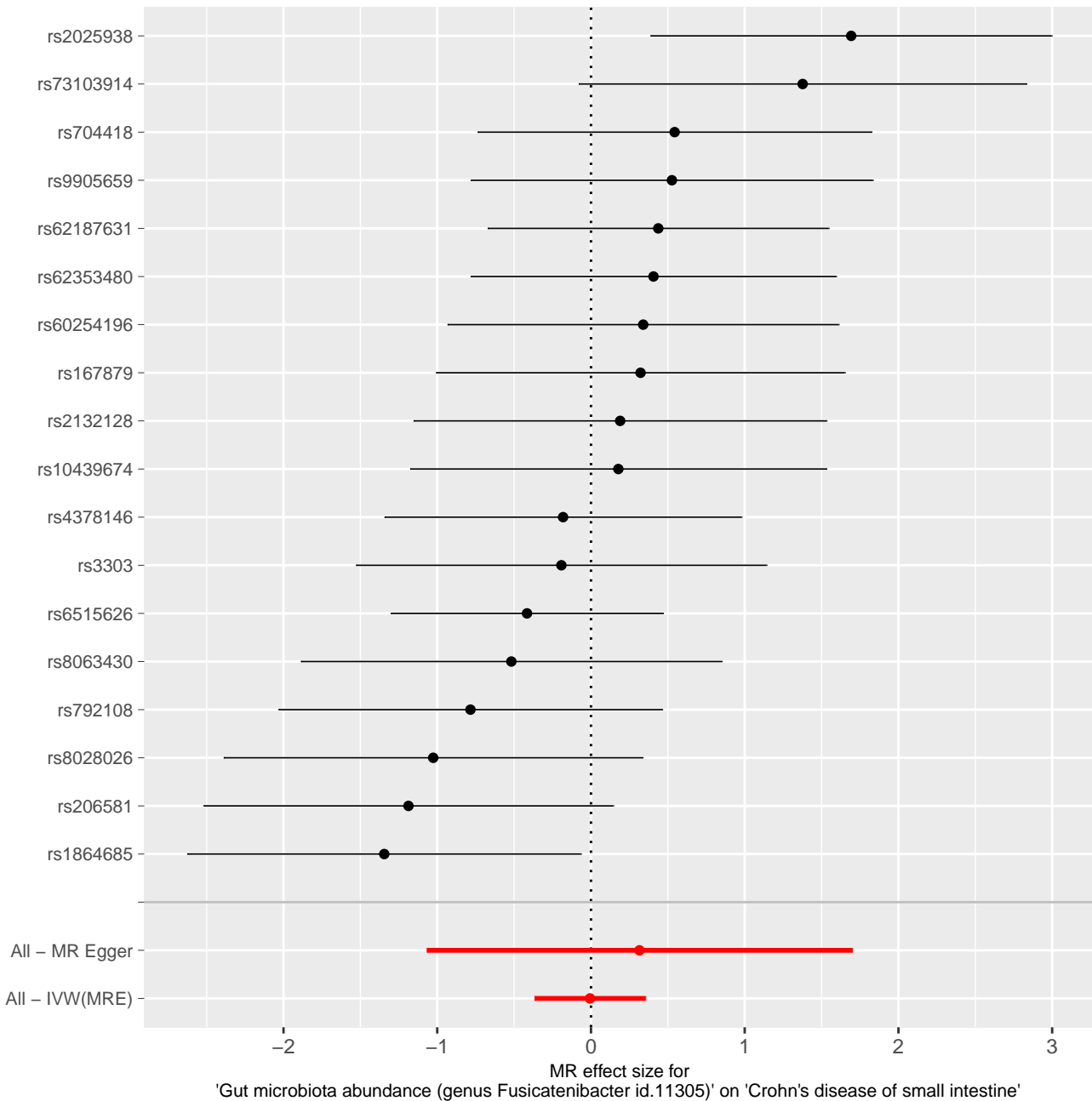


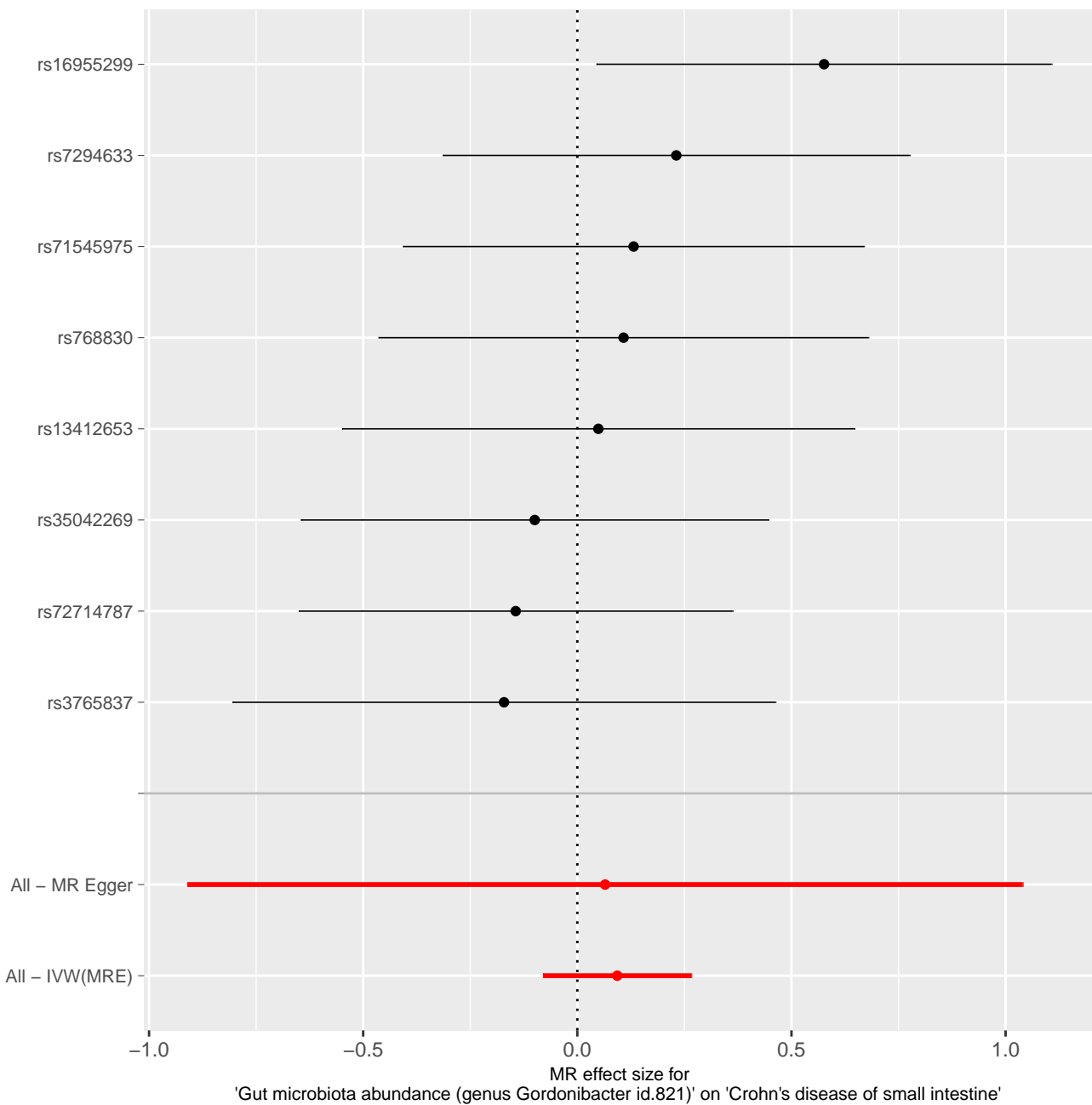
MR effect size for
'Gut microbiota abundance (genus Family XIII AD3011 group id.11293)' on 'Crohn's disease of small intestine'

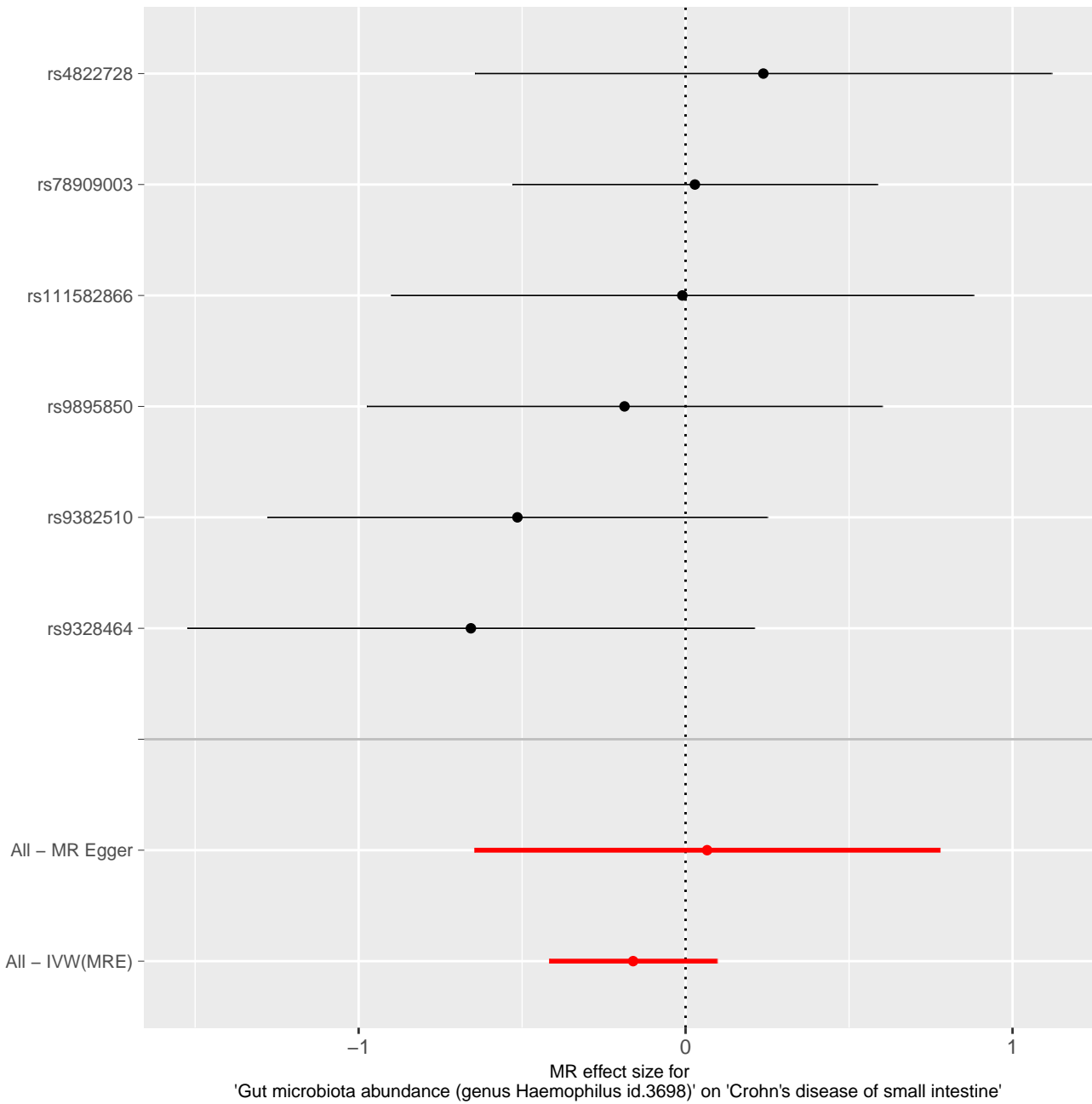


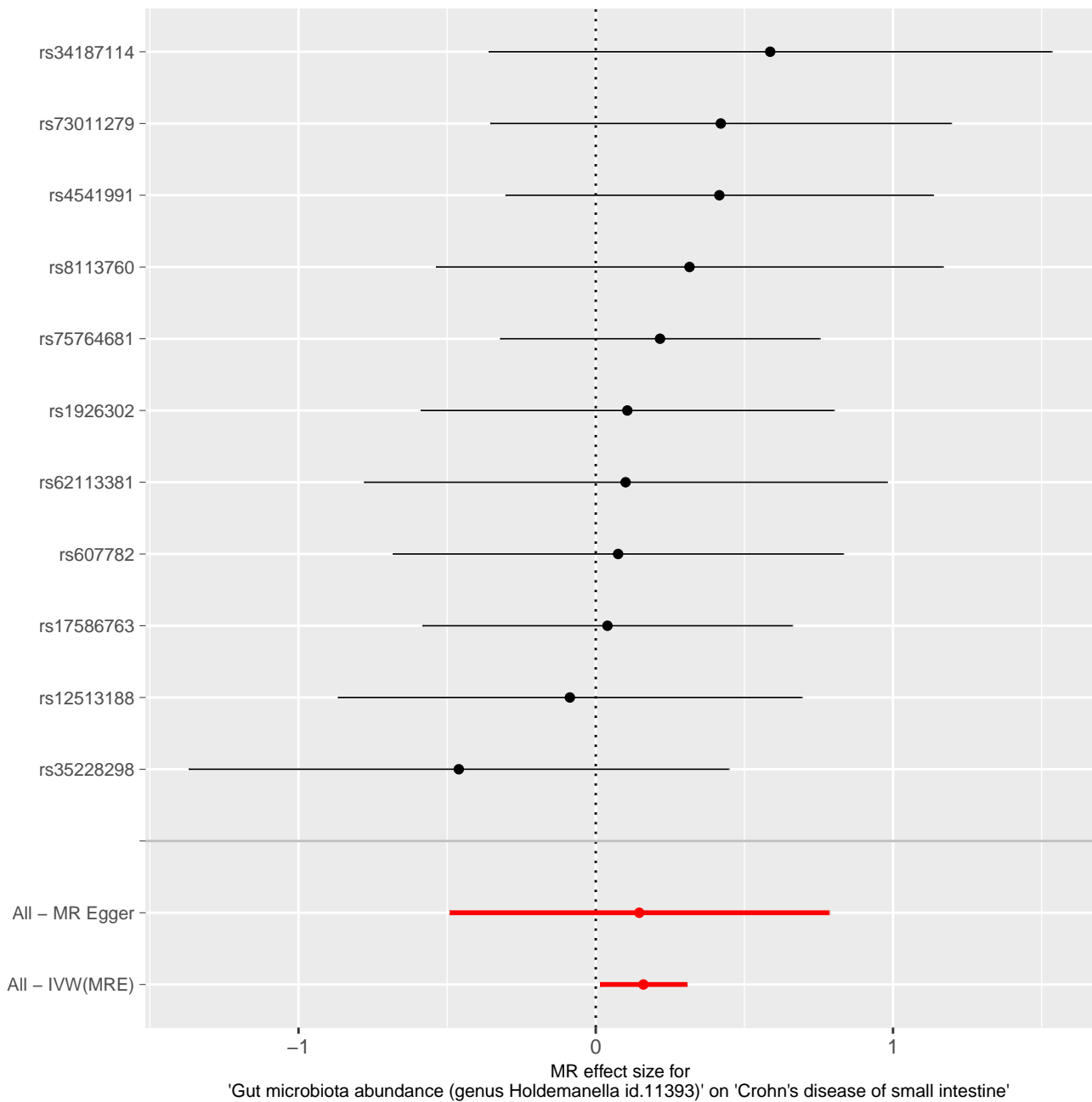


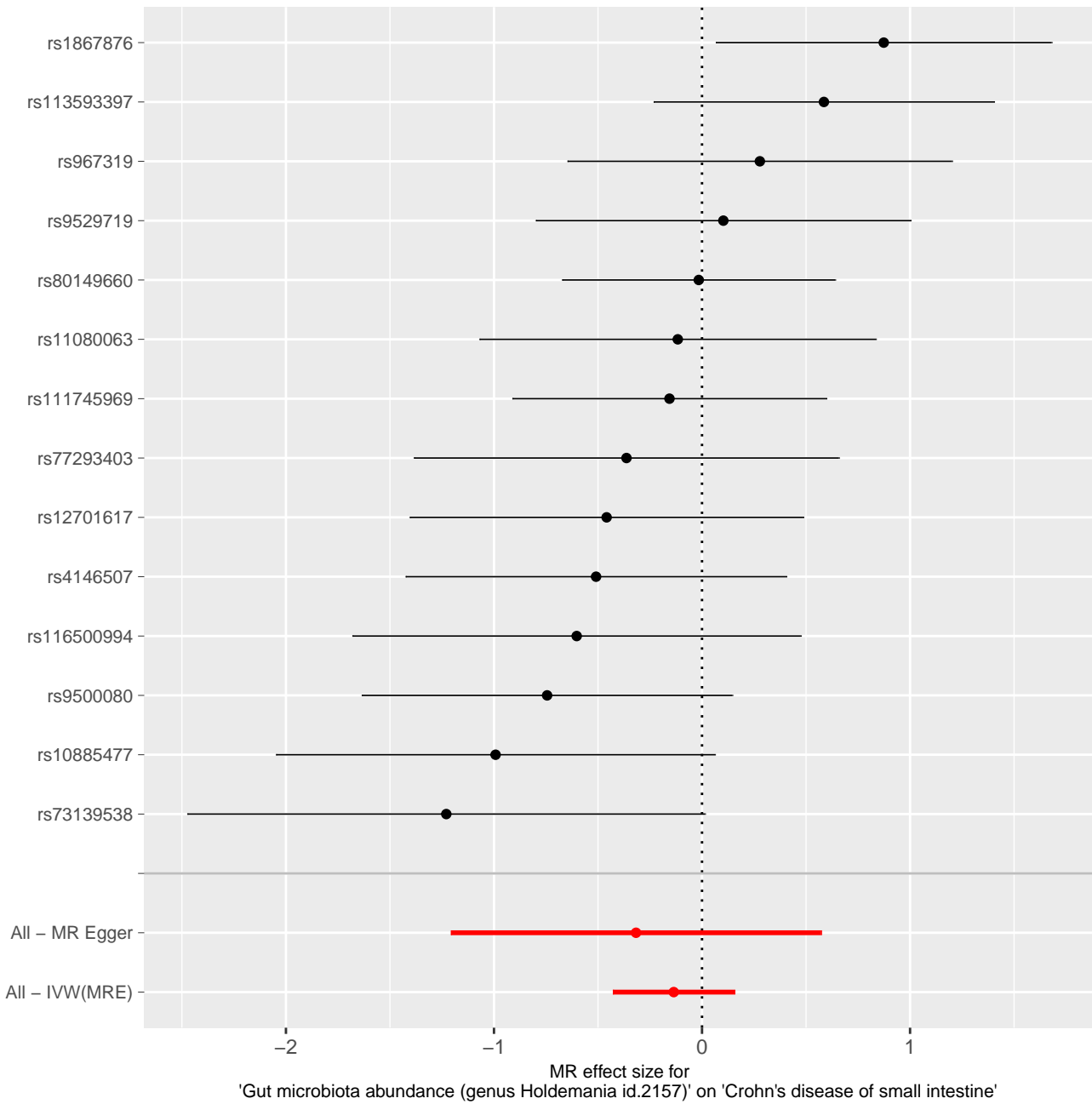
Batch 385 : Gut microbiota abundance (genus Fusicatenibacter id.11305) on Crohn's disease of small intestine



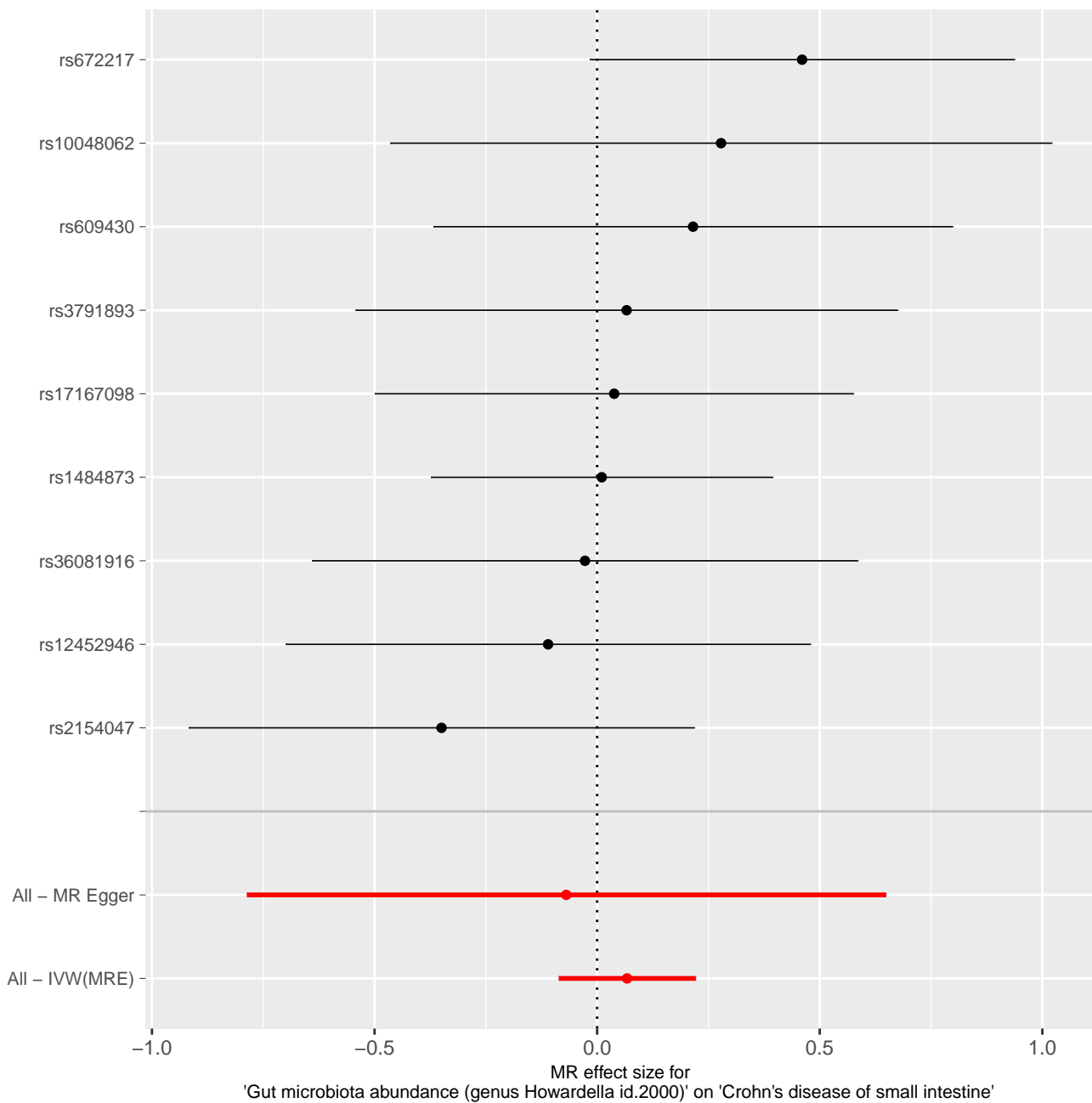


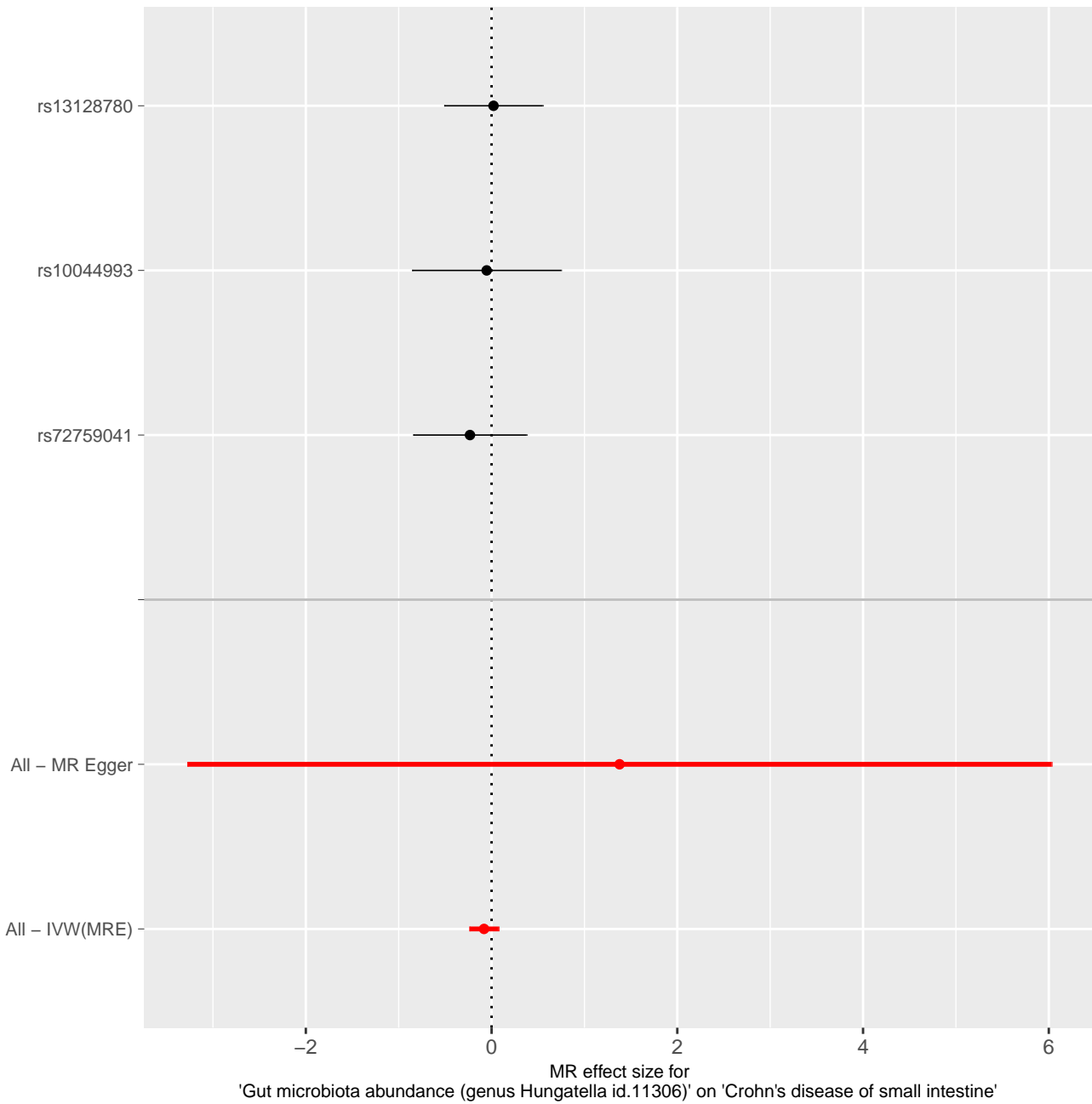


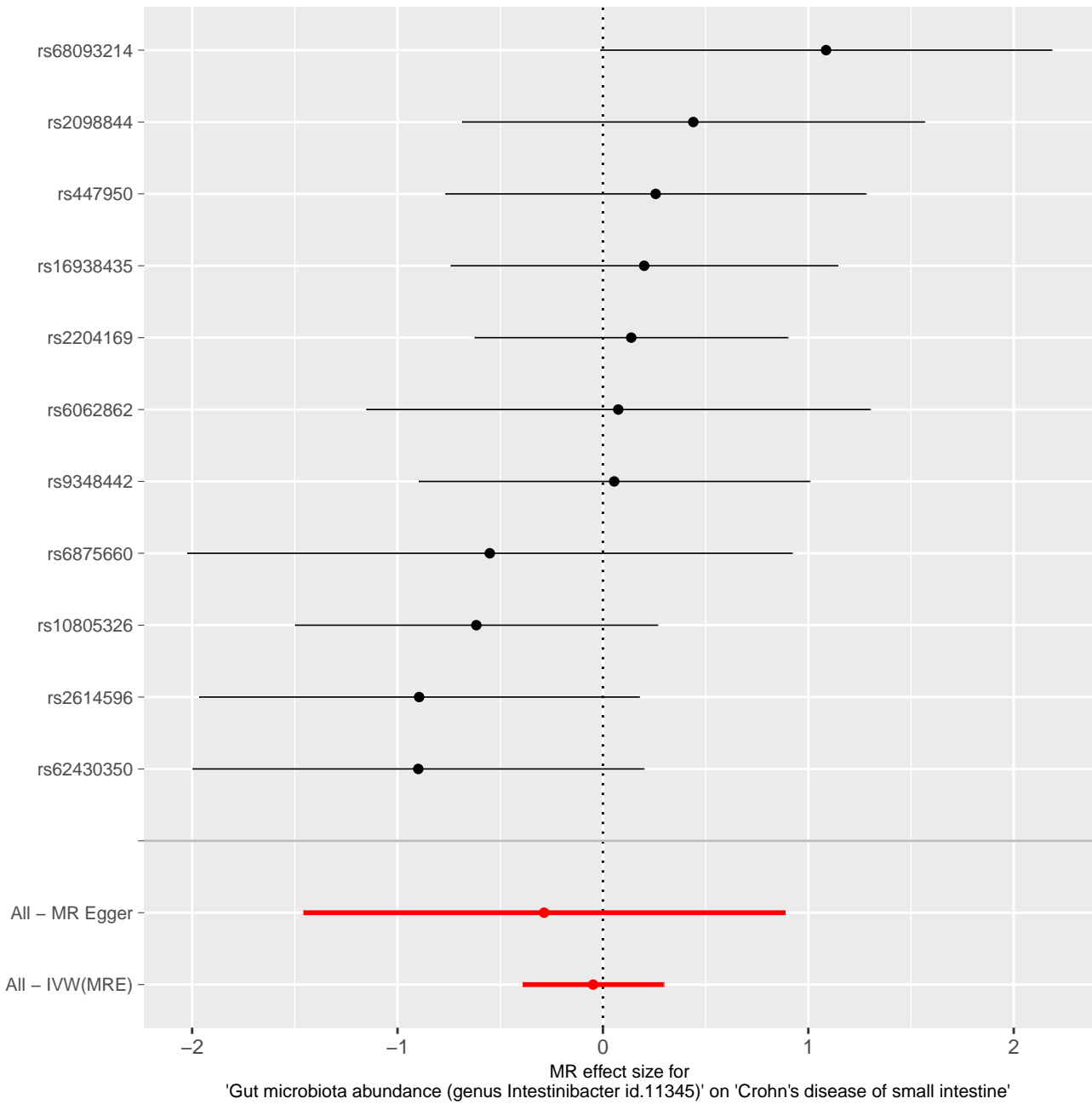


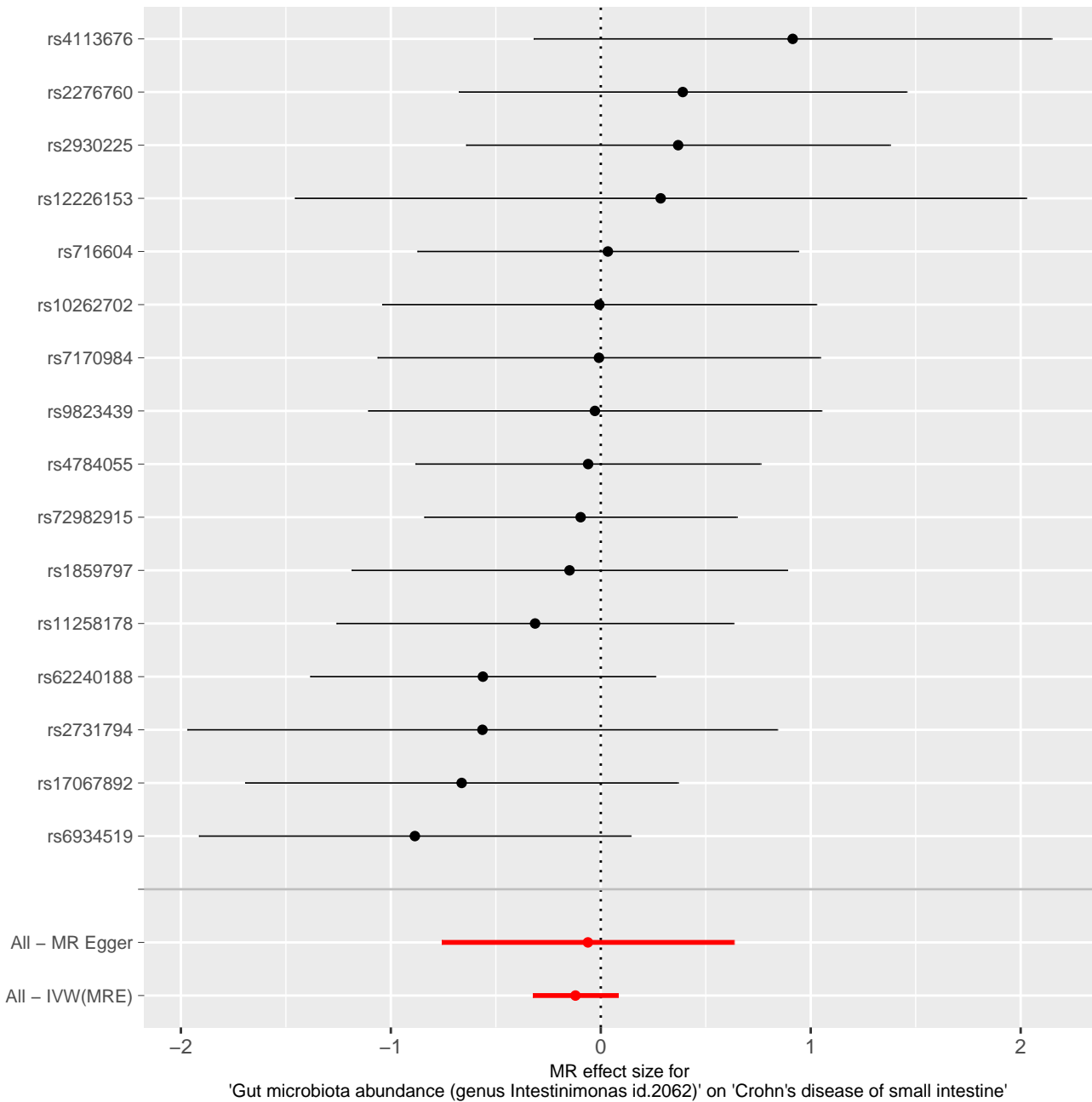


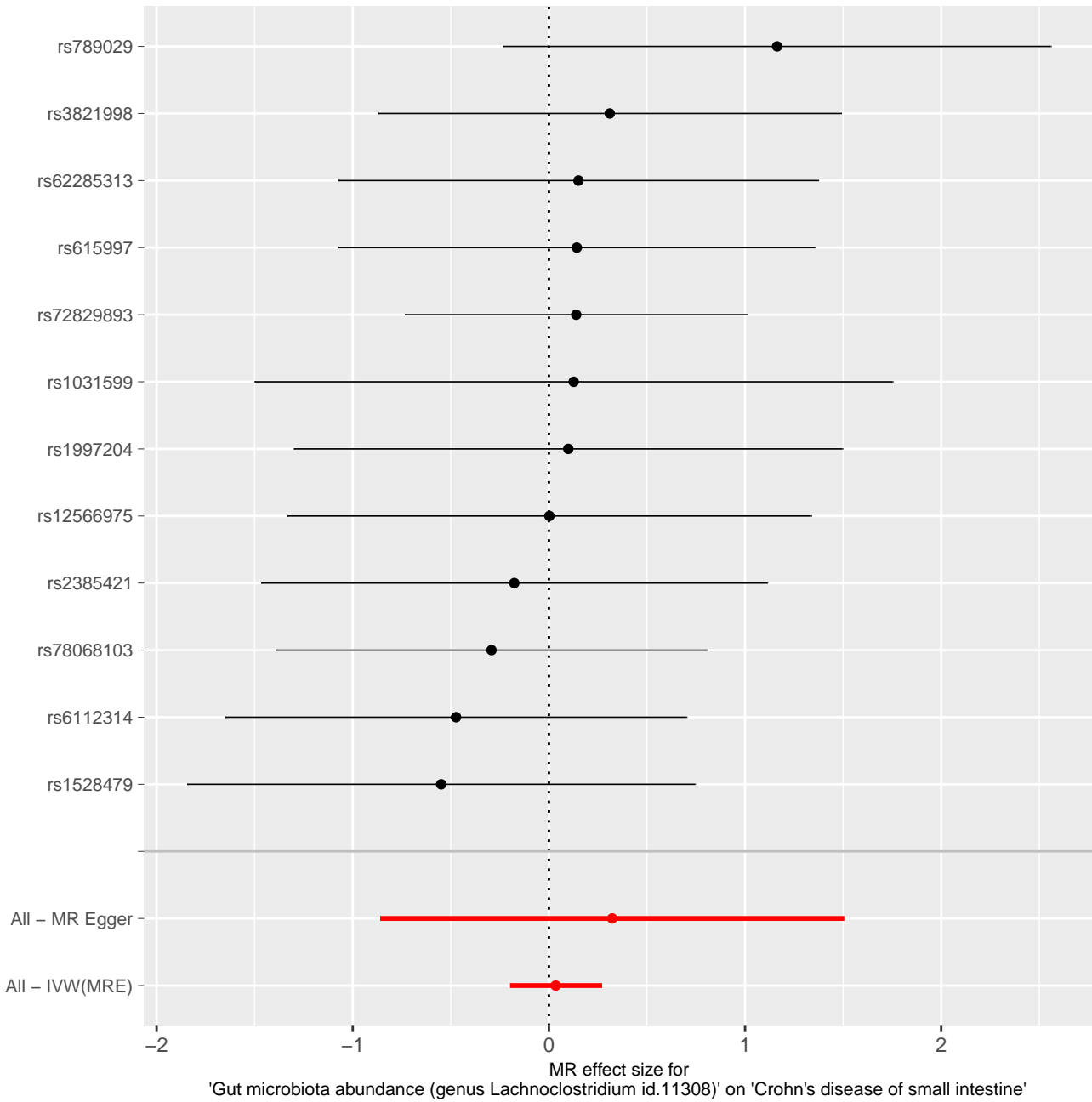
Batch 390 : Gut microbiota abundance (genus Howardella id.2000) on Crohn's disease of small intestine



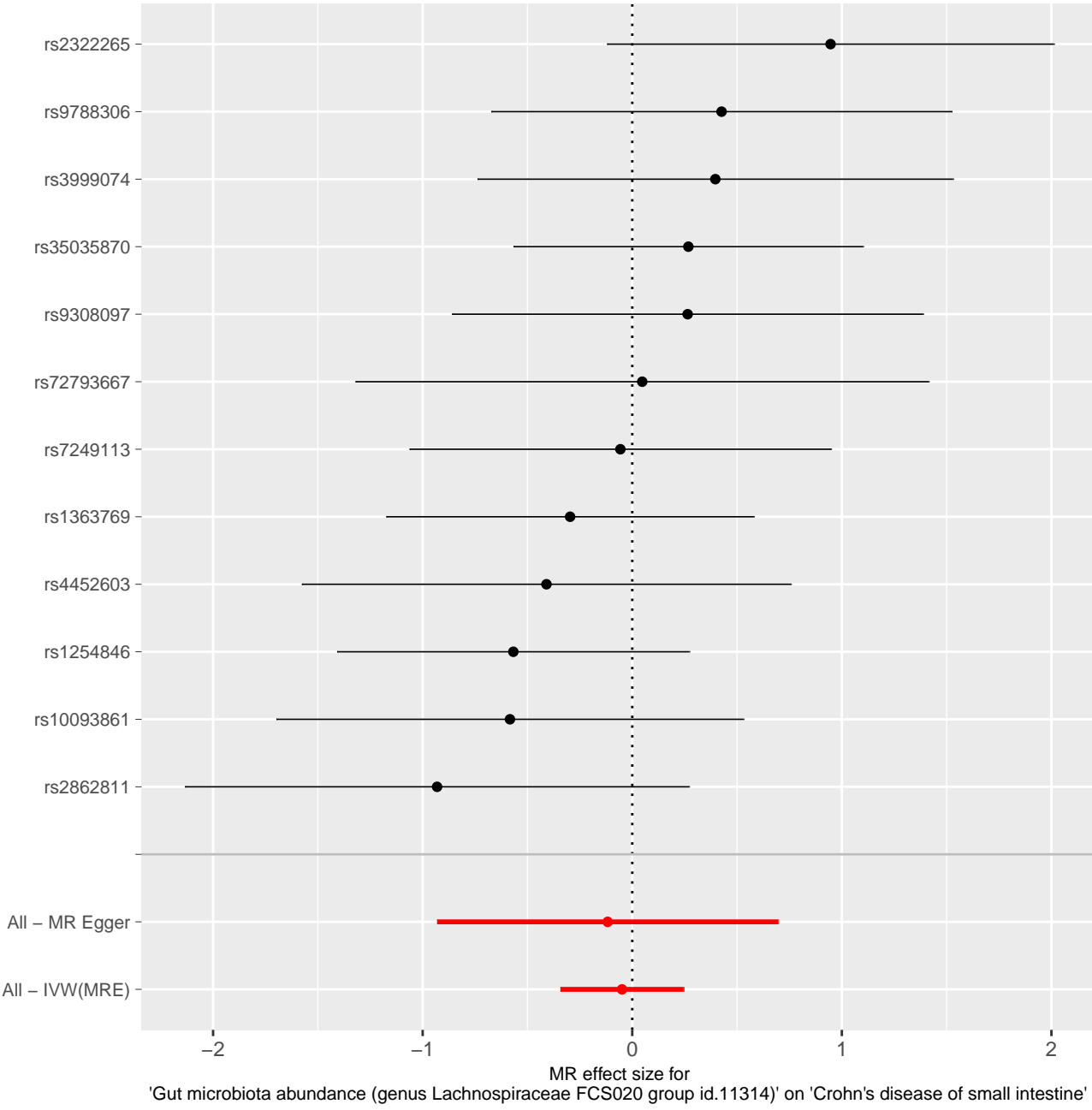


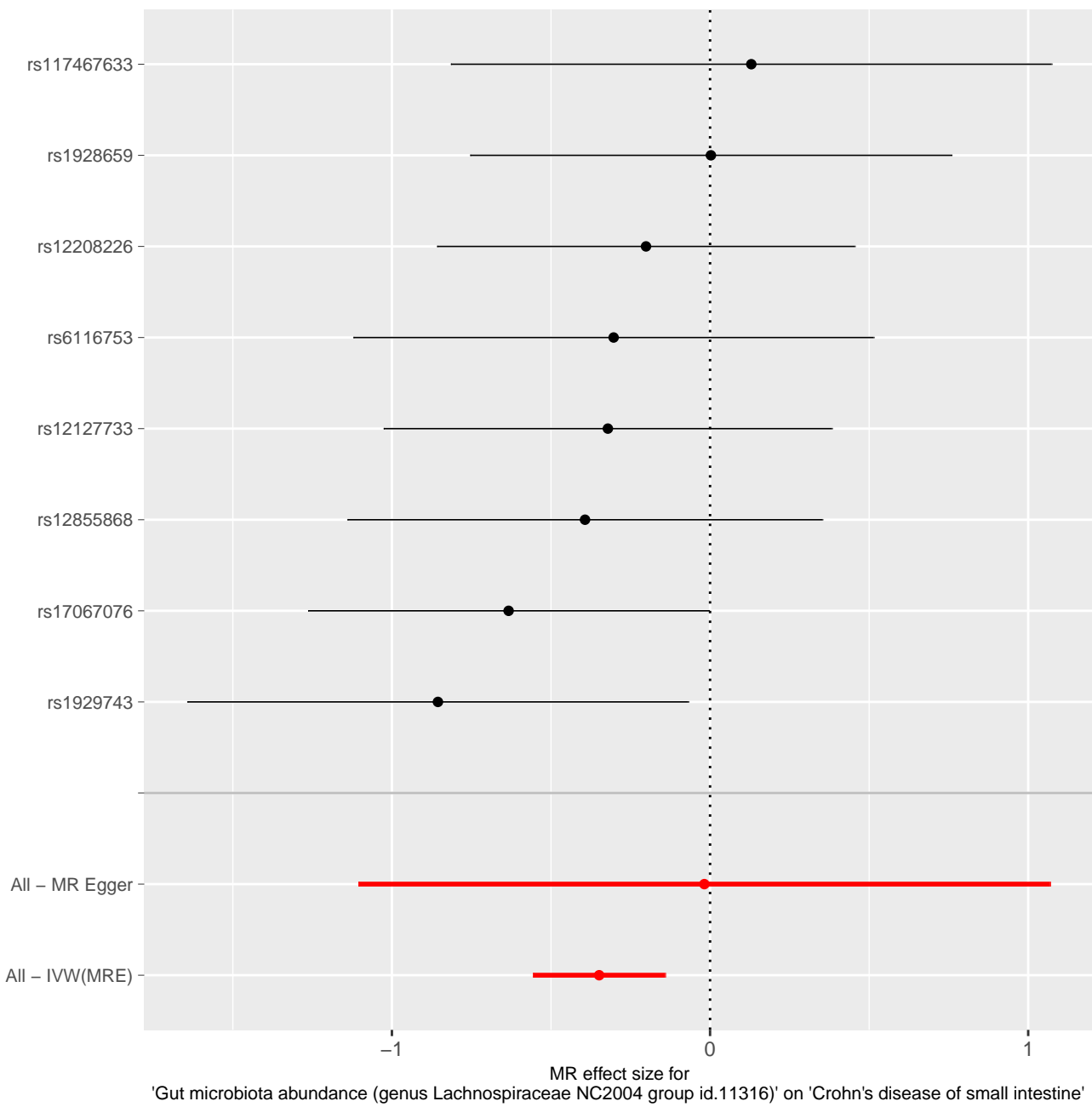




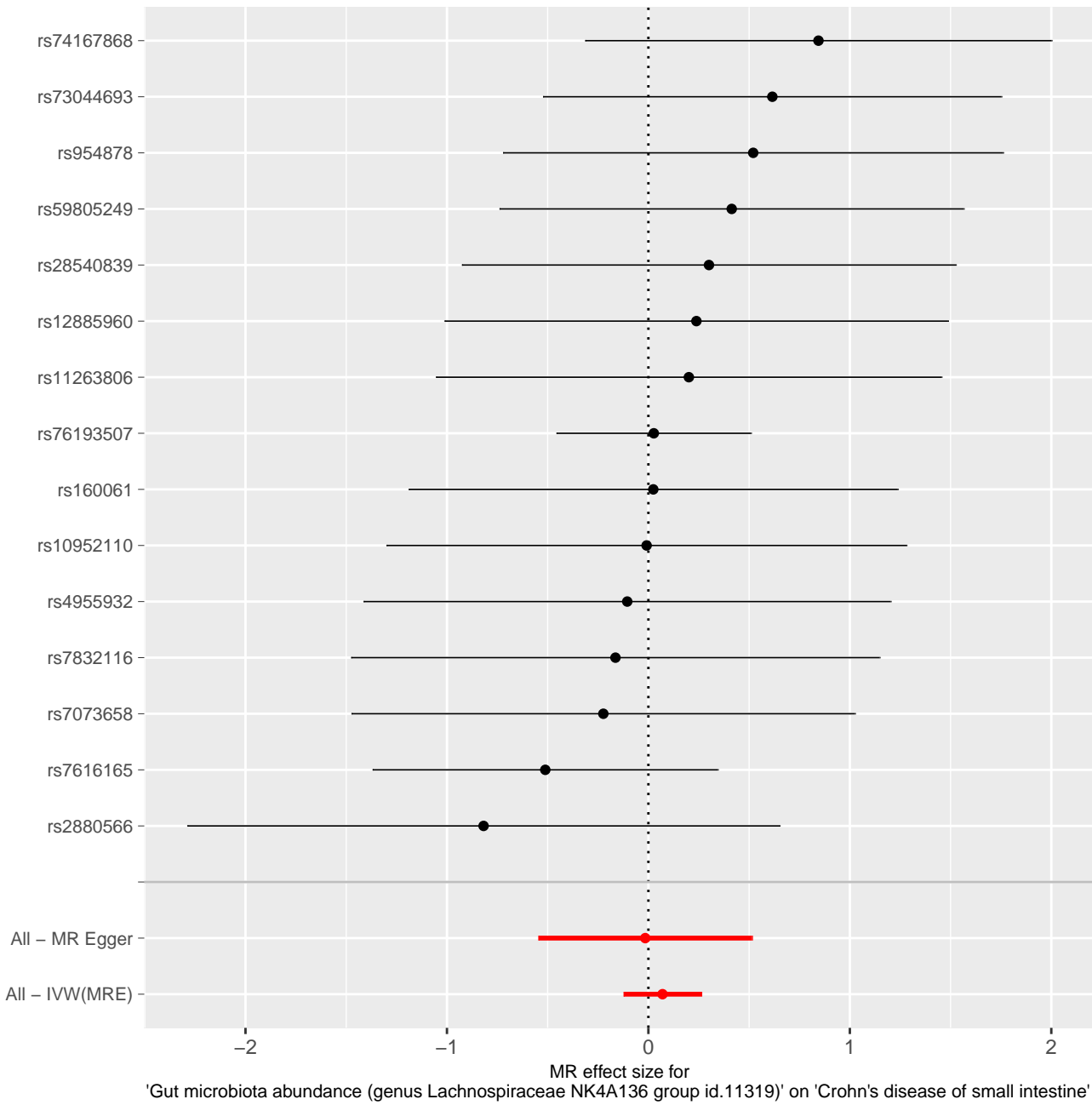


Batch 395 : Gut microbiota abundance (genus Lachnospiraceae FCS020 group id.11314) on Crohn's disease of small intestine

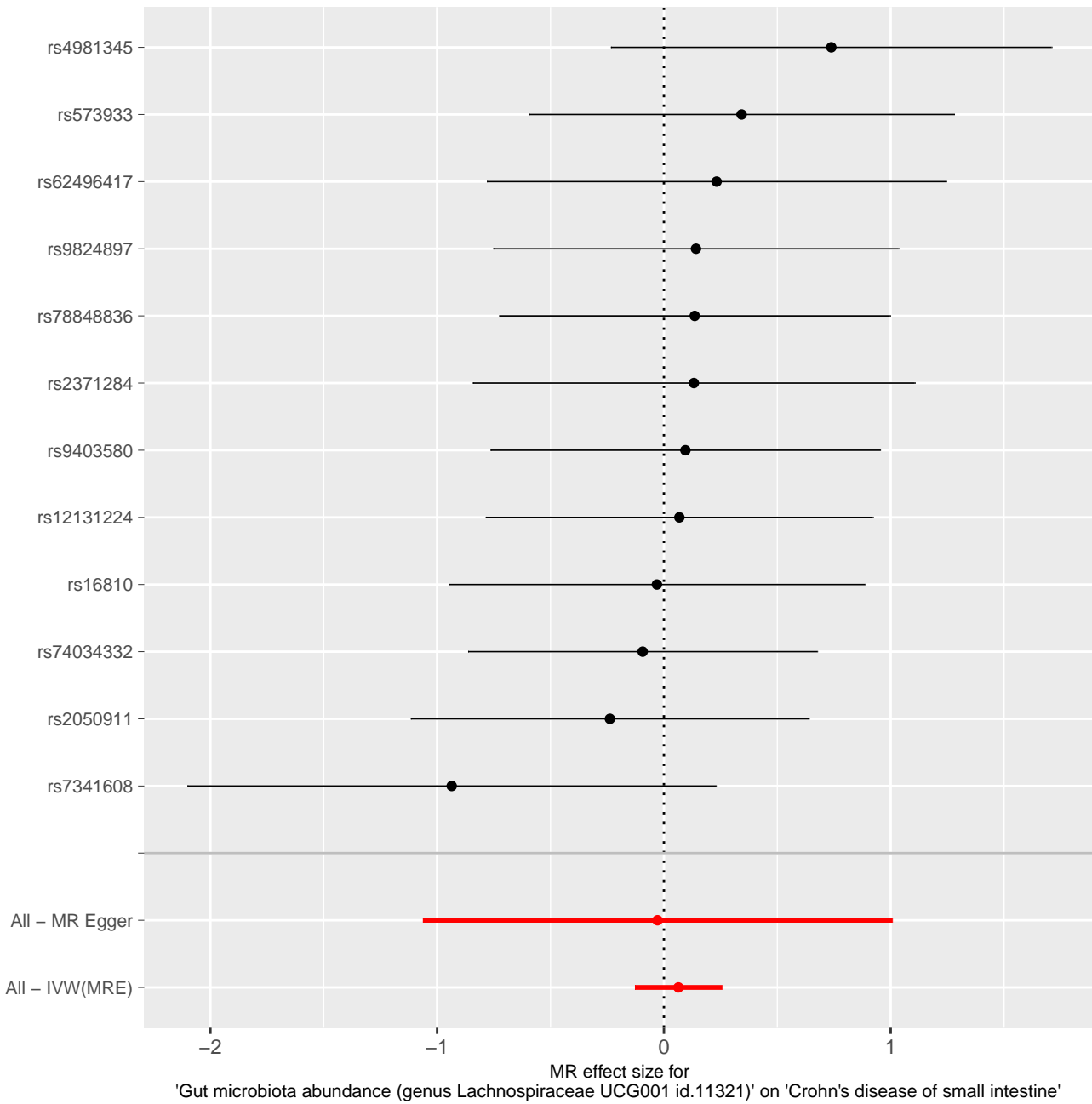


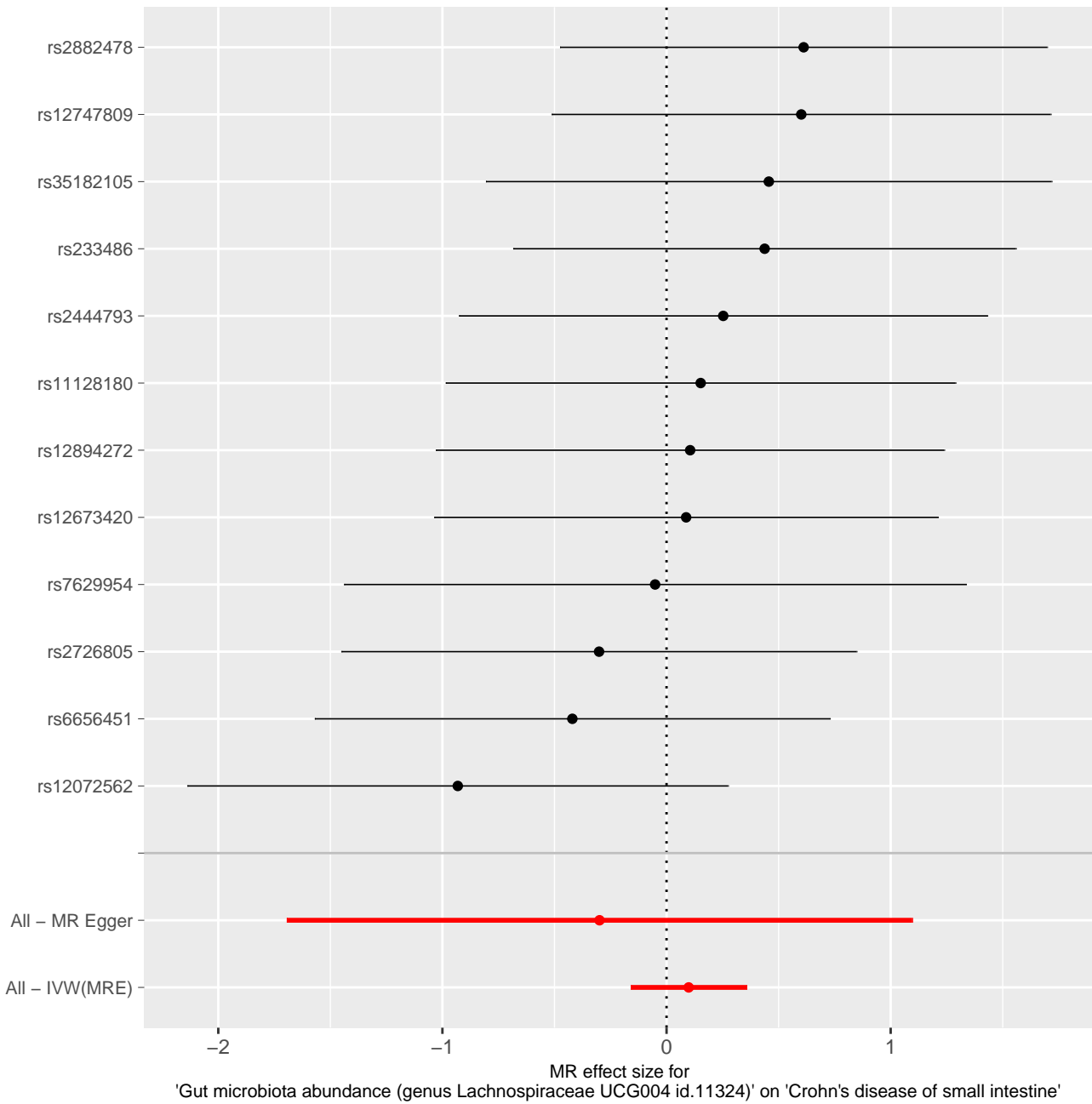


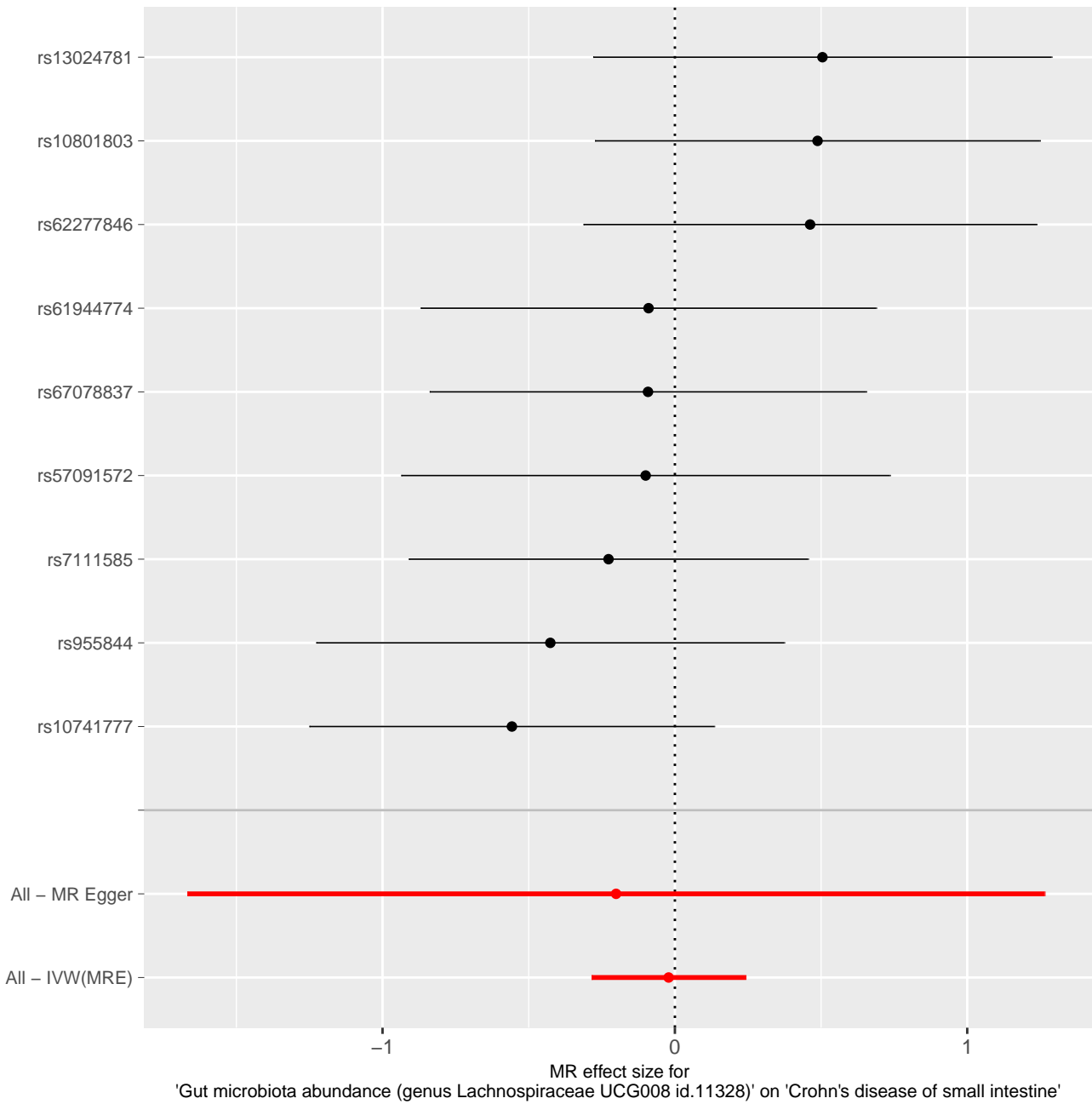
Batch 398 : Gut microbiota abundance (genus Lachnospiraceae NK4A136 group id.11319) on Crohn's disease of small intestine

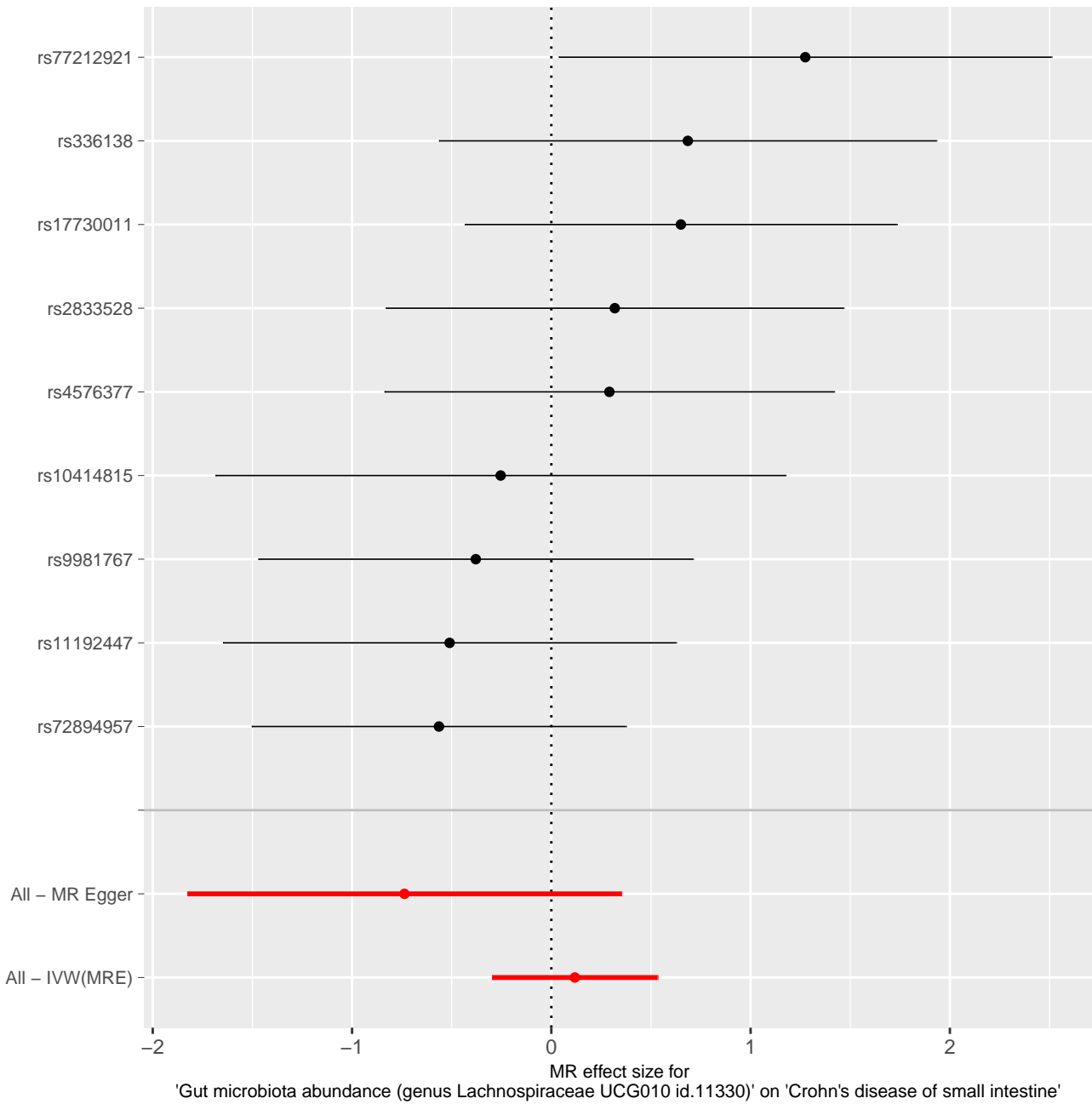


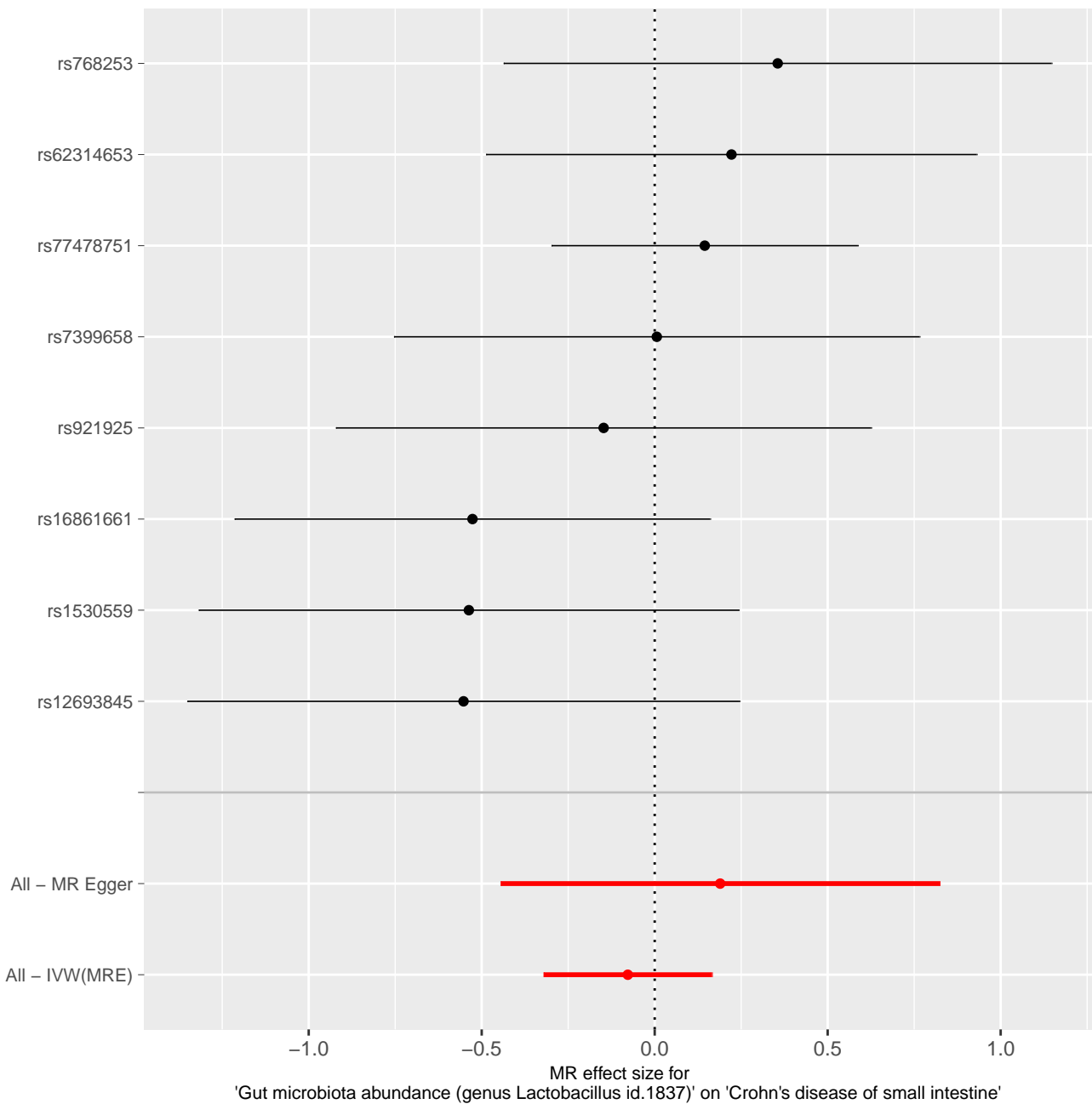
Batch 399 : Gut microbiota abundance (genus Lachnospiraceae UCG001 id.11321) on Crohn's disease of small intestine

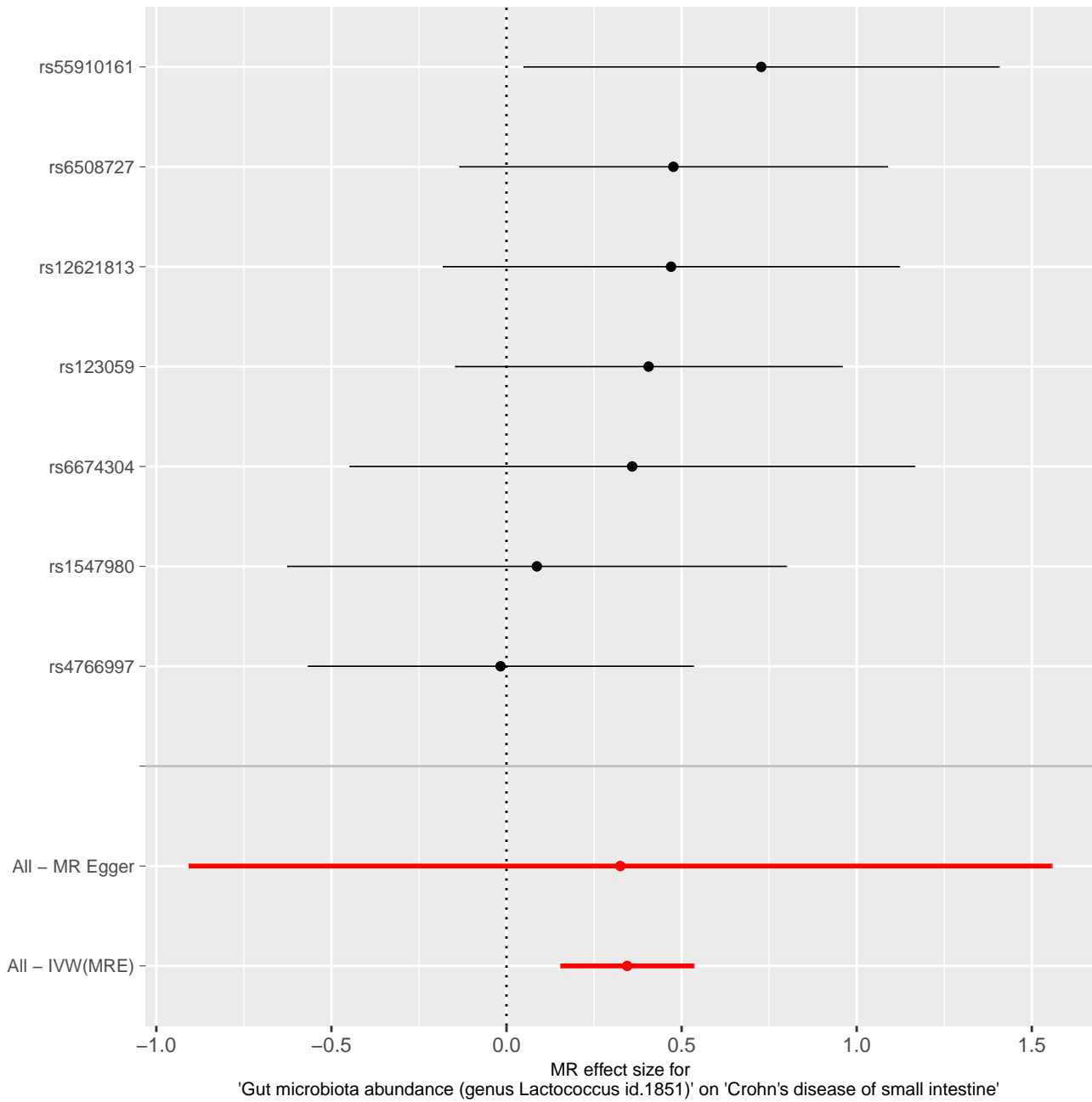




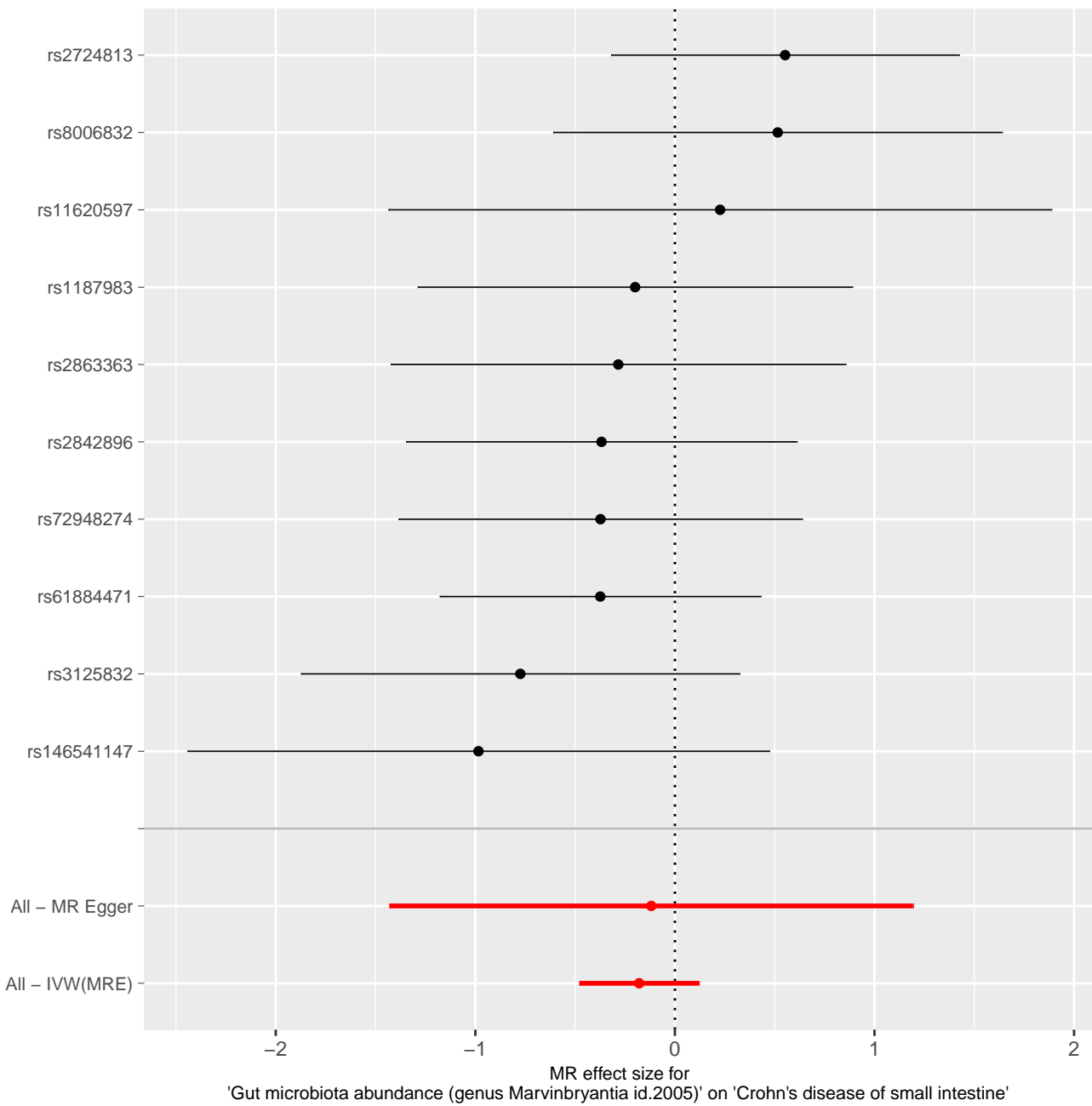


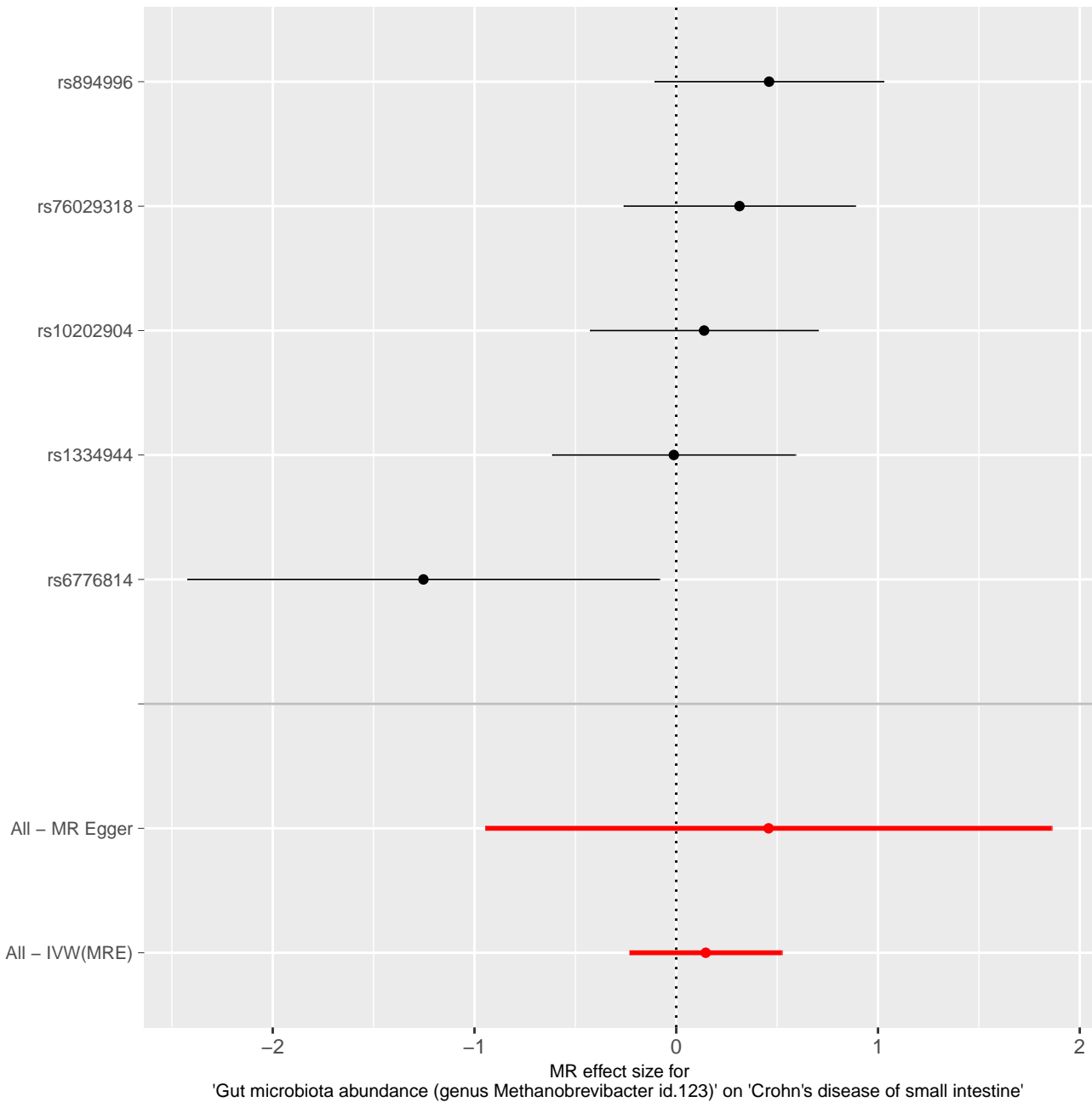




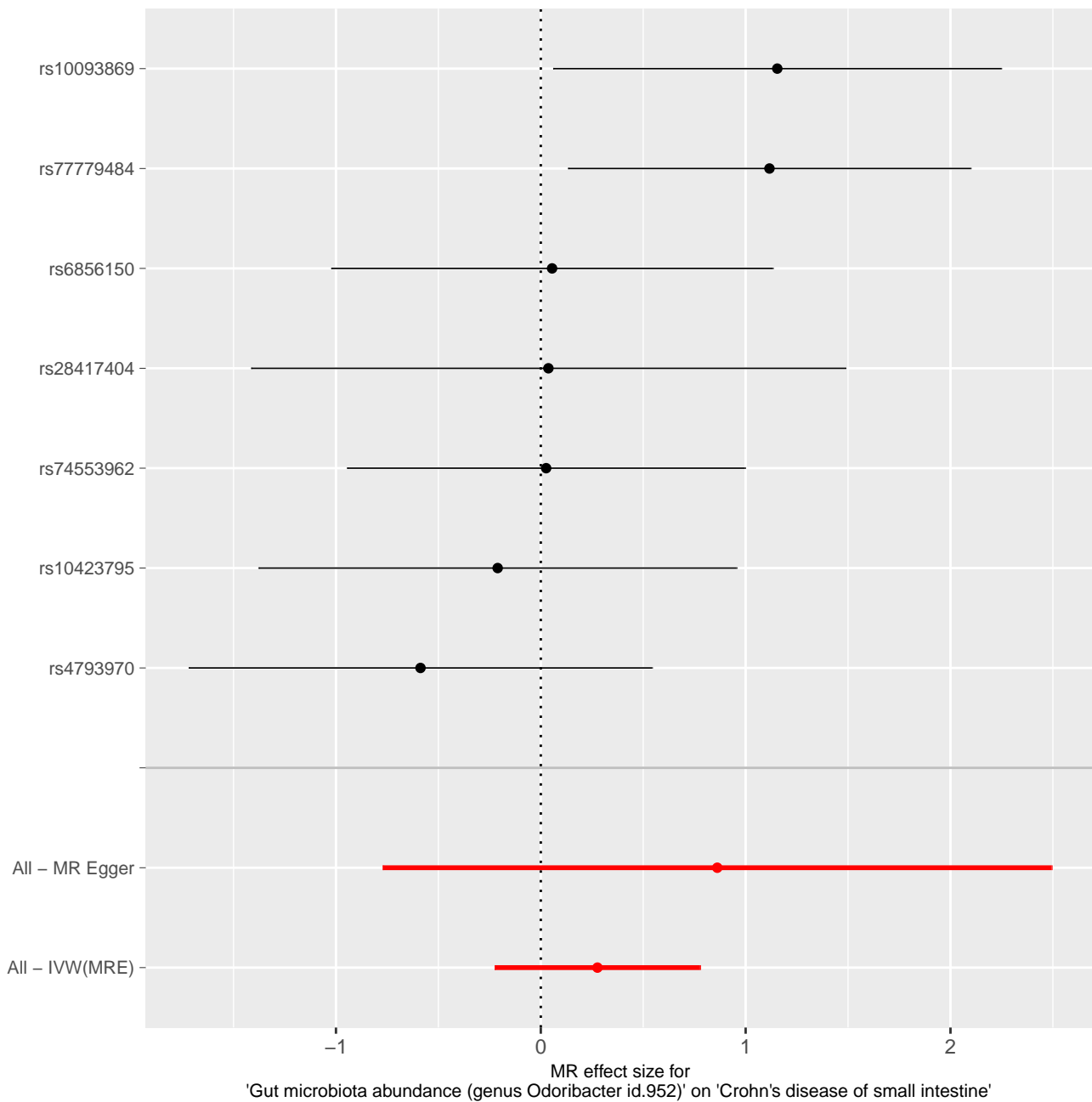


Batch 406 : Gut microbiota abundance (genus Marvinbryantia id.2005) on Crohn's disease of small intestine

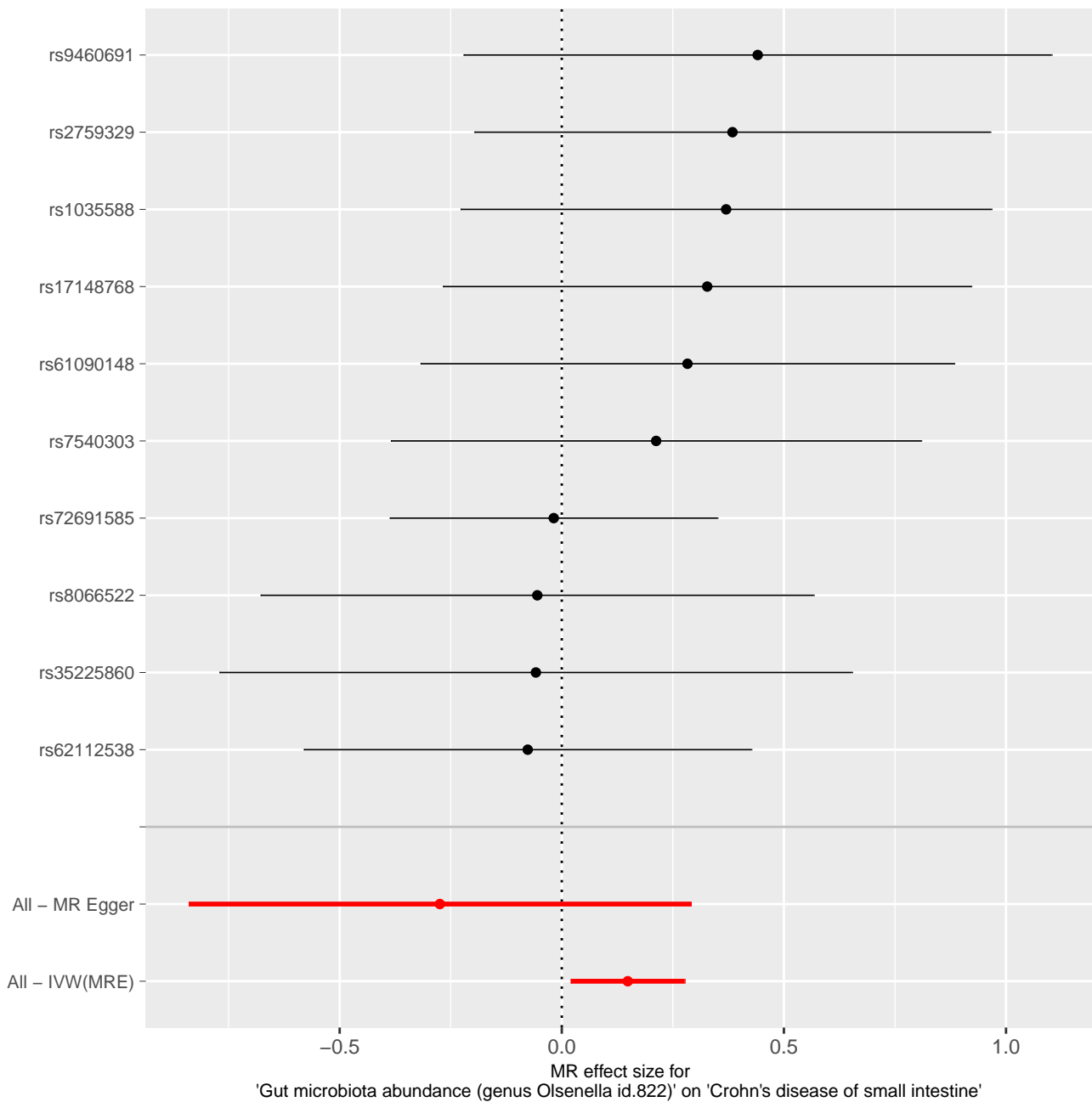


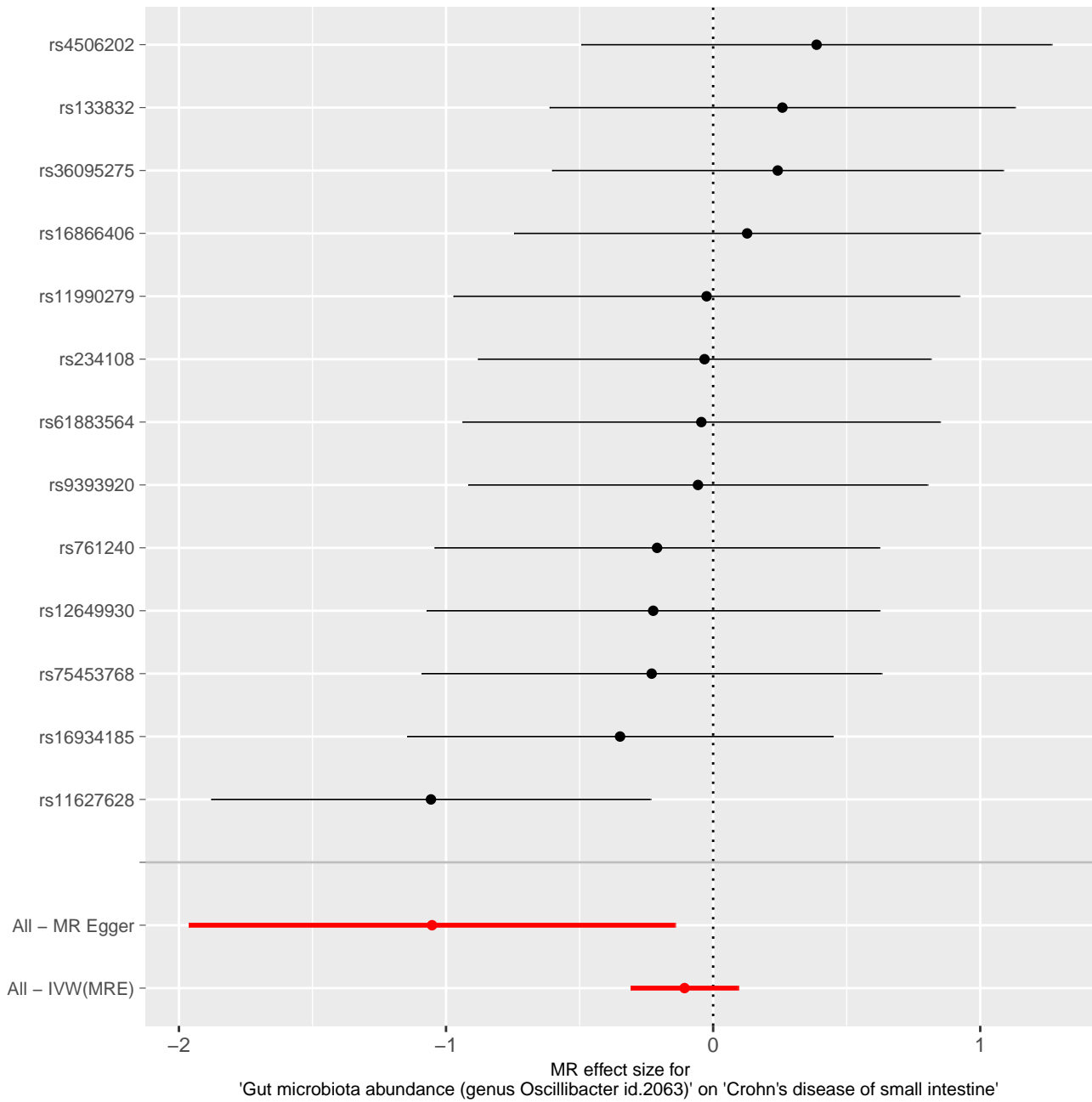


Batch 408 : Gut microbiota abundance (genus Odoribacter id.952) on Crohn's disease of small intestine

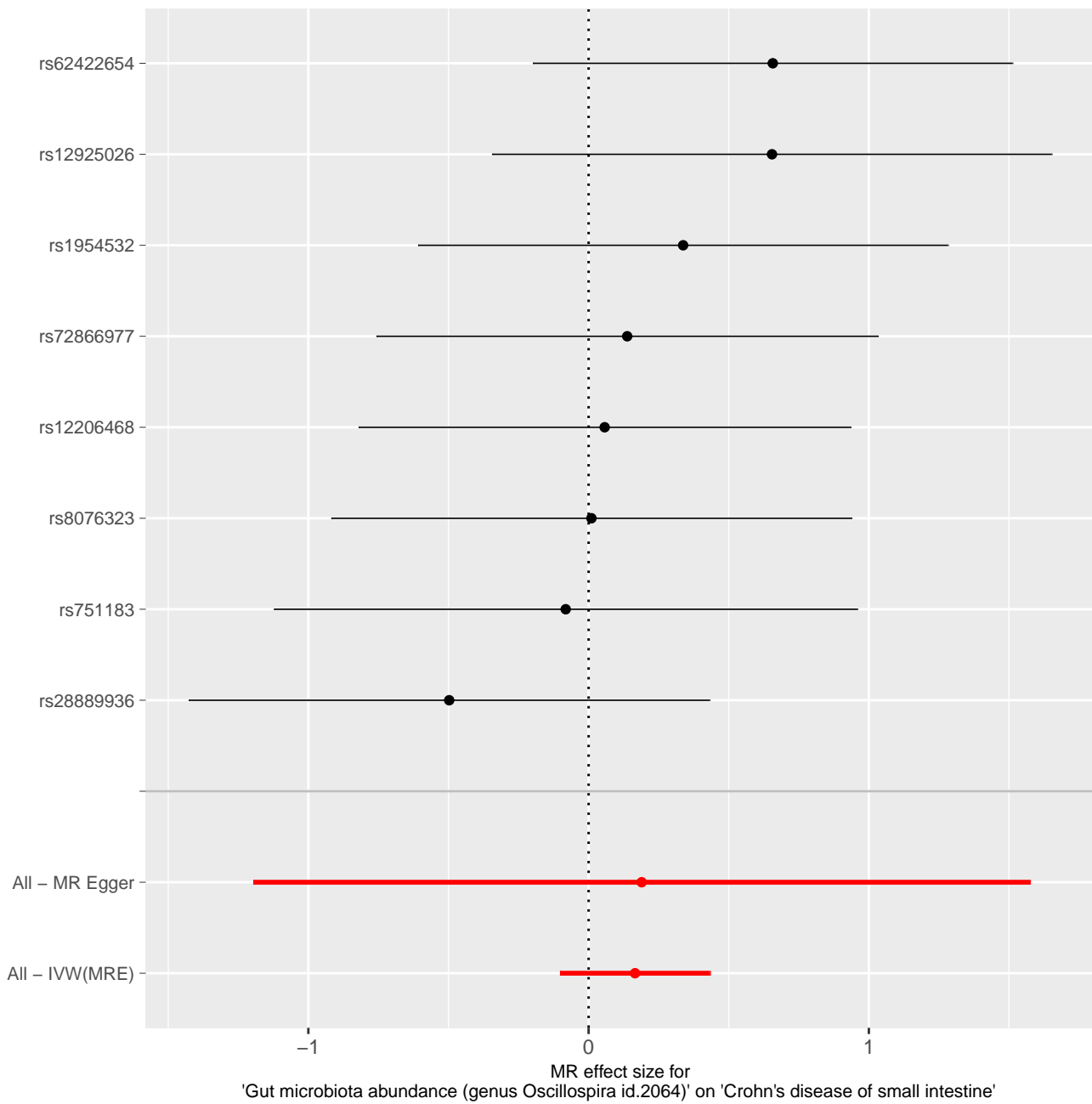


Batch 409 : Gut microbiota abundance (genus Olsenella id.822) on Crohn's disease of small intestine

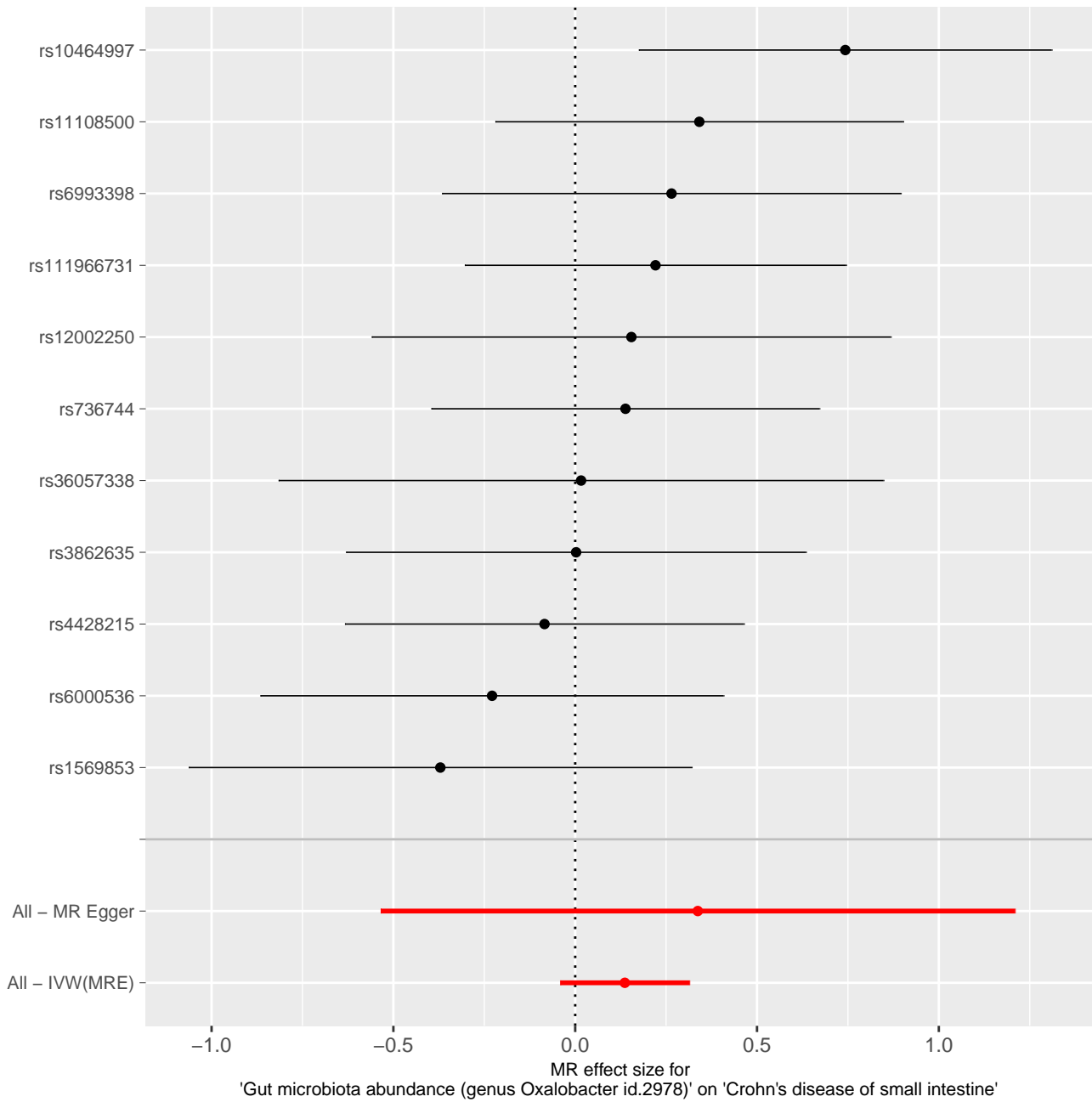


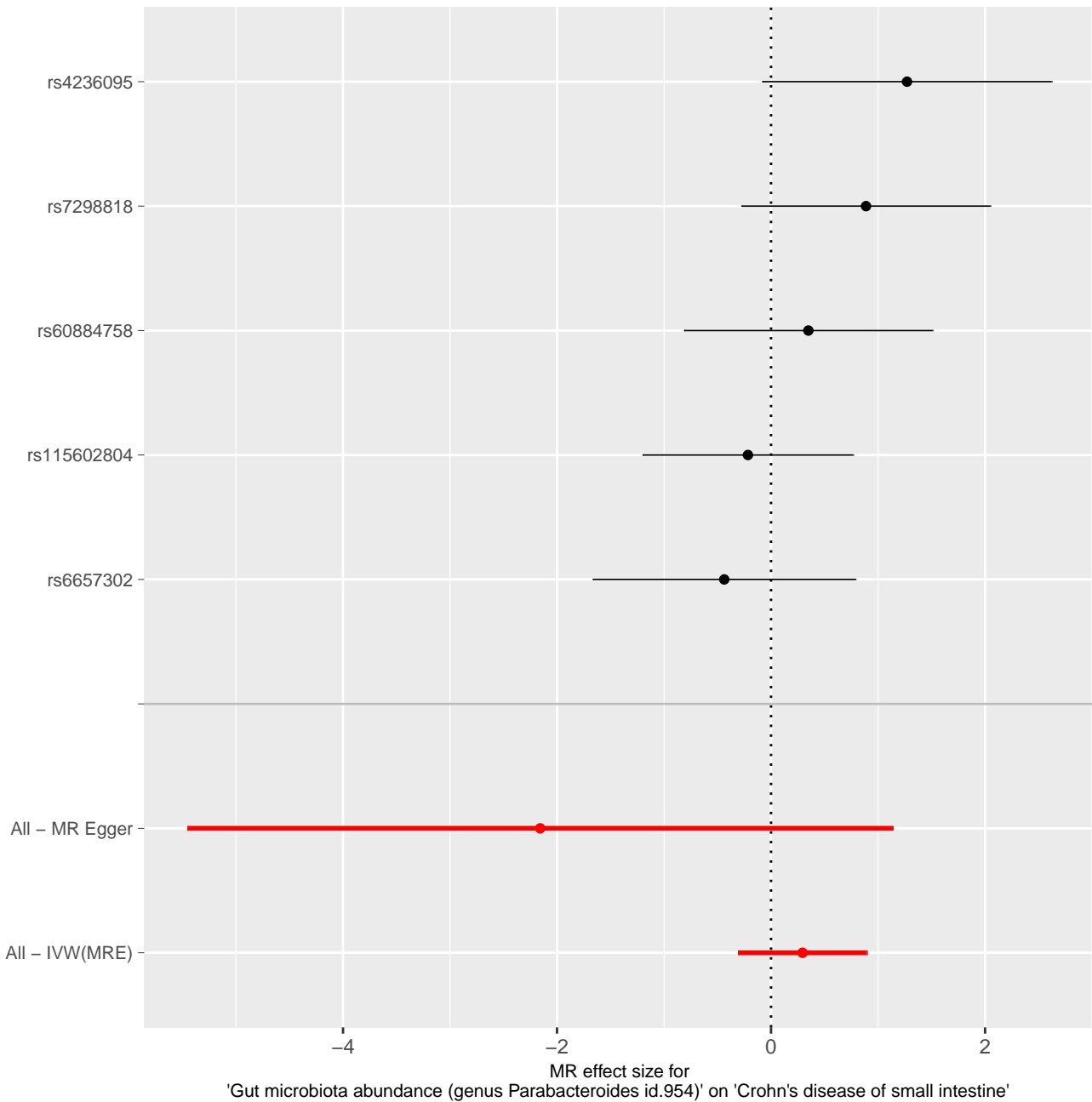


Batch 411 : Gut microbiota abundance (genus Oscillospira id.2064) on Crohn's disease of small intestine

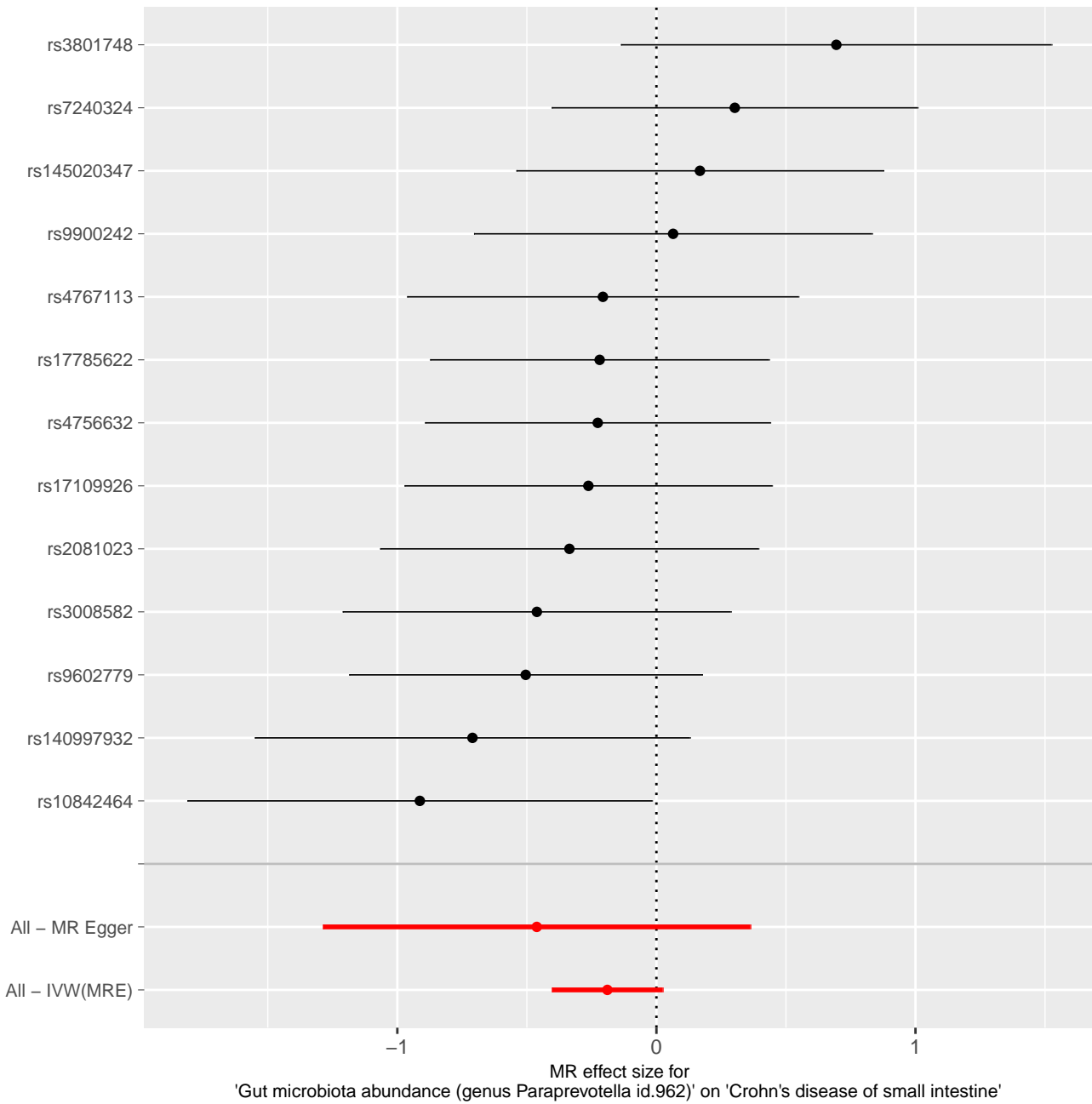


Batch 412 : Gut microbiota abundance (genus Oxalobacter id.2978) on Crohn's disease of small intestine

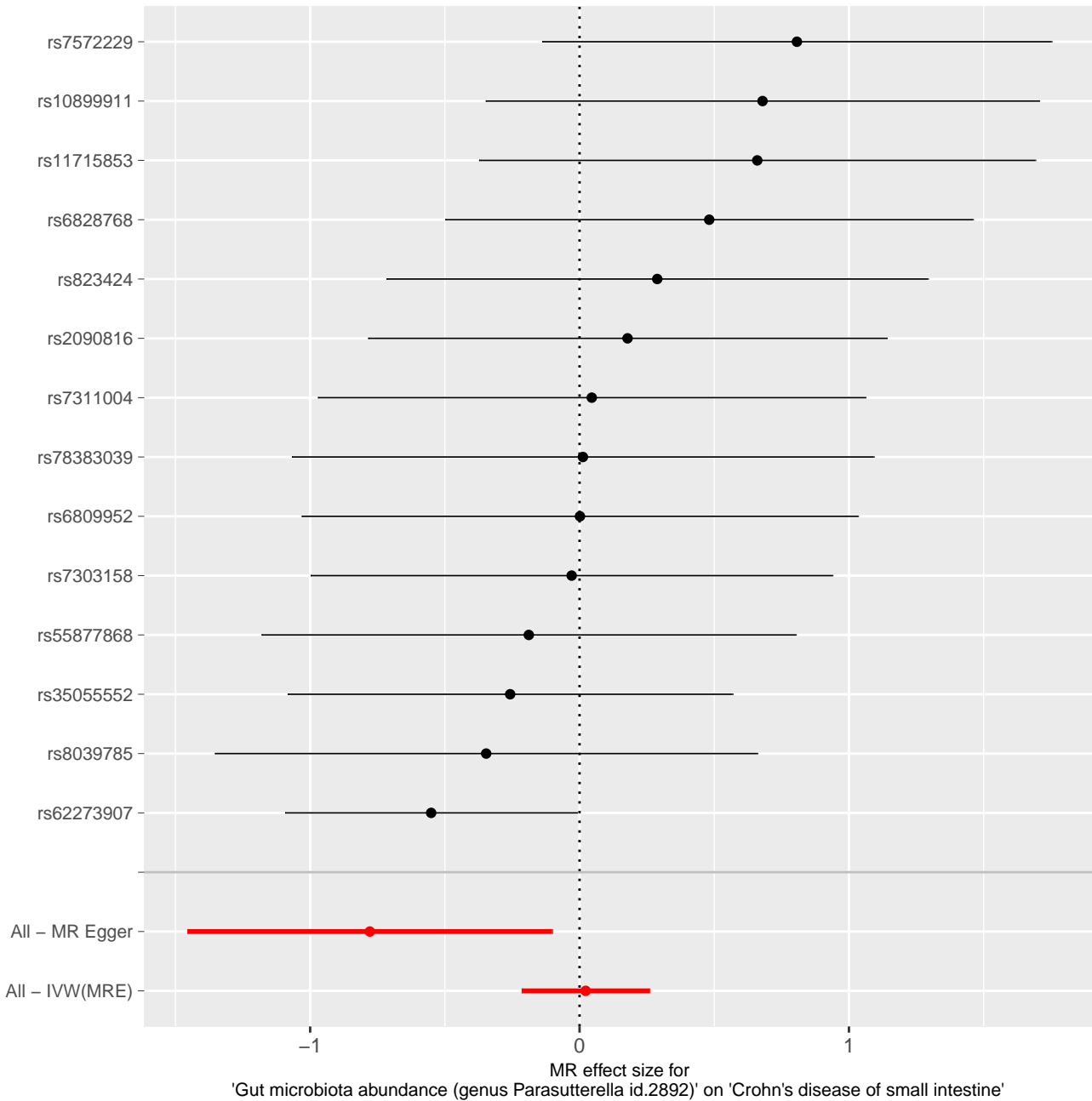


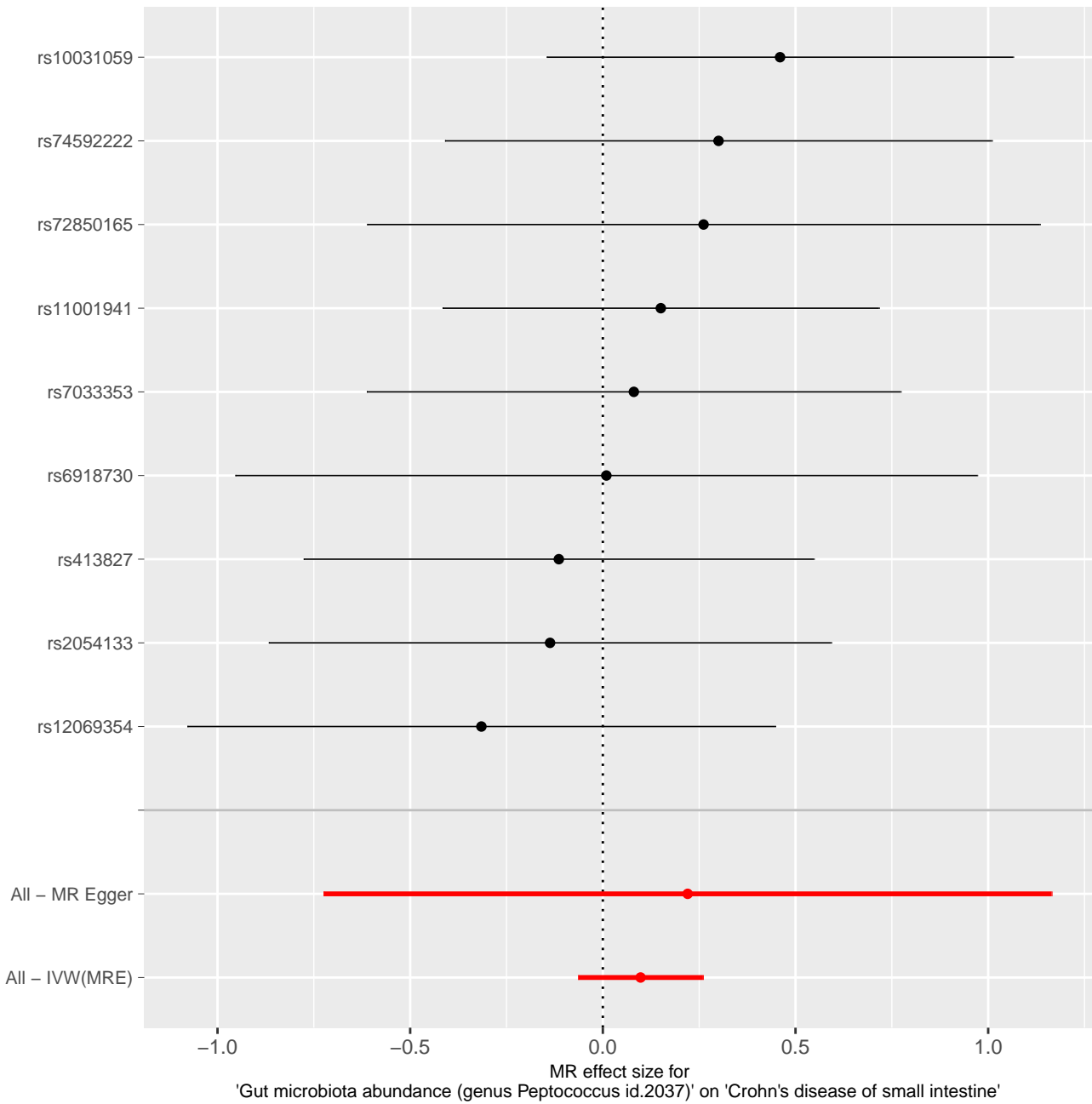


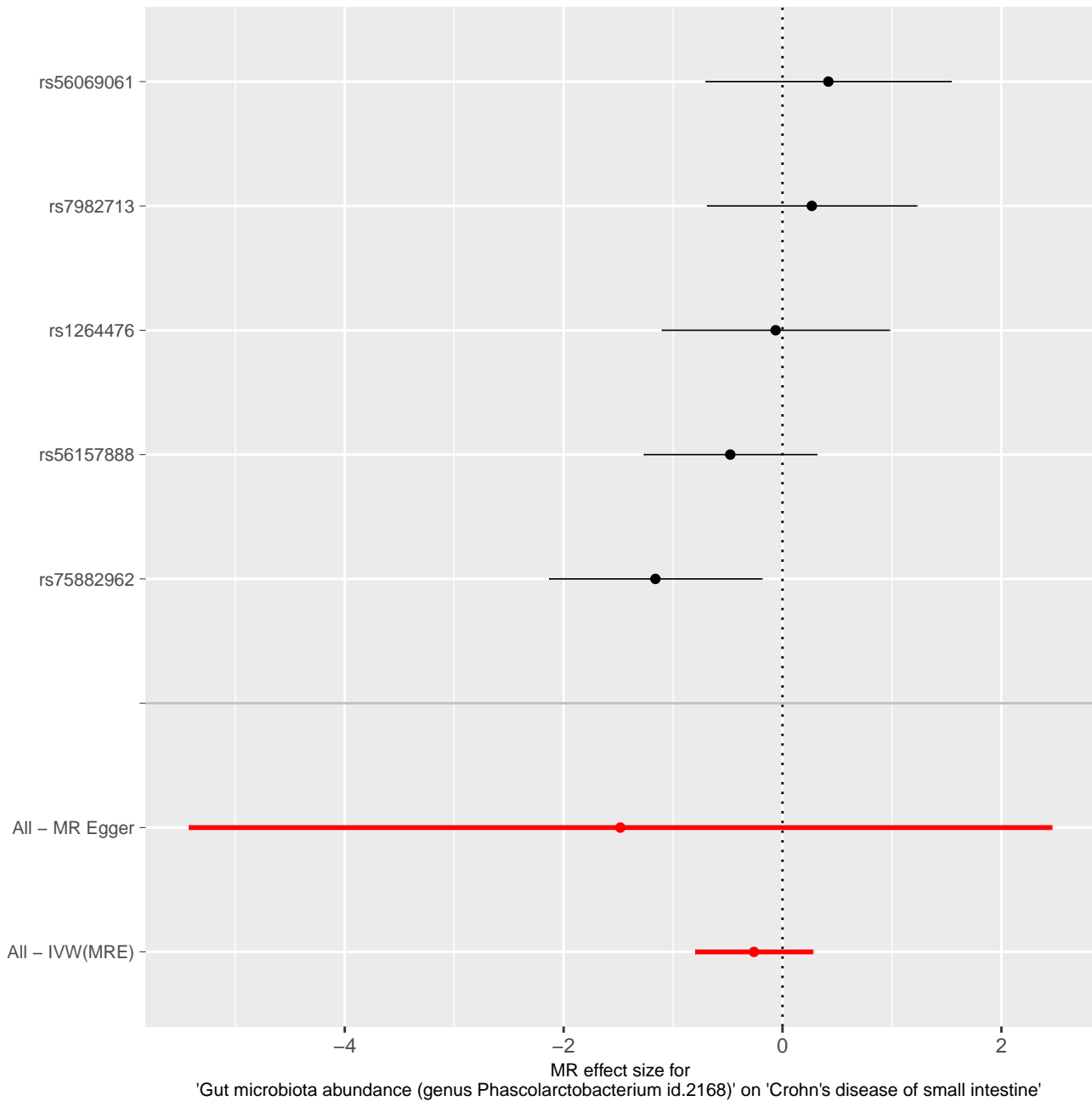
Batch 414 : Gut microbiota abundance (genus Paraprevotella id.962) on Crohn's disease of small intestine



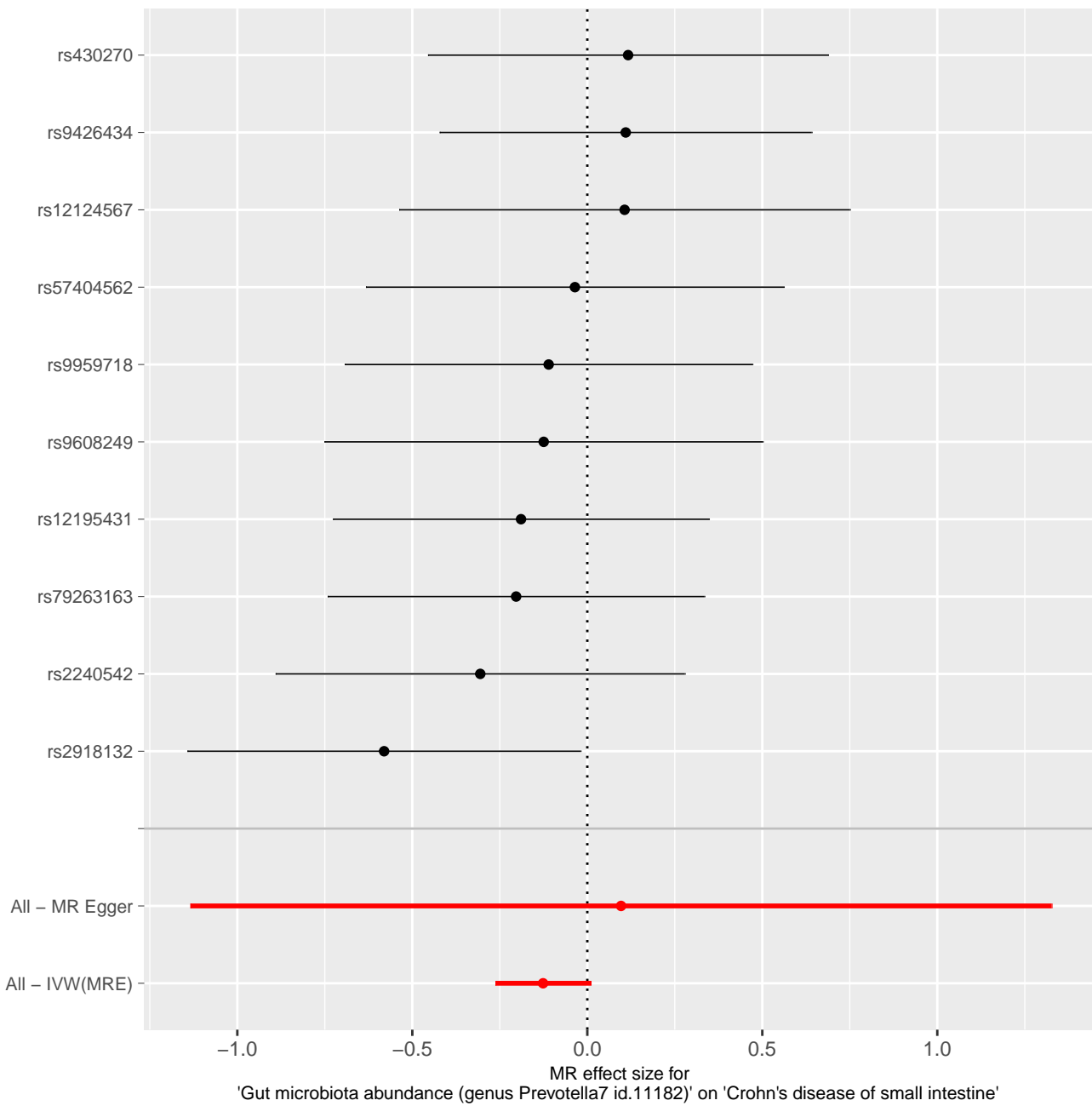
Batch 415 : Gut microbiota abundance (genus Parasutterella id.2892) on Crohn's disease of small intestine



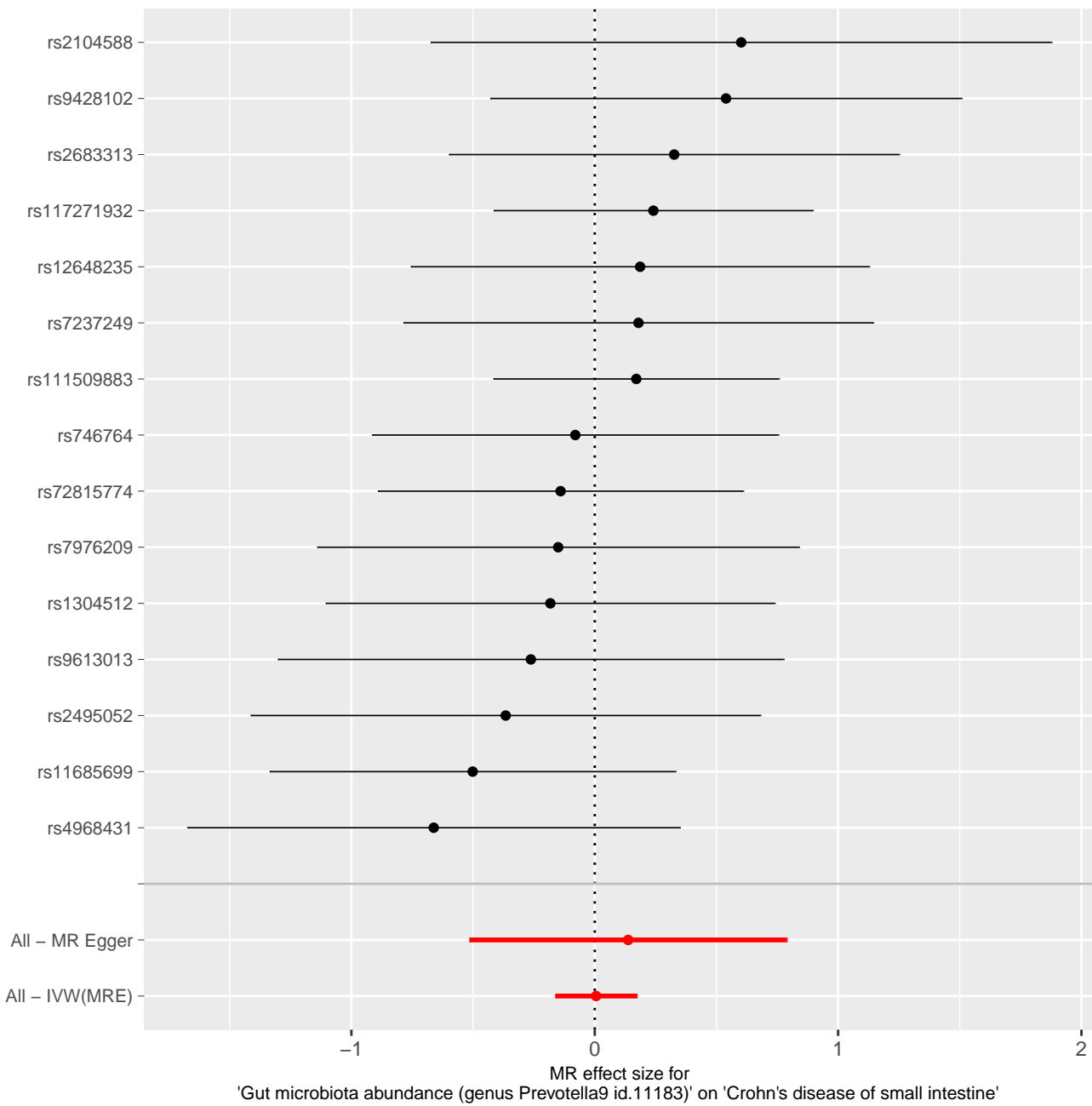


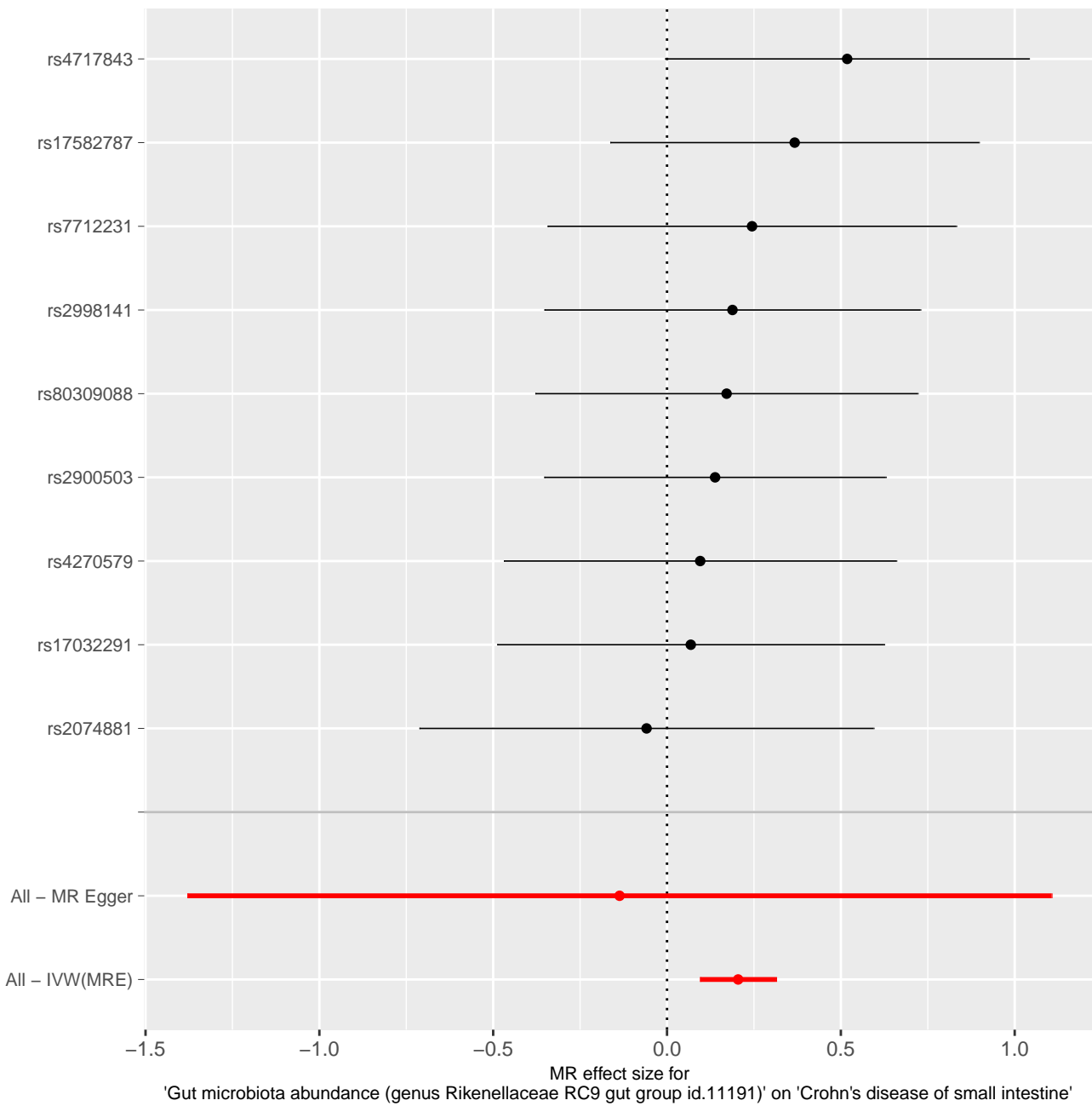


Batch 418 : Gut microbiota abundance (genus Prevotella7 id.11182) on Crohn's disease of small intestine

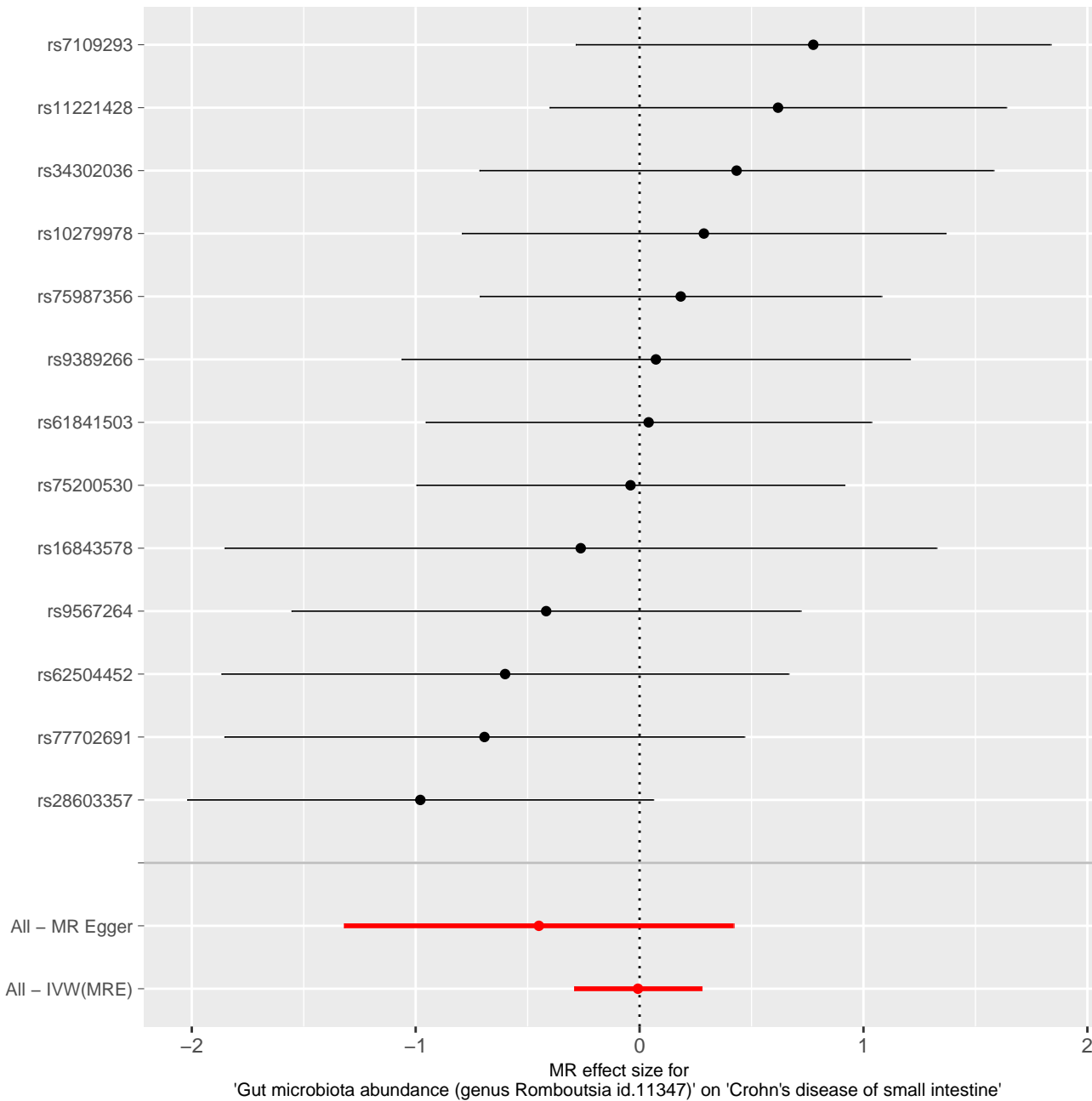


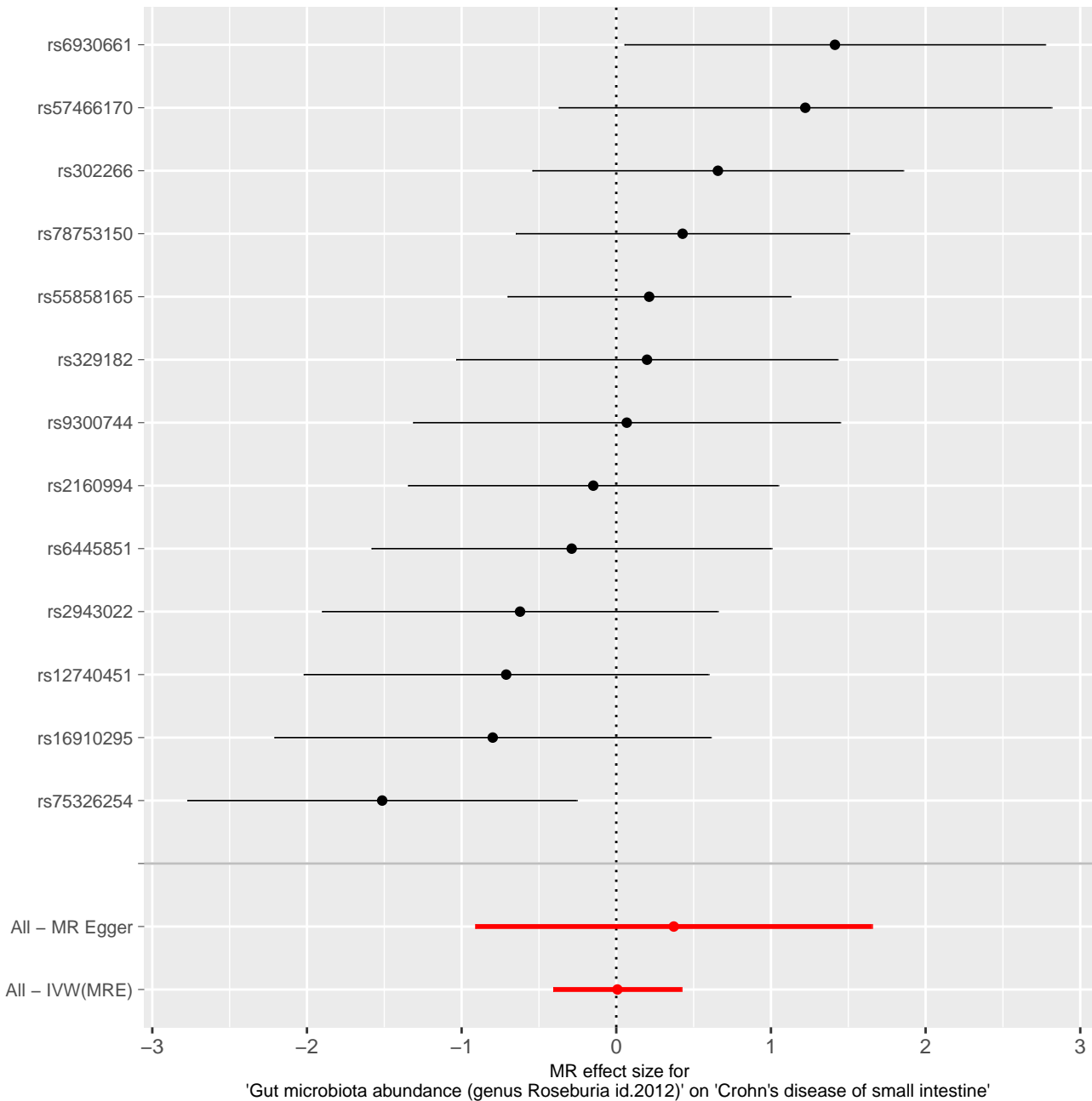
Batch 419 : Gut microbiota abundance (genus Prevotella9 id.11183) on Crohn's disease of small intestine

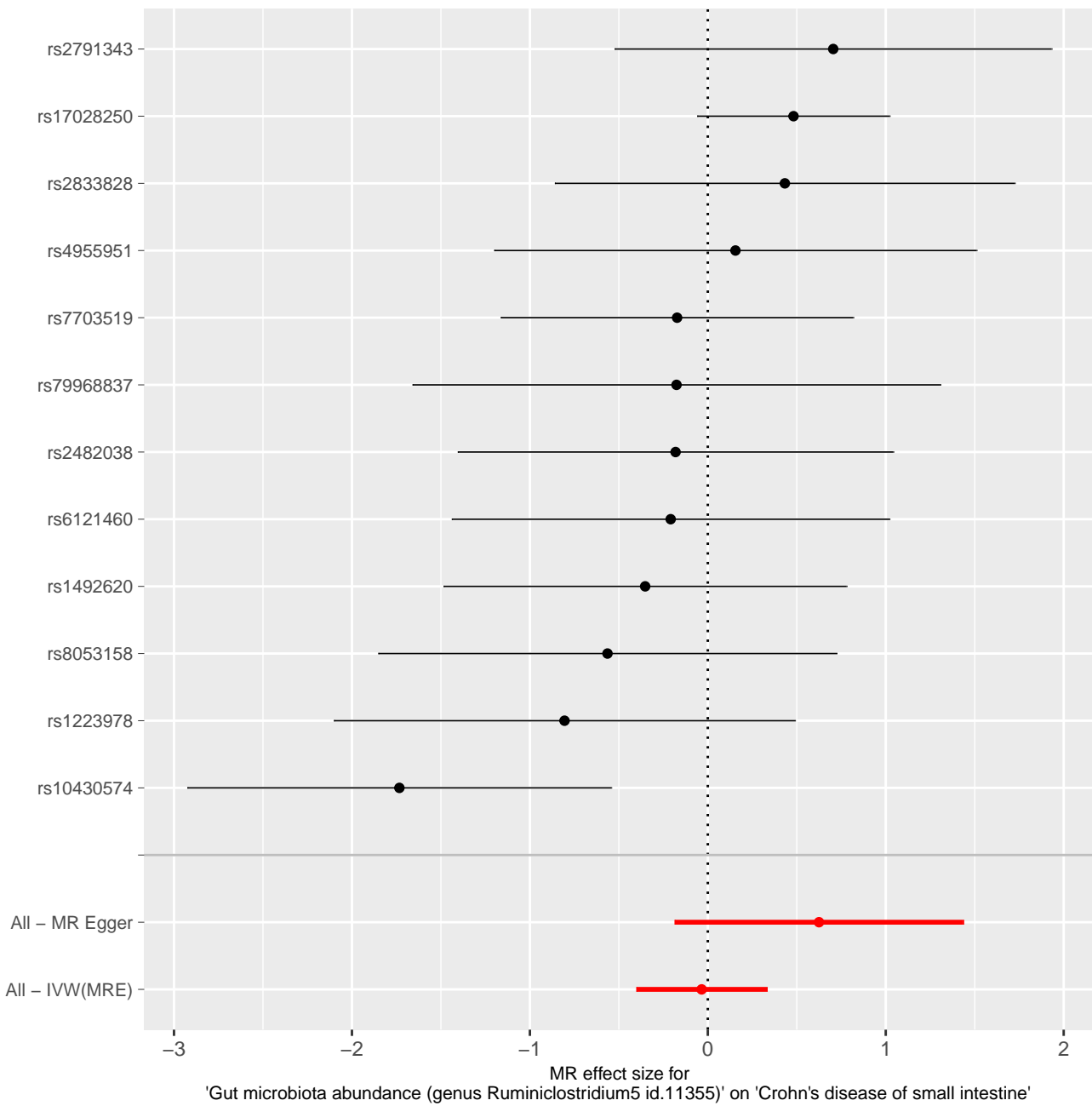


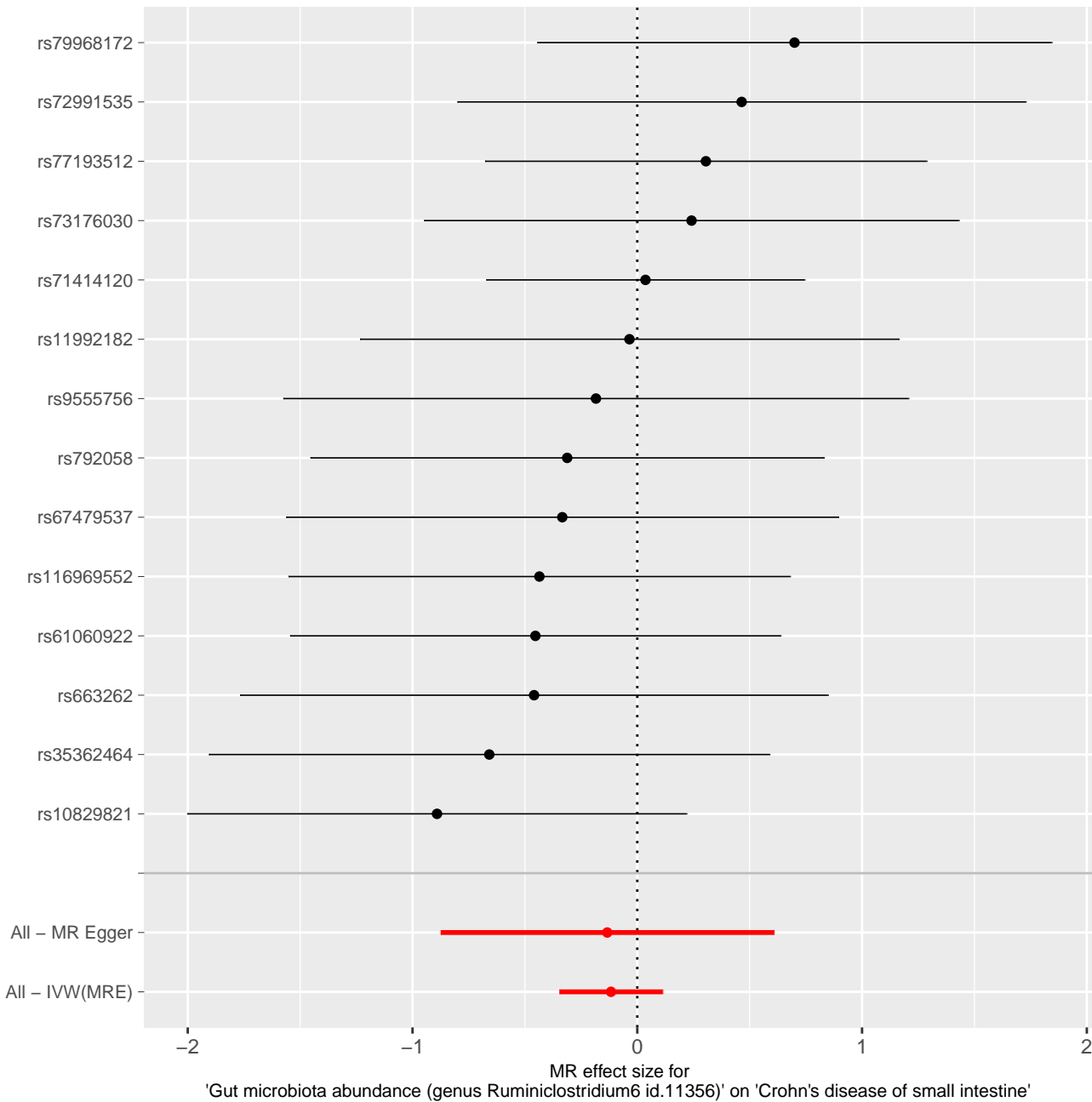


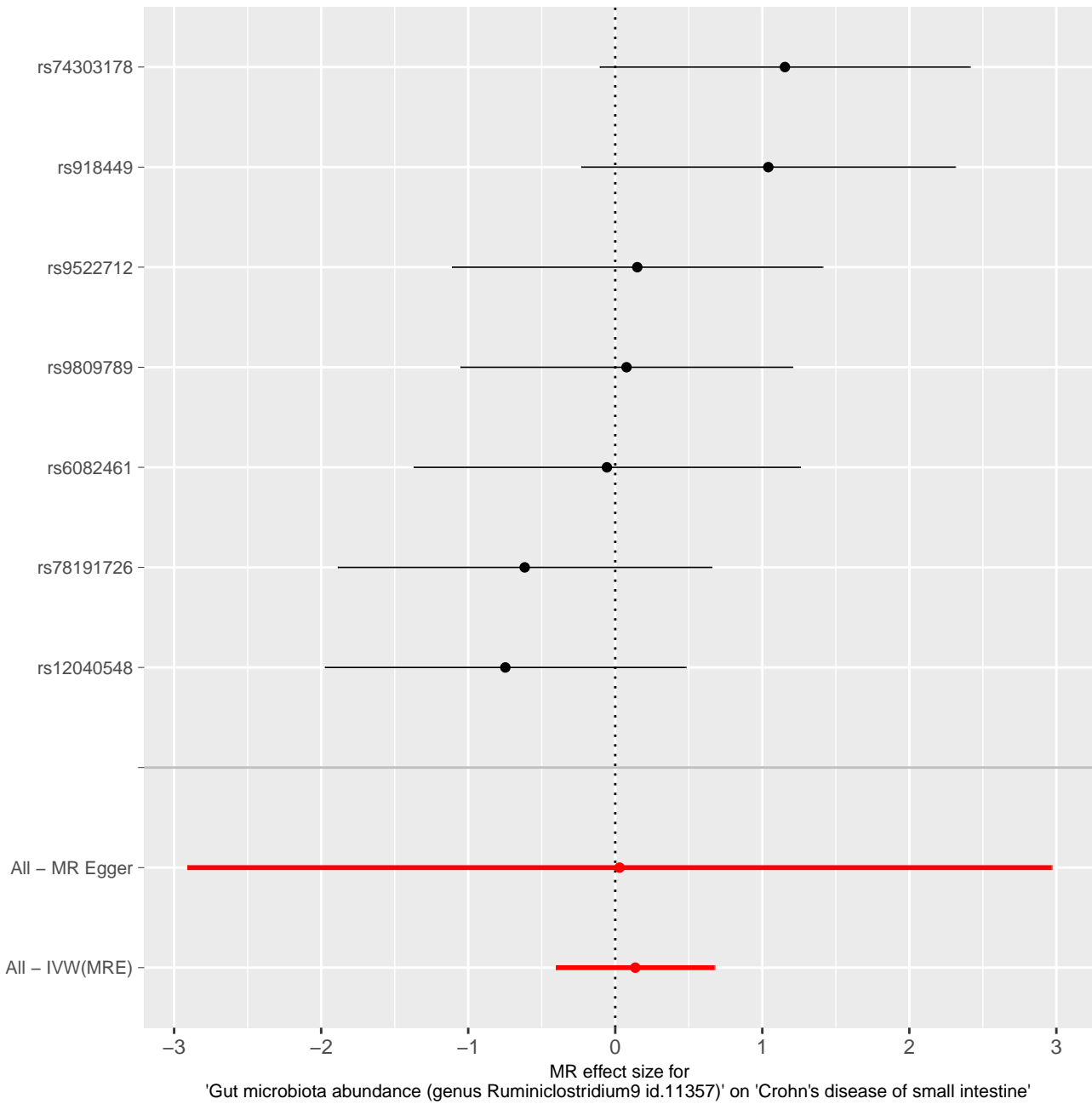
Batch 421 : Gut microbiota abundance (genus Romboutsia id.11347) on Crohn's disease of small intestine

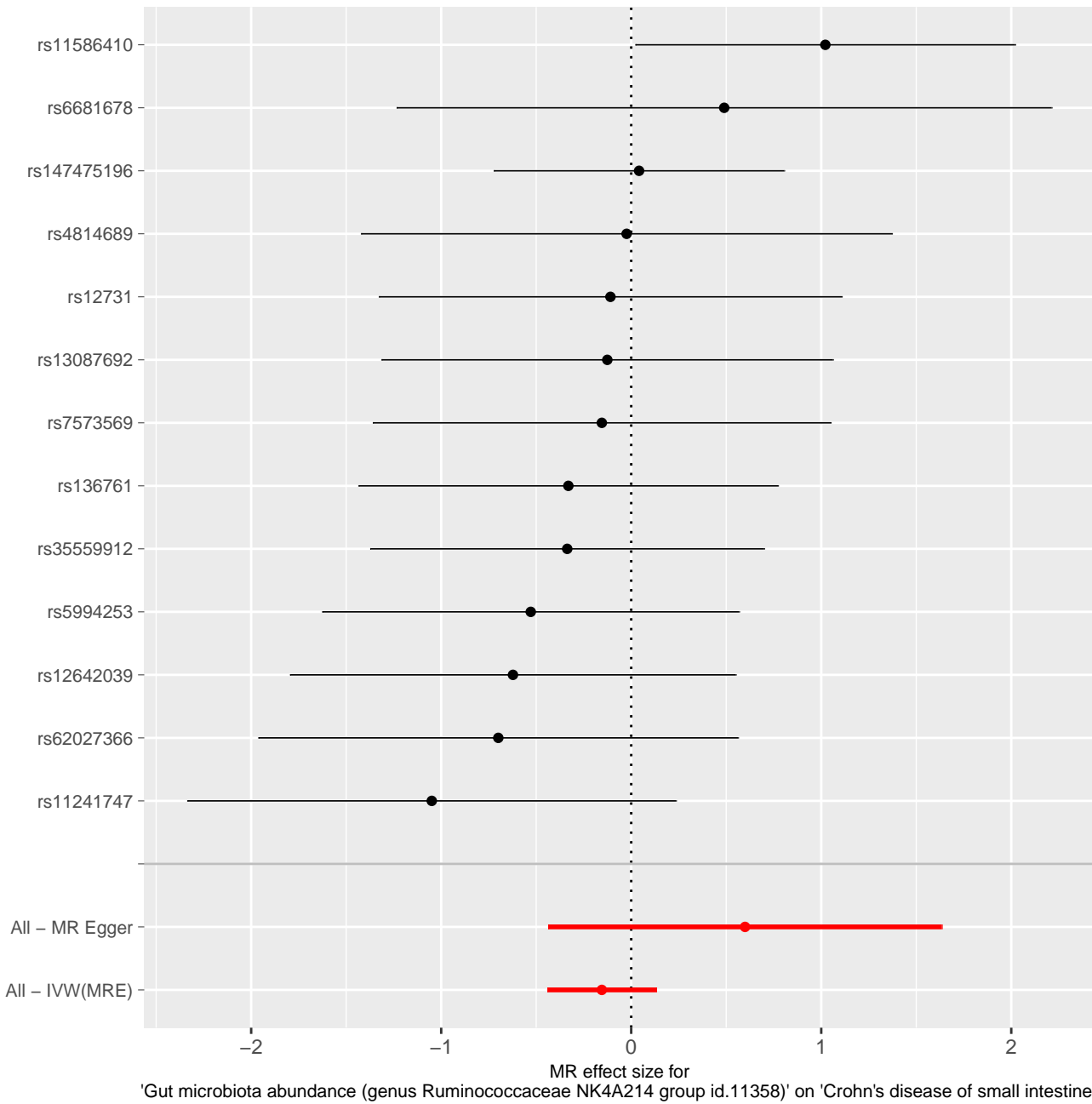




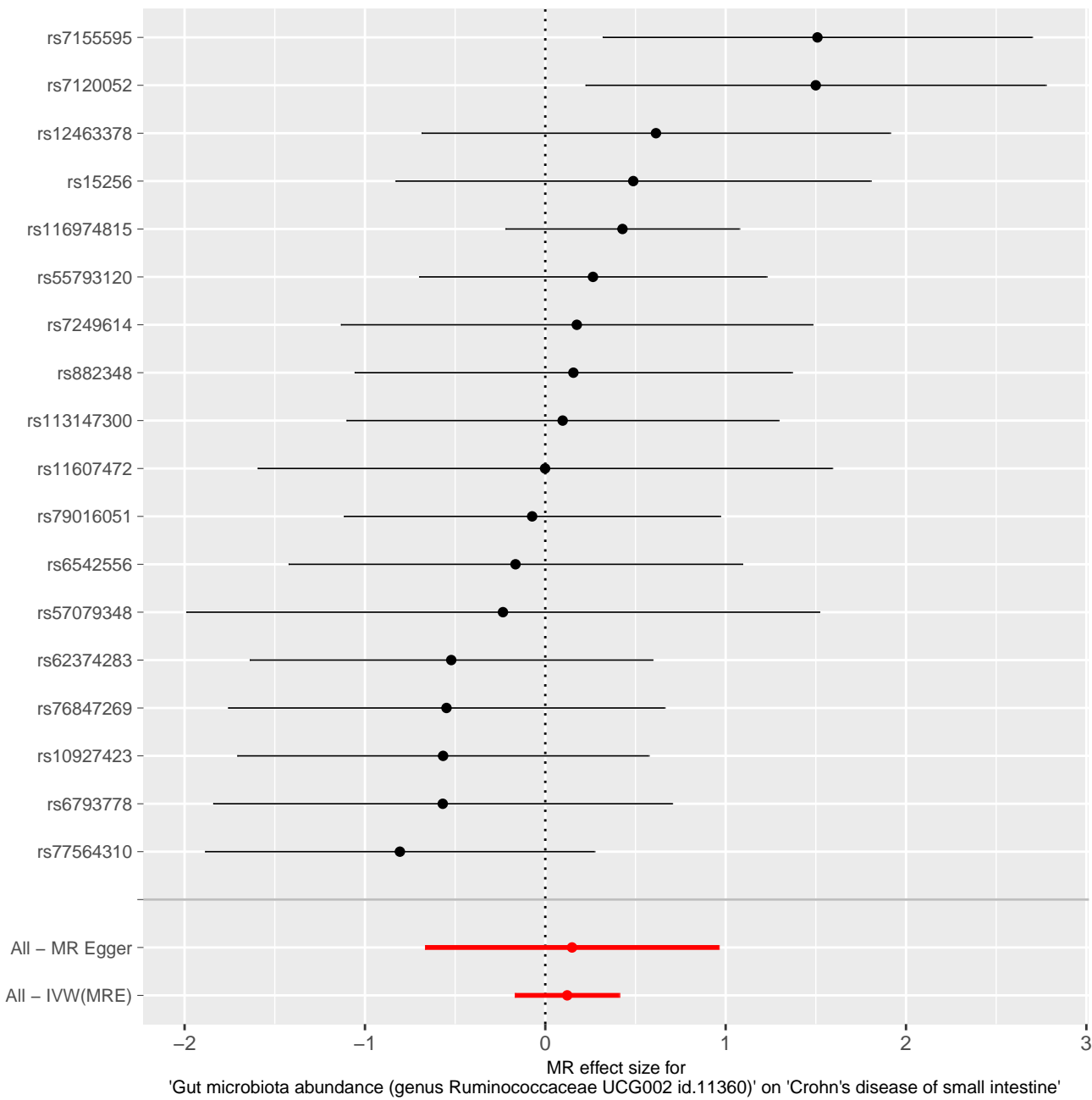


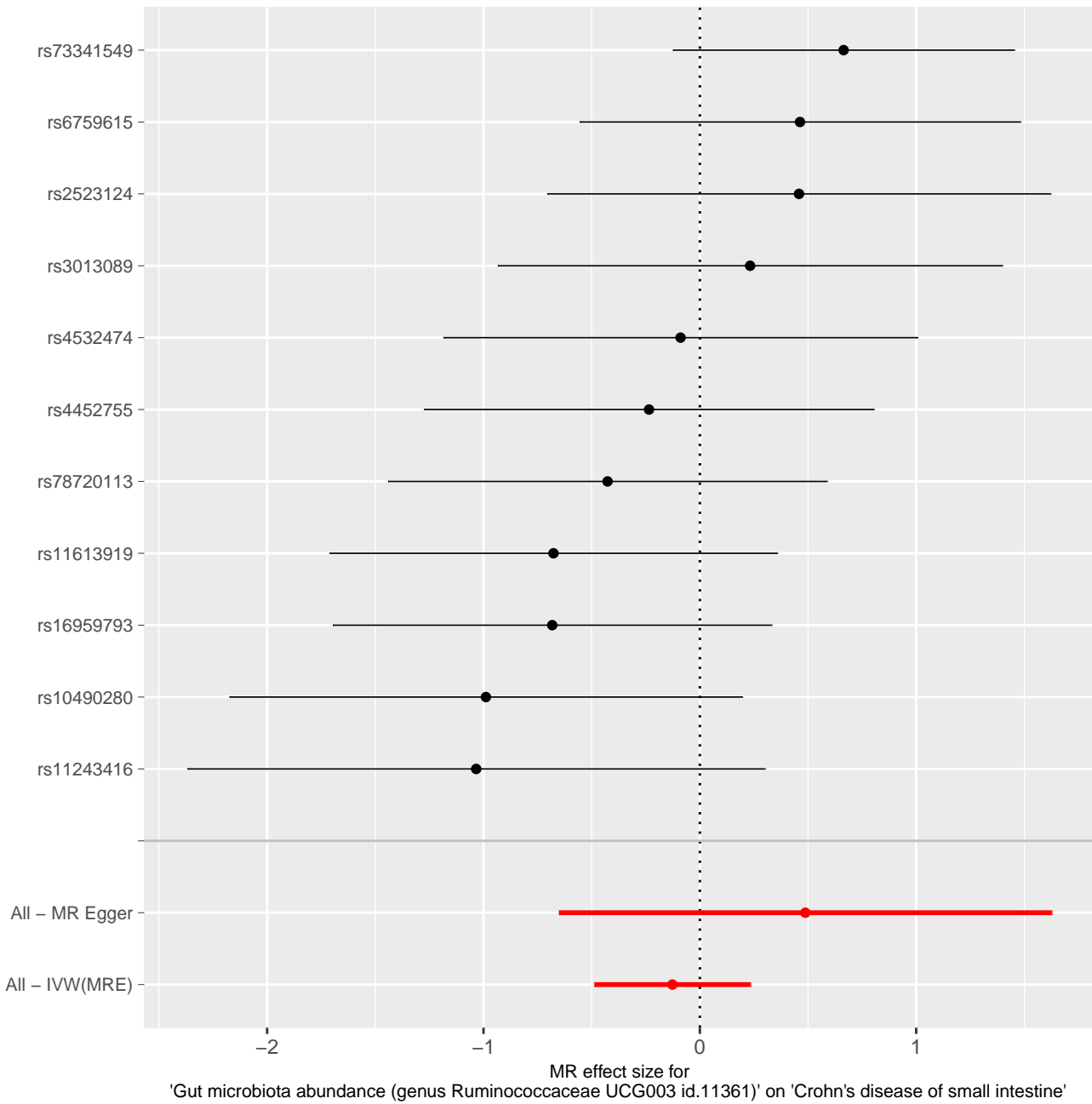


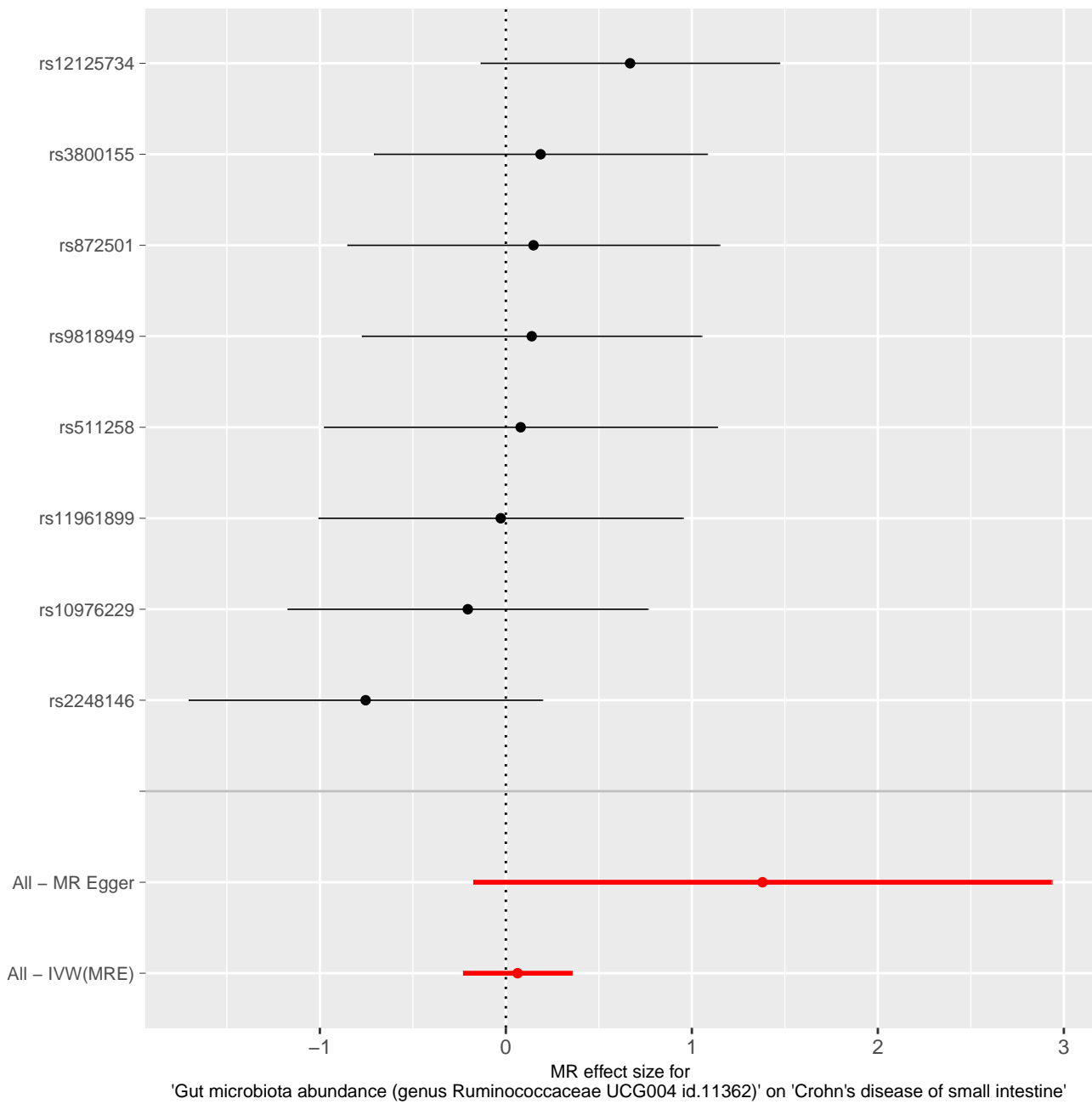


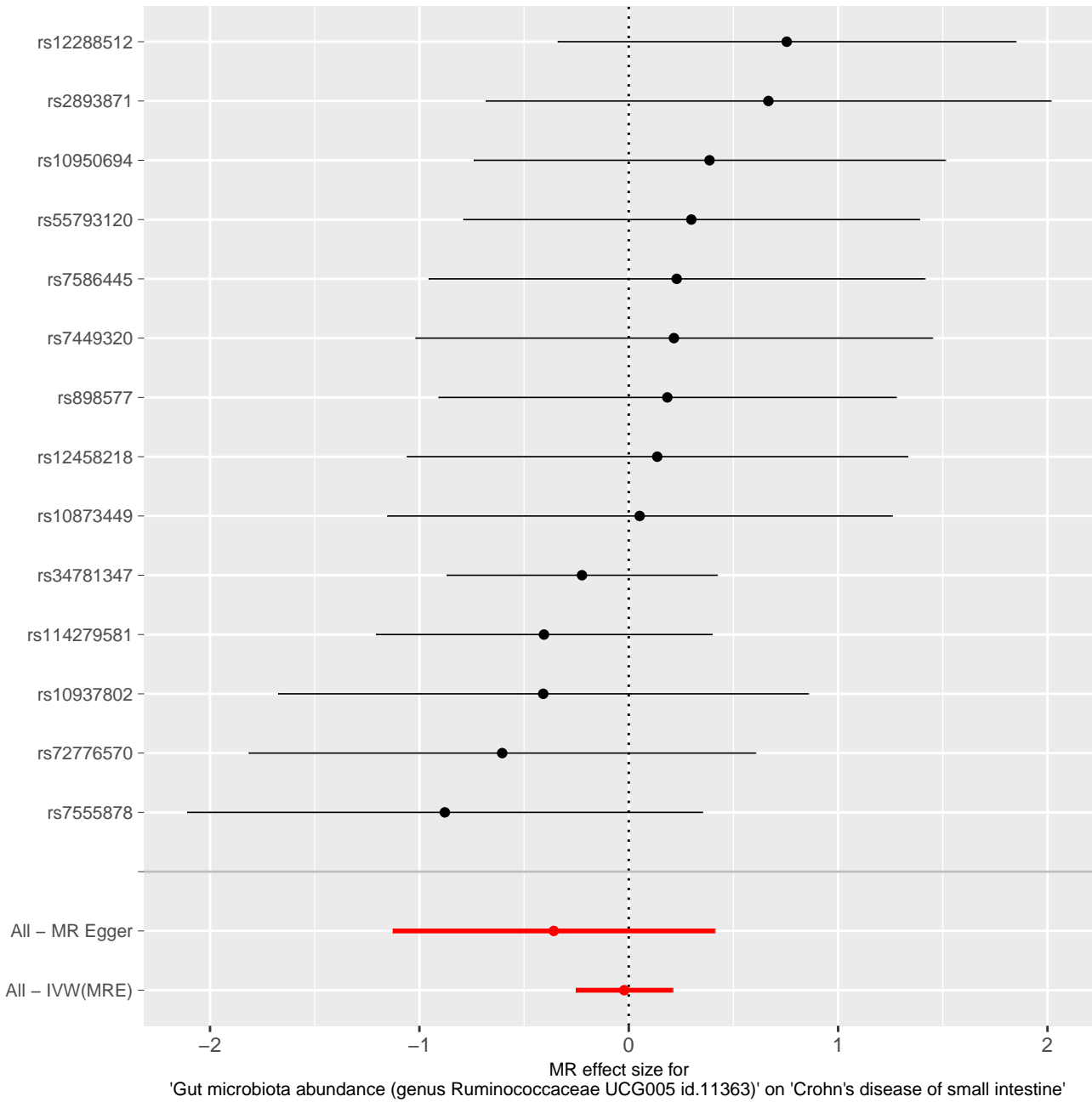


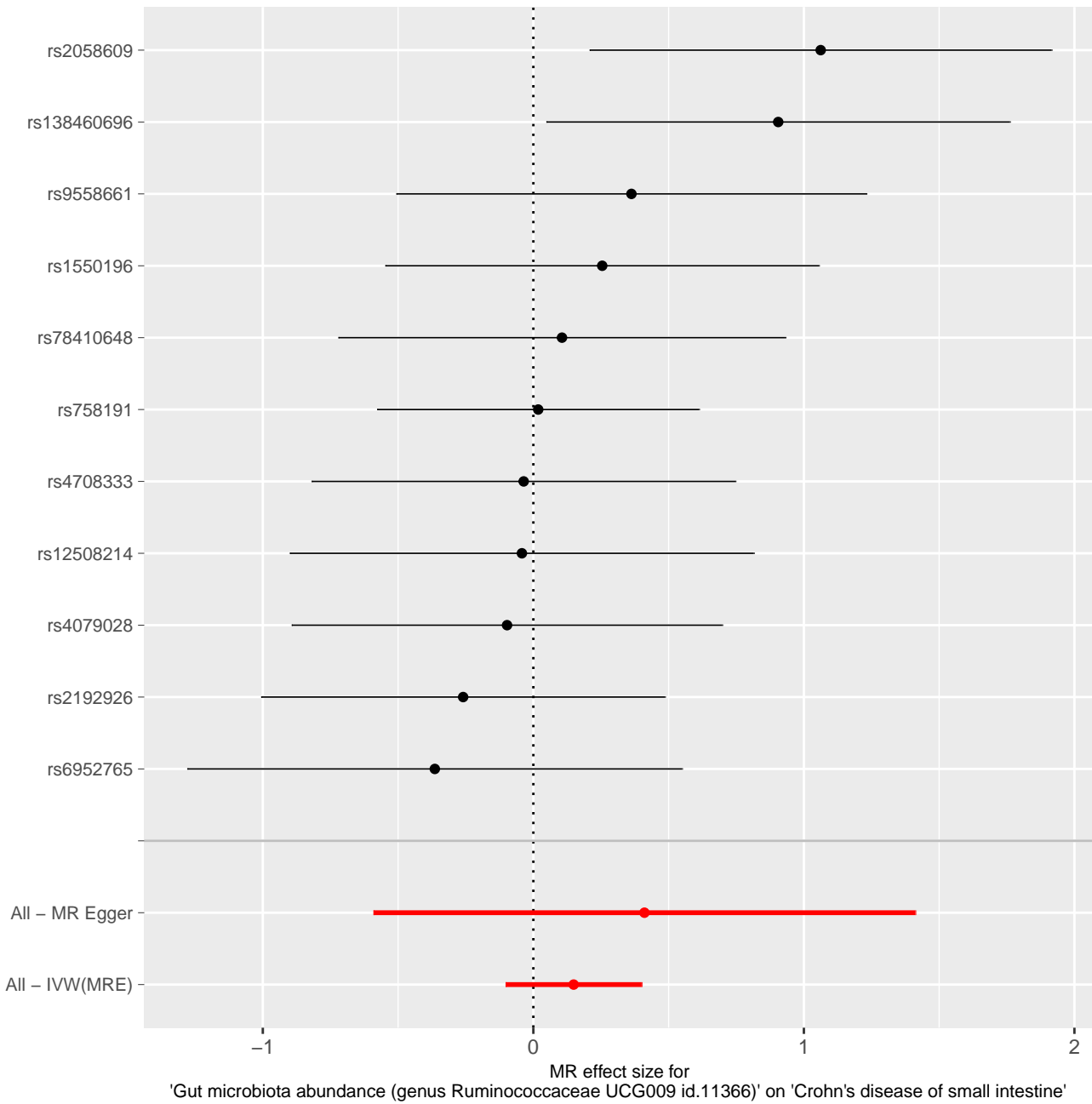
Batch 427 : Gut microbiota abundance (genus Ruminococcaceae UCG002 id.11360) on Crohn's disease of small intestine

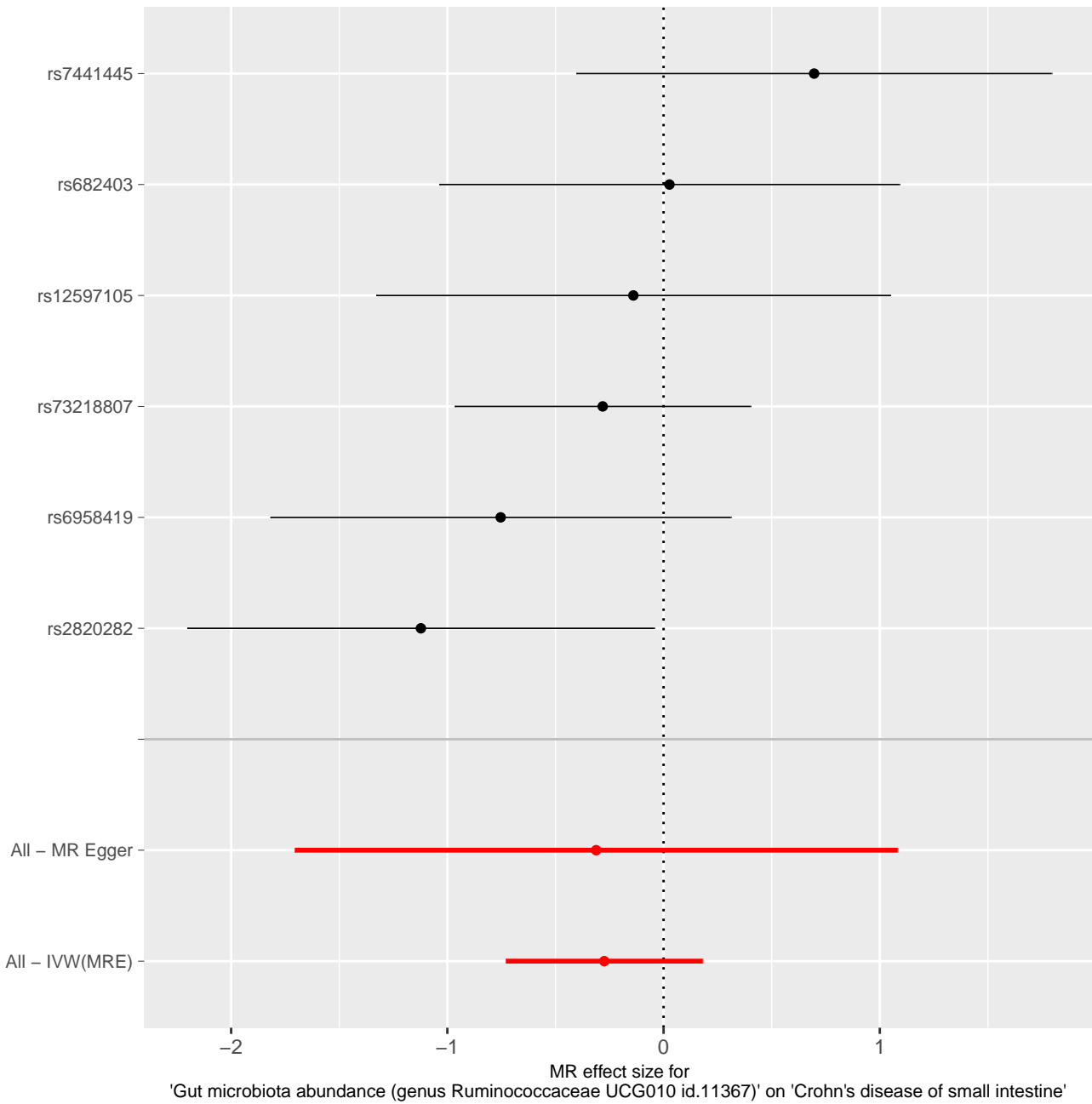


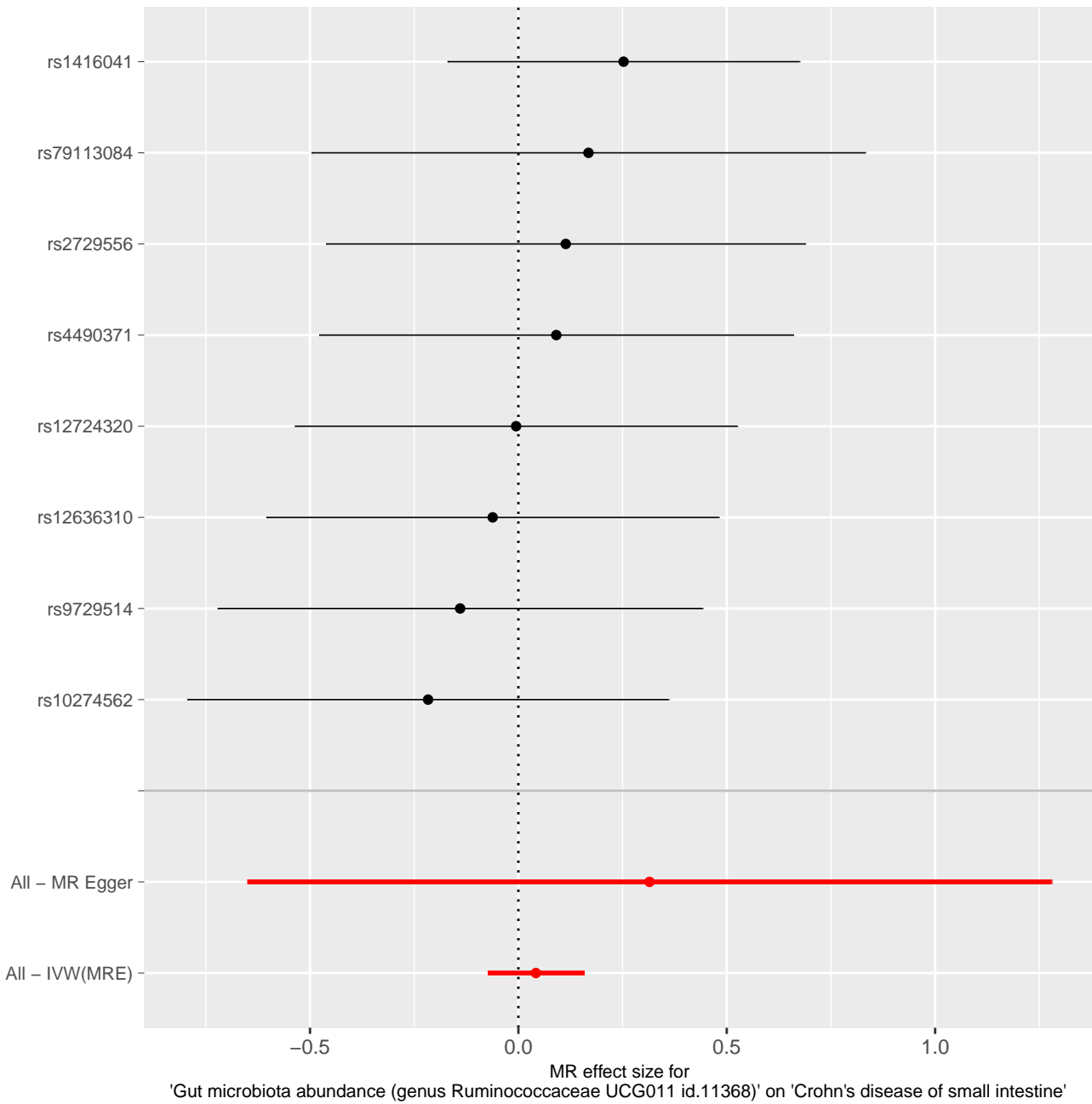


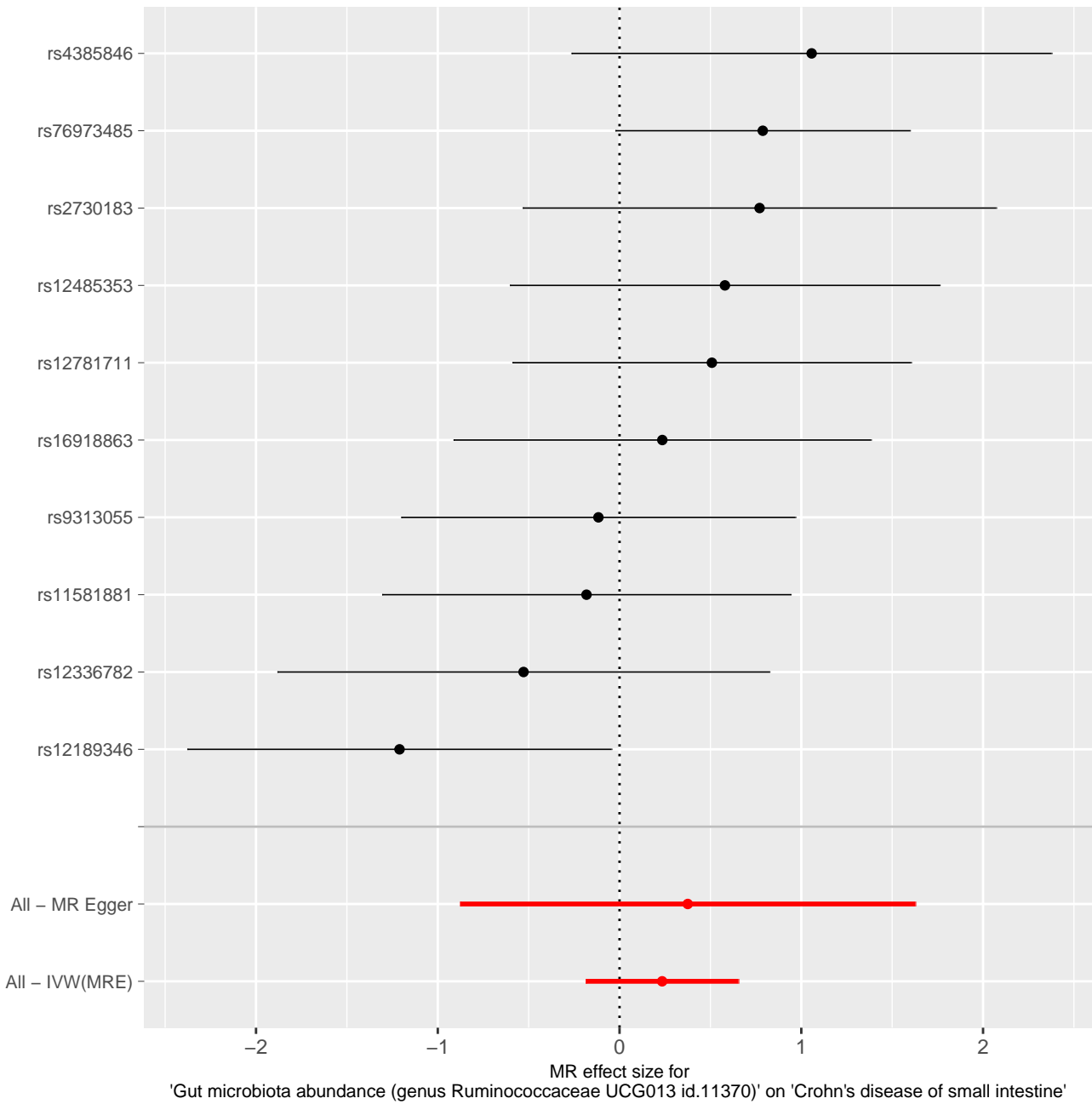


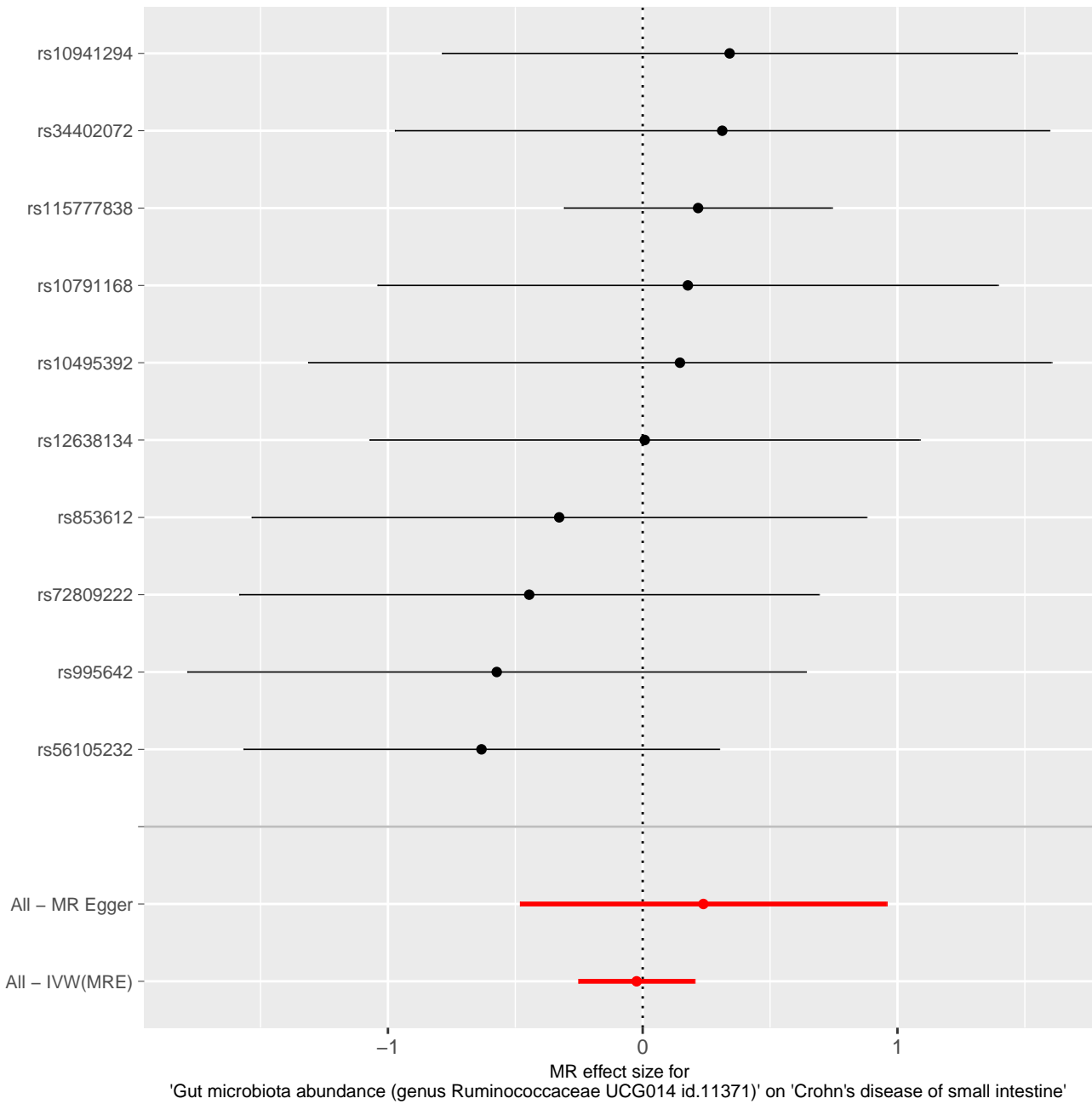




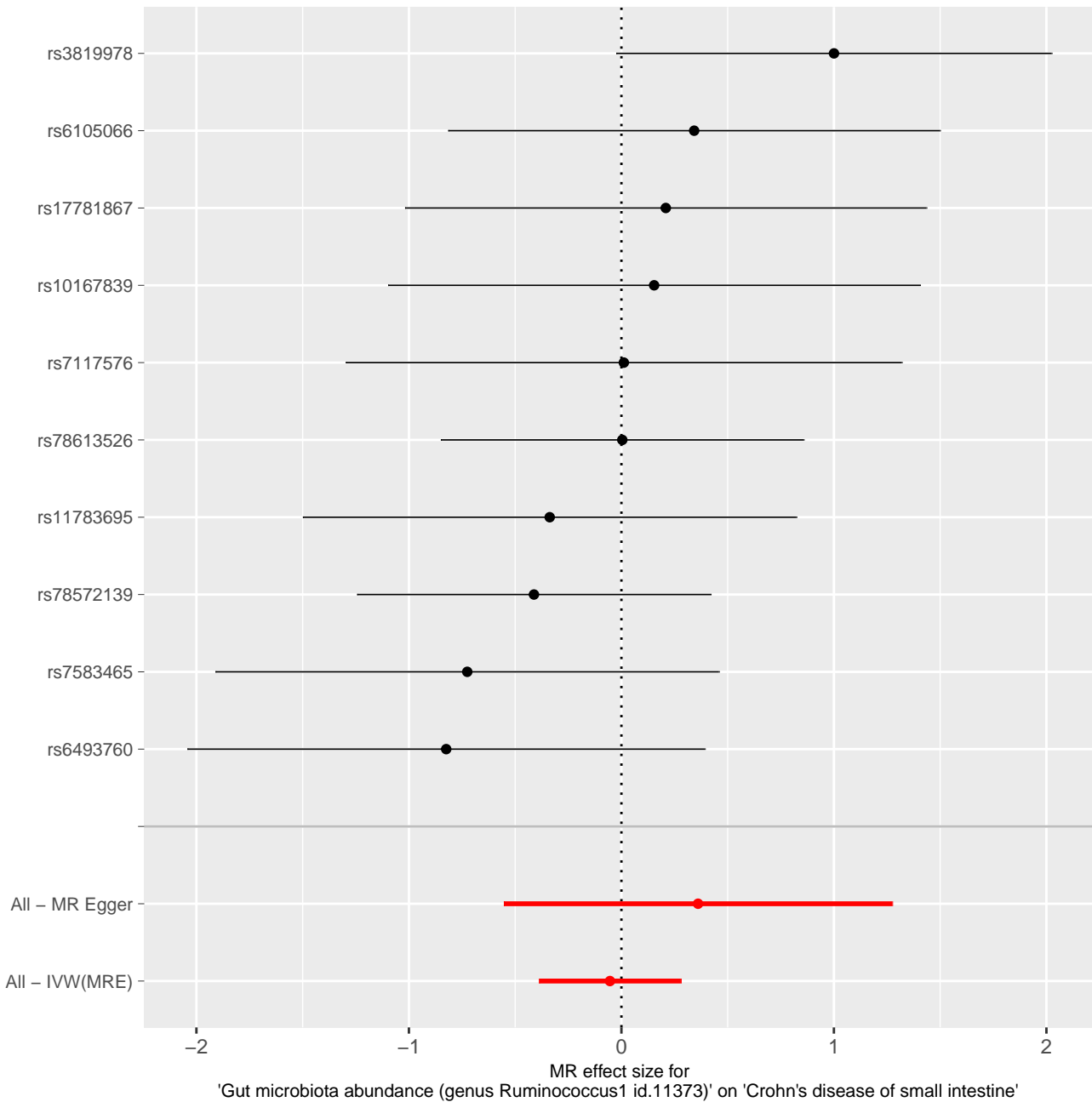




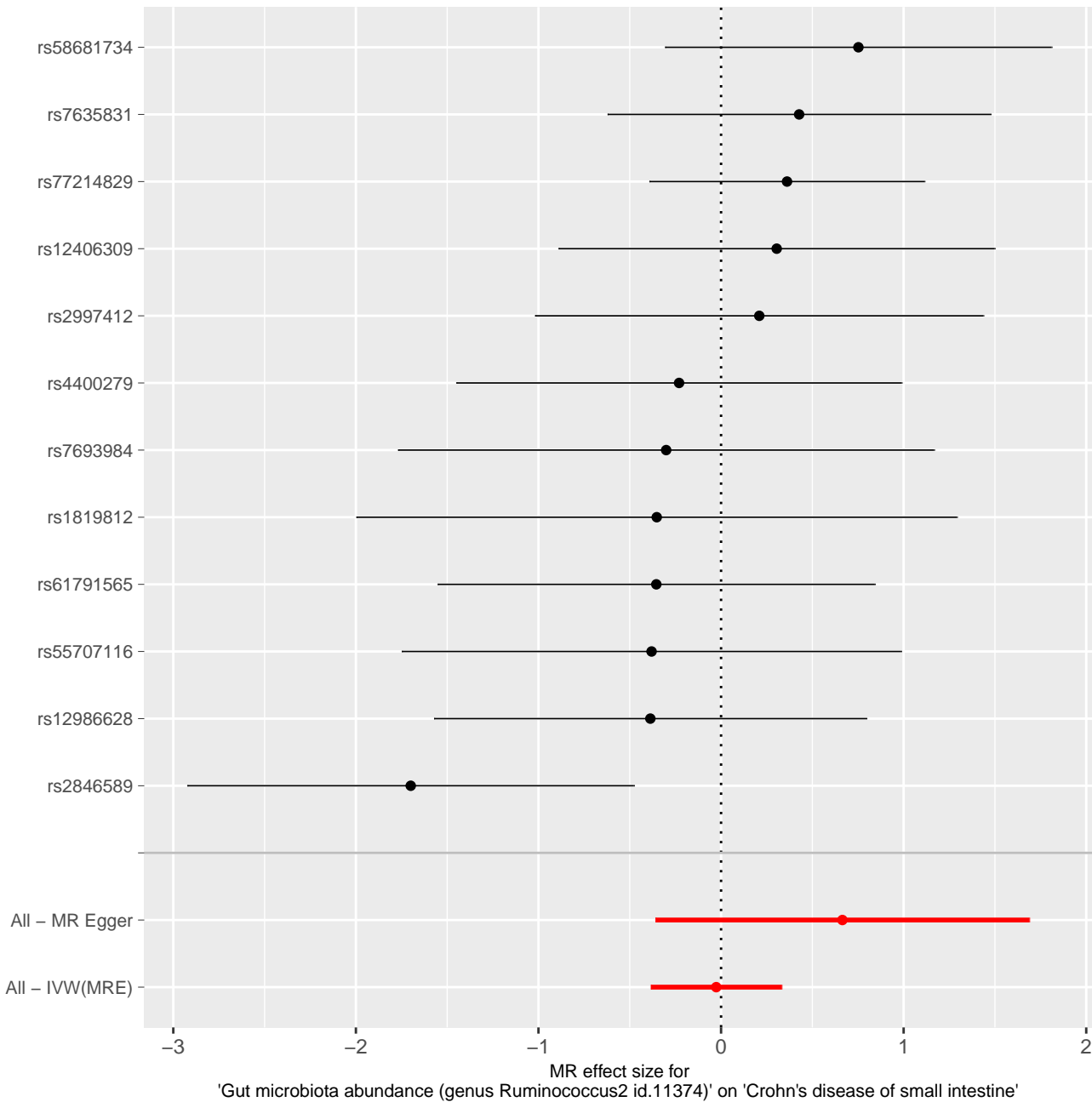




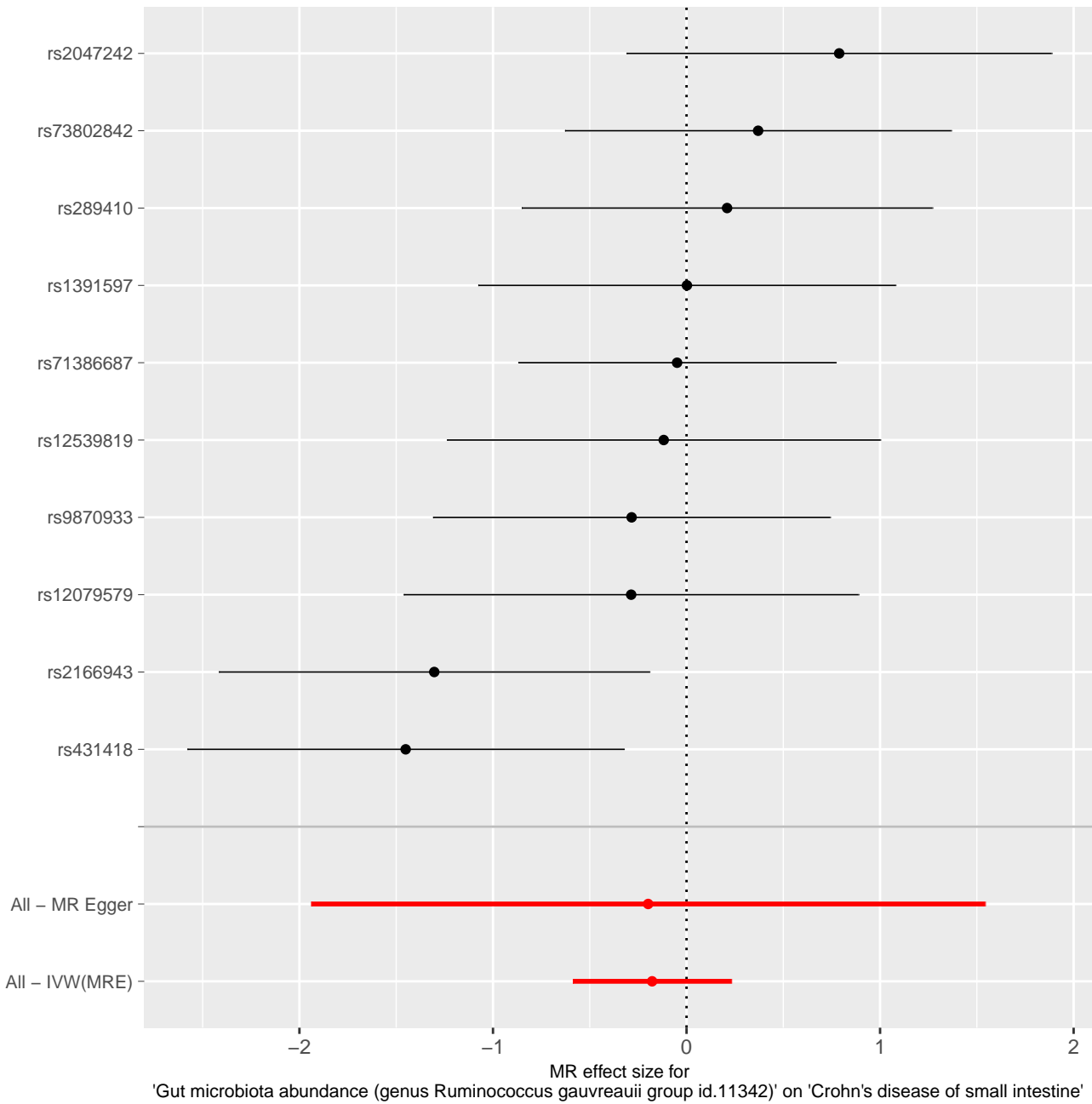
Batch 436 : Gut microbiota abundance (genus Ruminococcus1 id.11373) on Crohn's disease of small intestine



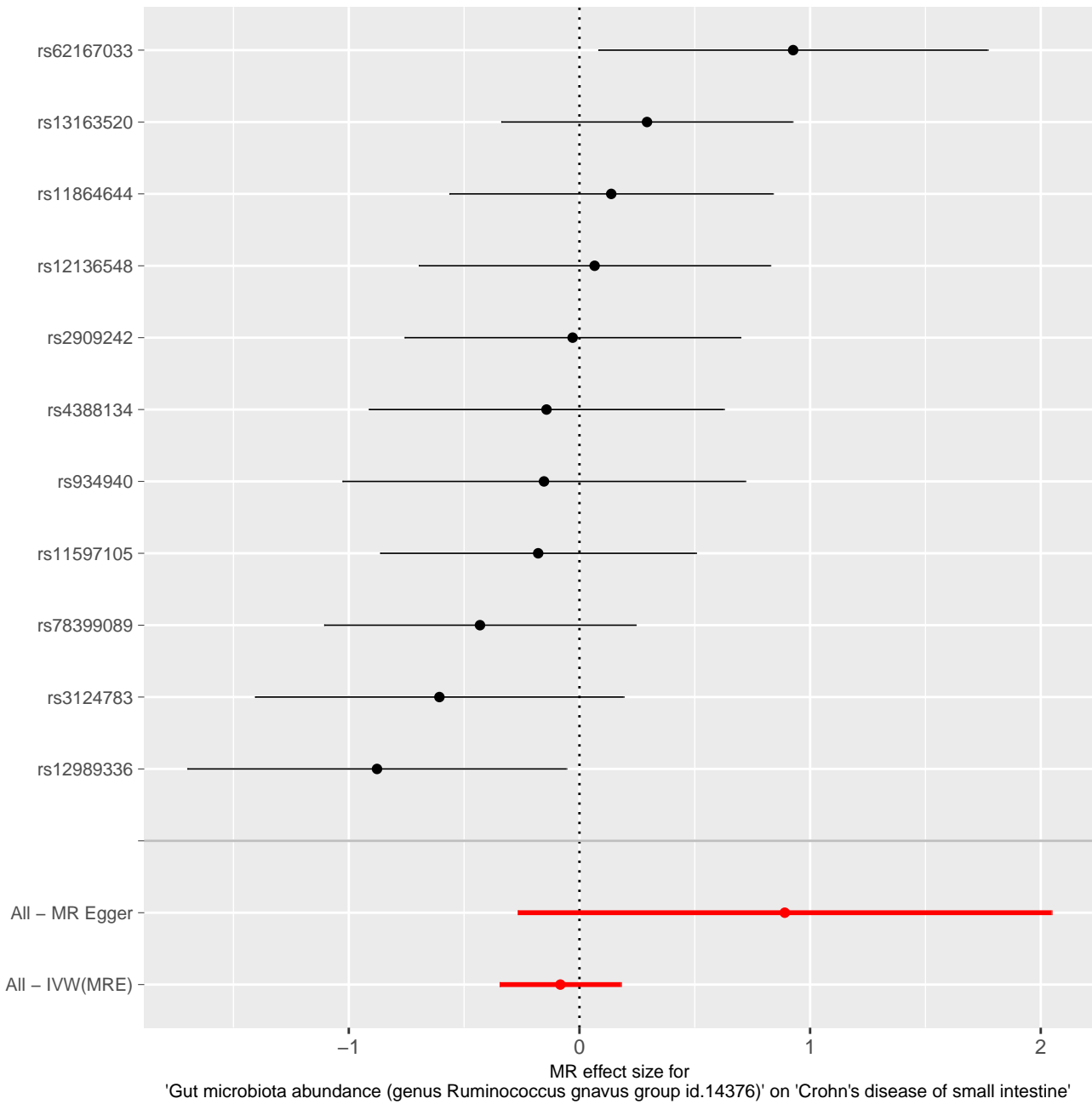
Batch 437 : Gut microbiota abundance (genus Ruminococcus2 id.11374) on Crohn's disease of small intestine



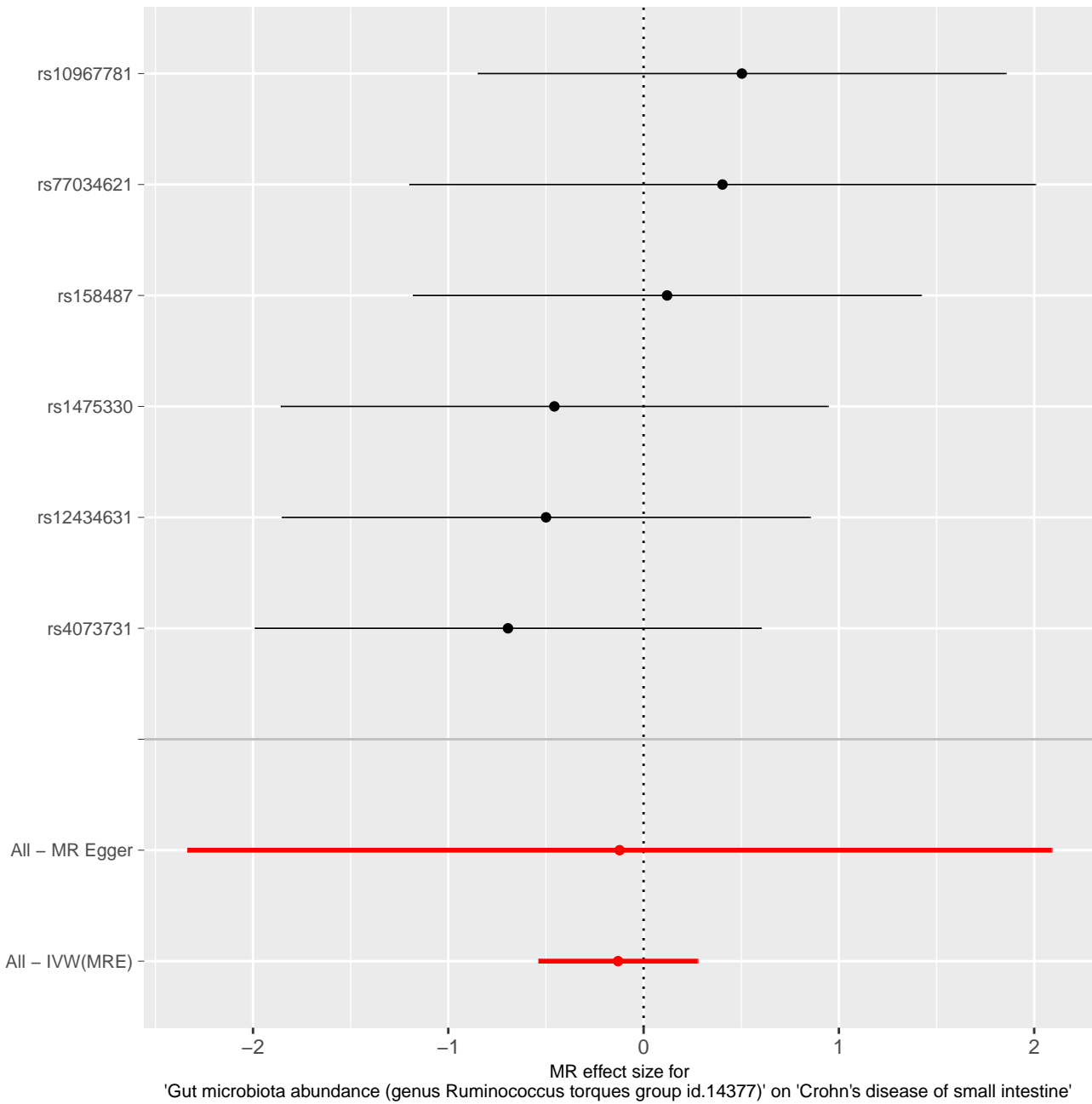
Batch 438 : Gut microbiota abundance (genus Ruminococcus gauvreauii group id.11342) on Crohn's disease of small intestine

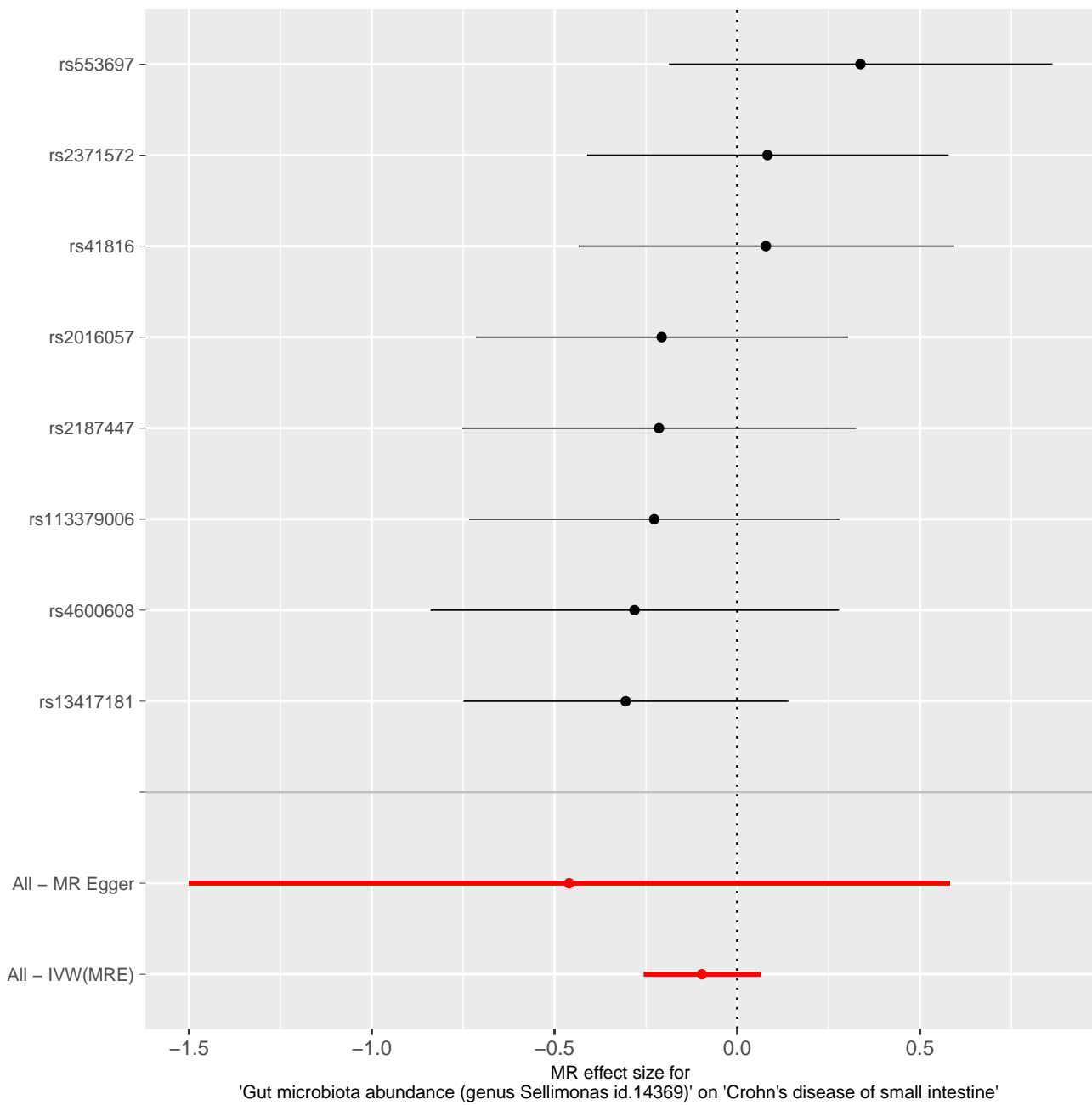


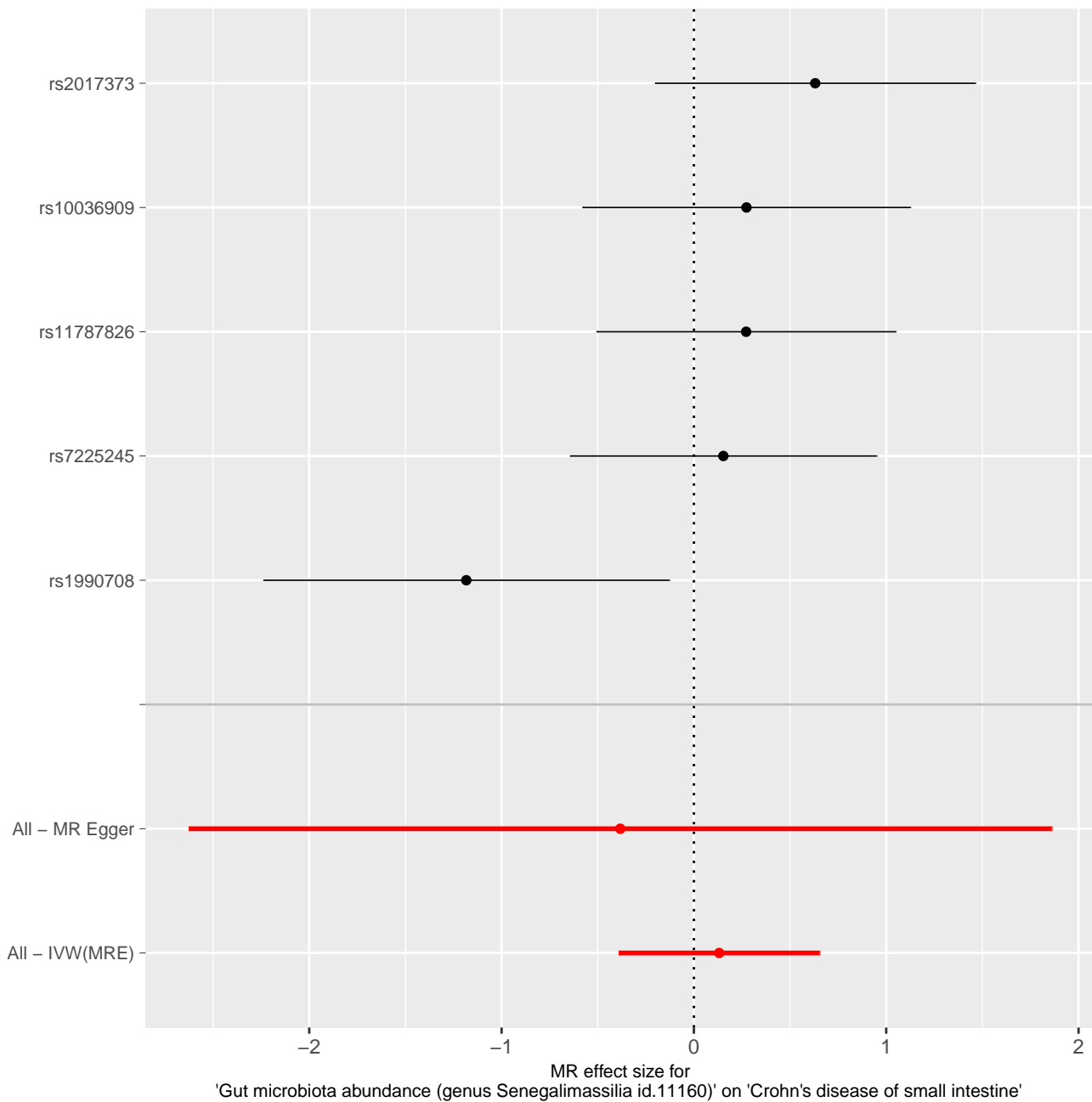
Batch 439 : Gut microbiota abundance (genus Ruminococcus gnnavus group id.14376) on Crohn's disease of small intestine

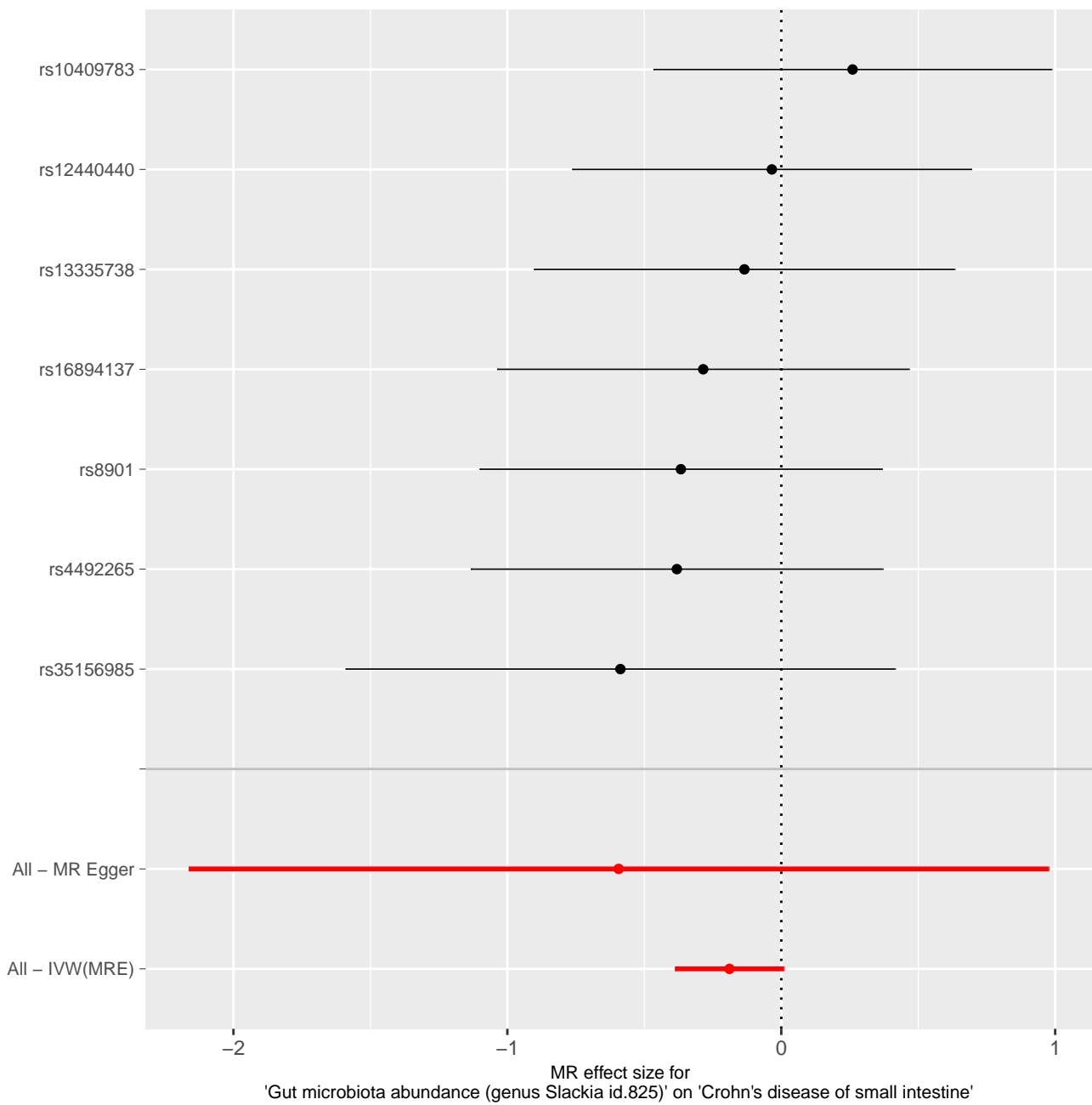


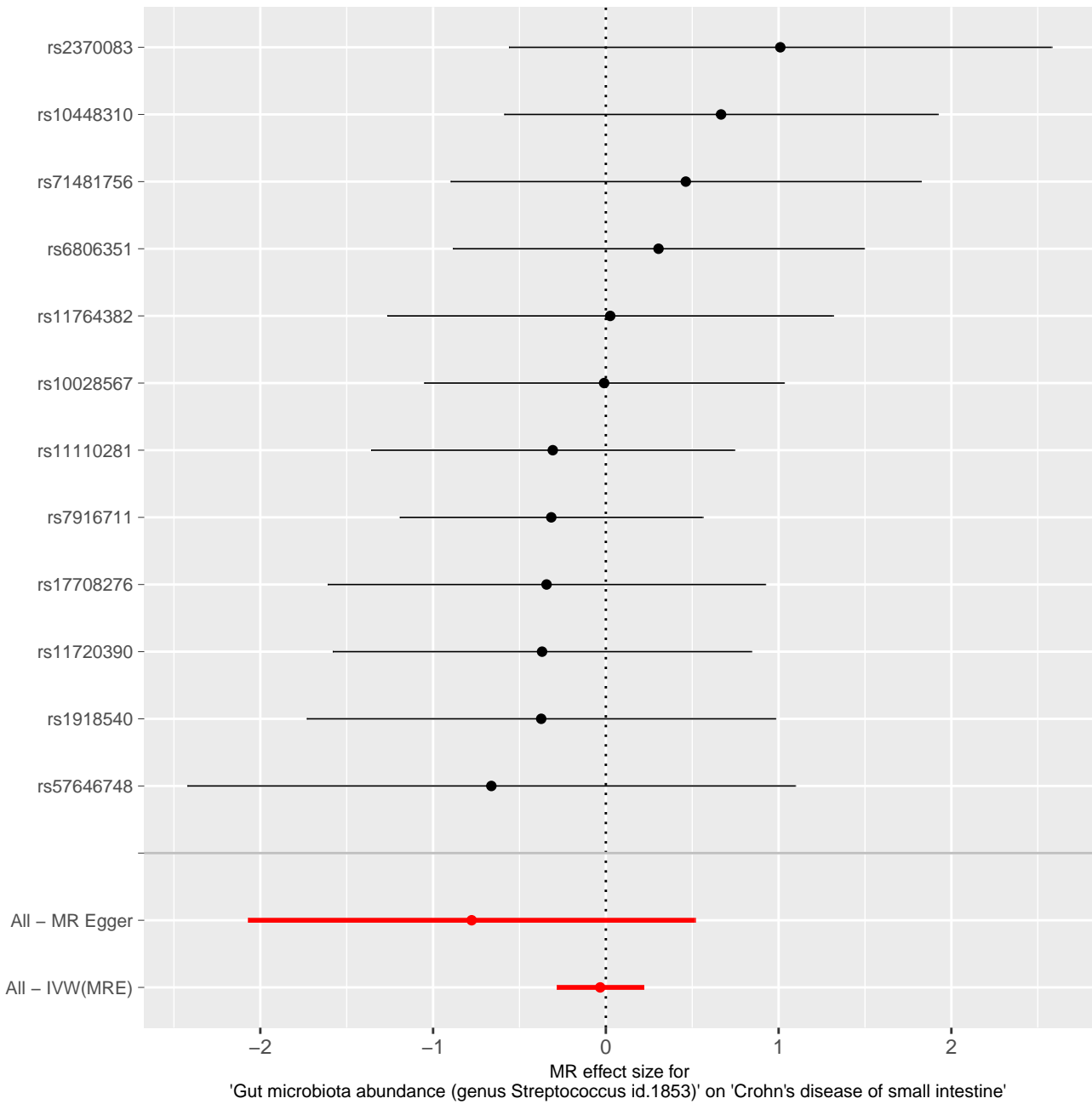
MR effect size for 'Gut microbiota abundance (genus Ruminococcus gnnavus group id.14376)' on 'Crohn's disease of small intestine'

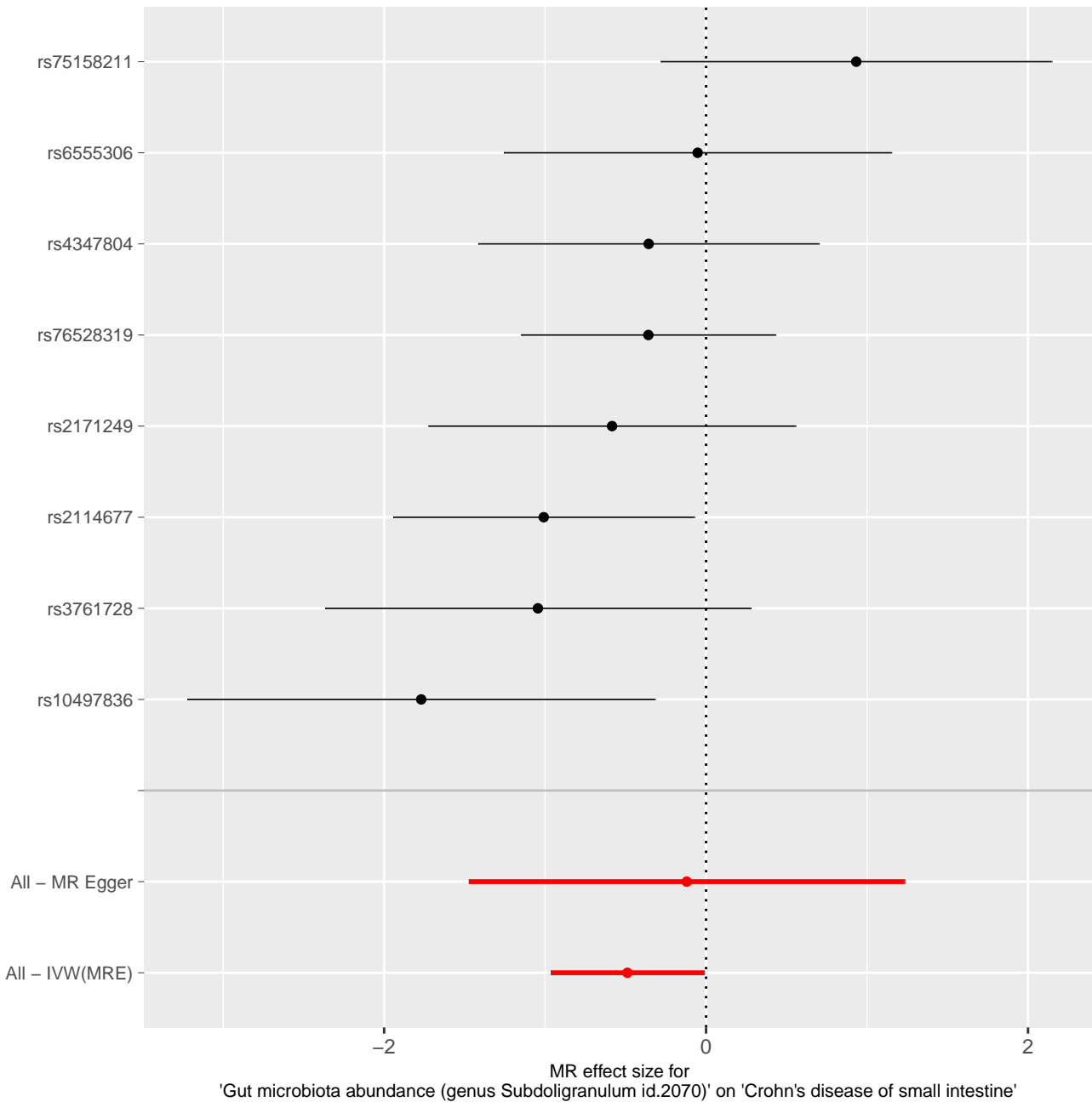




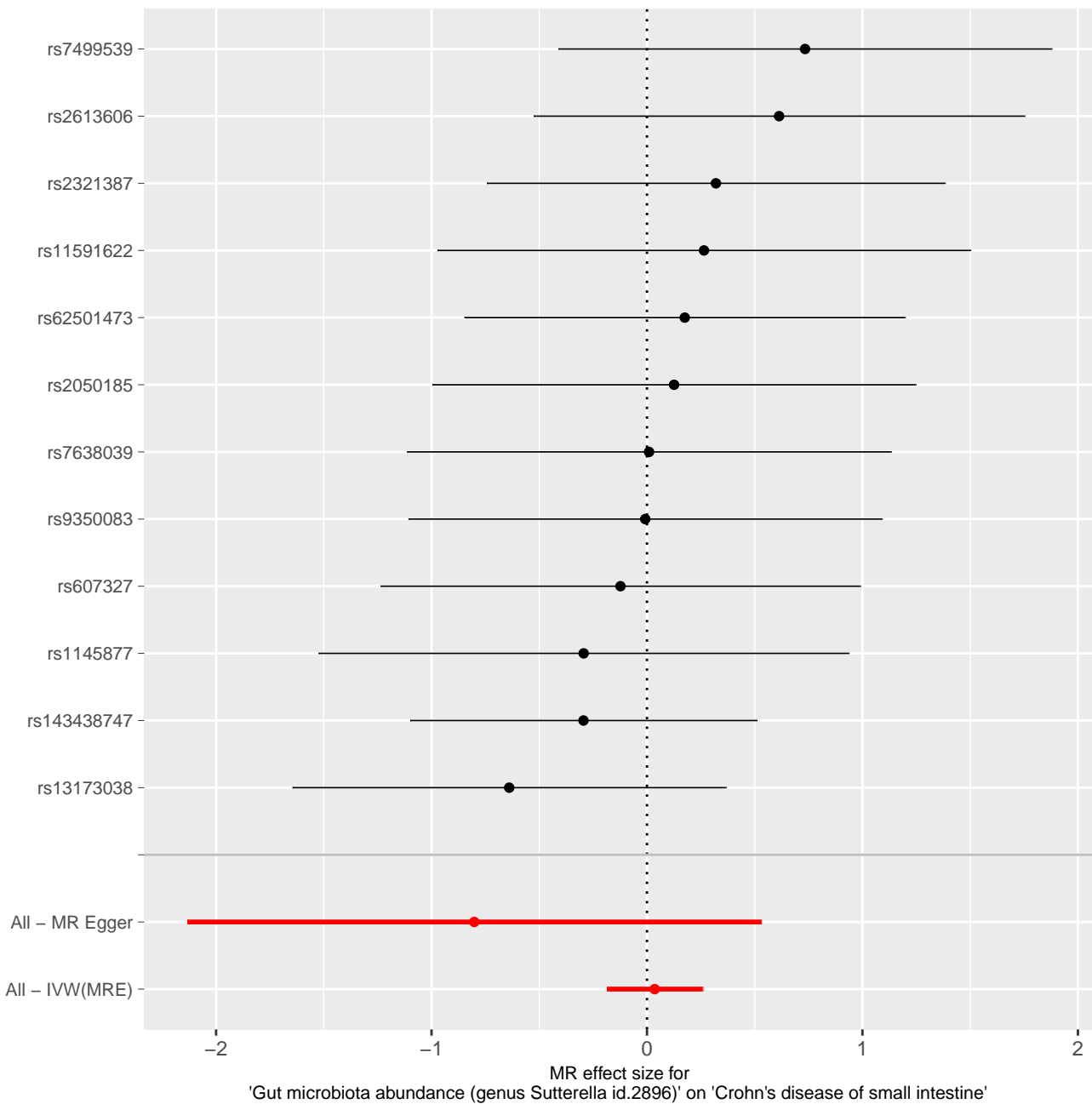


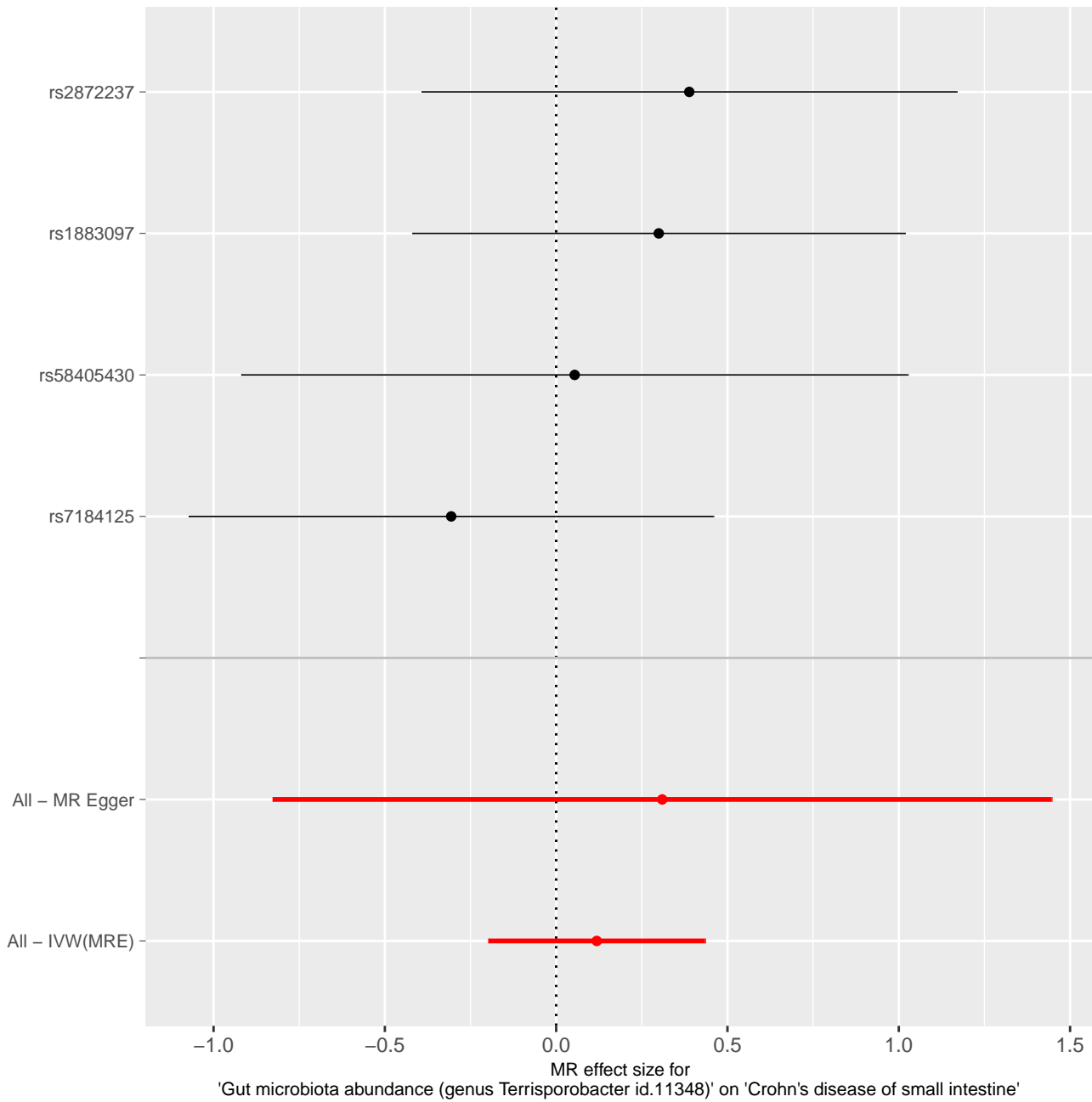


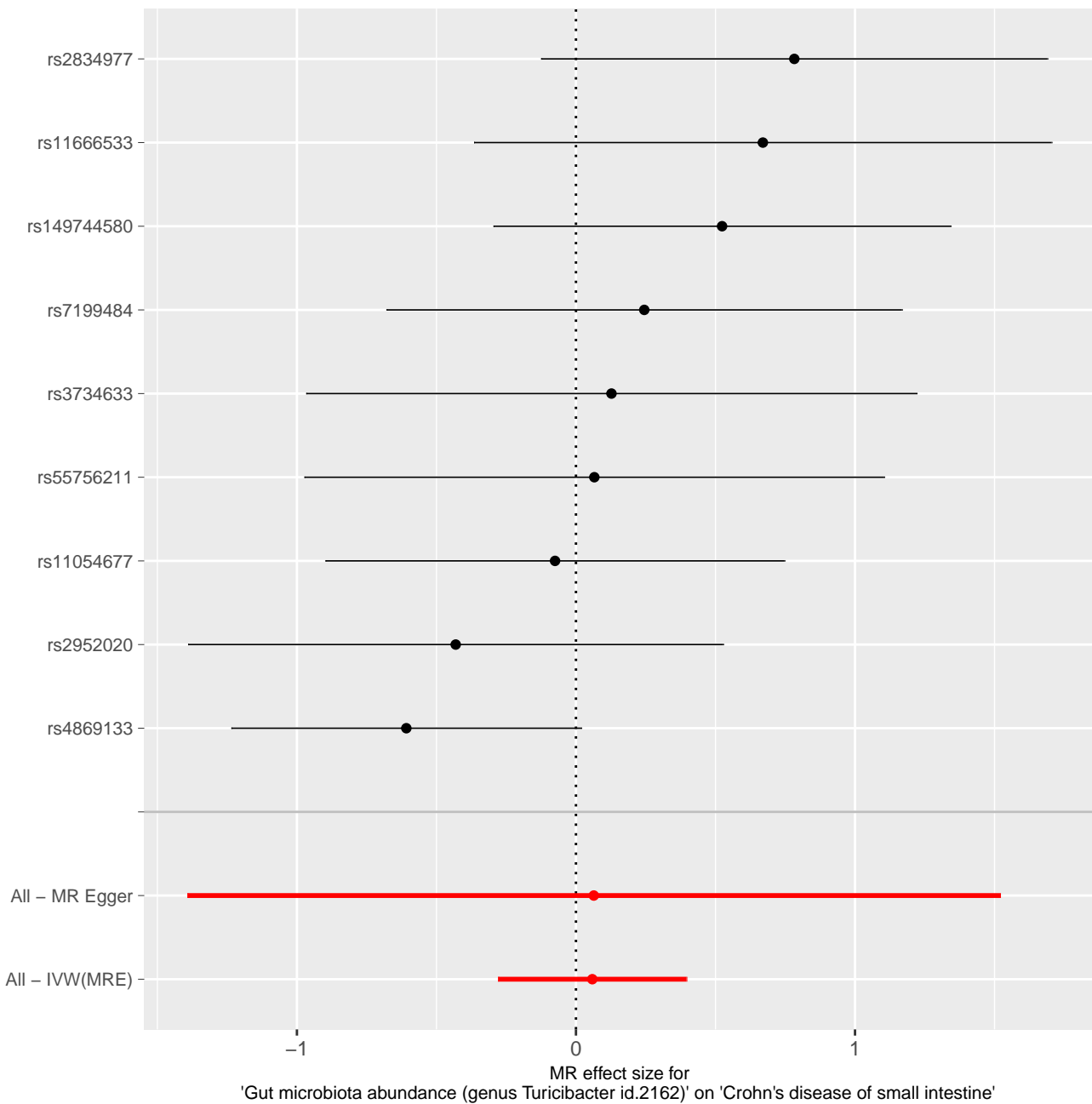




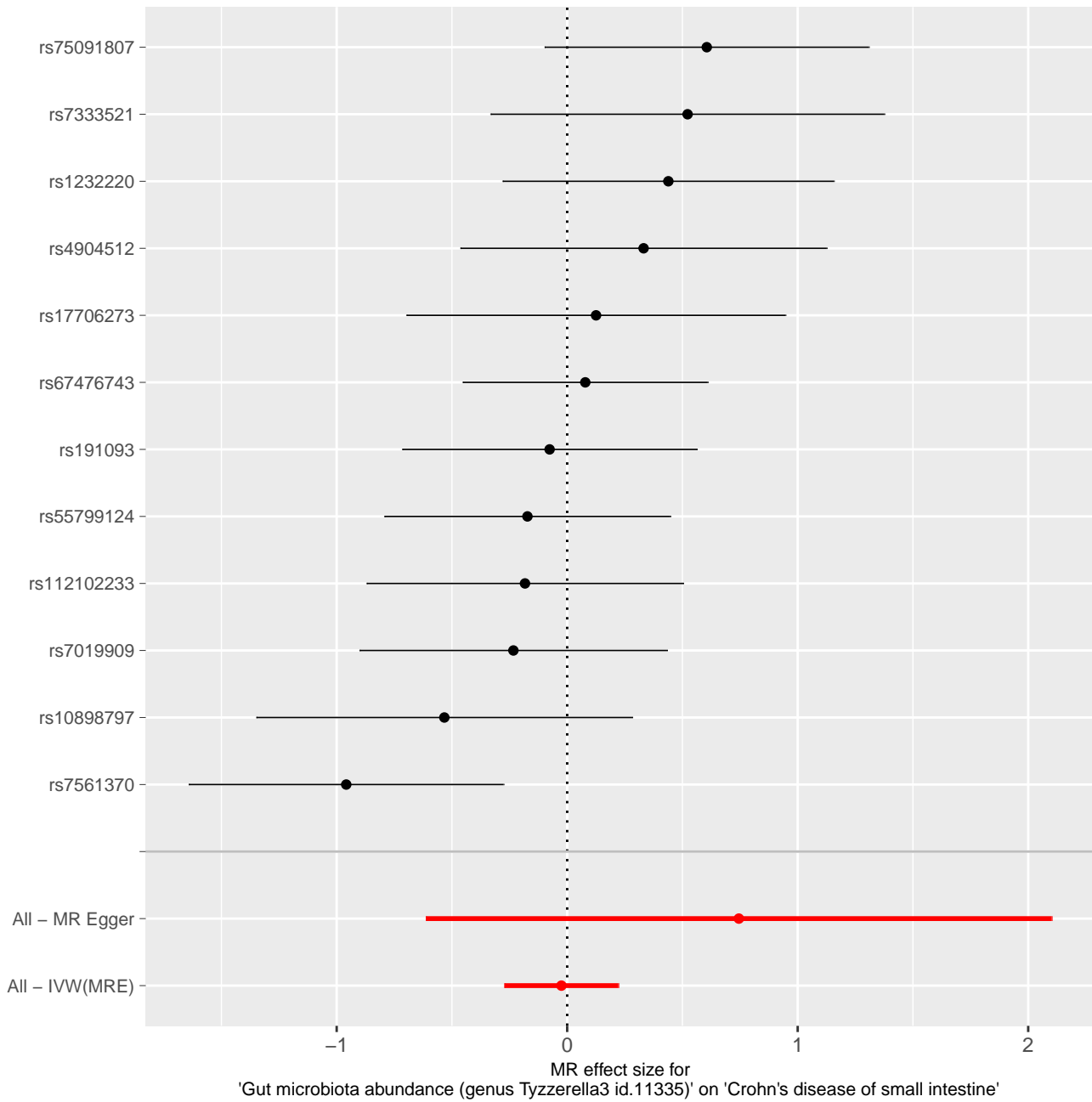
Batch 446 : Gut microbiota abundance (genus Sutterella id.2896) on Crohn's disease of small intestine

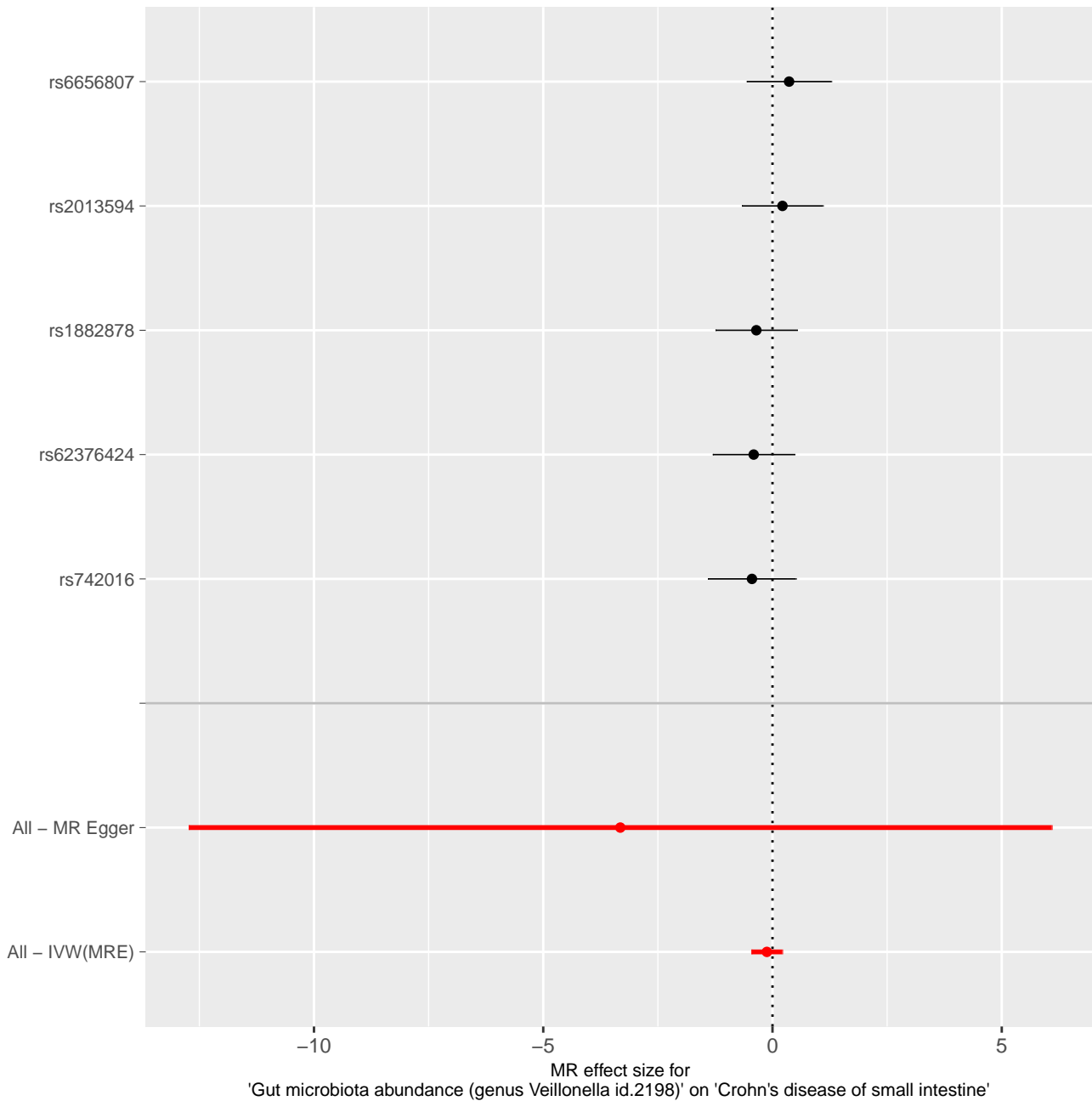


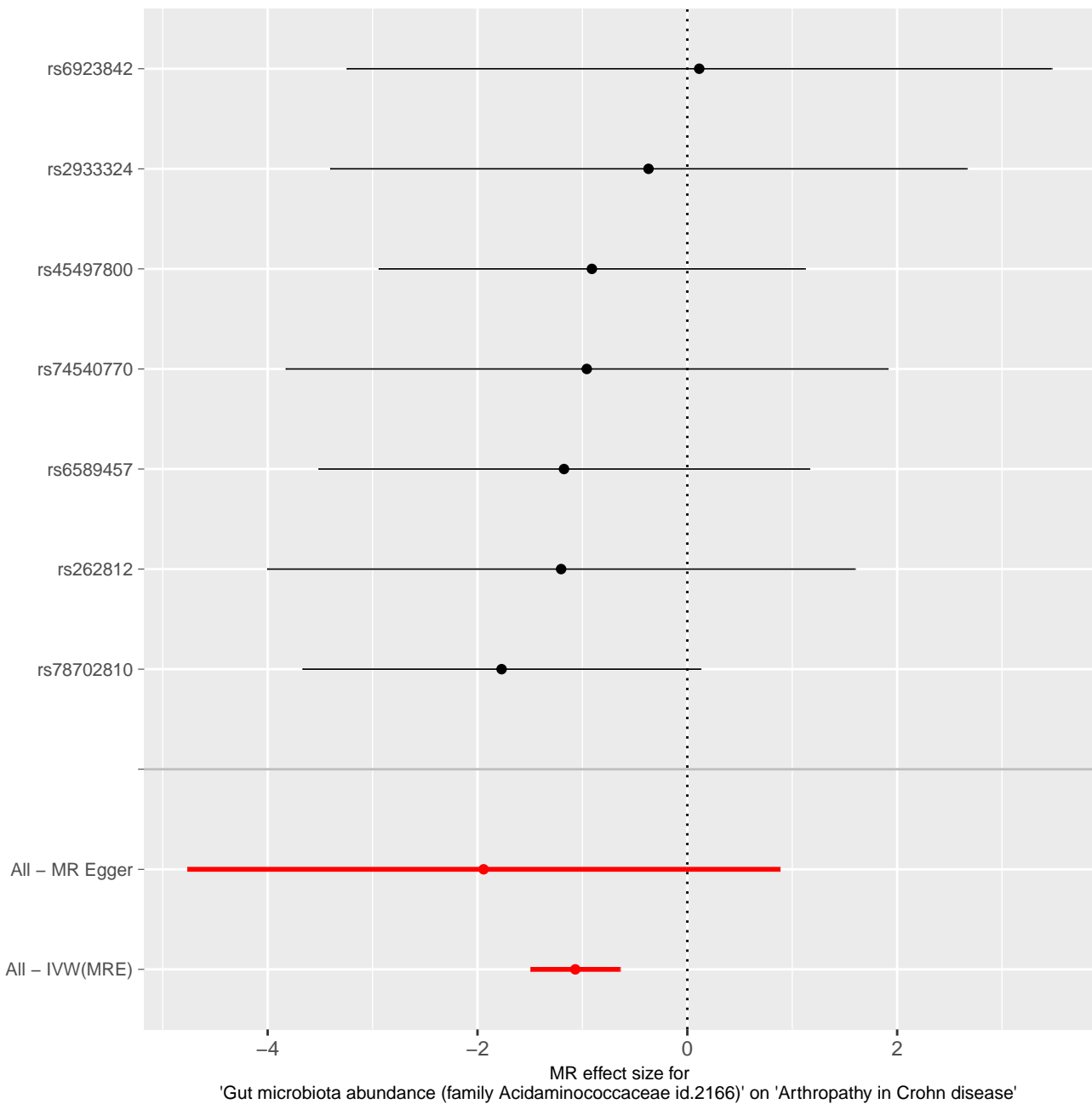


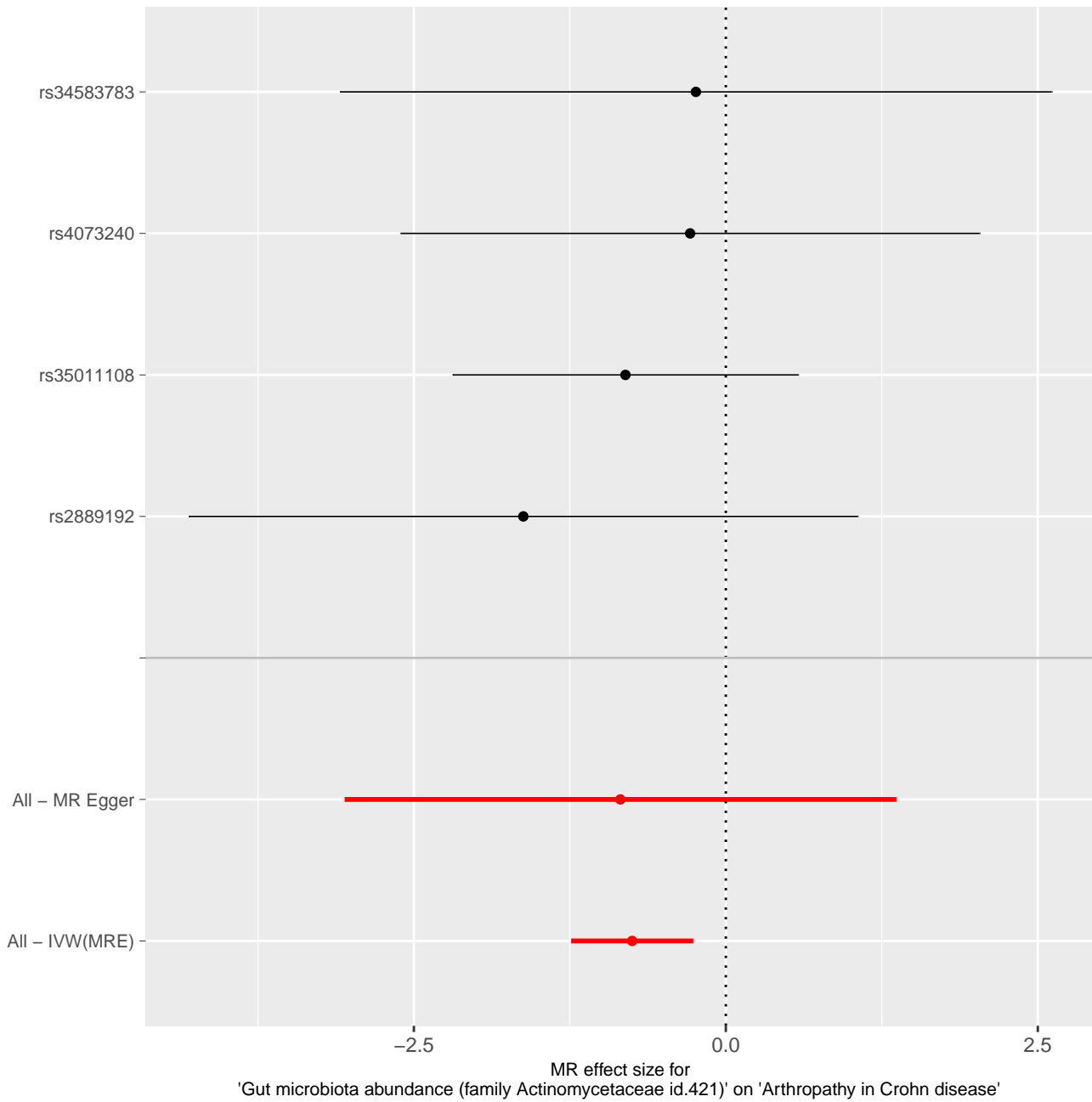


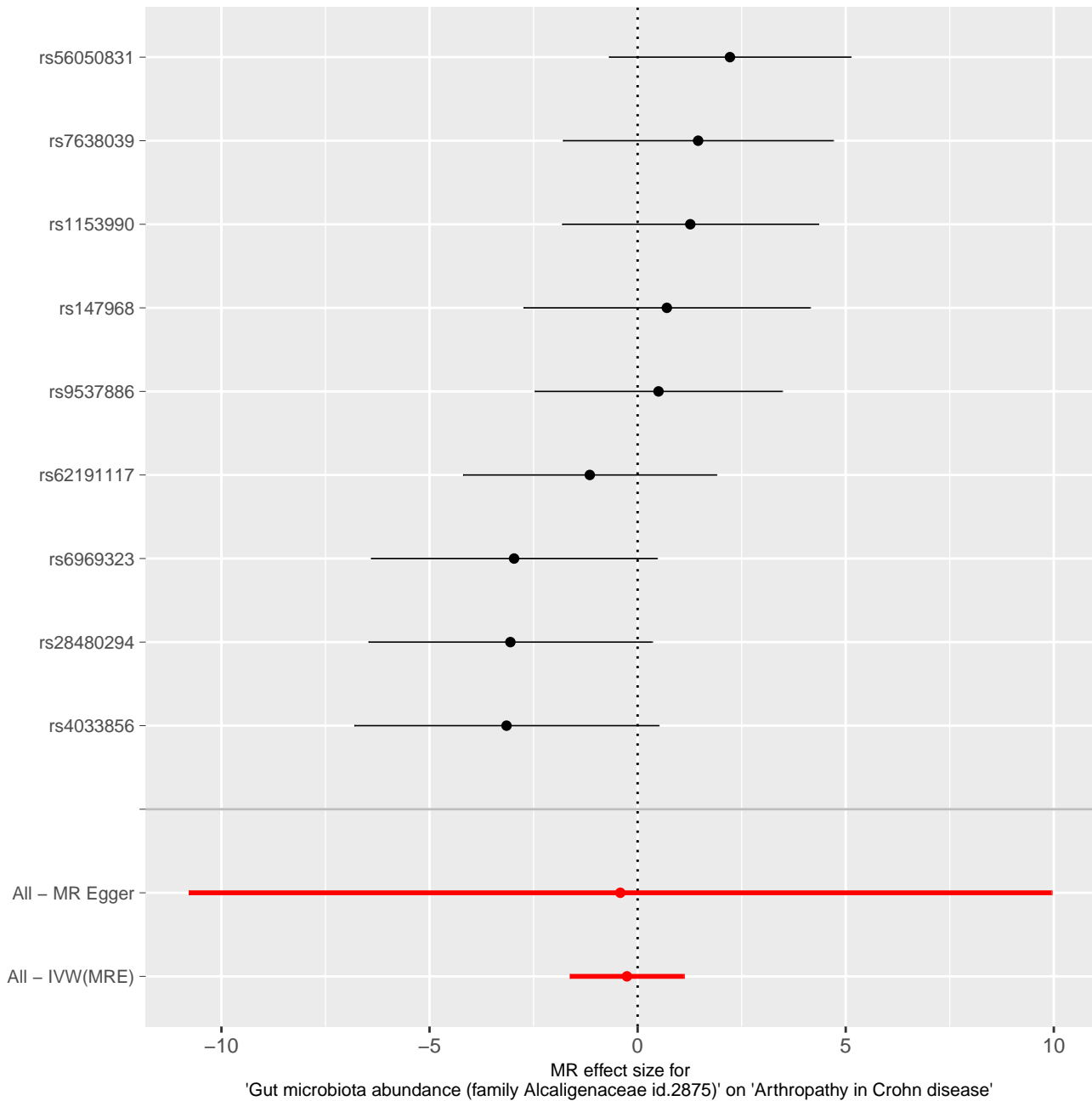
Batch 449 : Gut microbiota abundance (genus Tyzzerella3 id.11335) on Crohn's disease of small intestine

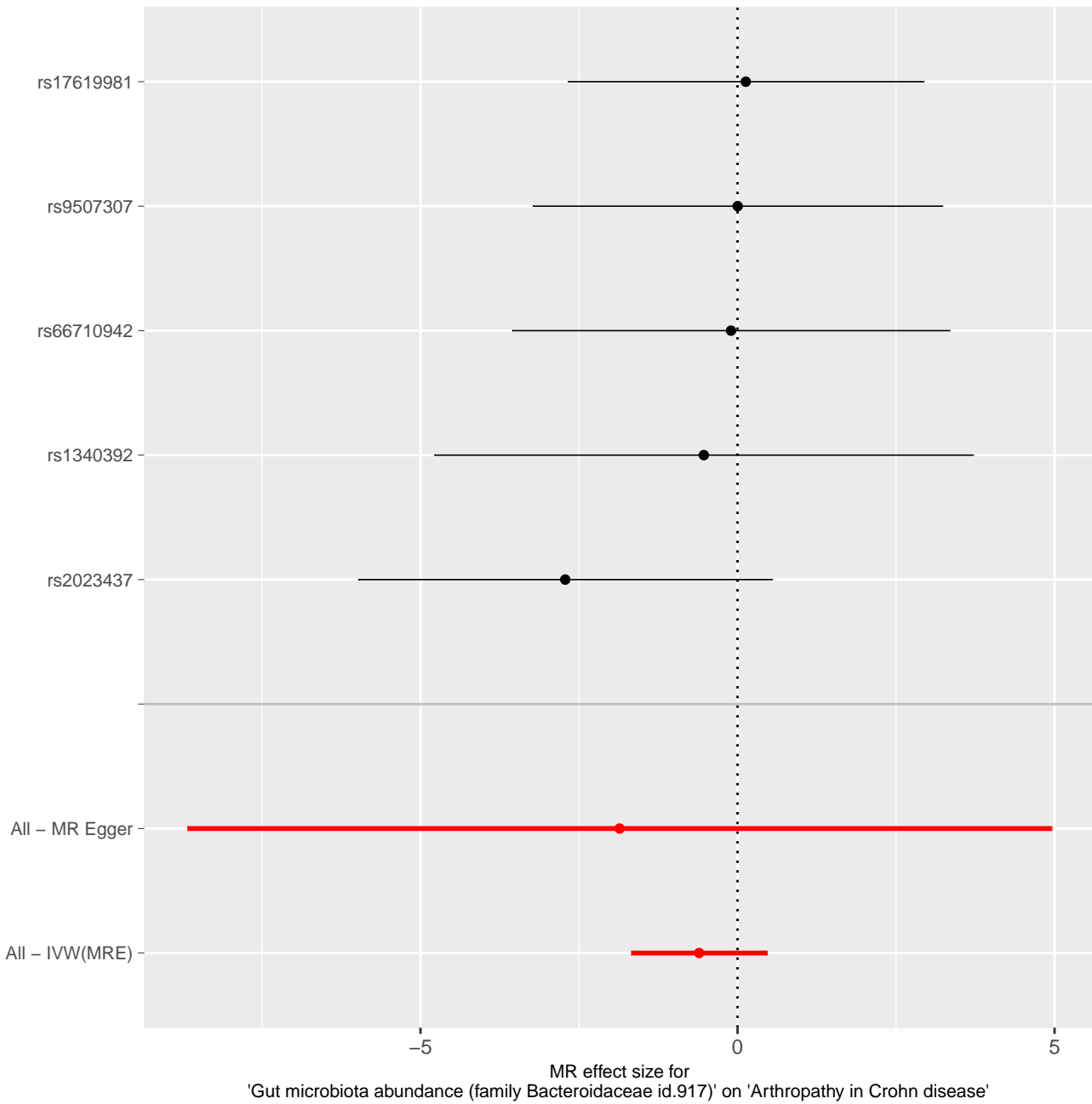


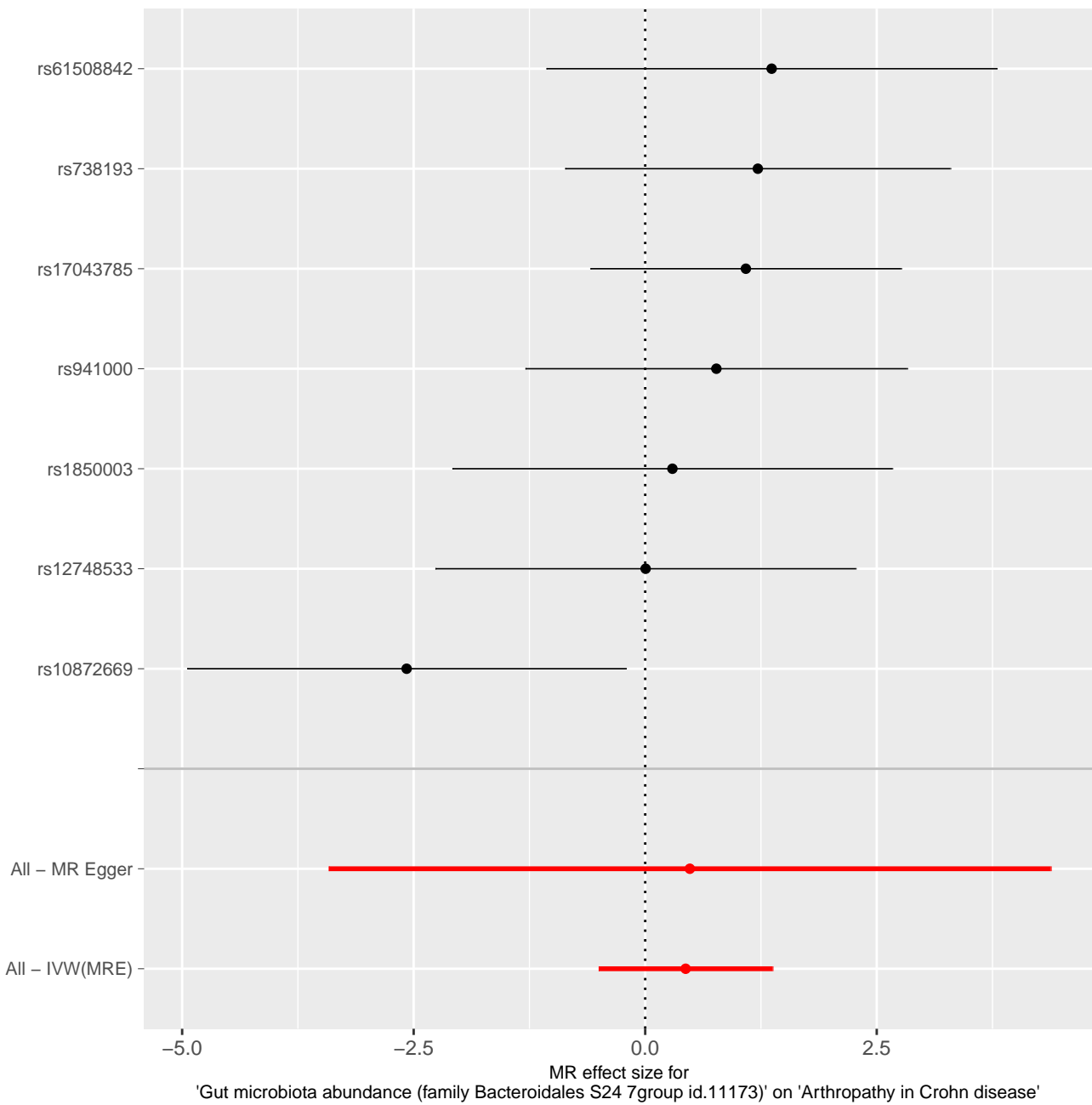


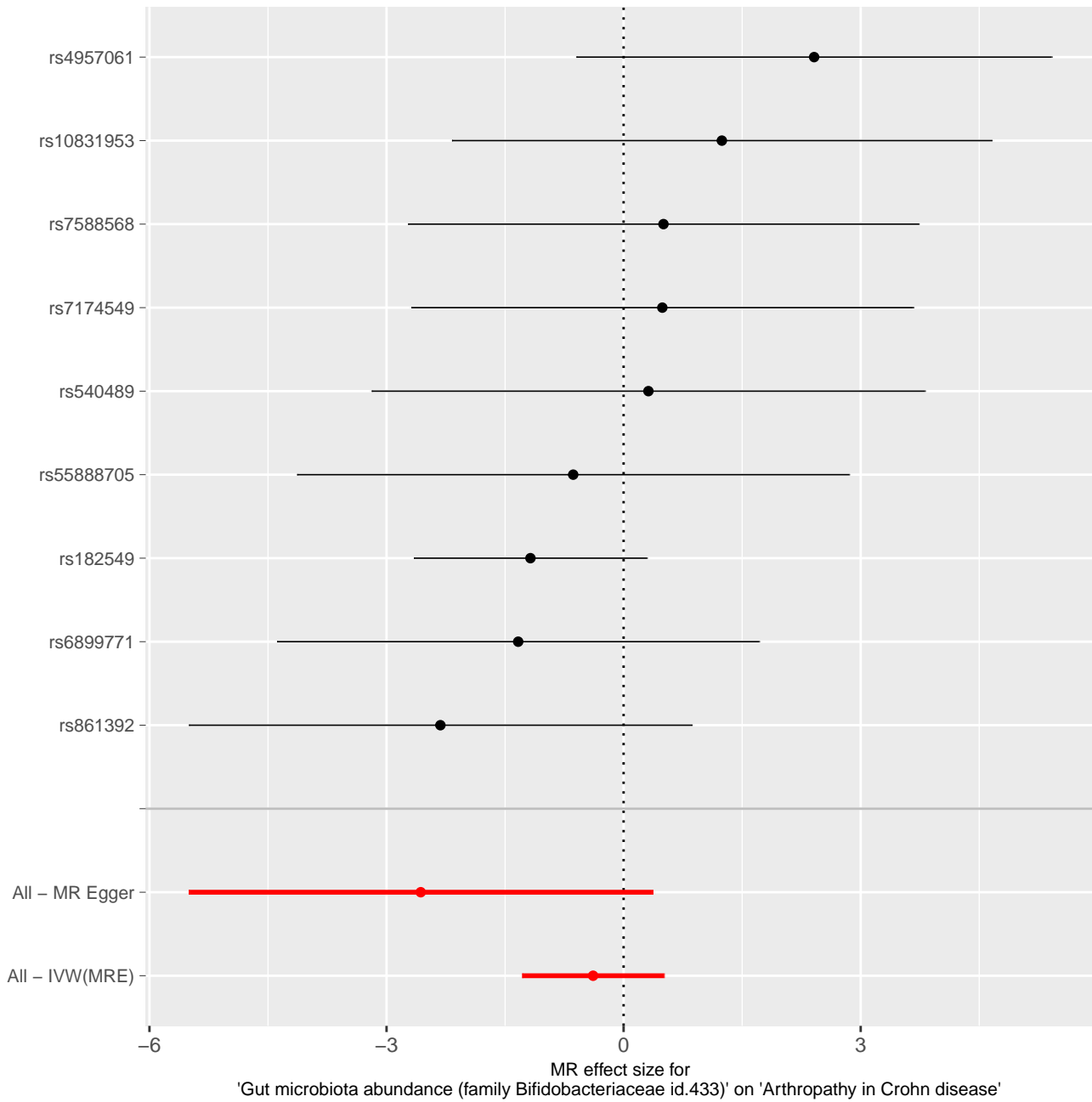


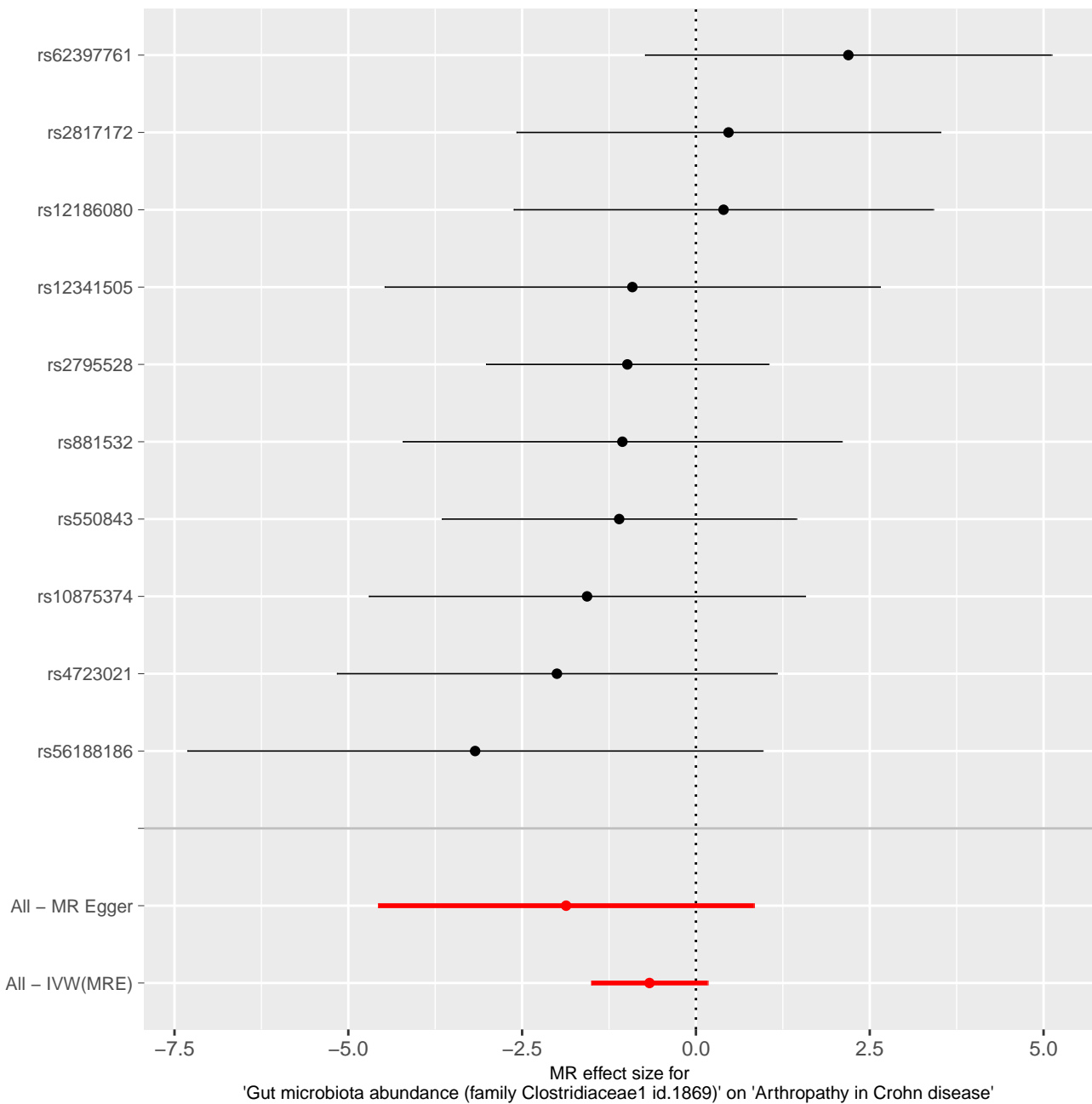


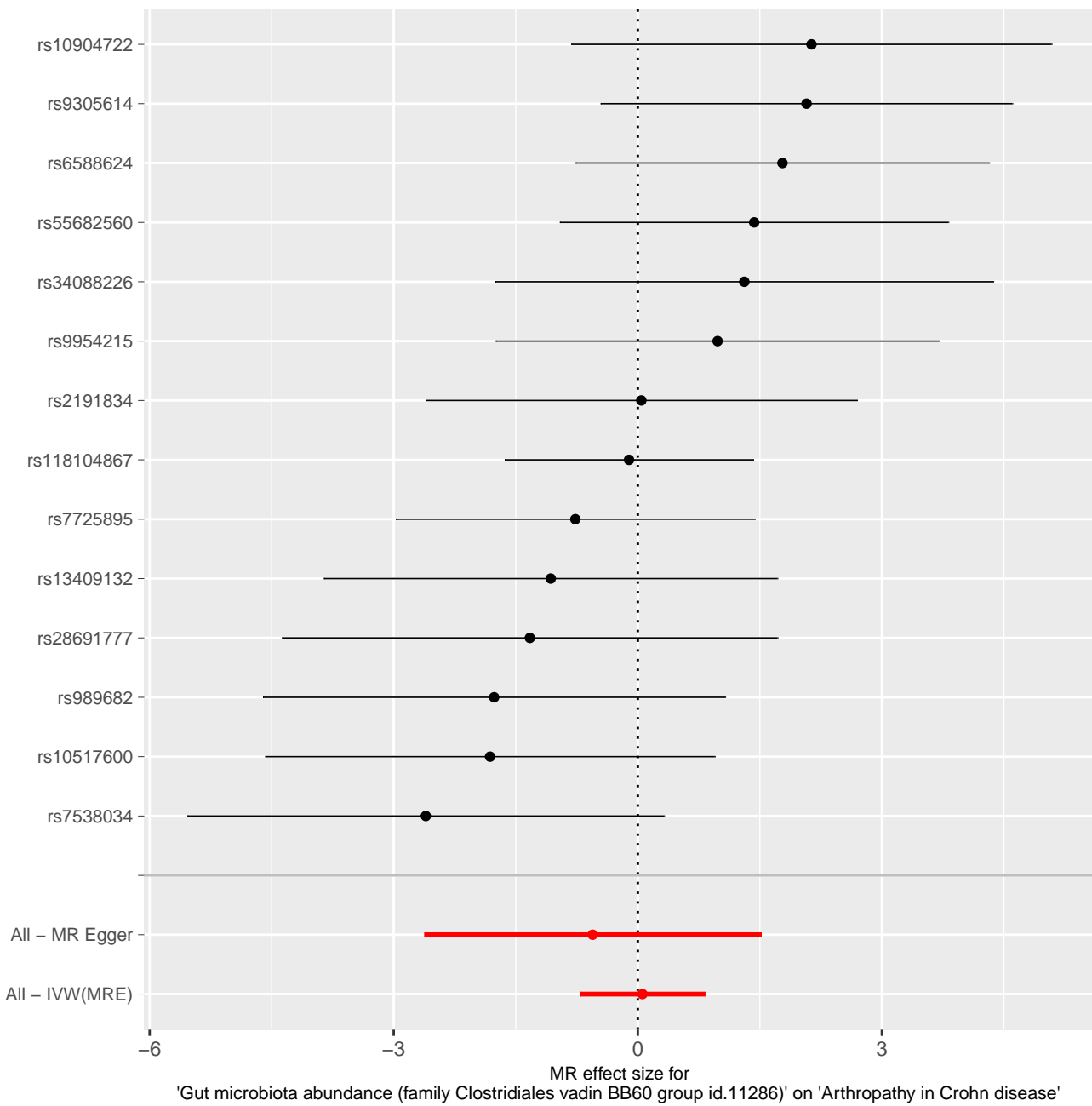




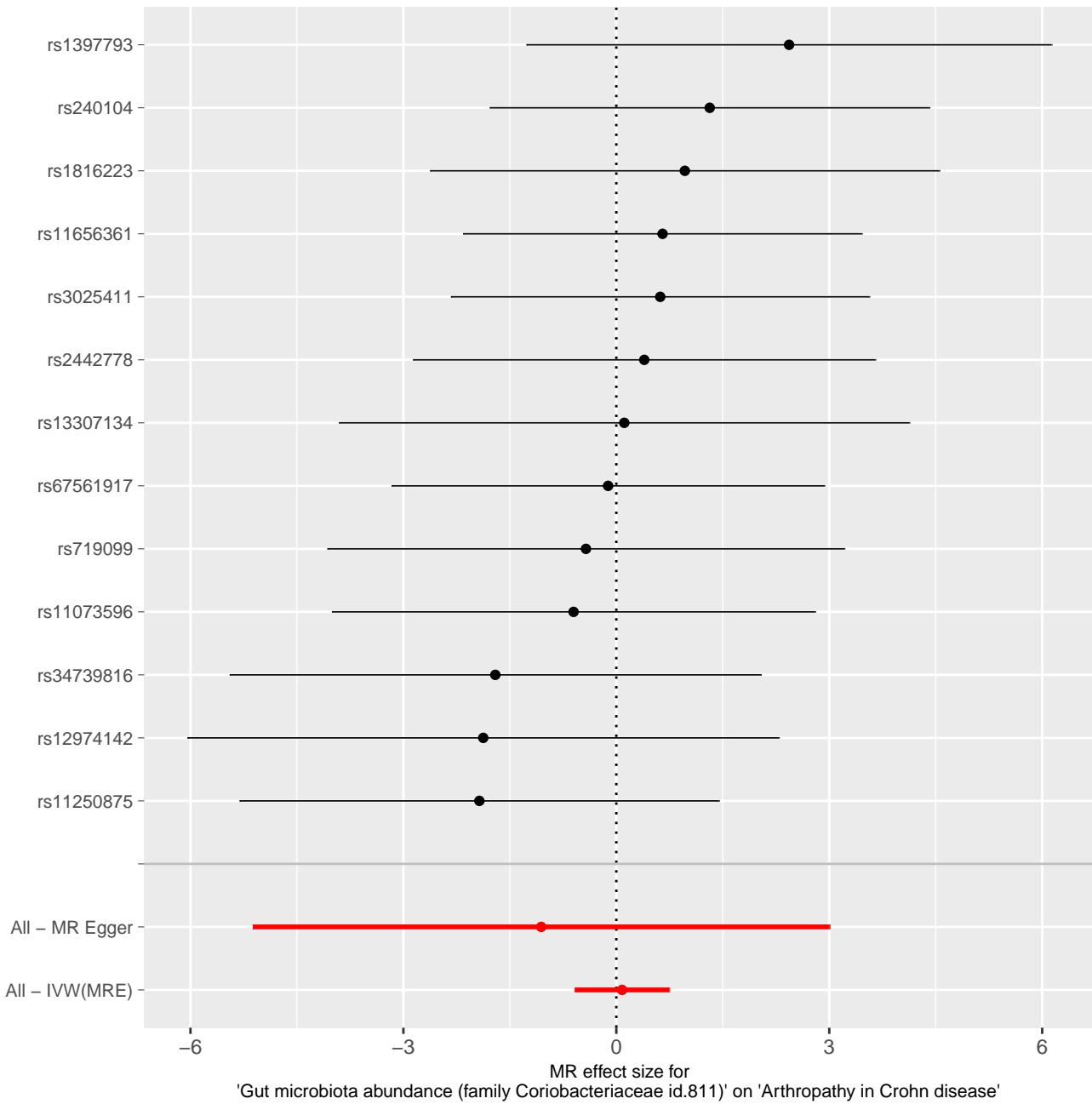




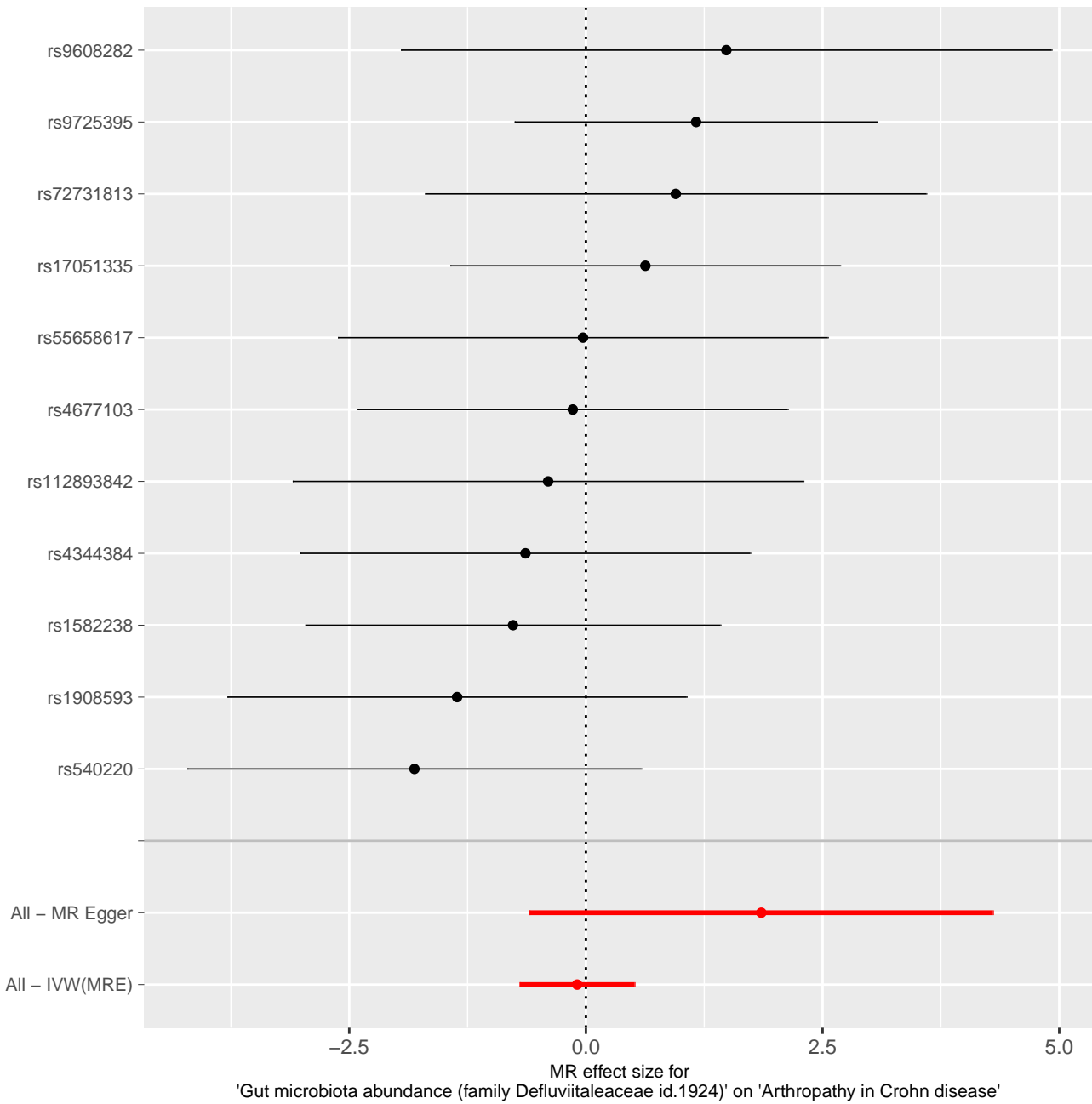




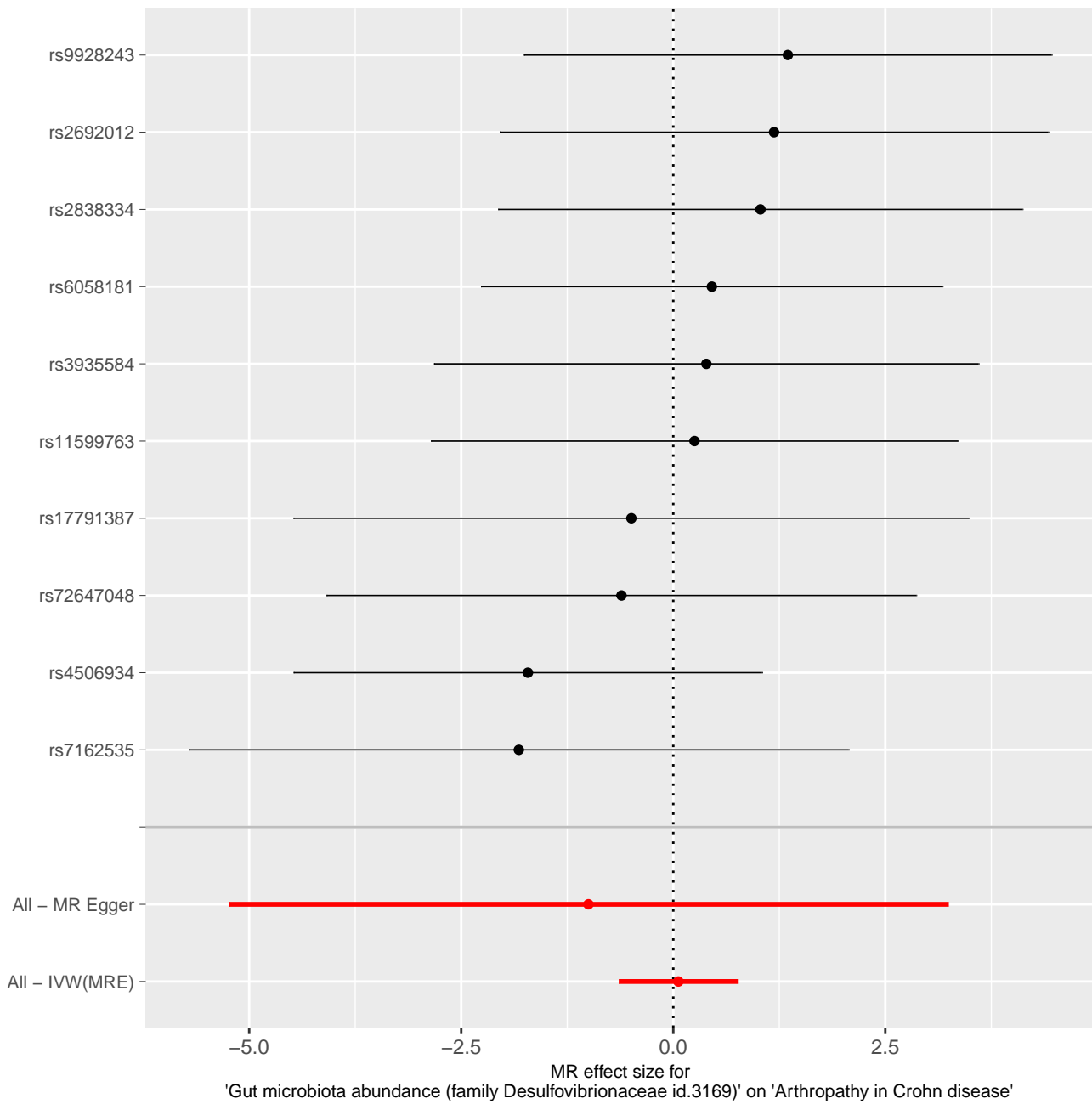
Batch 460 : Gut microbiota abundance (family Coriobacteriaceae id.811) on Arthropathy in Crohn disease

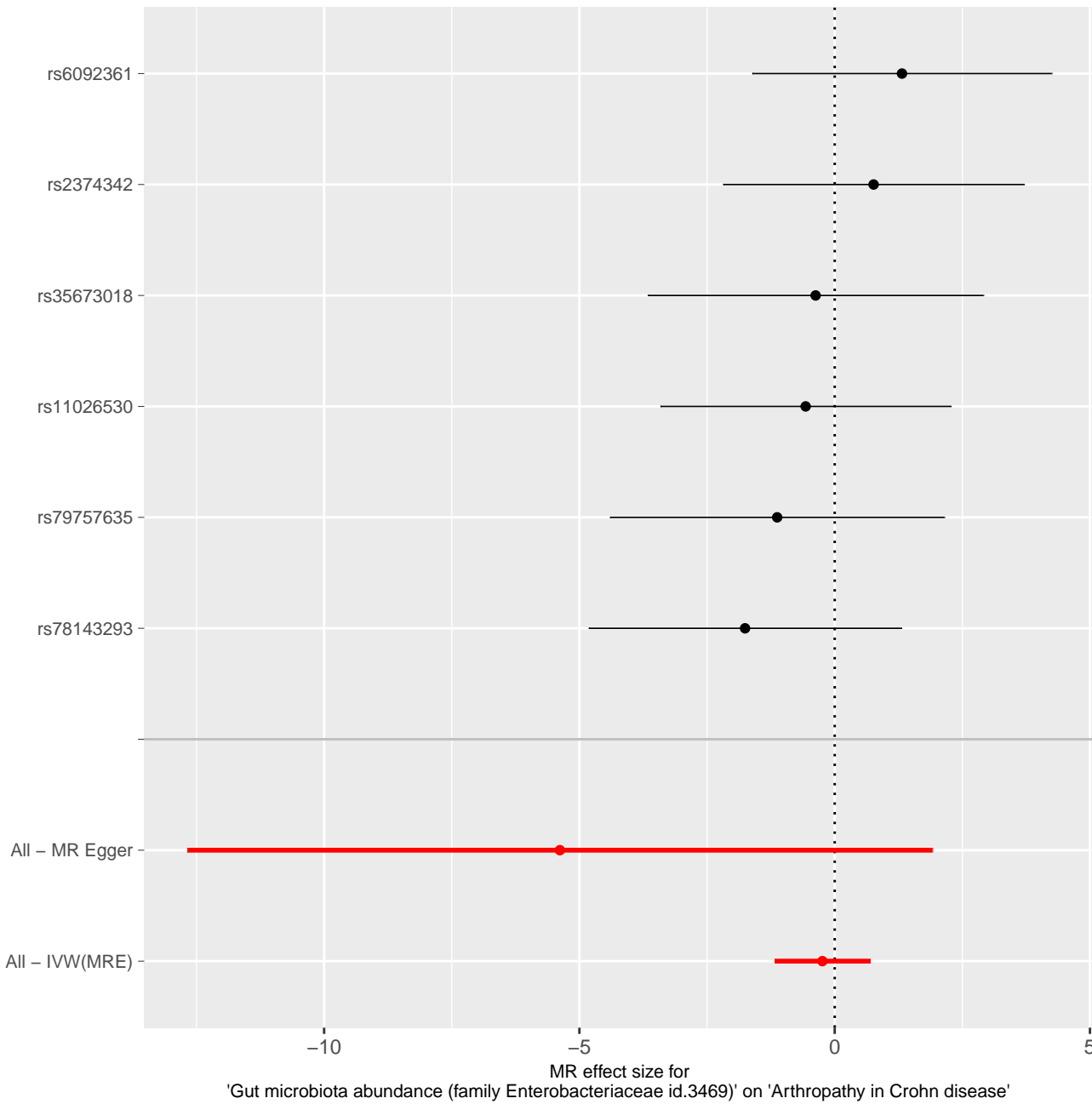


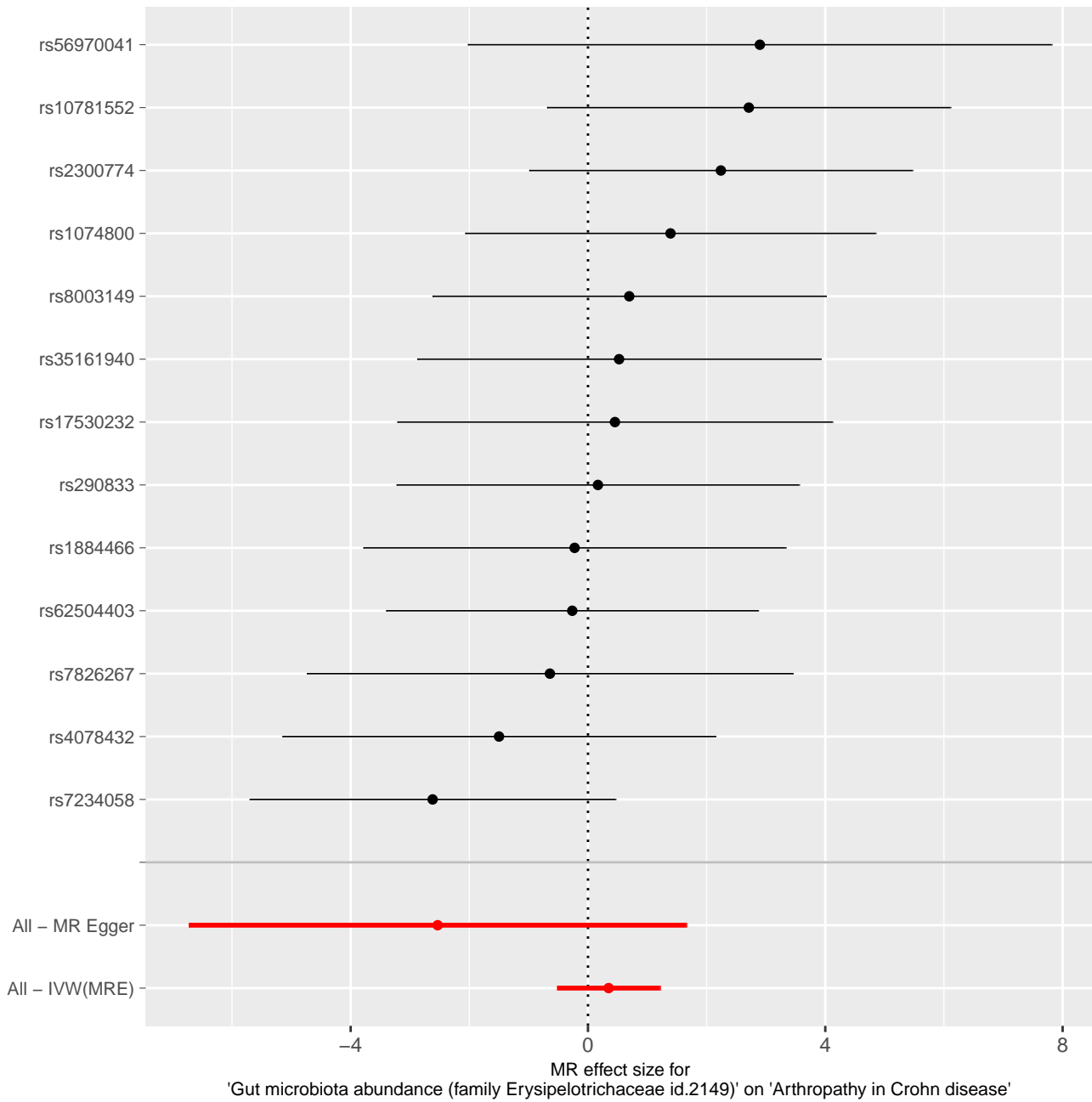
Batch 461 : Gut microbiota abundance (family Defluviitaleaceae id.1924) on Arthropathy in Crohn disease

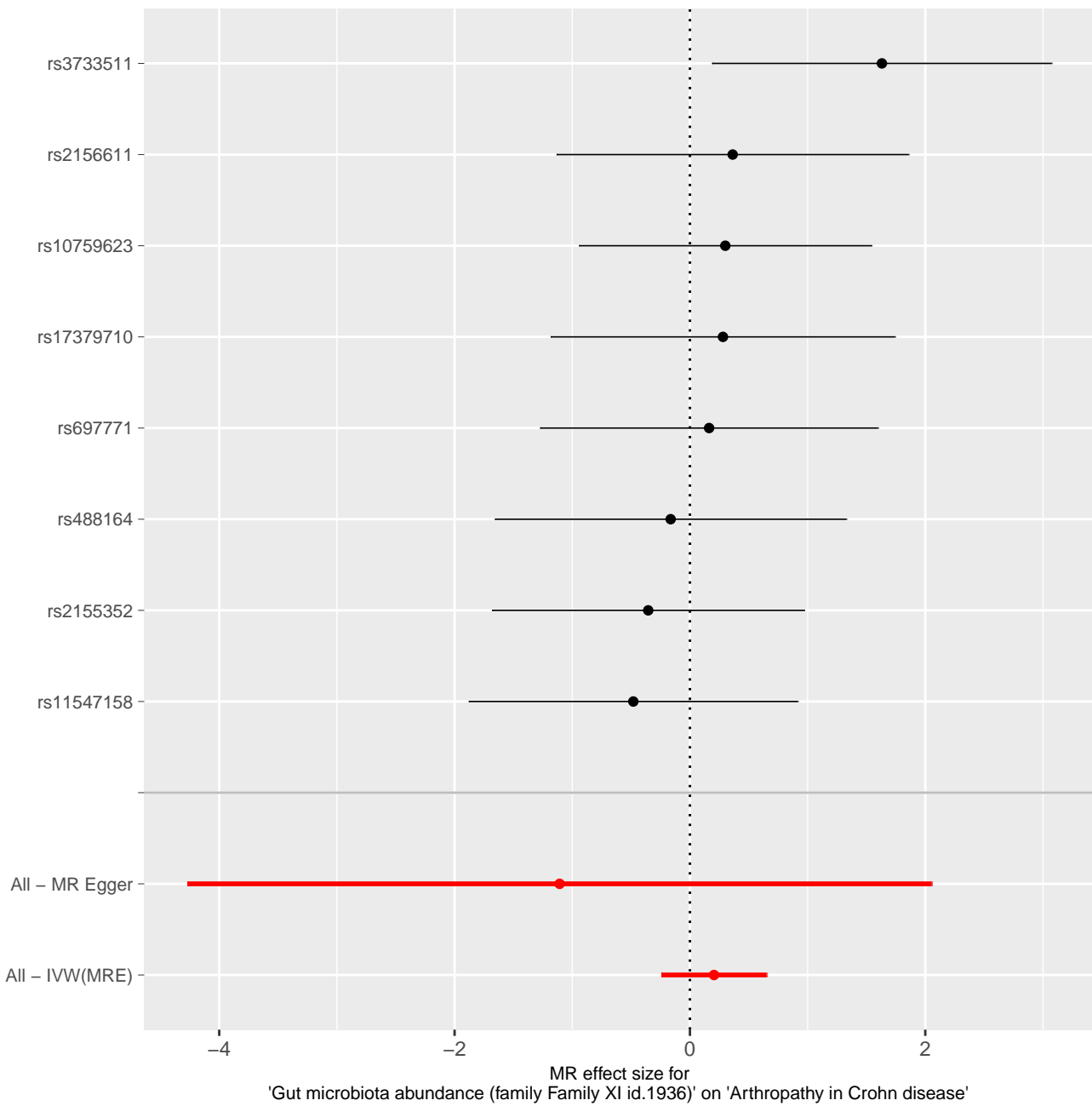


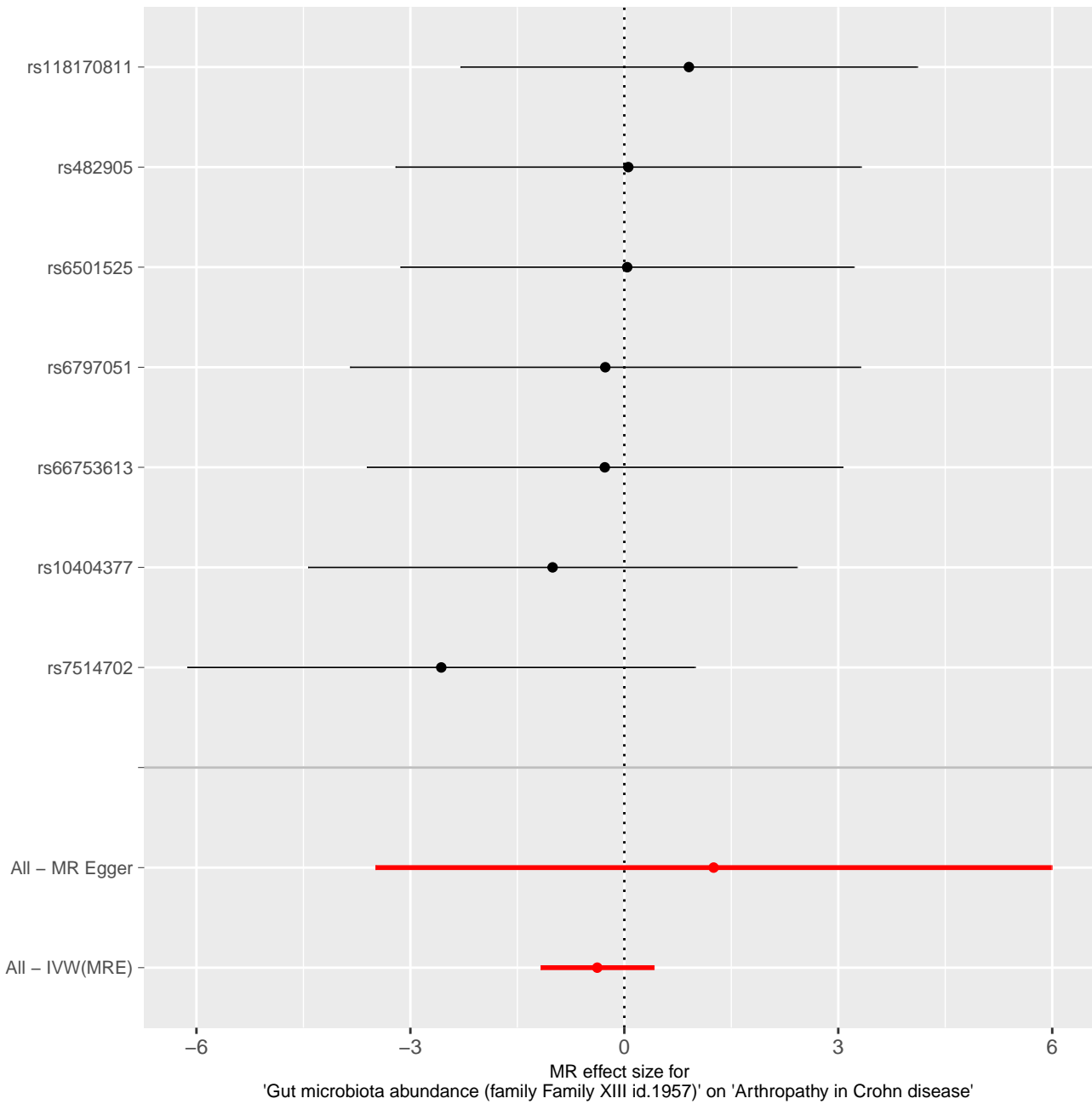
Batch 462 : Gut microbiota abundance (family Desulfovibrionaceae id.3169) on Arthropathy in Crohn disease



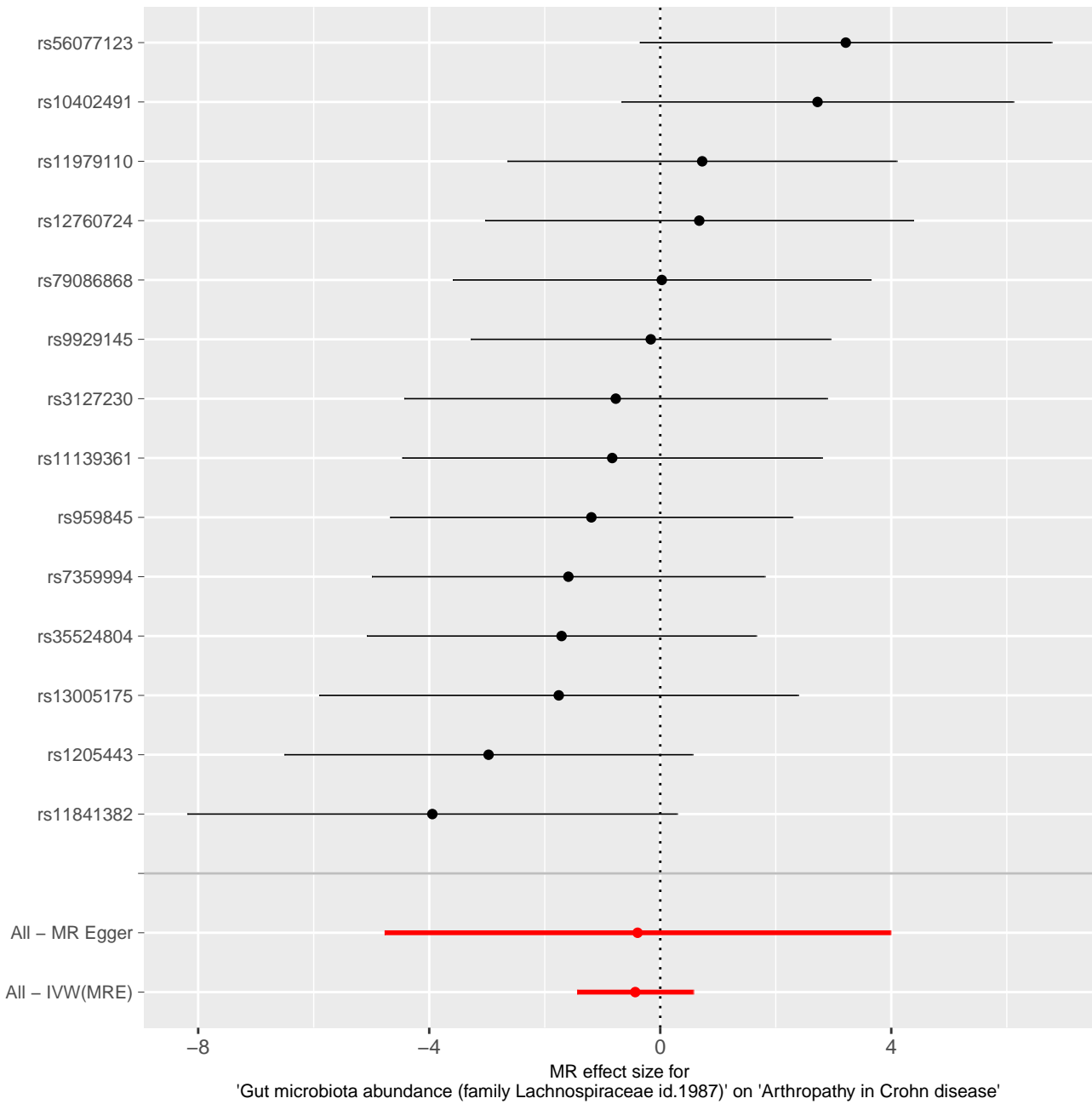


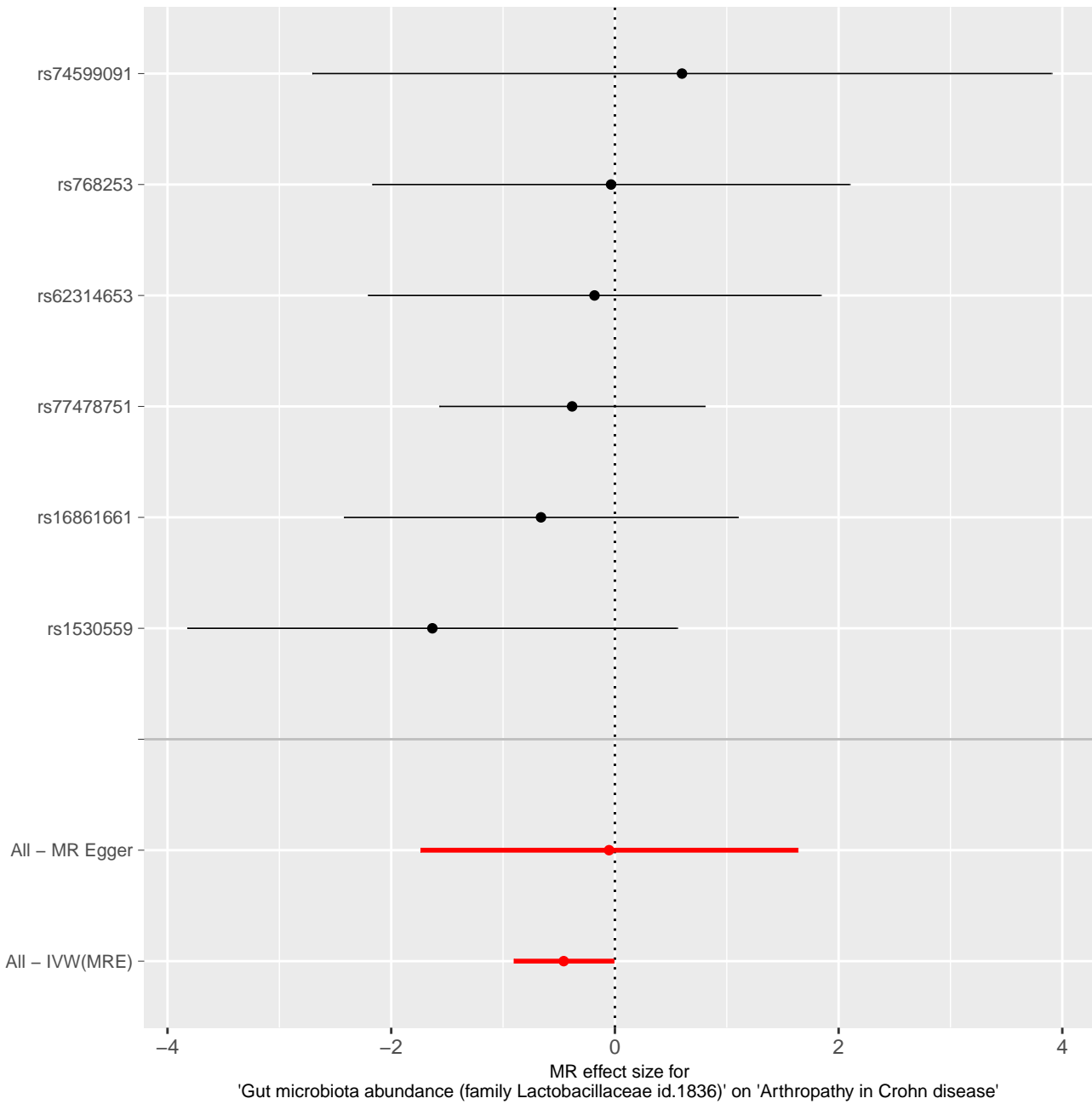


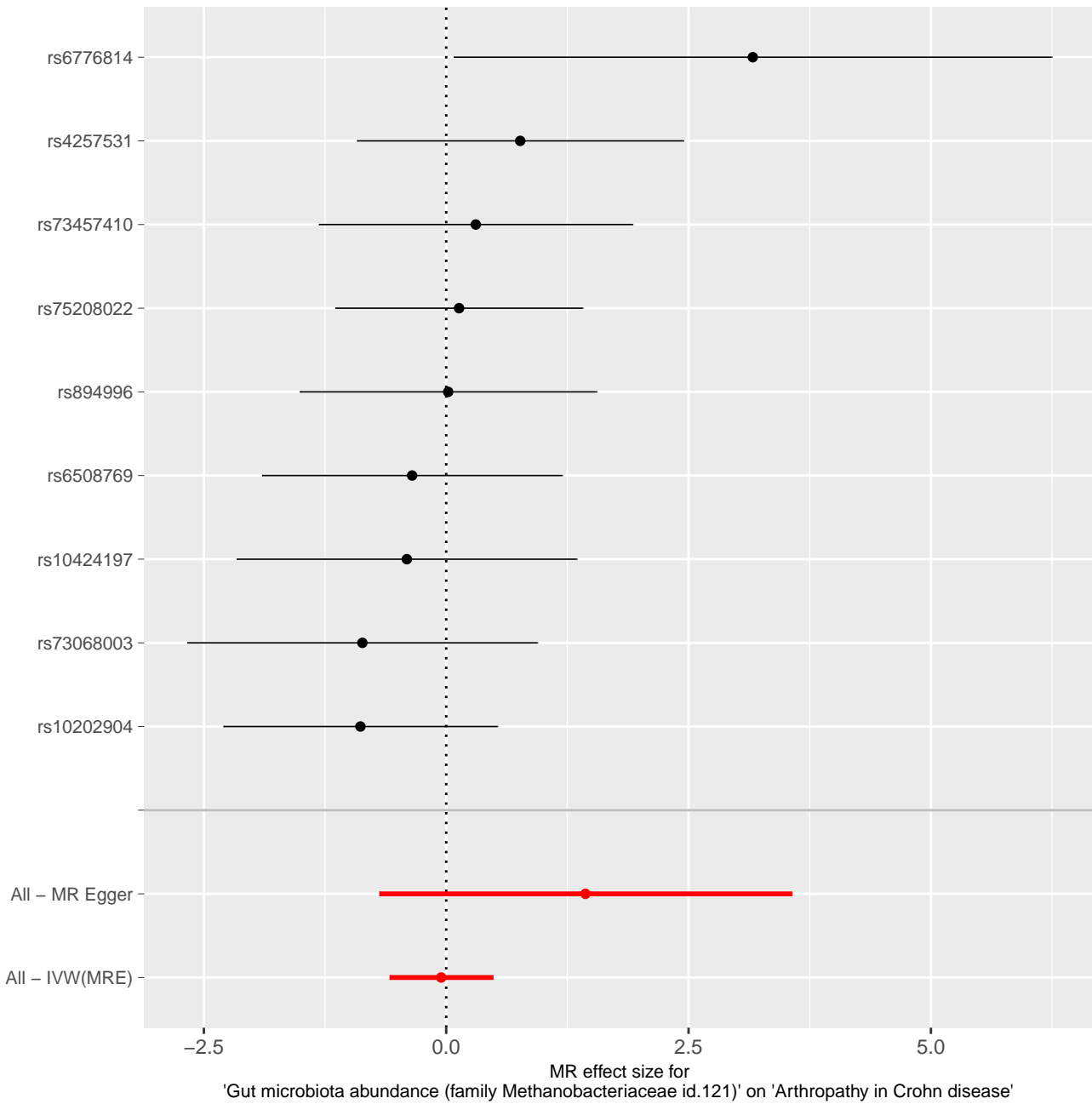


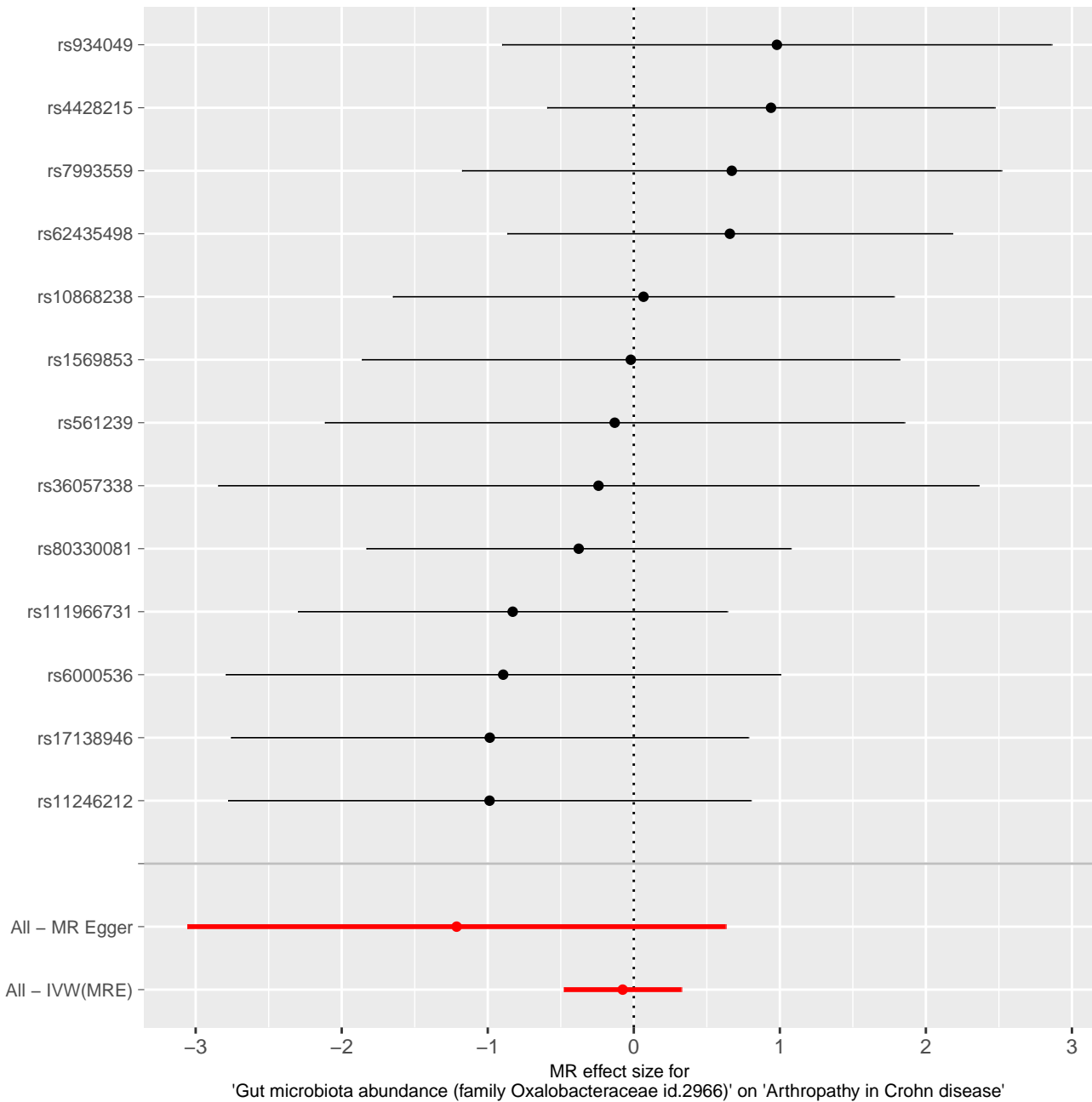


Batch 467 : Gut microbiota abundance (family Lachnospiraceae id.1987) on Arthropathy in Crohn disease

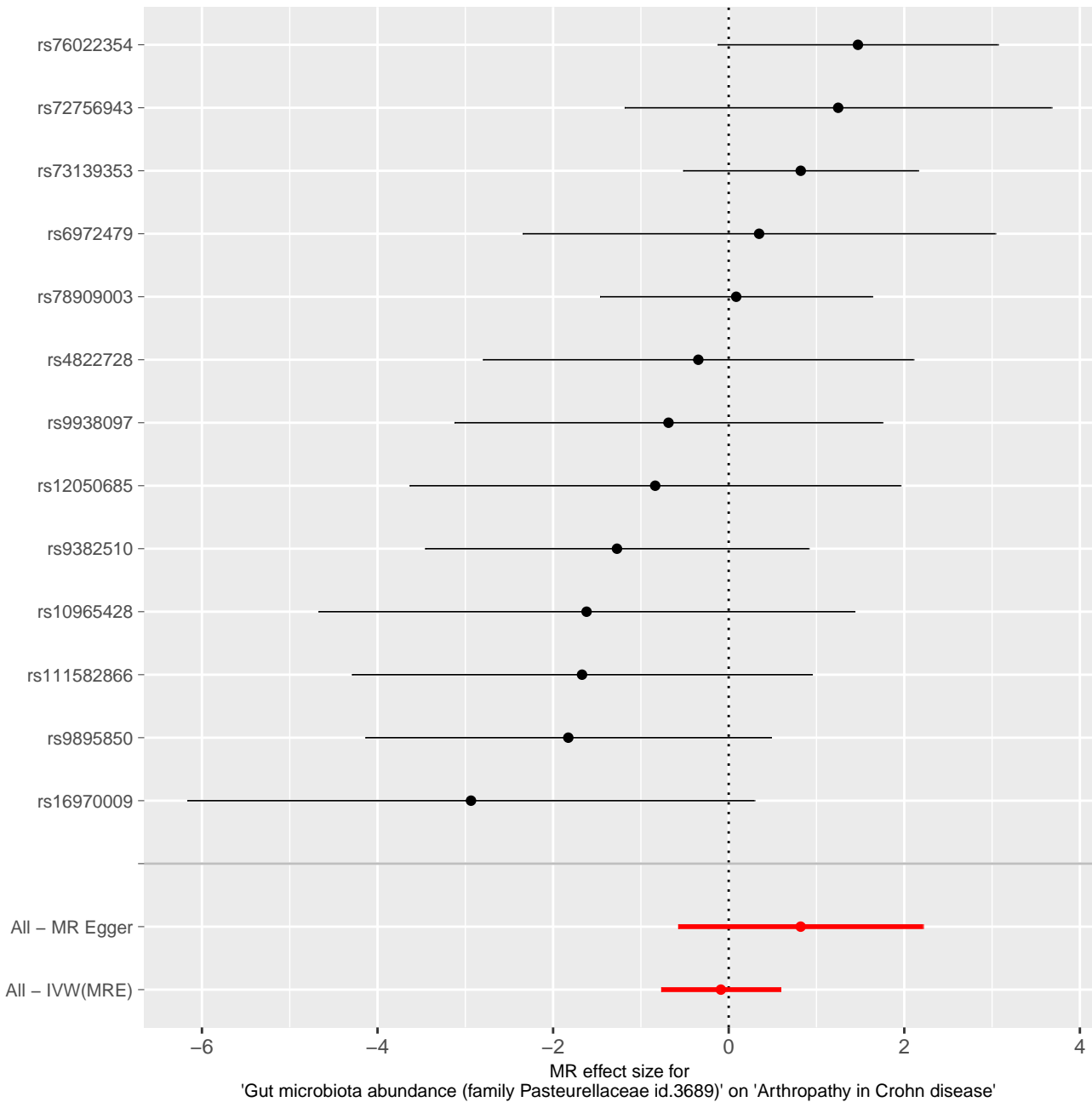




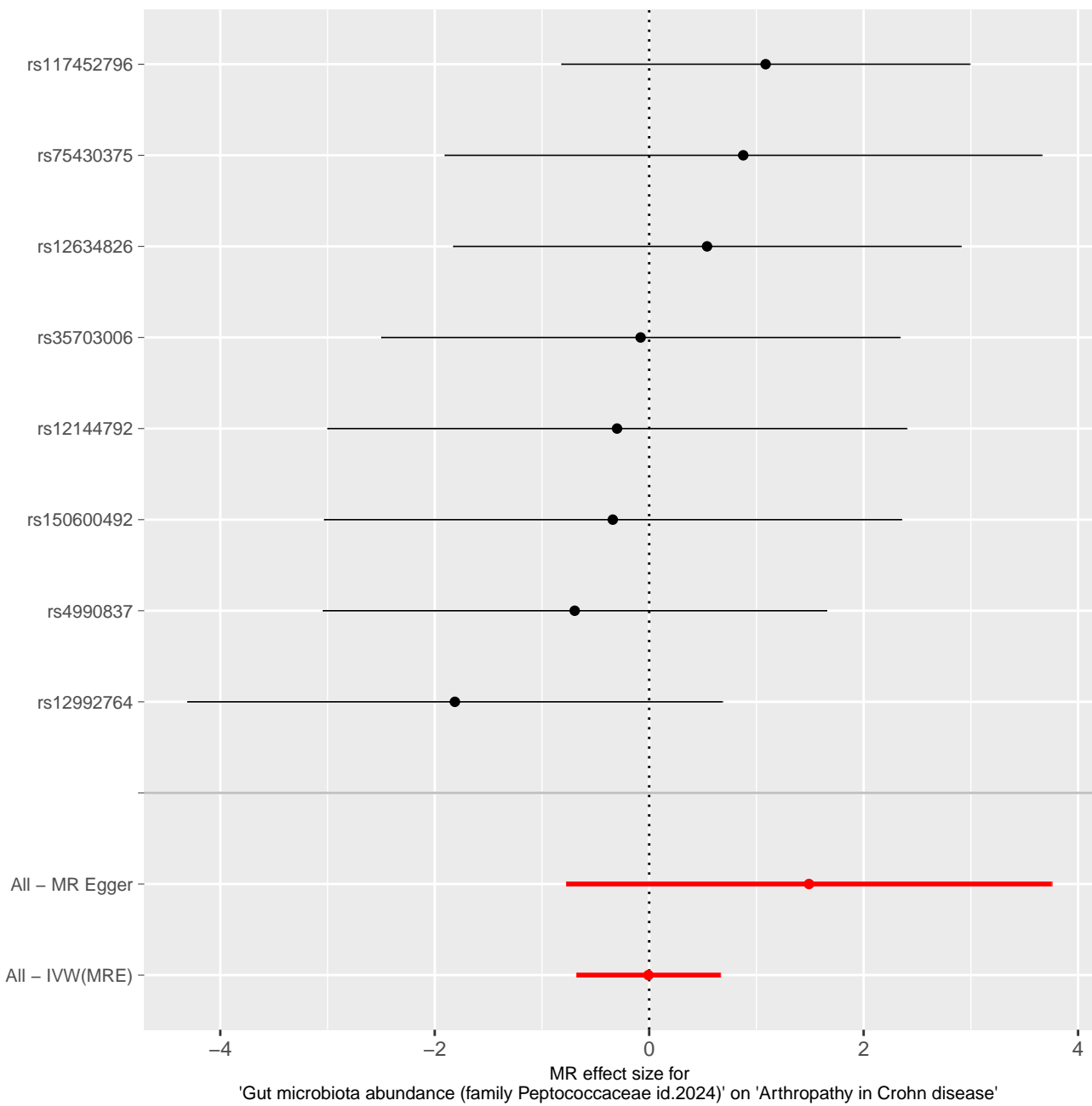


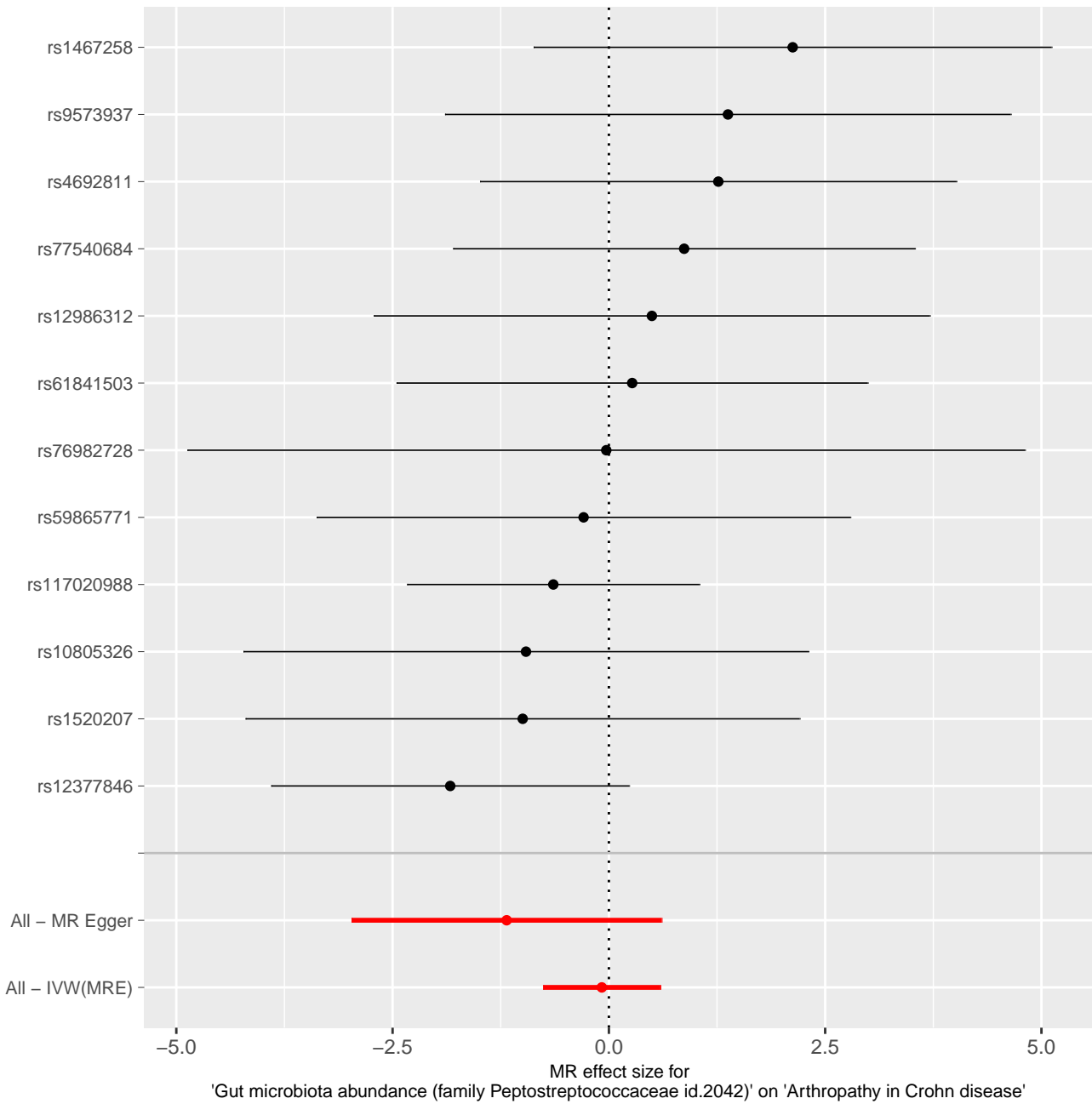


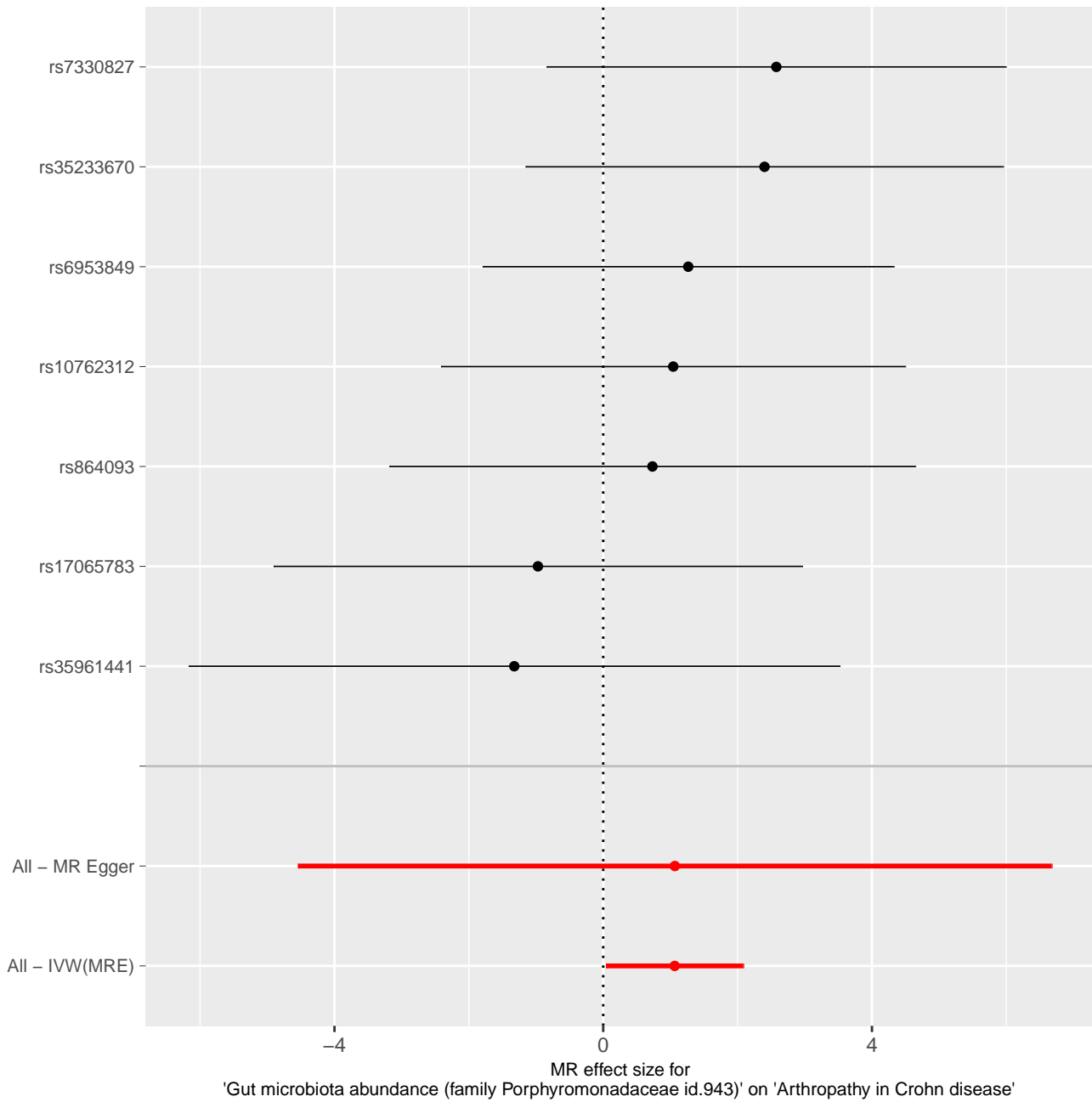
Batch 471 : Gut microbiota abundance (family Pasteurellaceae id.3689) on Arthropathy in Crohn disease



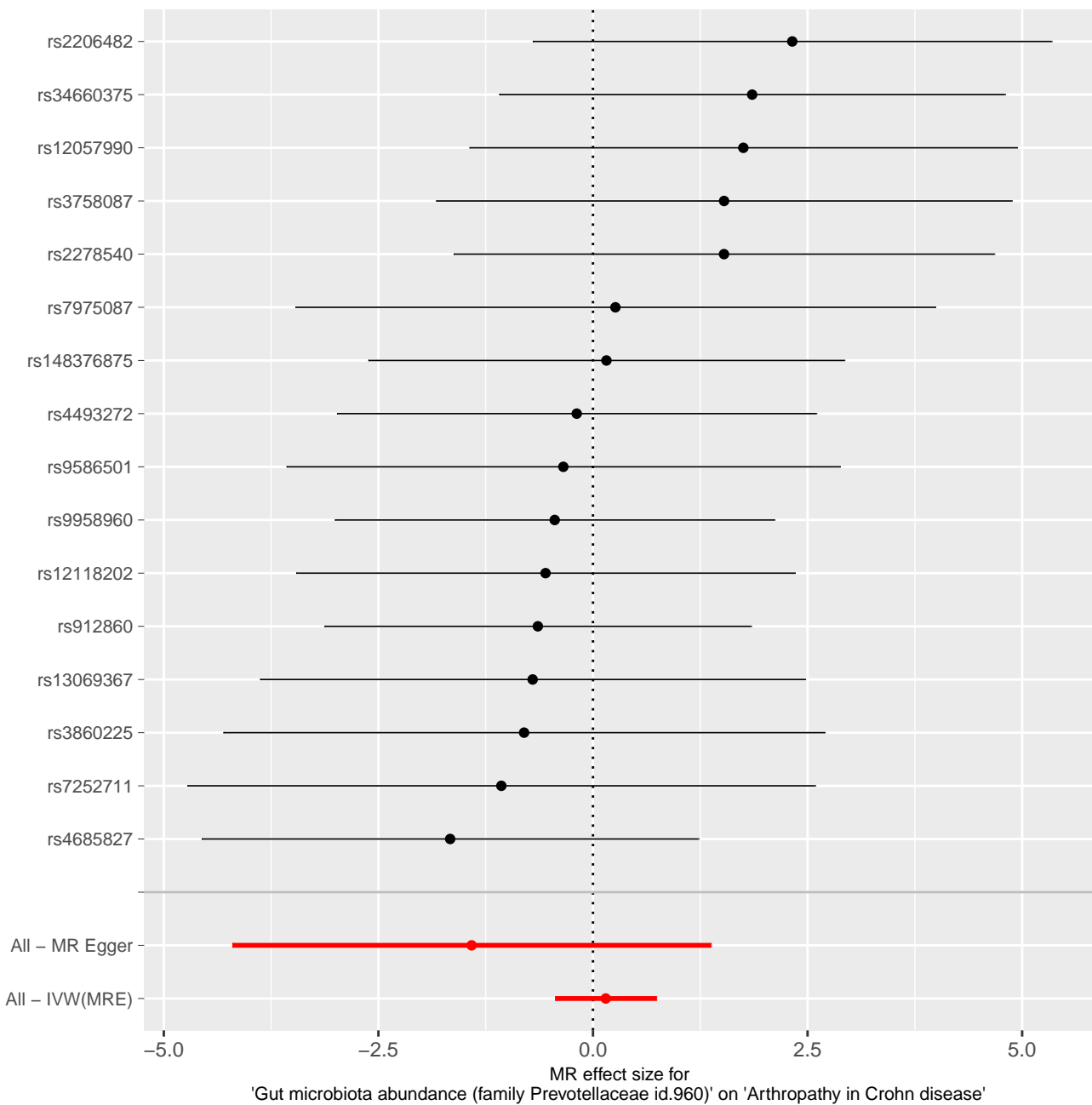
Batch 472 : Gut microbiota abundance (family Peptococcaceae id.2024) on Arthropathy in Crohn disease



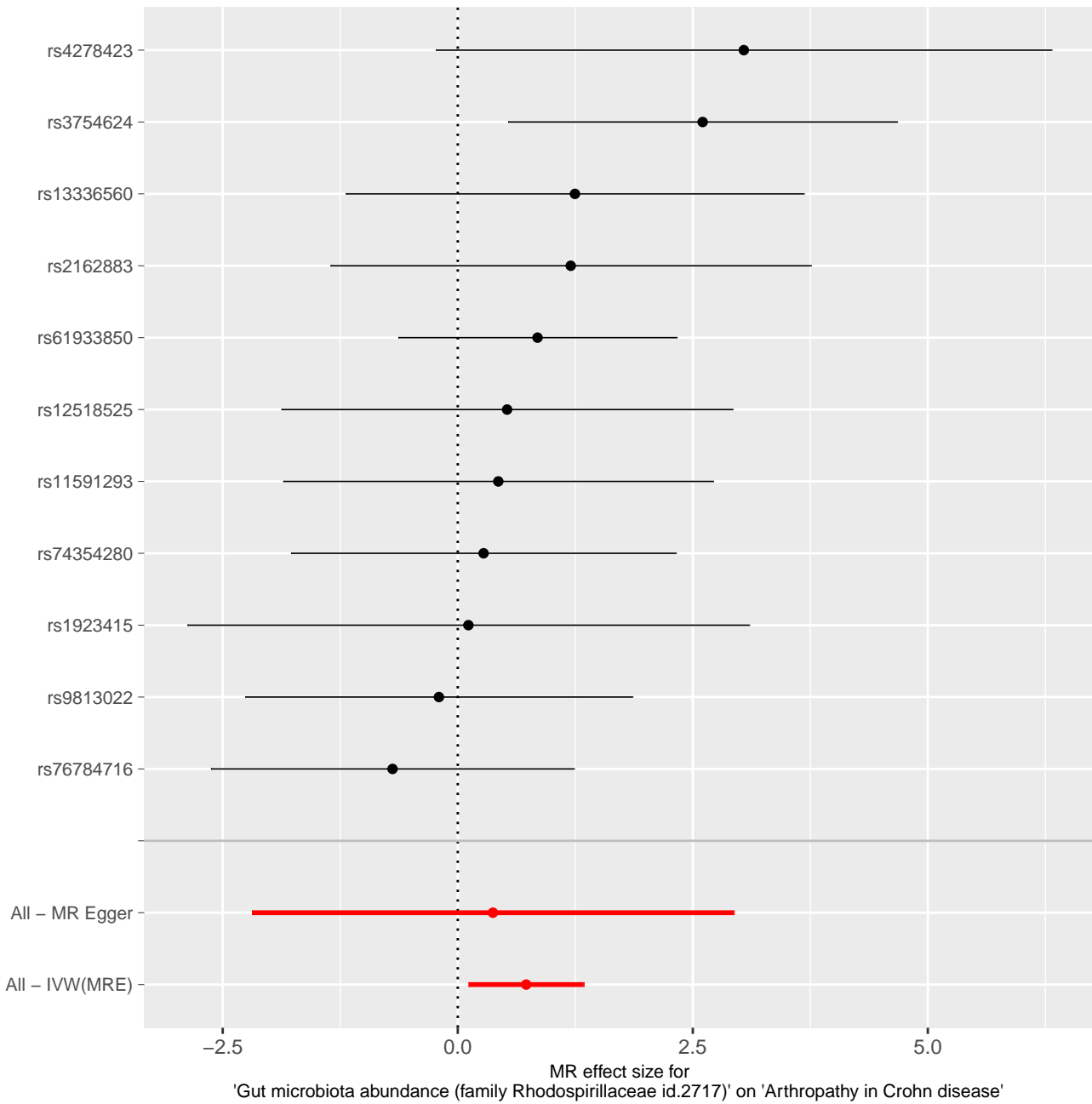




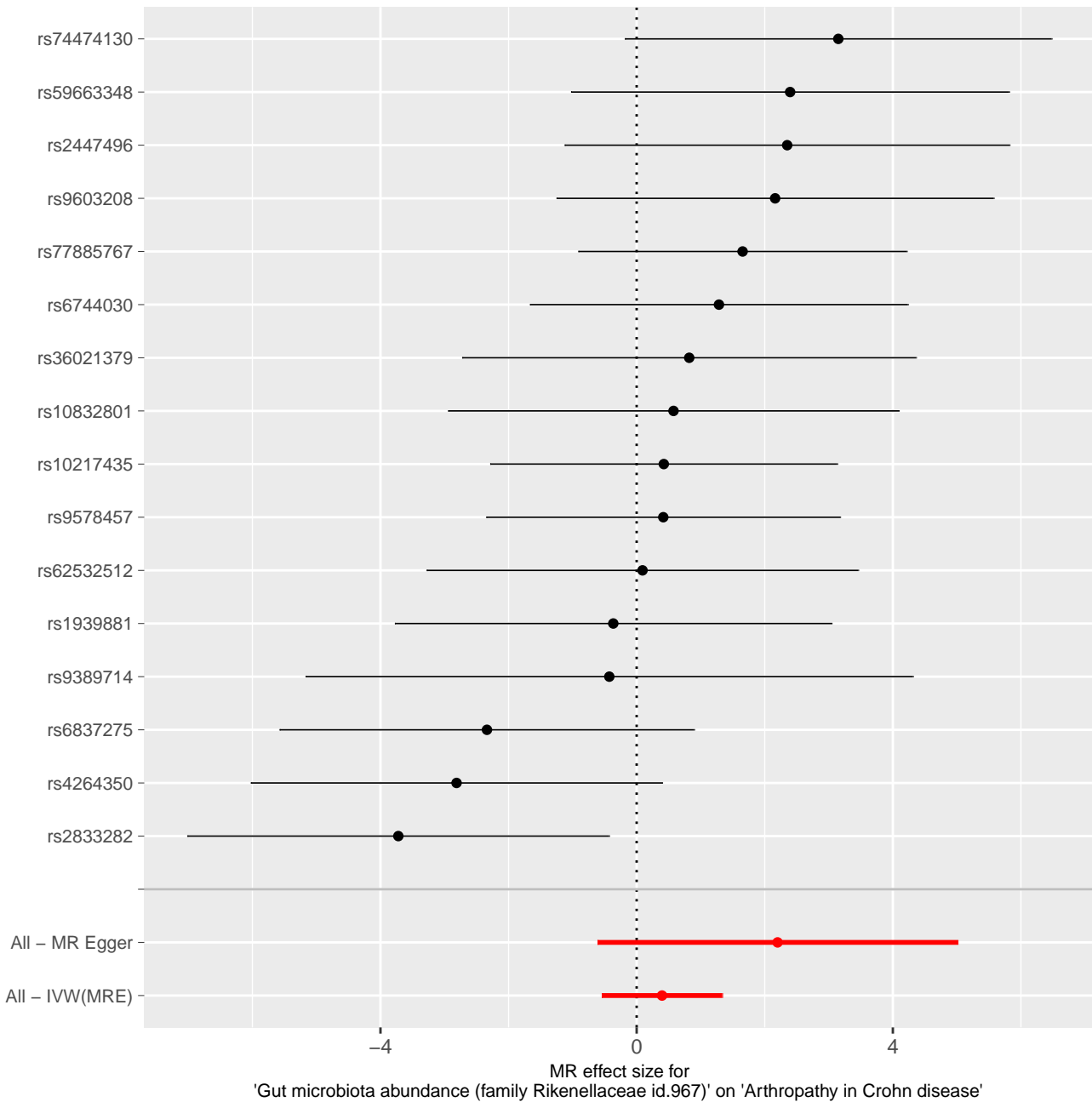
Batch 475 : Gut microbiota abundance (family Prevotellaceae id.960) on Arthropathy in Crohn disease



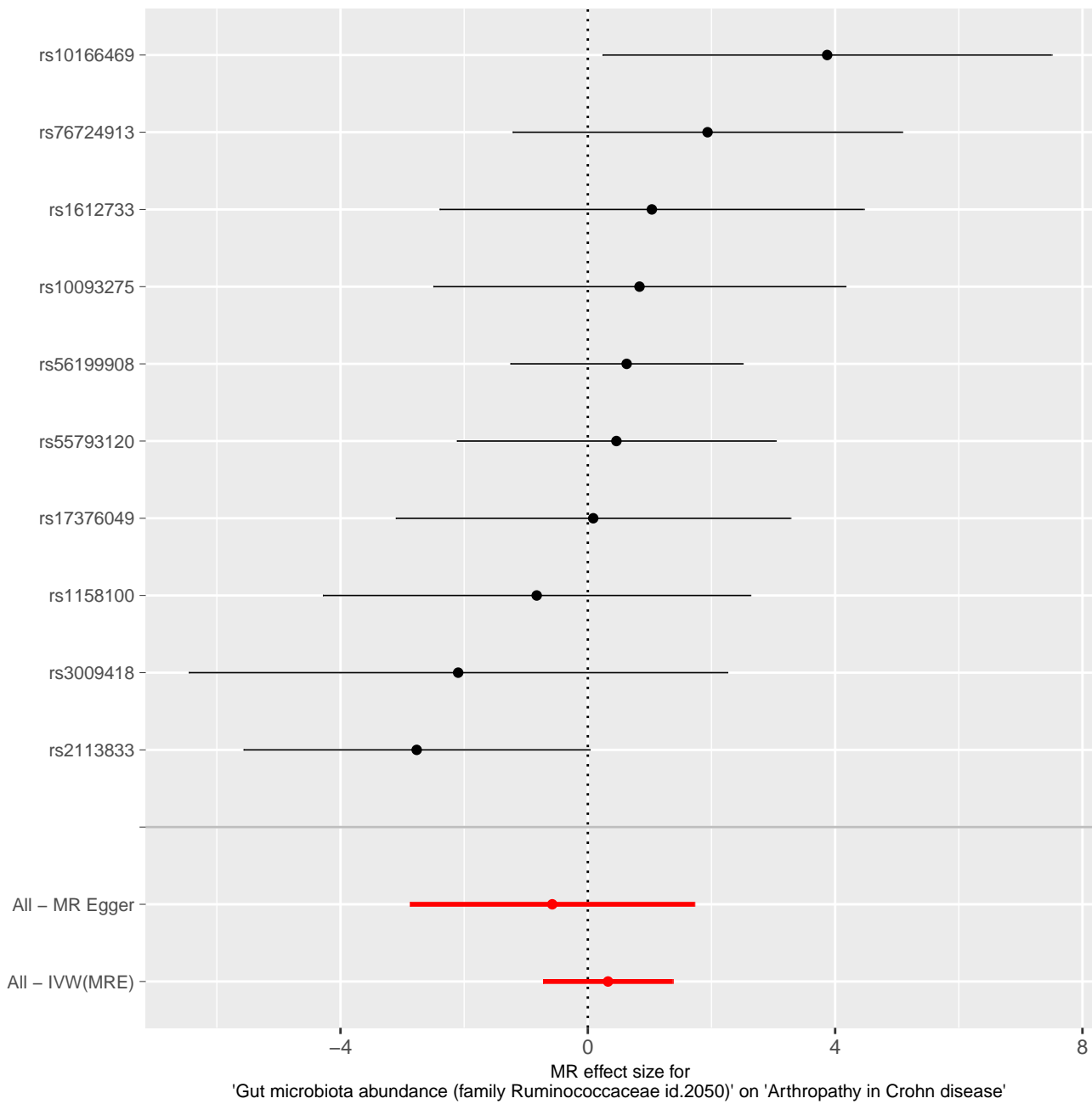
Batch 476 : Gut microbiota abundance (family Rhodospirillaceae id.2717) on Arthropathy in Crohn disease

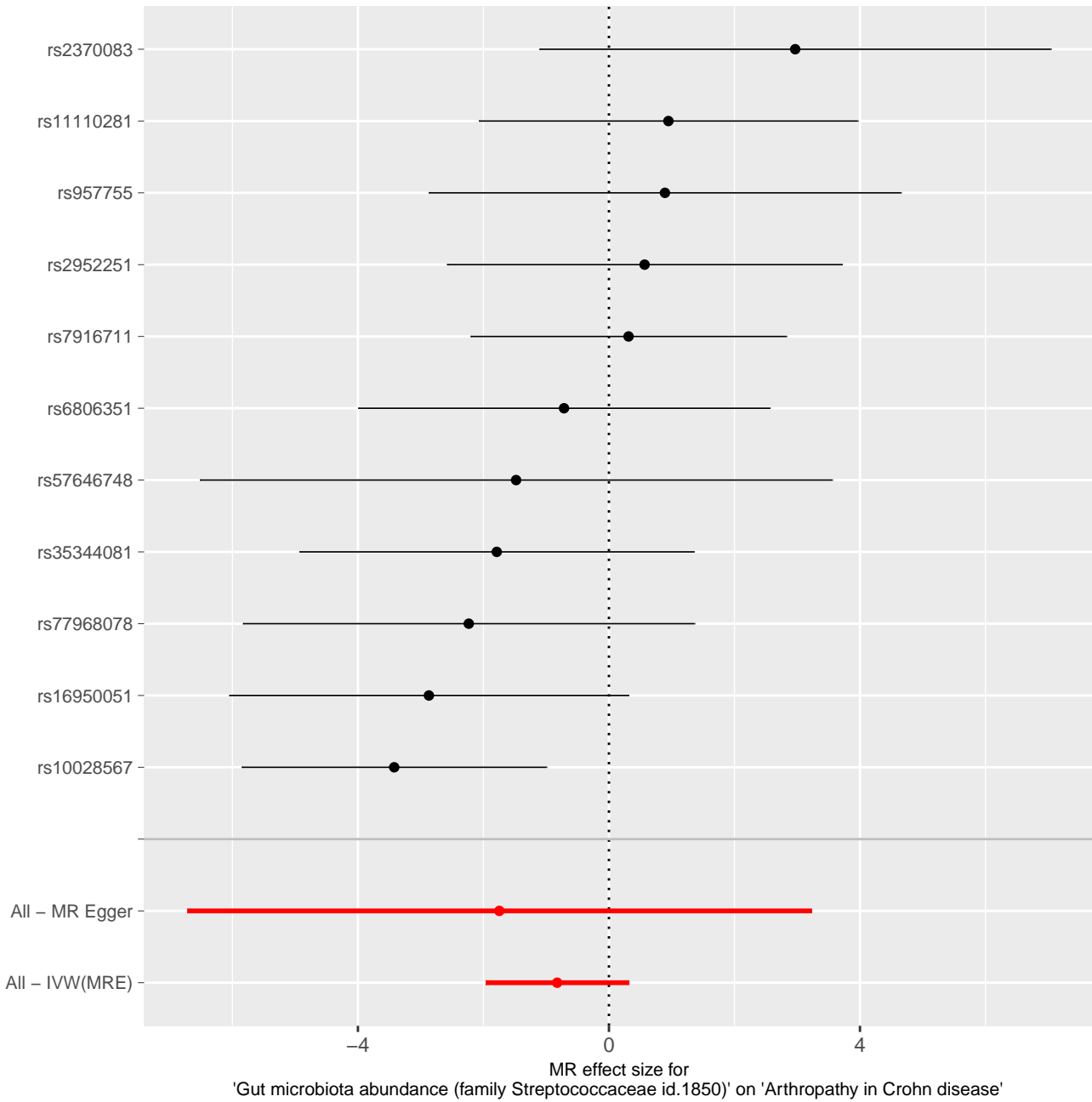


Batch 477 : Gut microbiota abundance (family Rikenellaceae id.967) on Arthropathy in Crohn disease

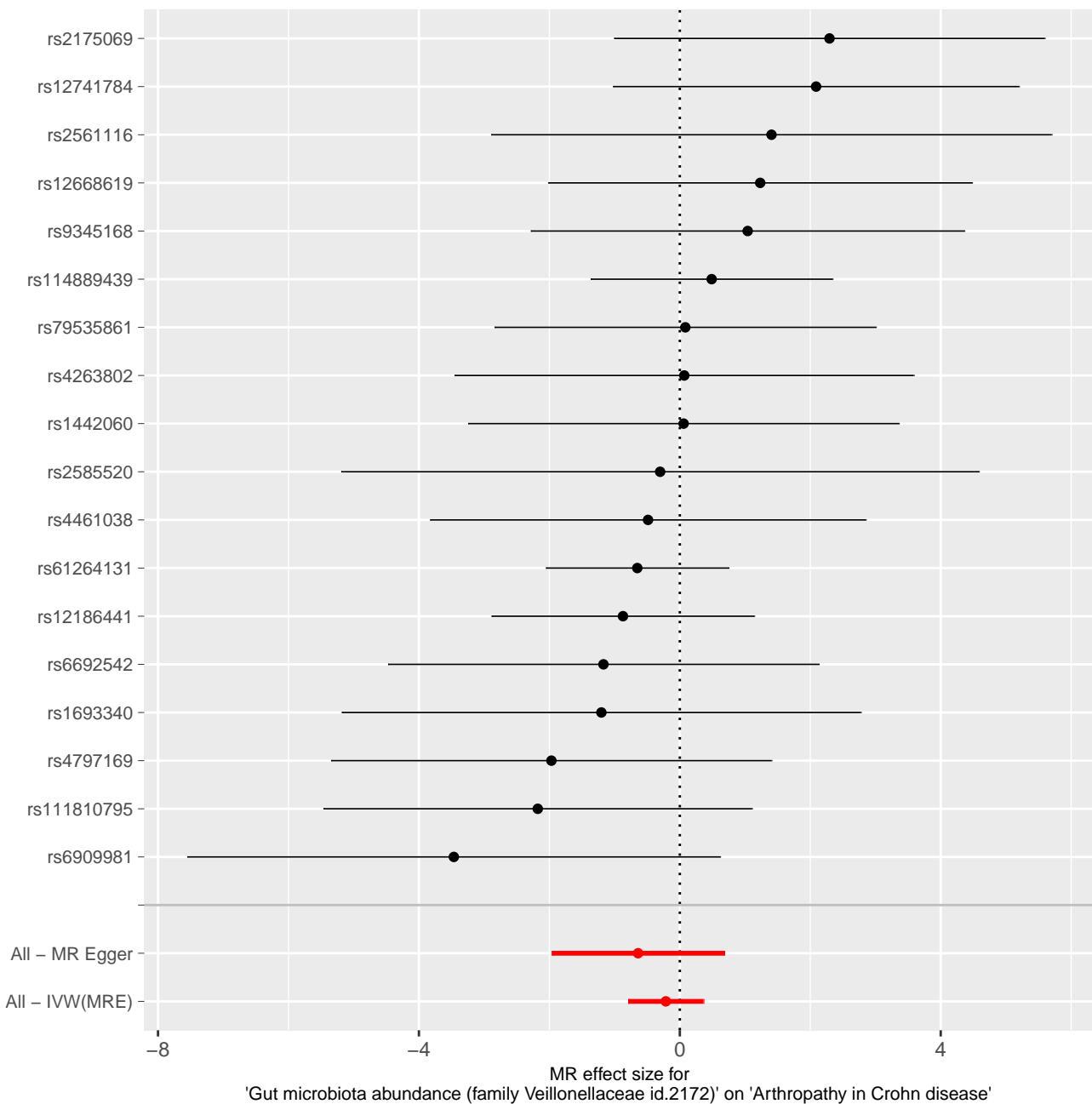


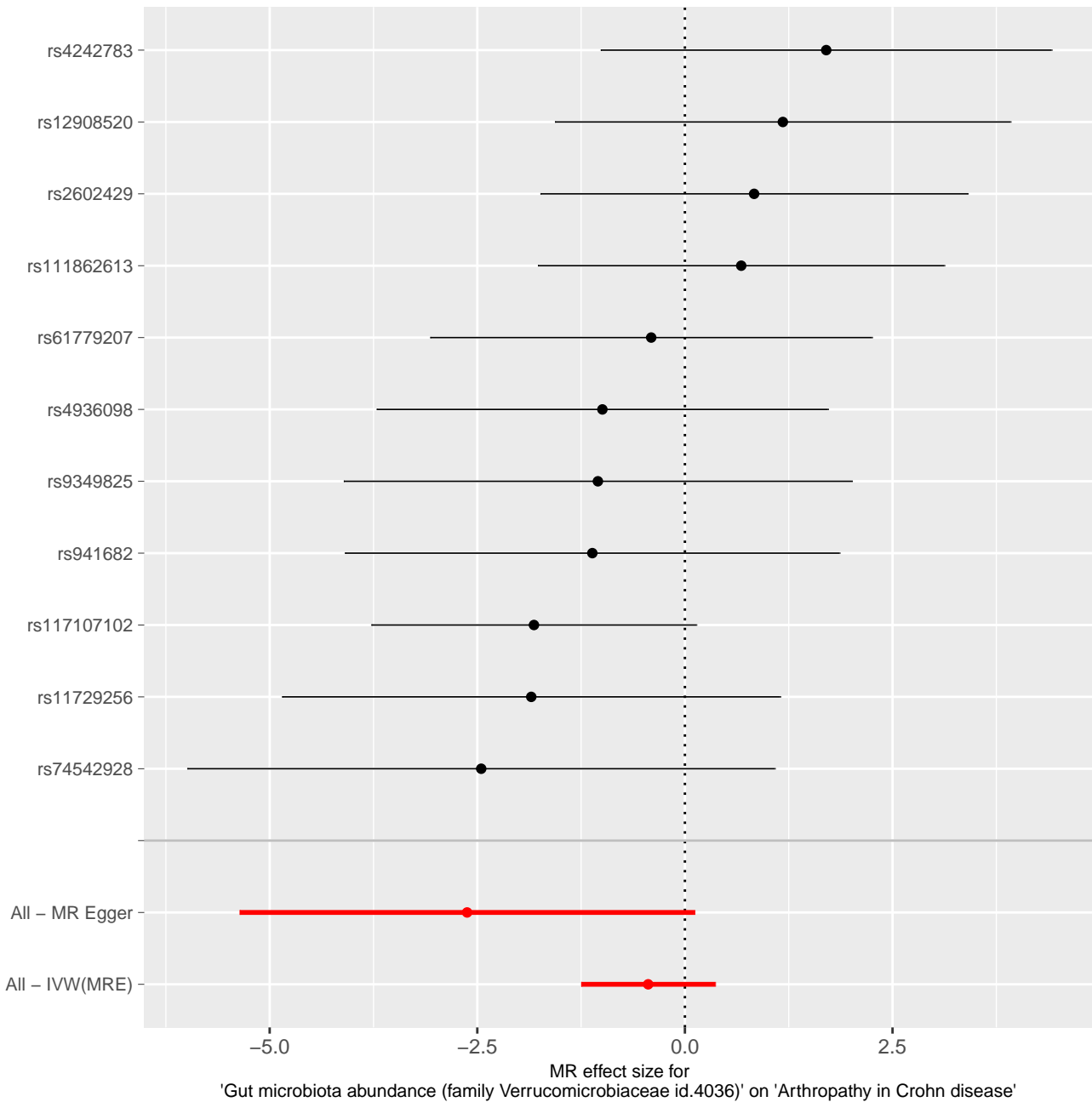
Batch 478 : Gut microbiota abundance (family Ruminococcaceae id.2050) on Arthropathy in Crohn disease



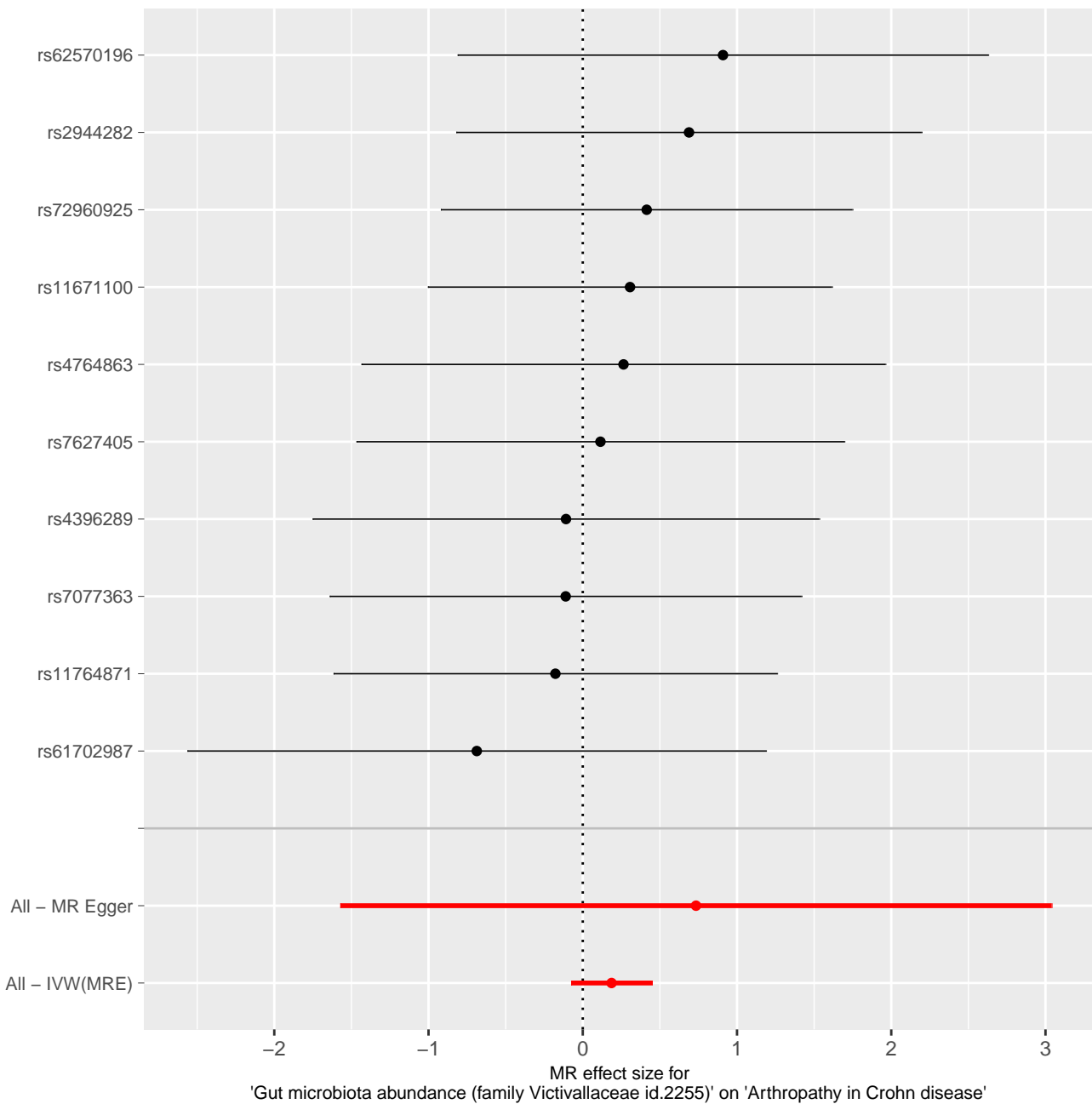


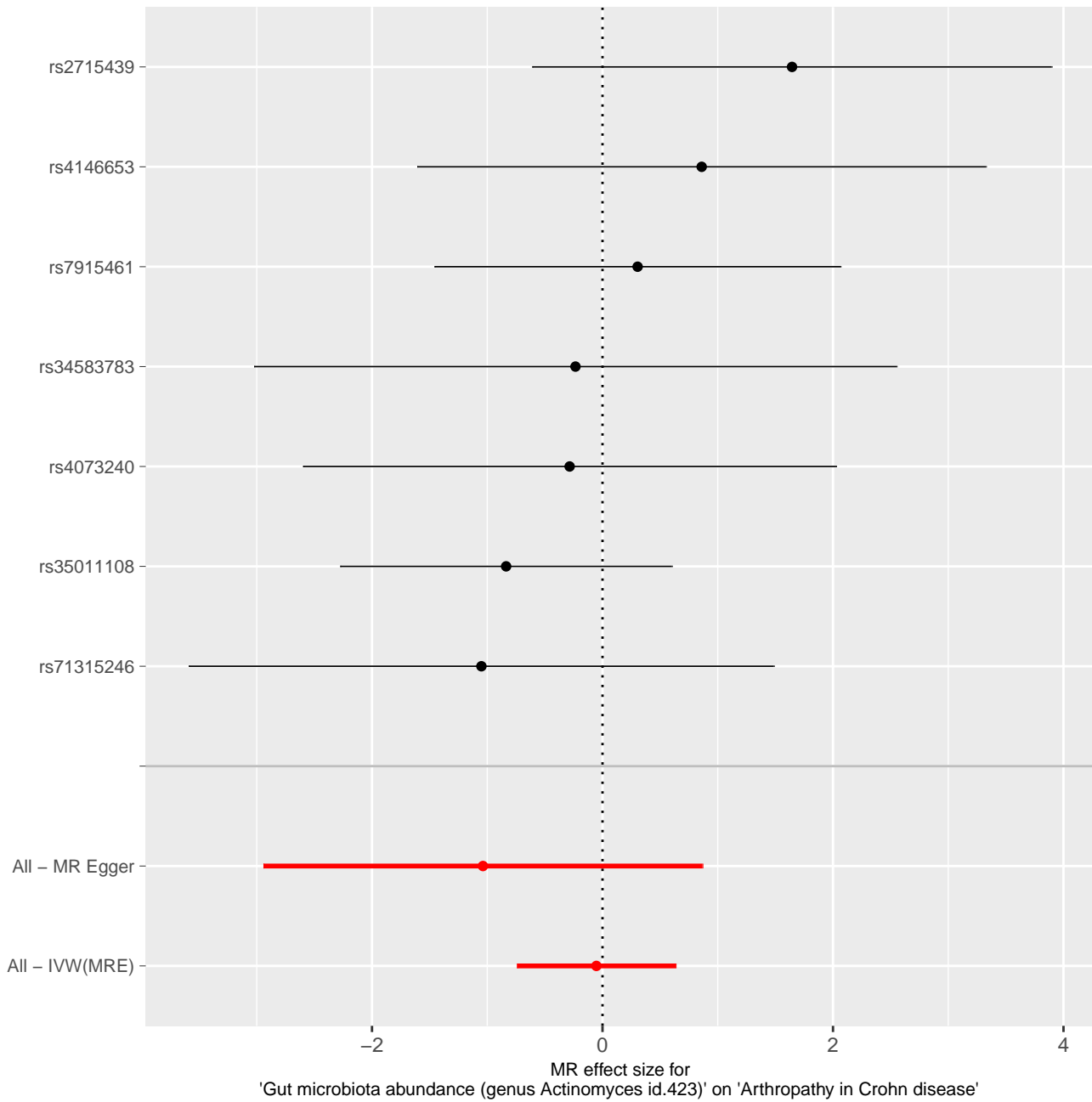
Batch 480 : Gut microbiota abundance (family Veillonellaceae id.2172) on Arthropathy in Crohn disease

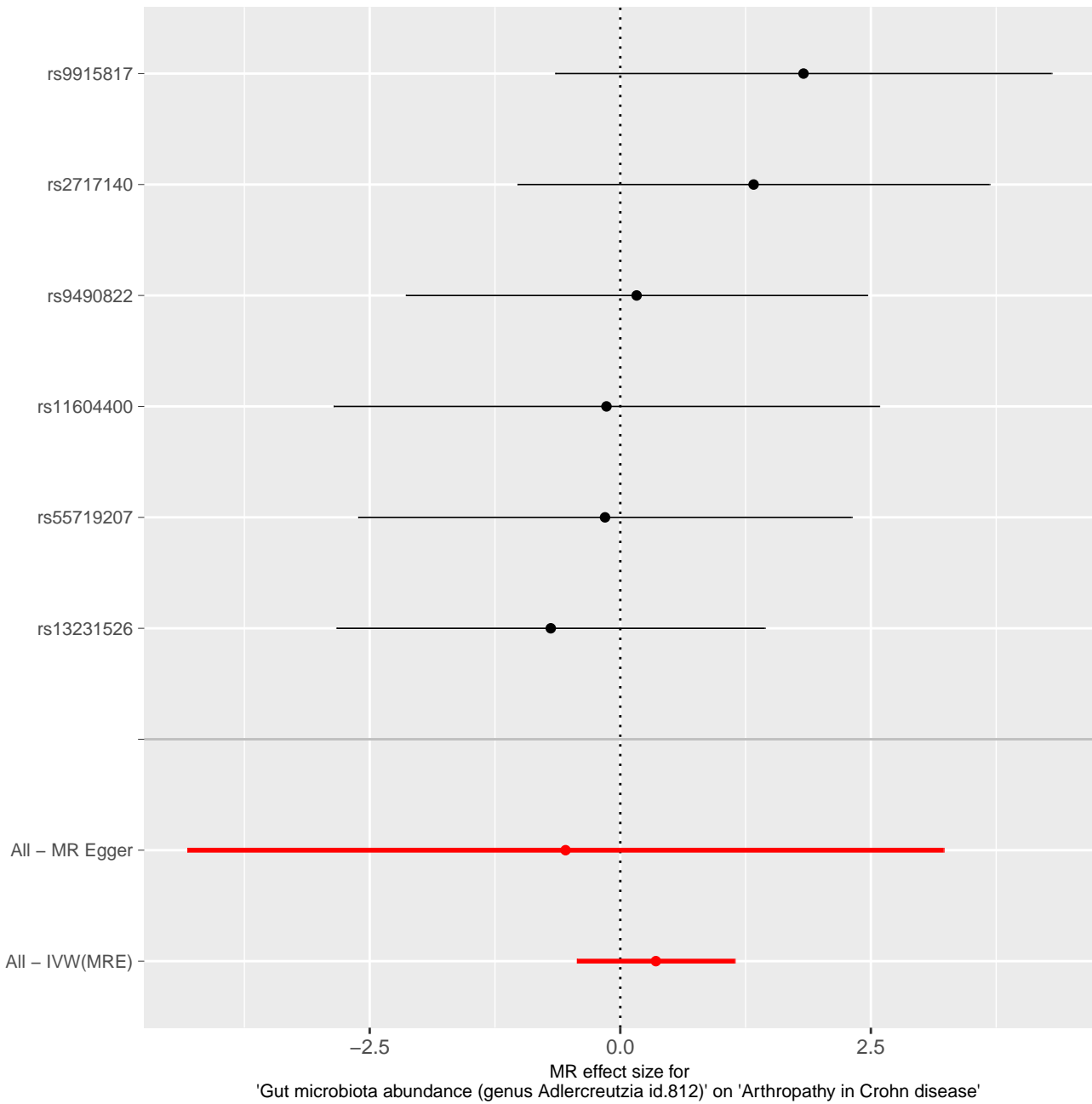


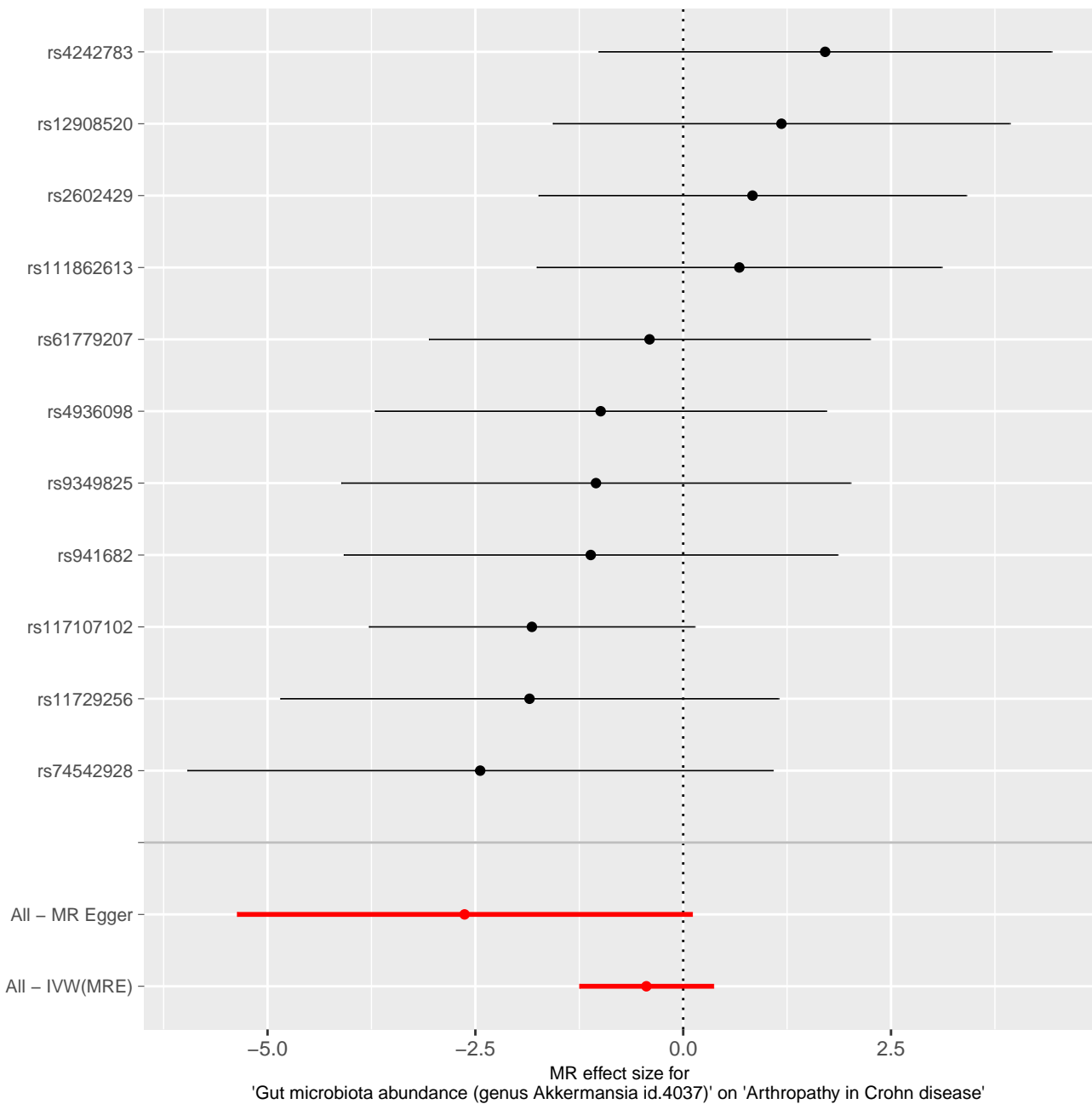


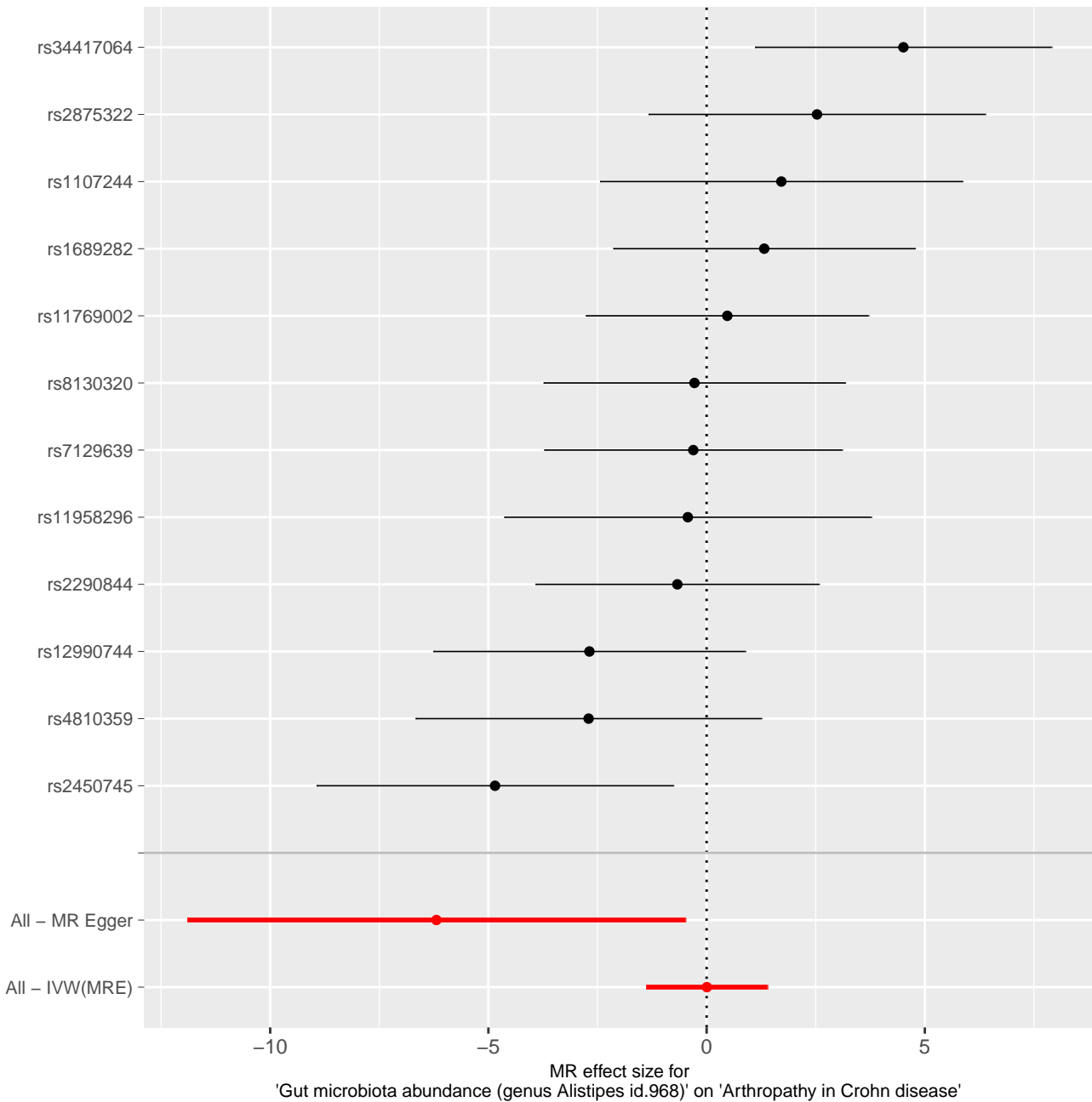
Batch 482 : Gut microbiota abundance (family Victivallaceae id.2255) on Arthropathy in Crohn disease

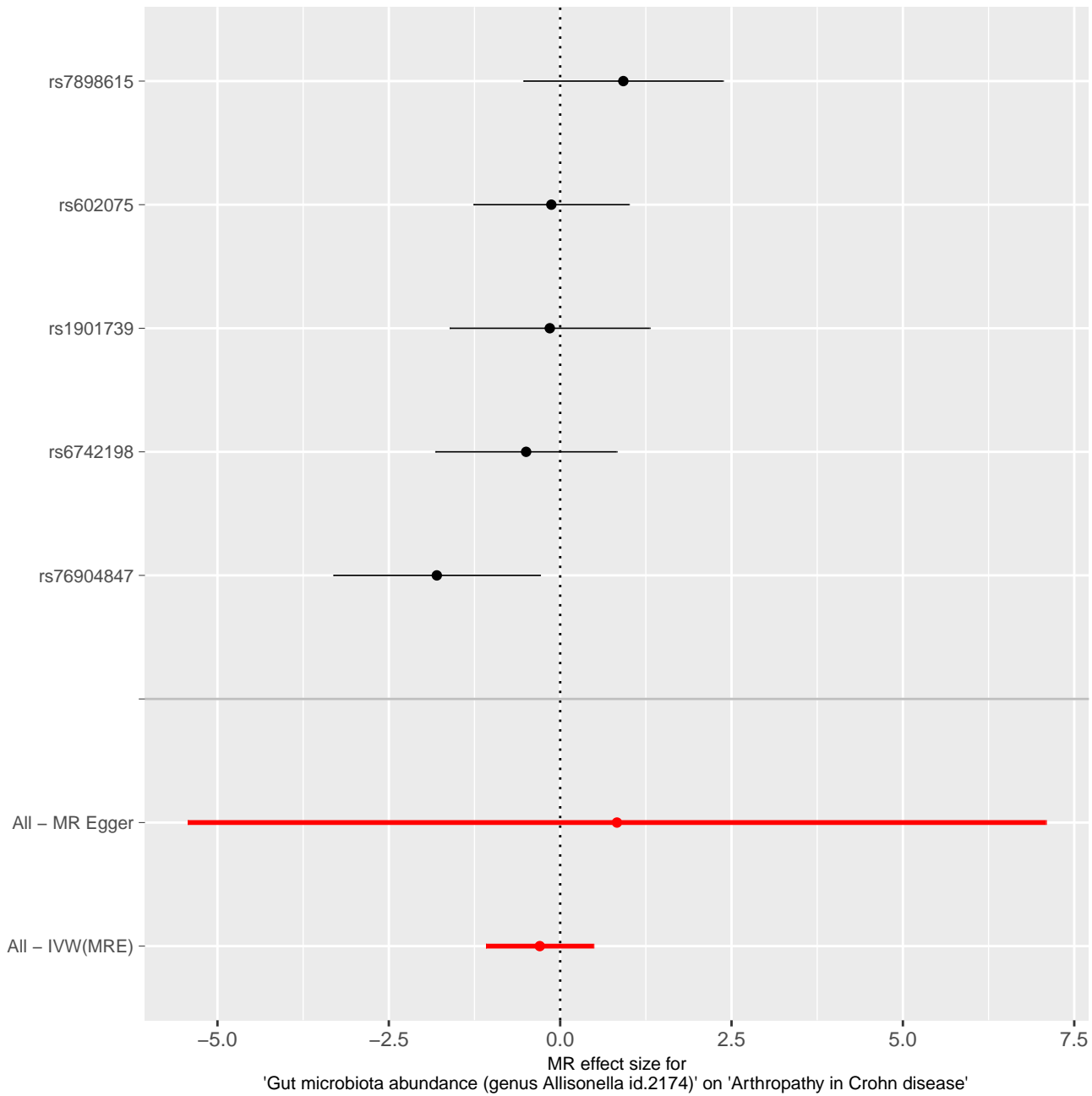


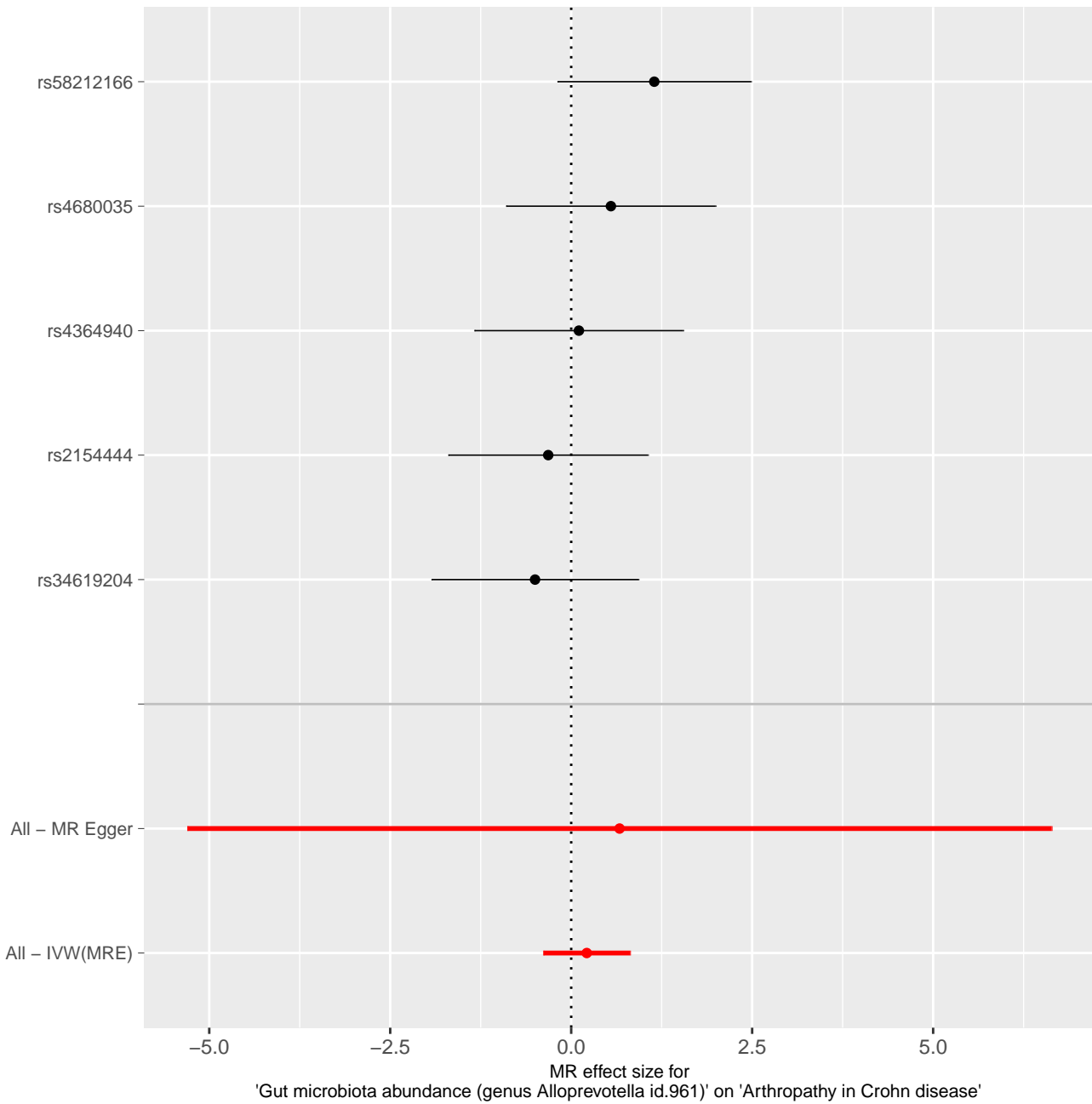




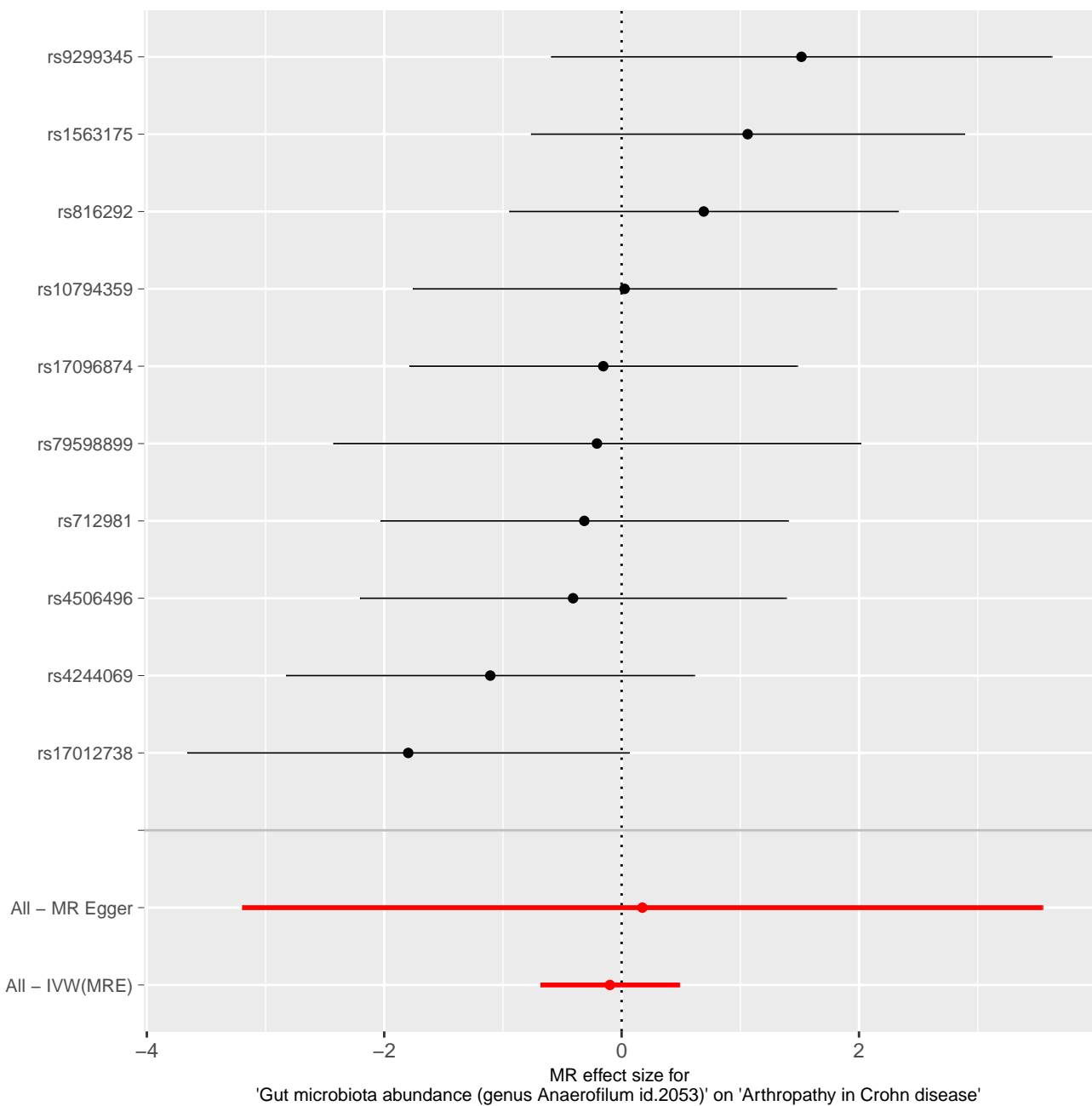


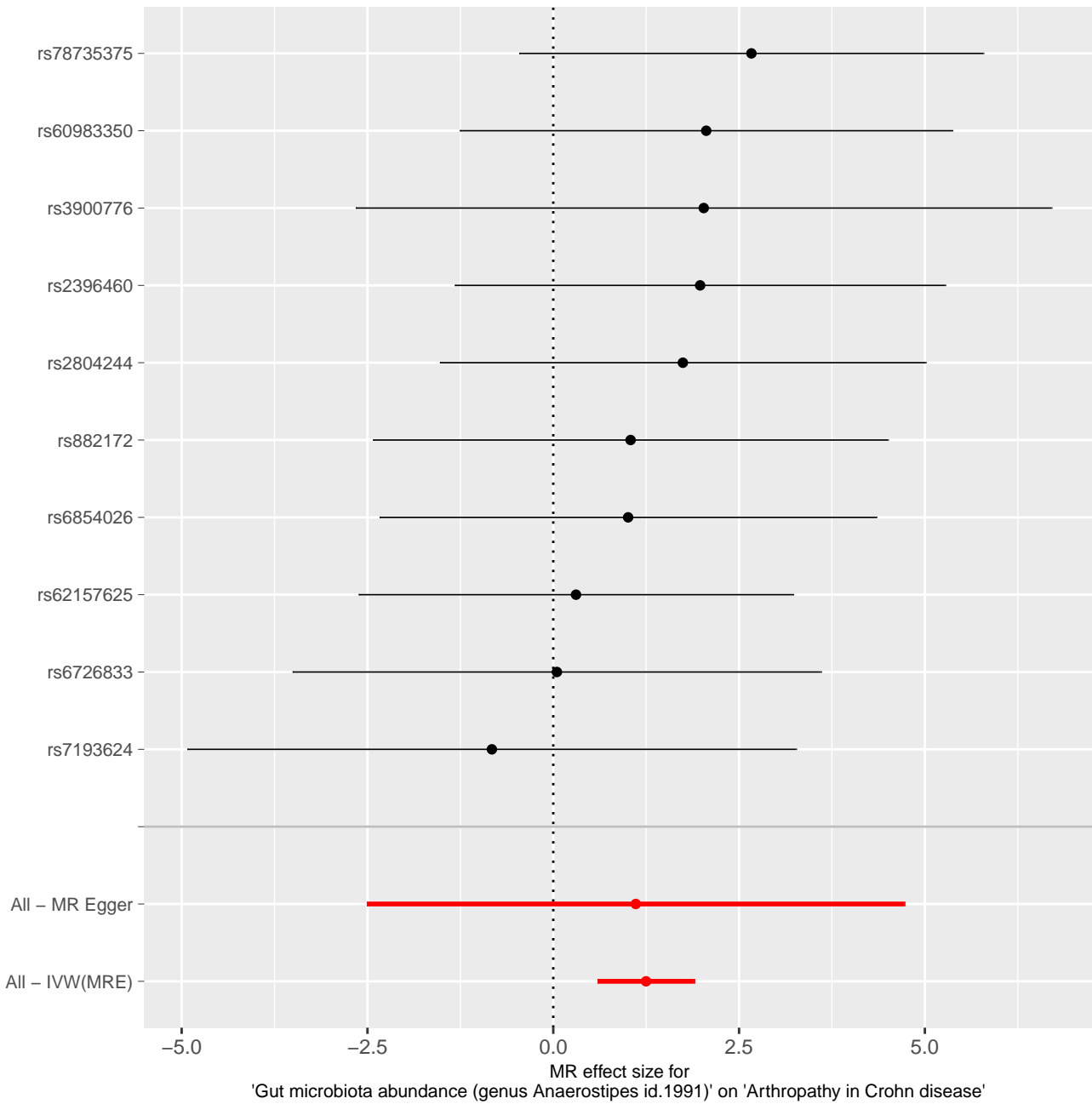


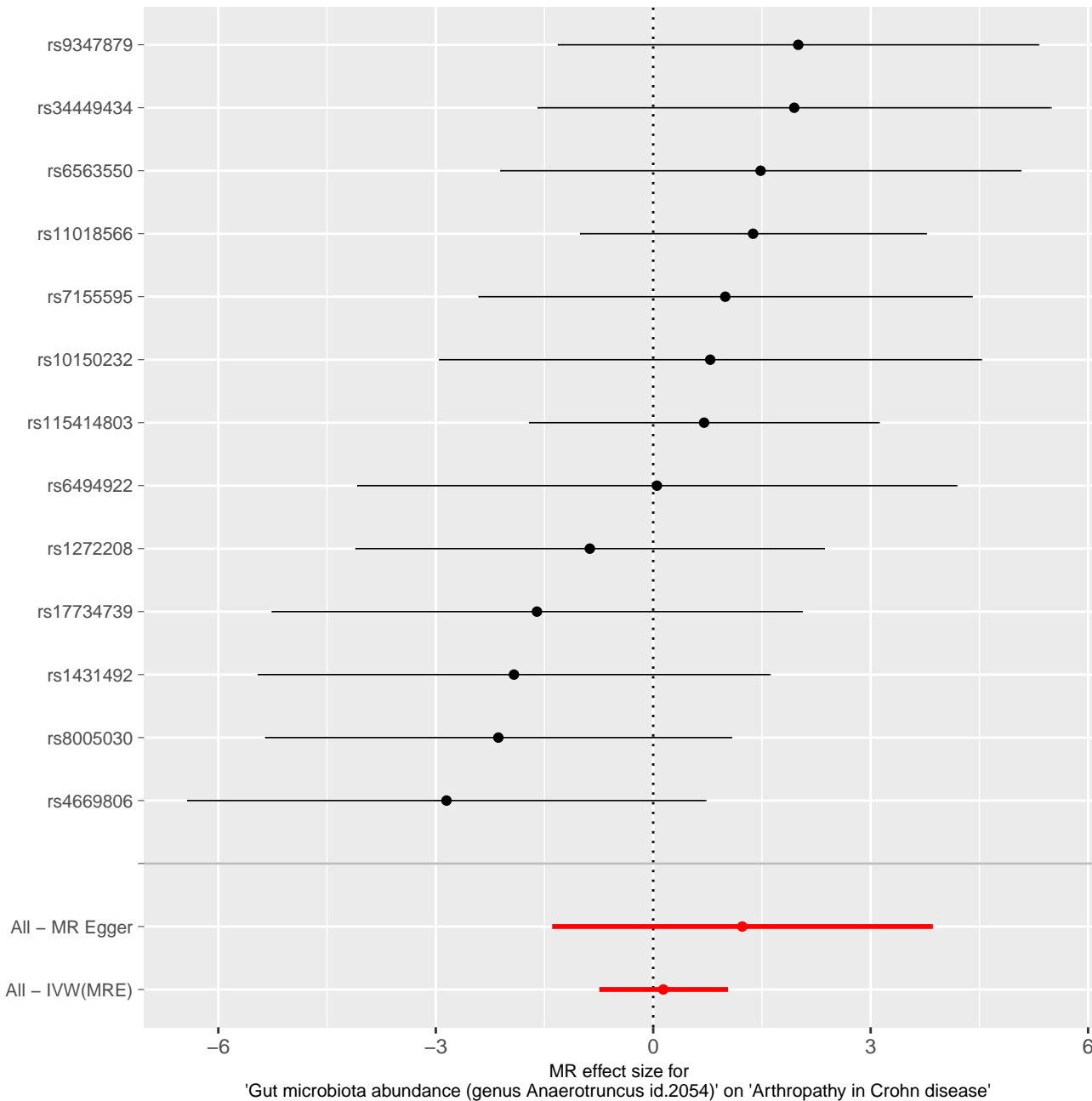


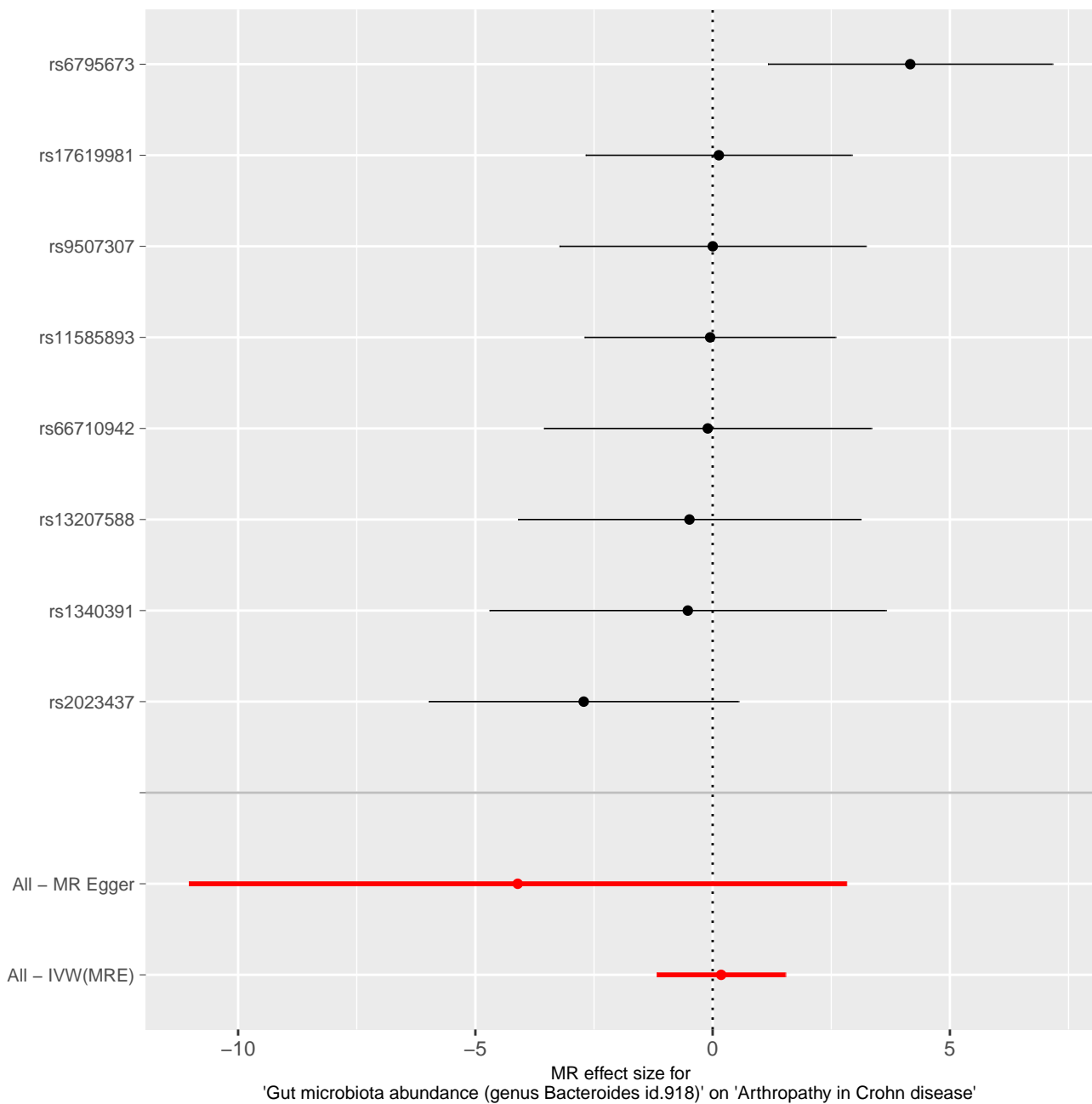


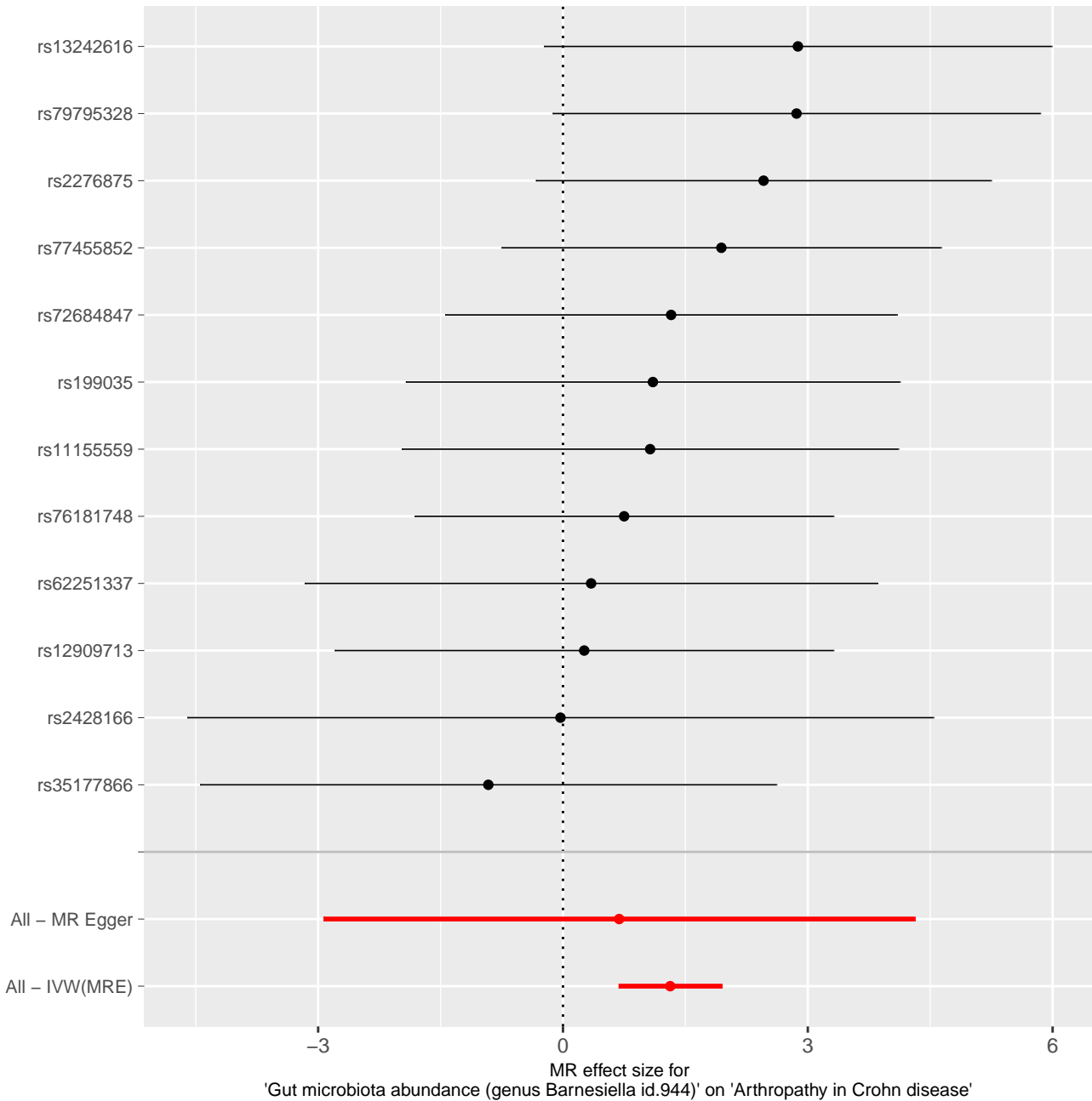
Batch 489 : Gut microbiota abundance (genus Anaerofilum id.2053) on Arthropathy in Crohn disease

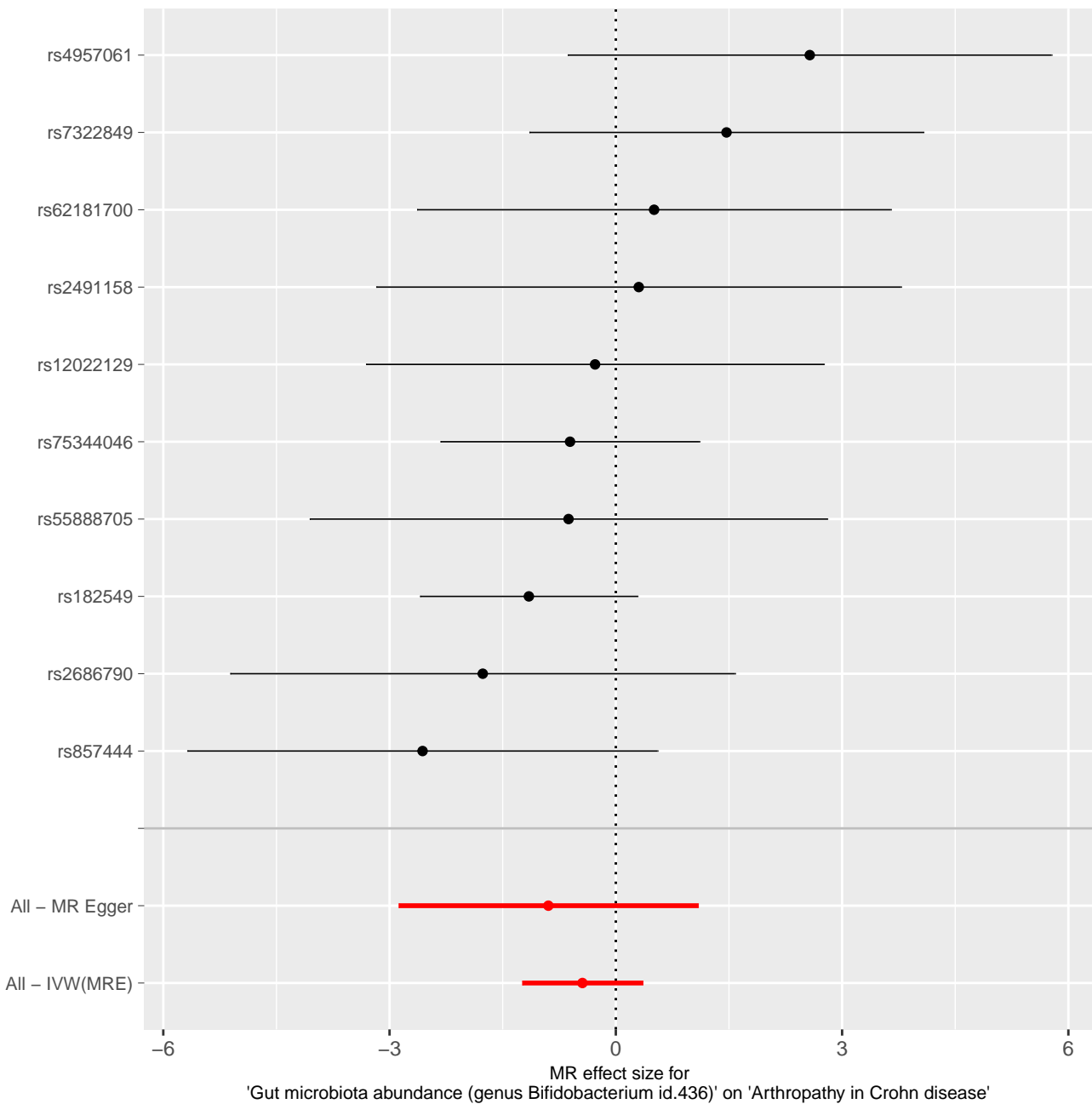




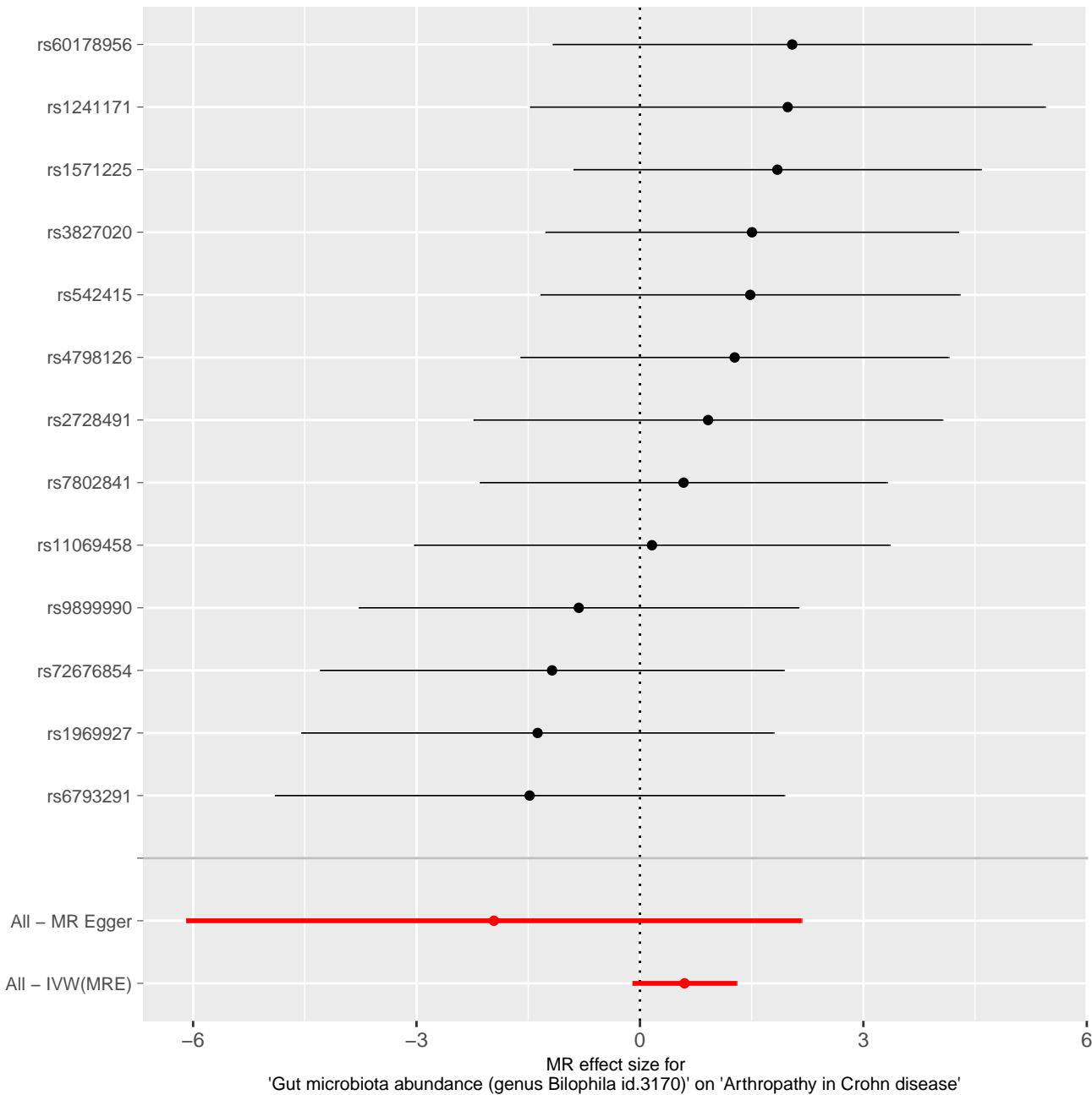


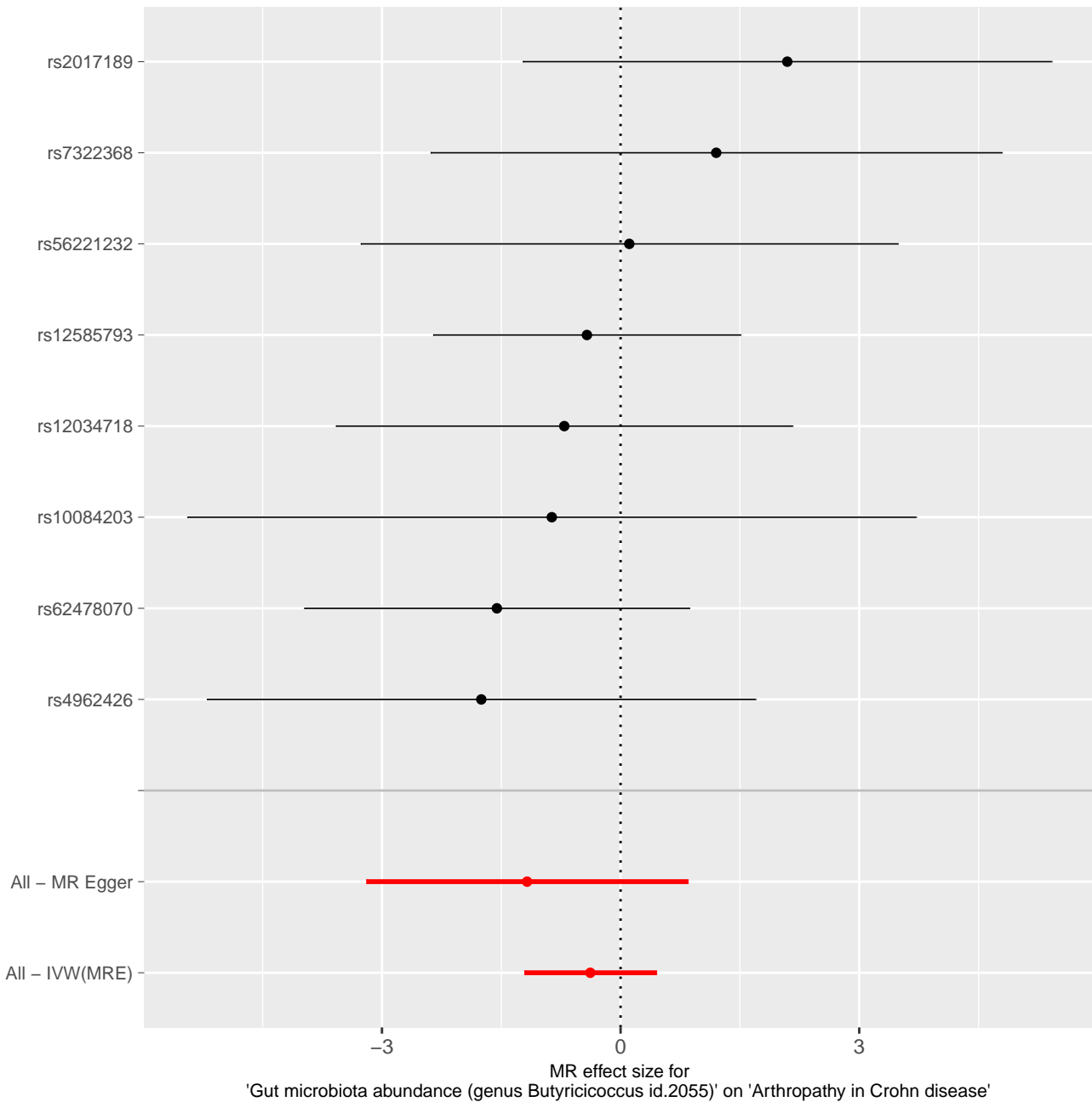






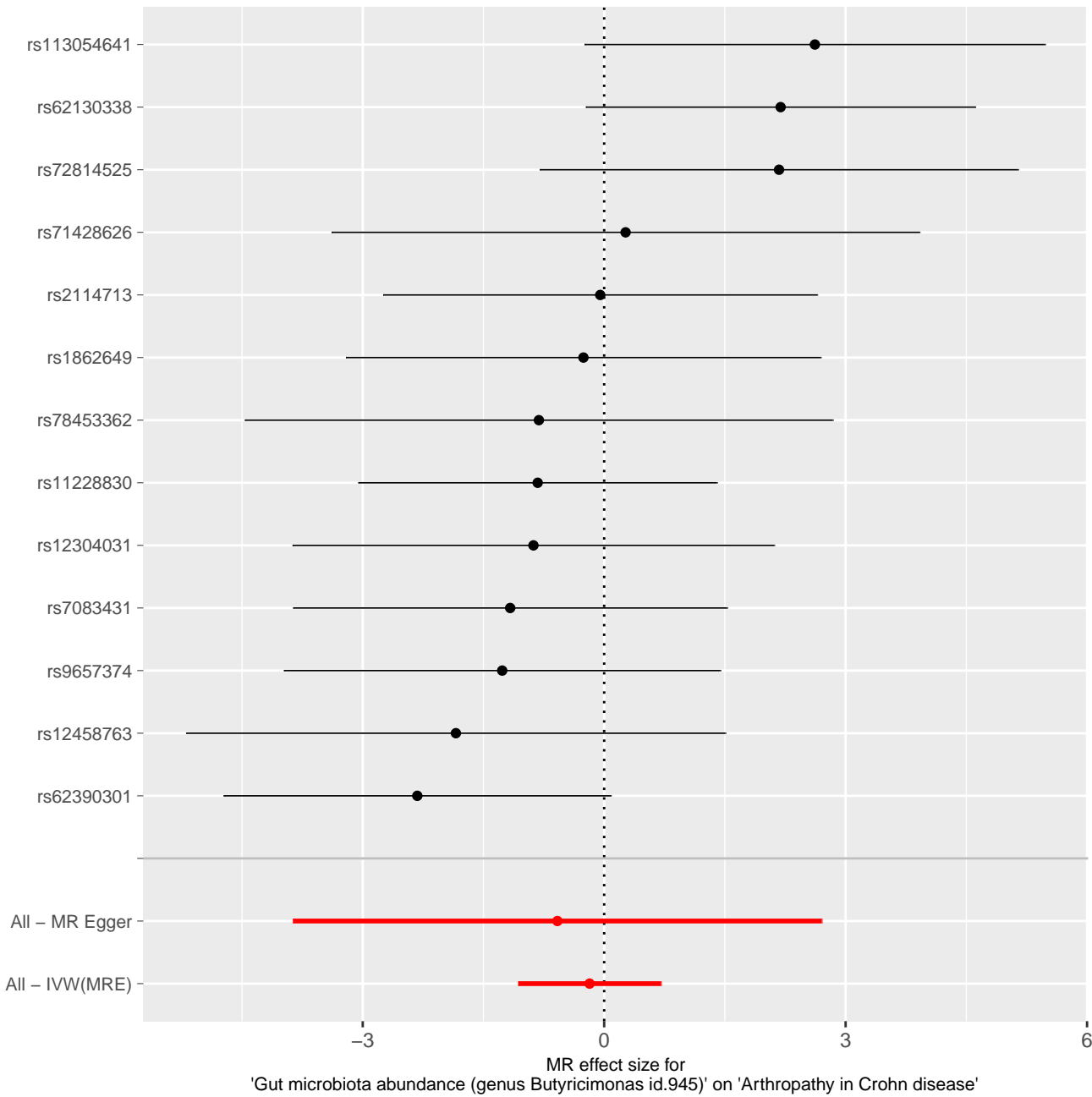
Batch 495 : Gut microbiota abundance (genus Bilophila id.3170) on Arthropathy in Crohn disease



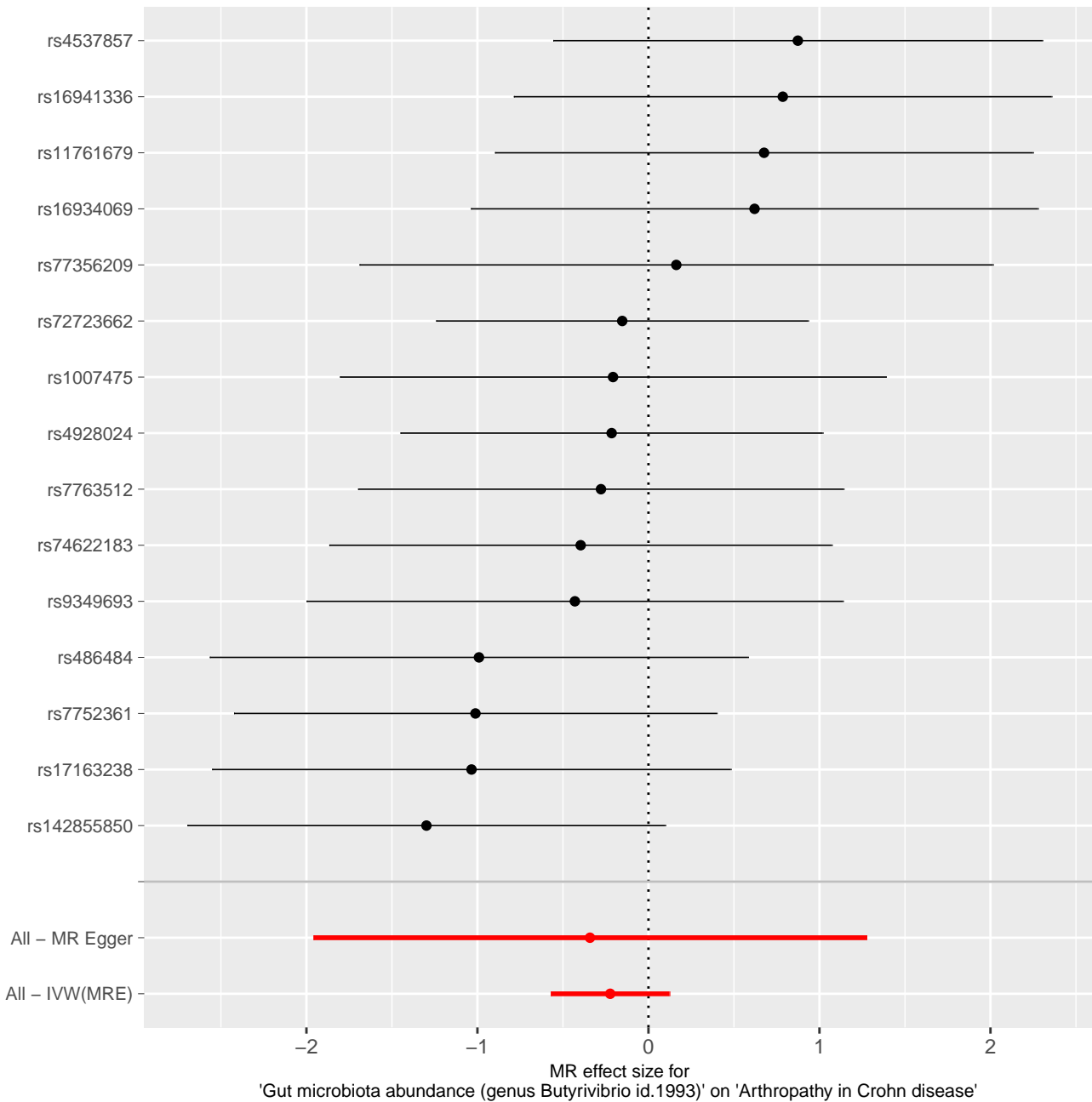


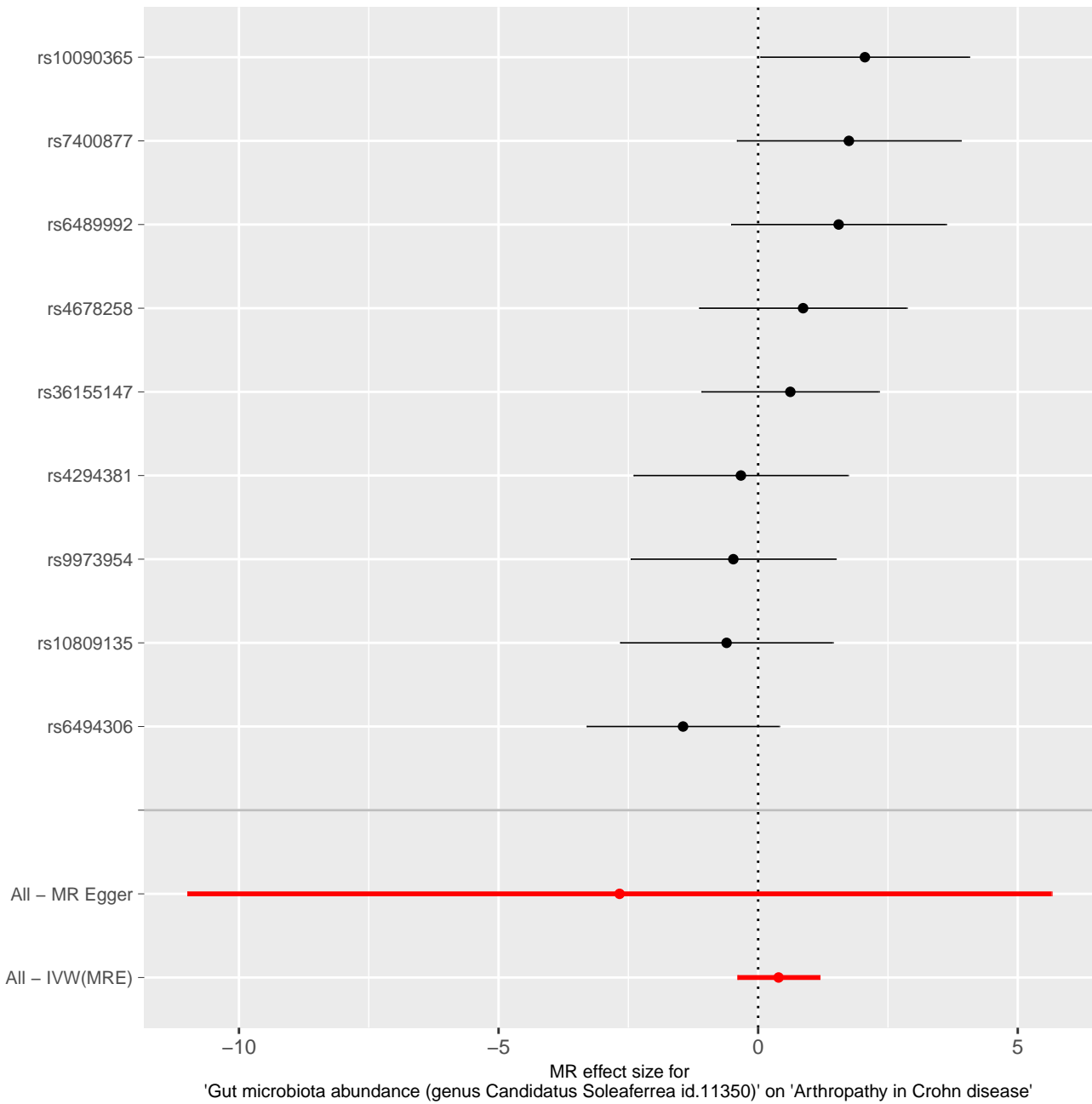
MR effect size for
'Gut microbiota abundance (genus Butyricoccus id.2055)' on 'Arthropathy in Crohn disease'

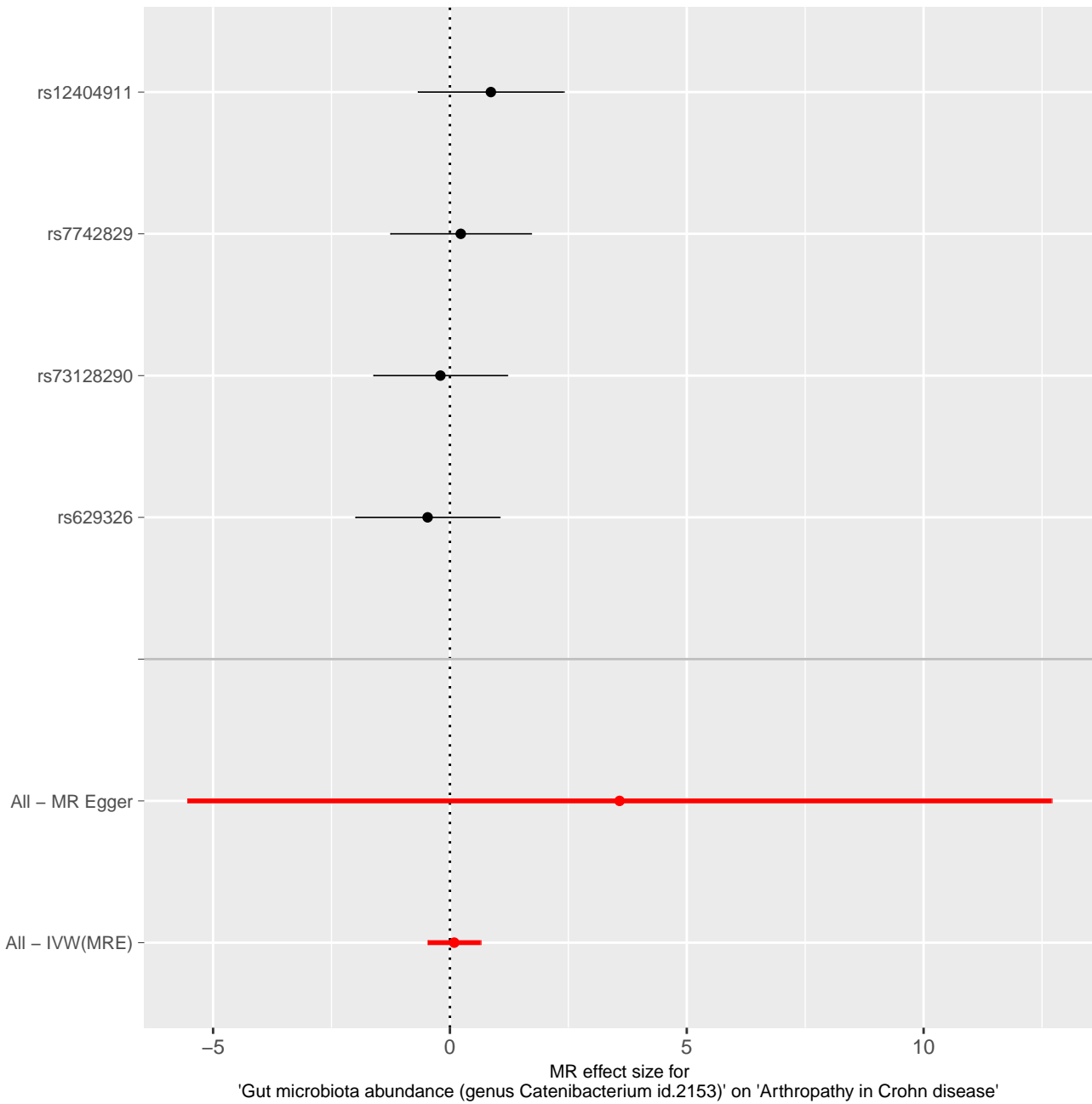
Batch 498 : Gut microbiota abundance (genus Butyricimonas id.945) on Arthropathy in Crohn disease

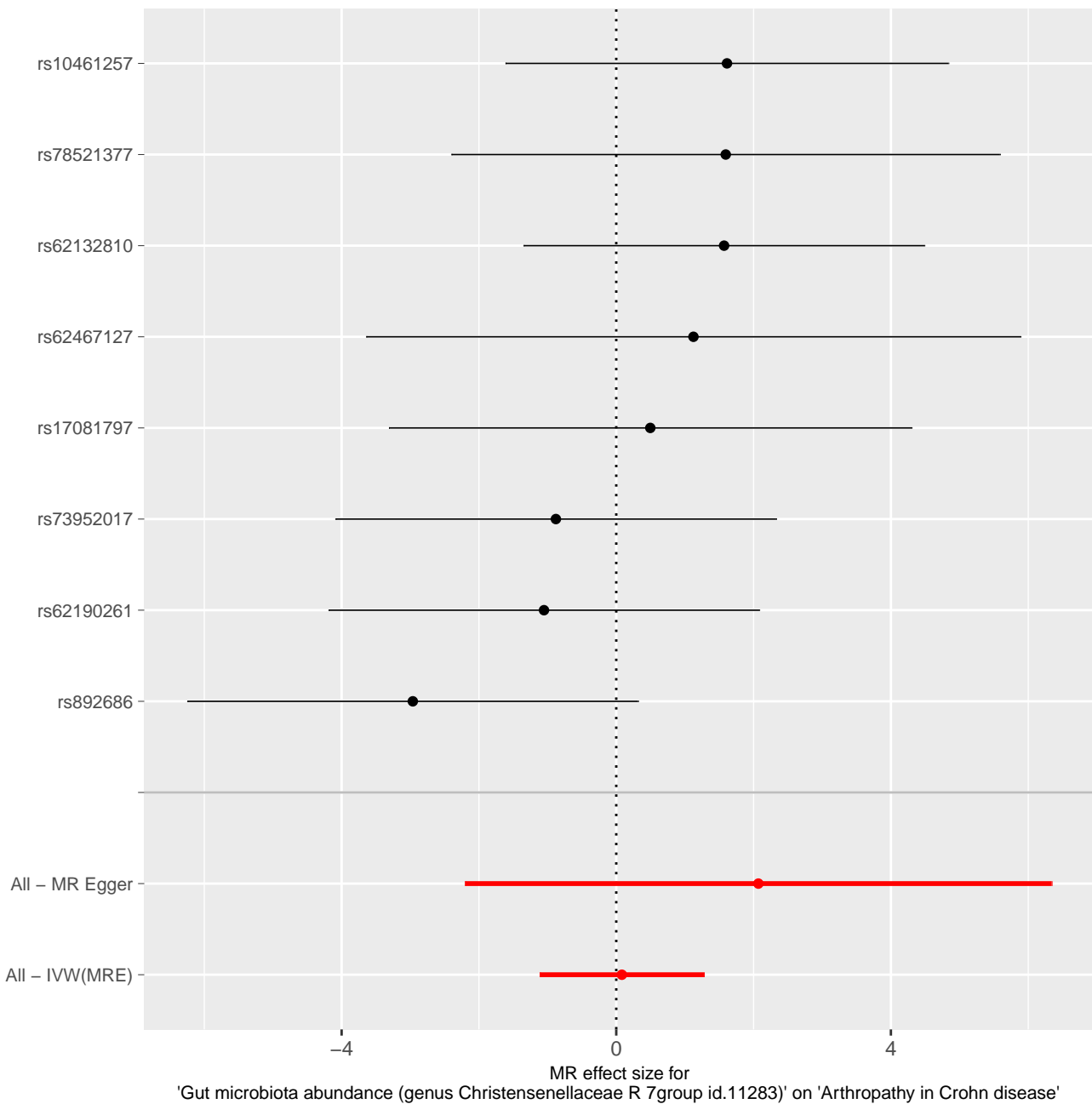


Batch 499 : Gut microbiota abundance (genus Butyrivibrio id.1993) on Arthropathy in Crohn disease

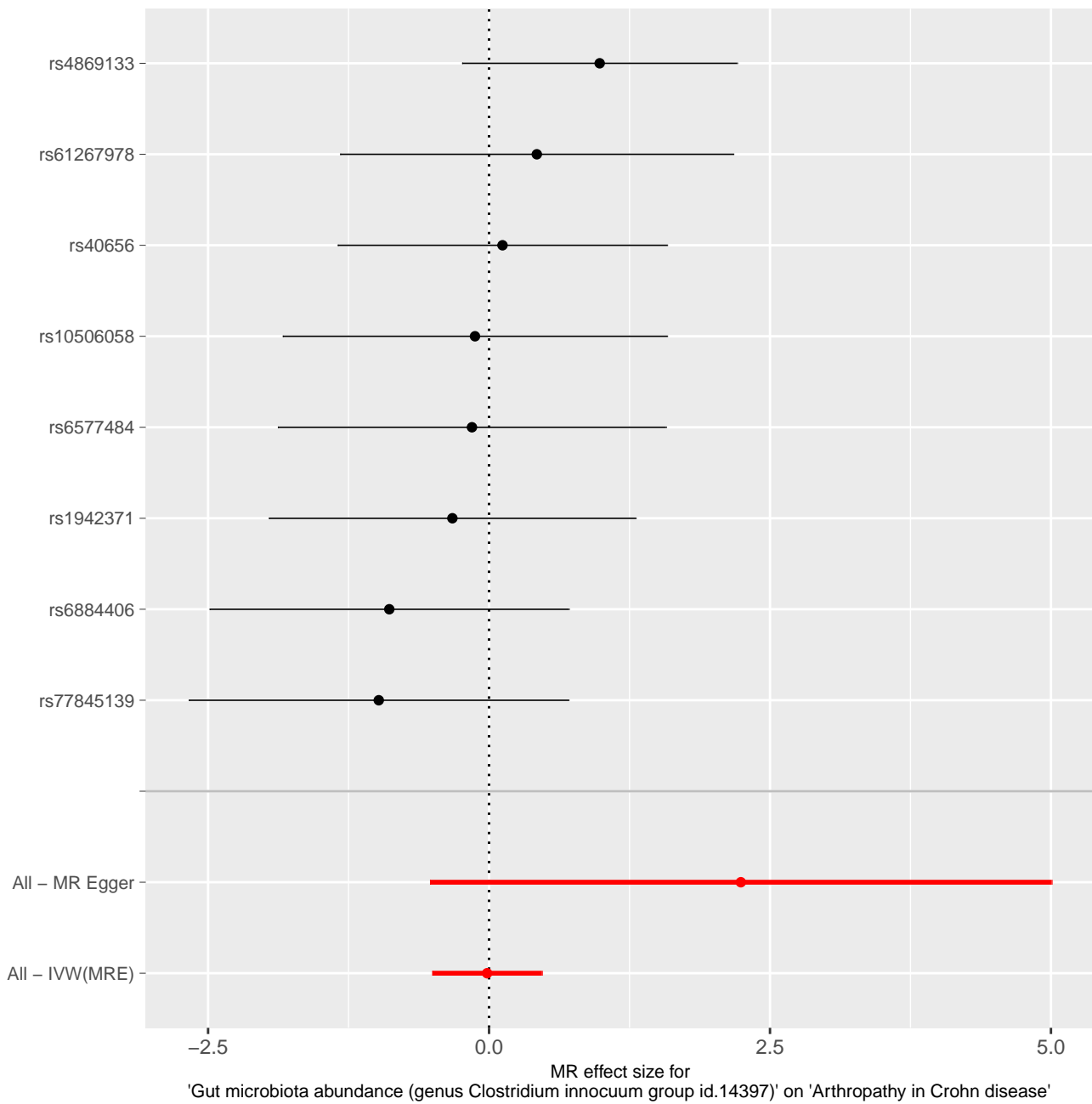


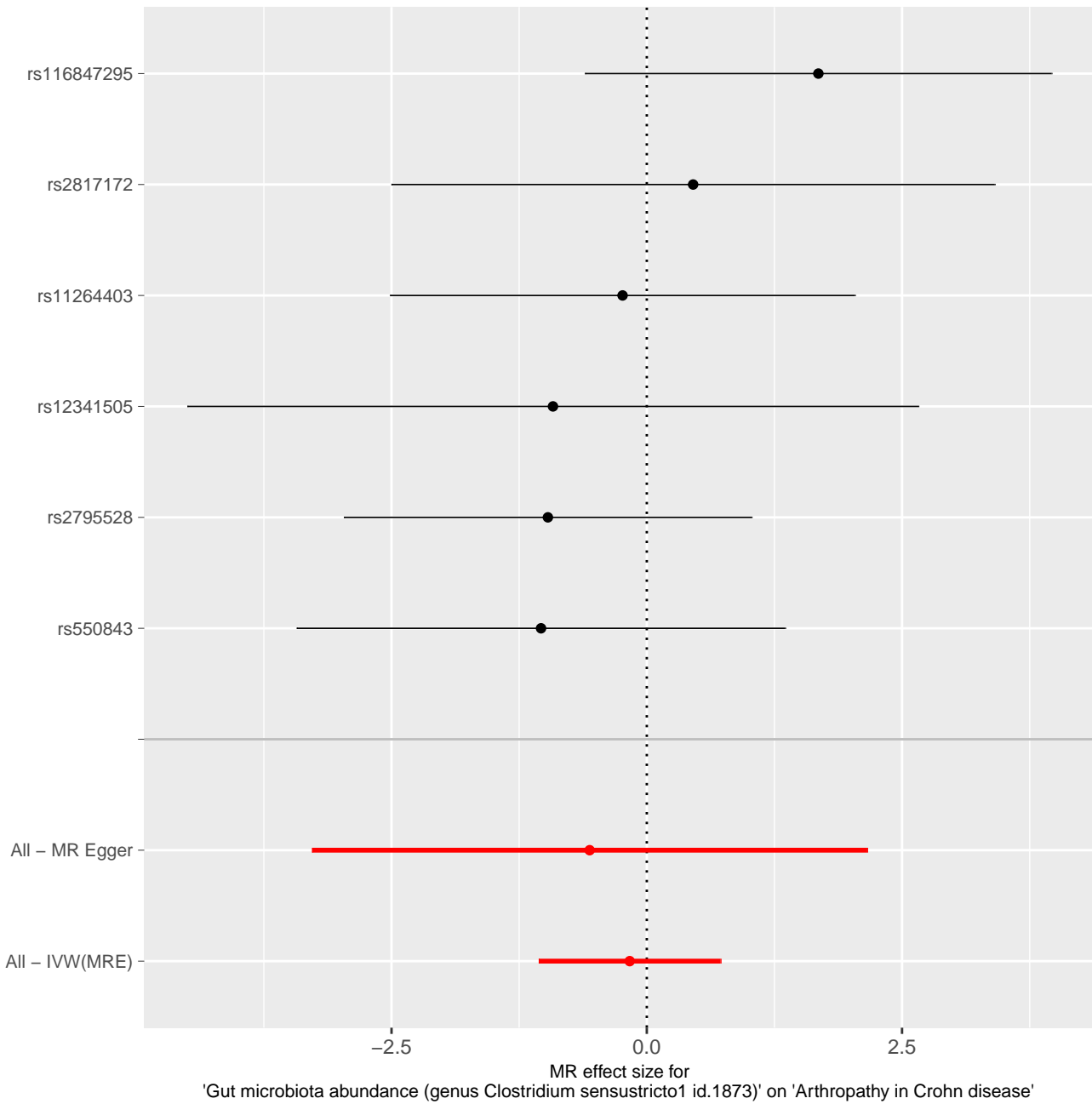


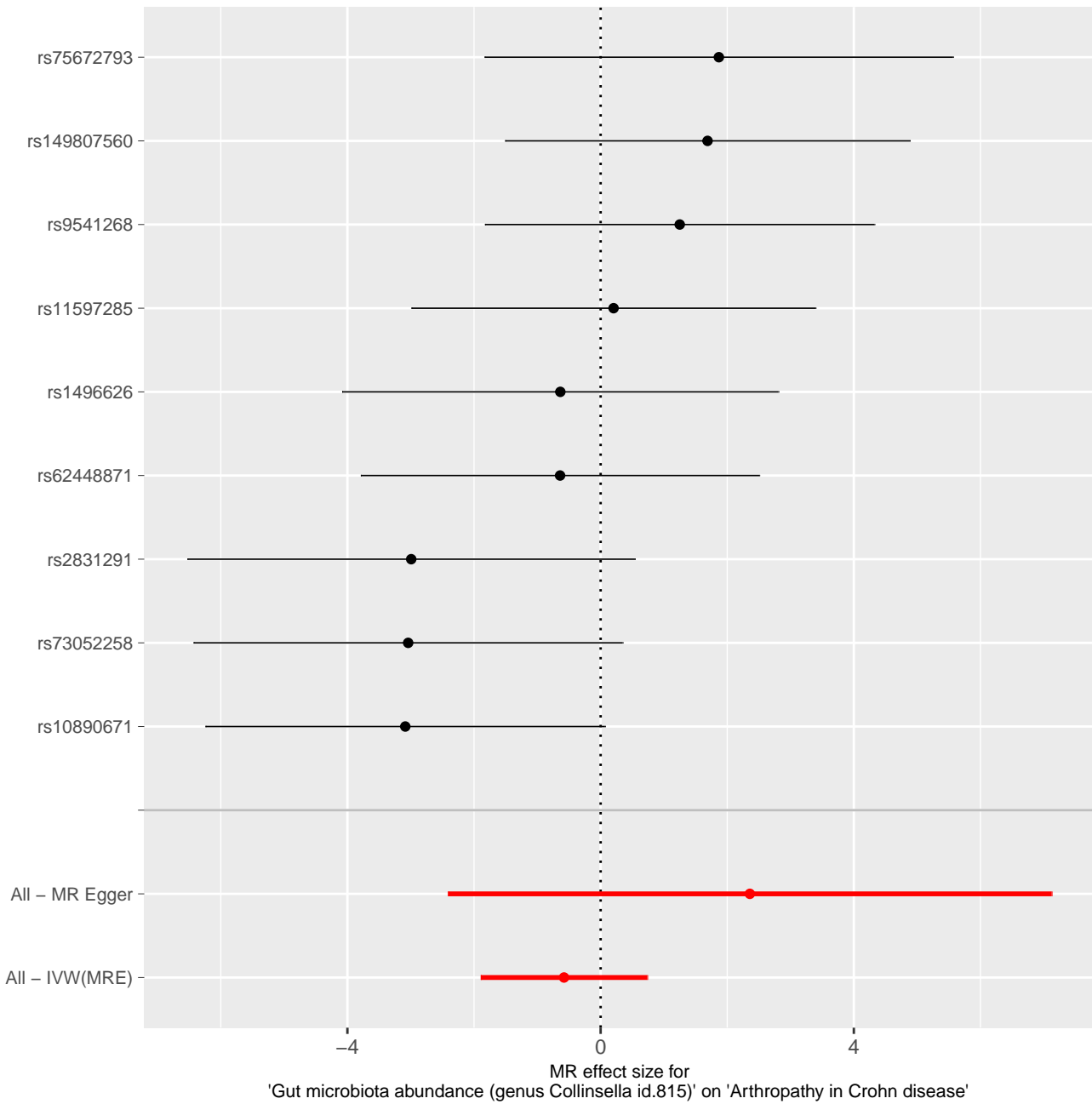




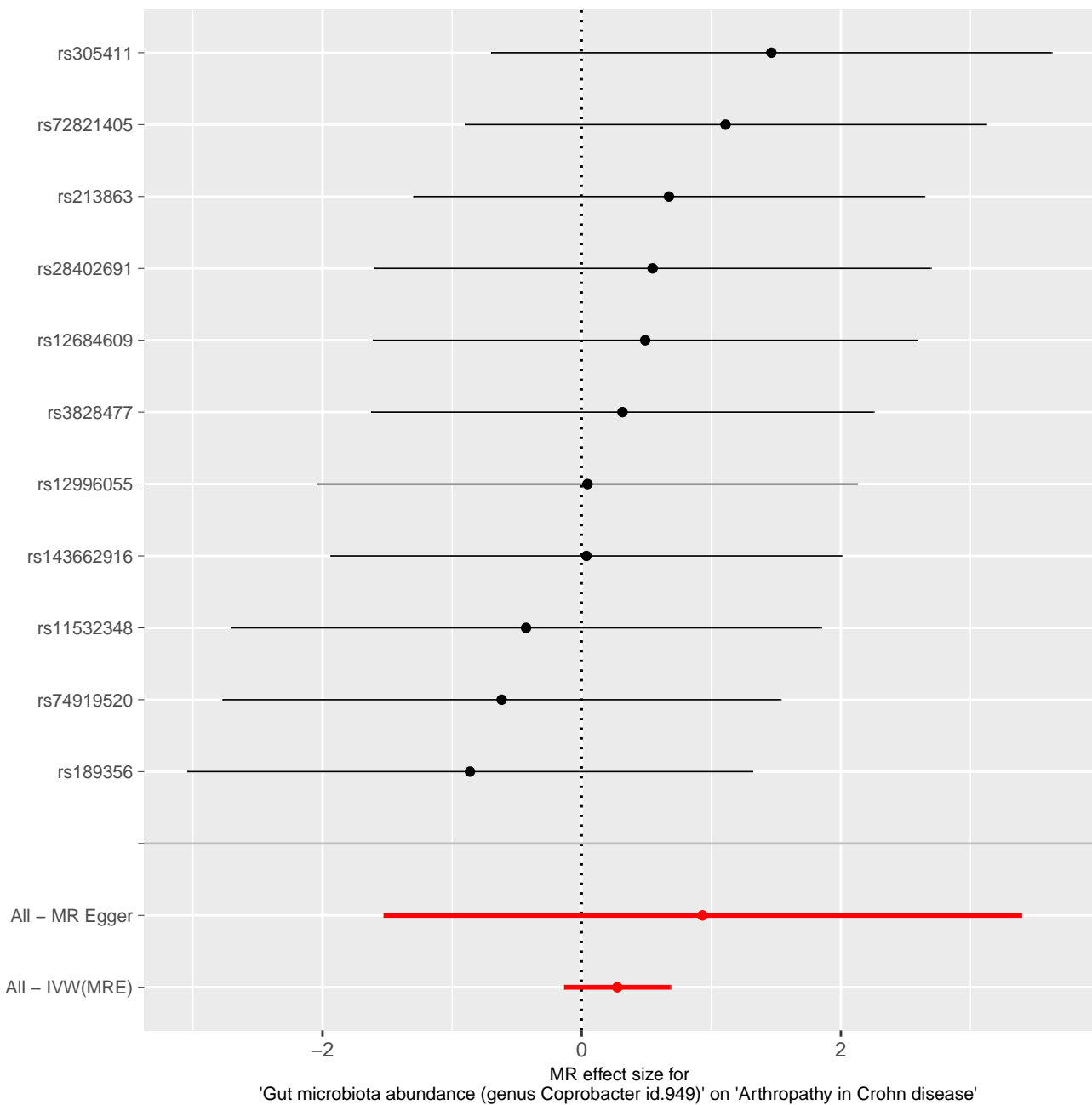
'Gut microbiota abundance (genus Christensenellaceae R 7group id.11283) on 'Arthropathy in Crohn disease'

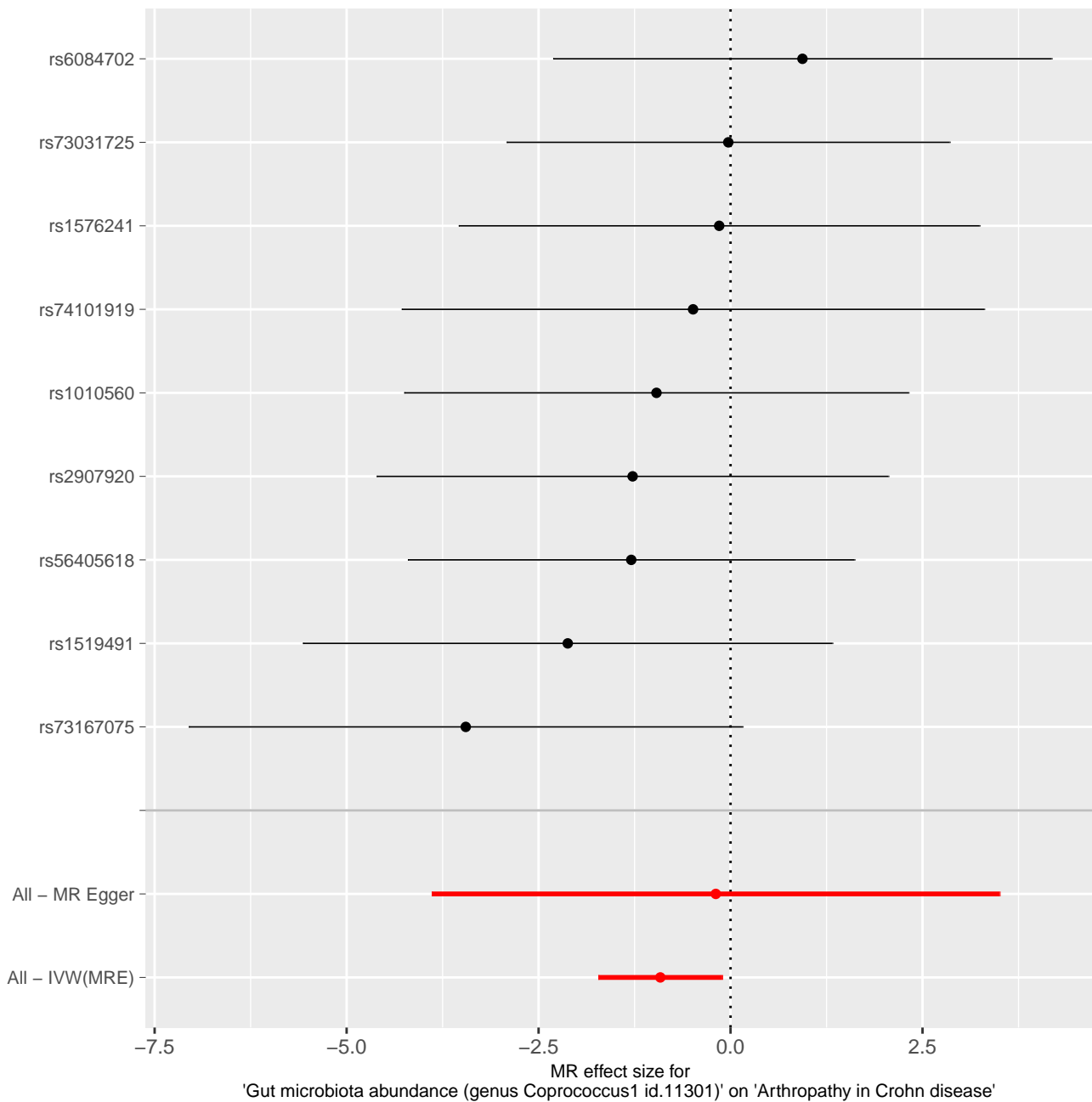


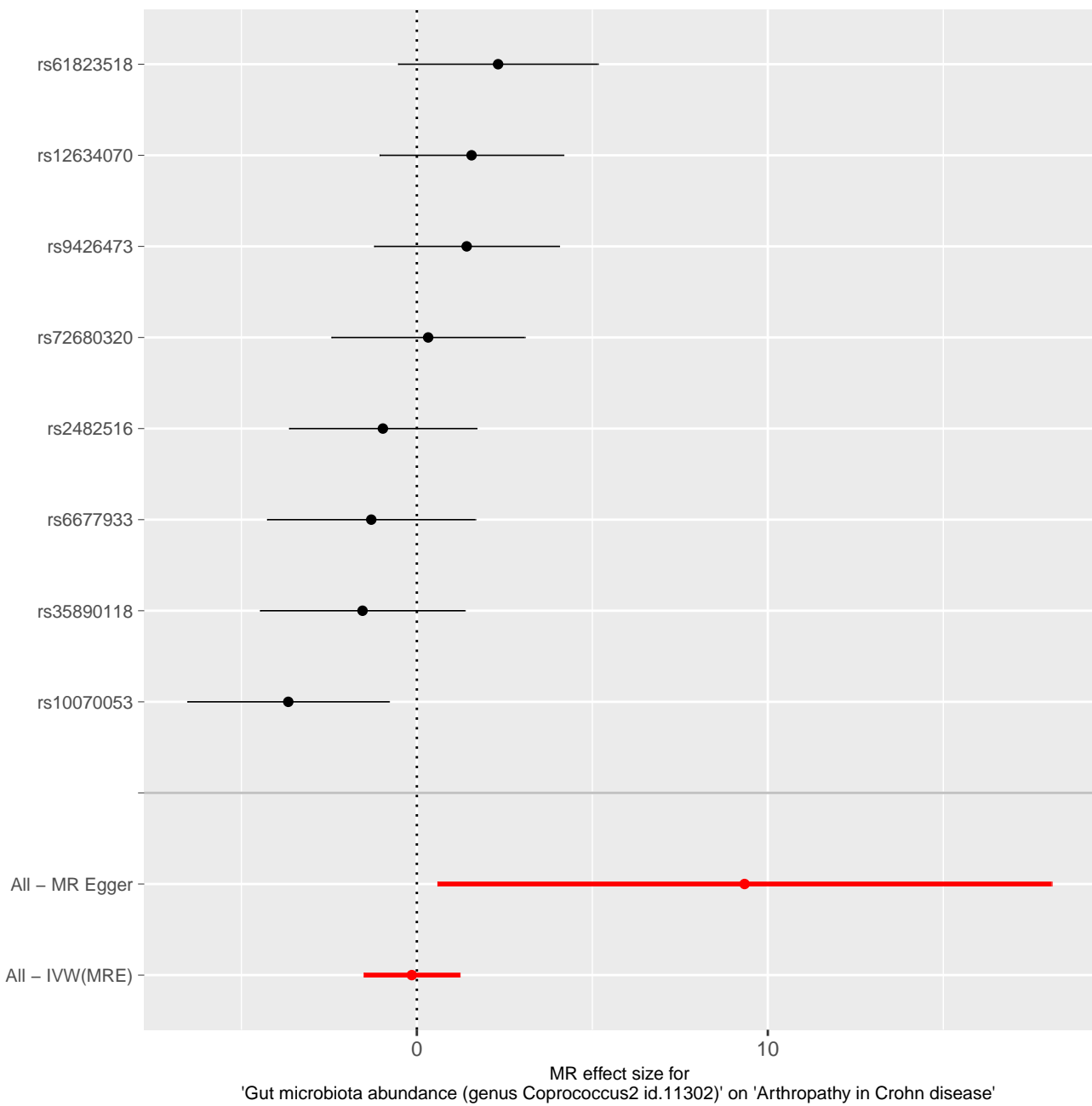


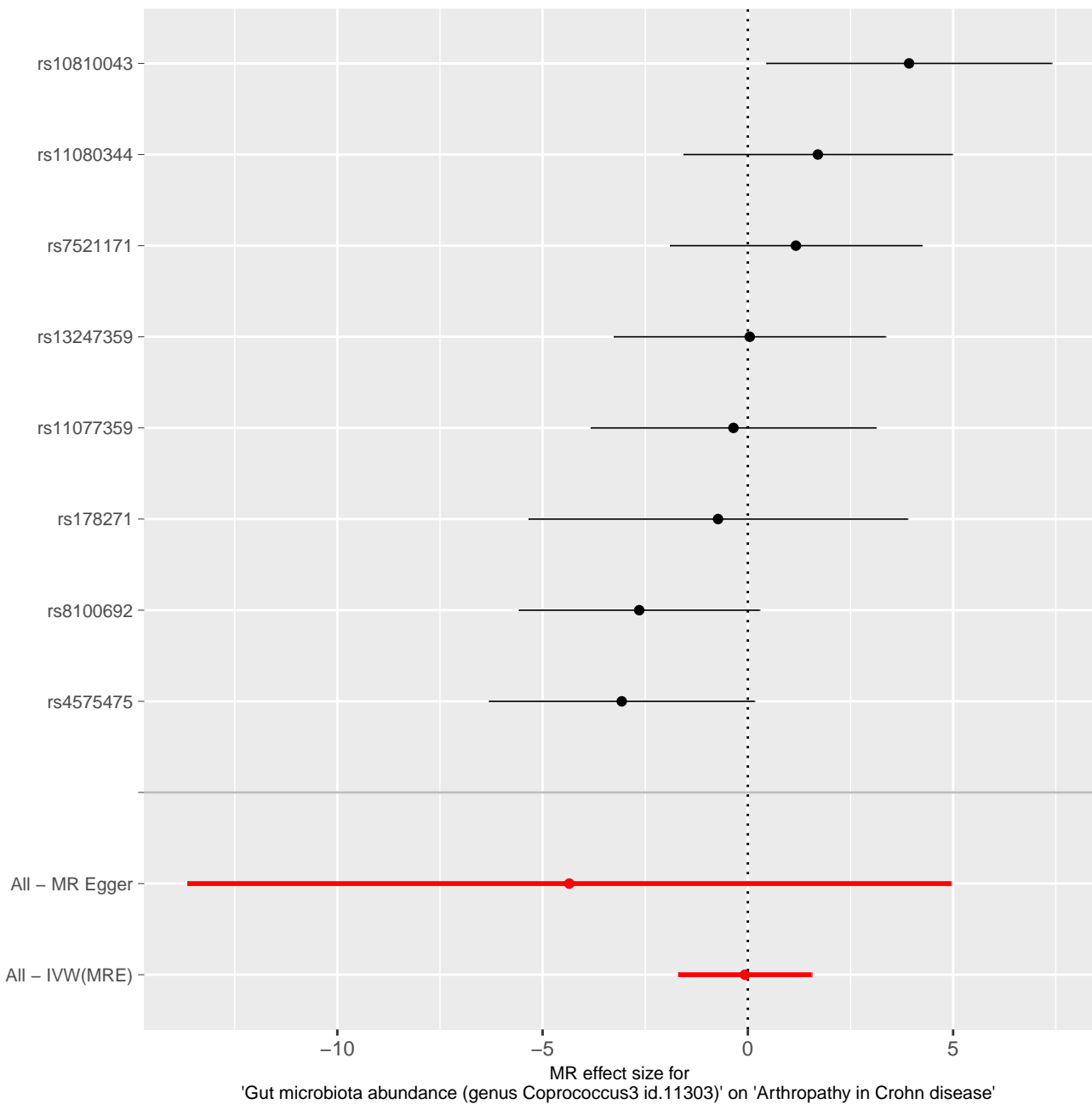


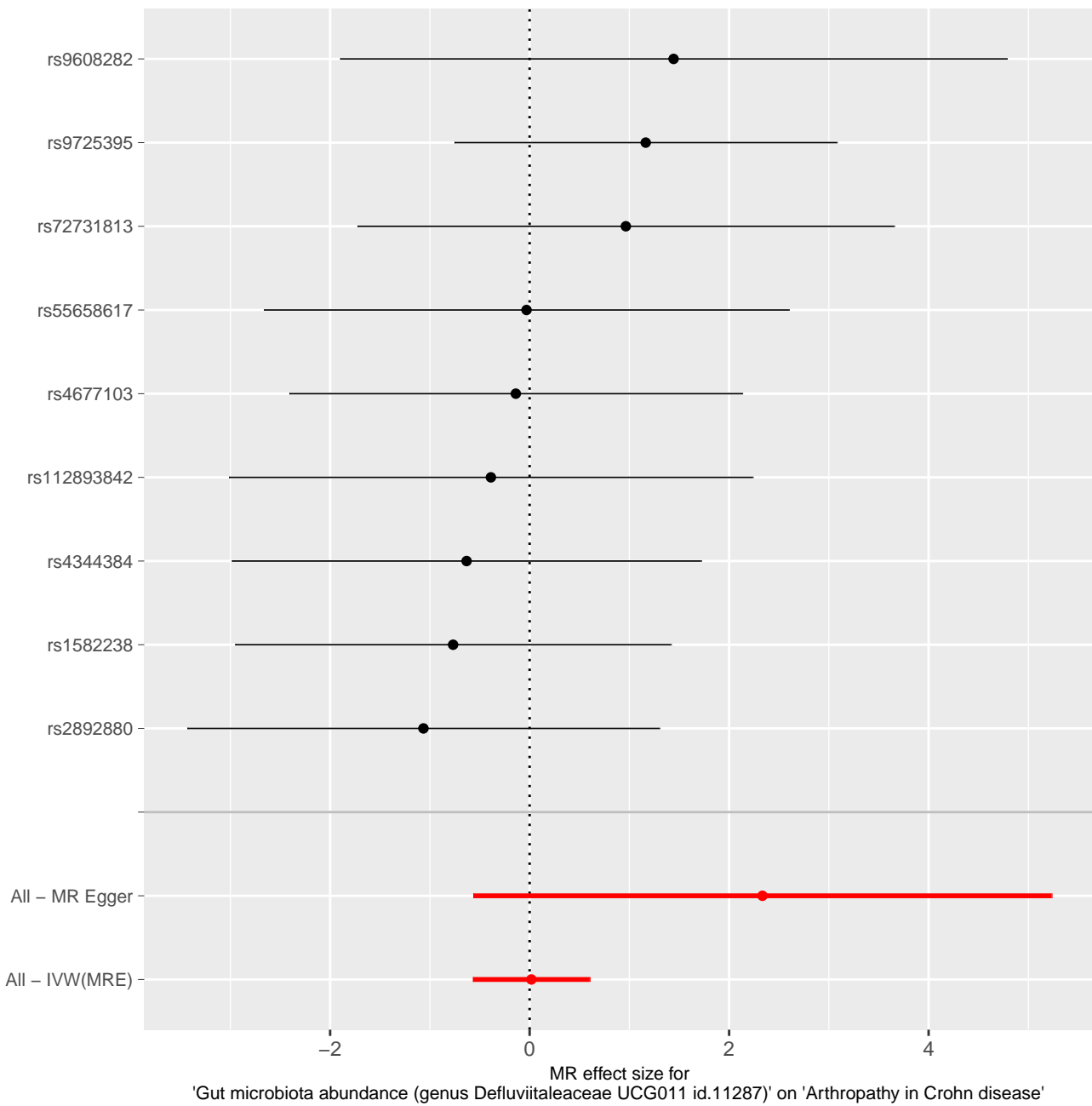
Batch 506 : Gut microbiota abundance (genus Coprobacter id.949) on Arthropathy in Crohn disease



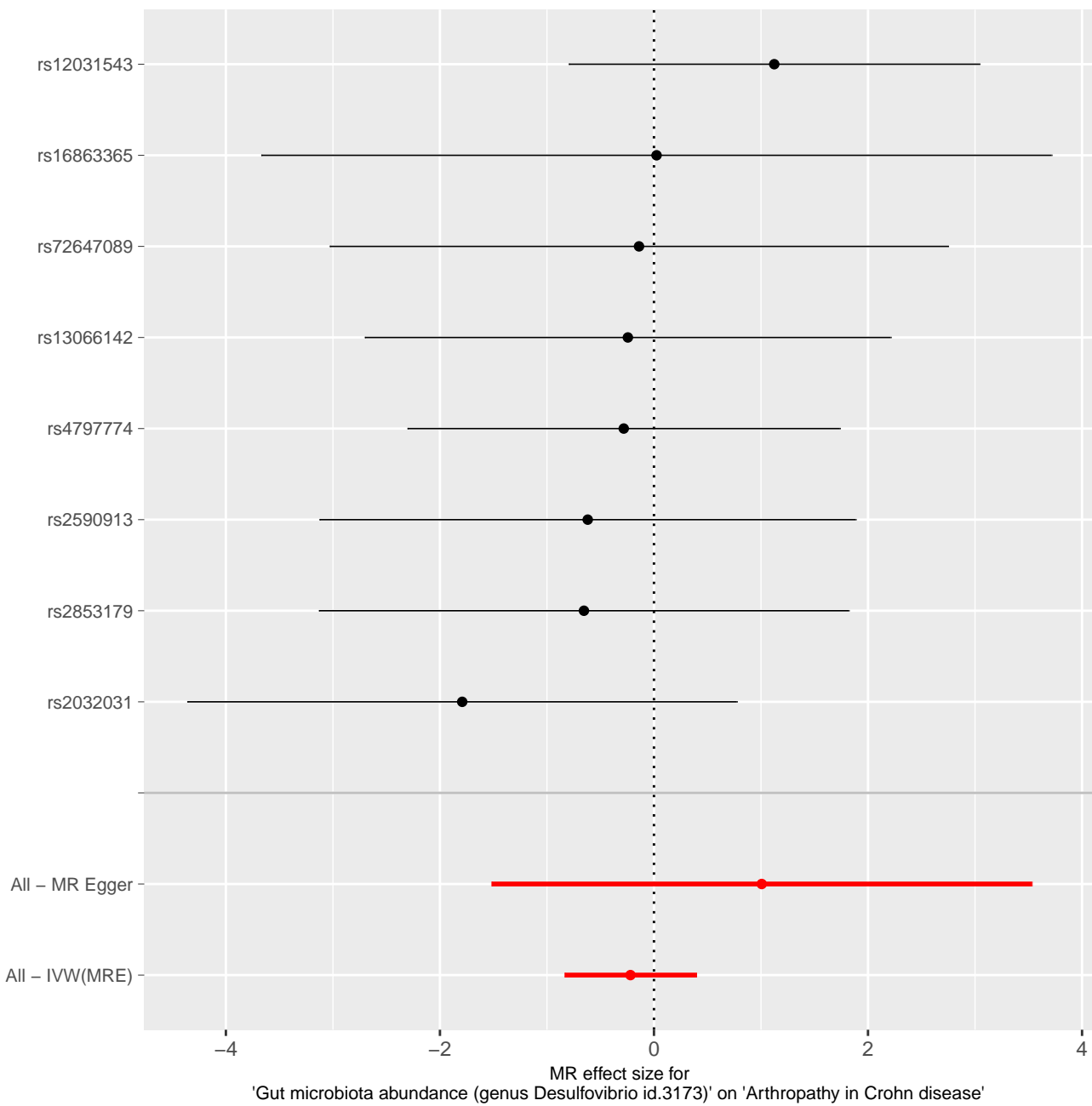




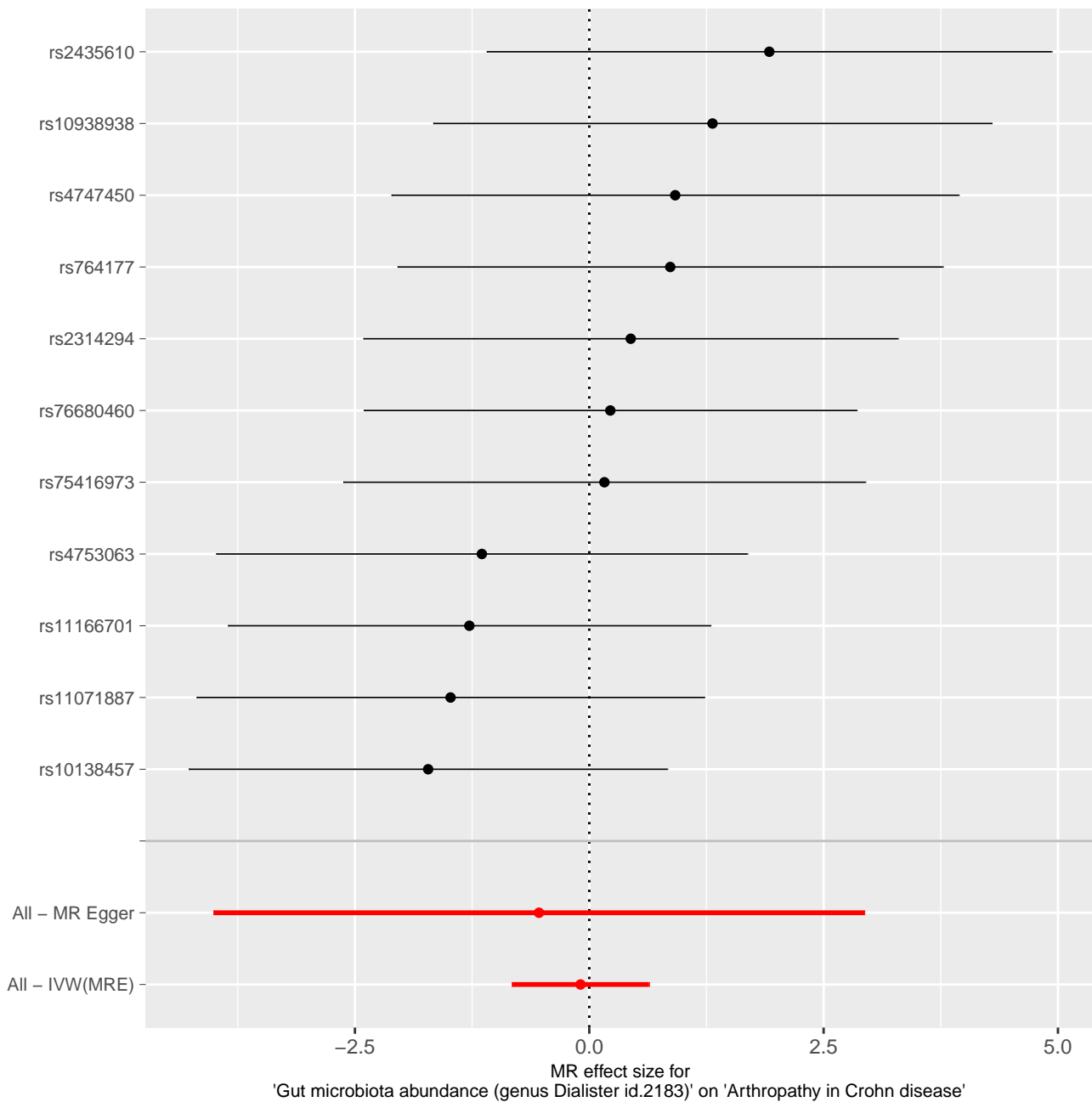


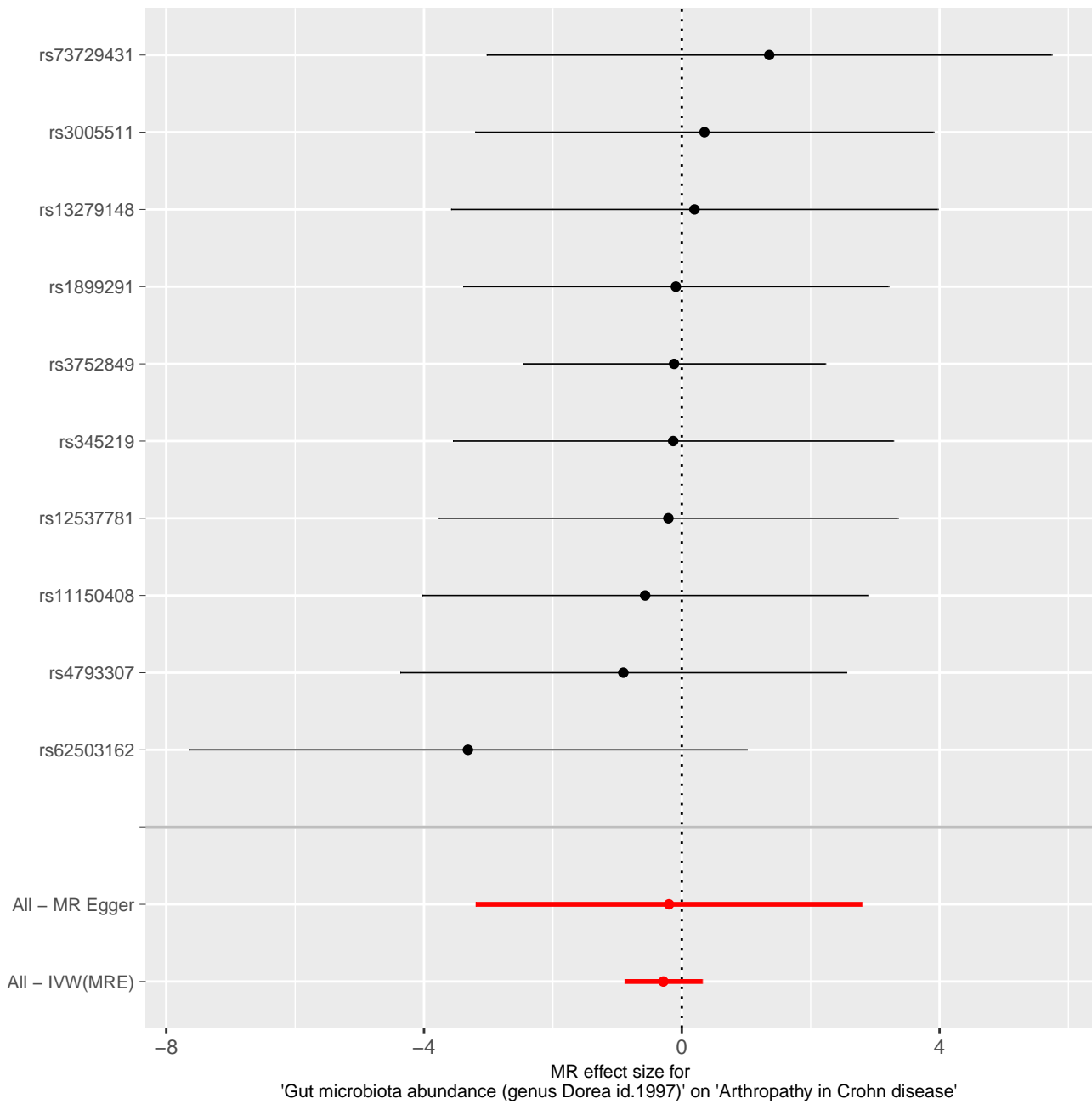


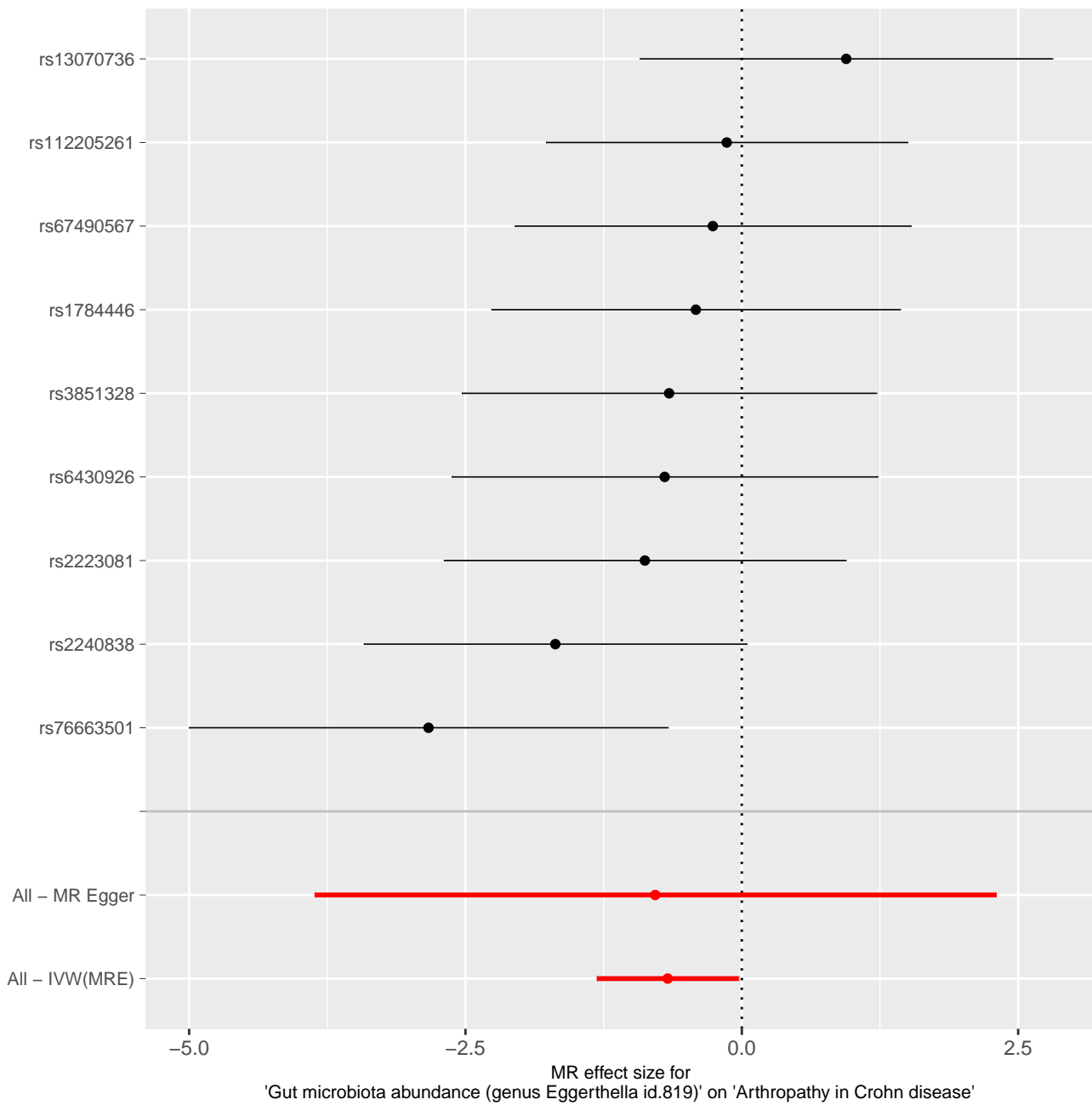
Batch 511 : Gut microbiota abundance (genus Desulfovibrio id.3173) on Arthropathy in Crohn disease



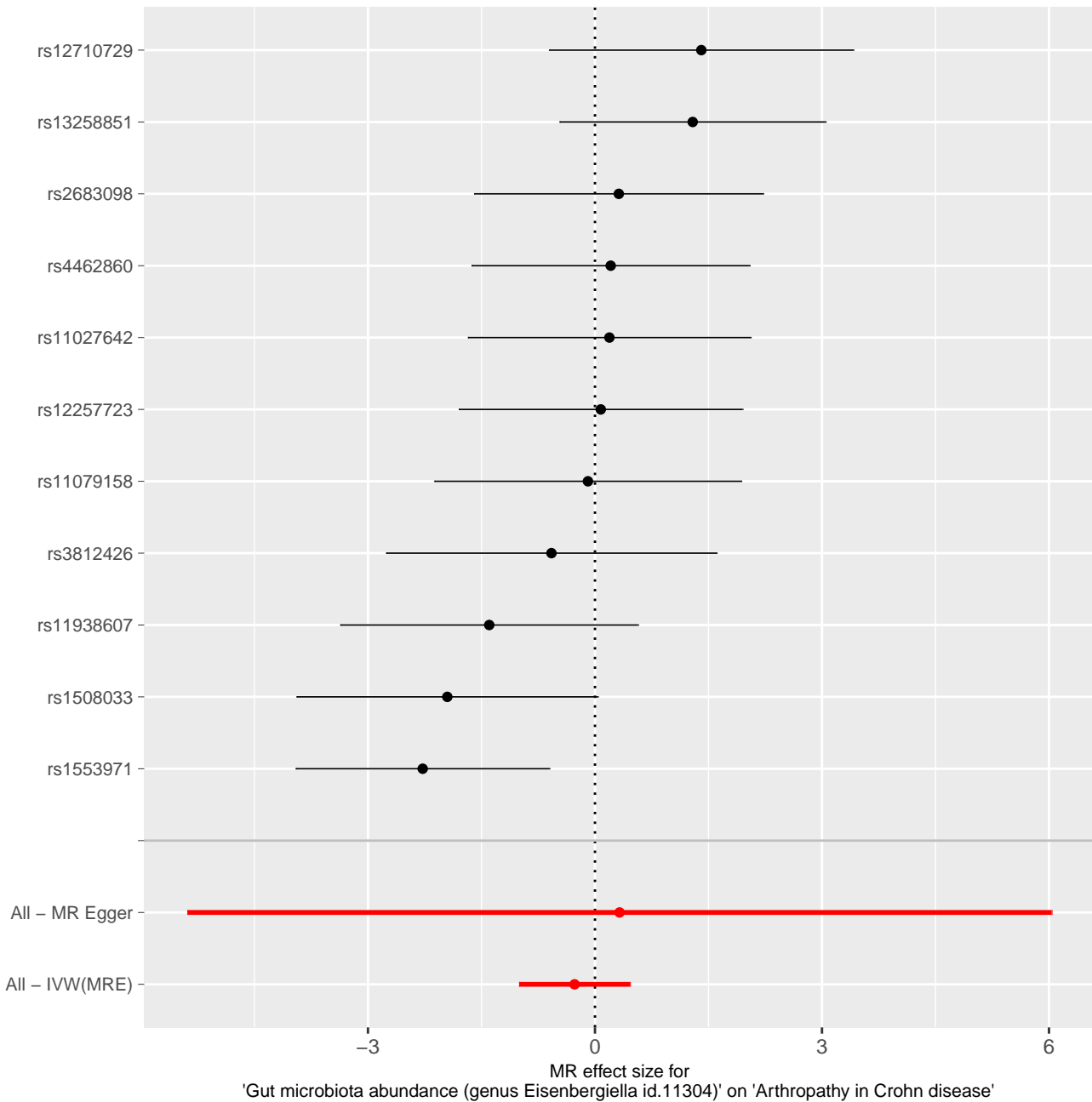
Batch 512 : Gut microbiota abundance (genus Dialister id.2183) on Arthropathy in Crohn disease

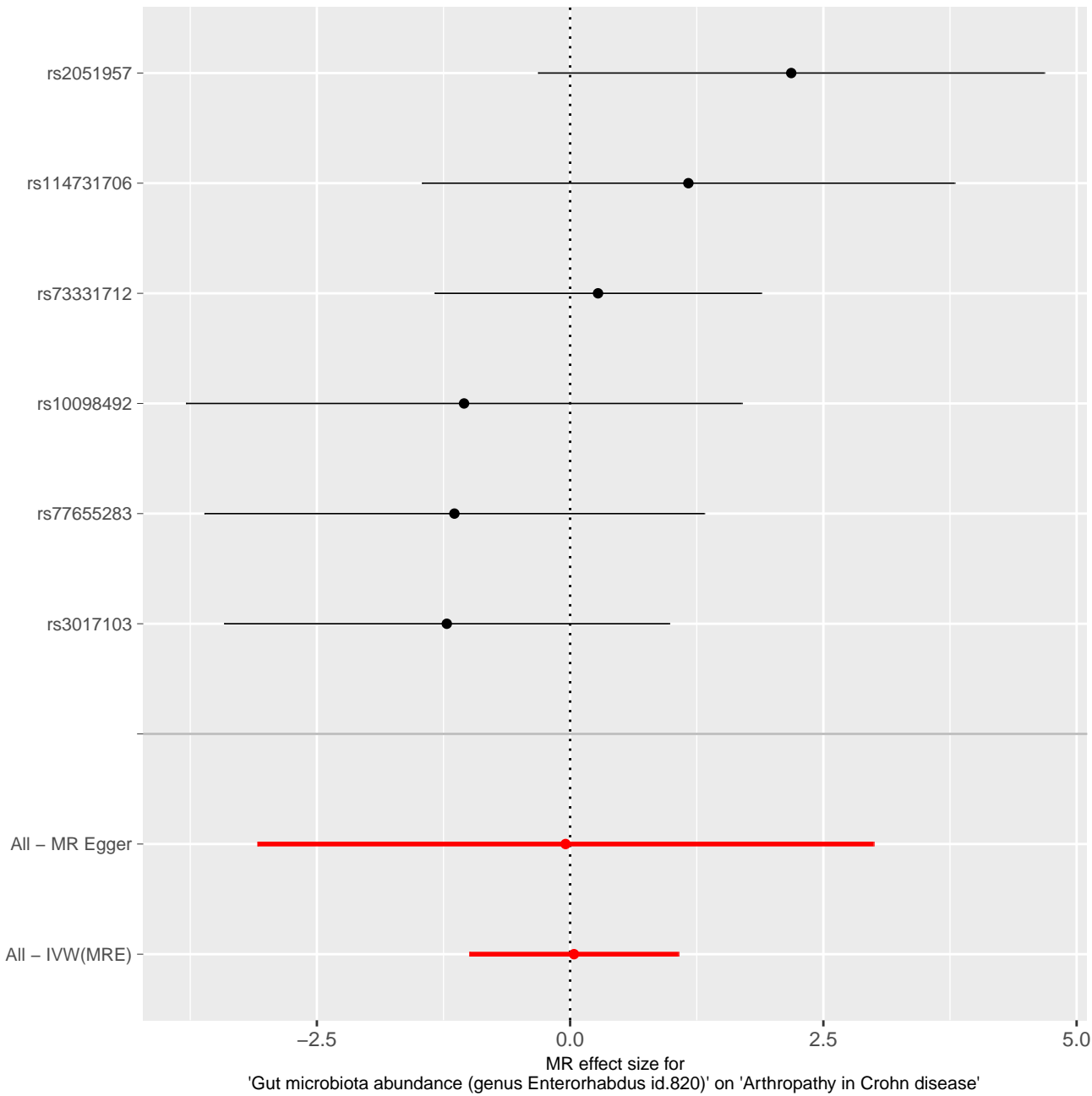




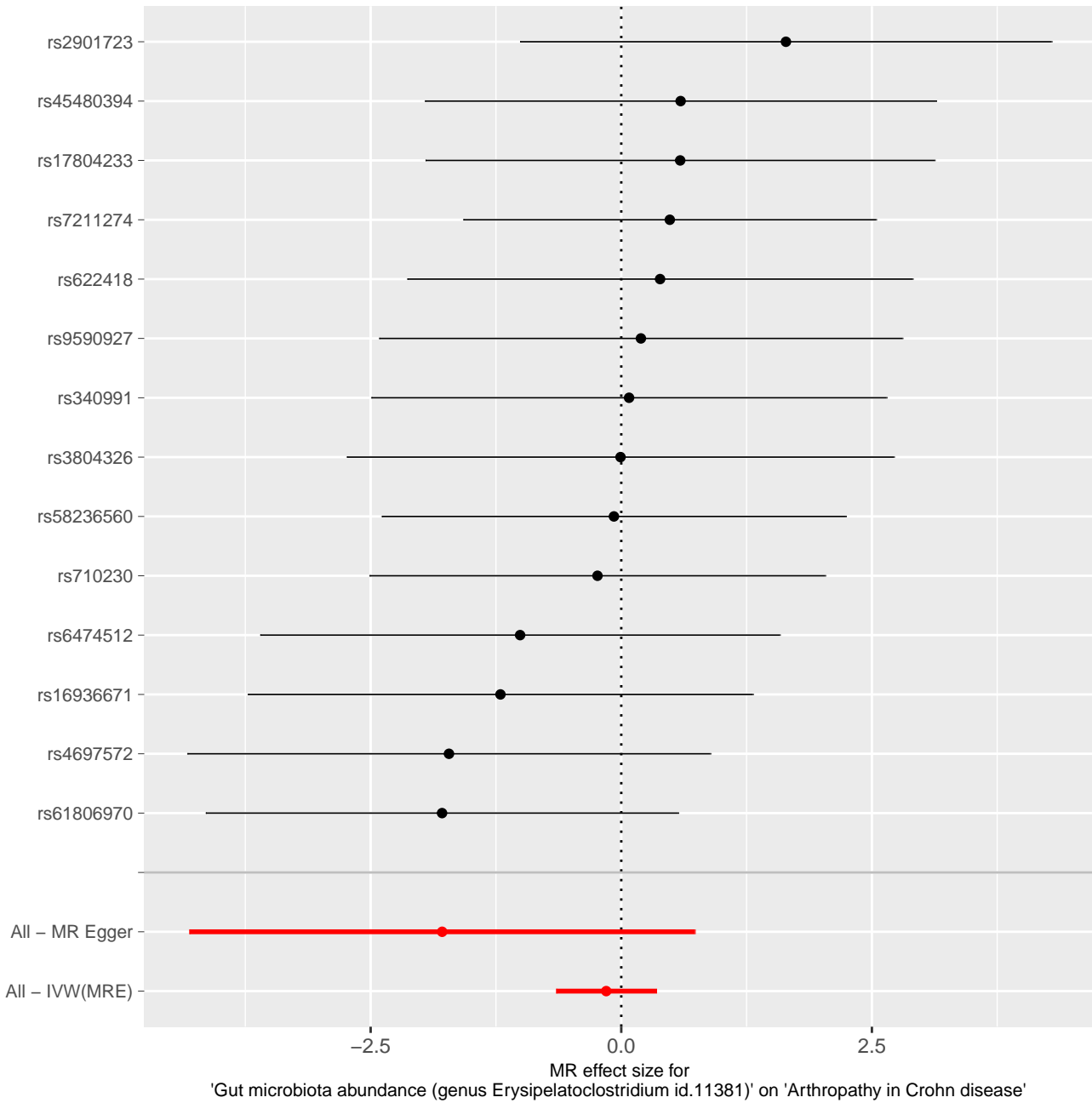


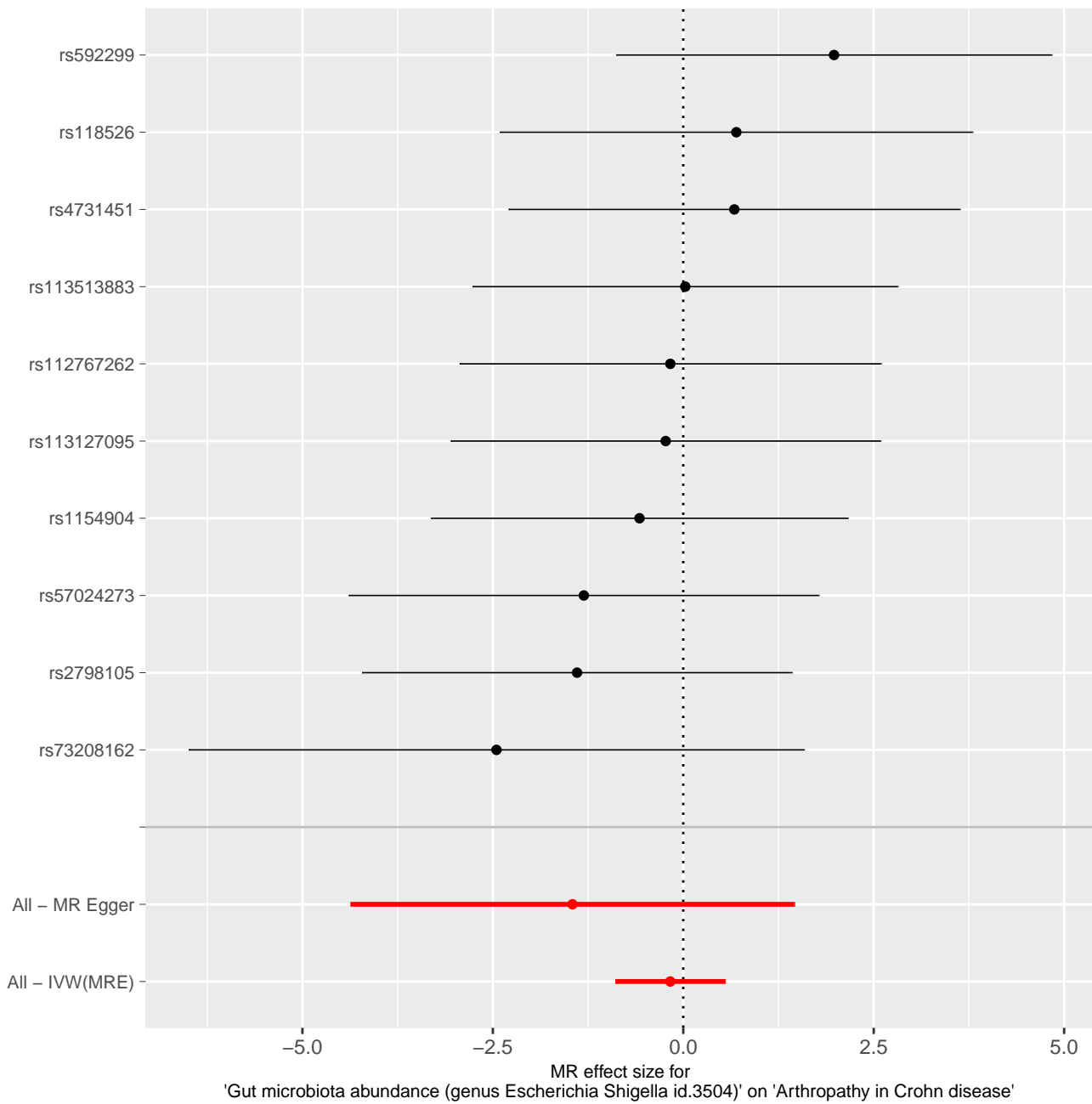
Batch 515 : Gut microbiota abundance (genus Eisenbergiella id.11304) on Arthropathy in Crohn disease

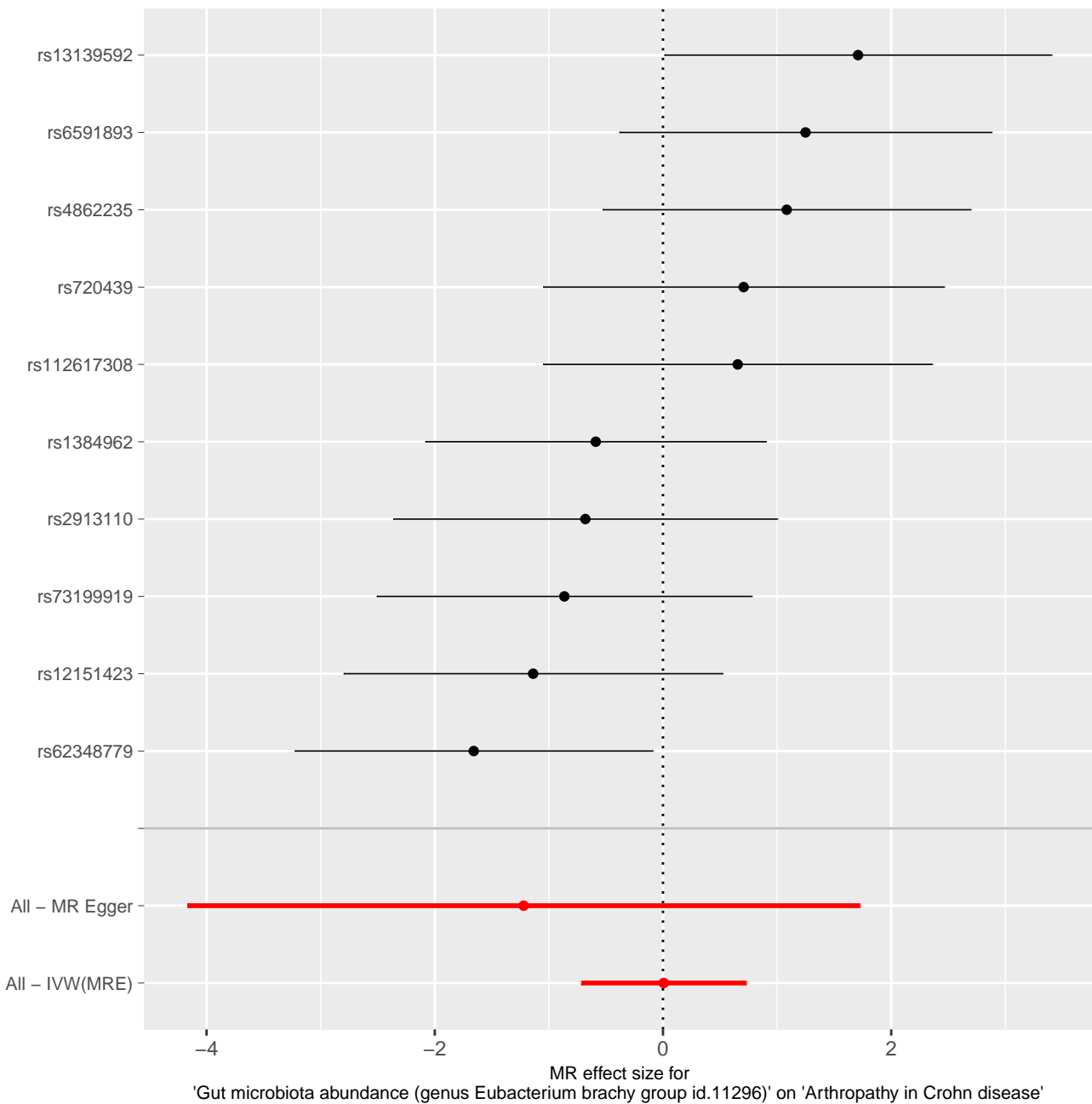


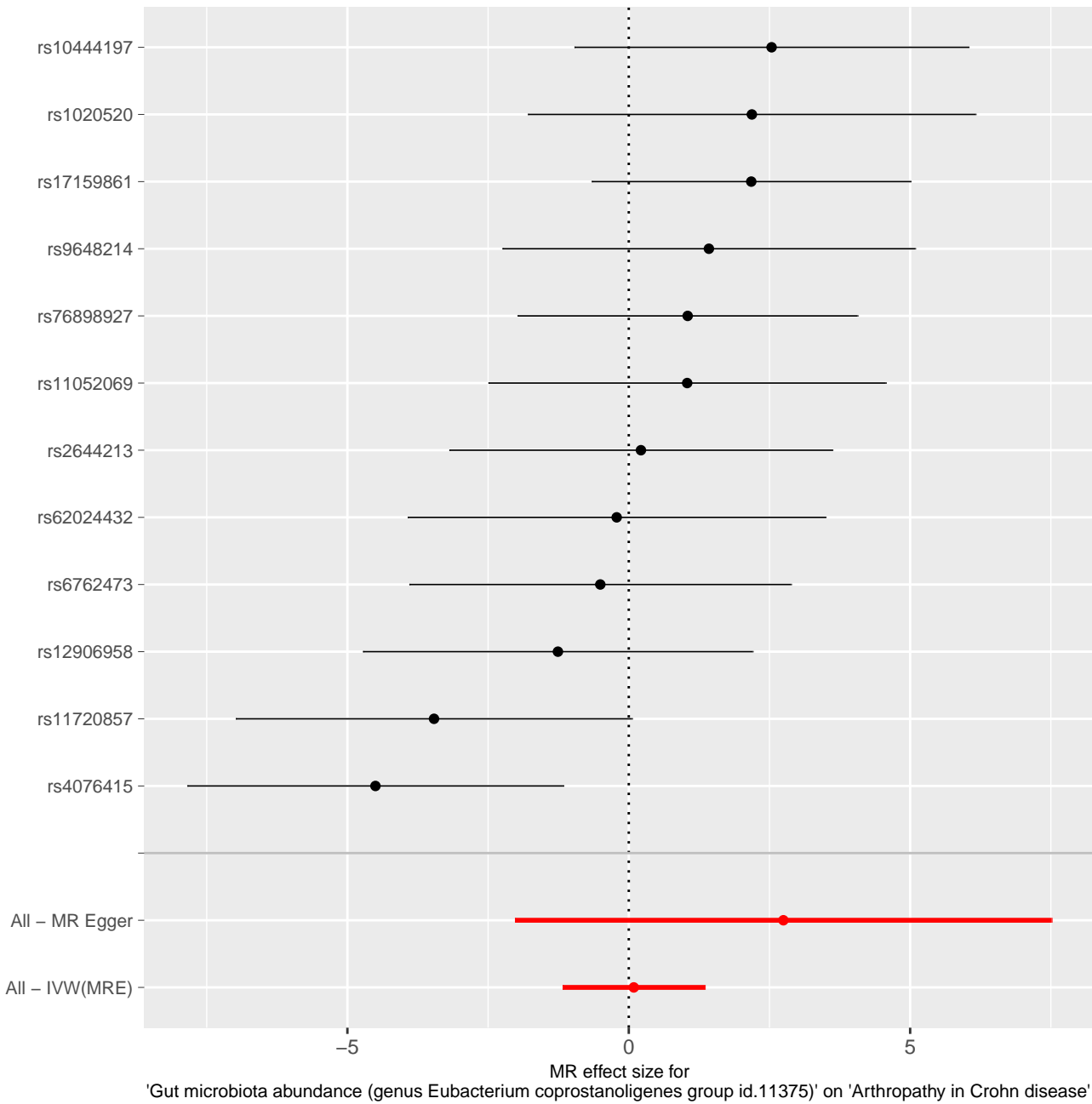


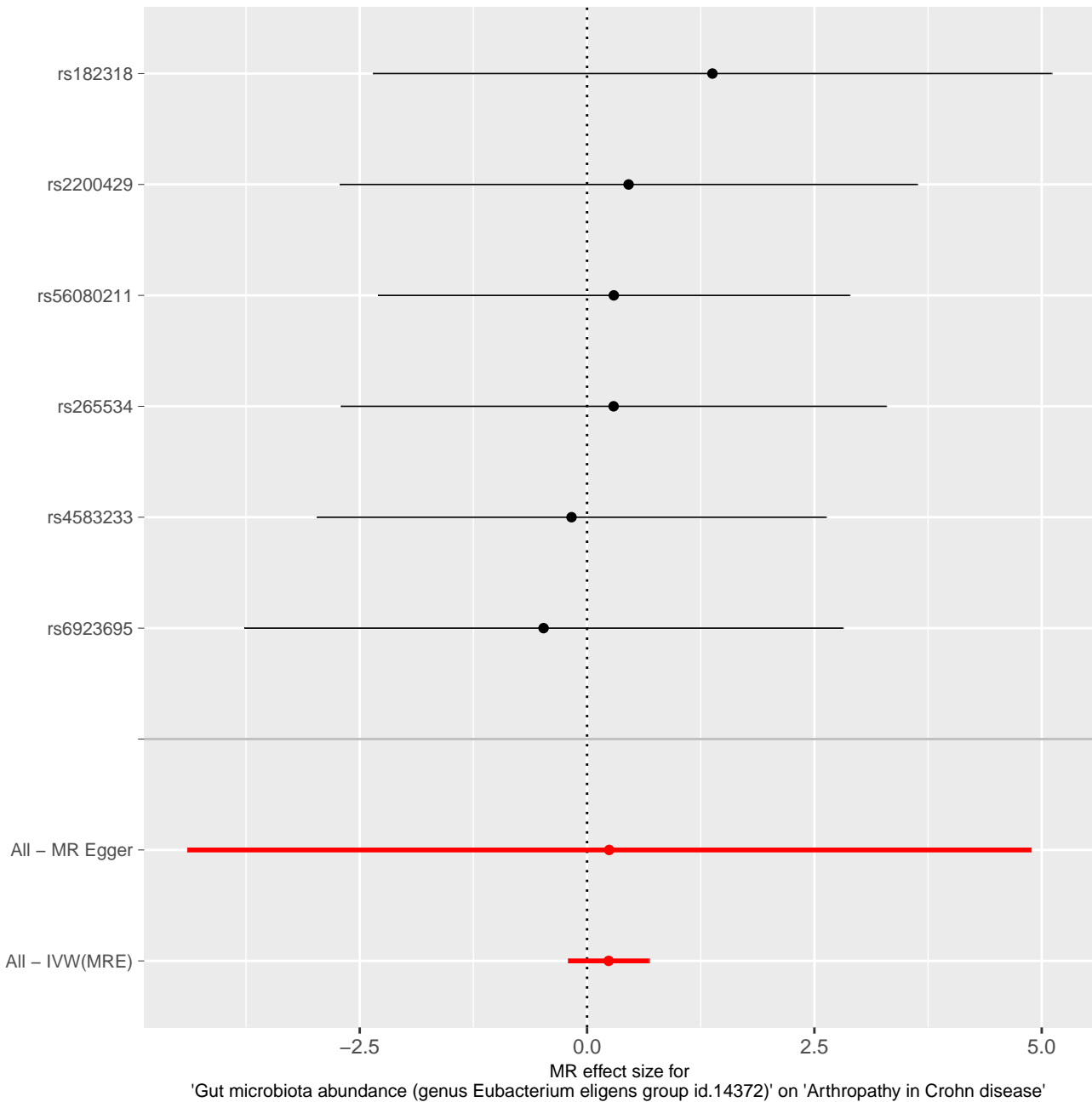
Batch 517 : Gut microbiota abundance (genus Erysipelatoclostridium id.11381) on Arthropathy in Crohn disease

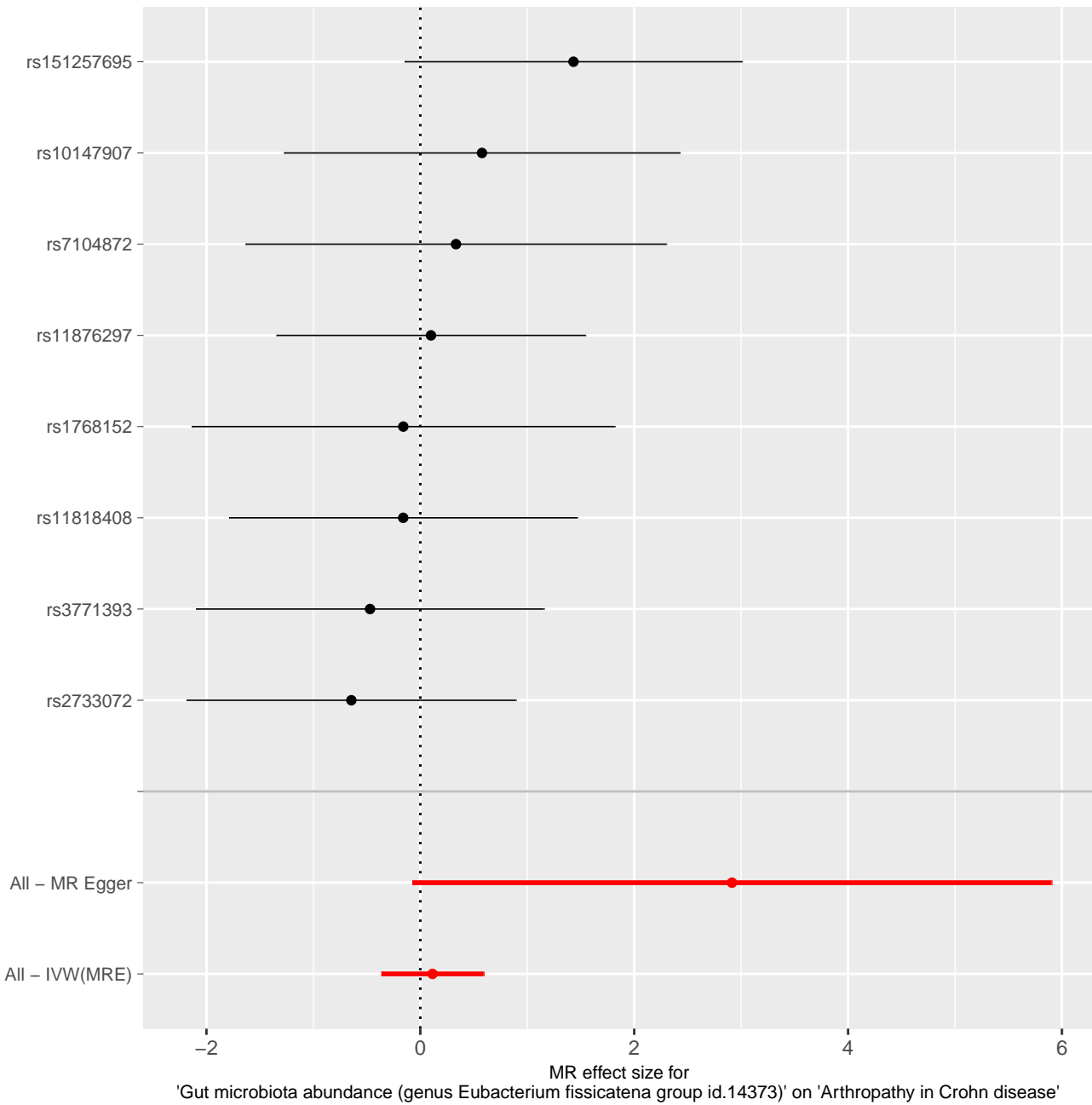


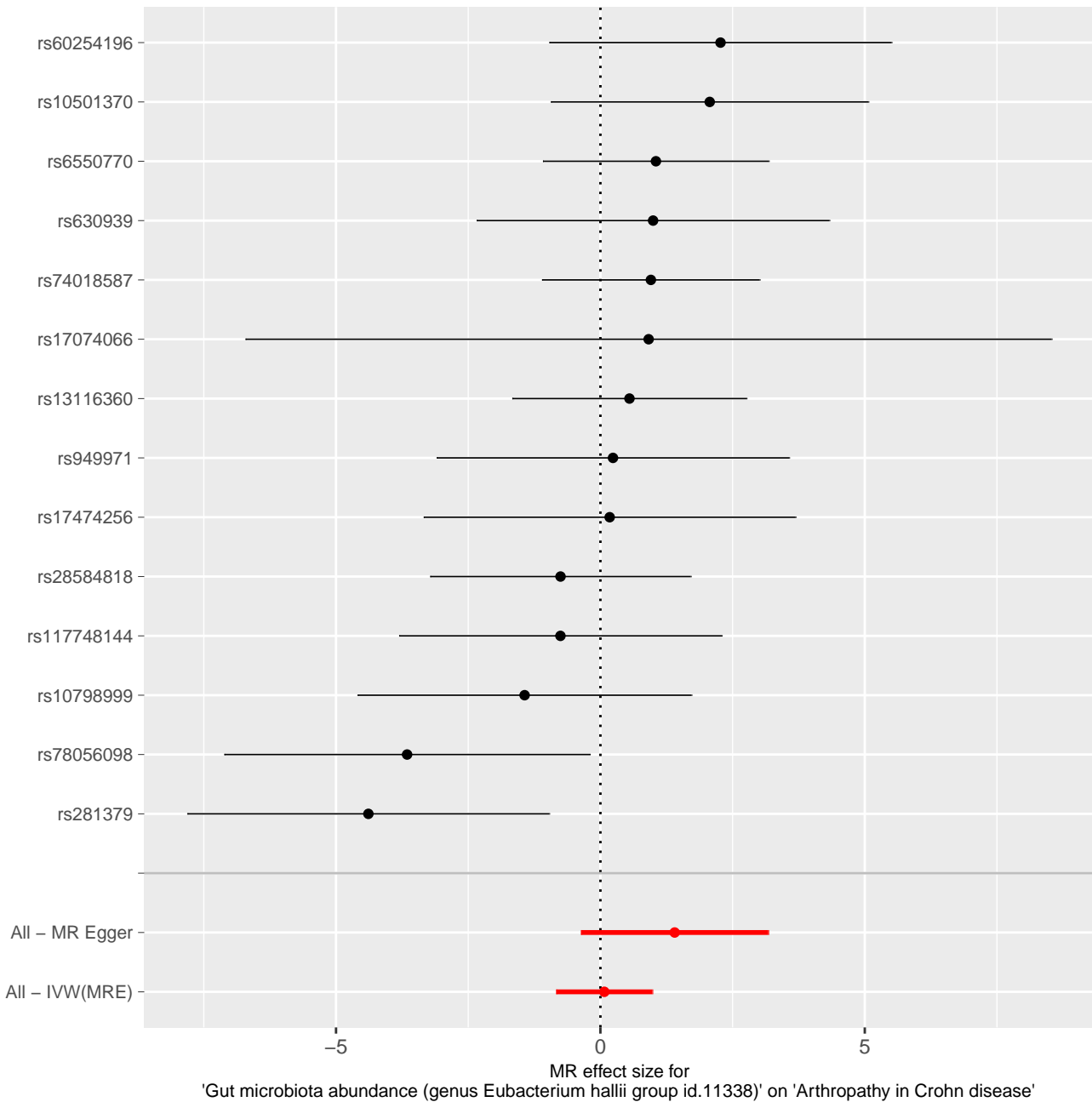


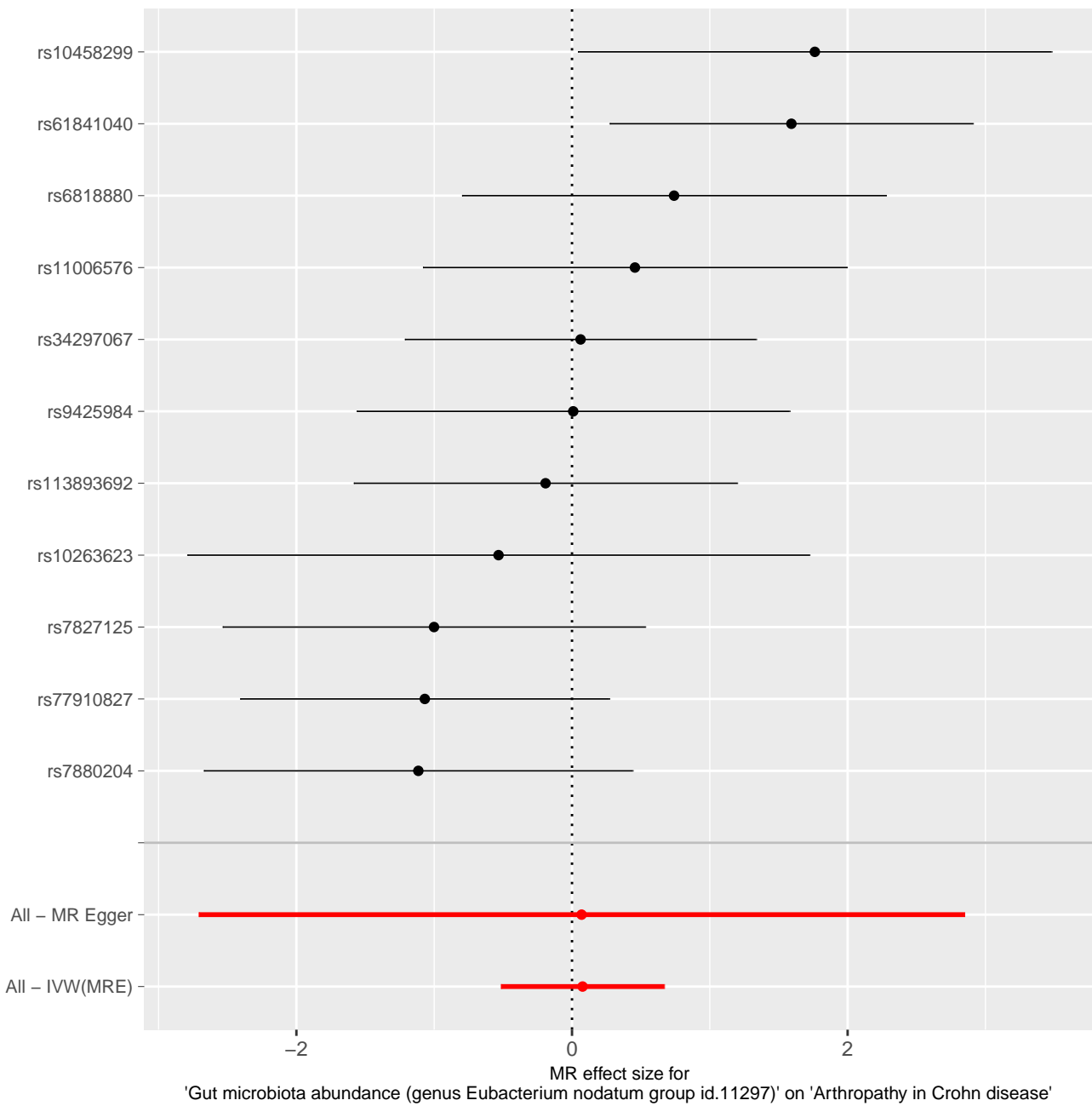


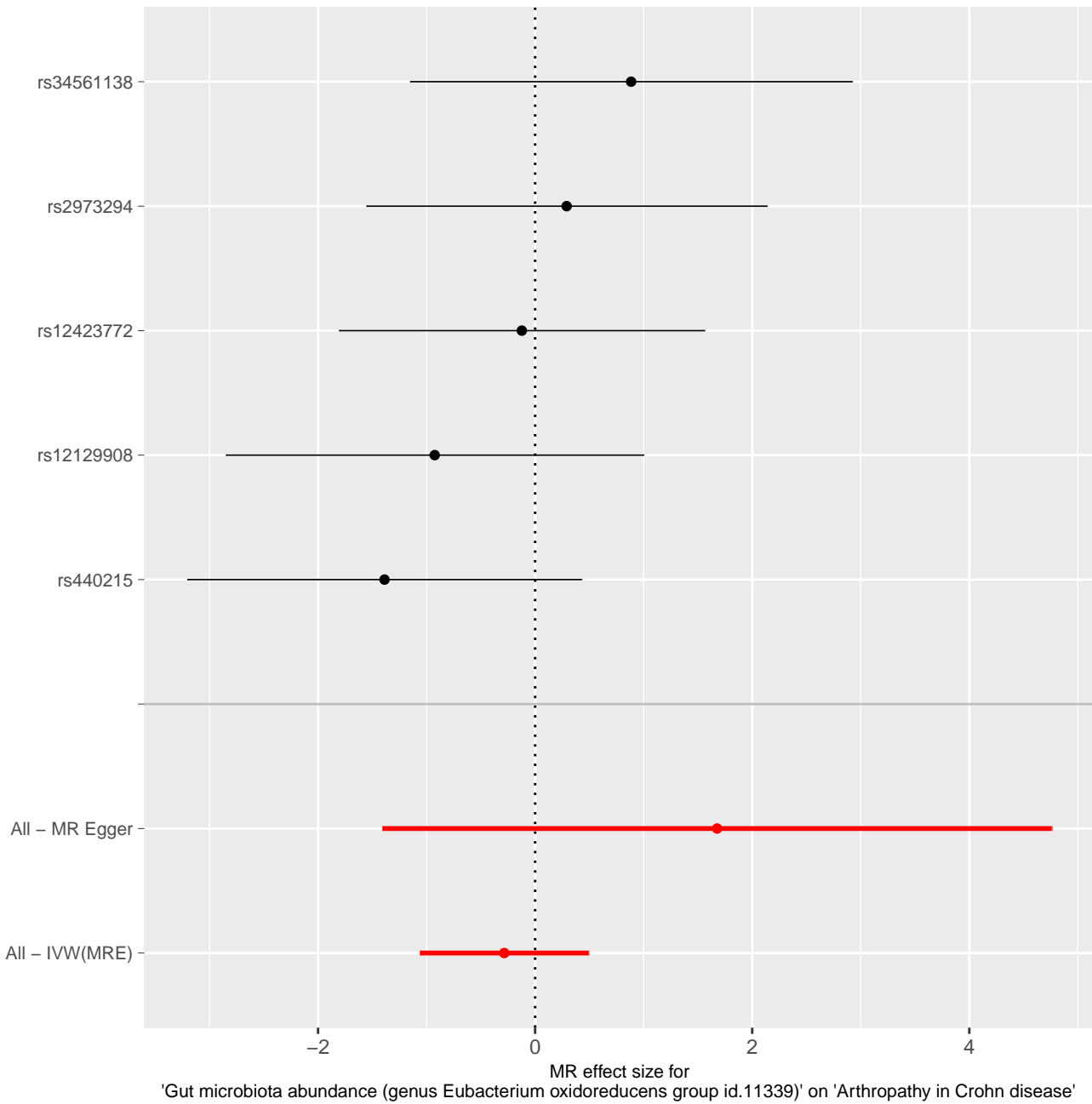


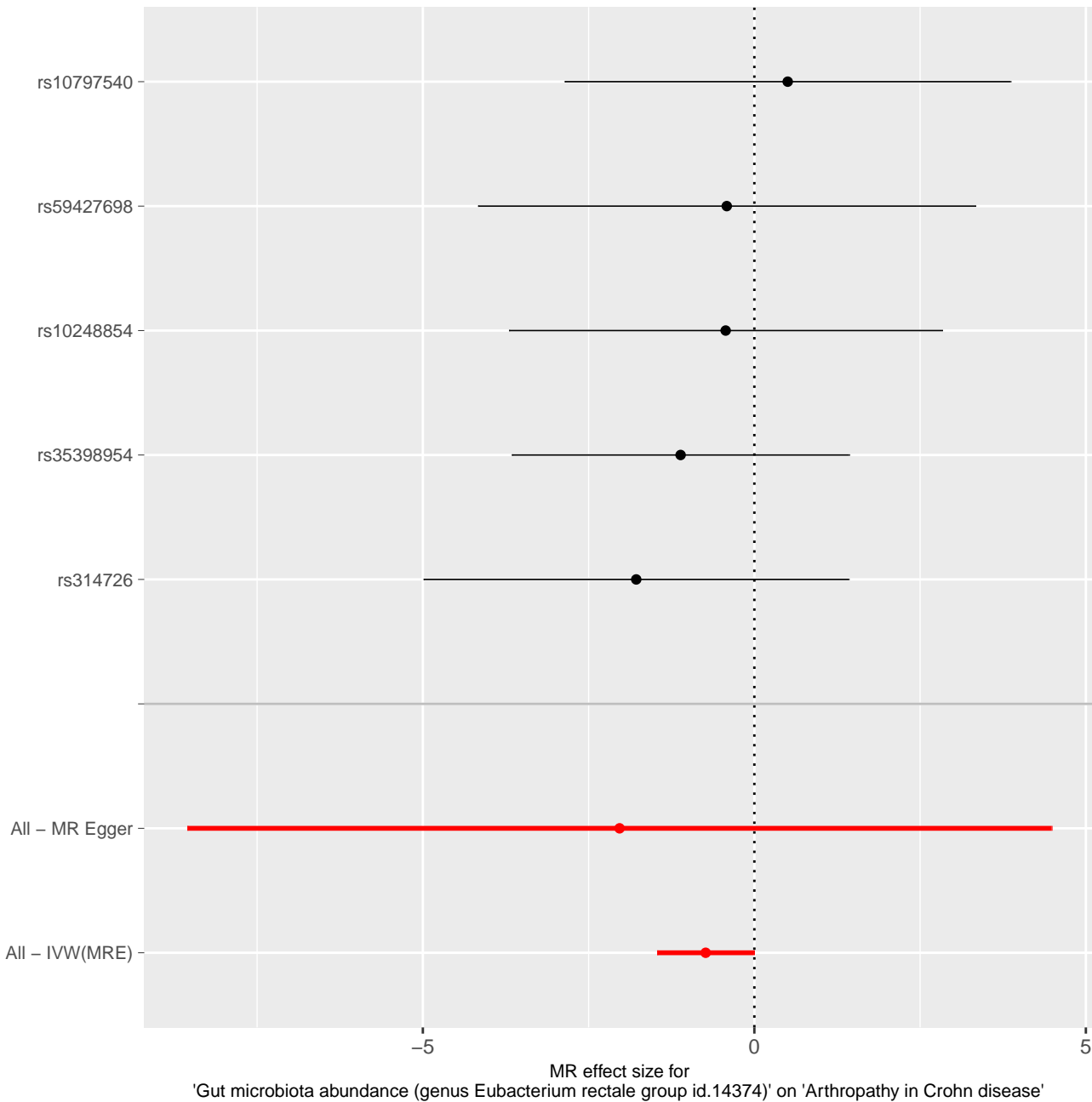






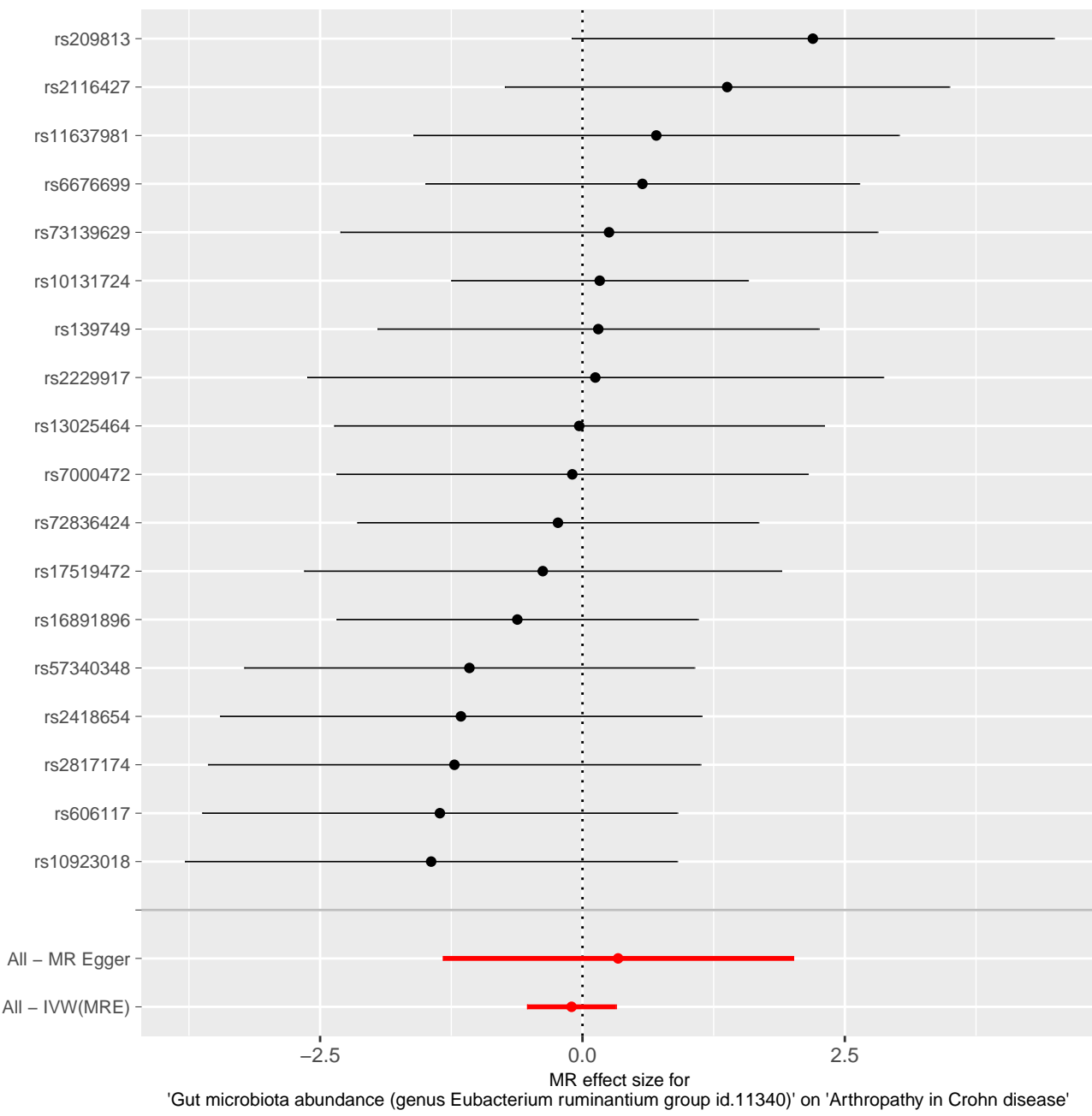




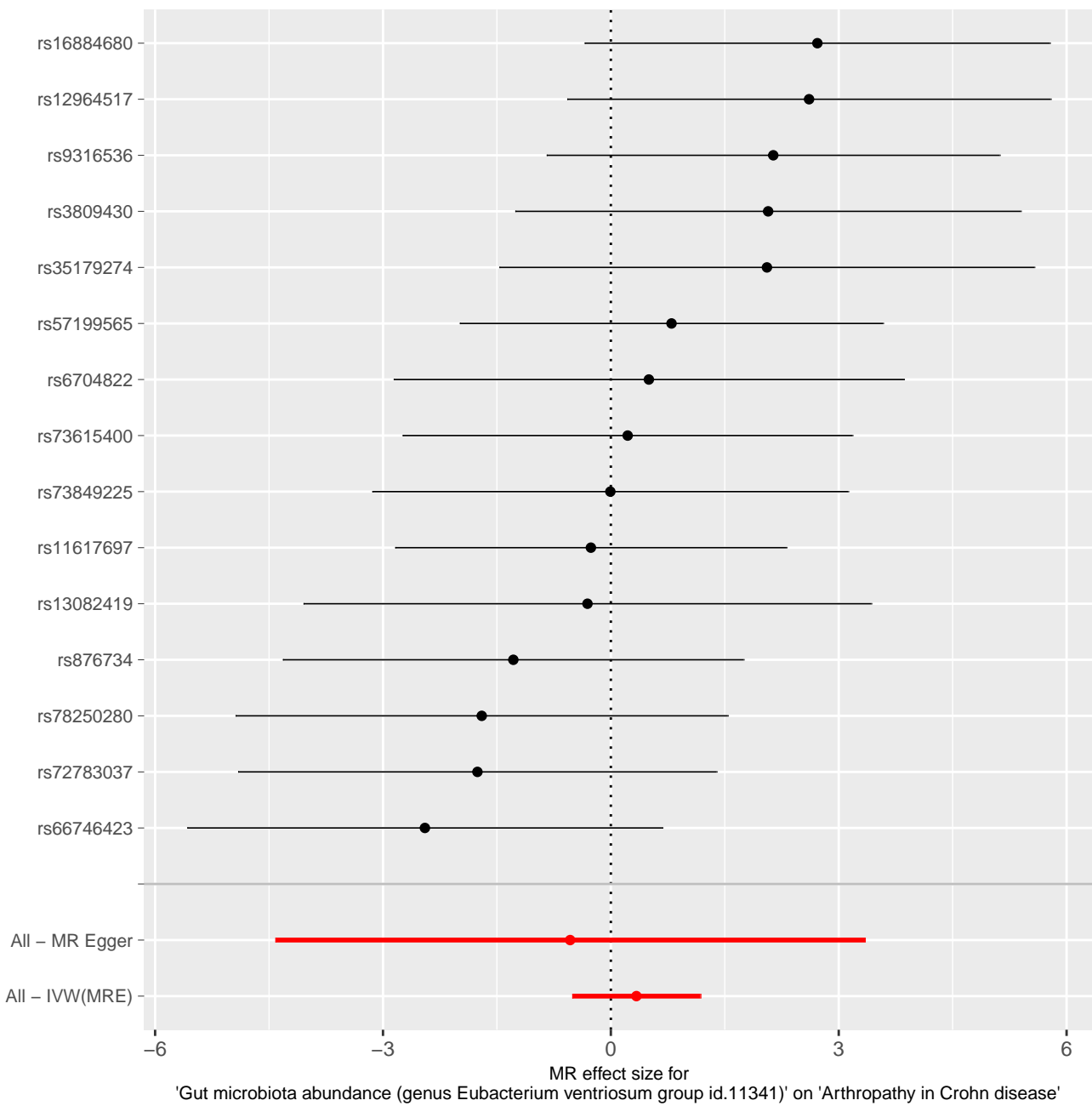


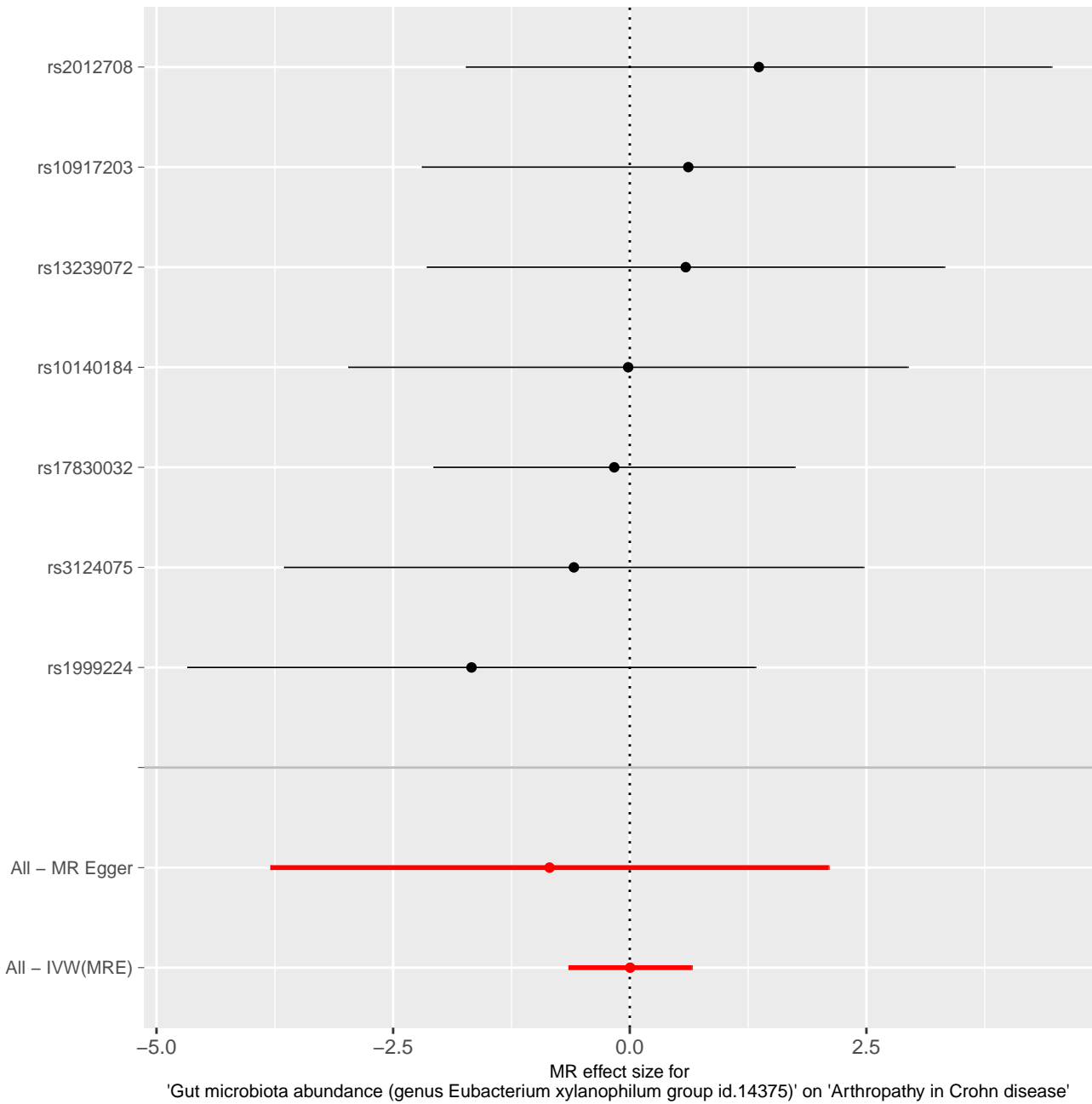
MR effect size for
'Gut microbiota abundance (genus Eubacterium rectale group id.14374)' on 'Arthropathy in Crohn disease'

Batch 528 : Gut microbiota abundance (genus Eubacterium ruminantium group id.11340) on Arthropathy in Crohn disease

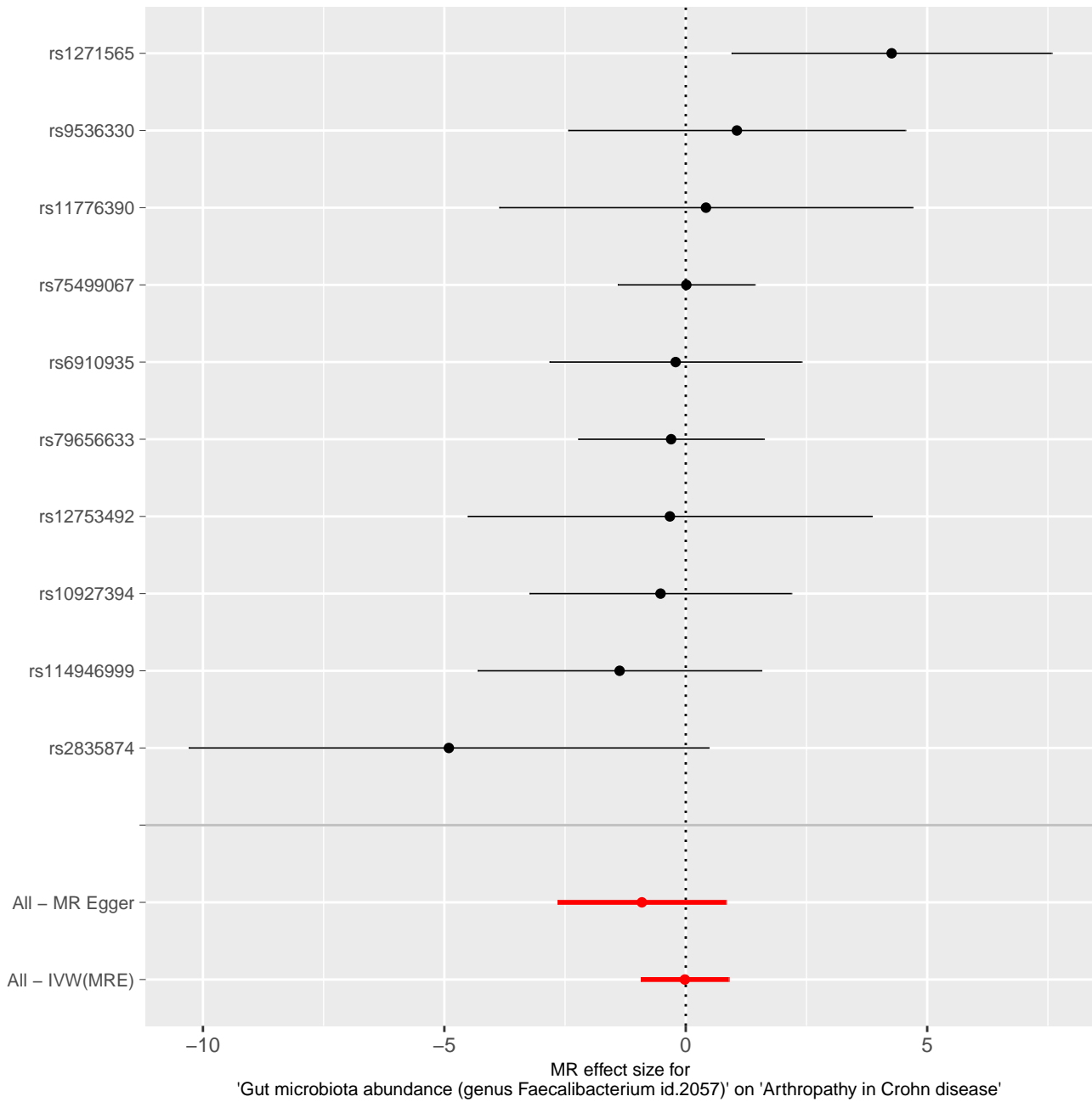


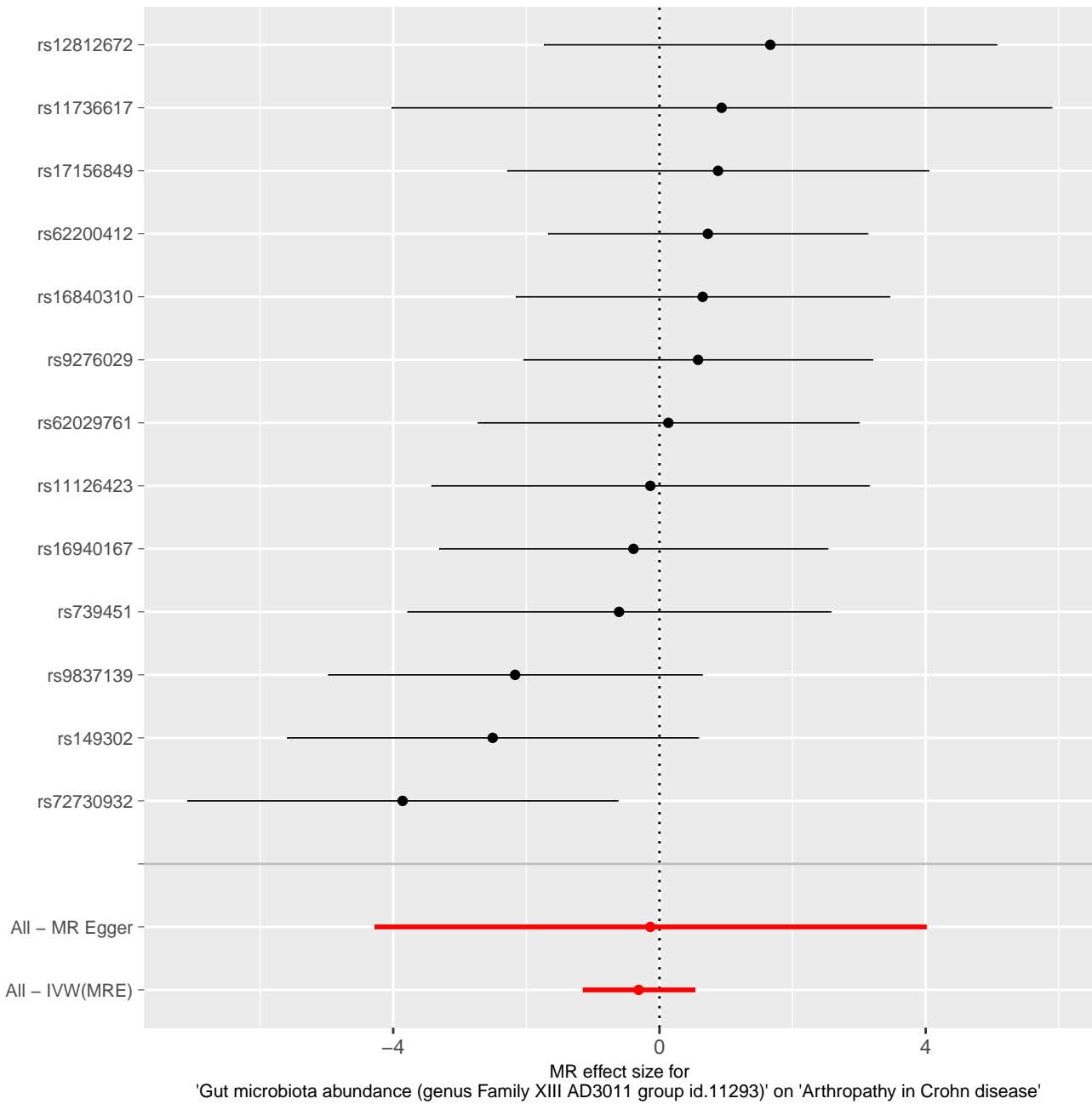
'Gut microbiota abundance (genus Eubacterium ruminantium group id.11340)' on 'Arthropathy in Crohn disease'

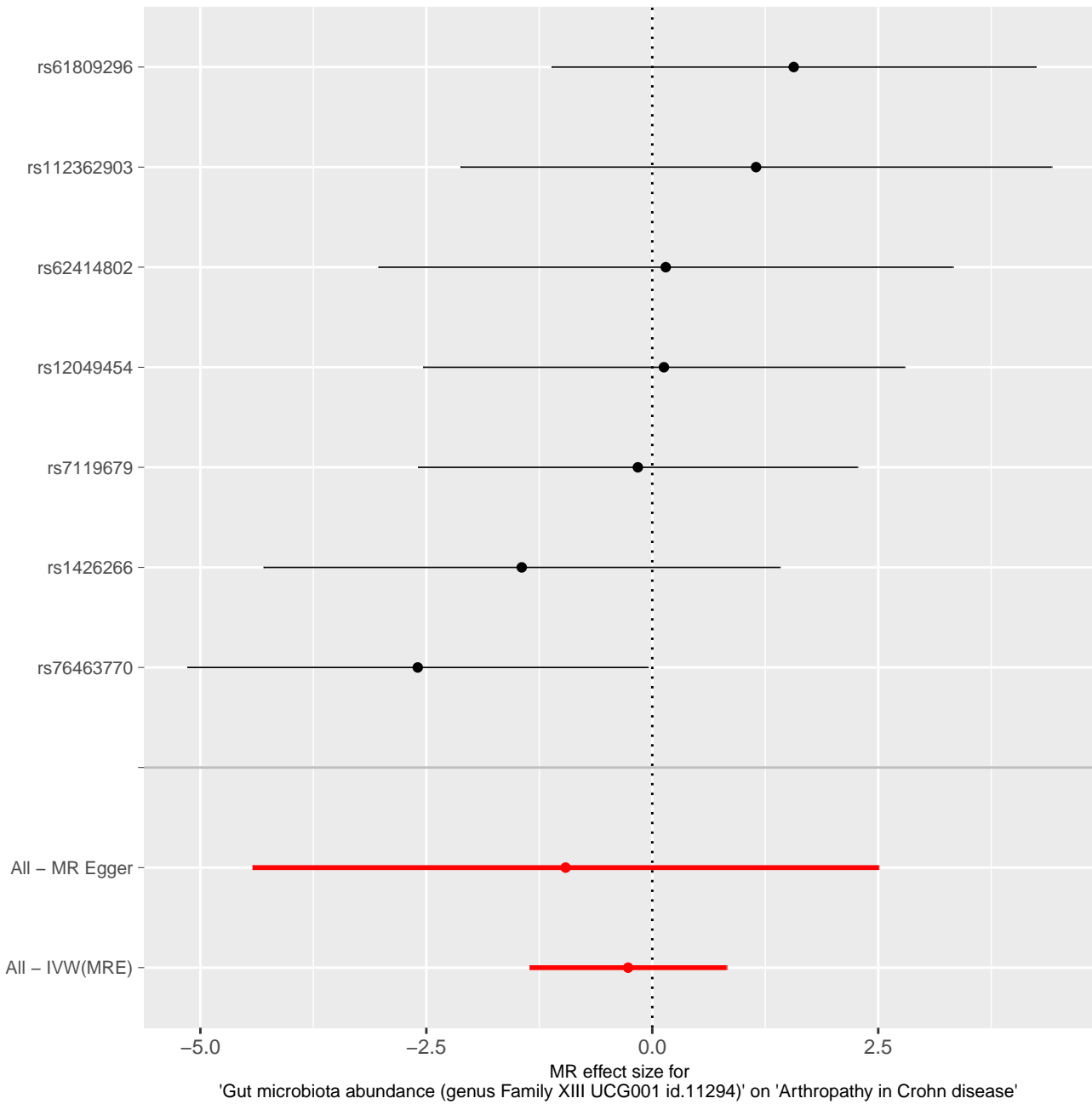




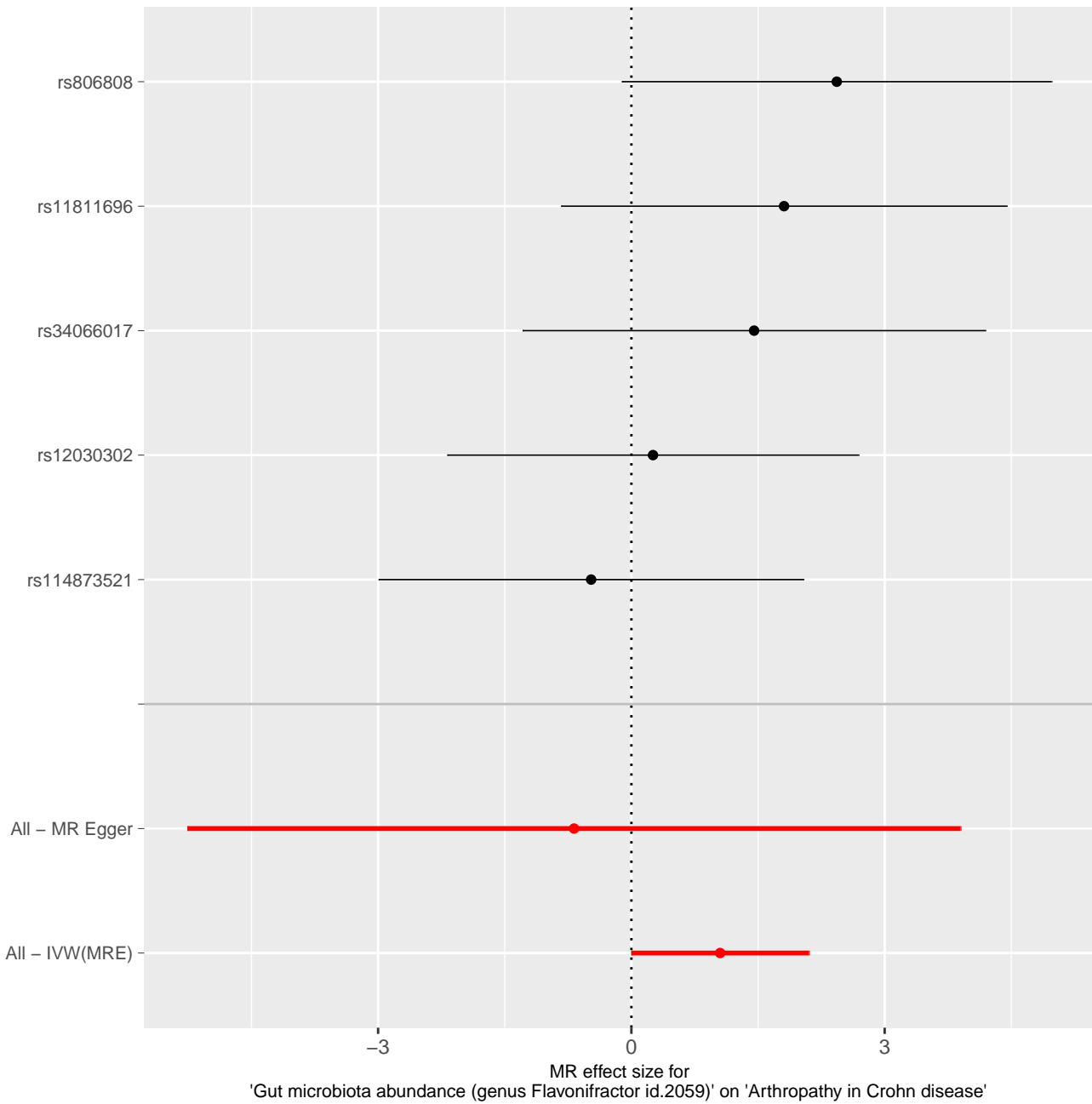
MR effect size for
'Gut microbiota abundance (genus Eubacterium xylanophilum group id.14375)' on 'Arthropathy in Crohn disease'



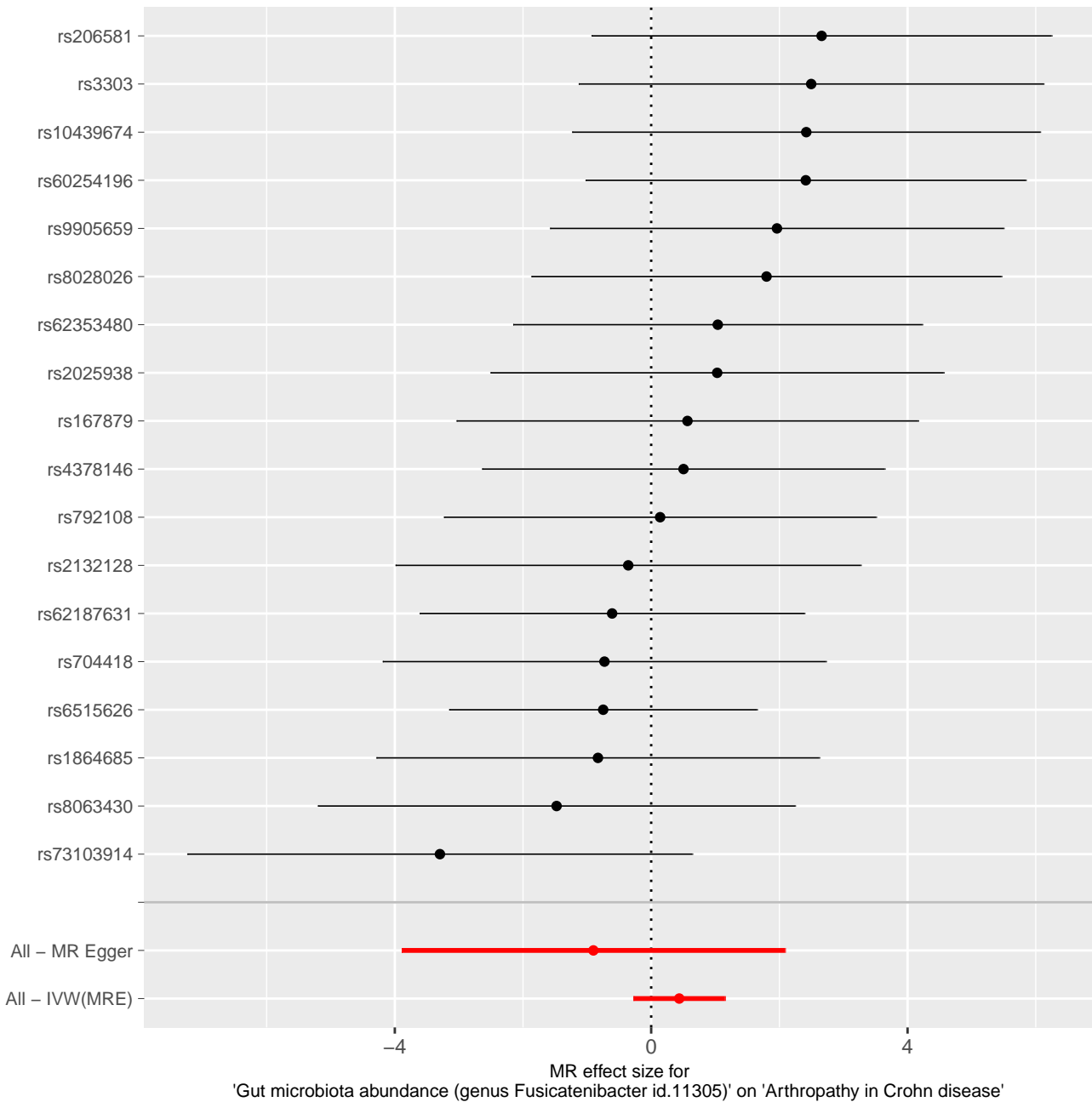


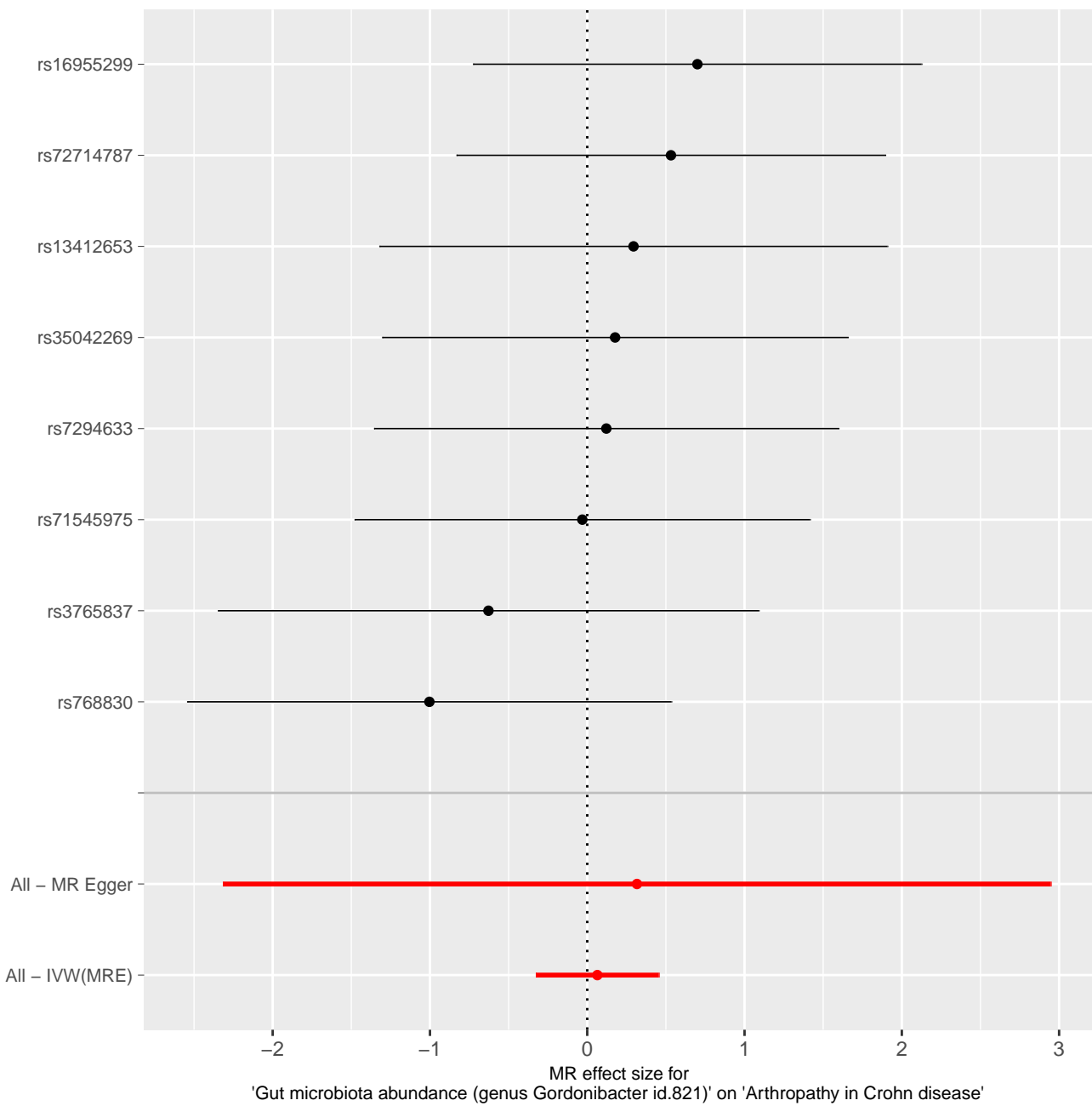


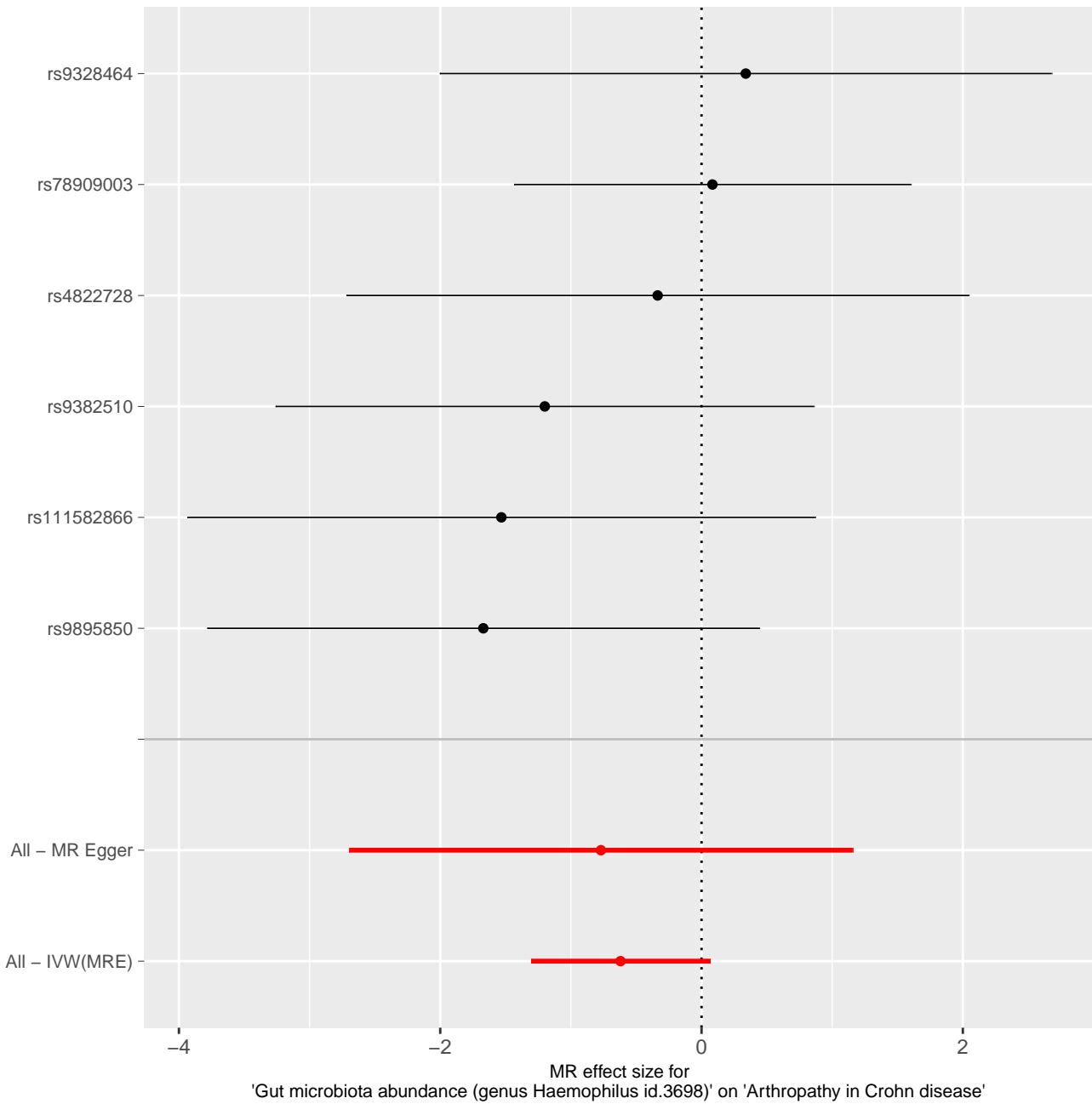
MR effect size for
'Gut microbiota abundance (genus Family XIII UCG001 id.11294)' on 'Arthropathy in Crohn disease'



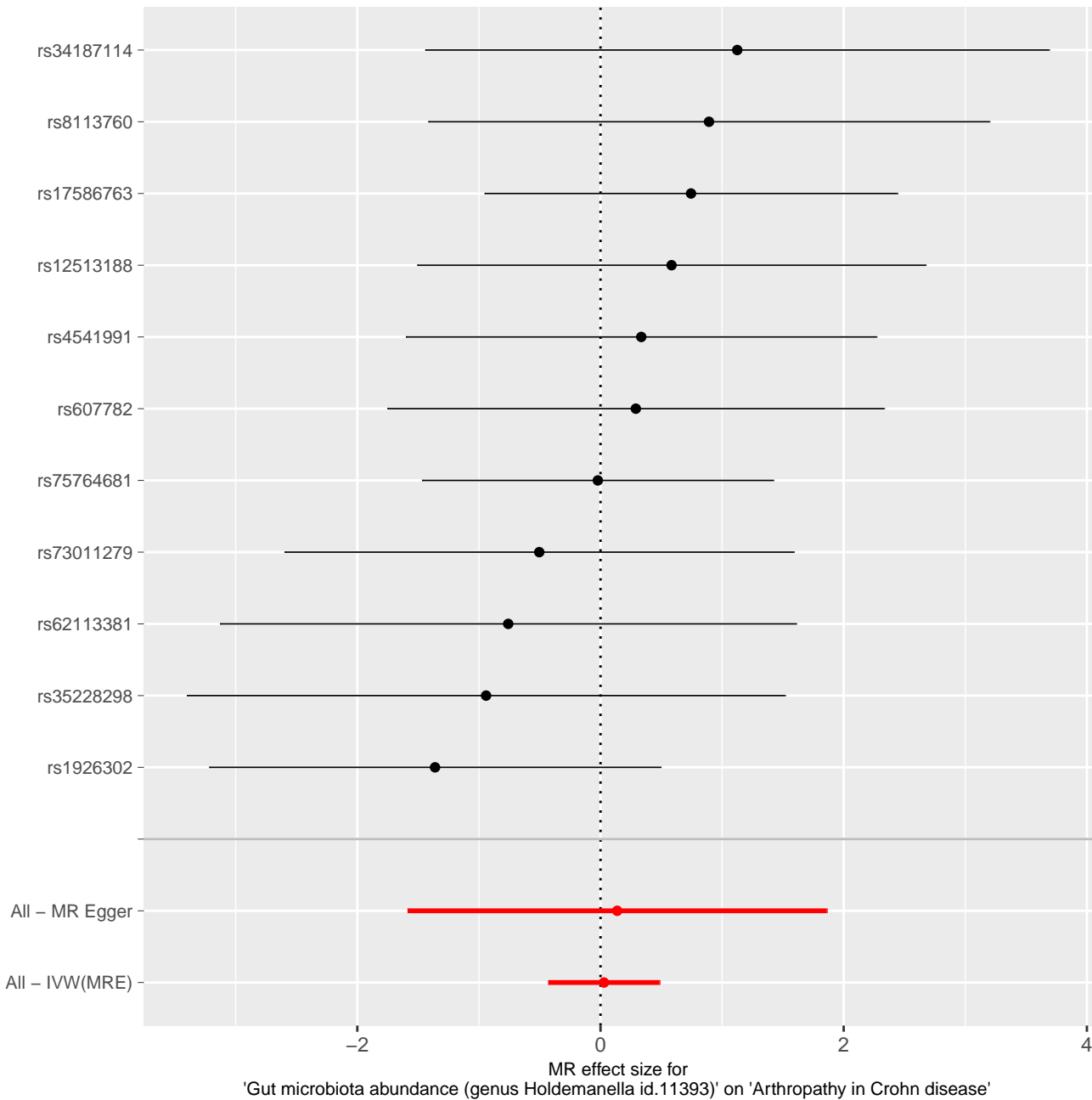
Batch 535 : Gut microbiota abundance (genus Fusicatenibacter id.11305) on Arthropathy in Crohn disease



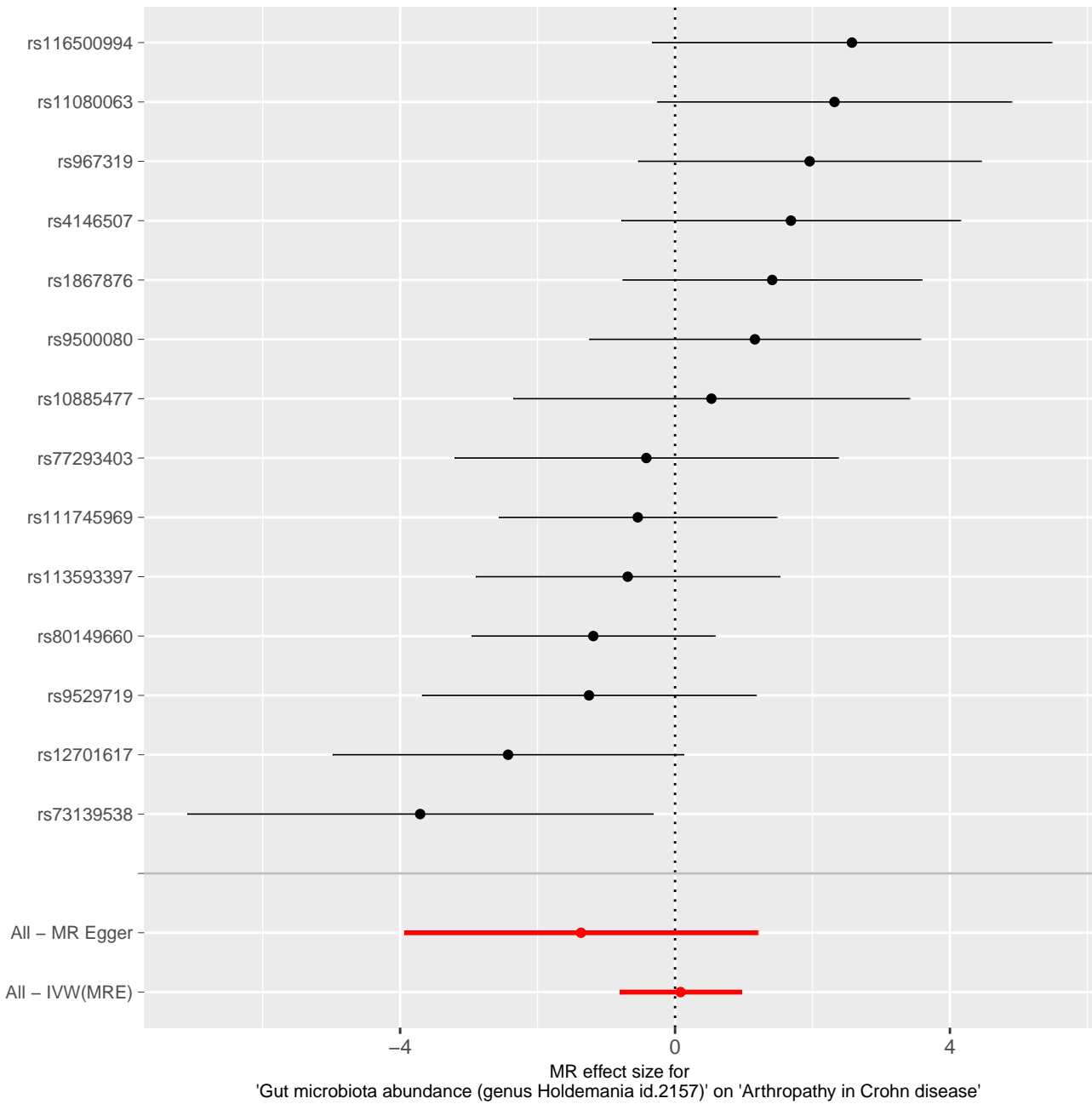


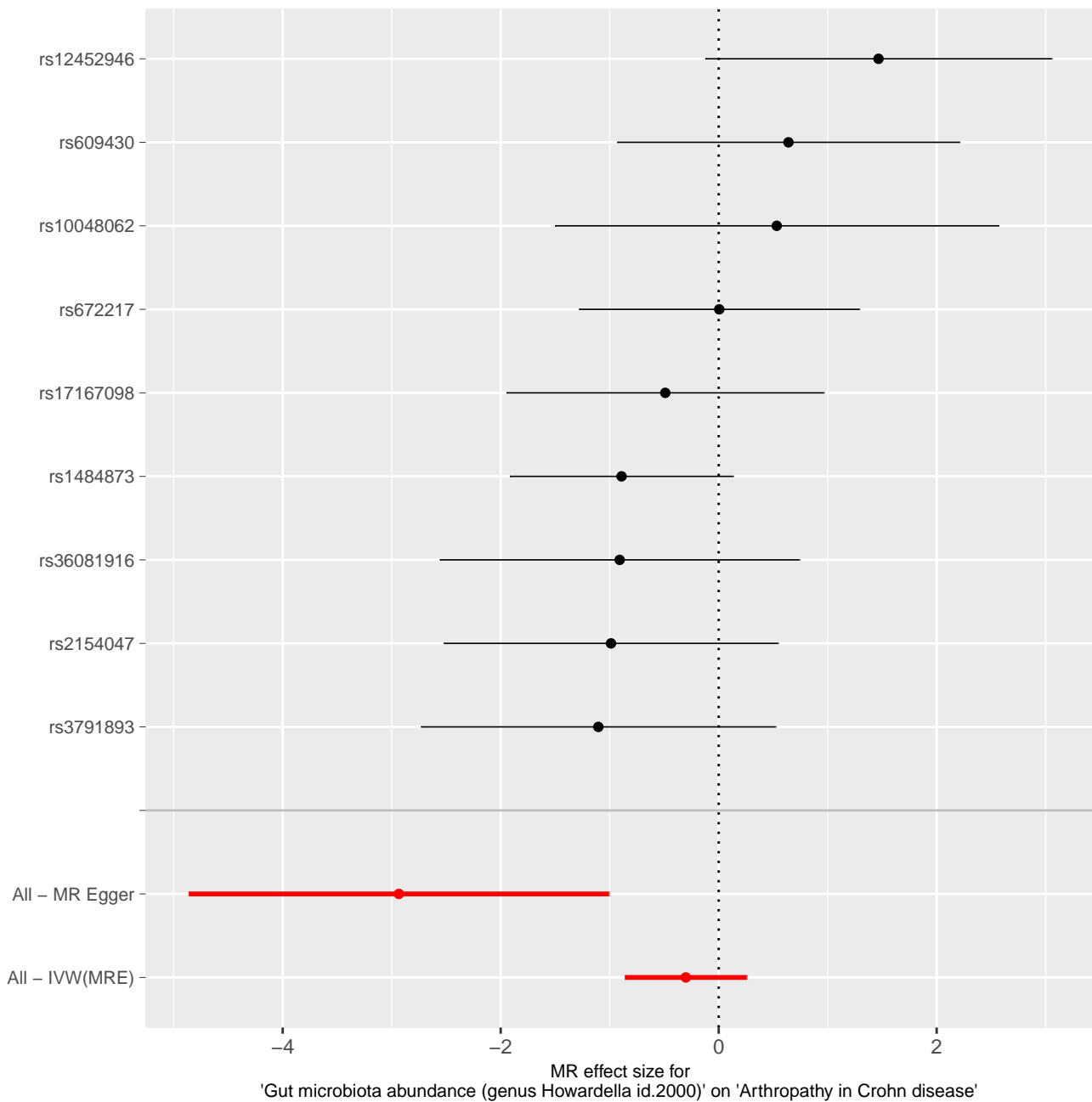


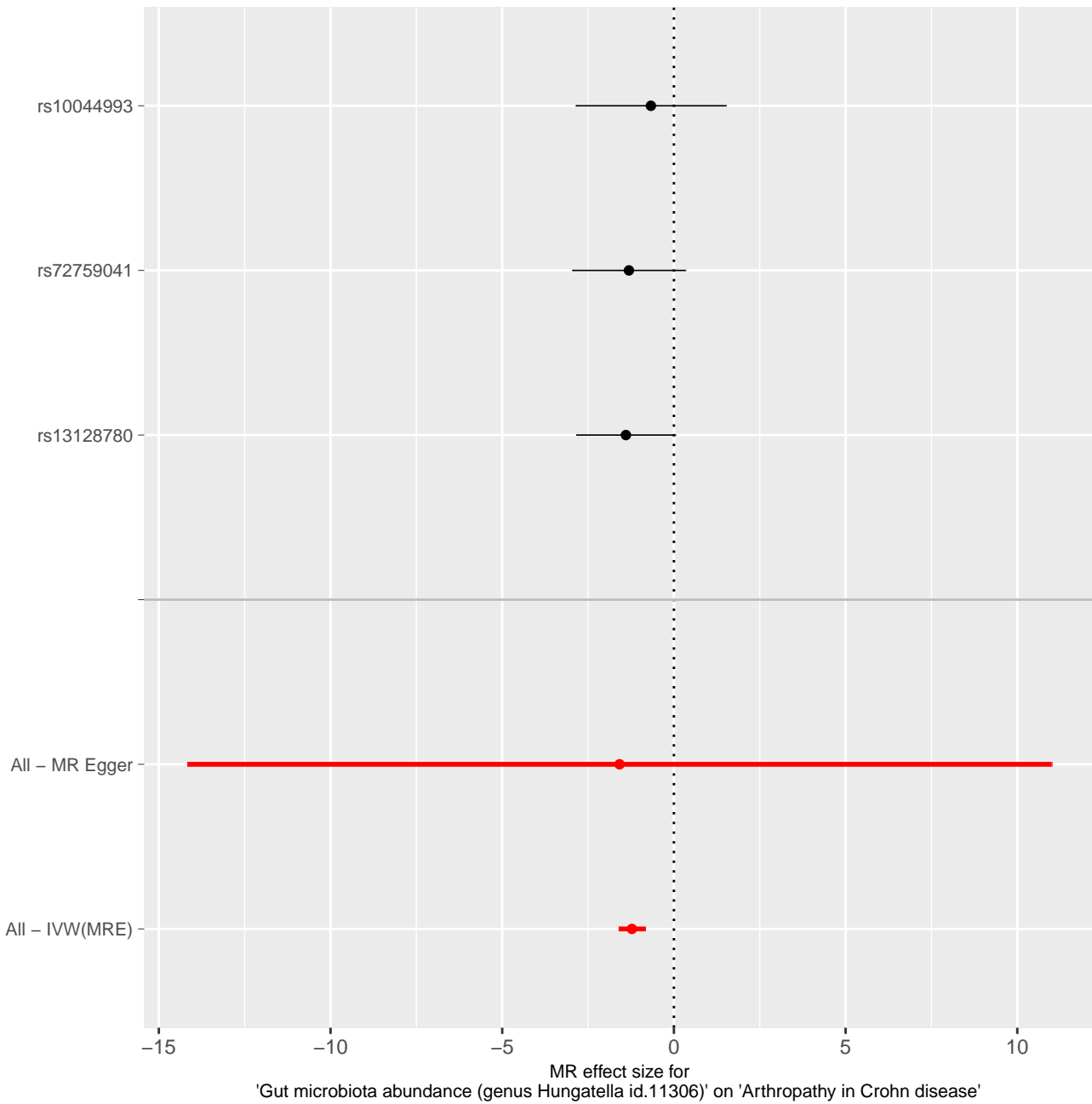
Batch 538 : Gut microbiota abundance (genus Holdemanella id.11393) on Arthropathy in Crohn disease



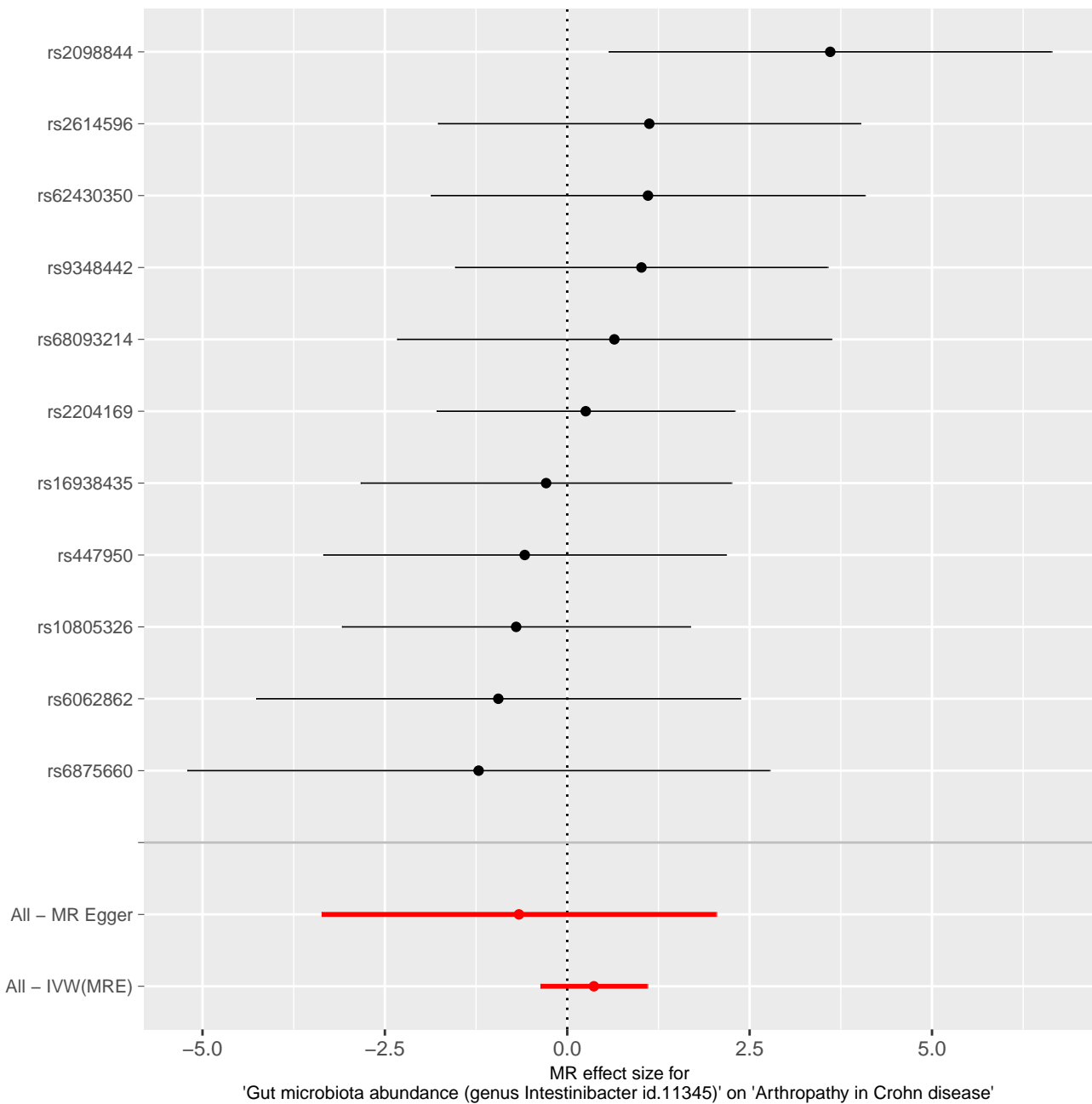
Batch 539 : Gut microbiota abundance (genus Holdemania id.2157) on Arthropathy in Crohn disease



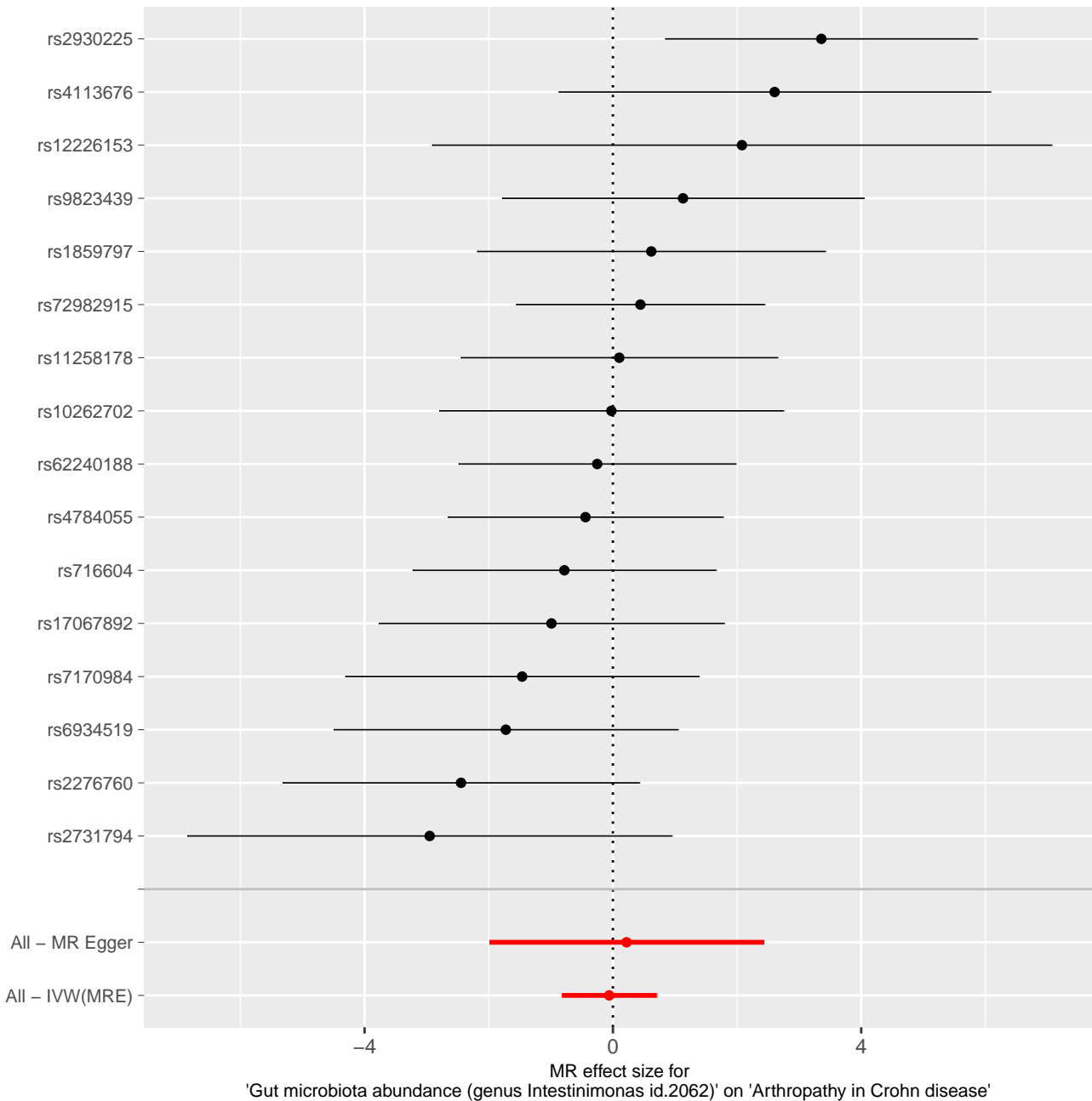




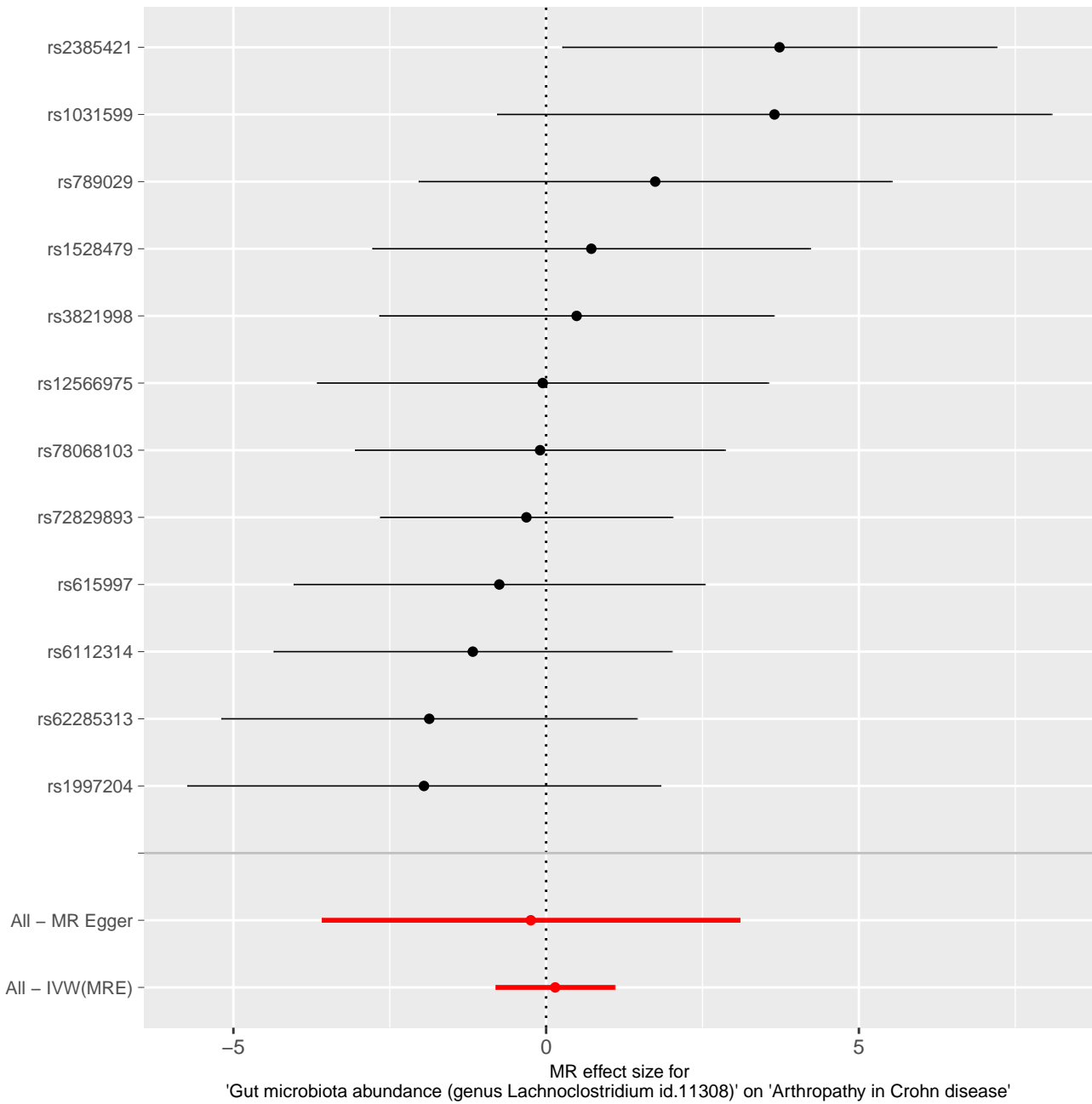
Batch 542 : Gut microbiota abundance (genus Intestinibacter id.11345) on Arthropathy in Crohn disease

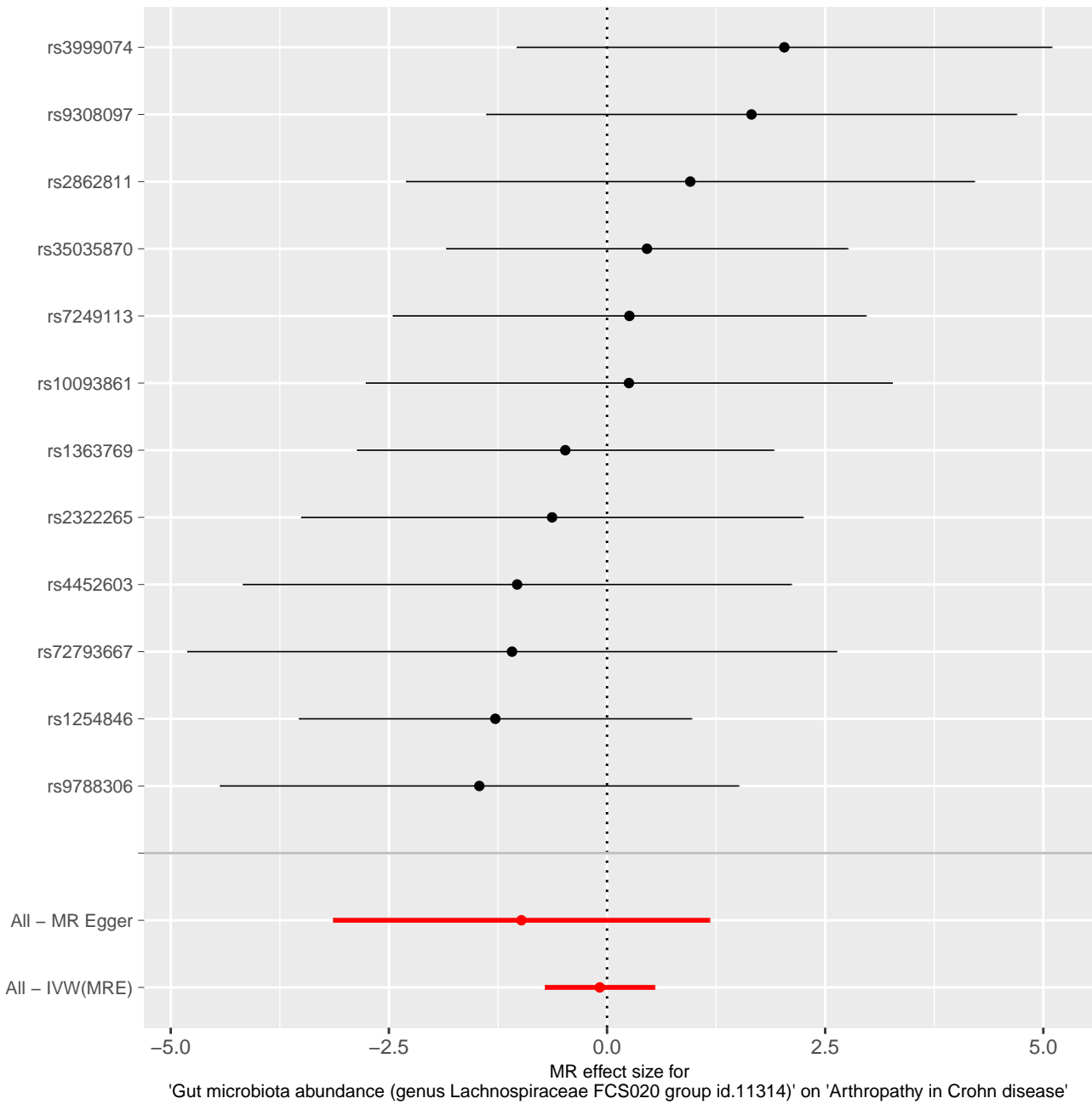


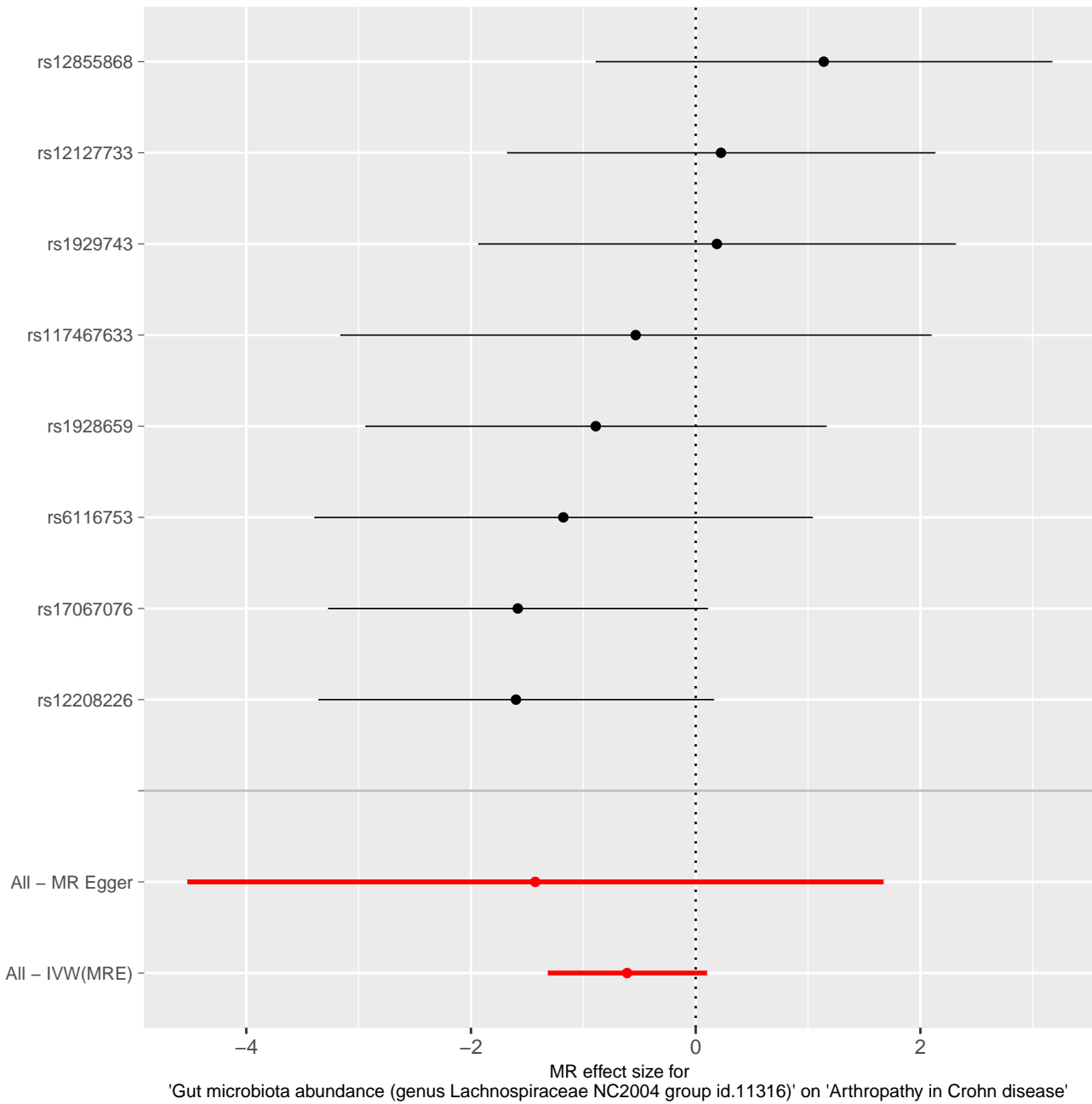
Batch 543 : Gut microbiota abundance (genus Intestinimonas id.2062) on Arthropathy in Crohn disease

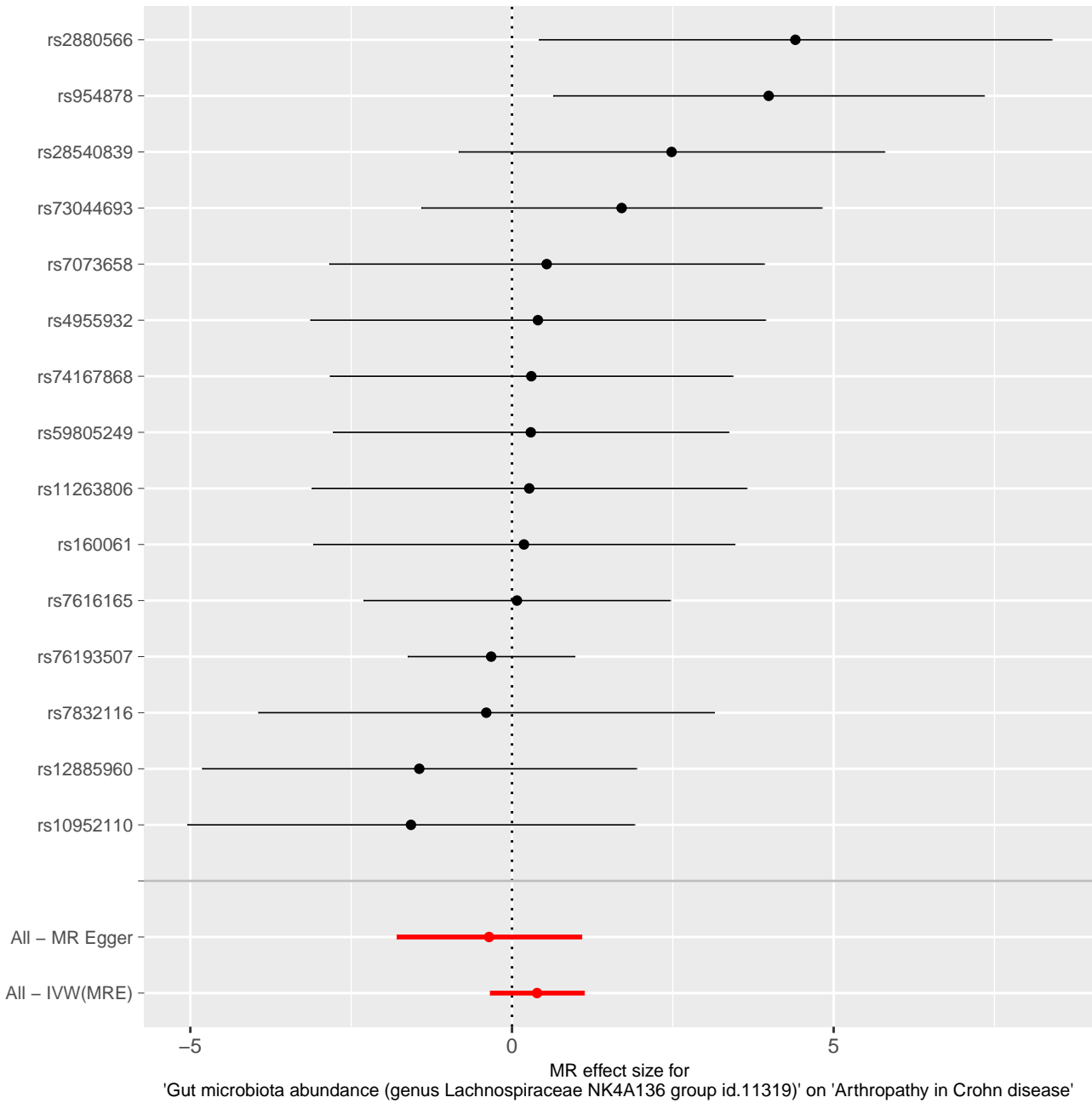


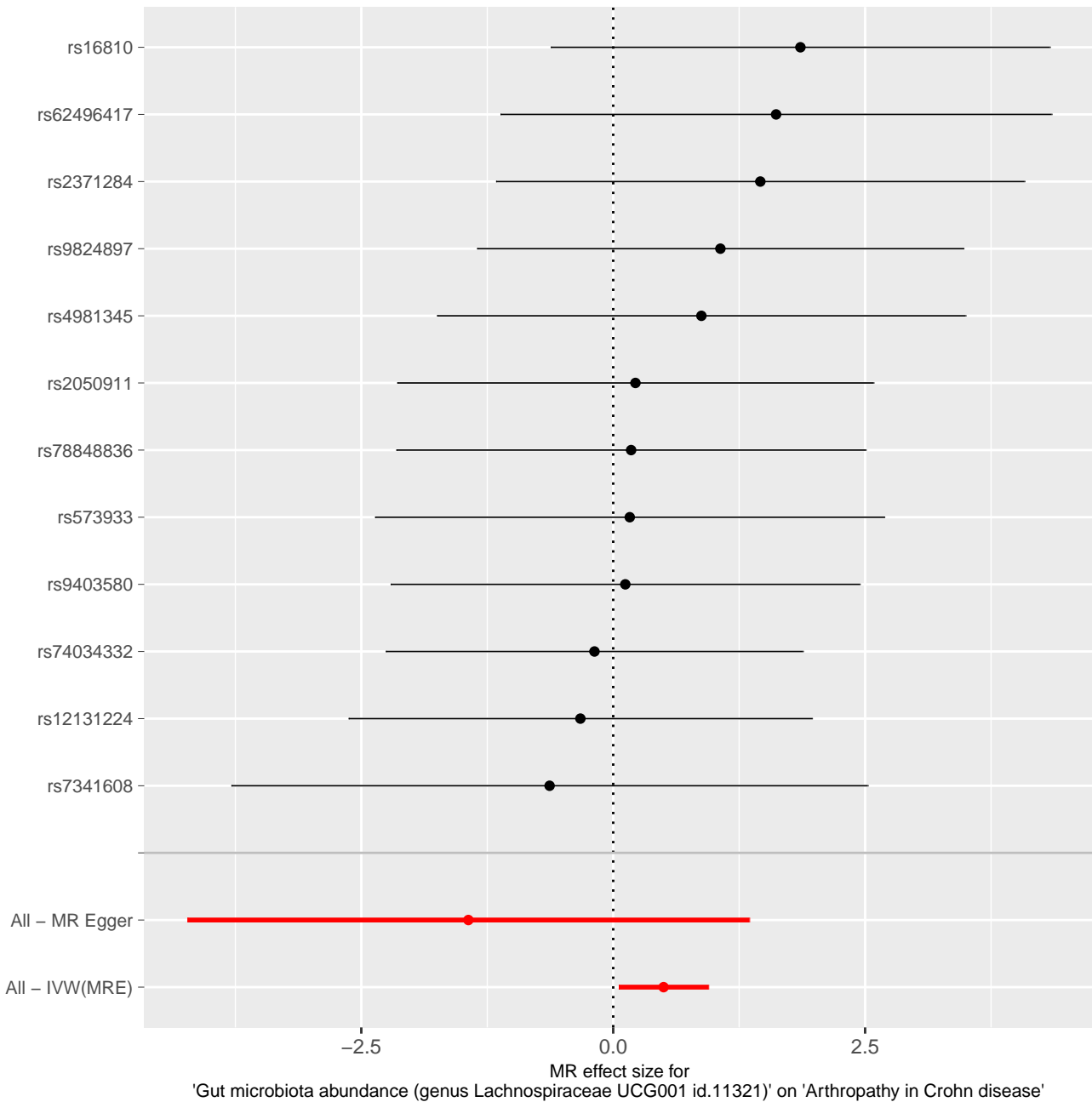
Batch 544 : Gut microbiota abundance (genus Lachnocostridium id.11308) on Arthropathy in Crohn disease

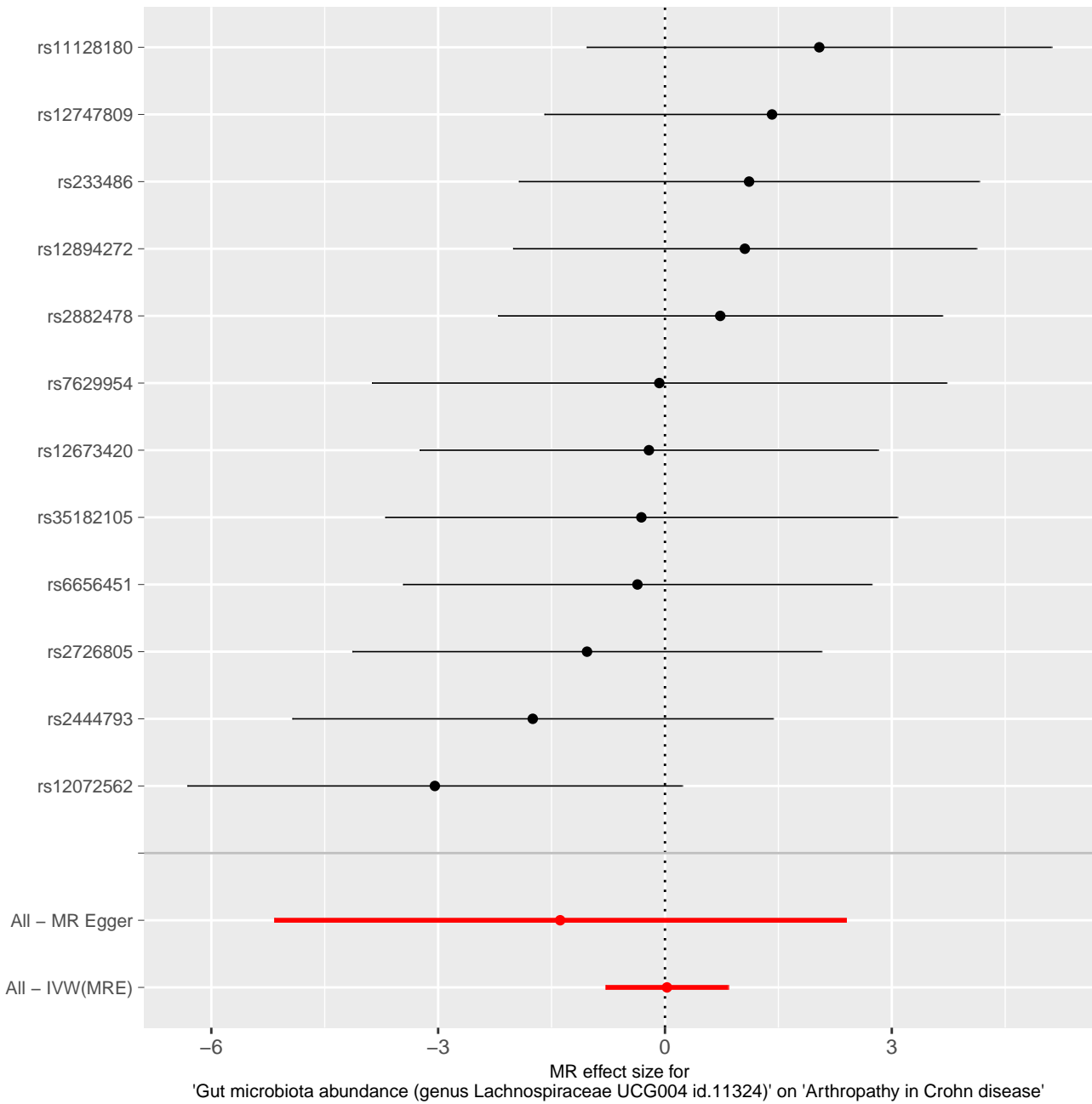


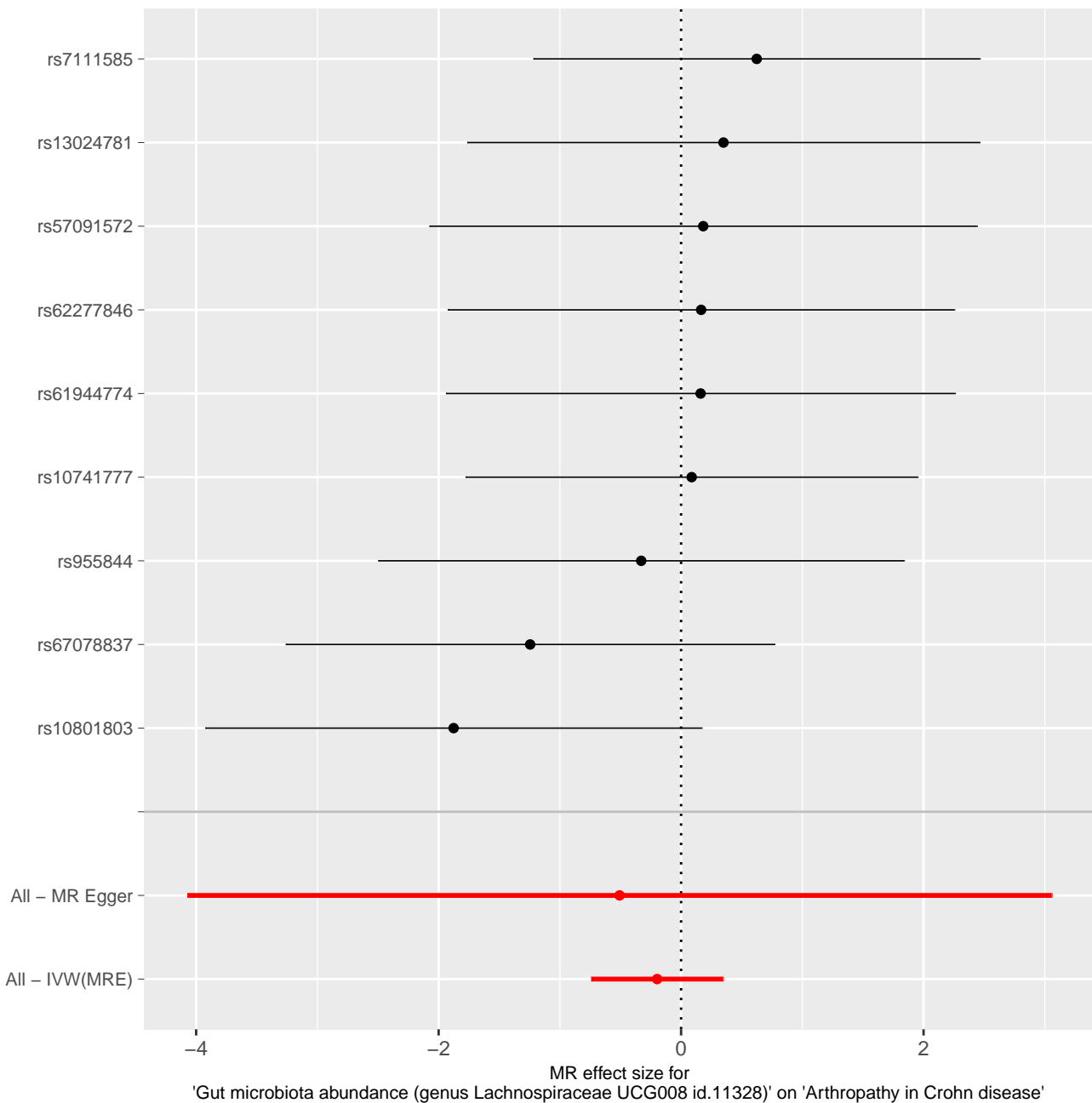


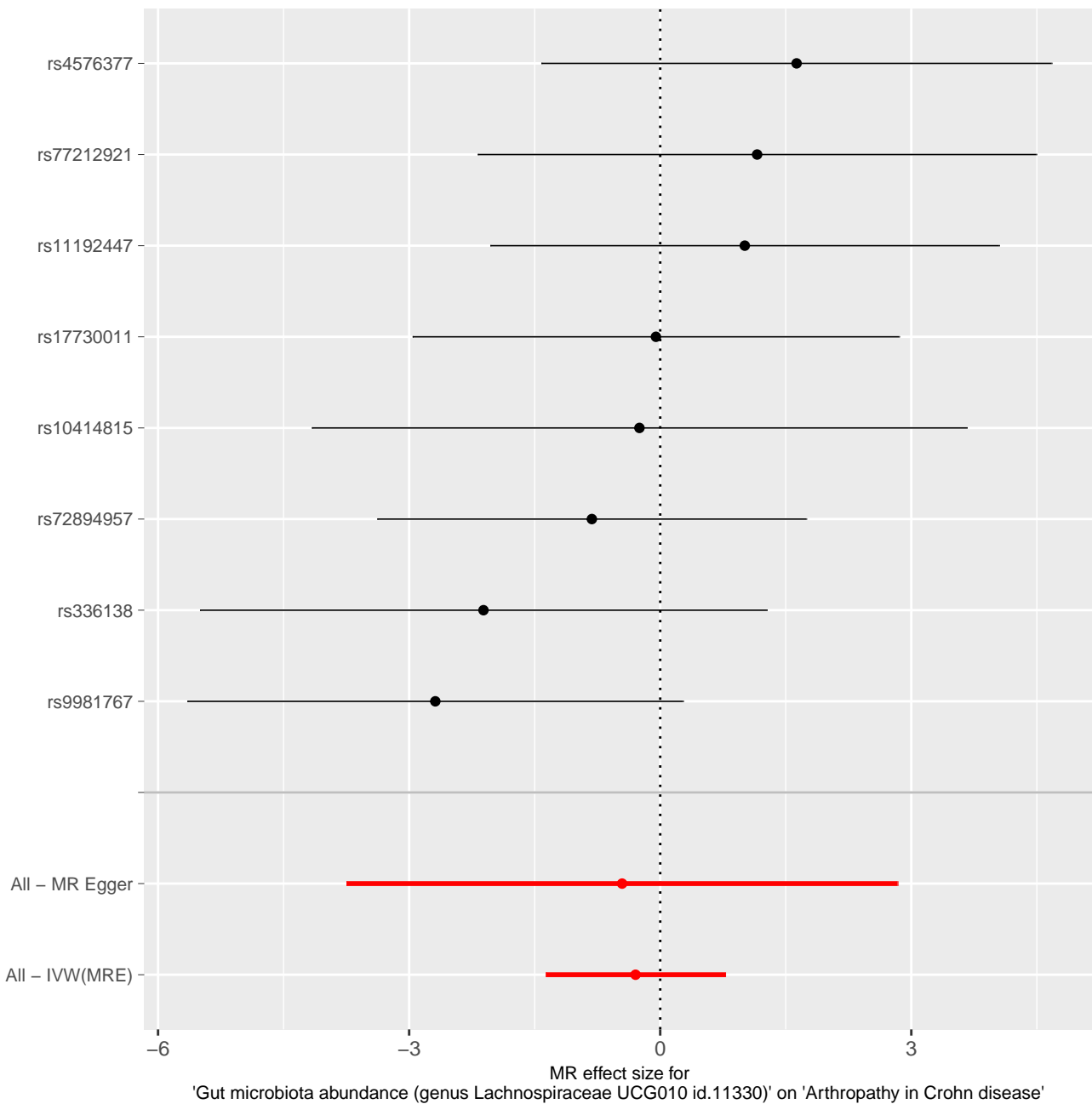




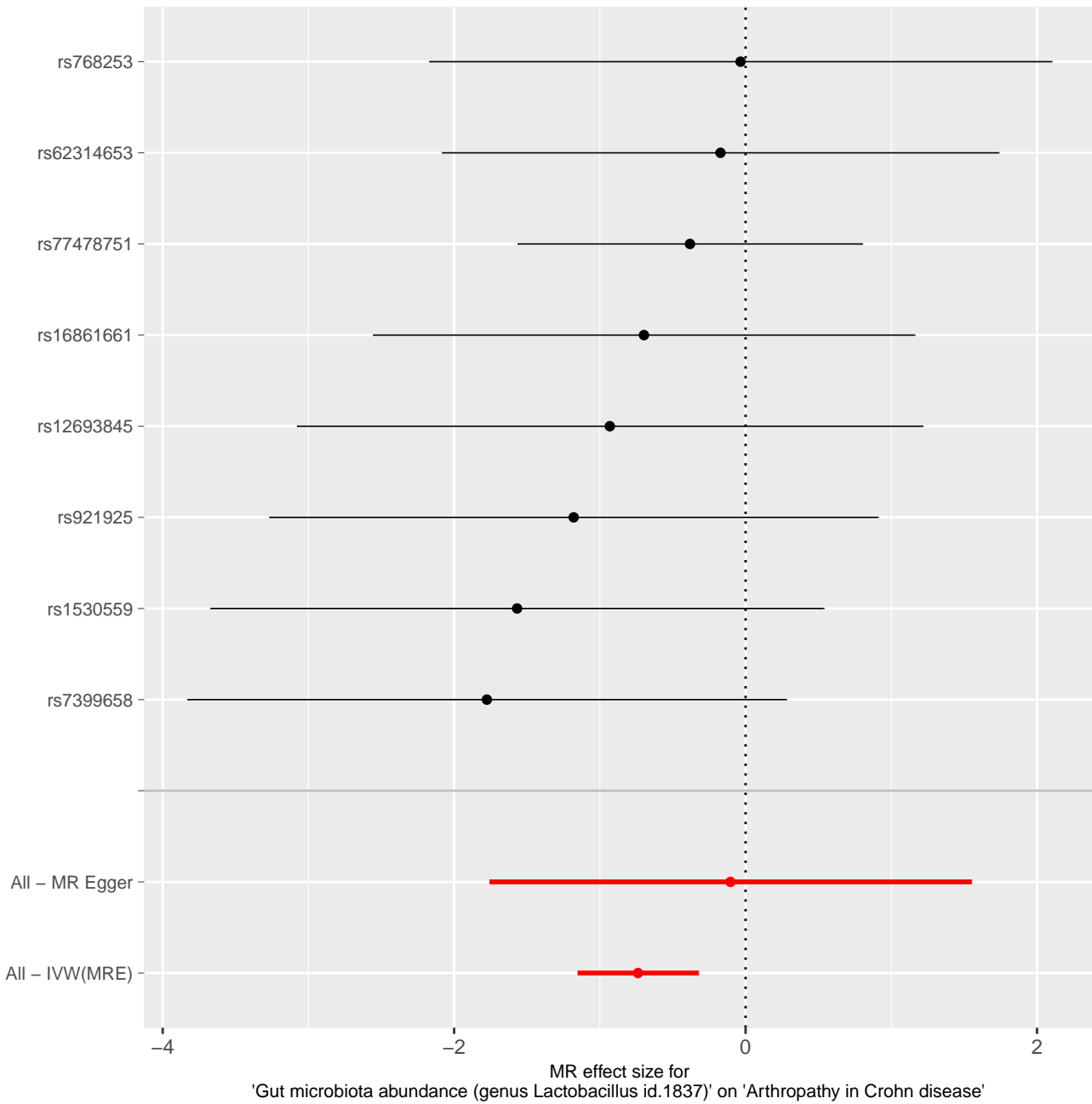


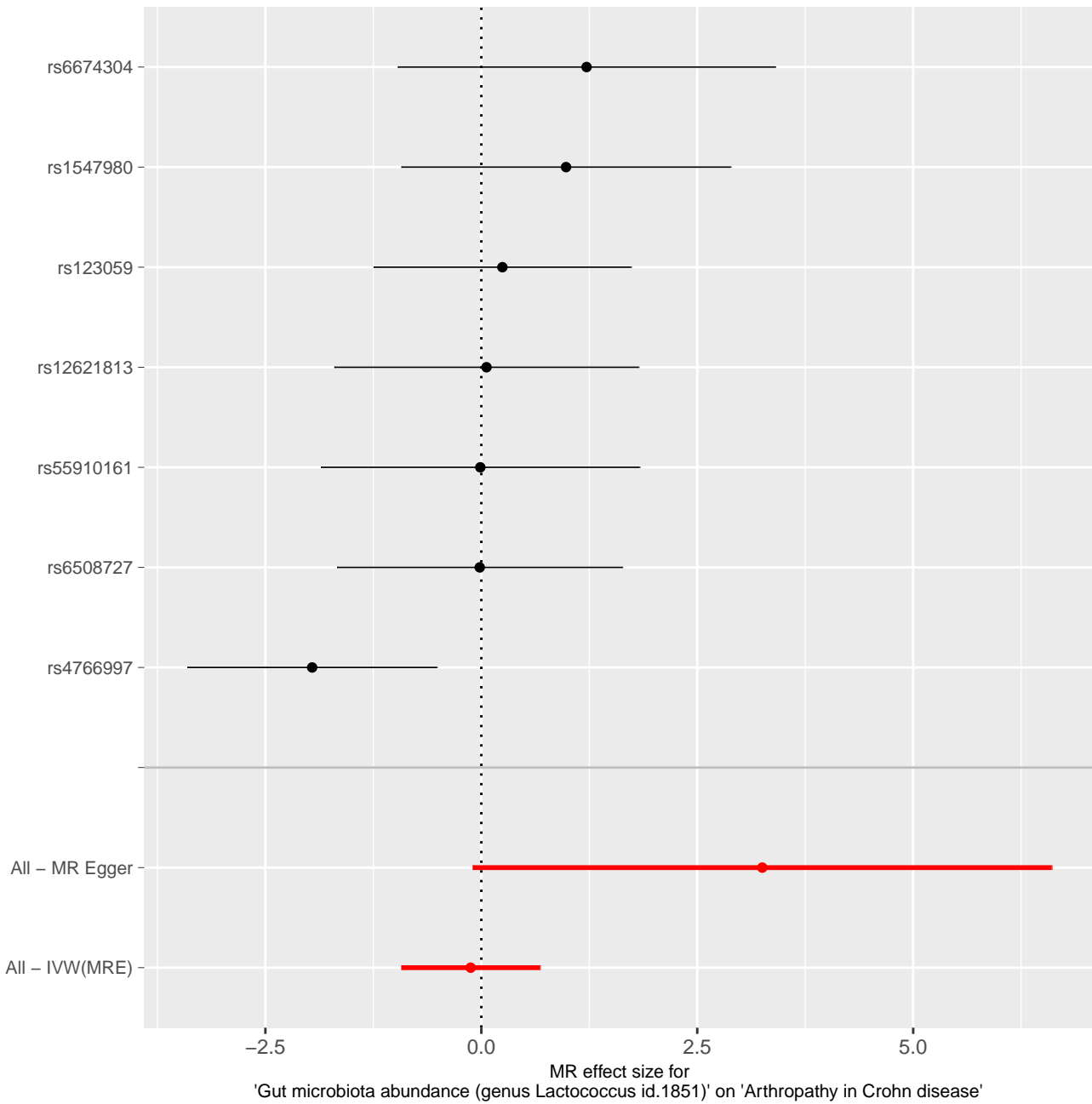


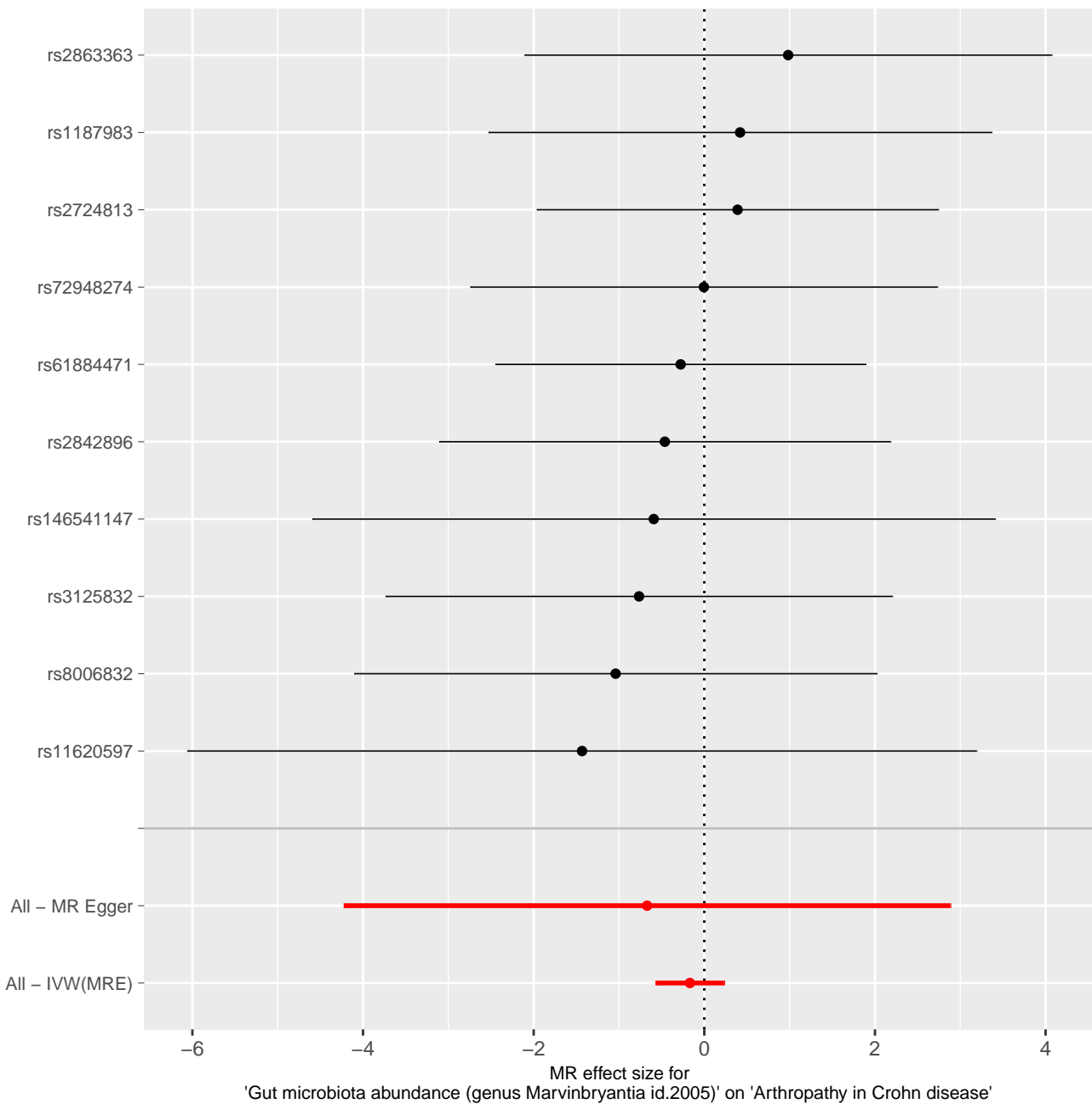


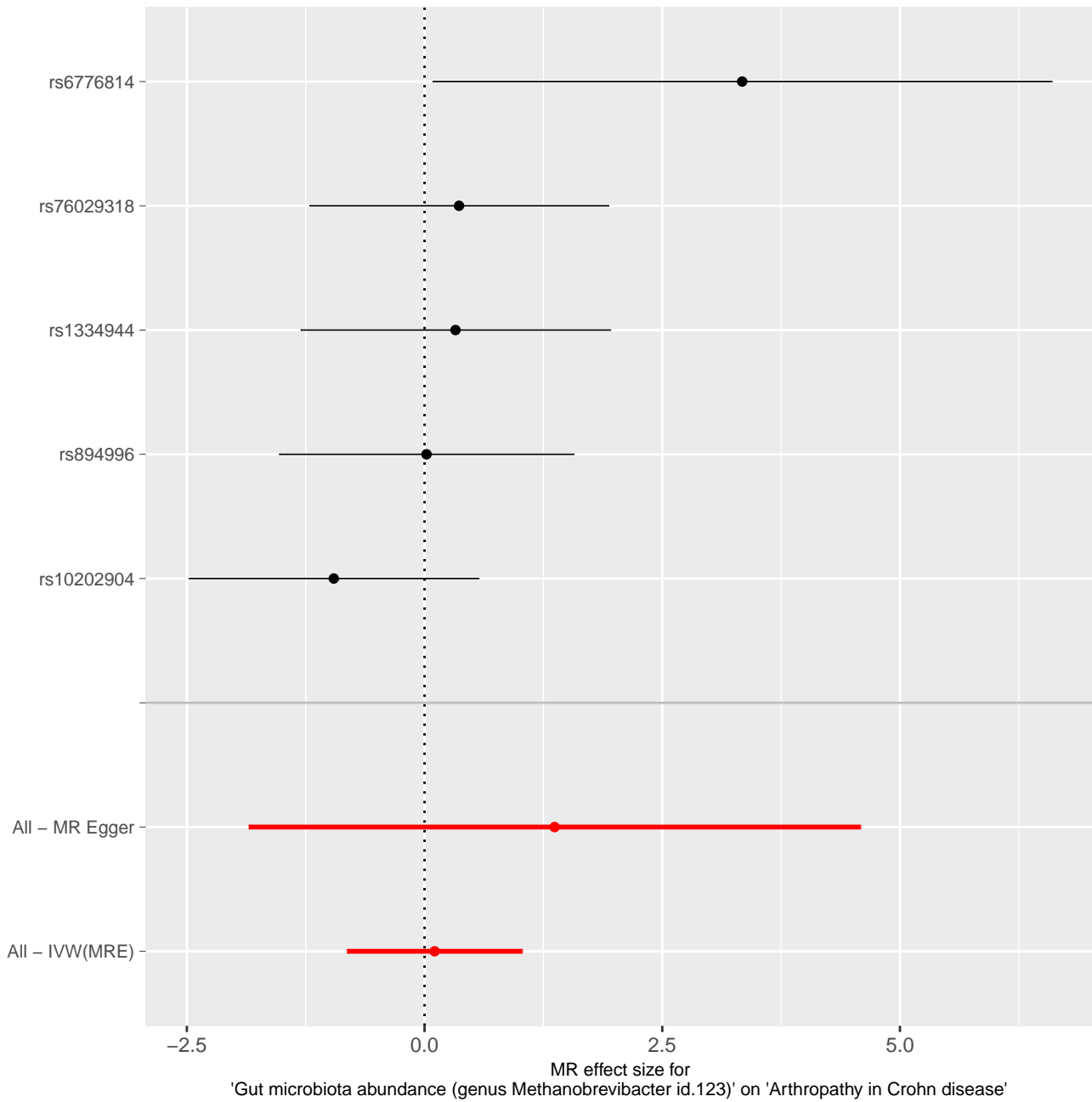


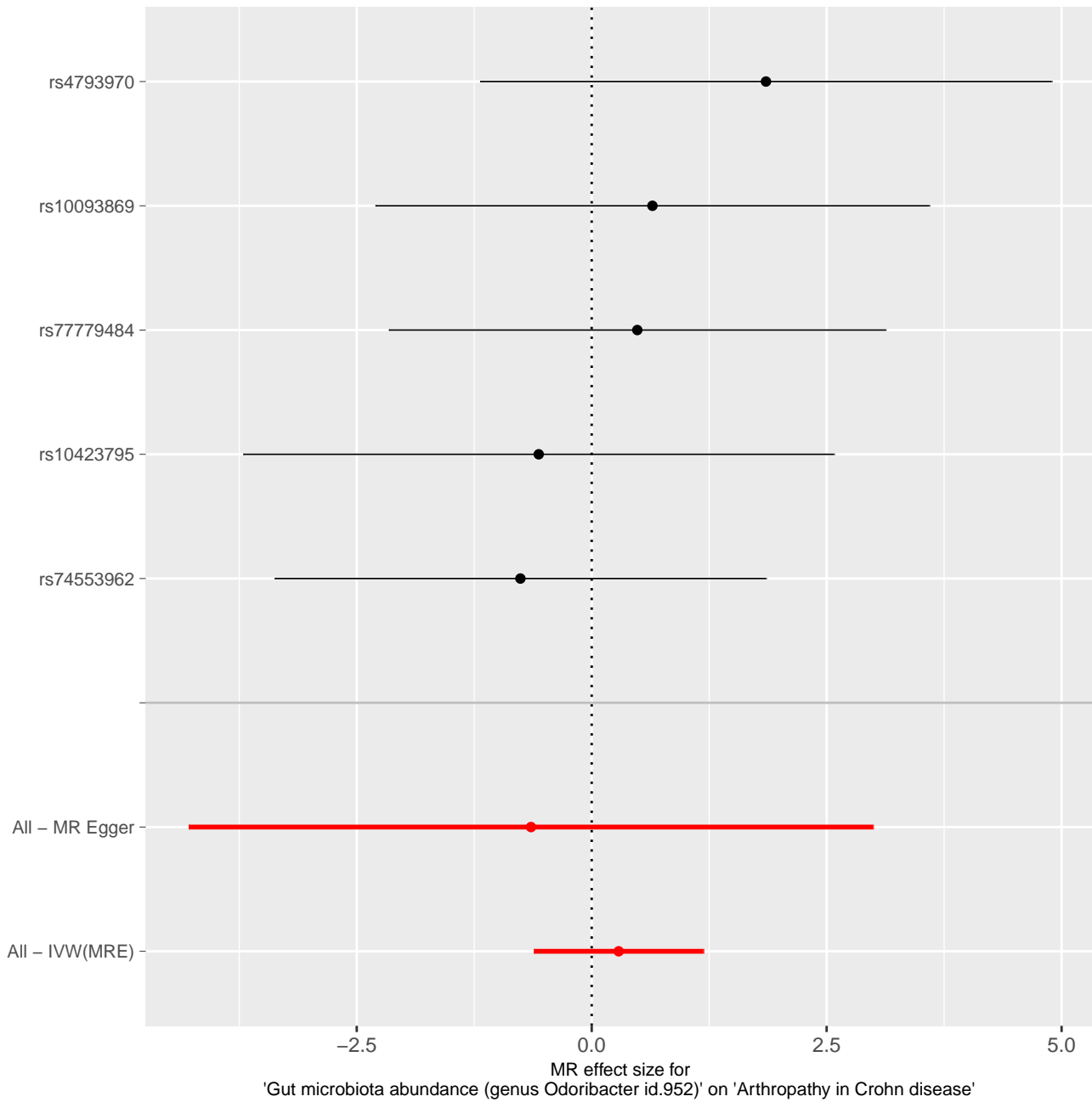
MR effect size for
'Gut microbiota abundance (genus Lachnospiraceae UCG010 id.11330)' on 'Arthropathy in Crohn disease'

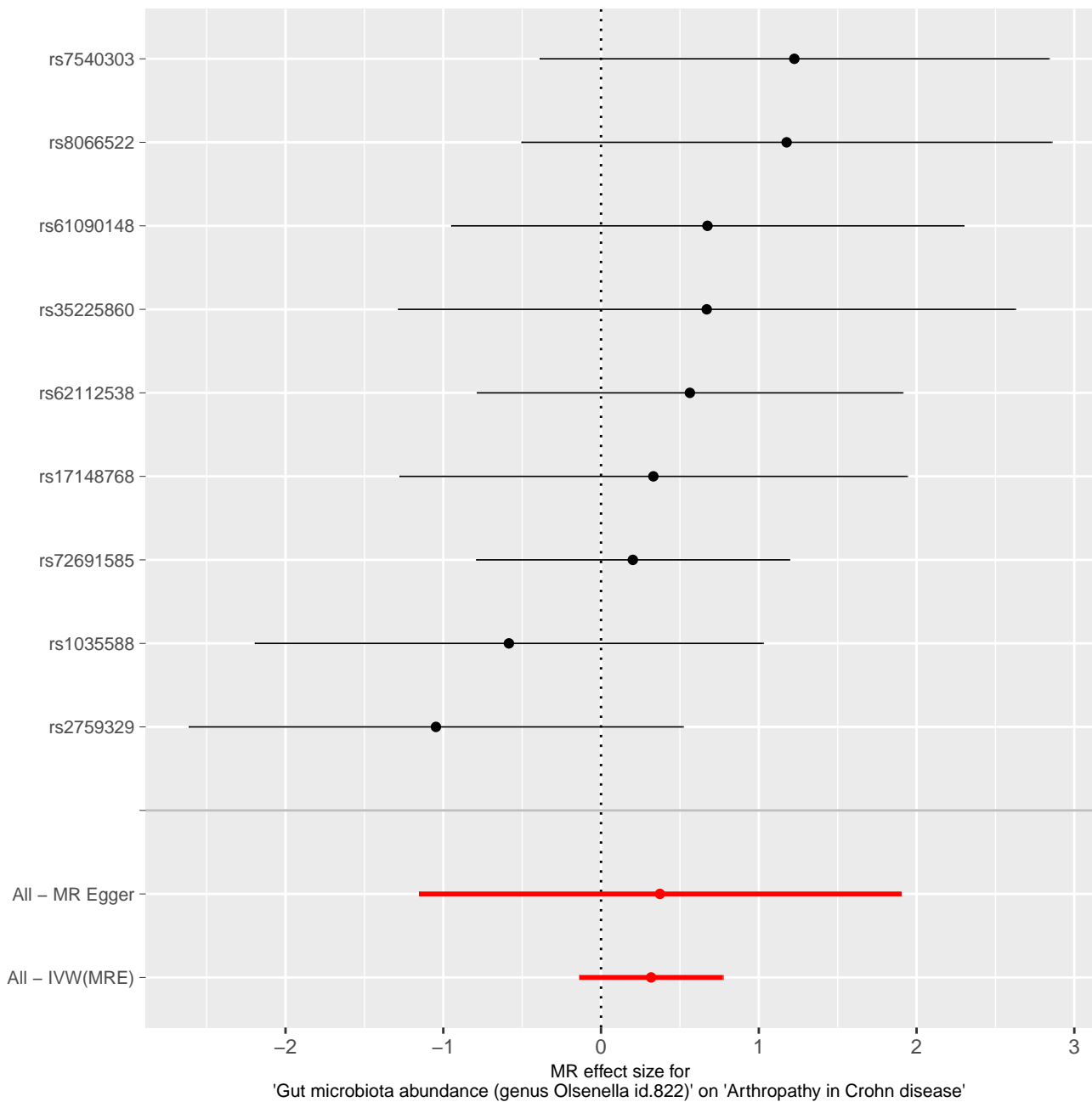


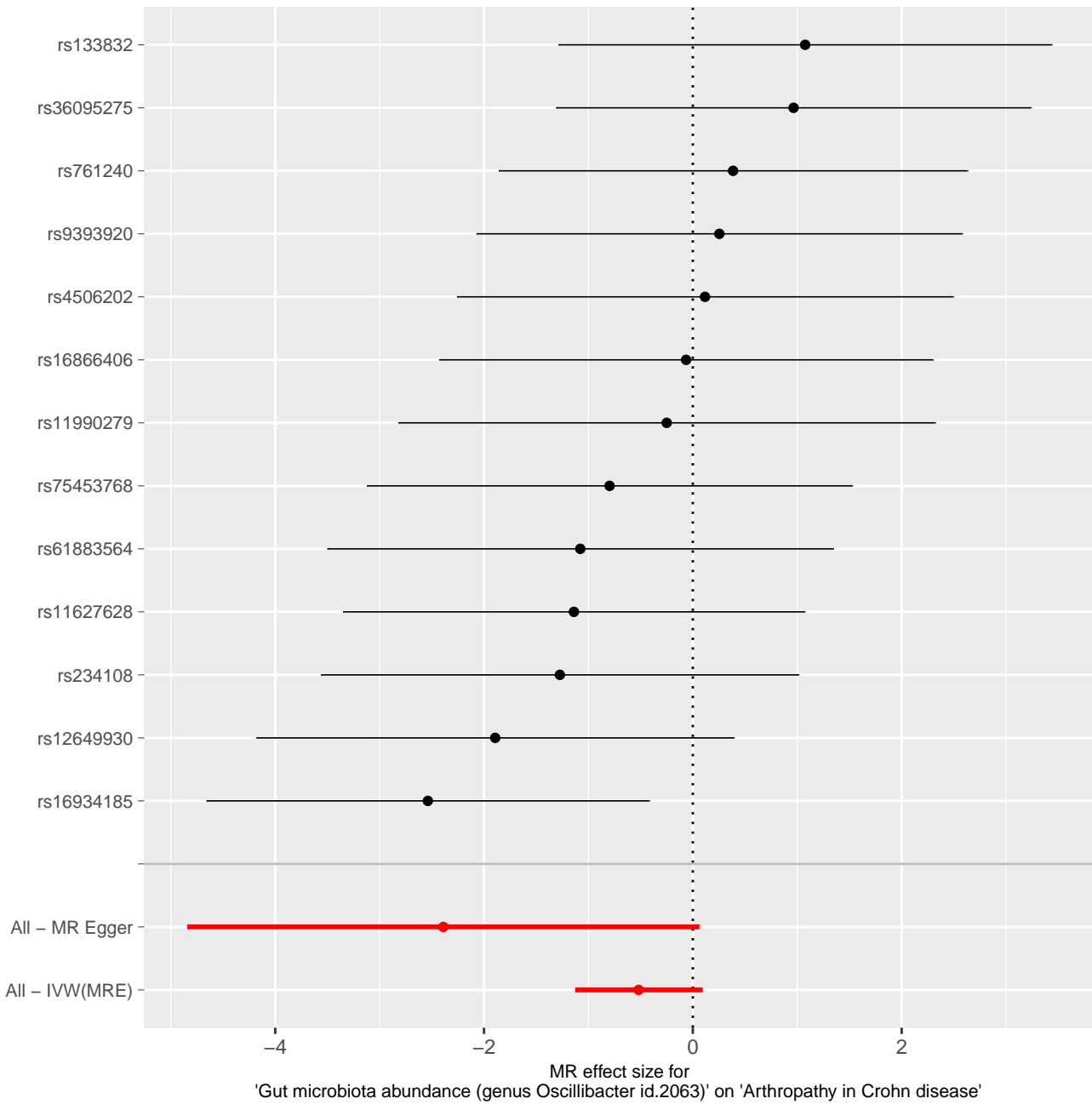


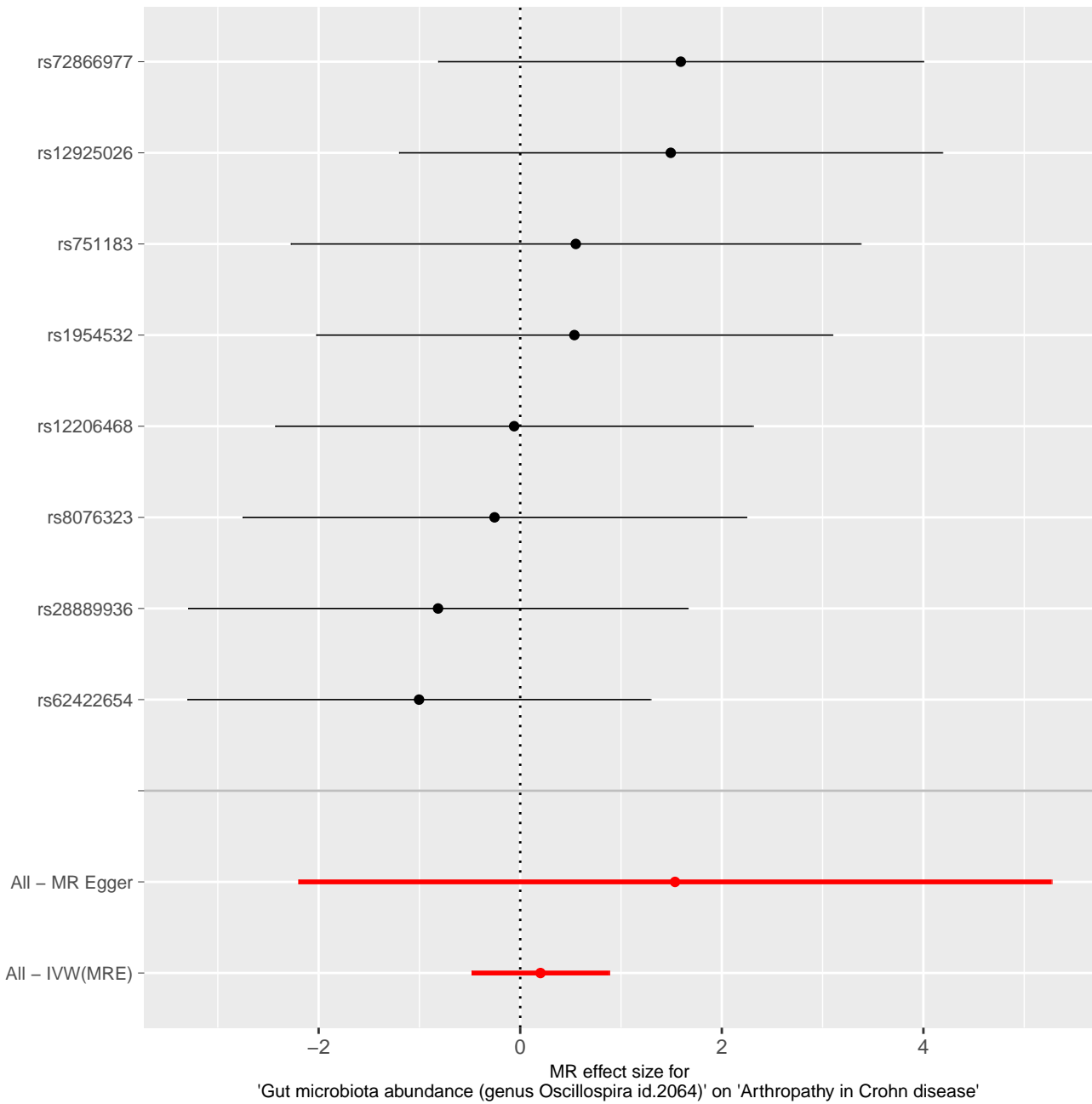




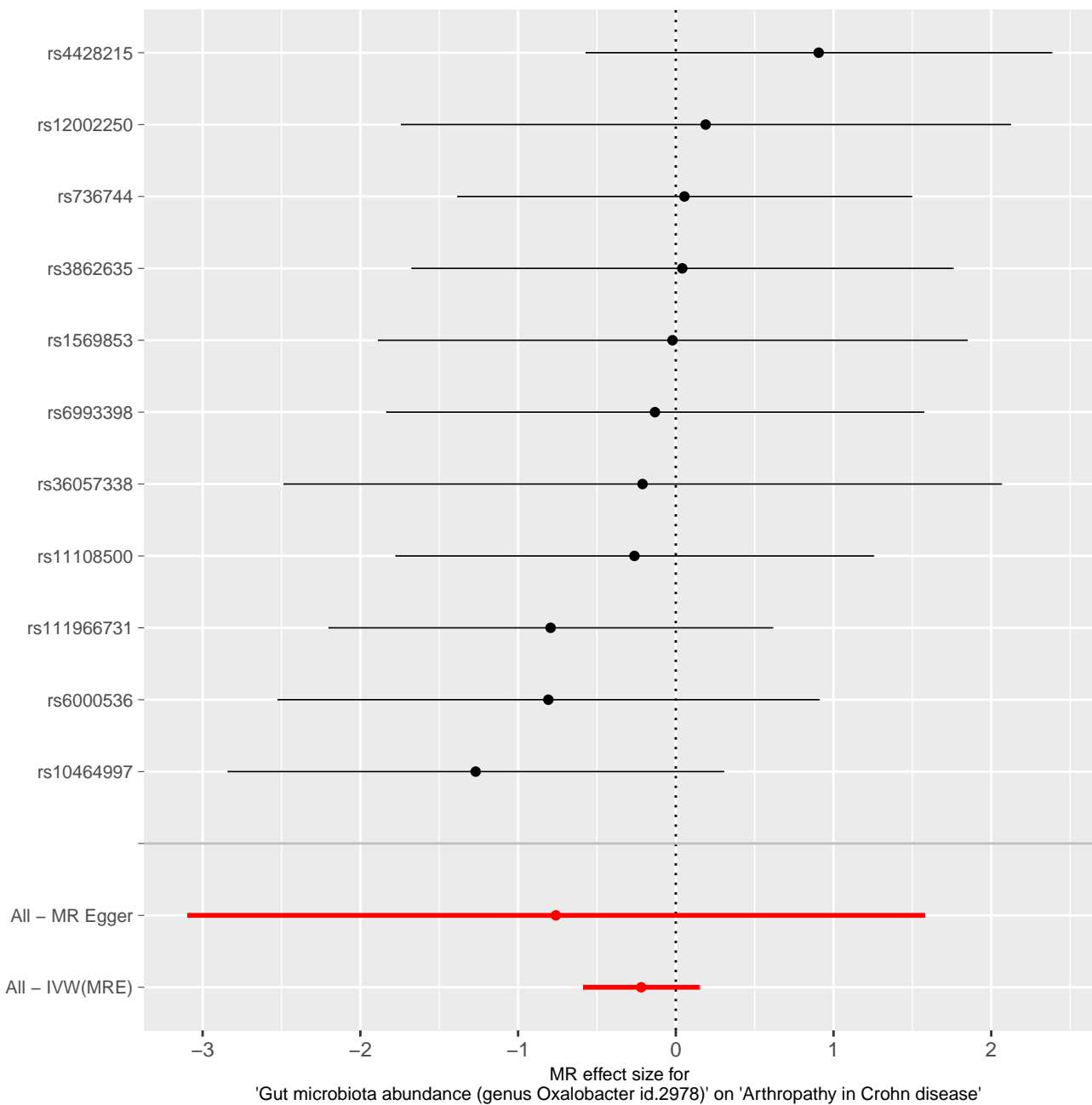




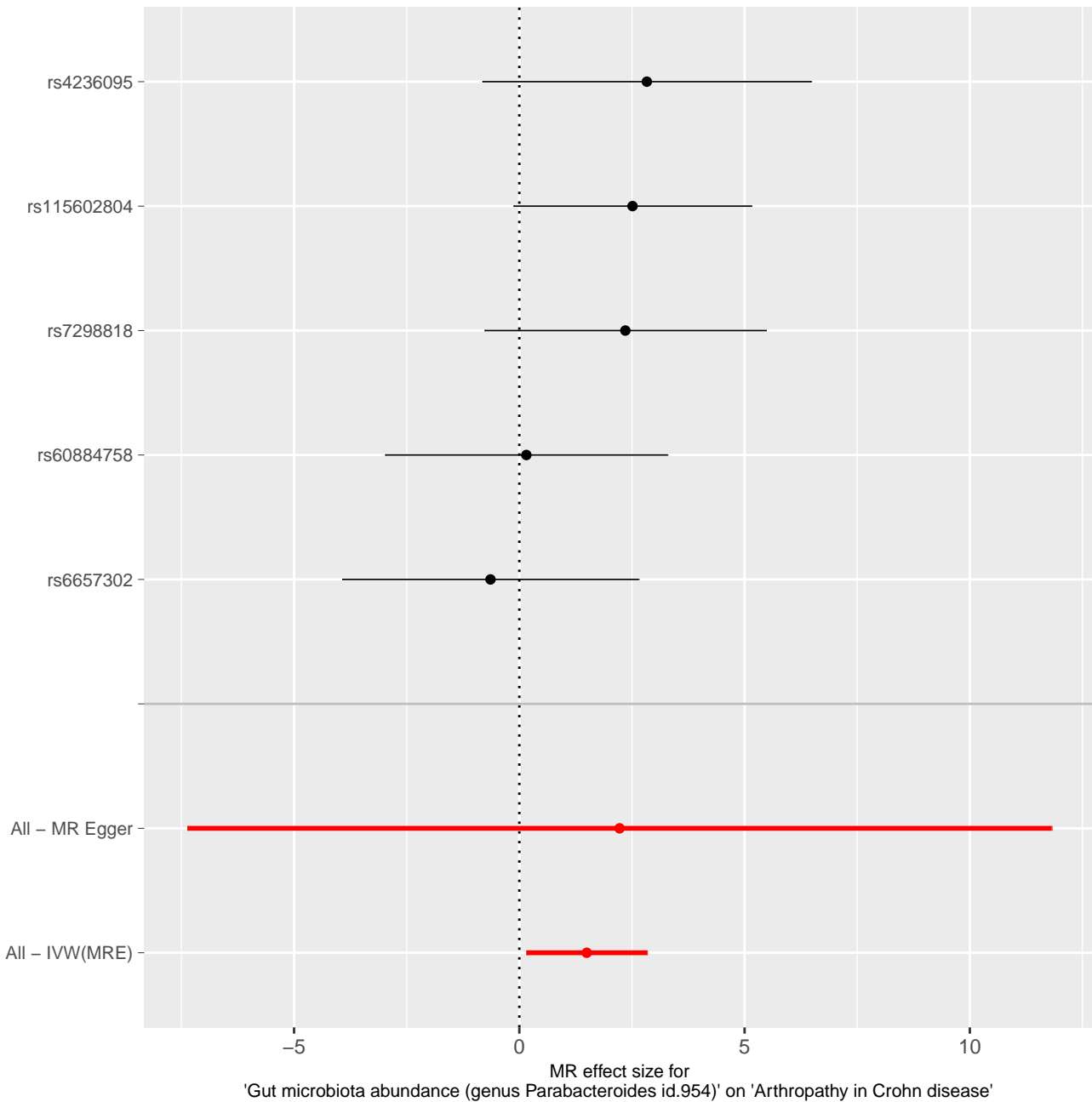




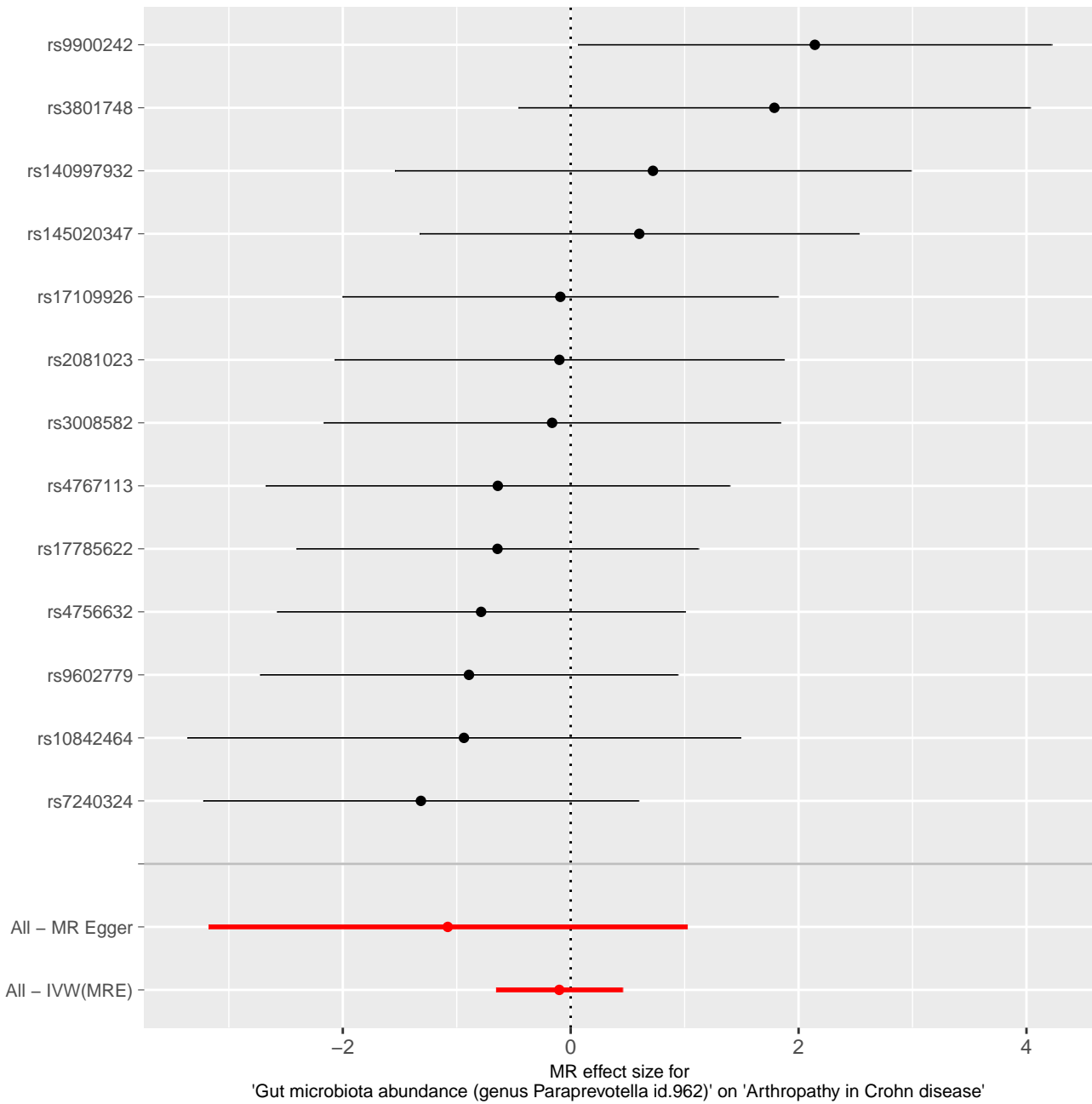
Batch 562 : Gut microbiota abundance (genus Oxalobacter id.2978) on Arthropathy in Crohn disease

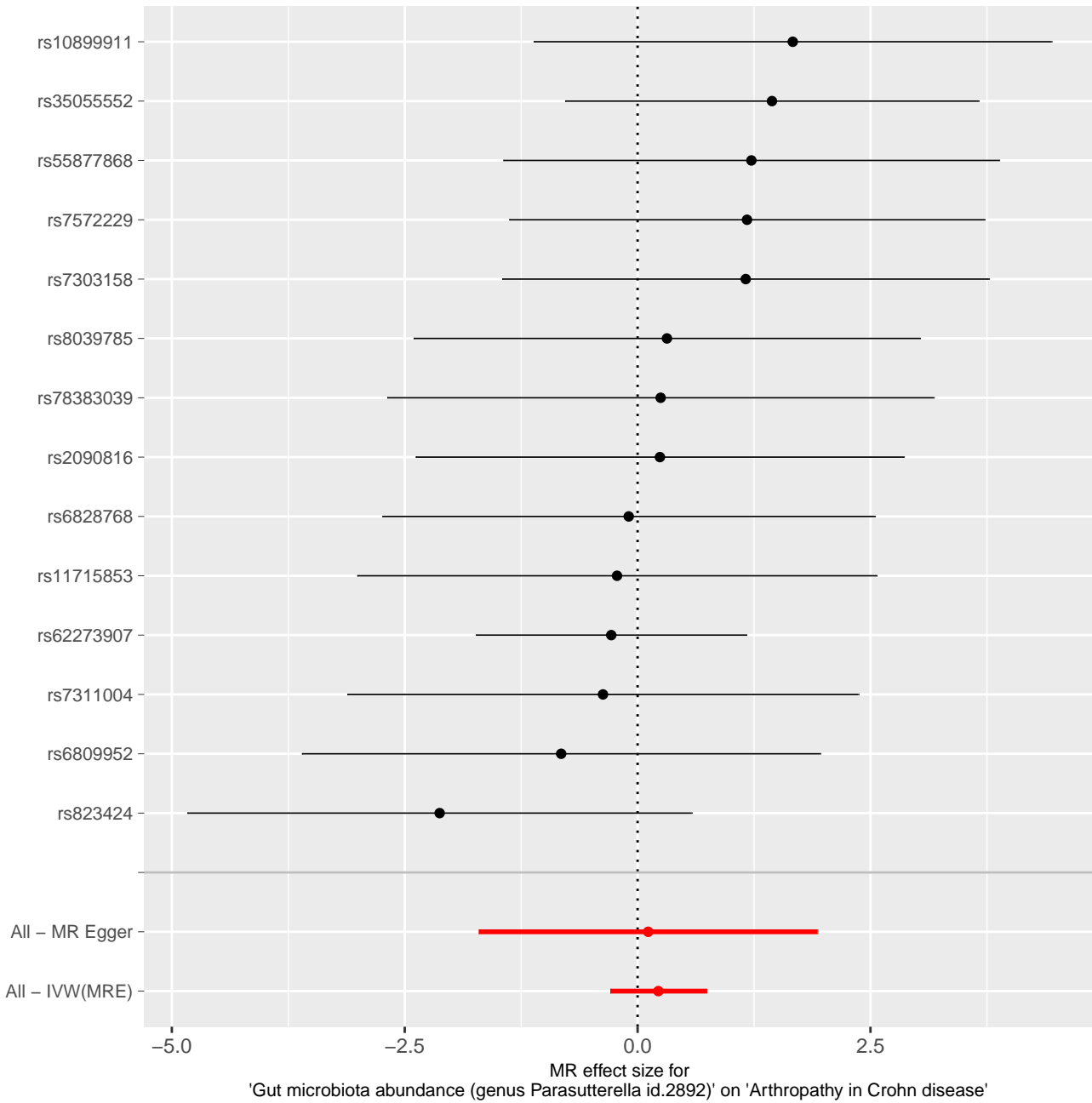


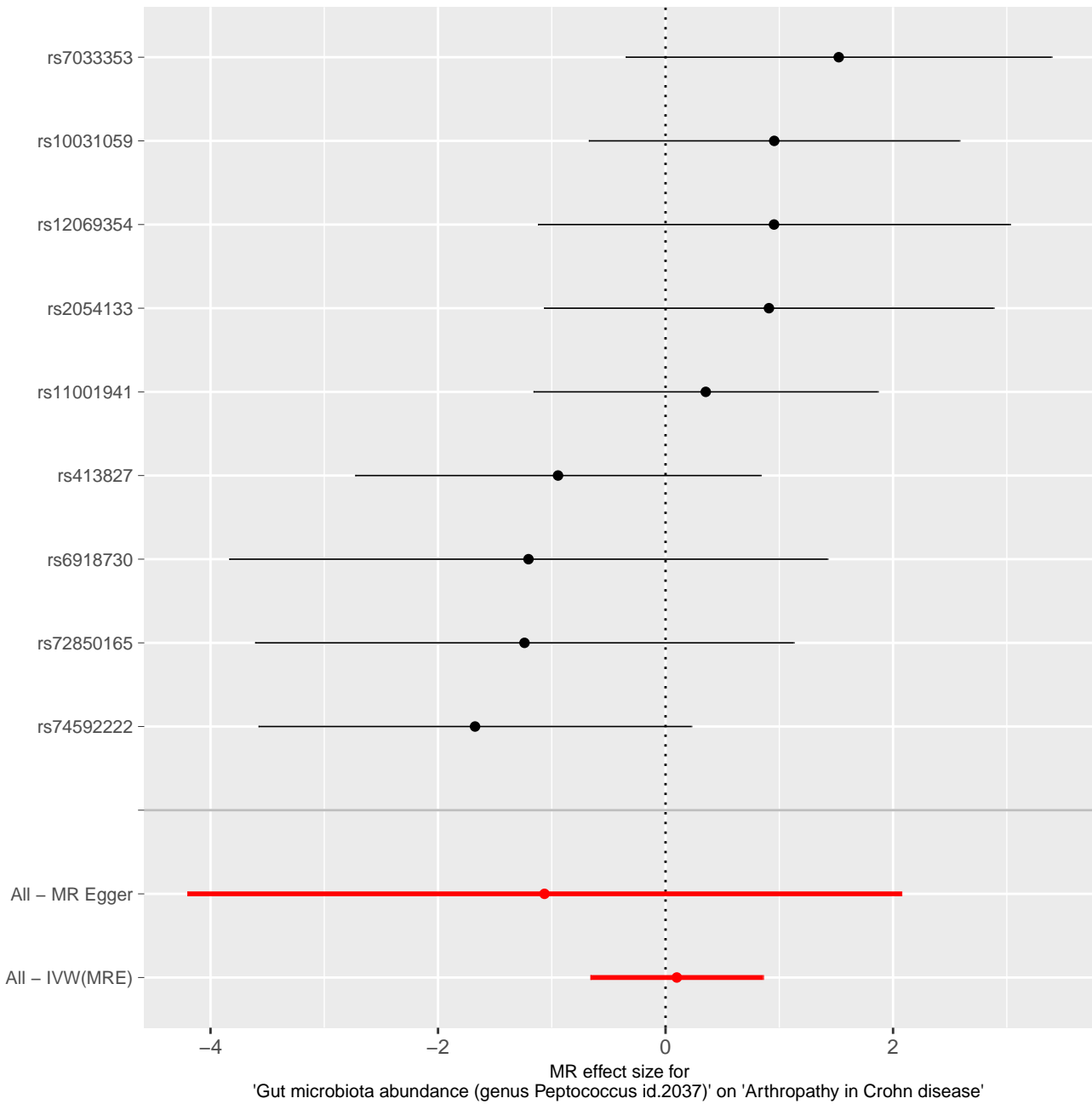
Batch 563 : Gut microbiota abundance (genus Parabacteroides id.954) on Arthropathy in Crohn disease

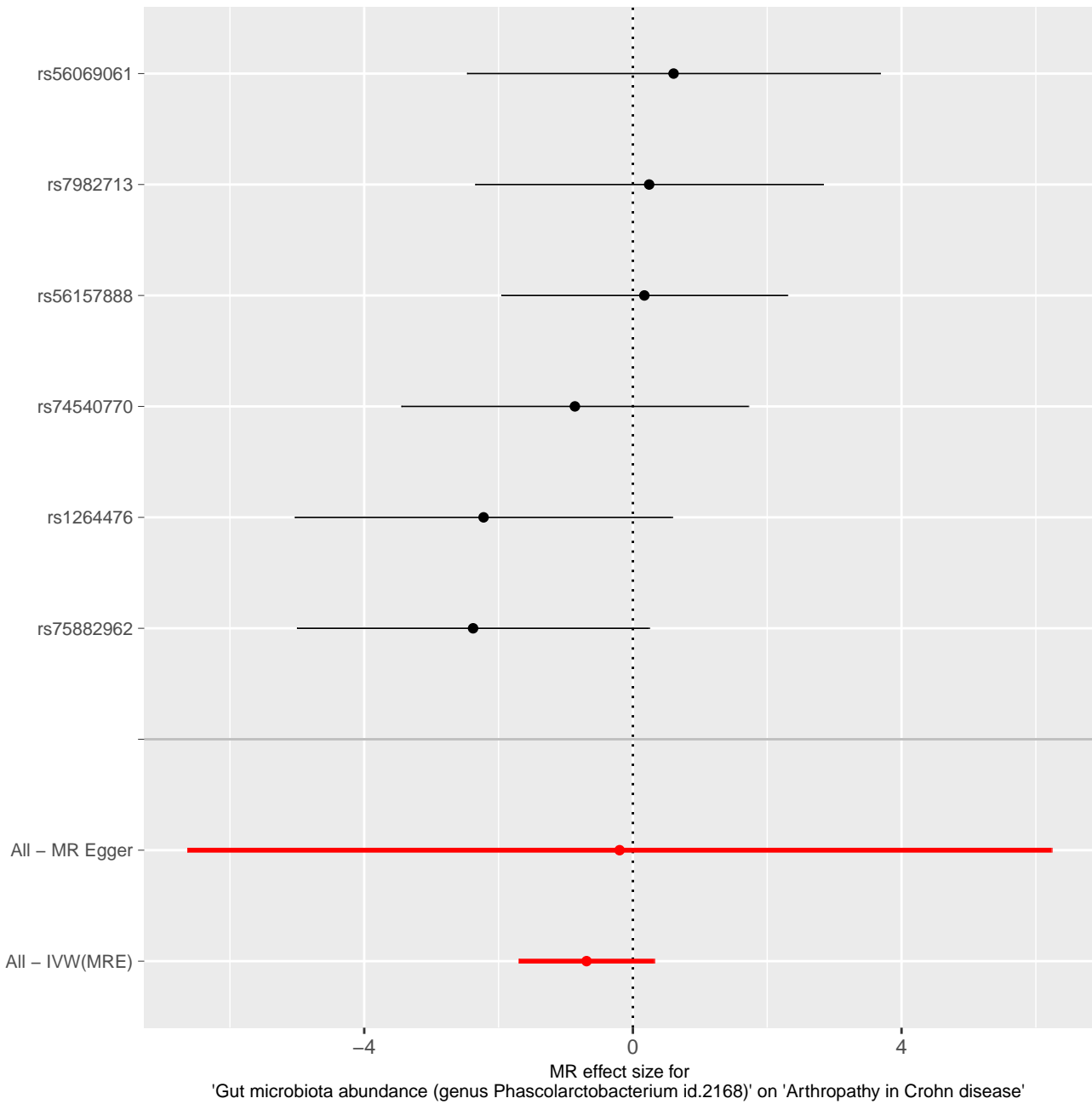


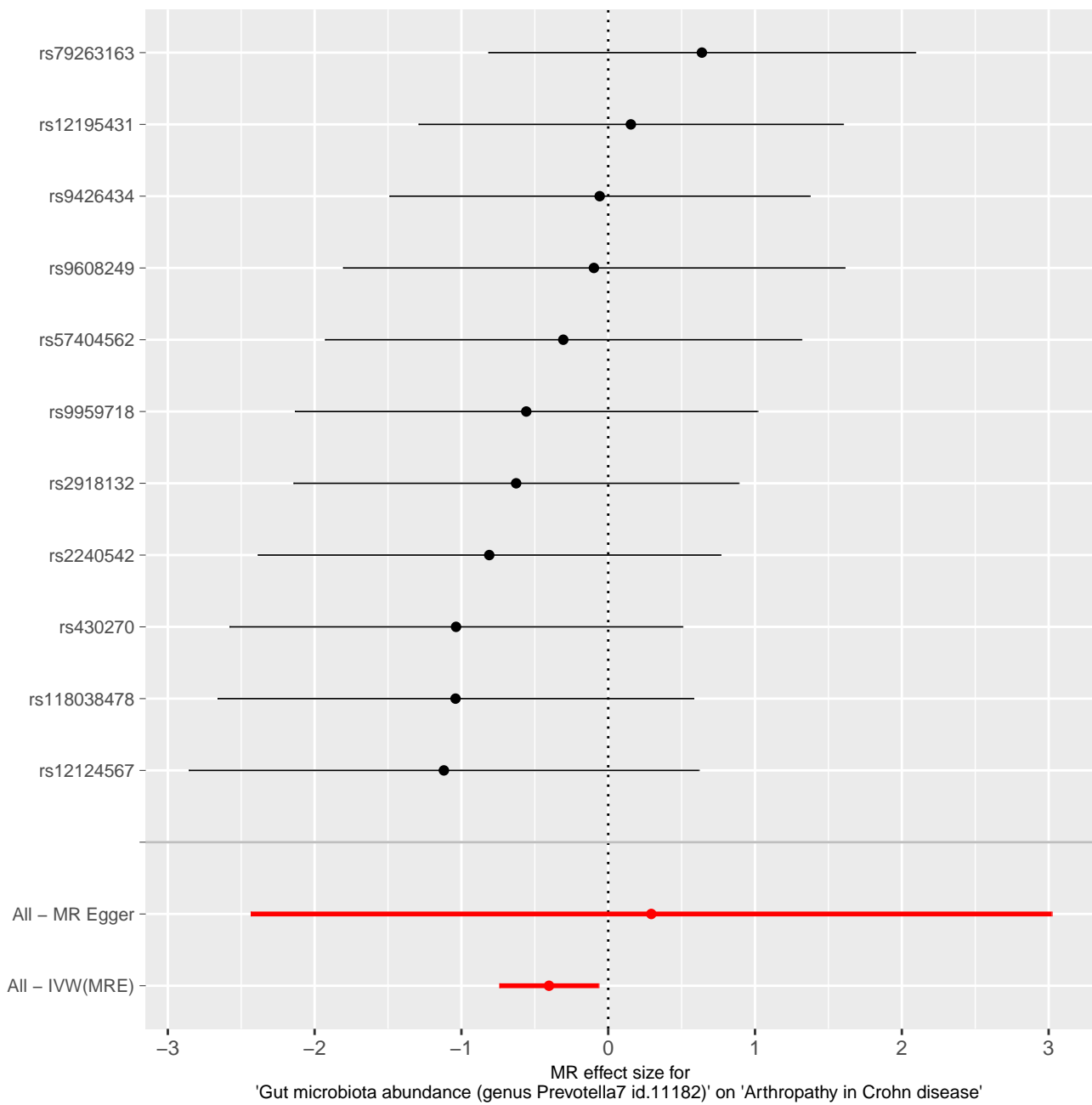
Batch 564 : Gut microbiota abundance (genus Paraprevotella id.962) on Arthropathy in Crohn disease

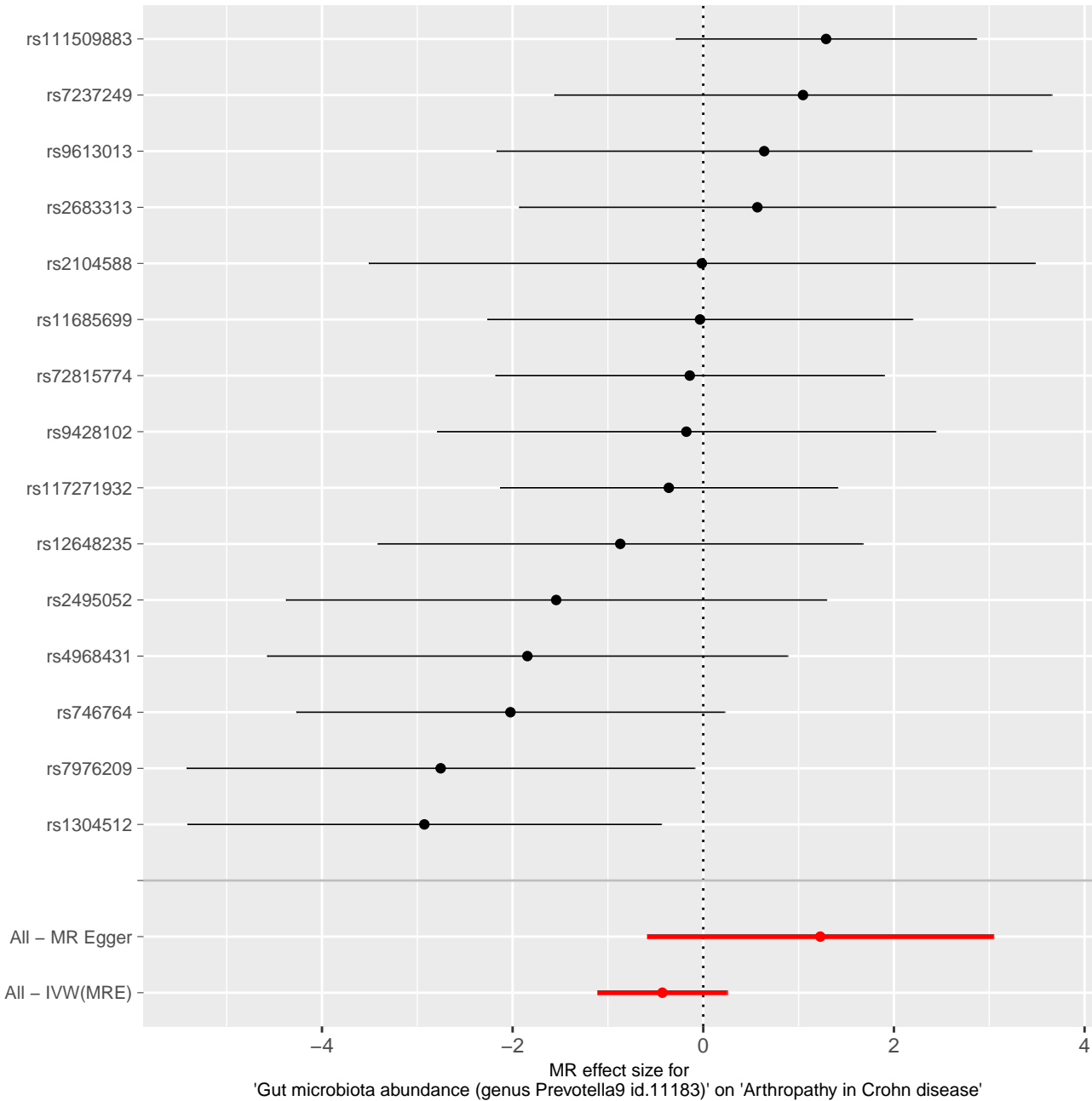


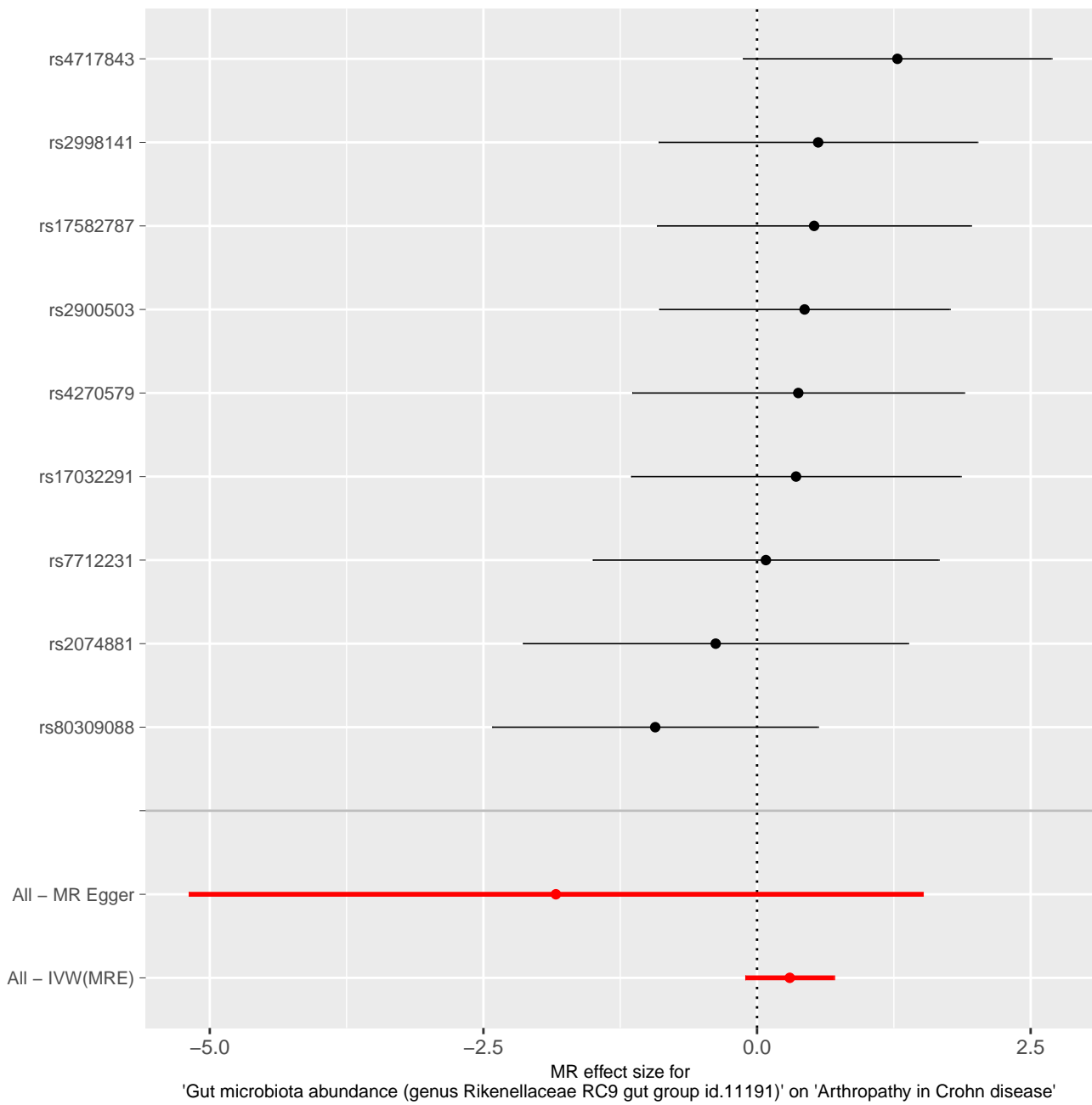




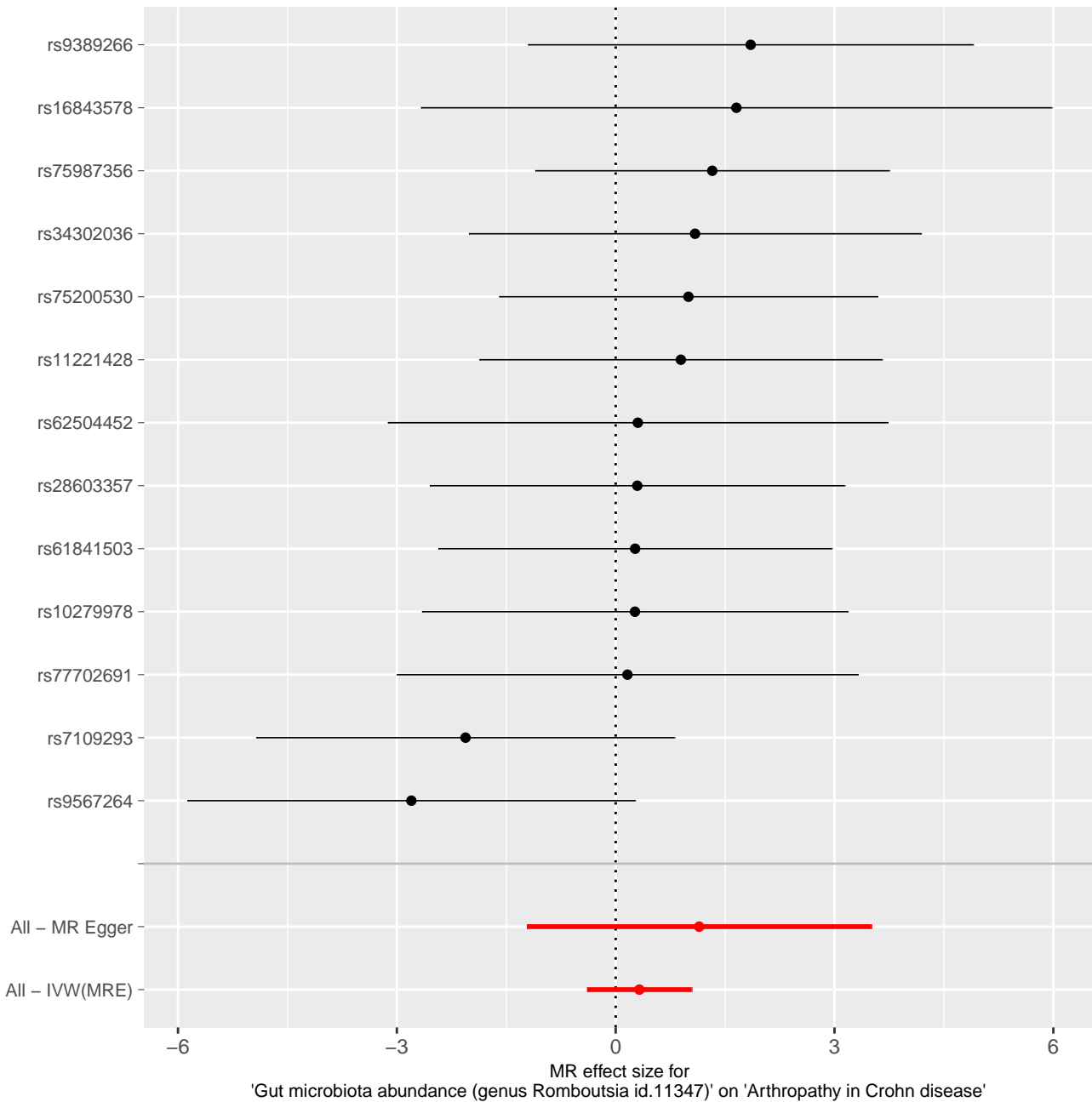




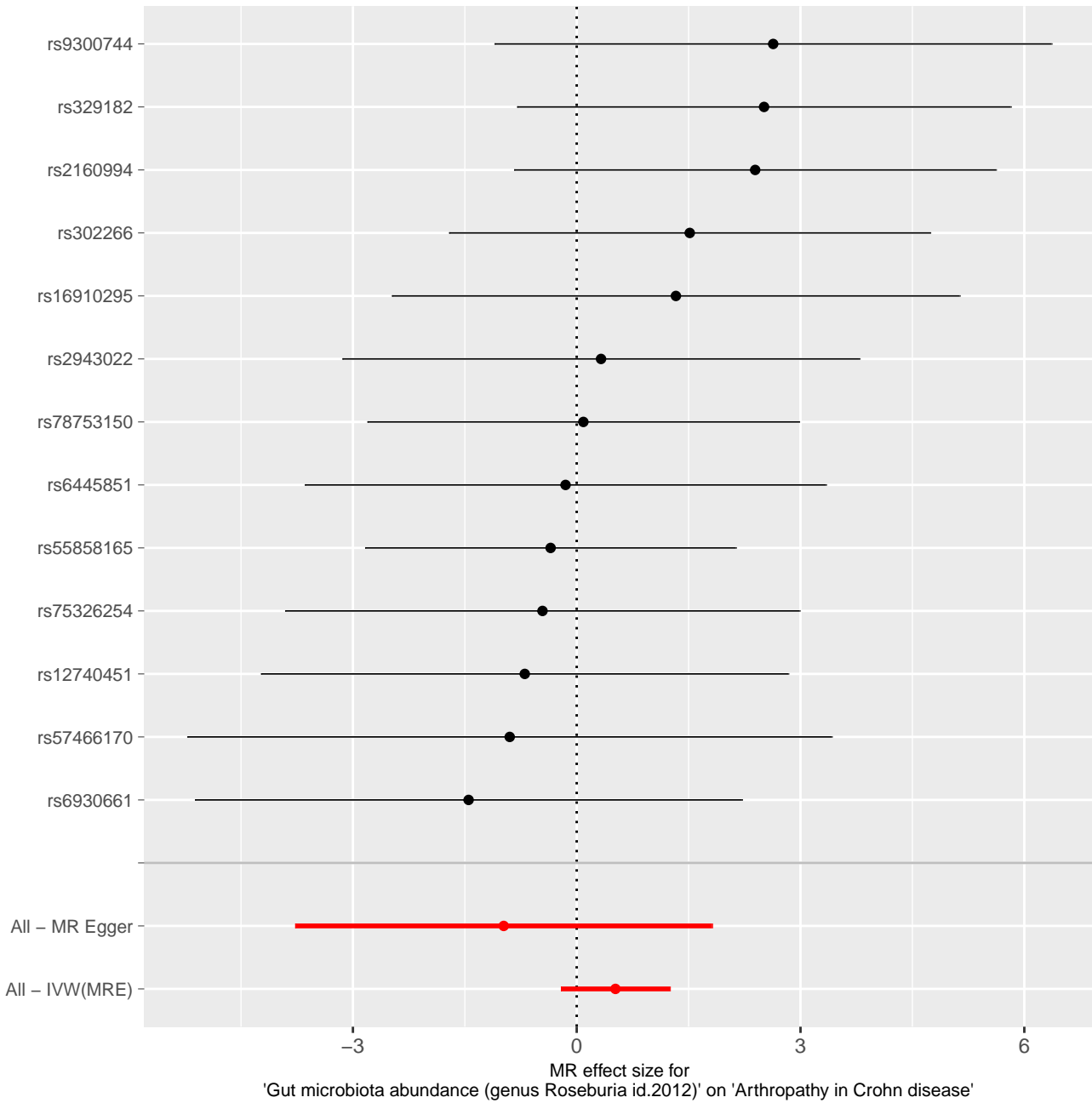




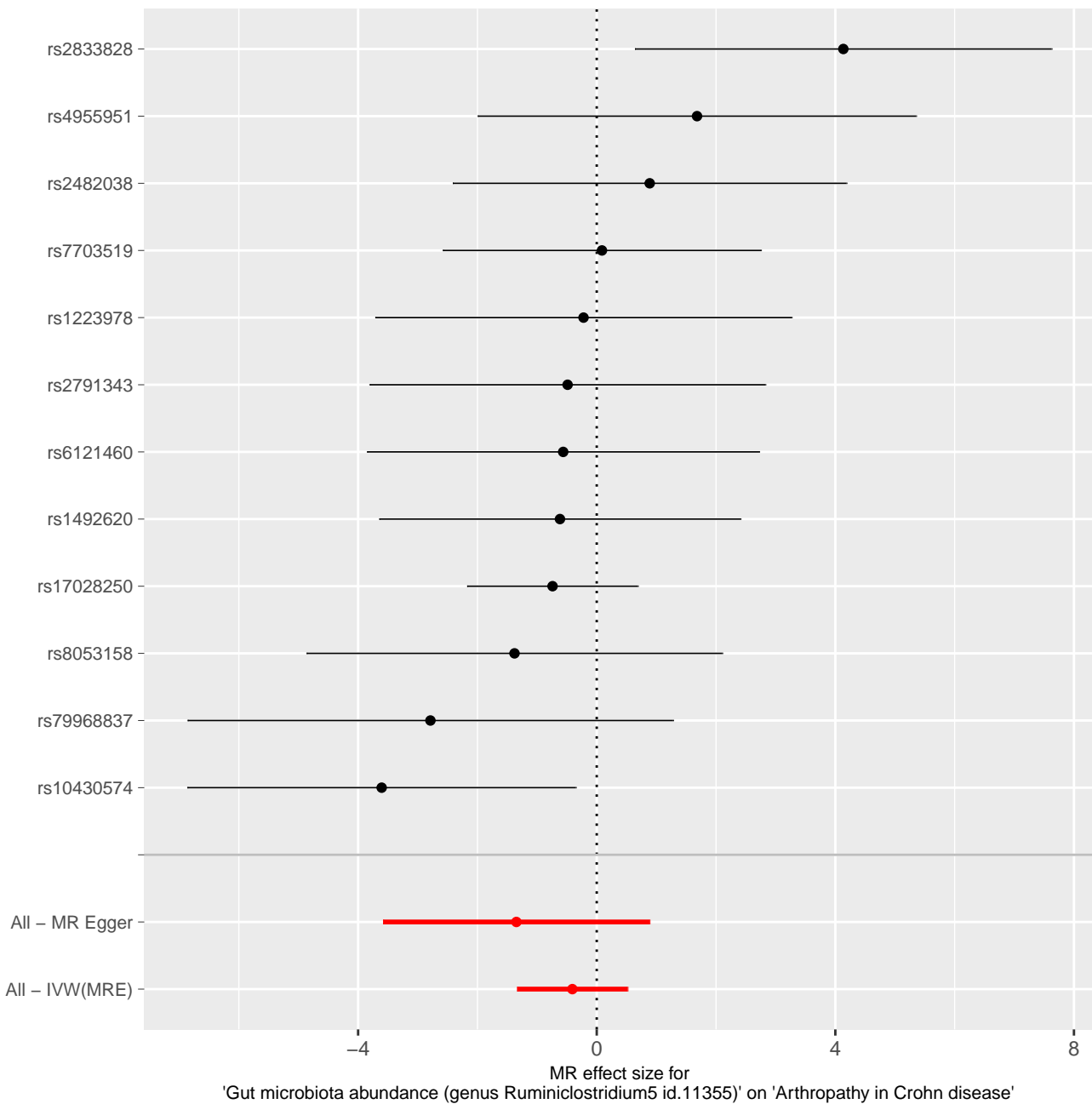
Batch 571 : Gut microbiota abundance (genus Romboutsia id.11347) on Arthropathy in Crohn disease



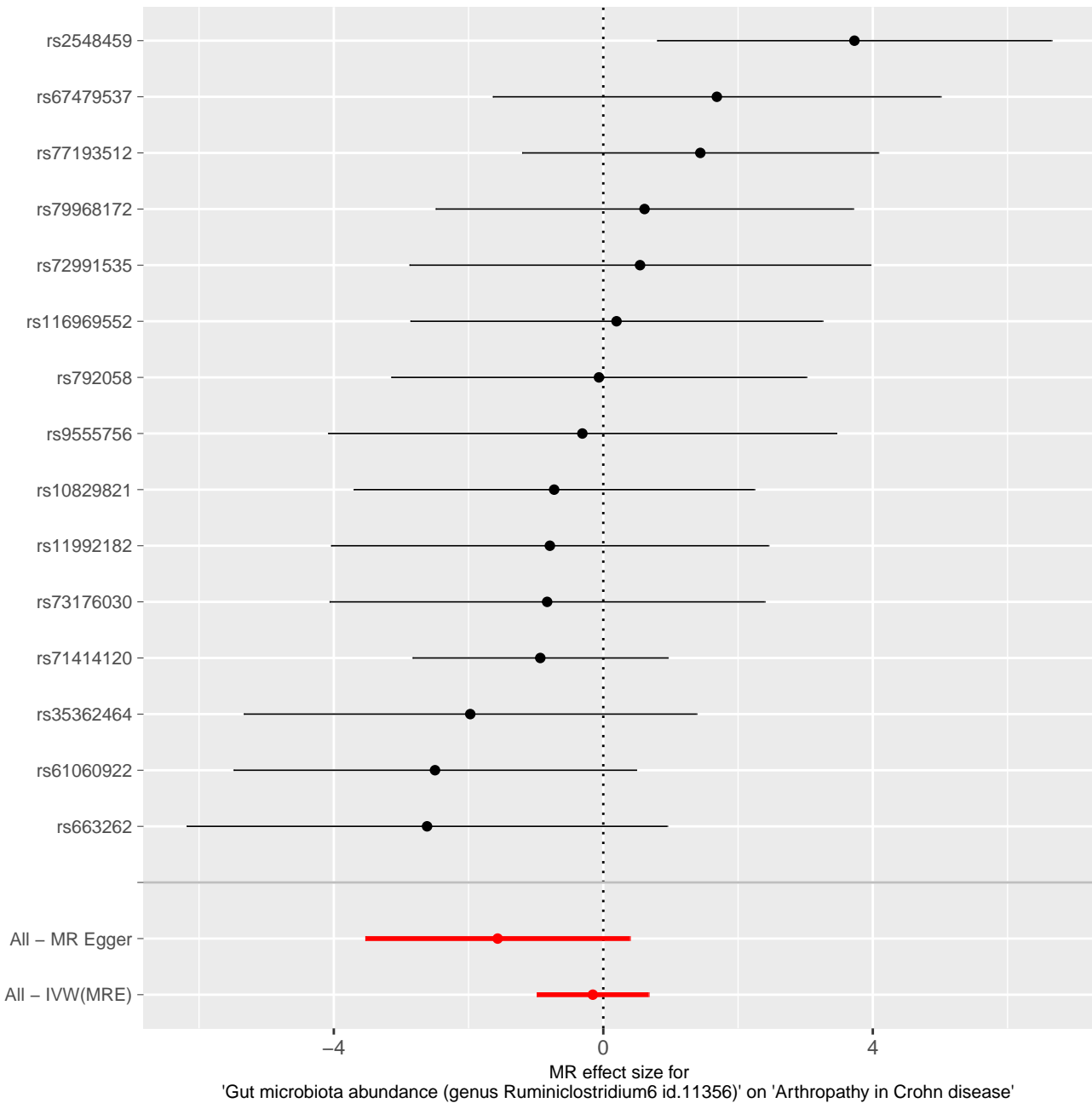
Batch 572 : Gut microbiota abundance (genus Roseburia id.2012) on Arthropathy in Crohn disease

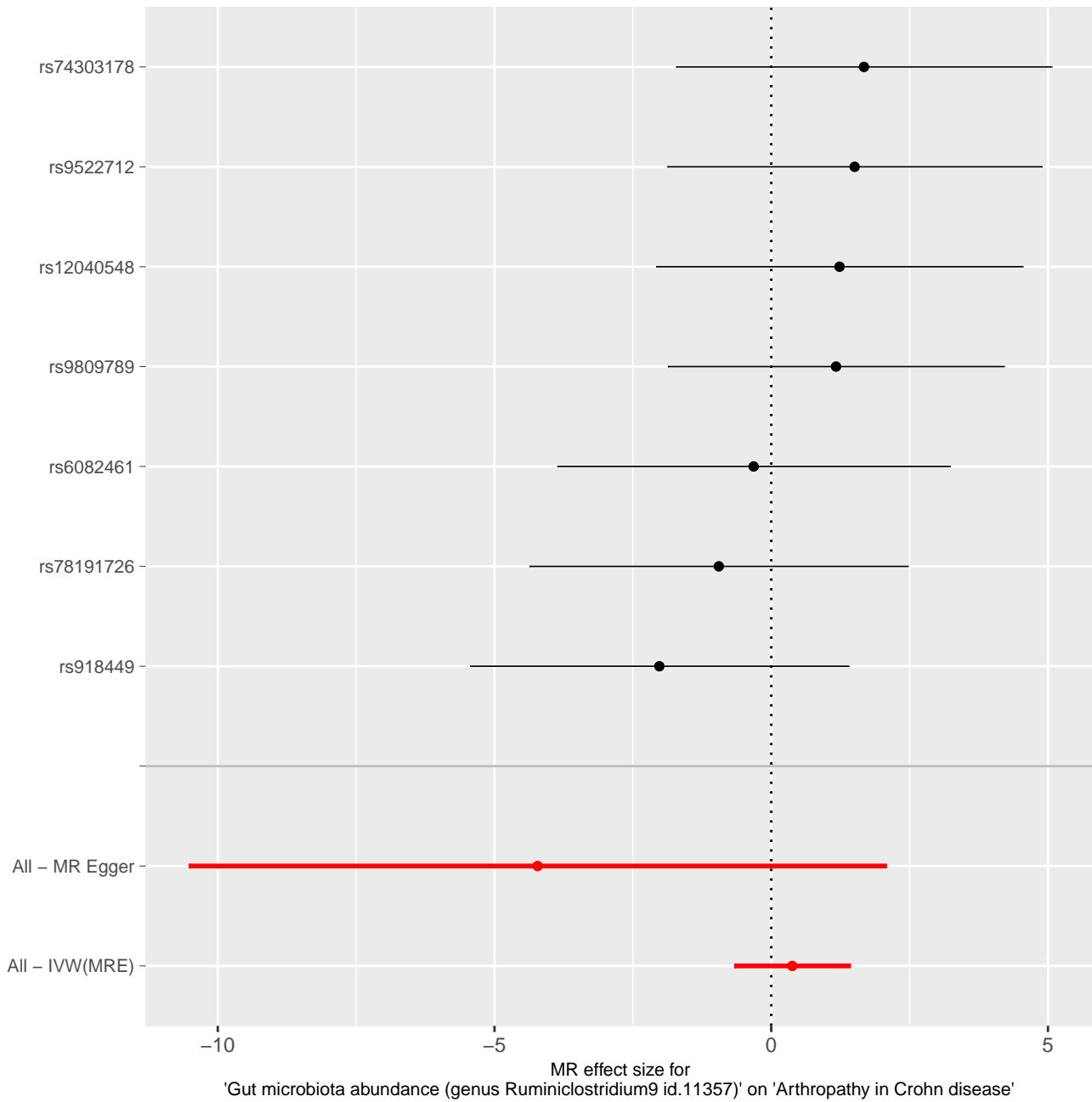


Batch 573 : Gut microbiota abundance (genus Ruminiclostridium5 id.11355) on Arthropathy in Crohn disease

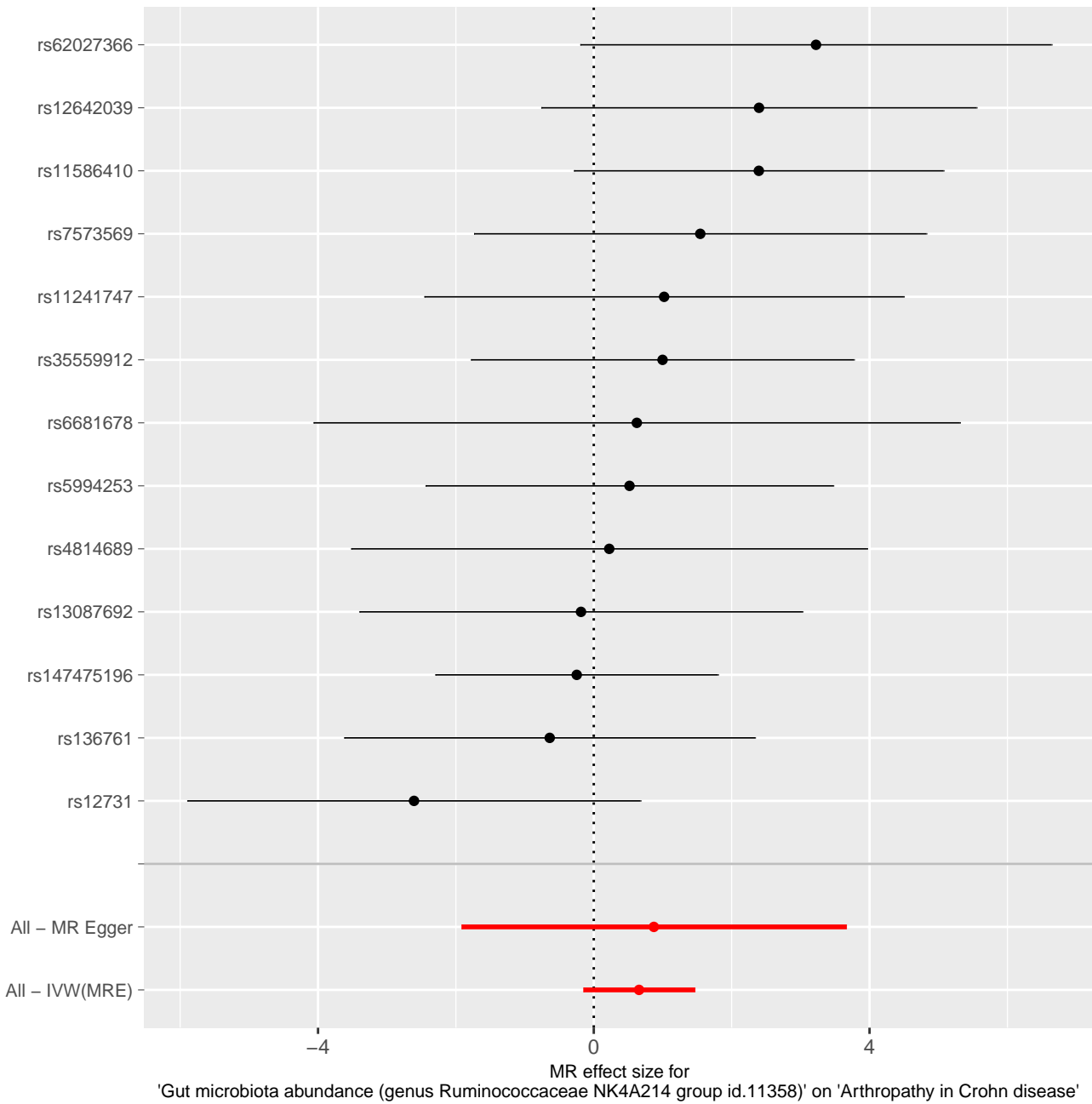


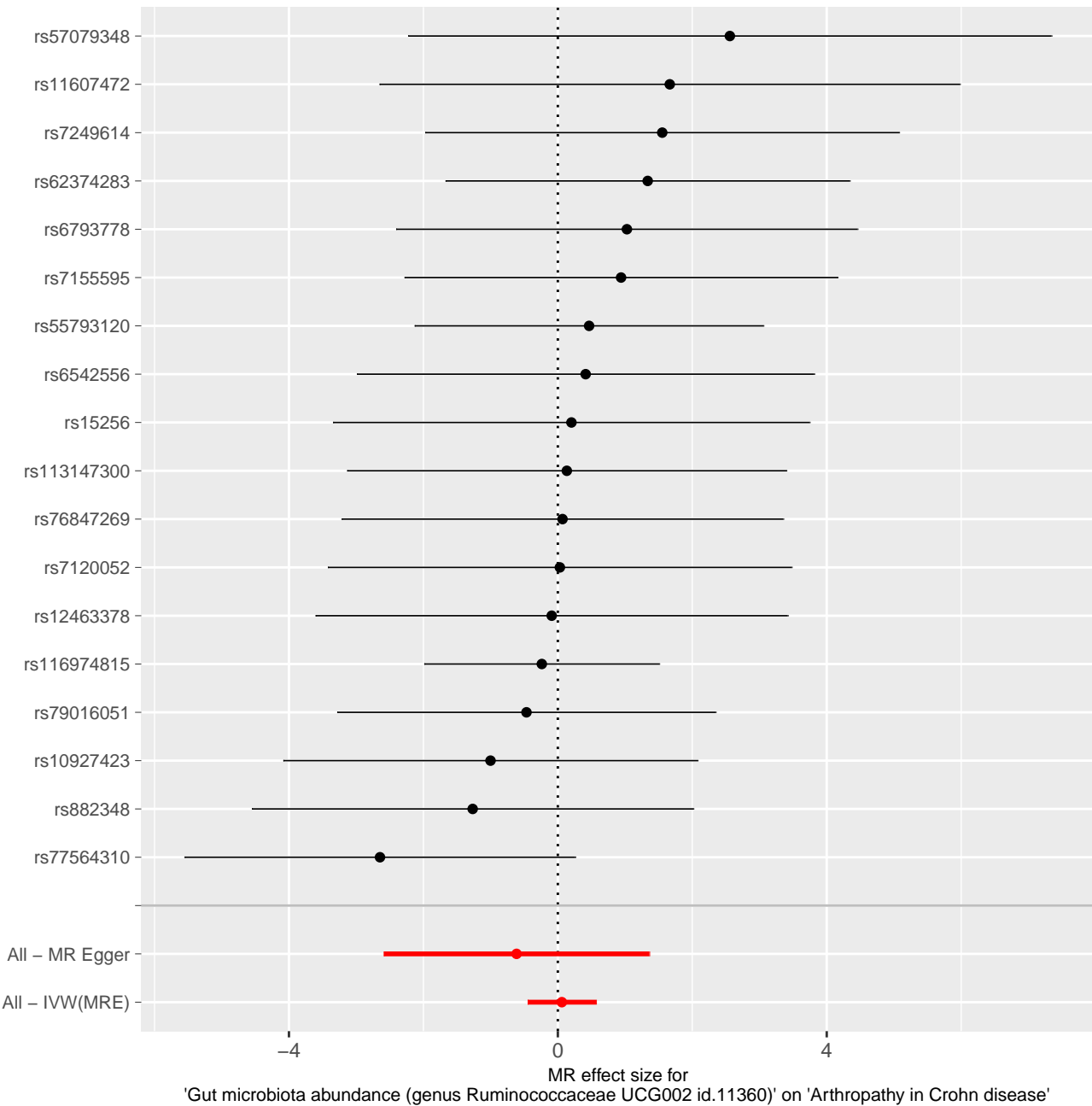
Batch 574 : Gut microbiota abundance (genus Ruminiclostridium6 id.11356) on Arthropathy in Crohn disease



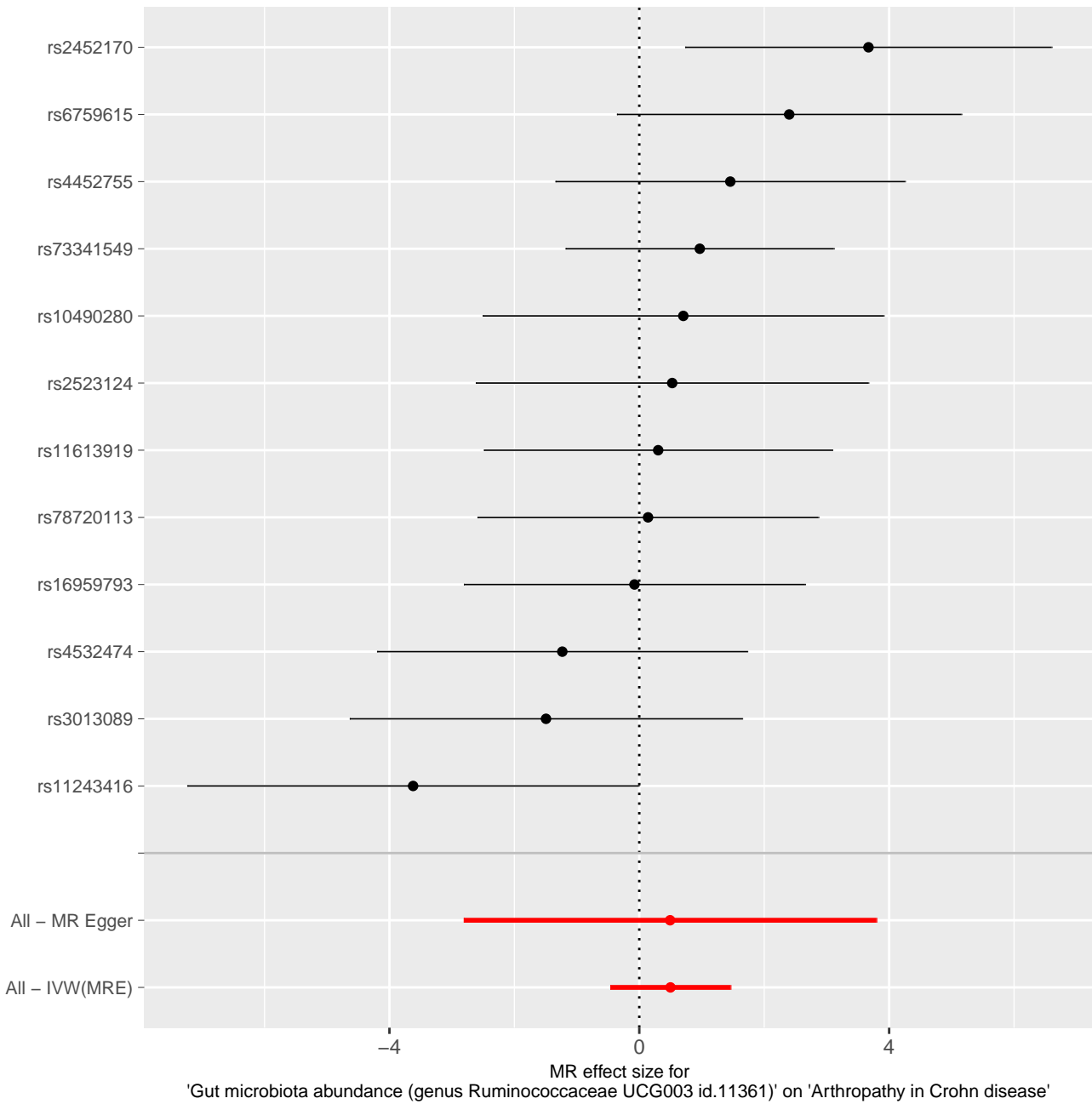


Batch 576 : Gut microbiota abundance (genus Ruminococcaceae NK4A214 group id.11358) on Arthropathy in Crohn disease

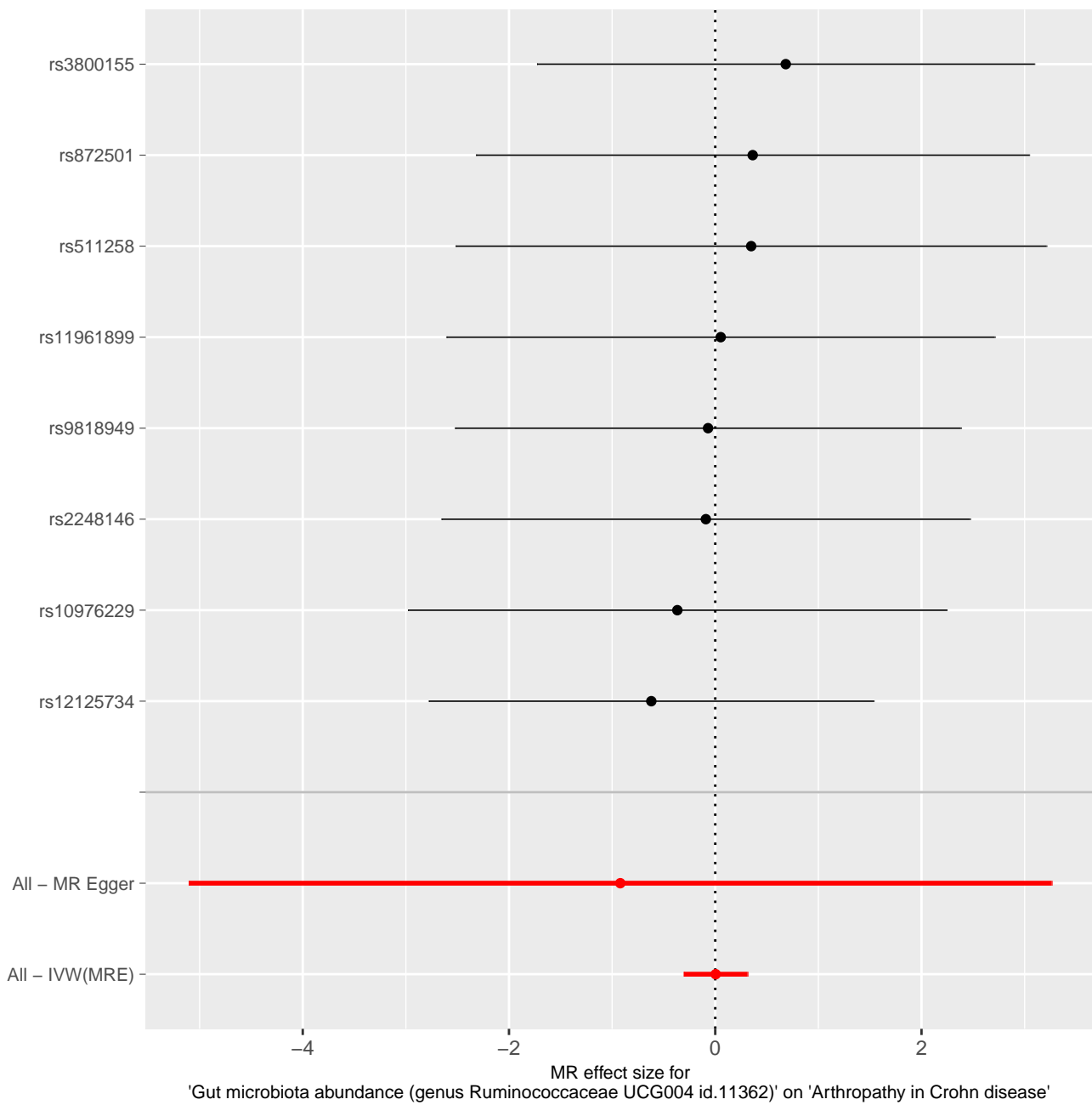


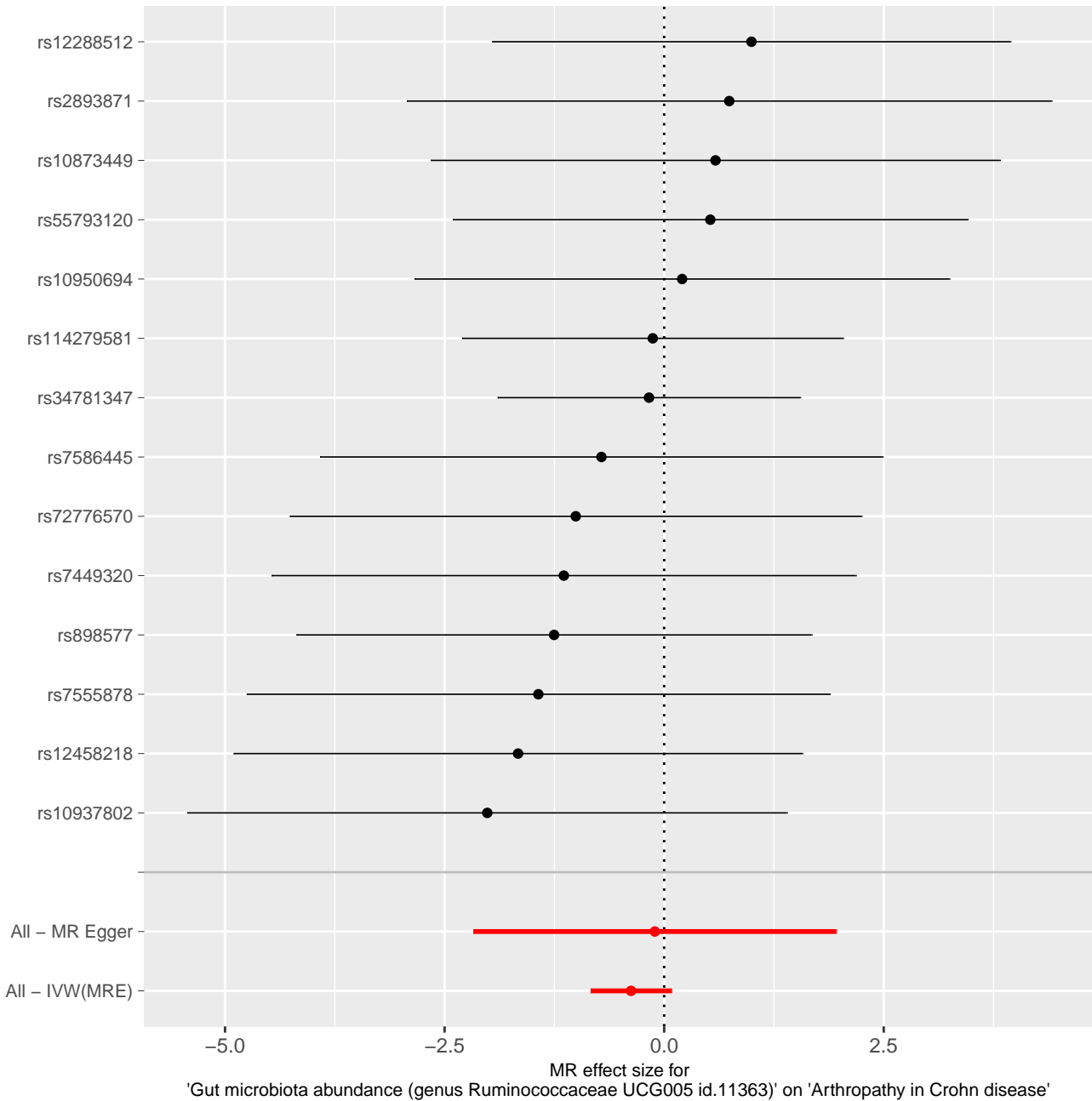


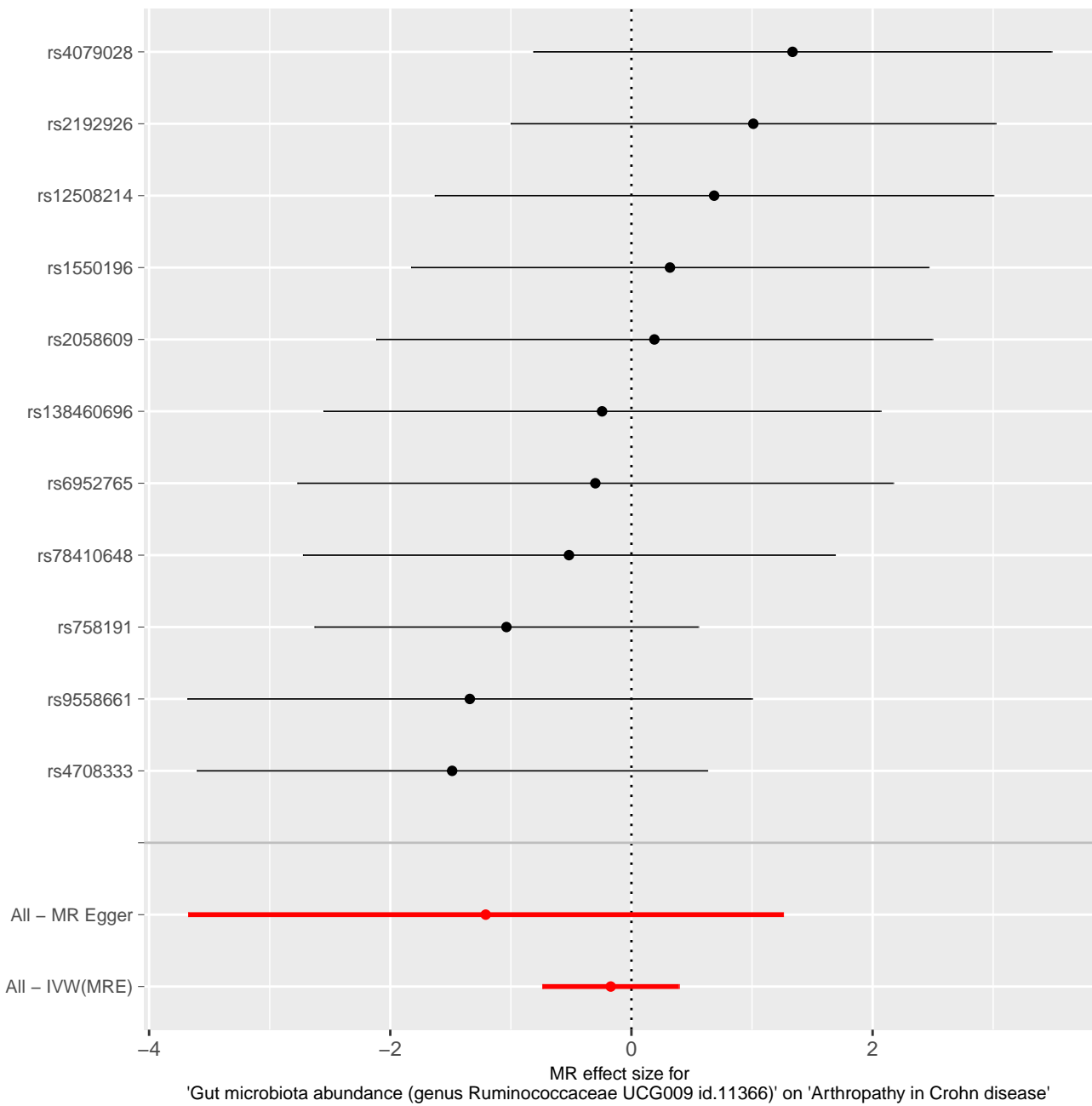
Batch 578 : Gut microbiota abundance (genus Ruminococcaceae UCG003 id.11361) on Arthropathy in Crohn disease

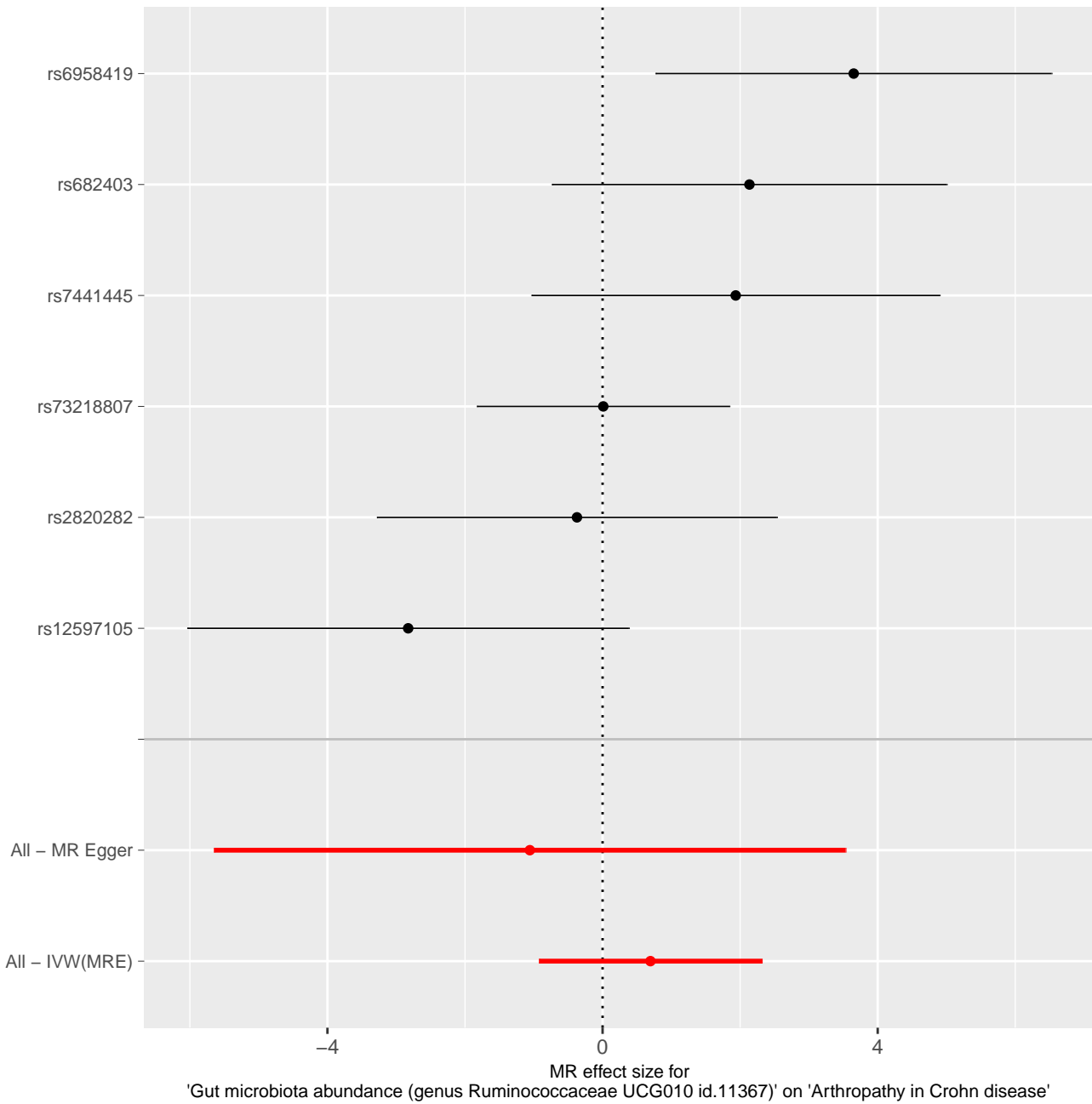


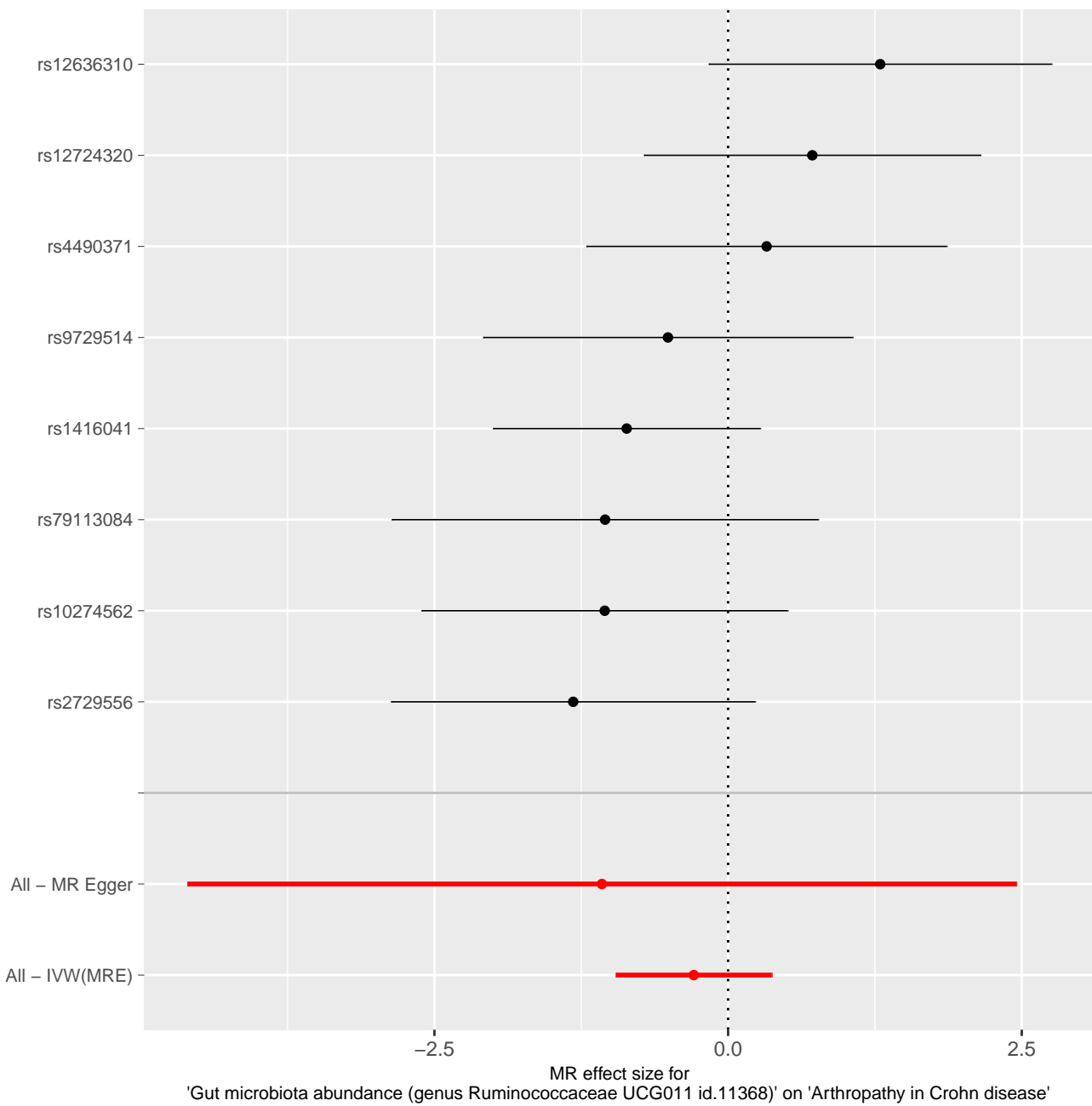
Batch 579 : Gut microbiota abundance (genus Ruminococcaceae UCG004 id.11362) on Arthropathy in Crohn disease



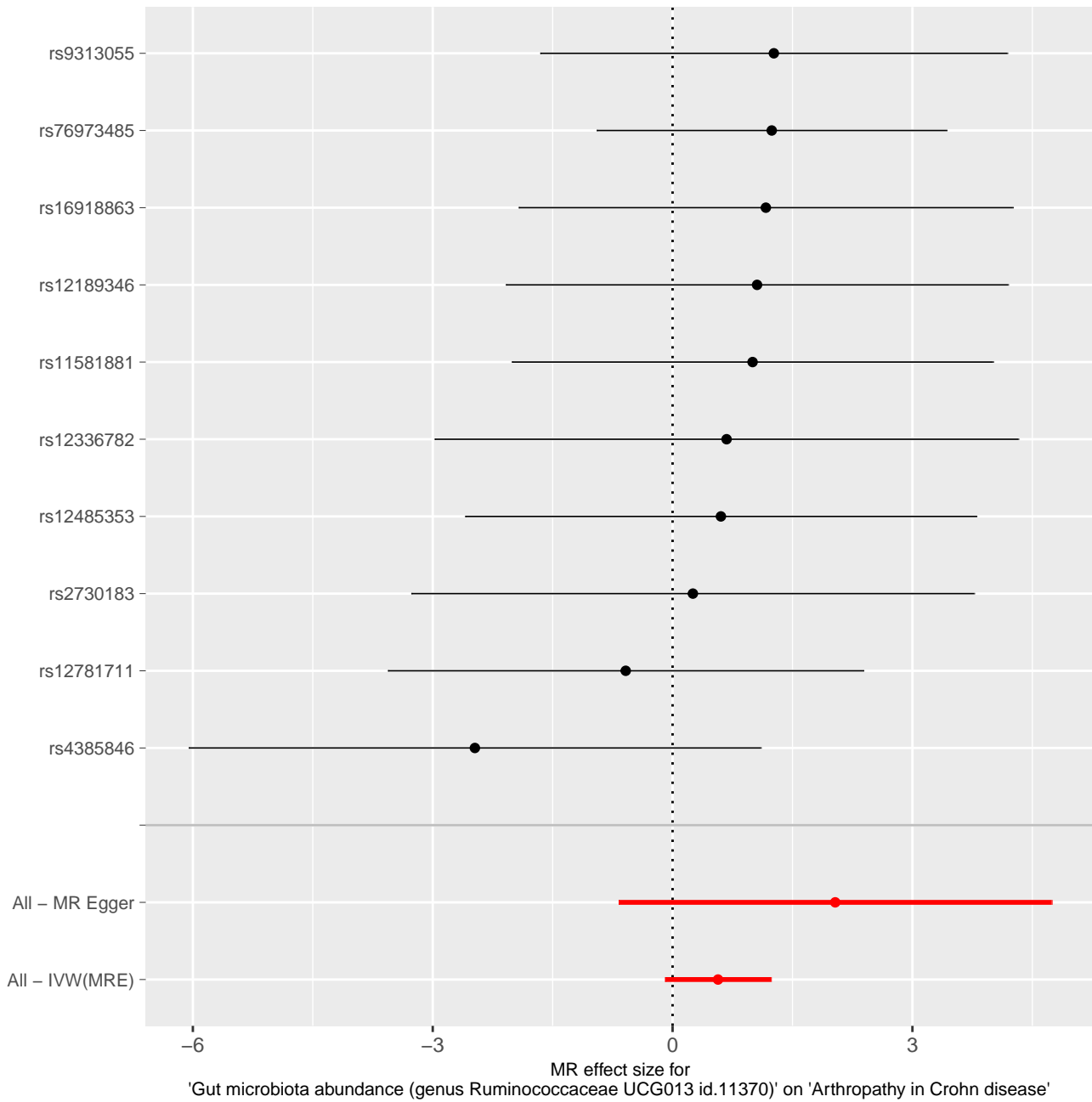


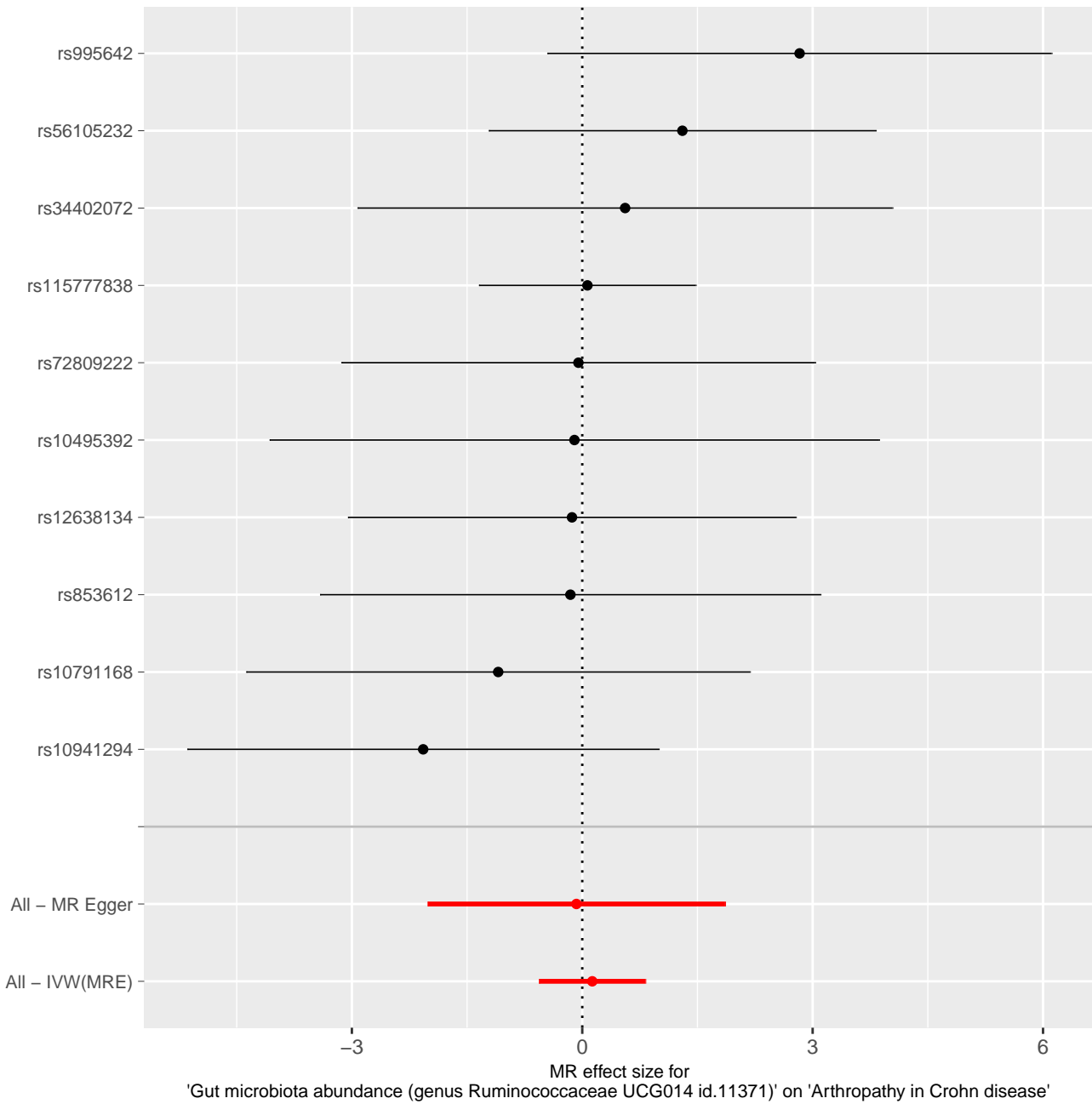


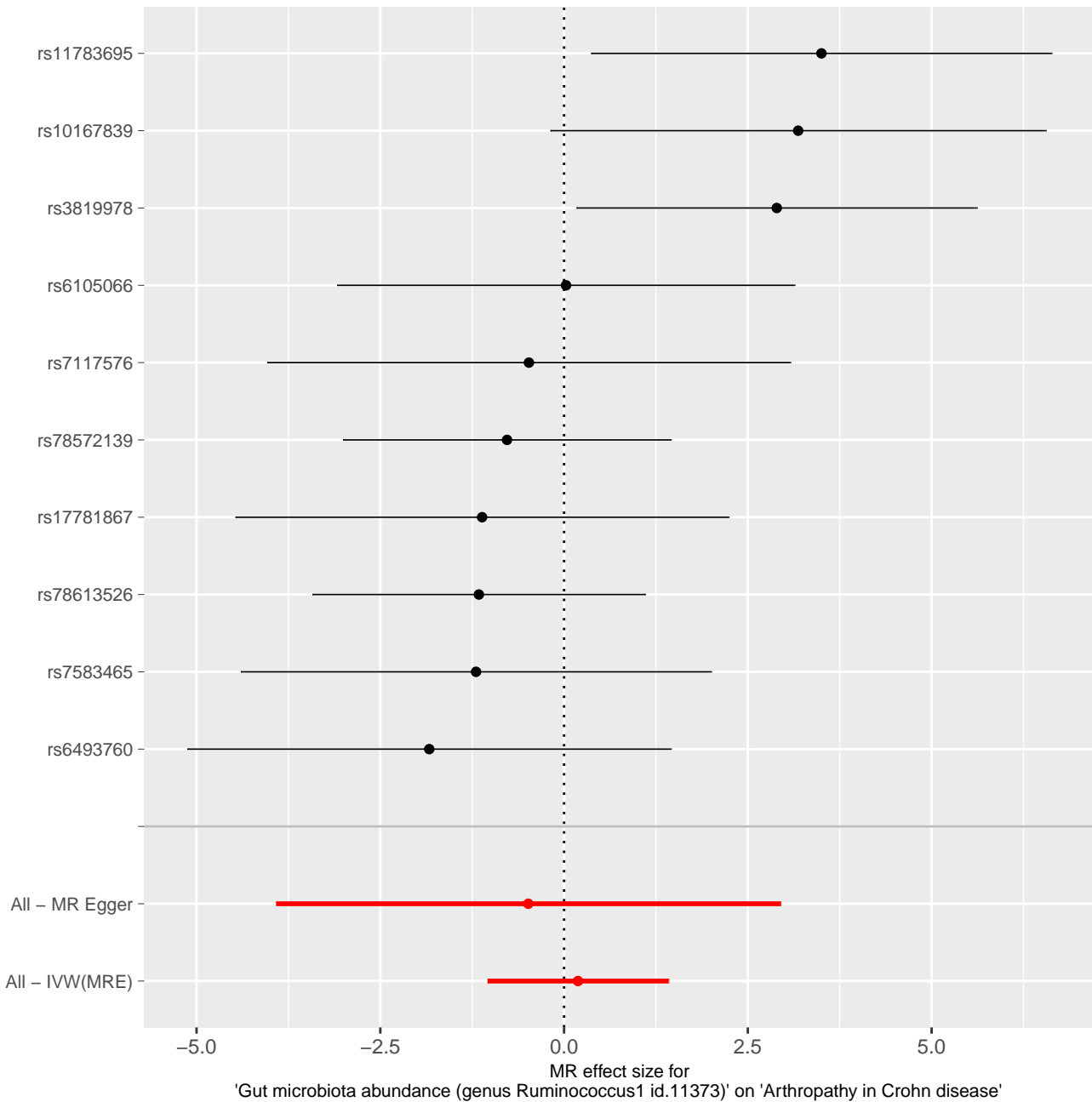




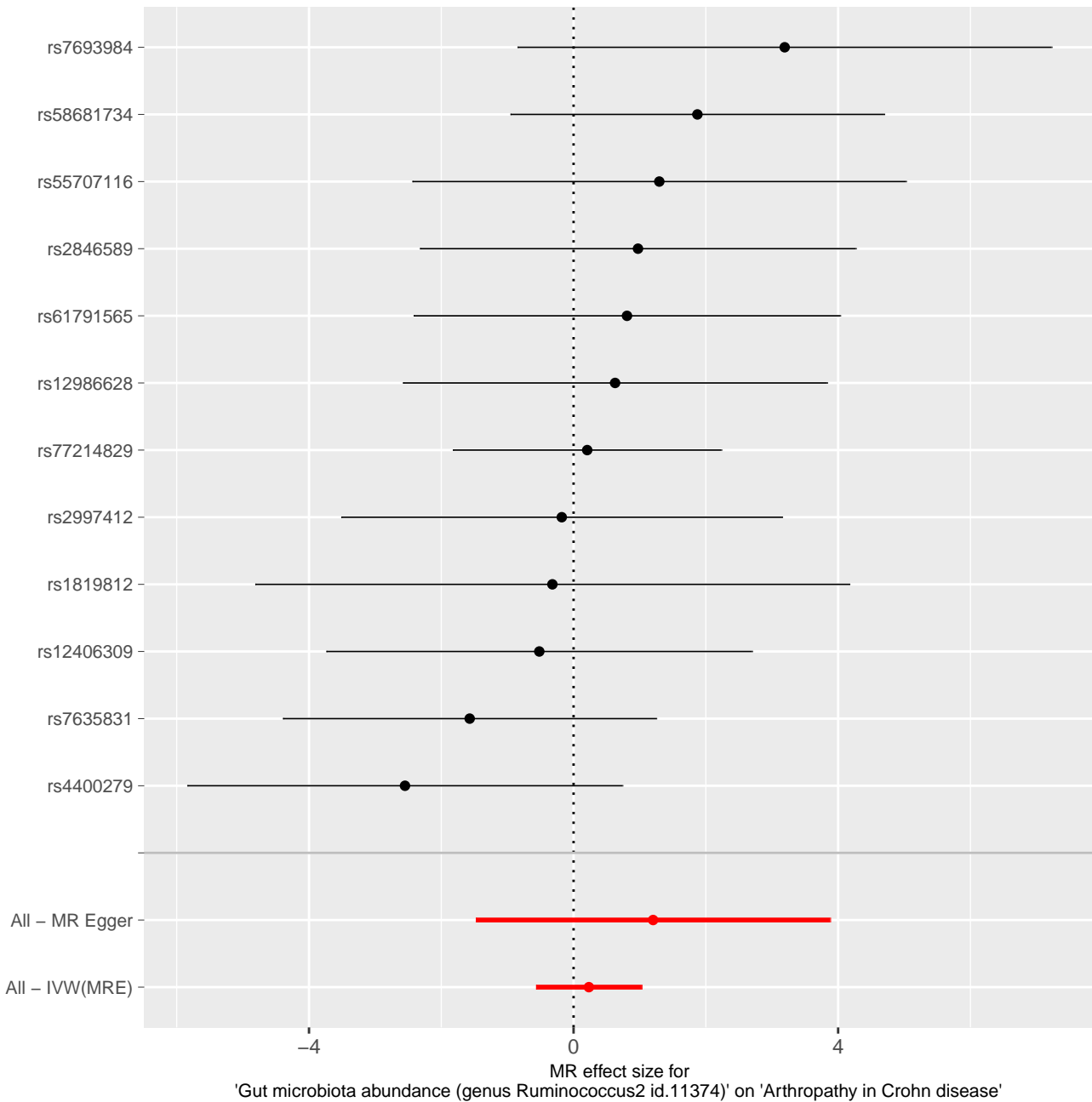
Batch 584 : Gut microbiota abundance (genus Ruminococcaceae UCG013 id.11370) on Arthropathy in Crohn disease

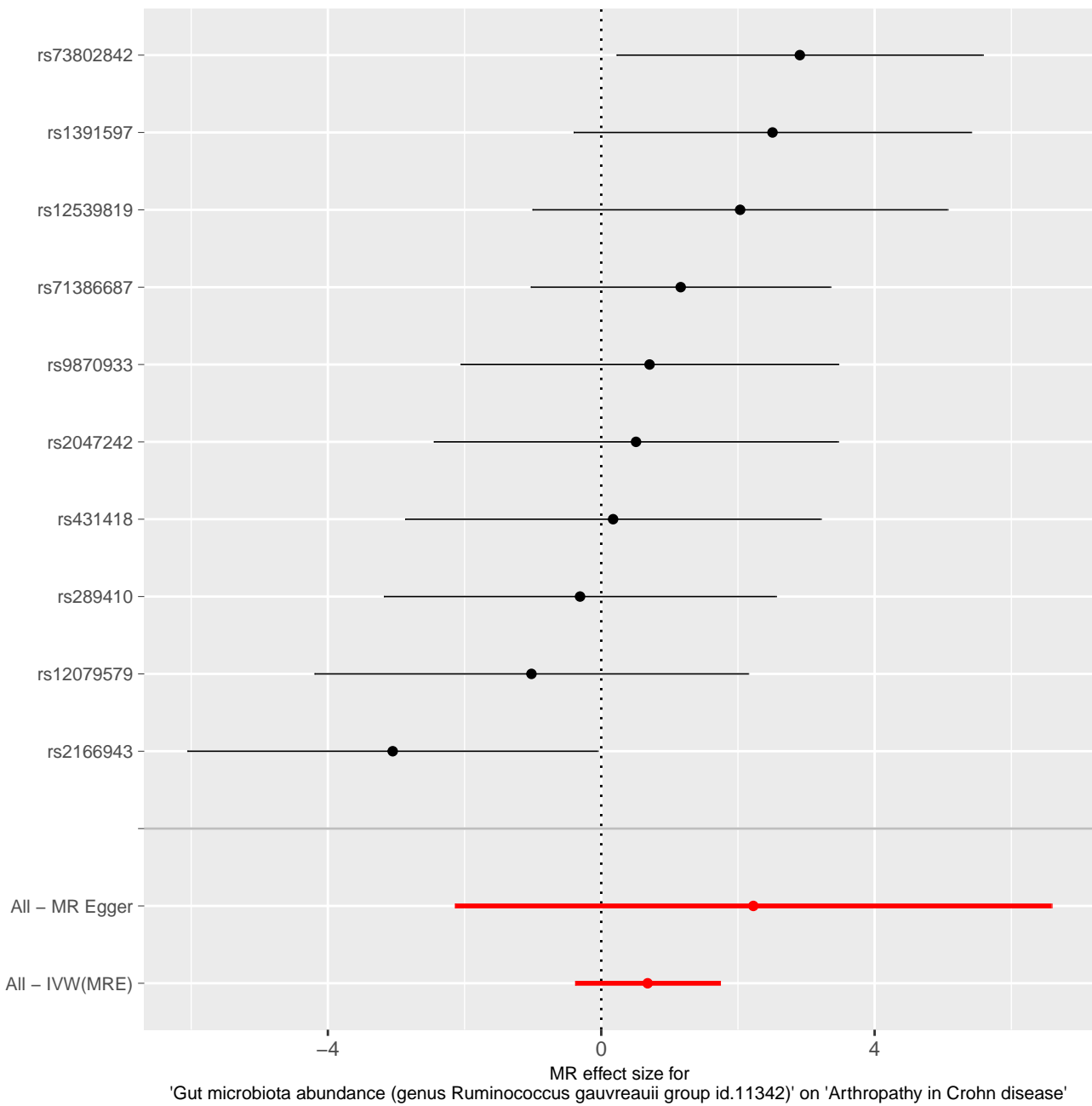




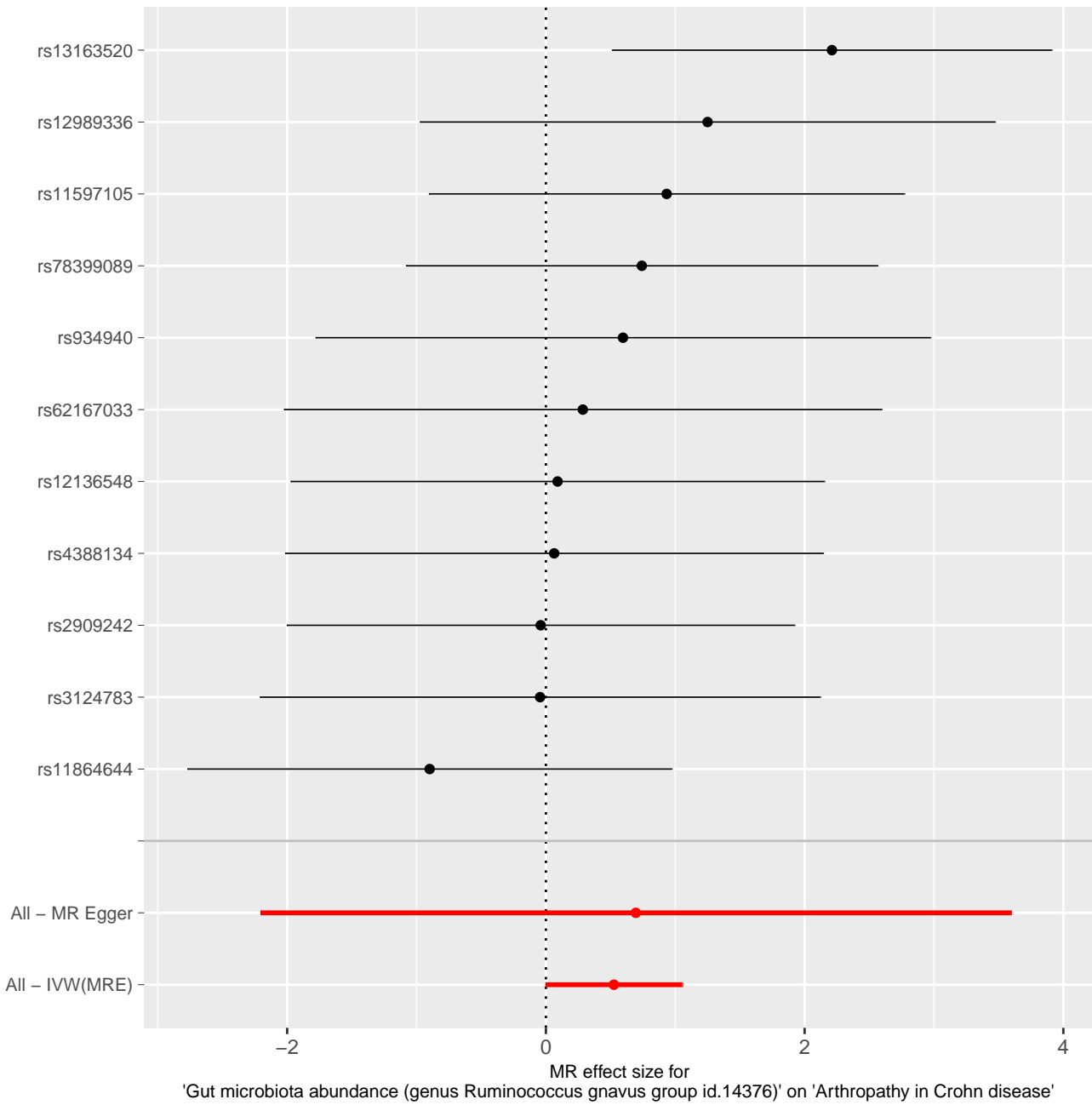


Batch 587 : Gut microbiota abundance (genus Ruminococcus2 id.11374) on Arthropathy in Crohn disease

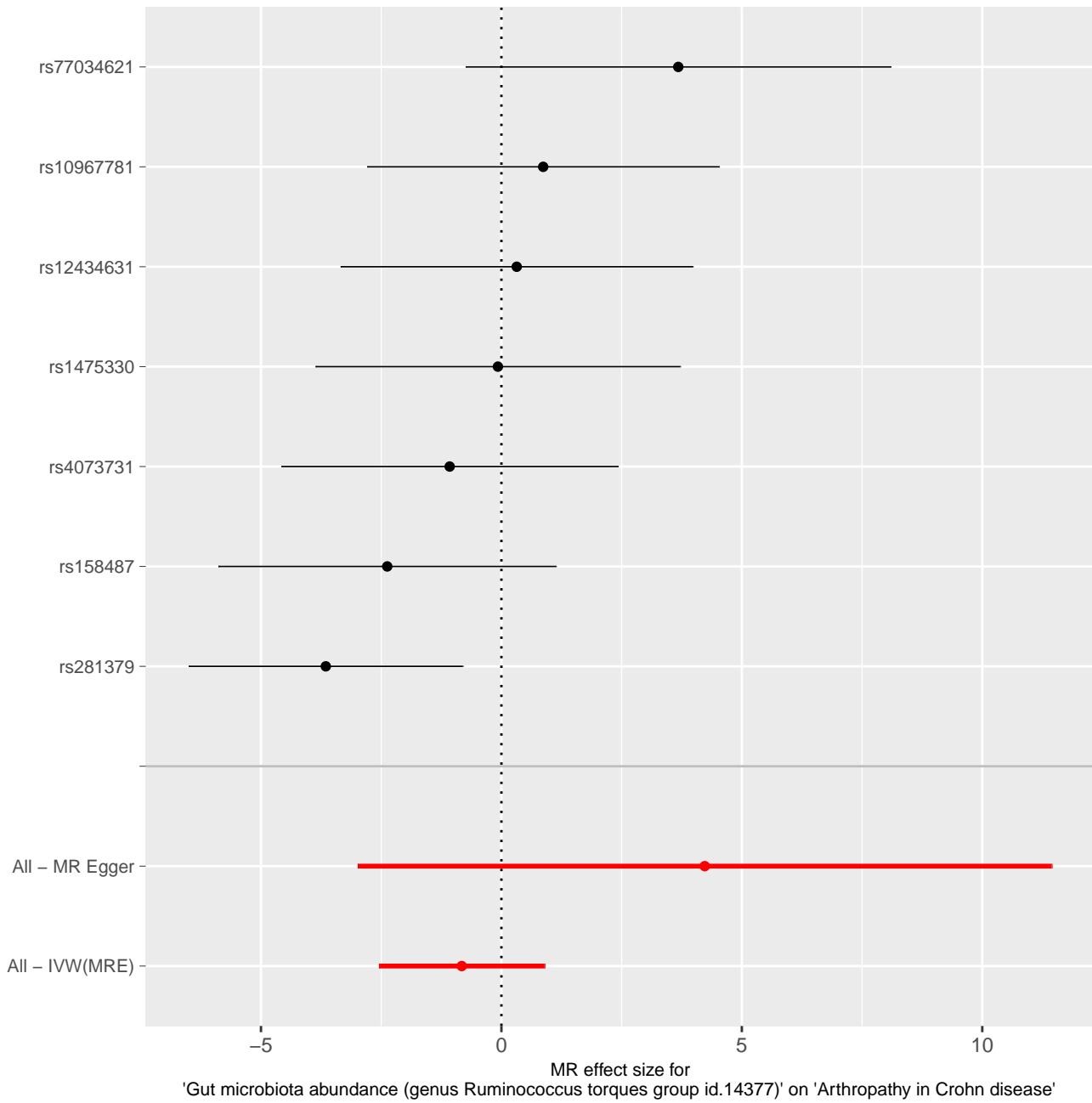


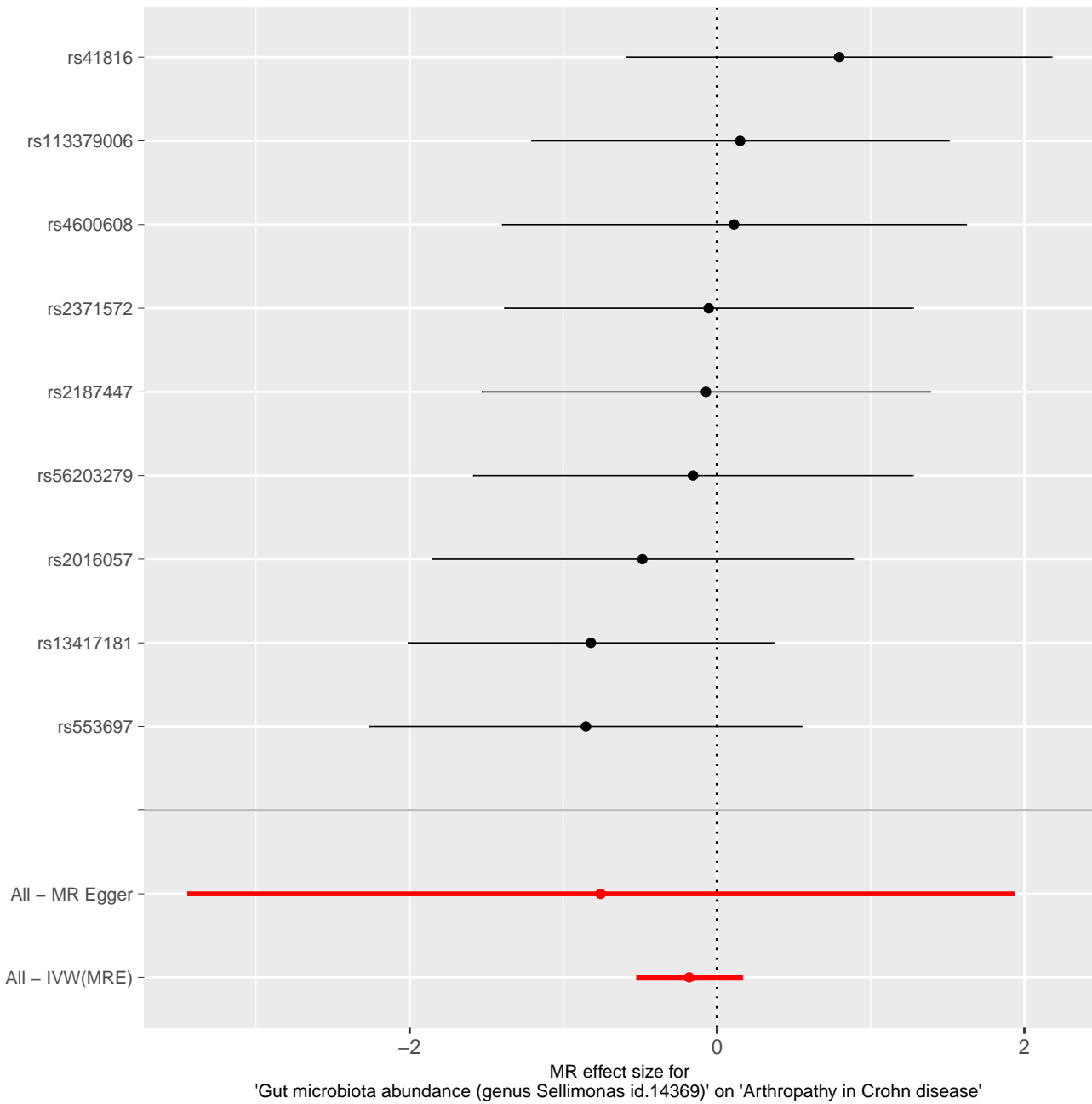


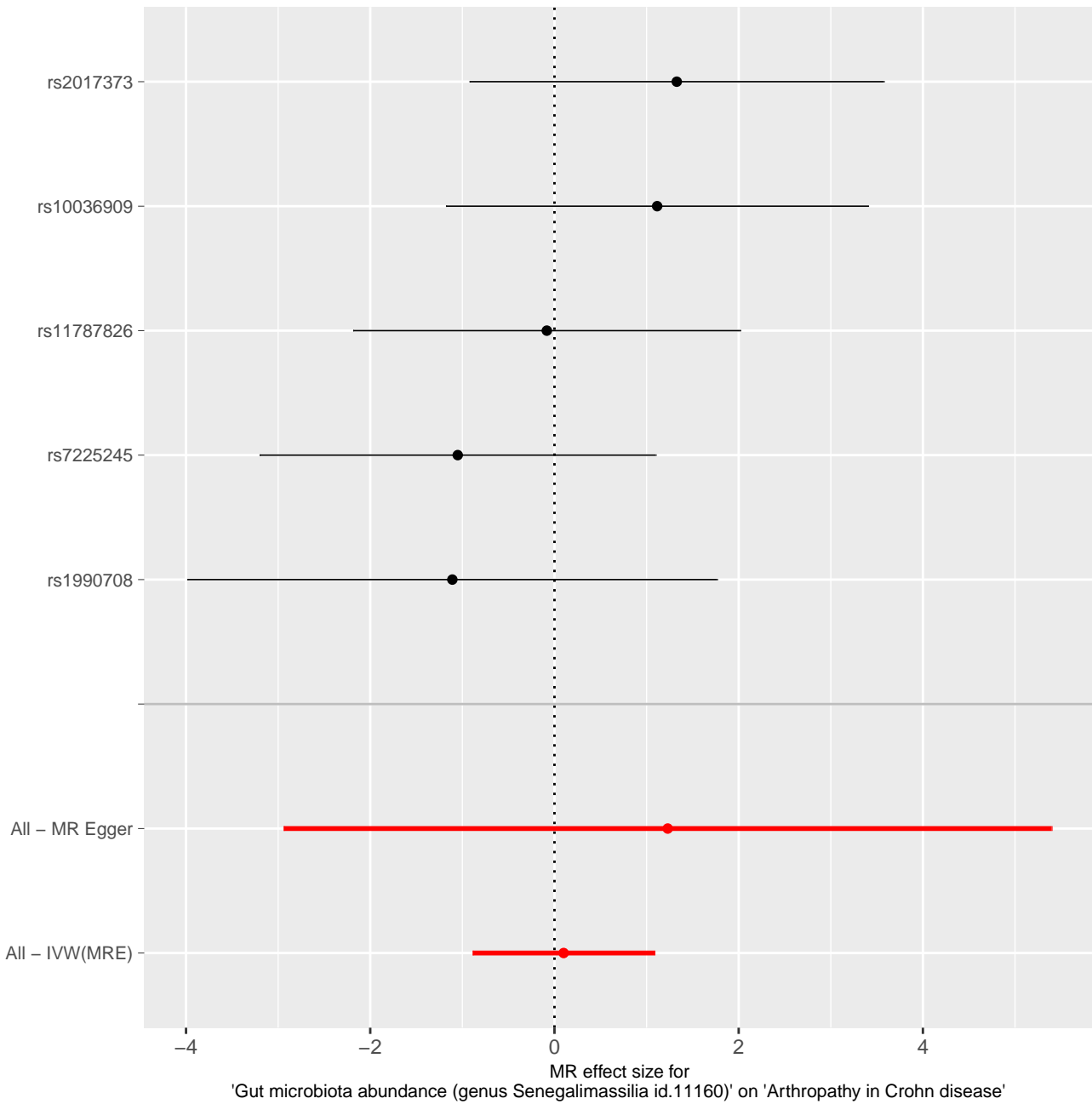
MR effect size for
'Gut microbiota abundance (genus Ruminococcus gauvreauii group id.11342)' on 'Arthropathy in Crohn disease'

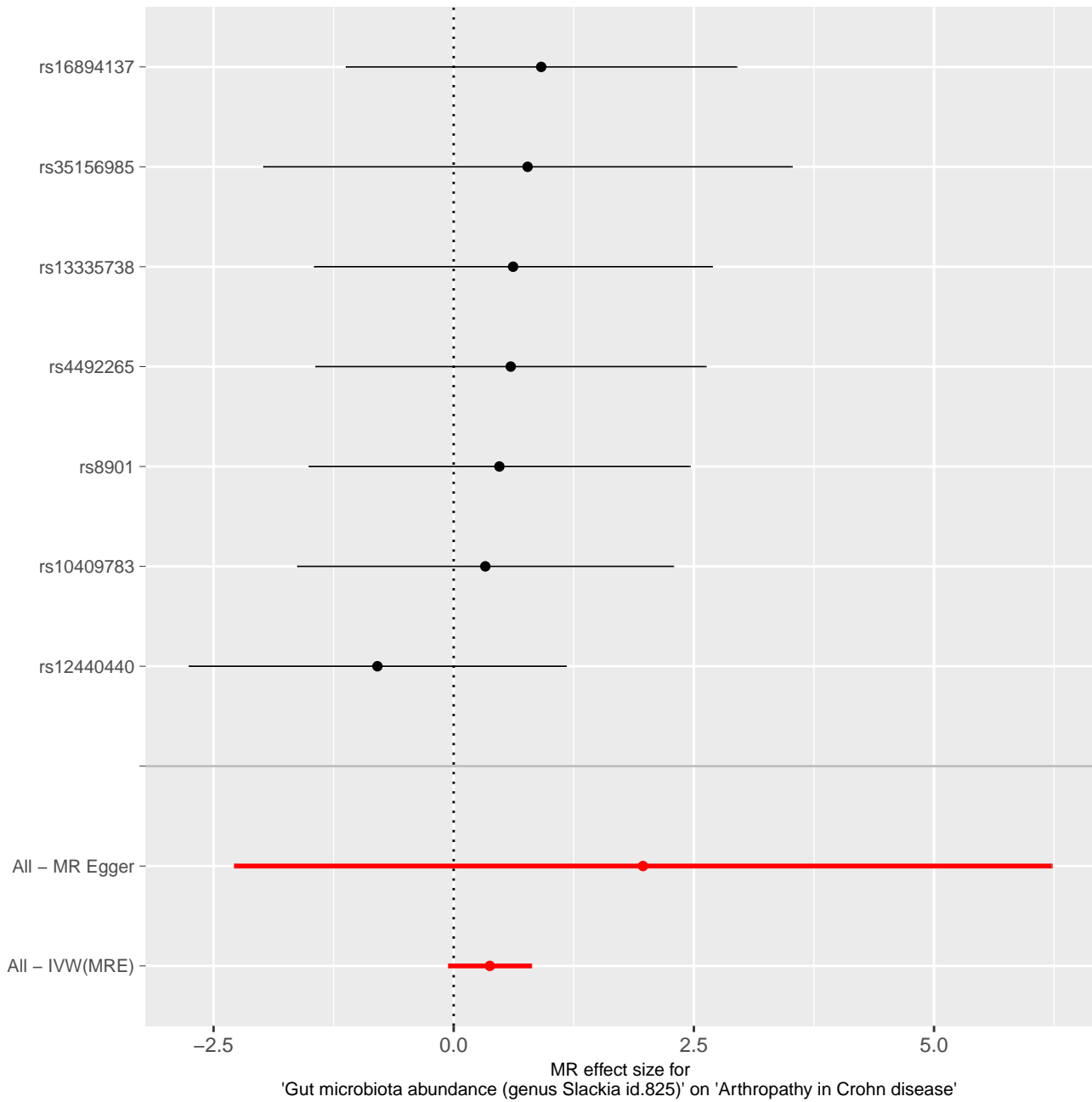


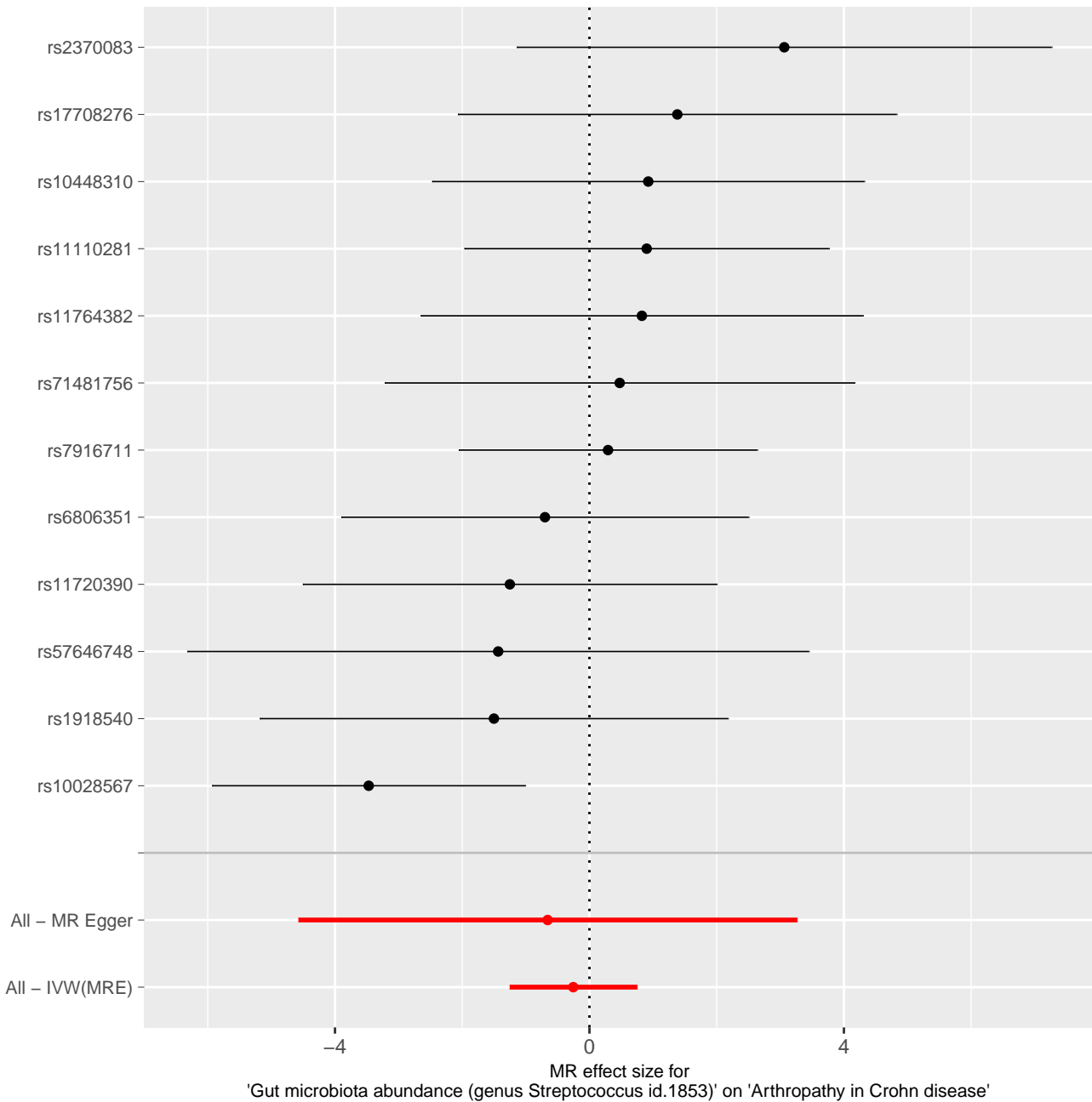
Batch 590 : Gut microbiota abundance (genus Ruminococcus torques group id.14377) on Arthropathy in Crohn disease

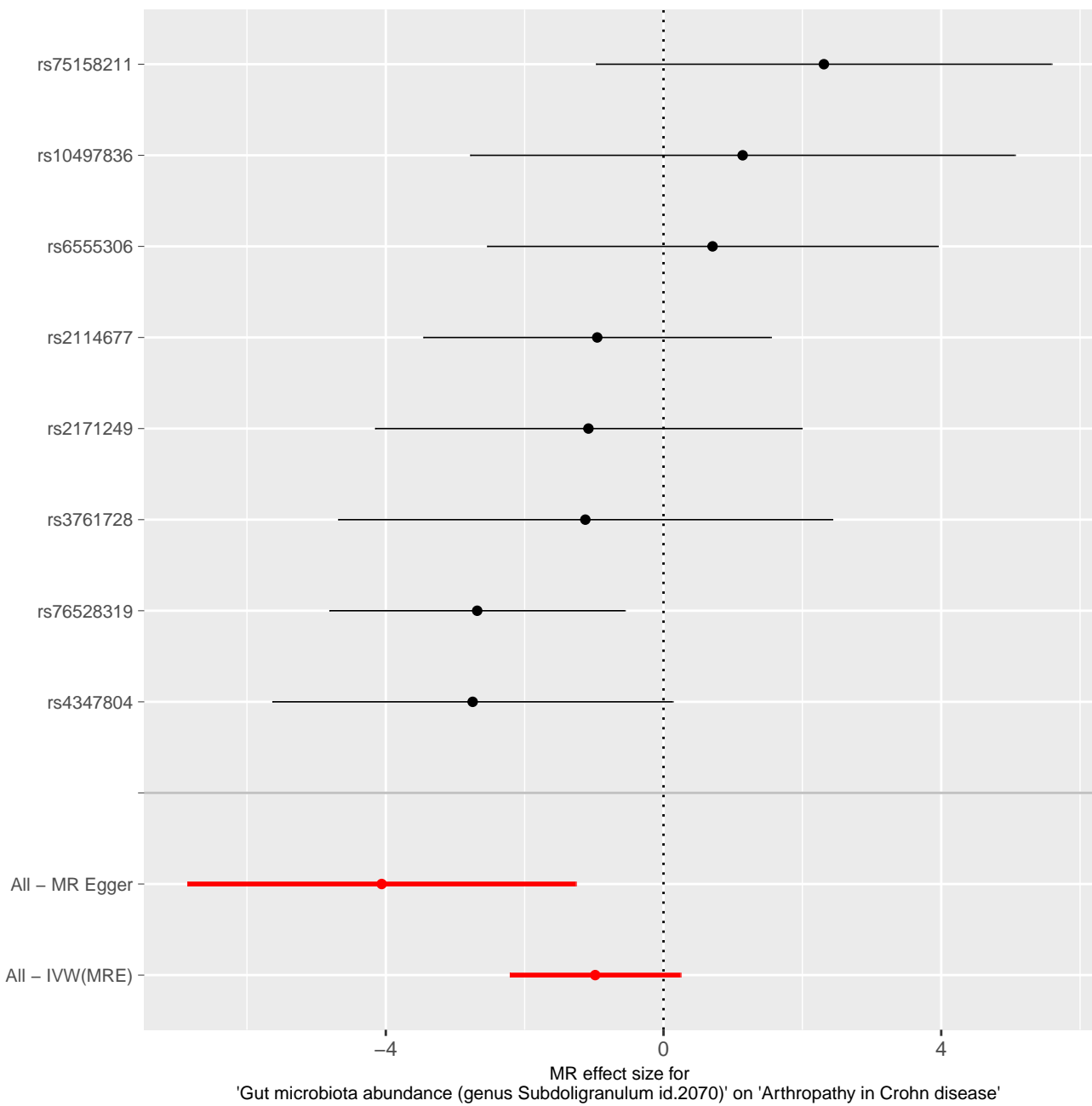




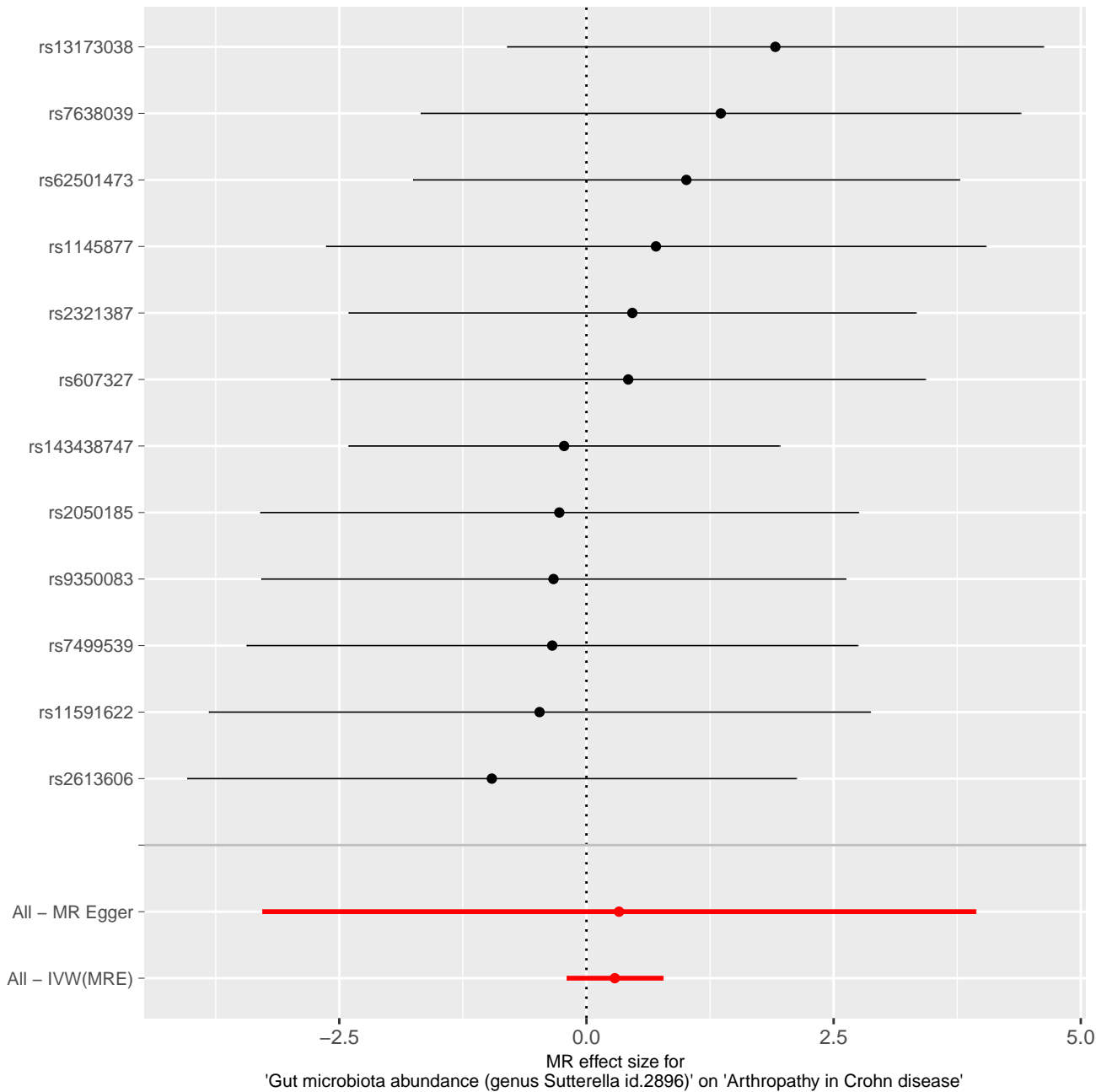


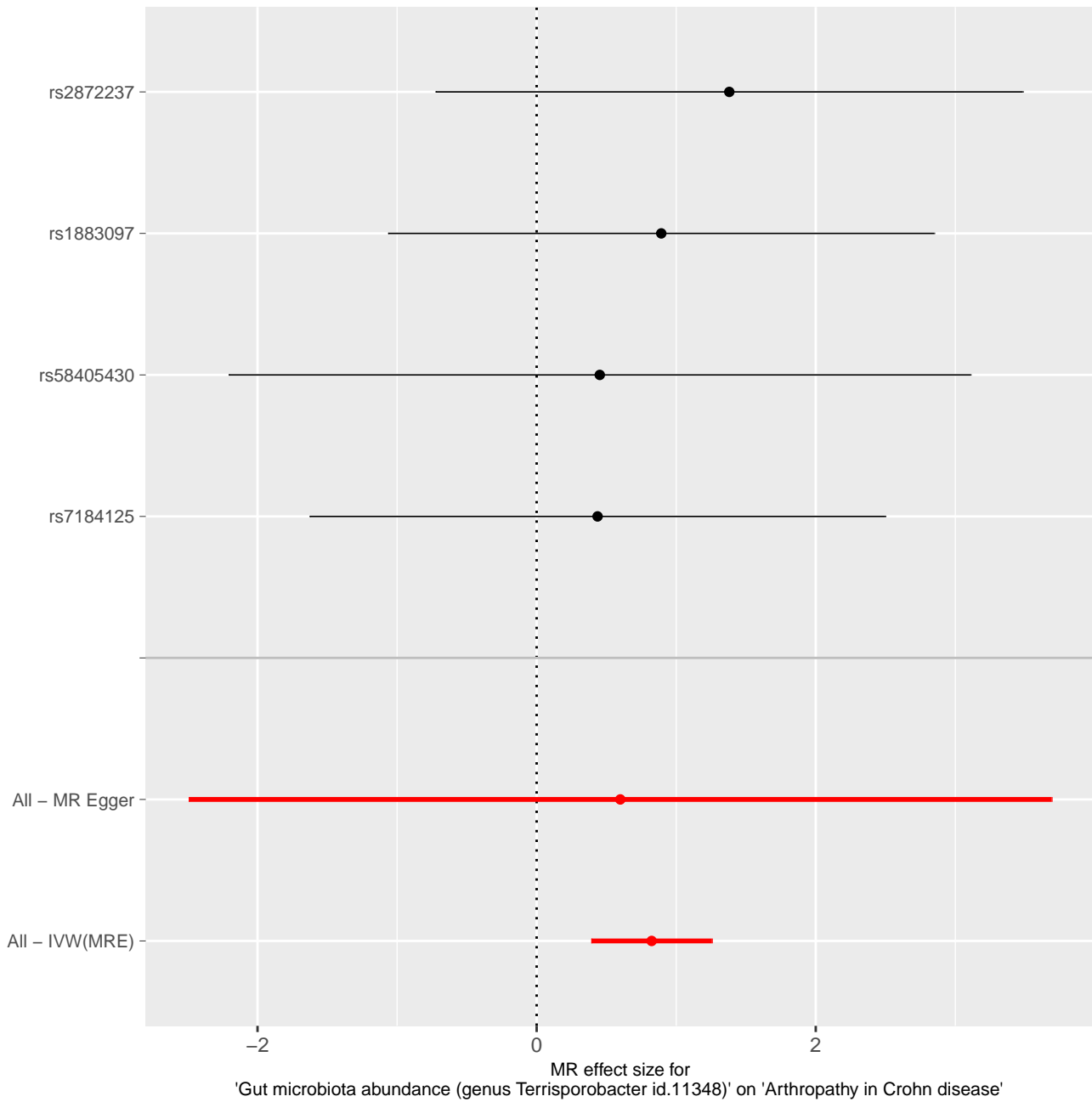


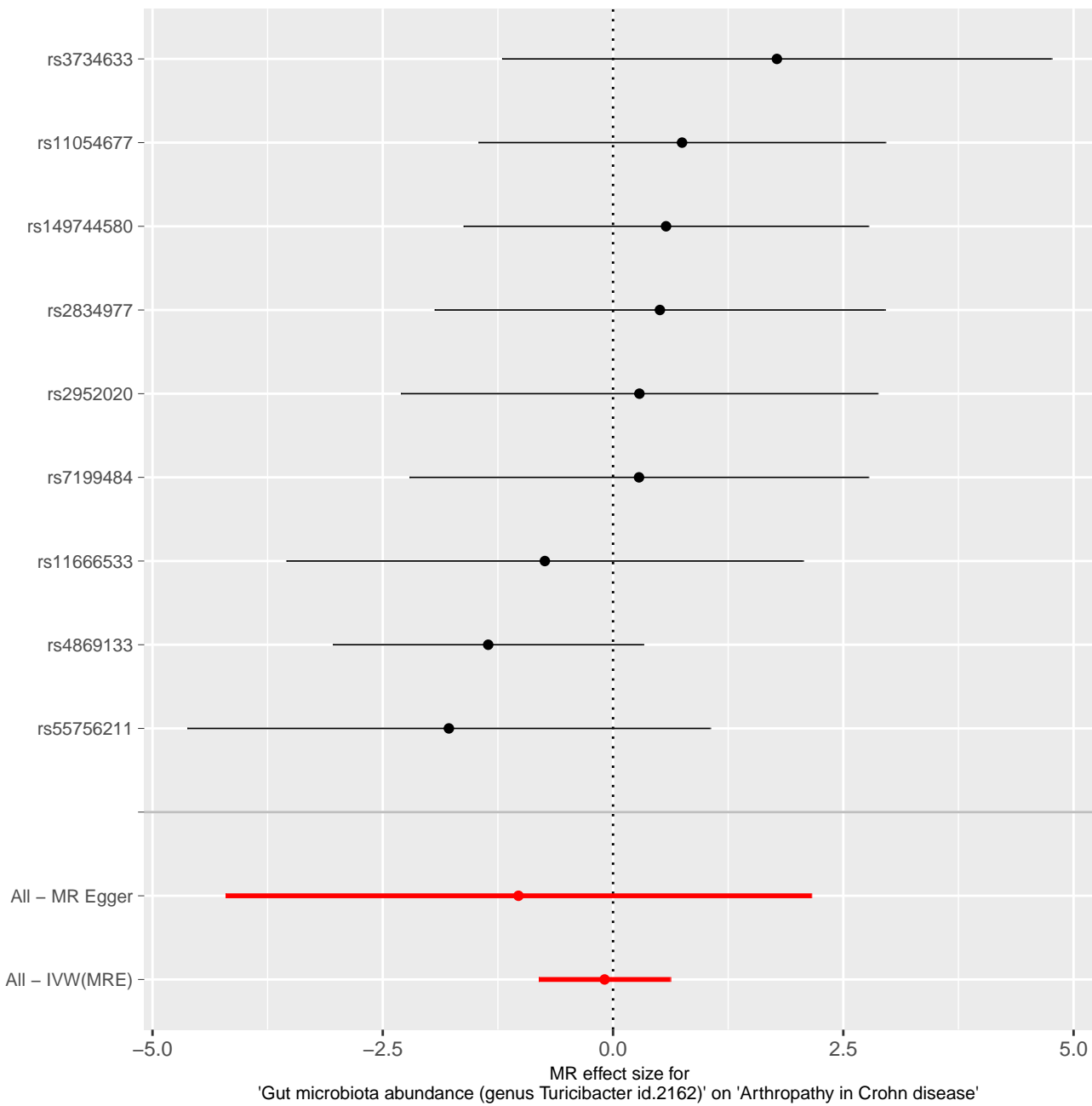




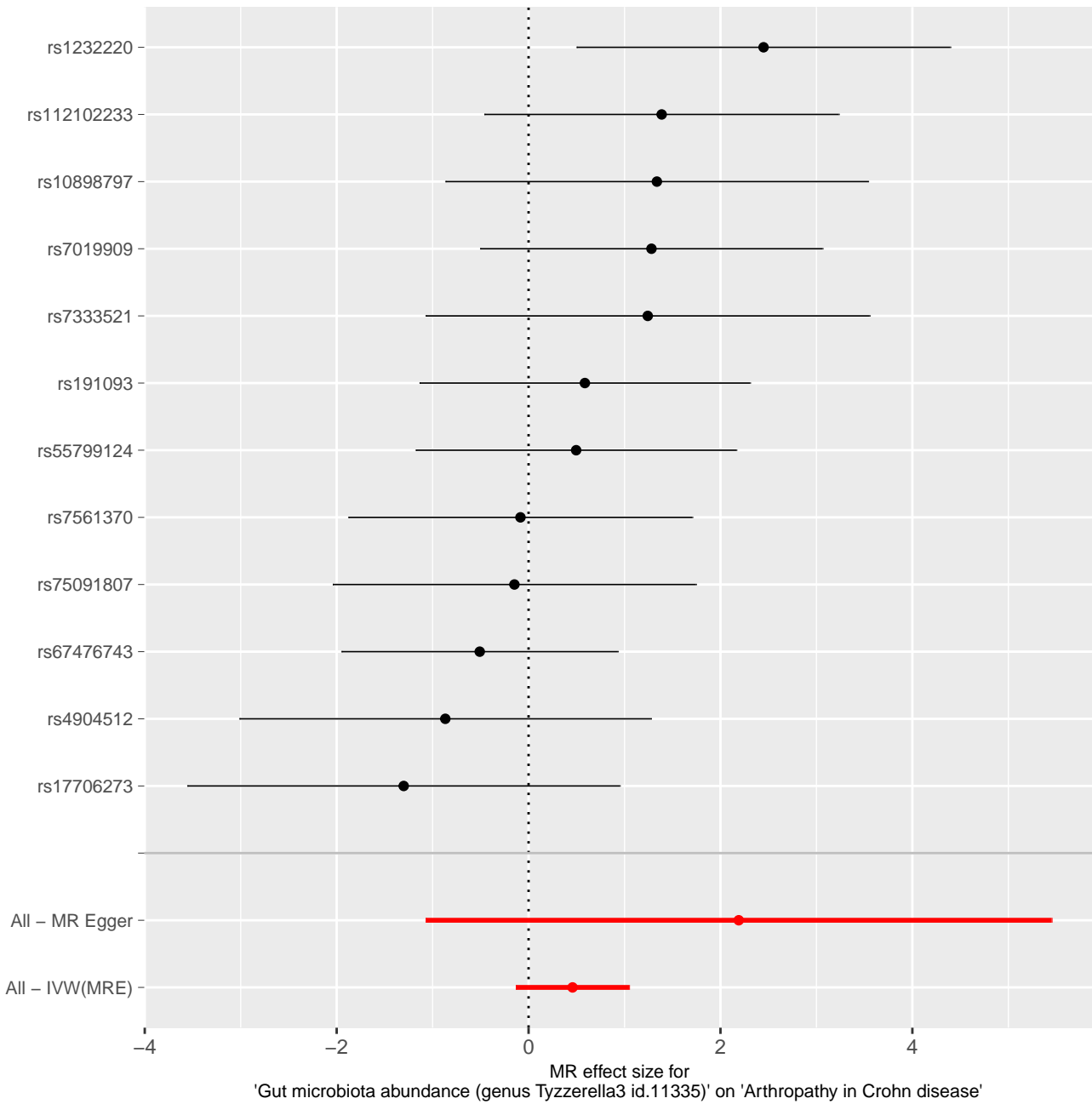
Batch 596 : Gut microbiota abundance (genus Sutterella id.2896) on Arthropathy in Crohn disease

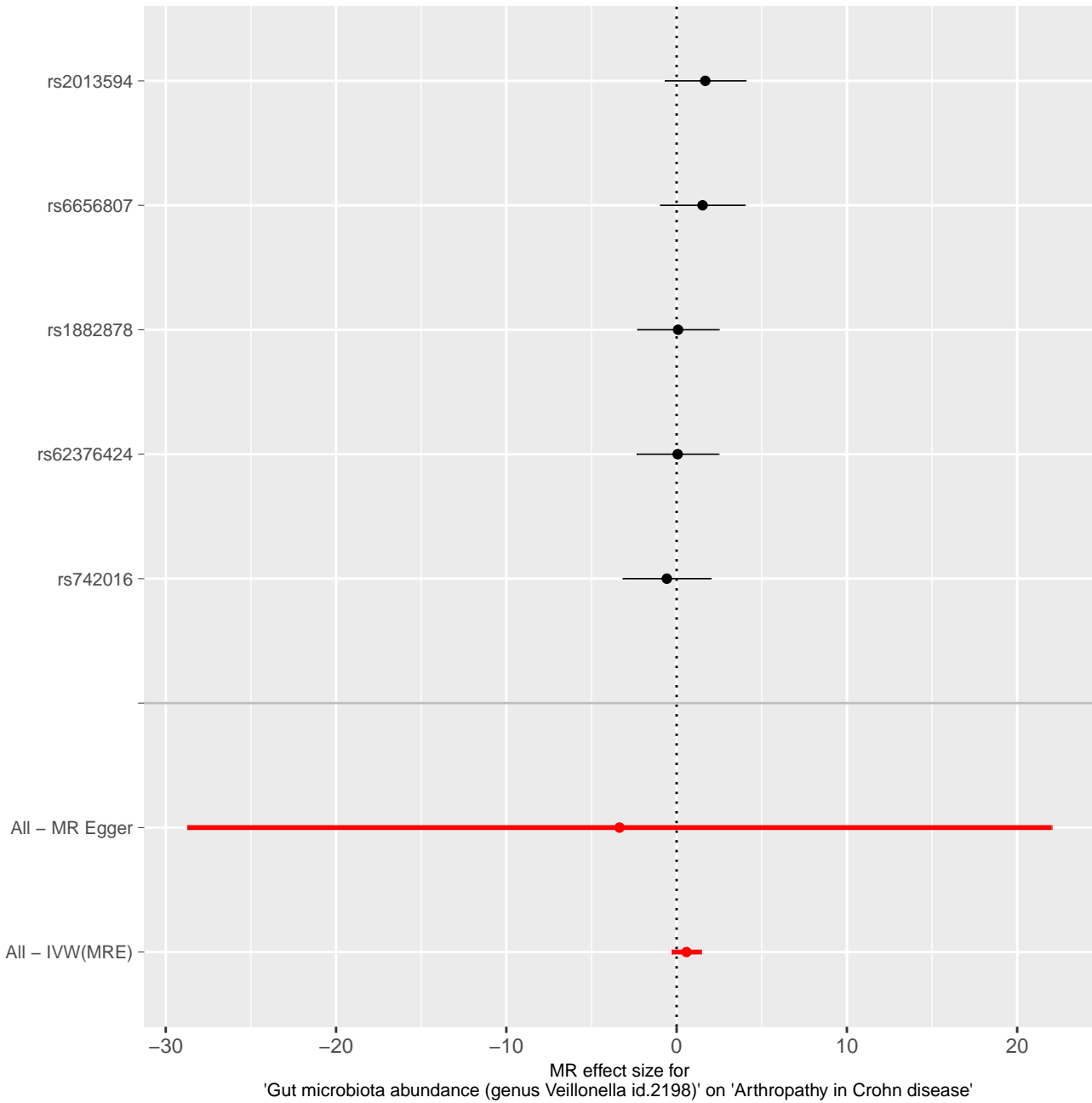


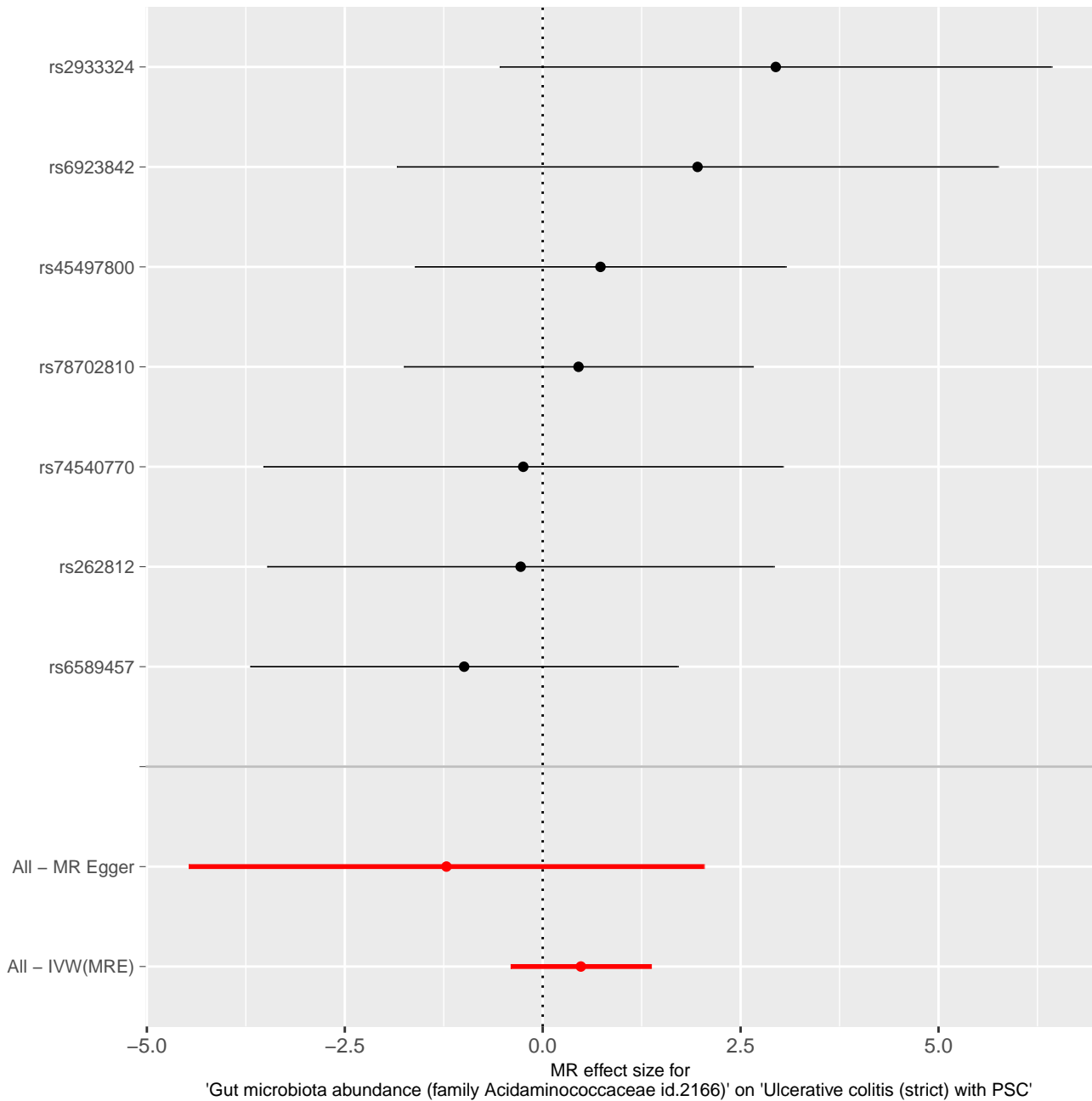


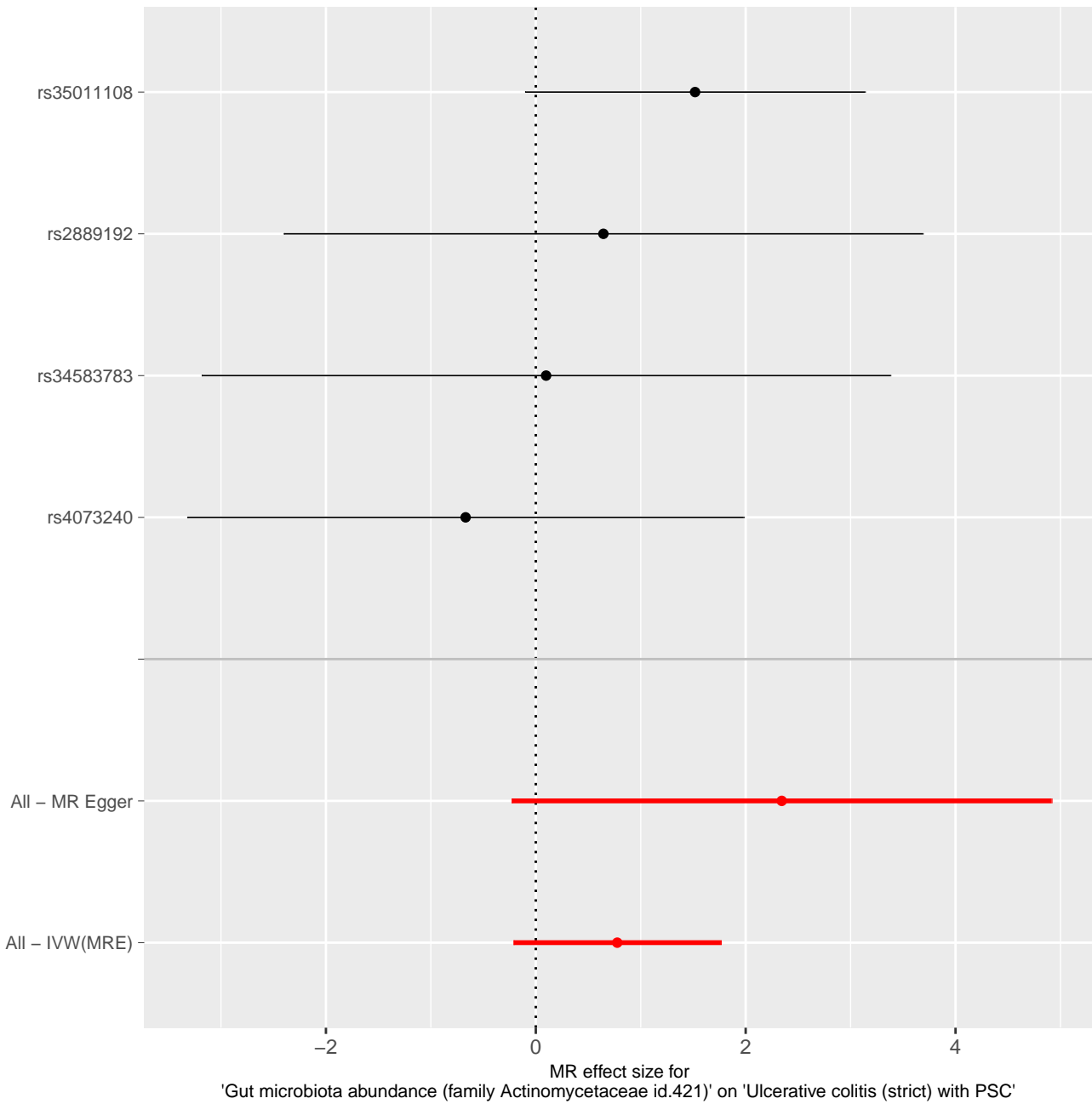


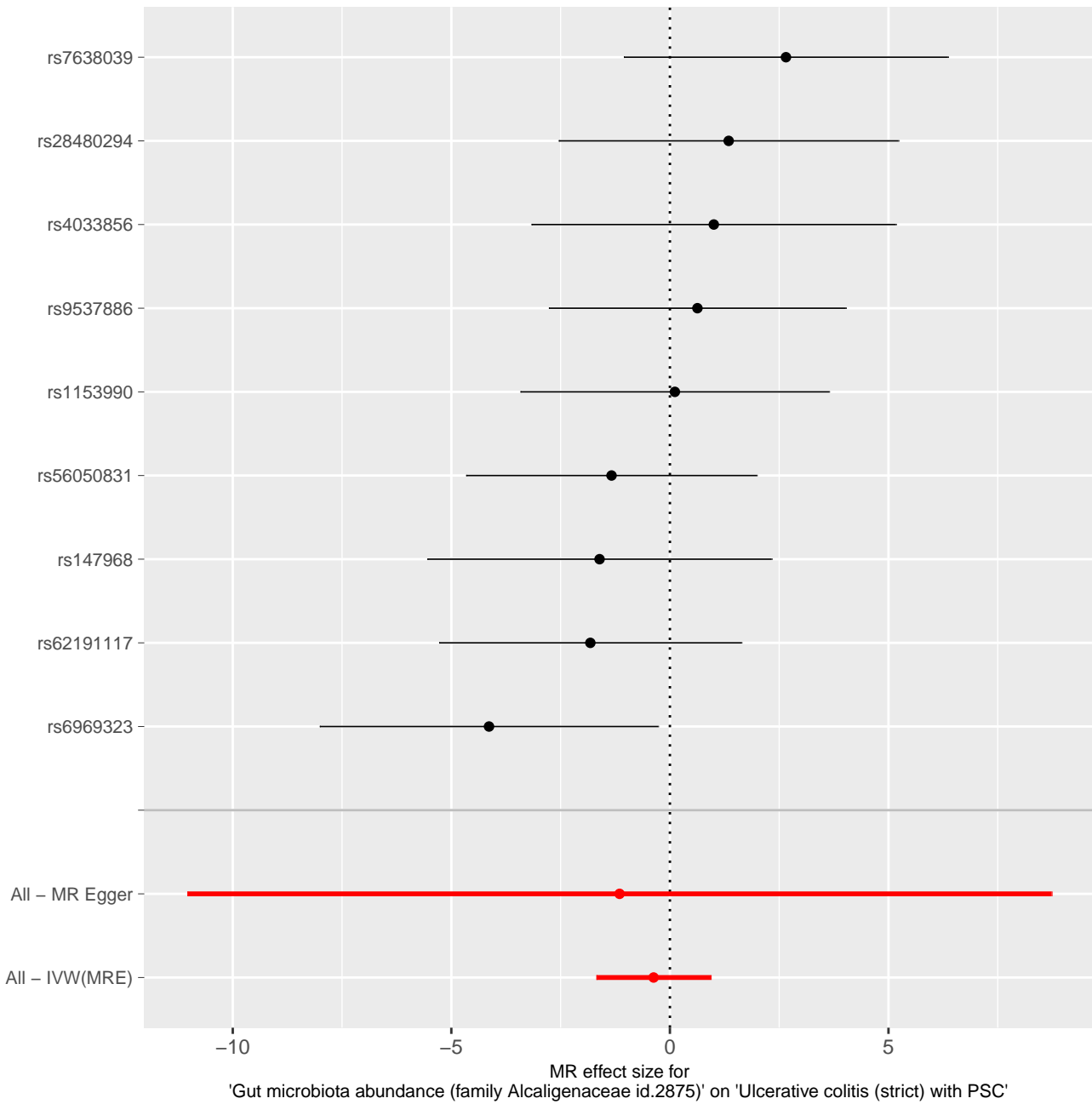
Batch 599 : Gut microbiota abundance (genus Tyzzerella3 id.11335) on Arthropathy in Crohn disease

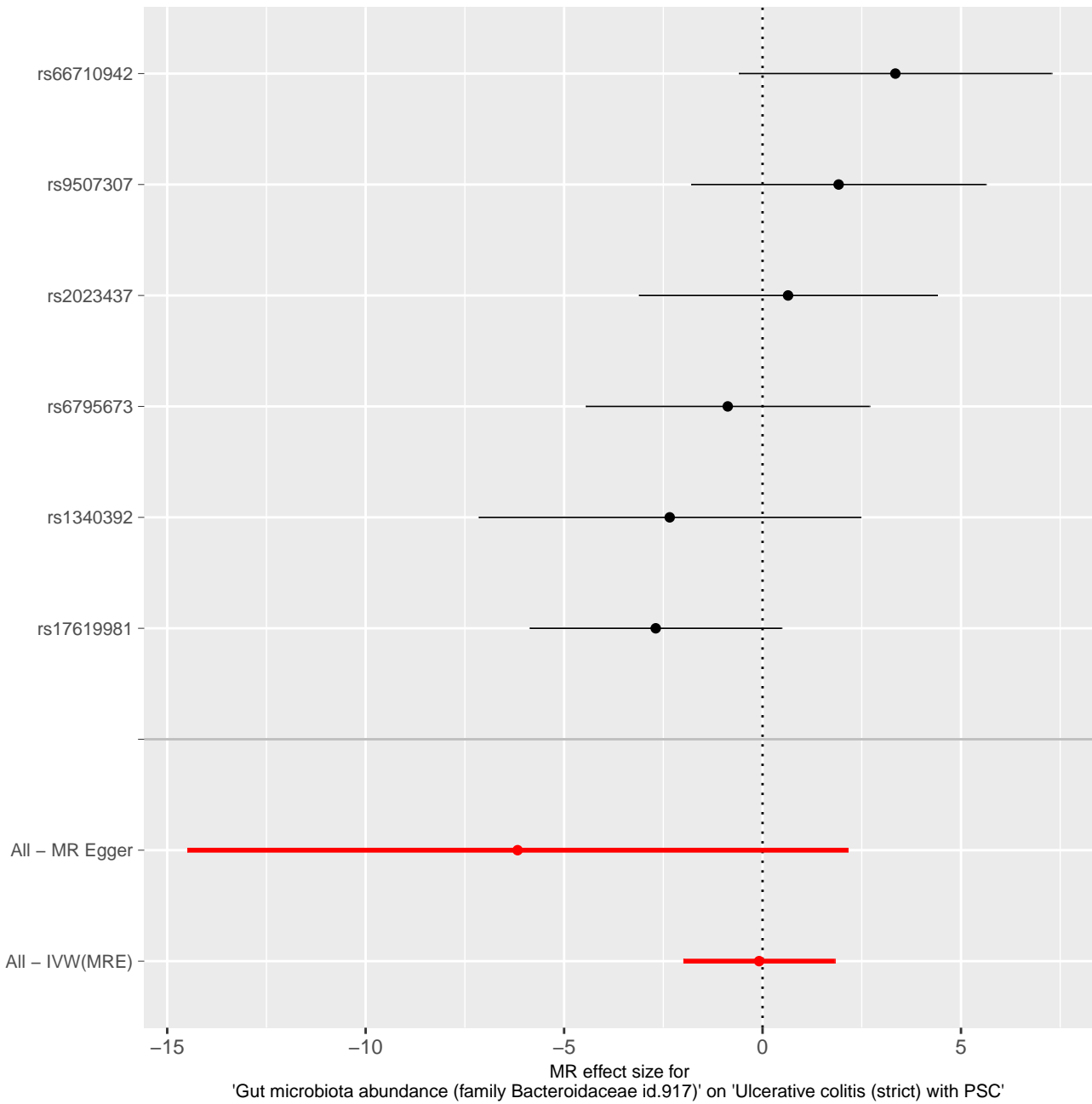


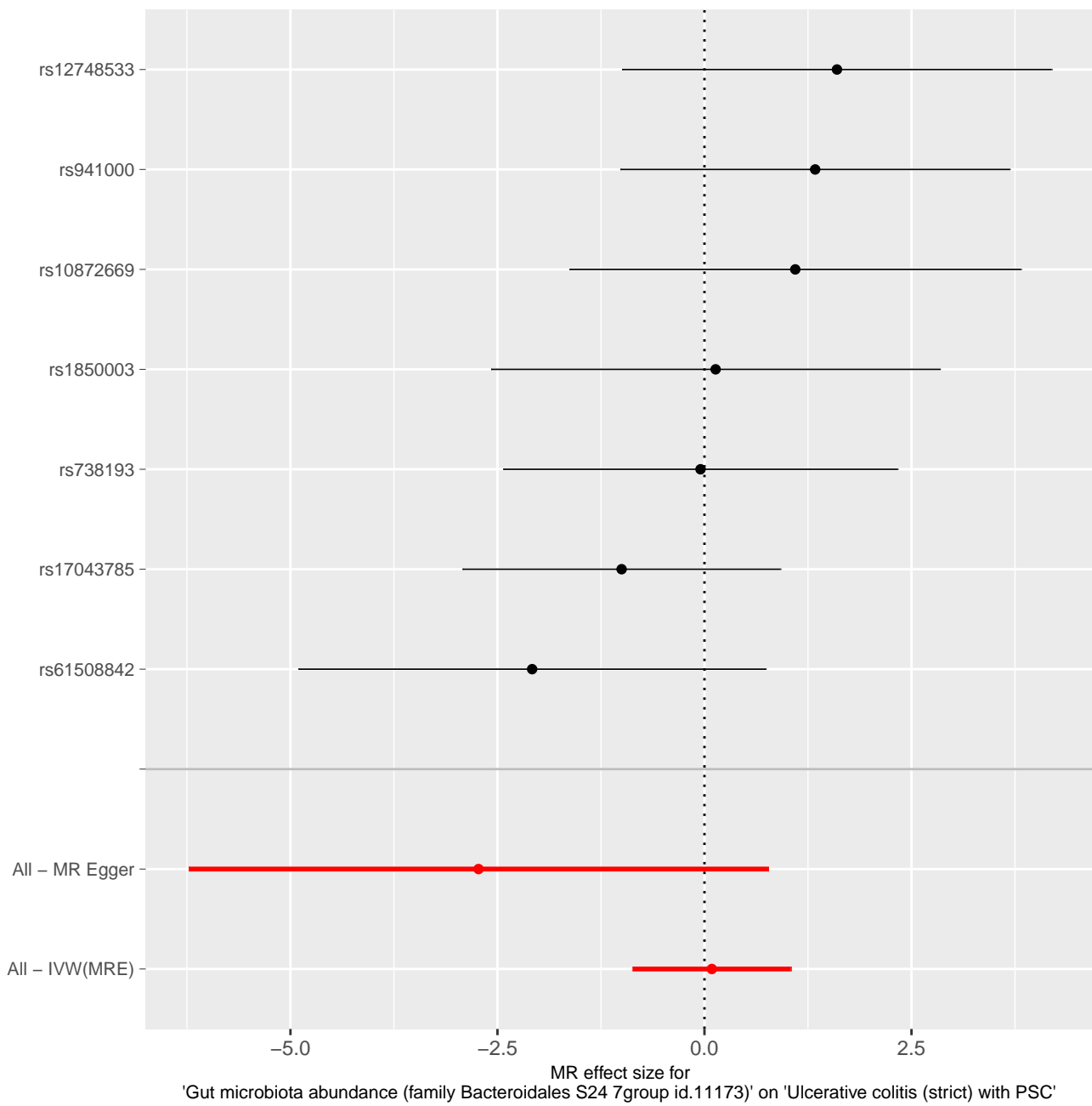


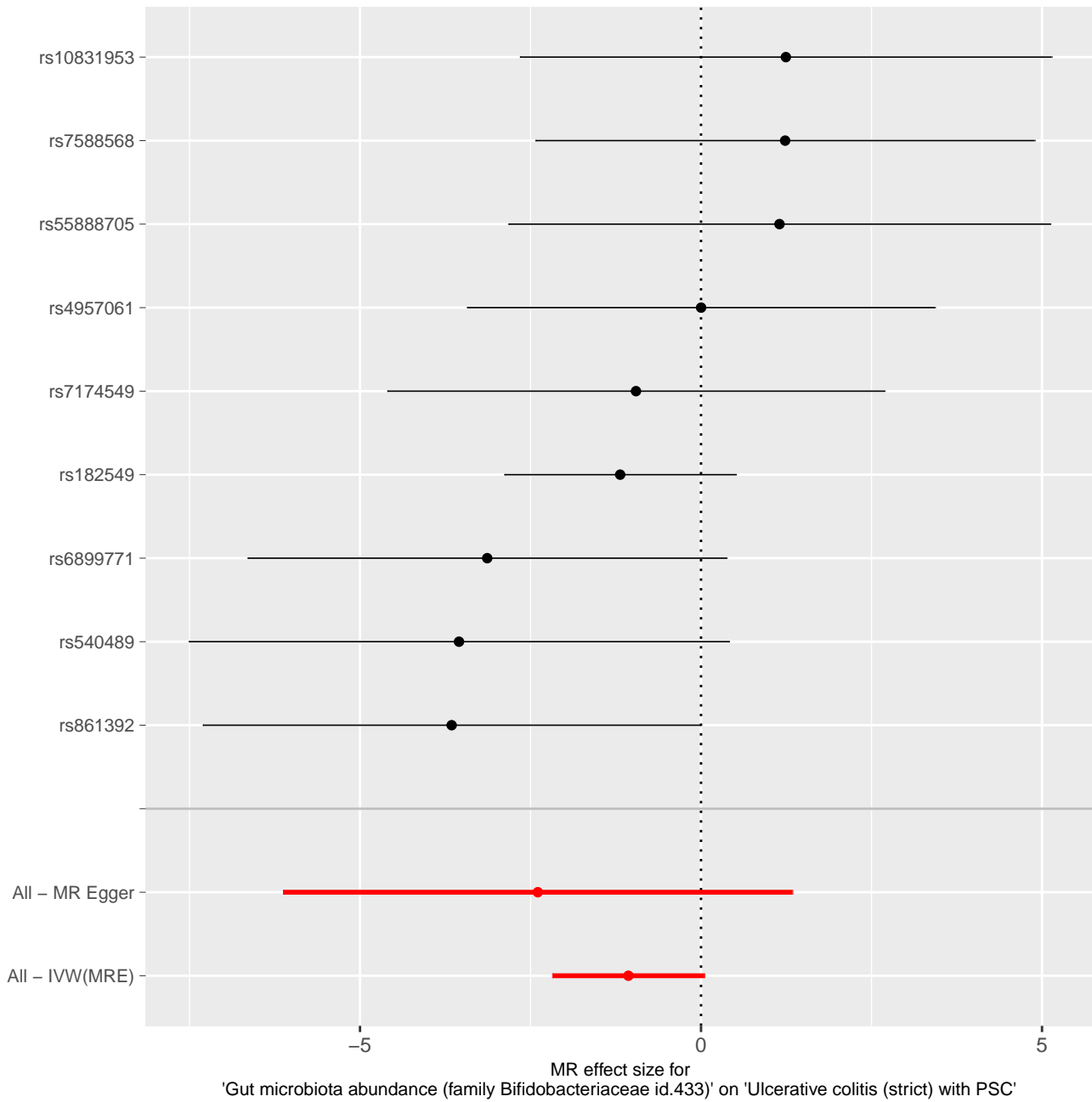


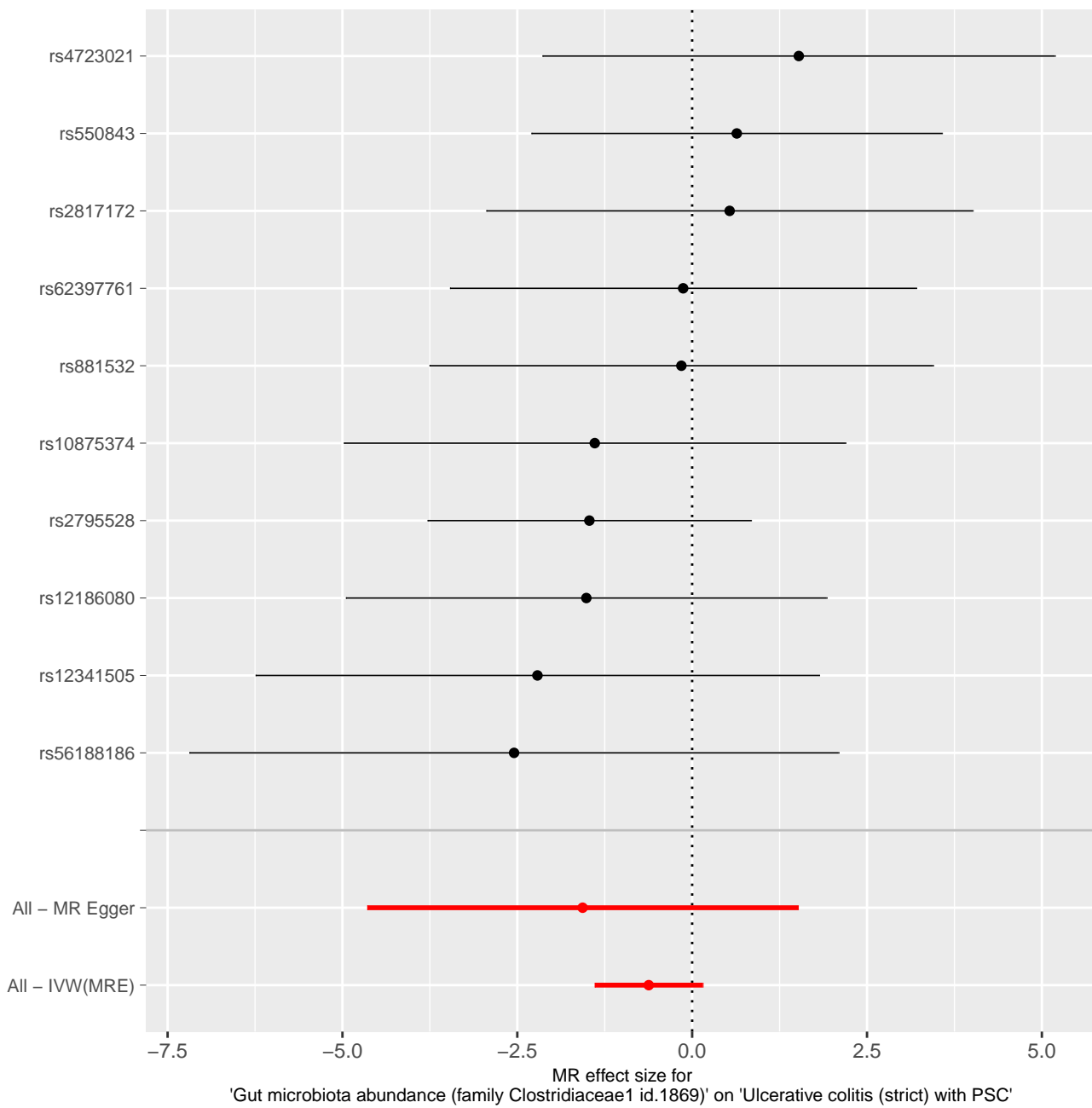


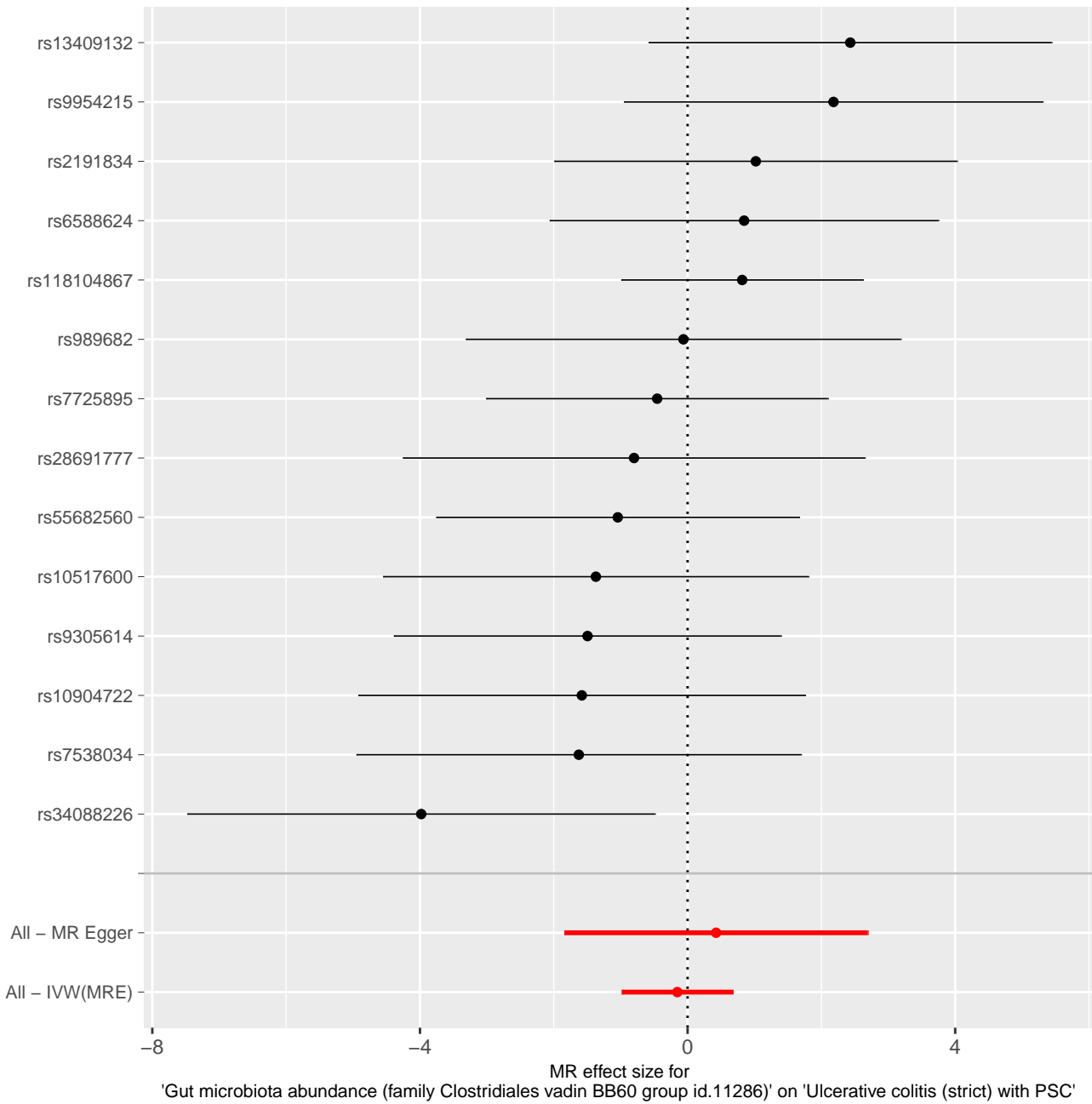


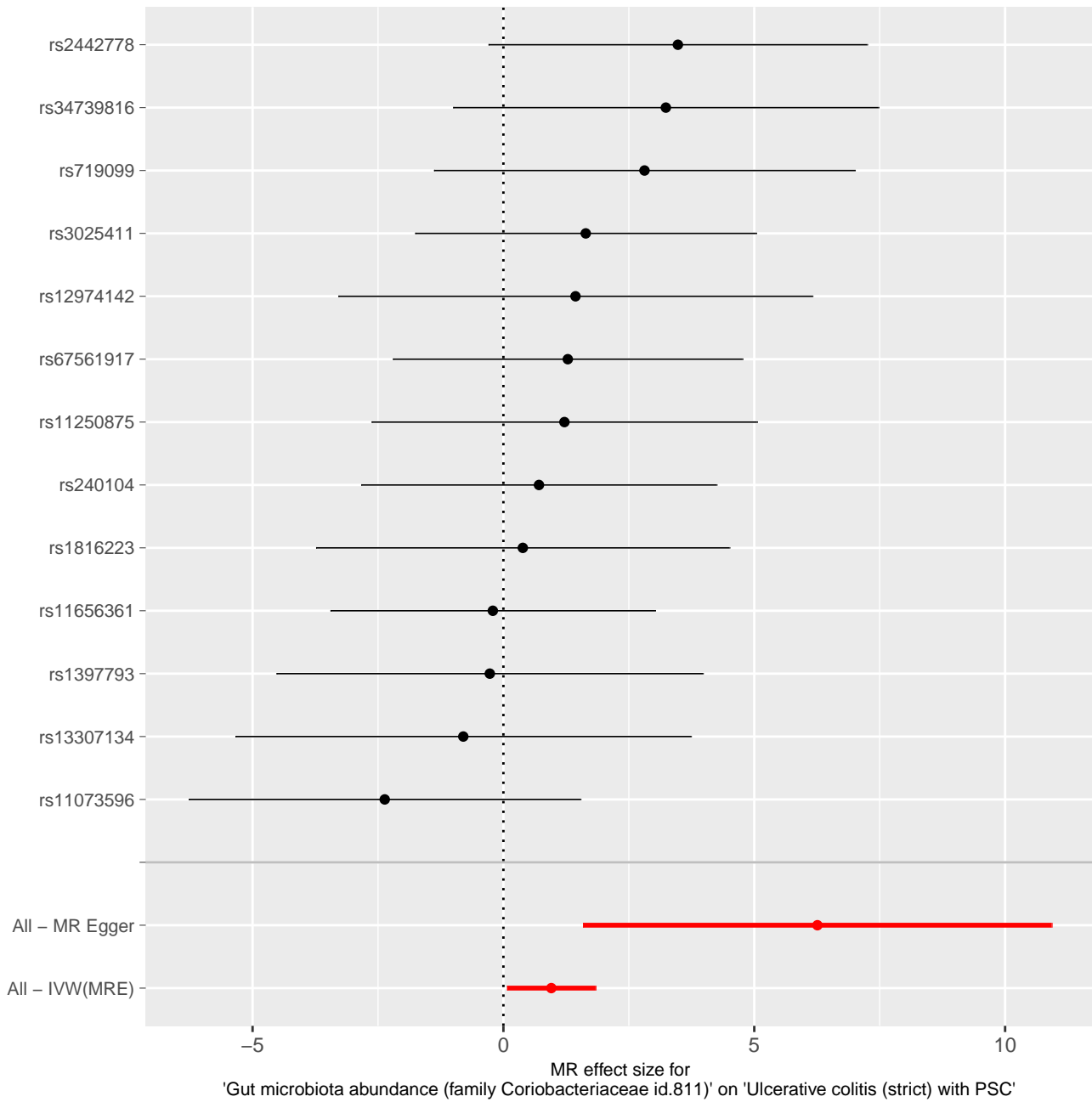


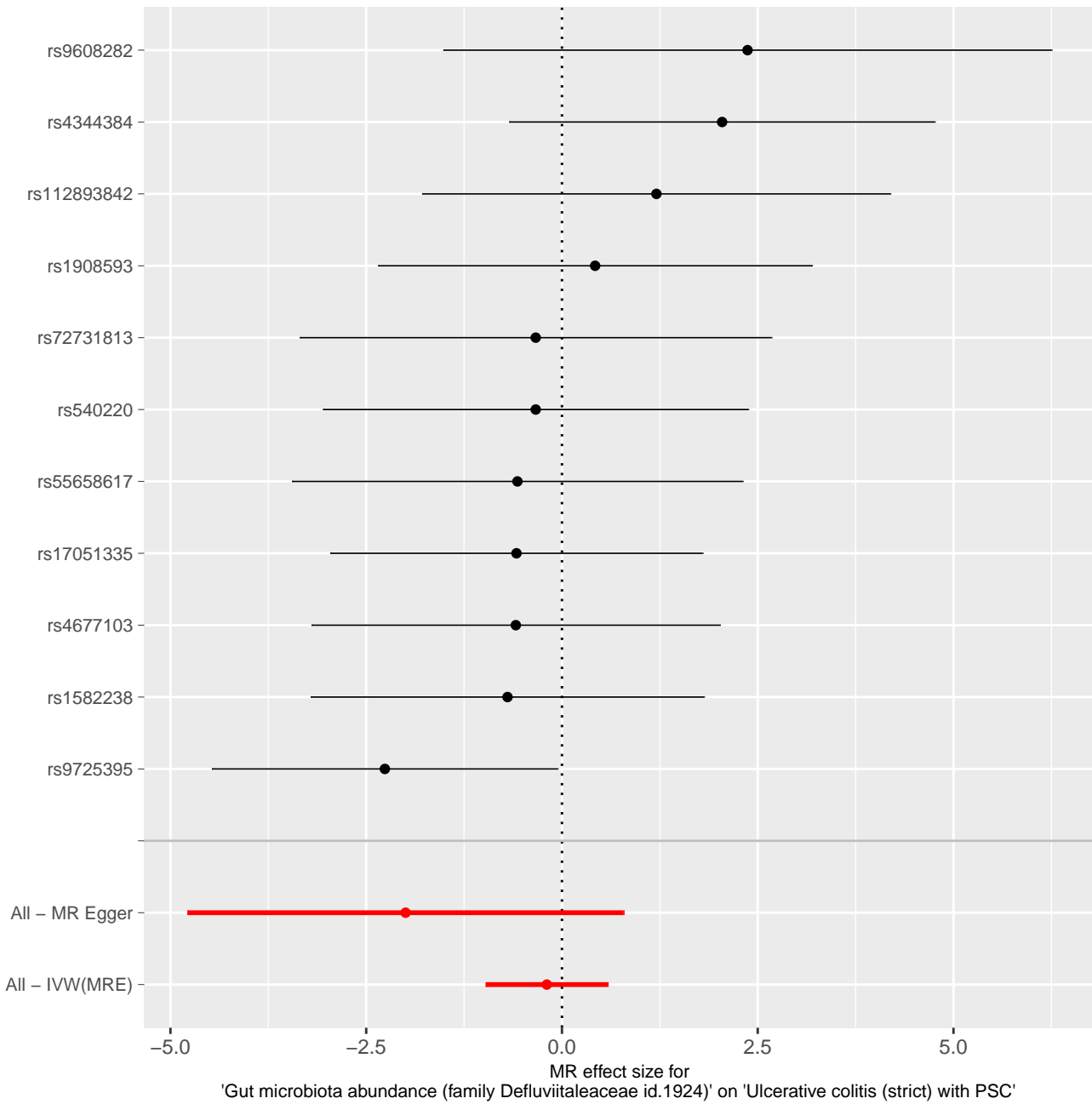


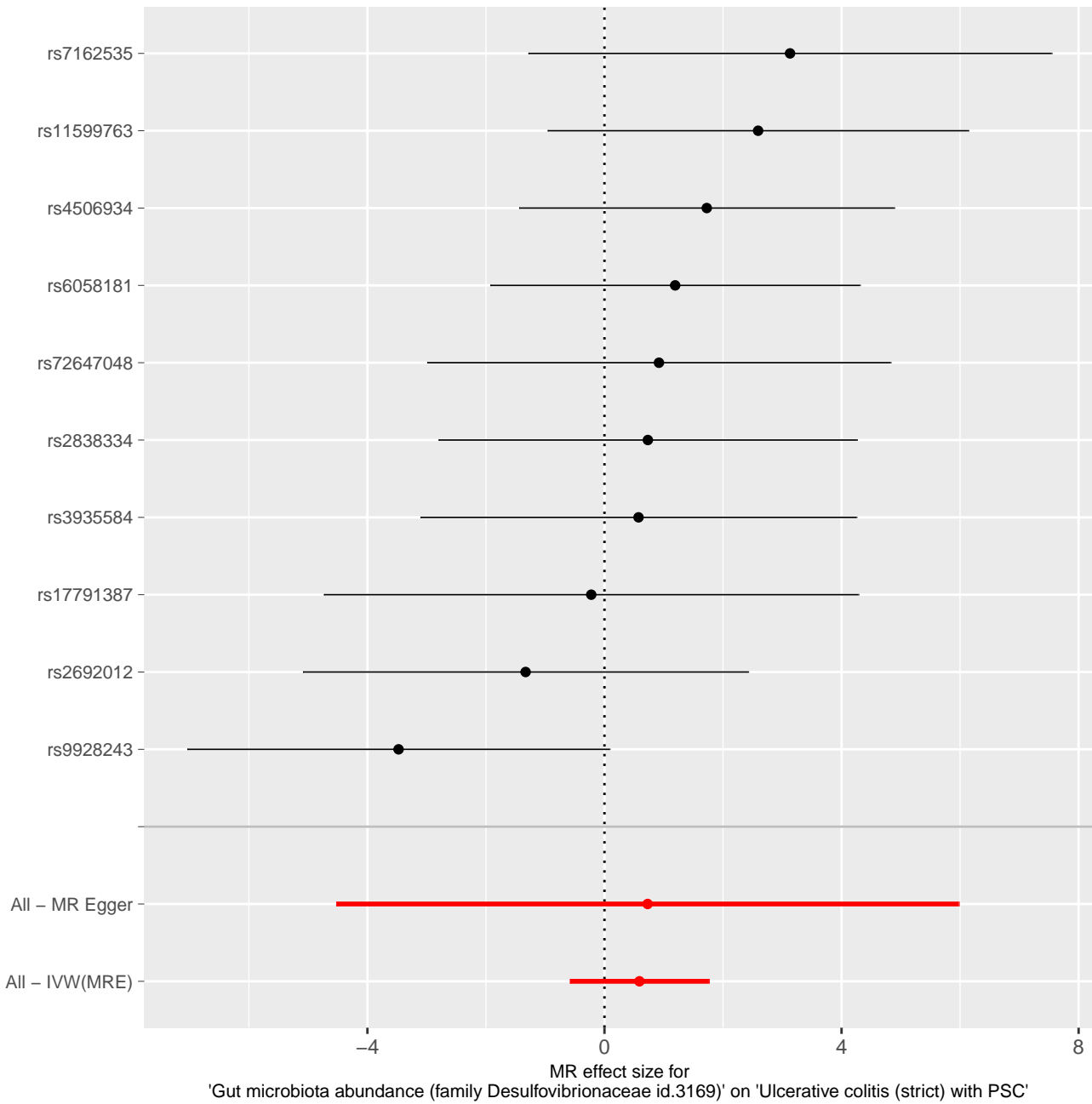




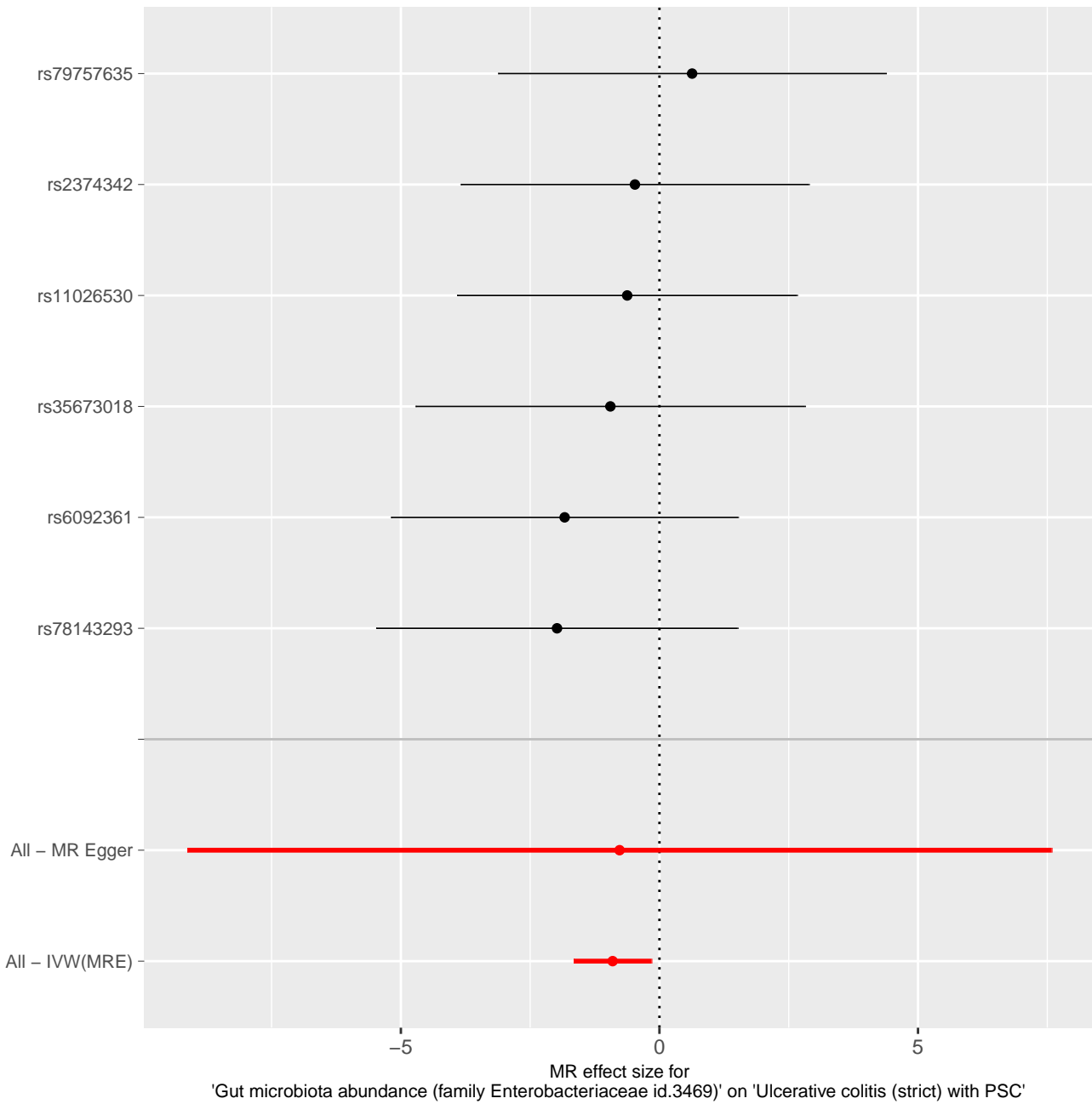


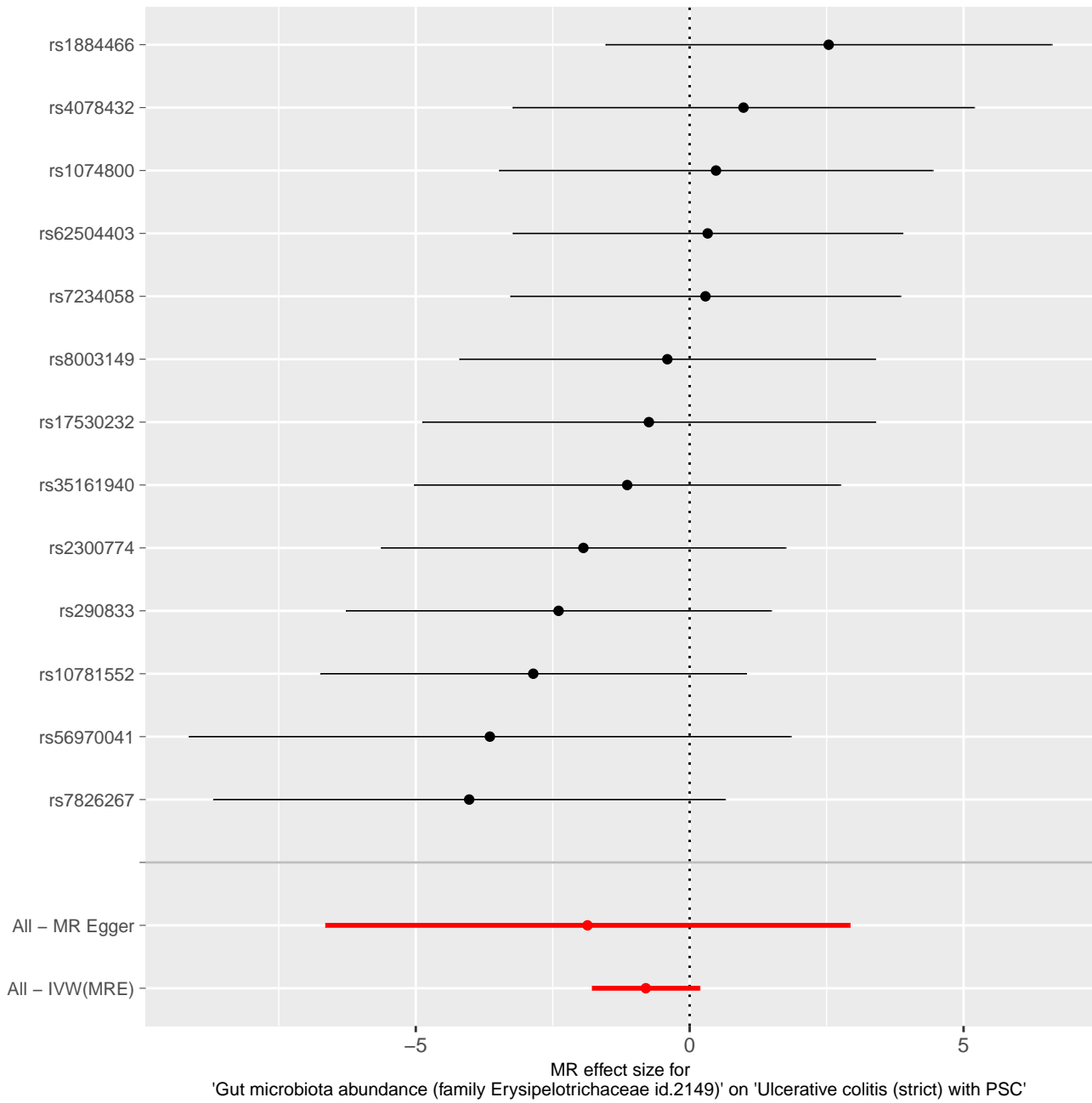


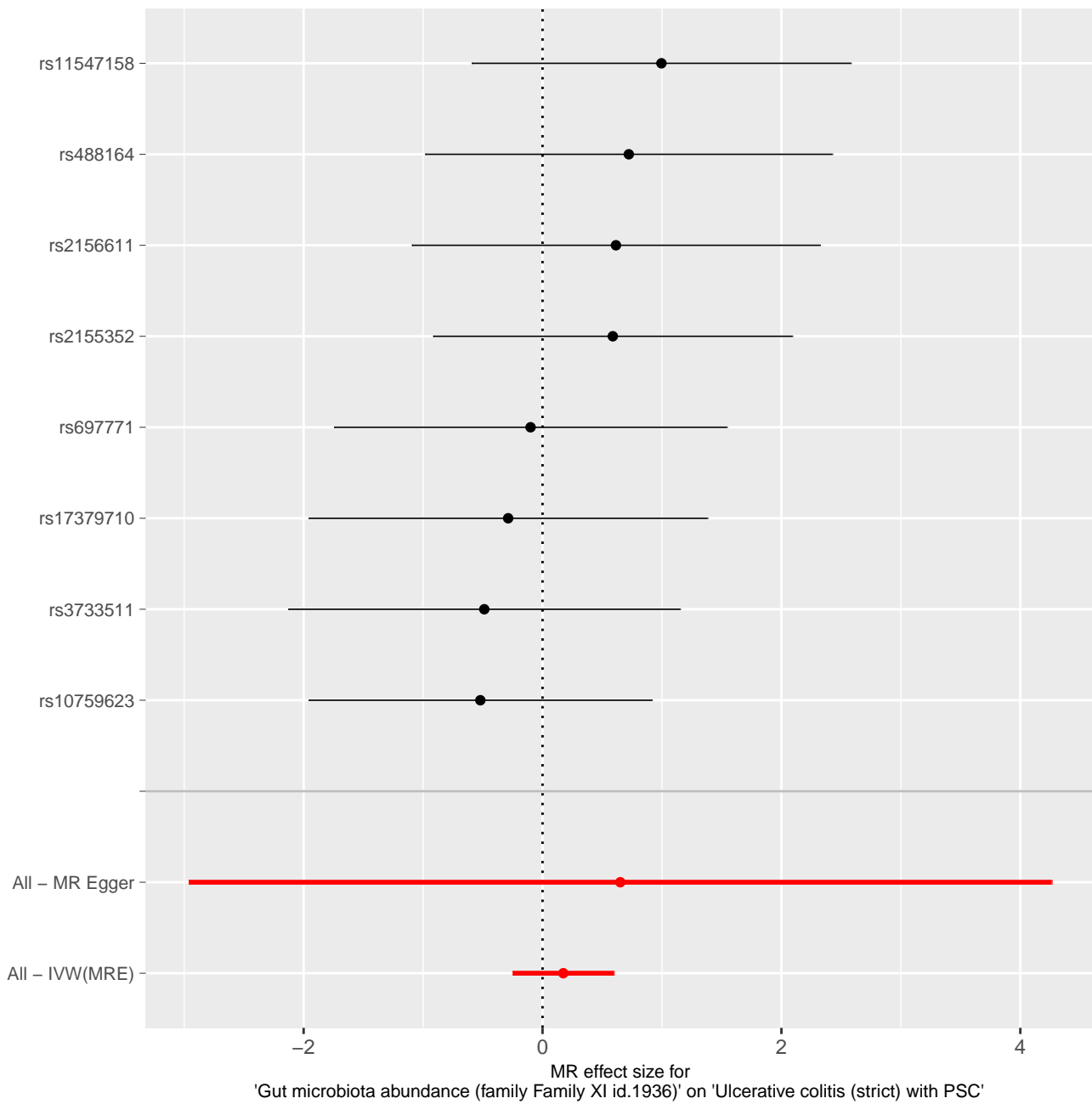


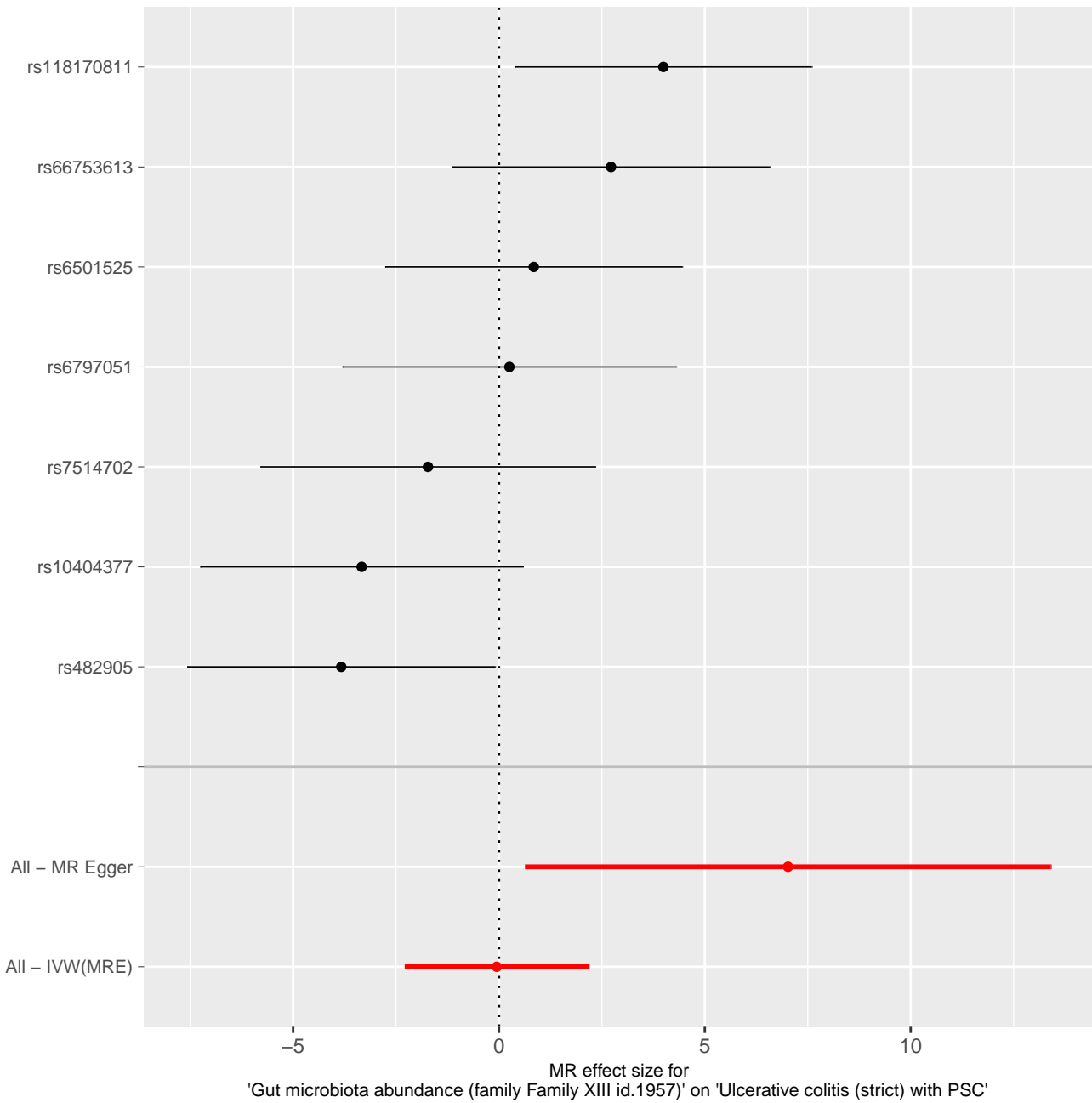


MR effect size for
'Gut microbiota abundance (family Desulfovibrionaceae id.3169) on 'Ulcerative colitis (strict) with PSC'

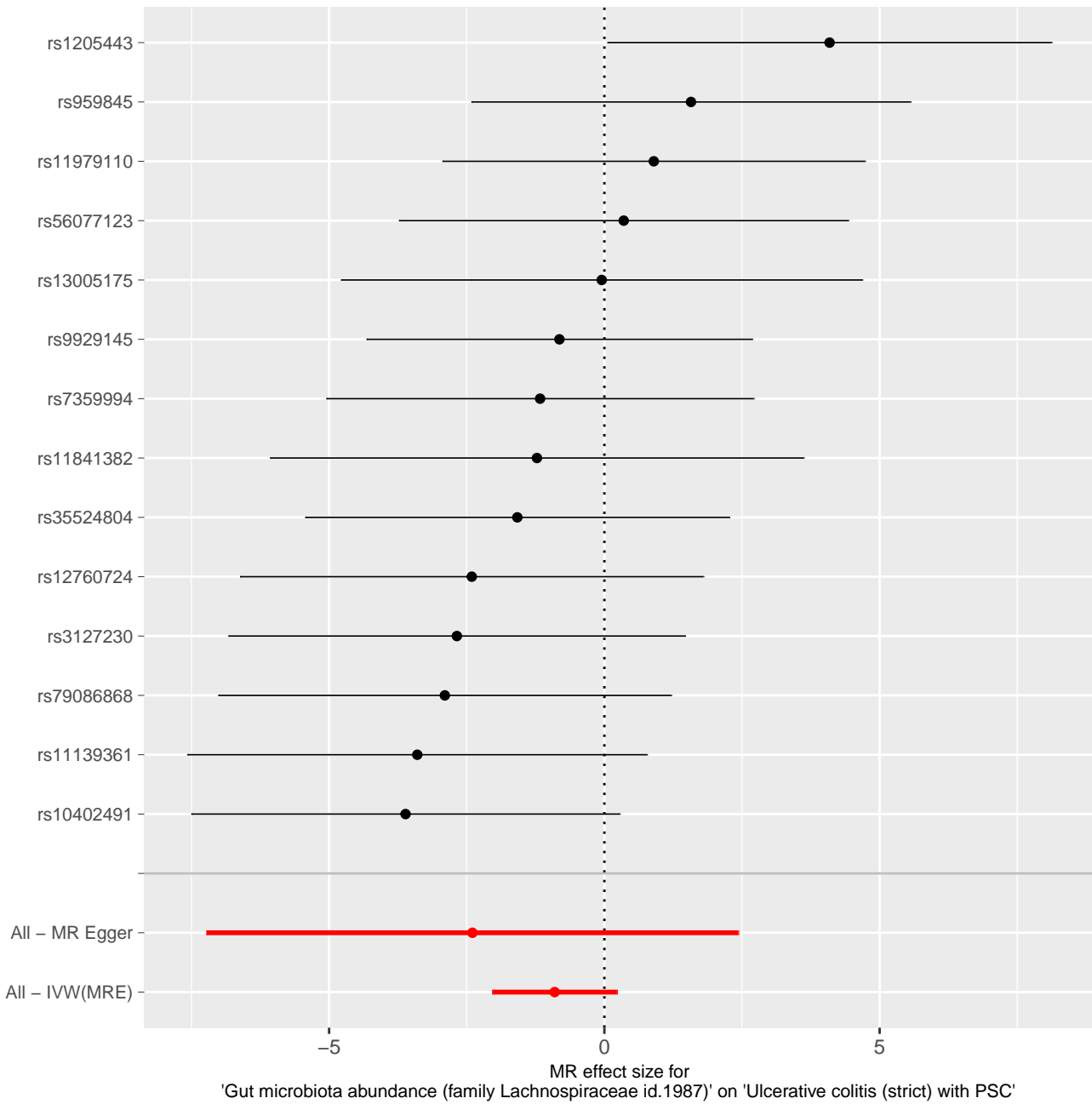


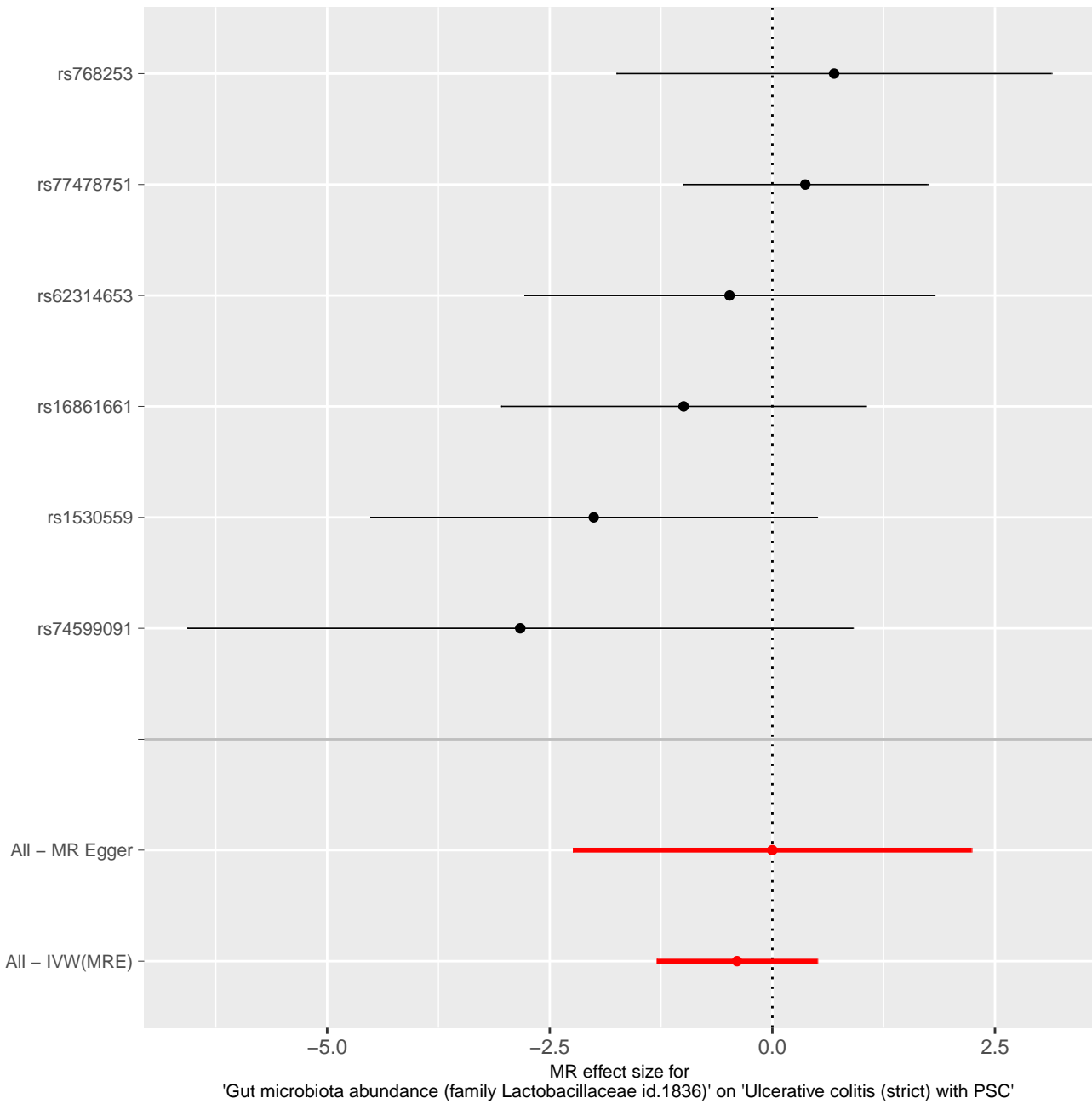


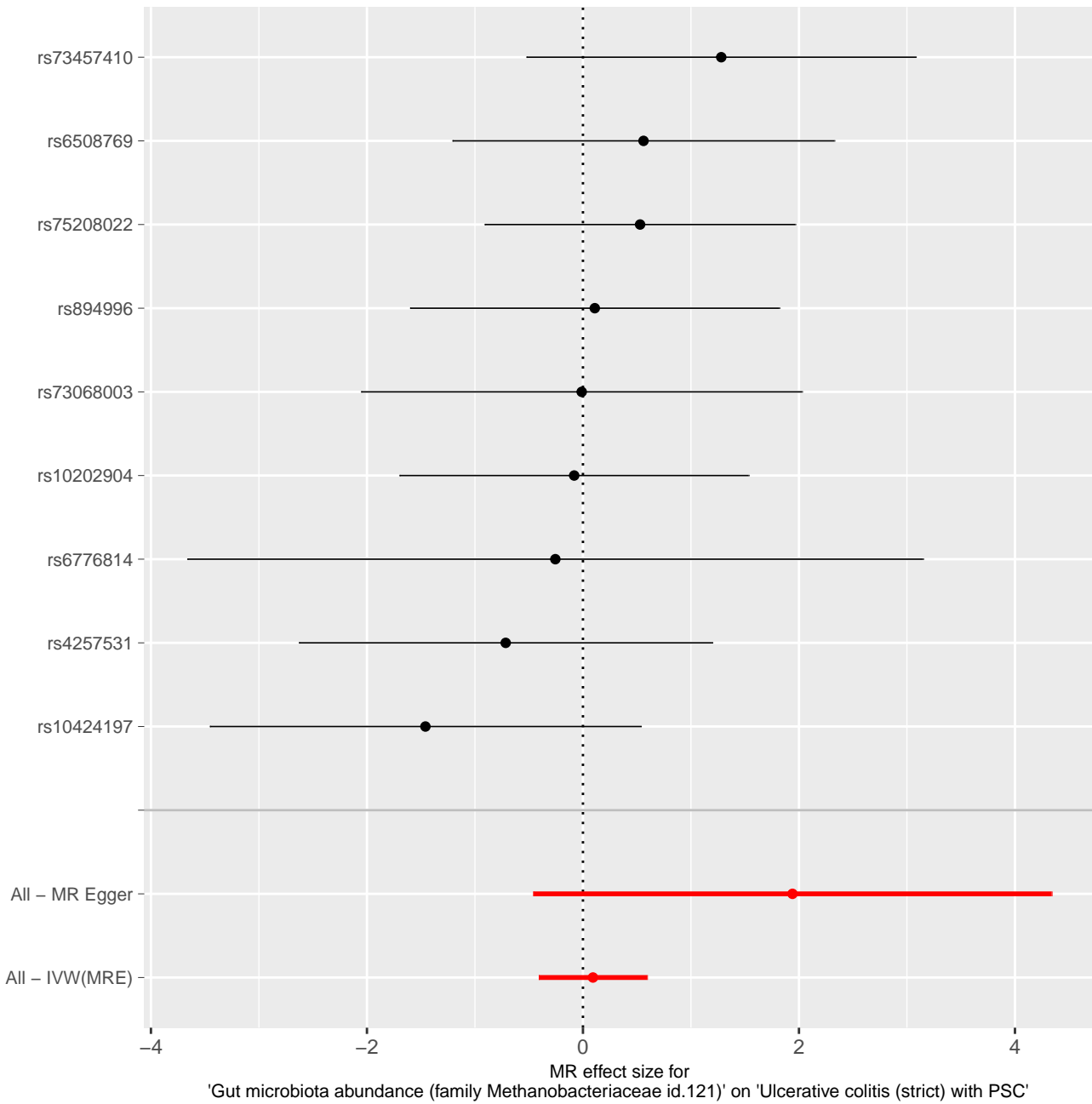


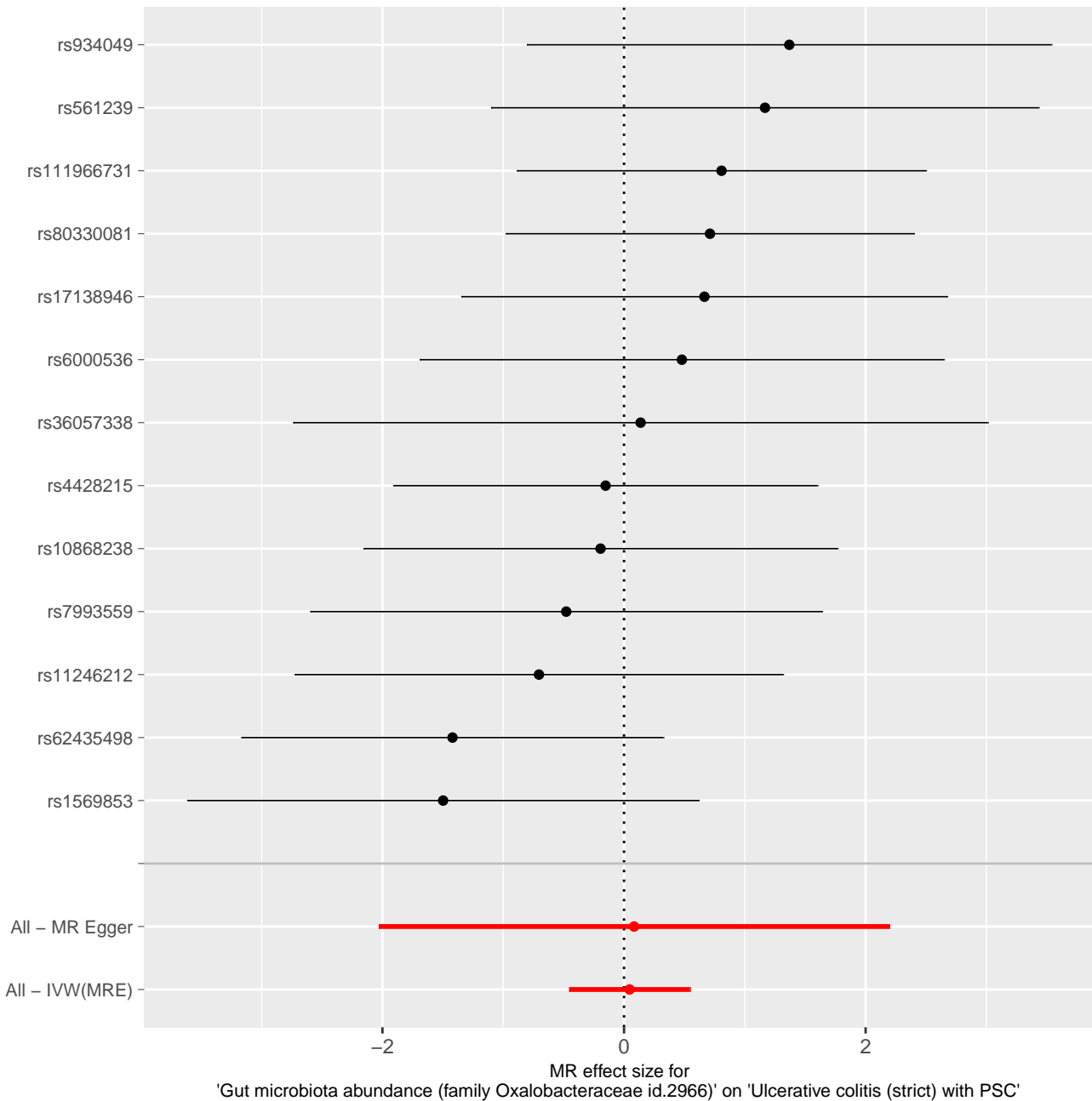


Batch 617 : Gut microbiota abundance (family Lachnospiraceae id.1987) on Ulcerative colitis (strict) with PSC

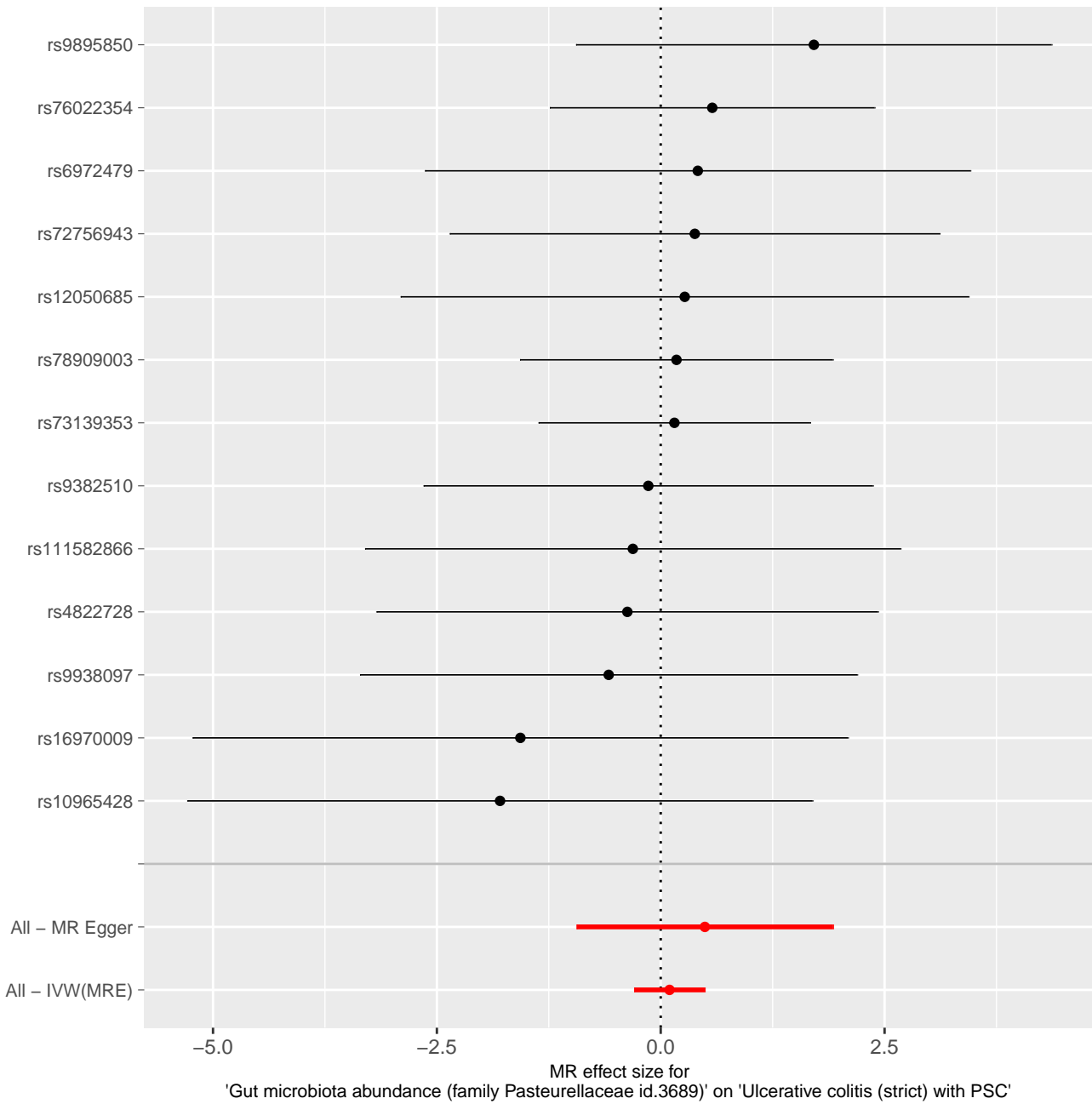


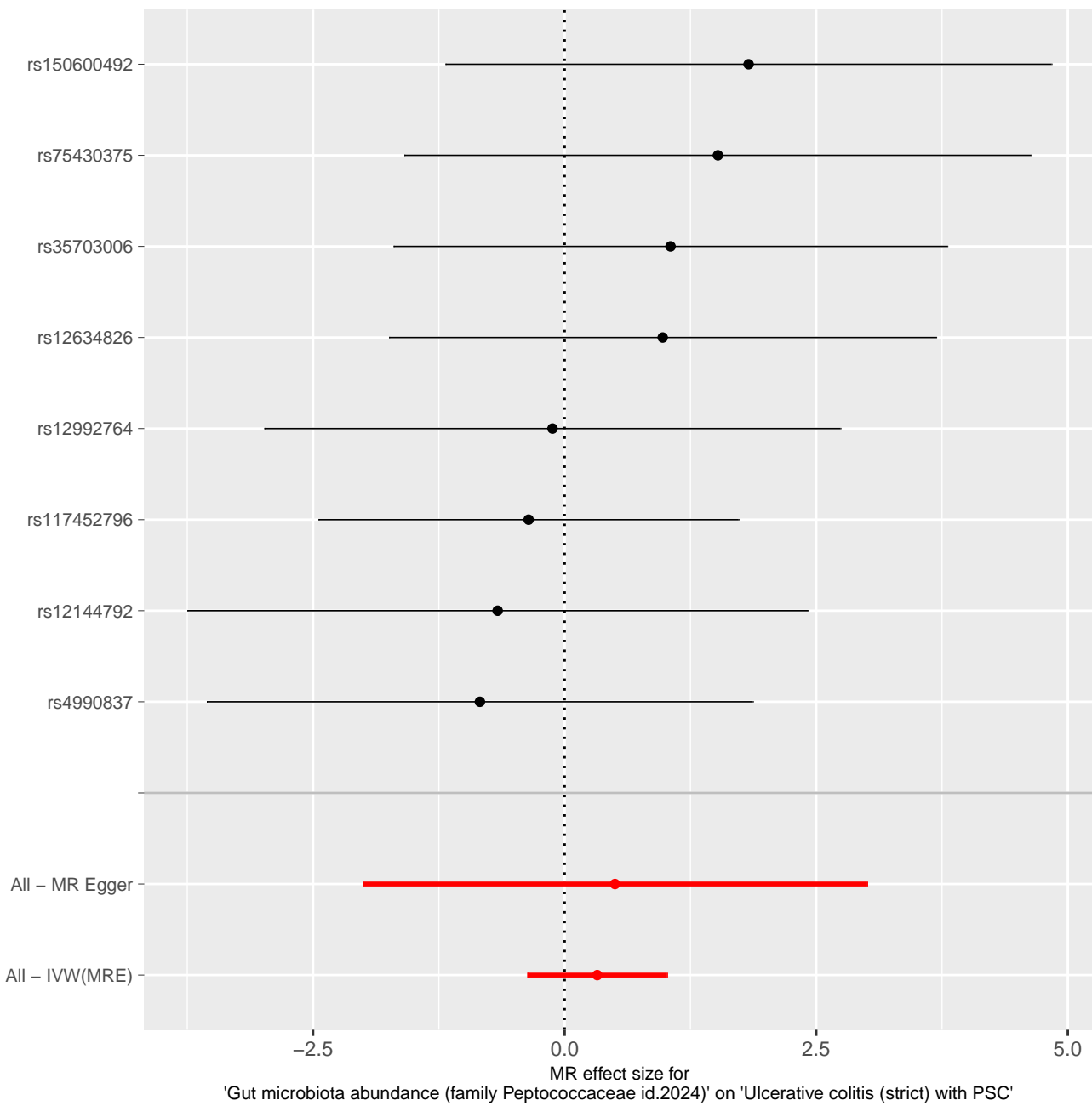


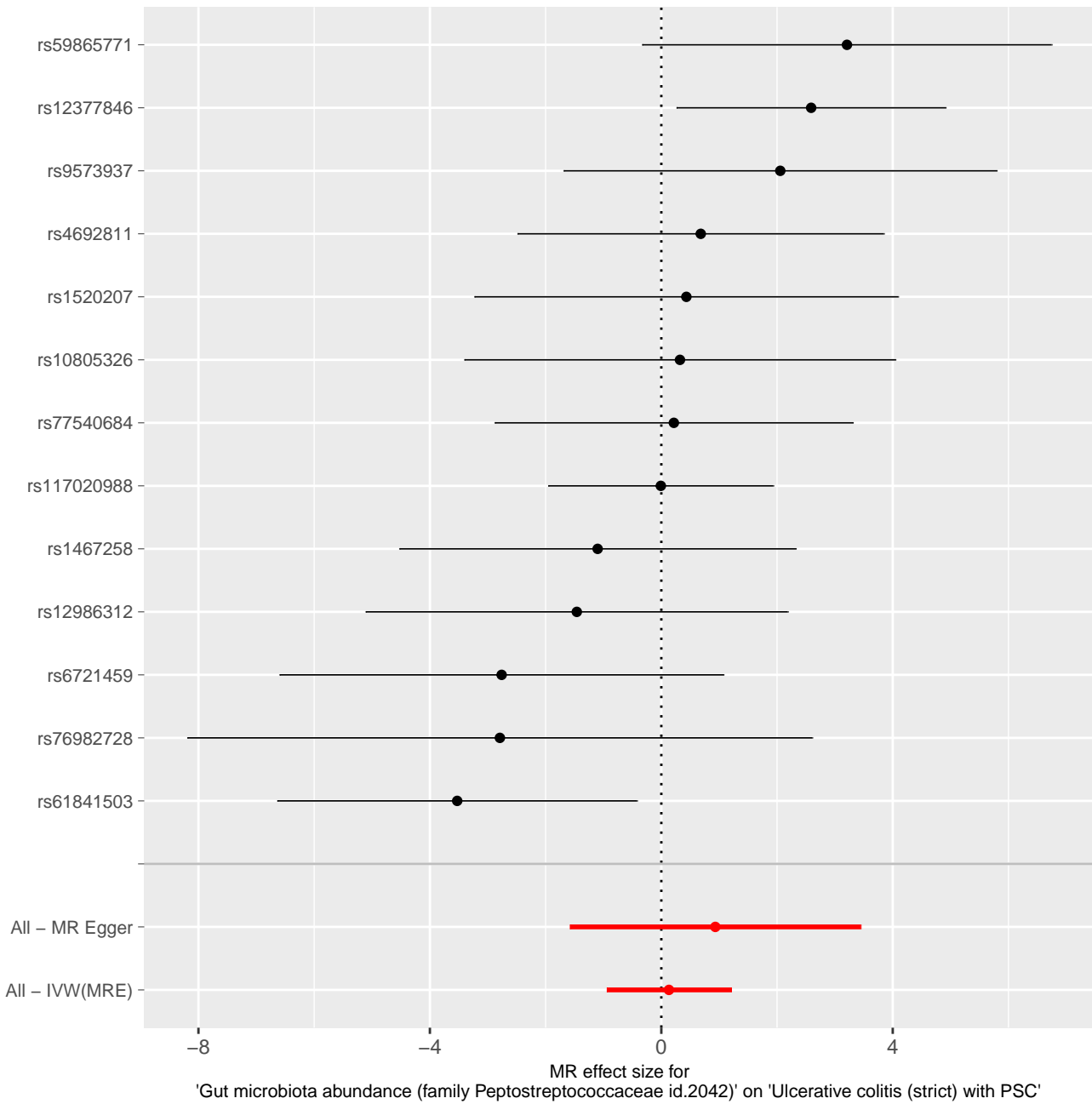


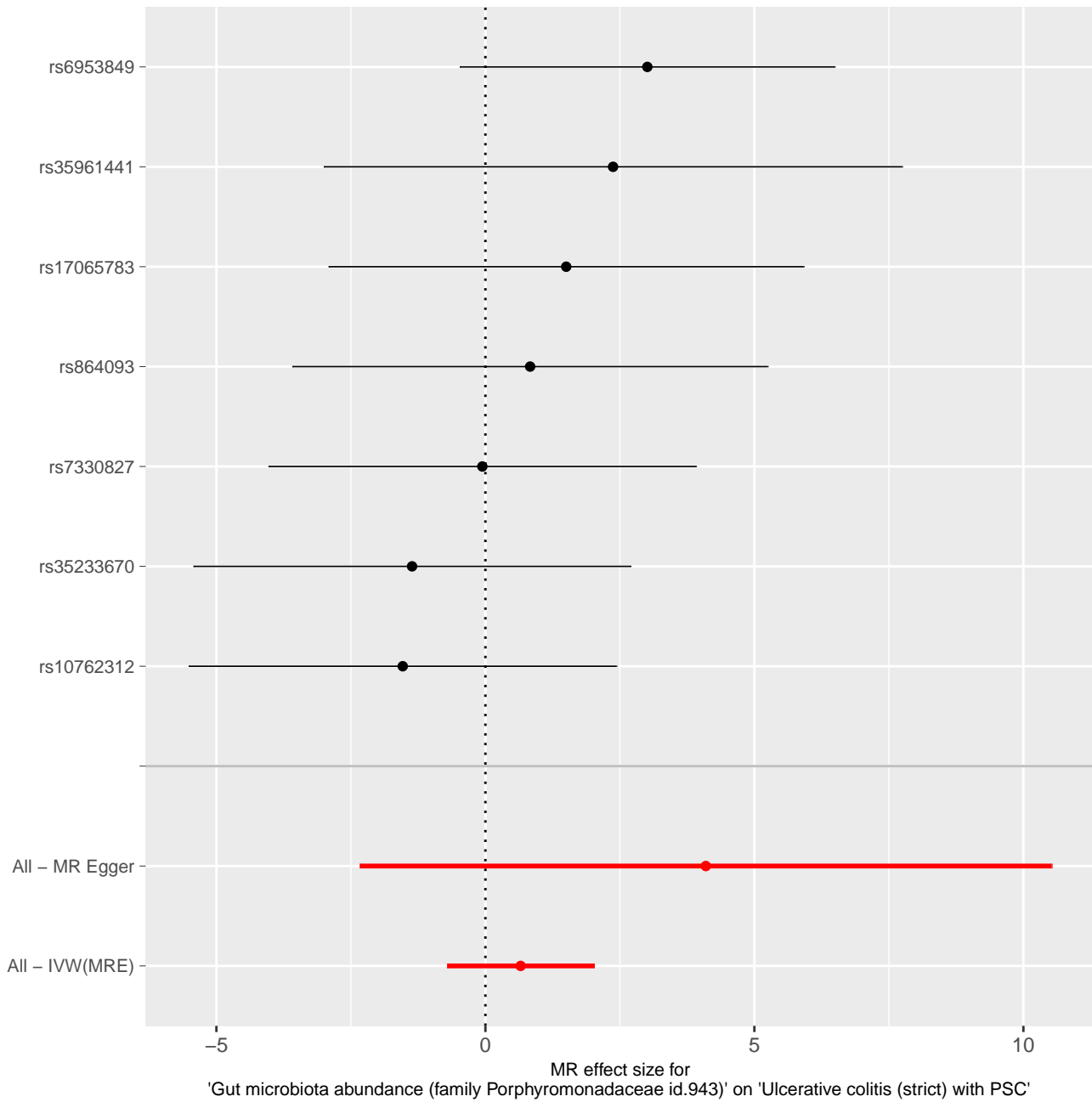


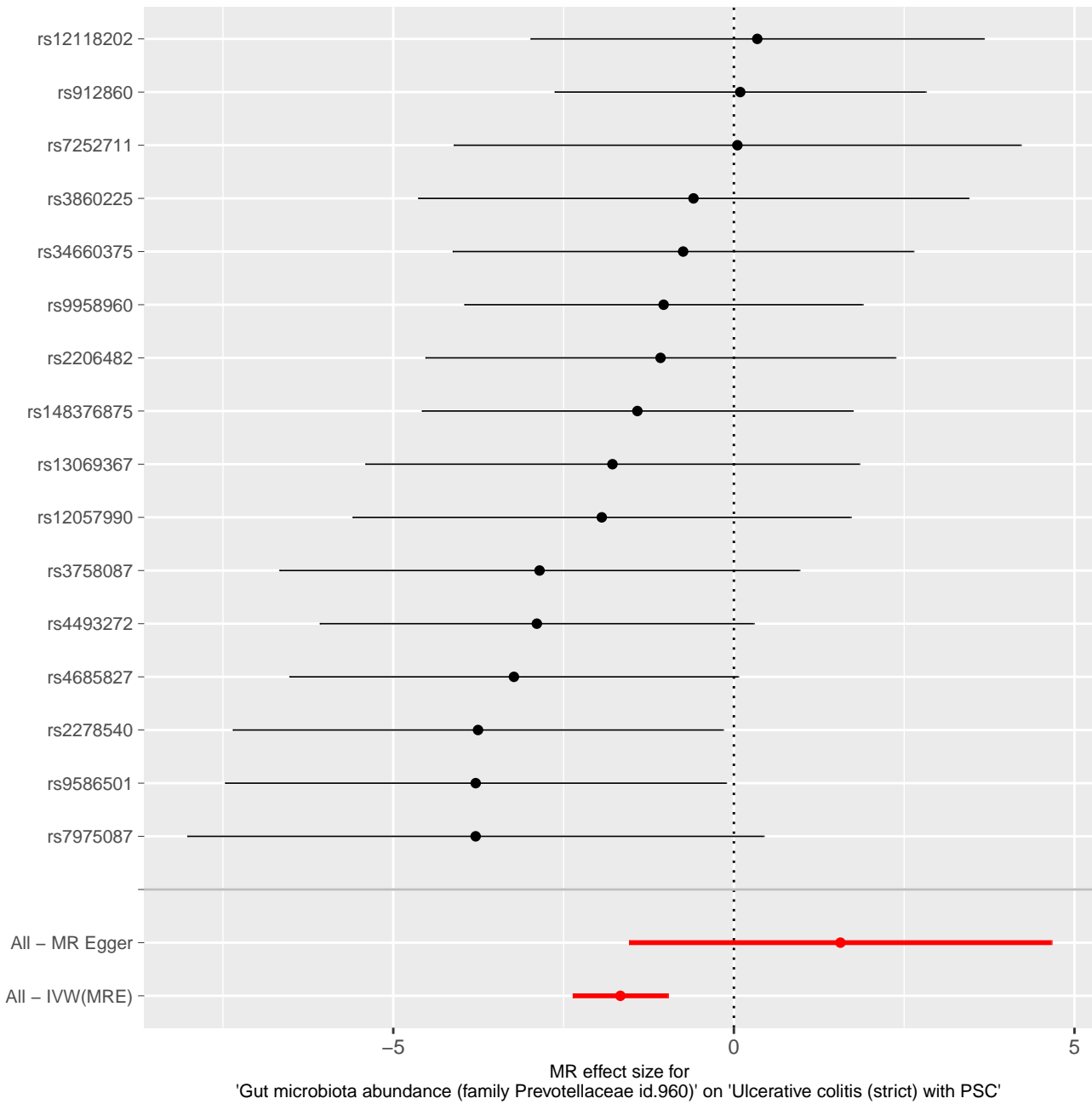
Batch 621 : Gut microbiota abundance (family Pasteurellaceae id.3689) on Ulcerative colitis (strict) with PSC

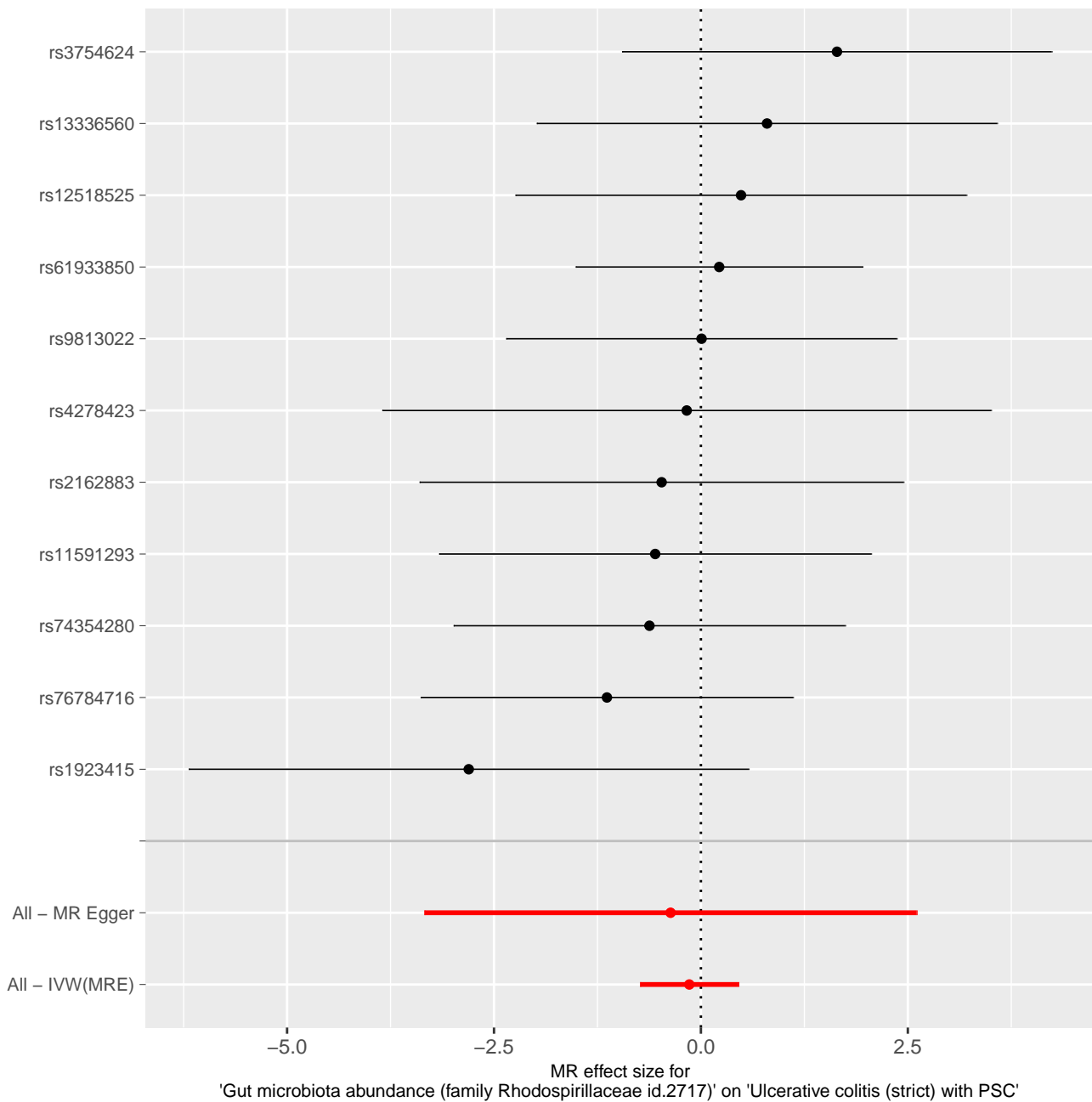




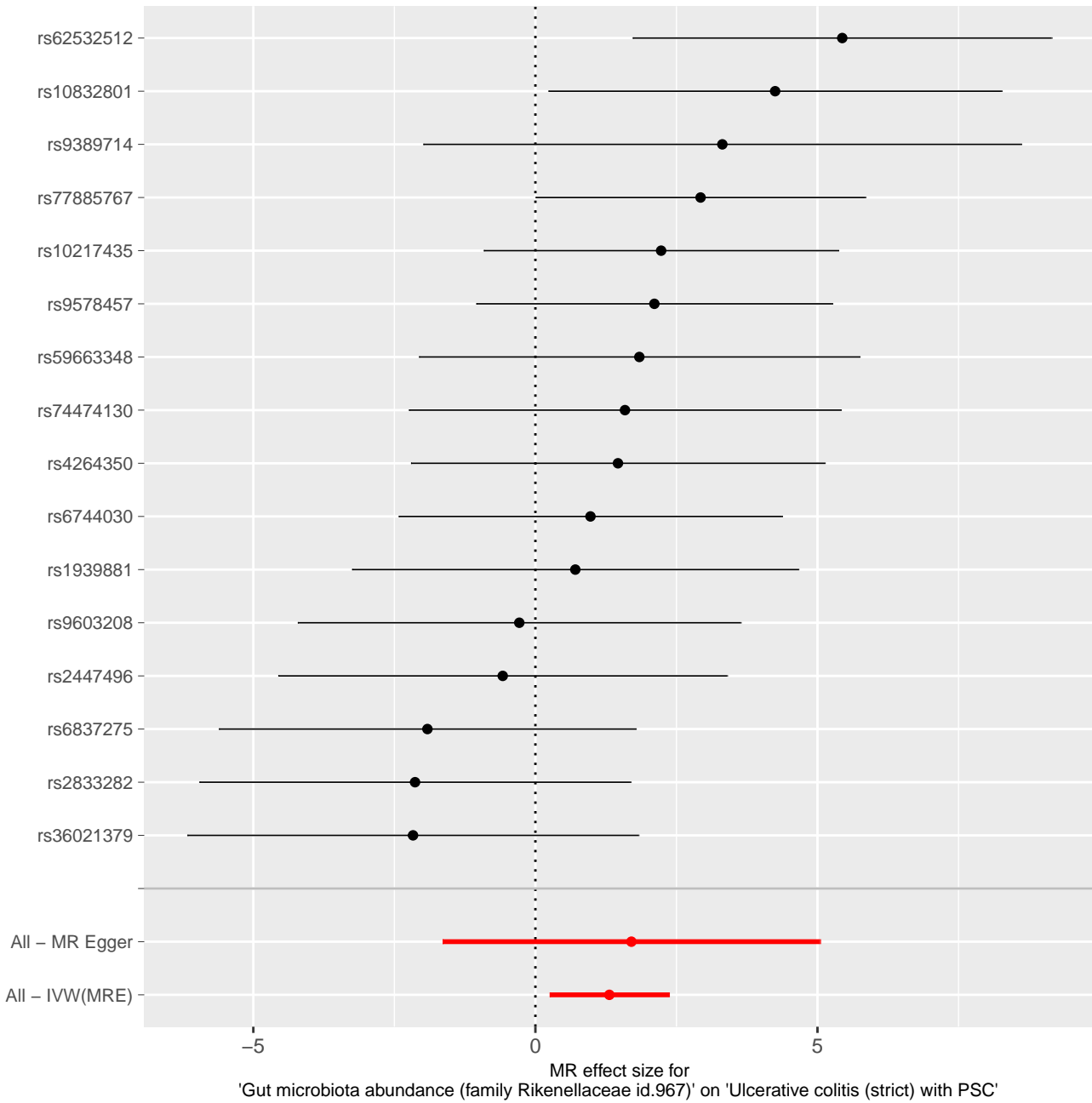


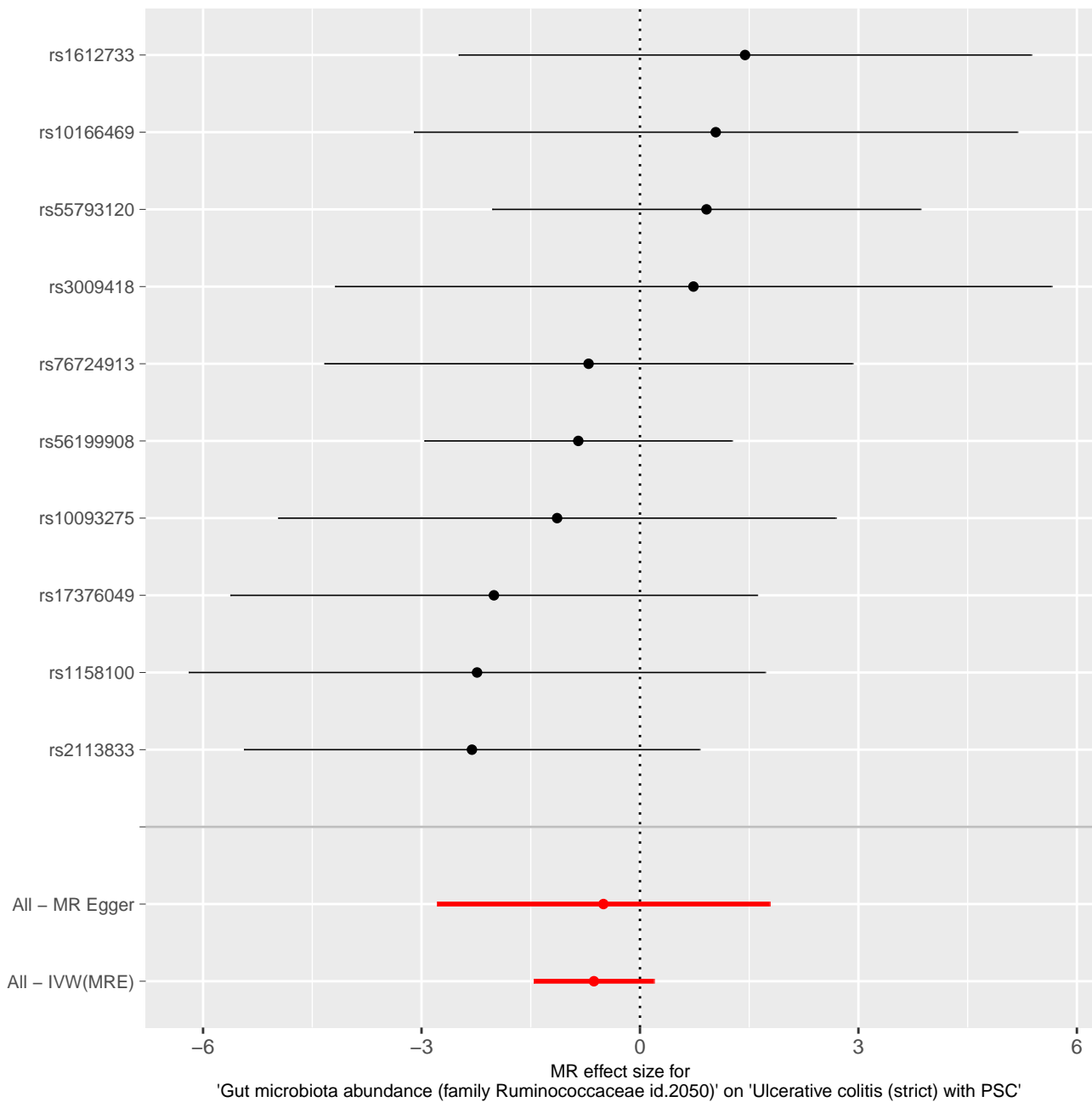


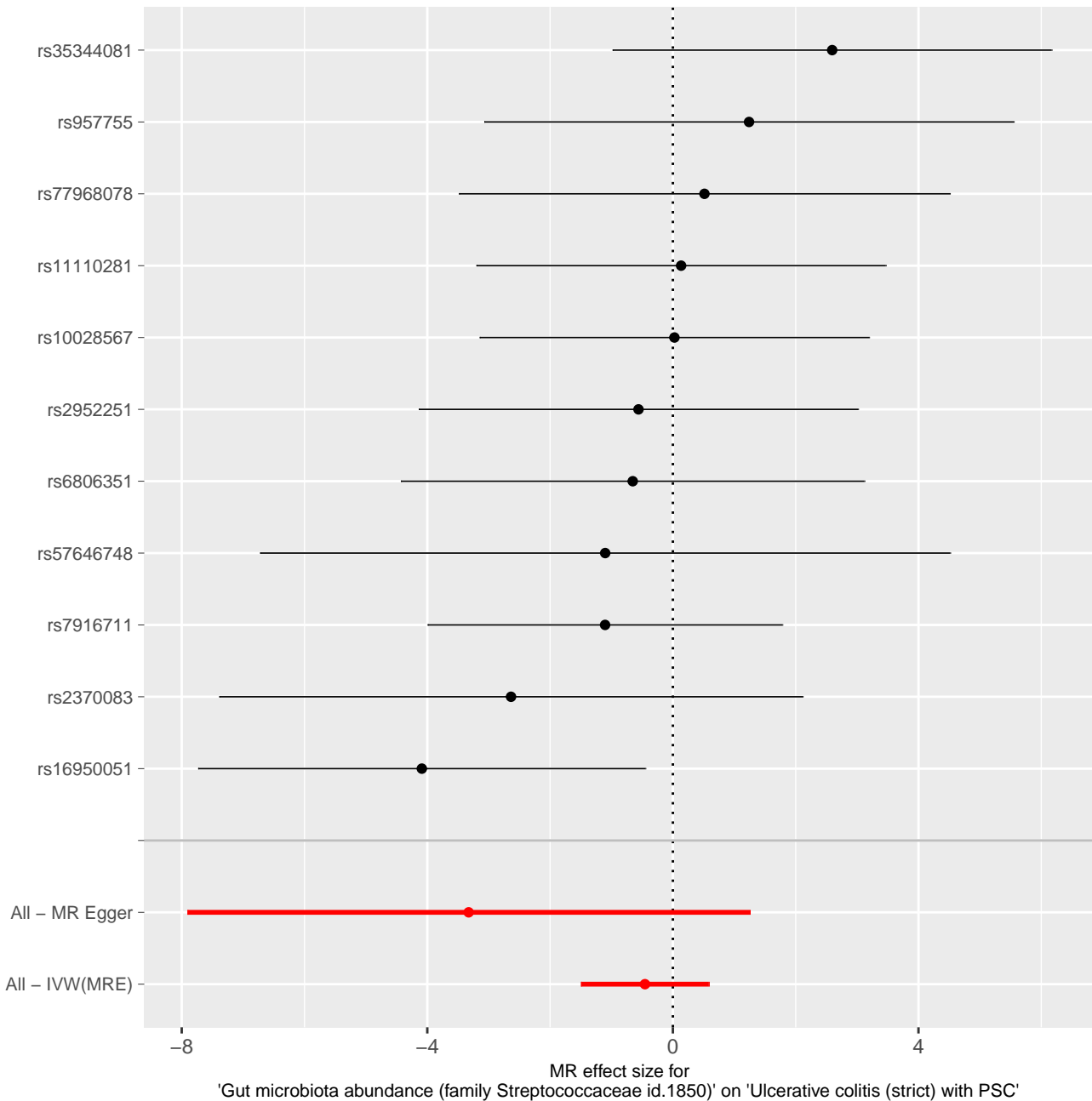


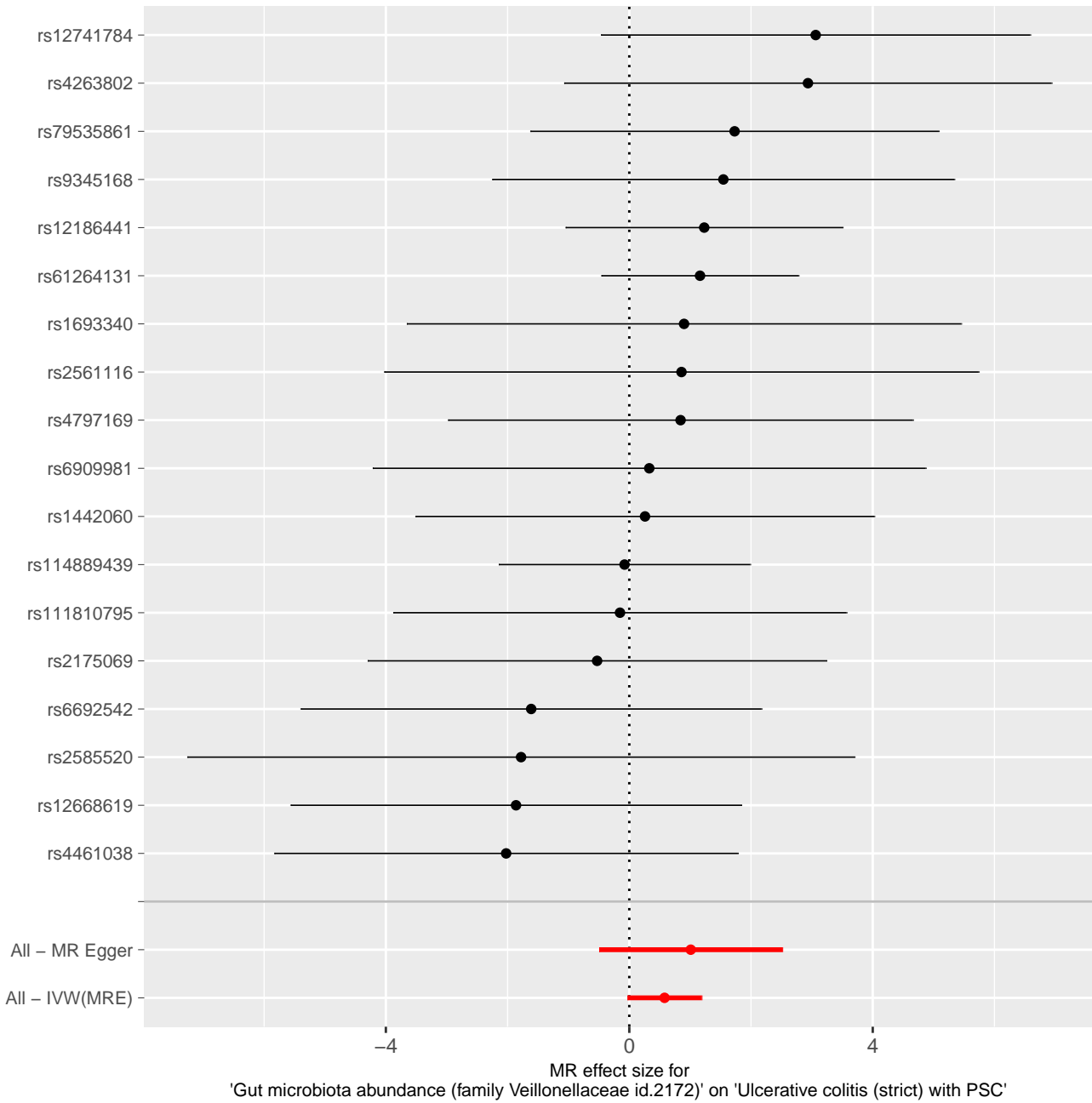


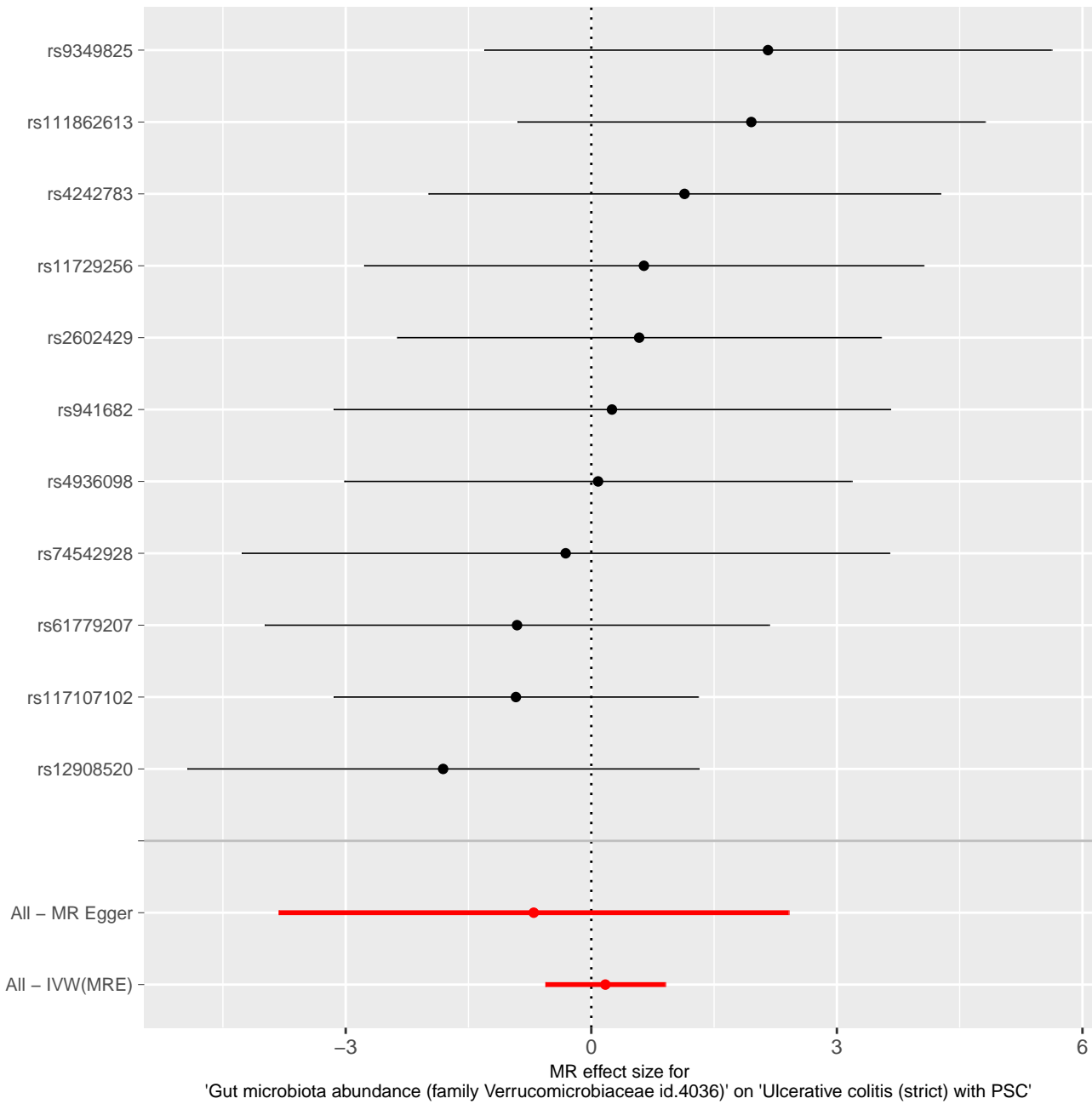
Batch 627 : Gut microbiota abundance (family Rikenellaceae id.967) on Ulcerative colitis (strict) with PSC

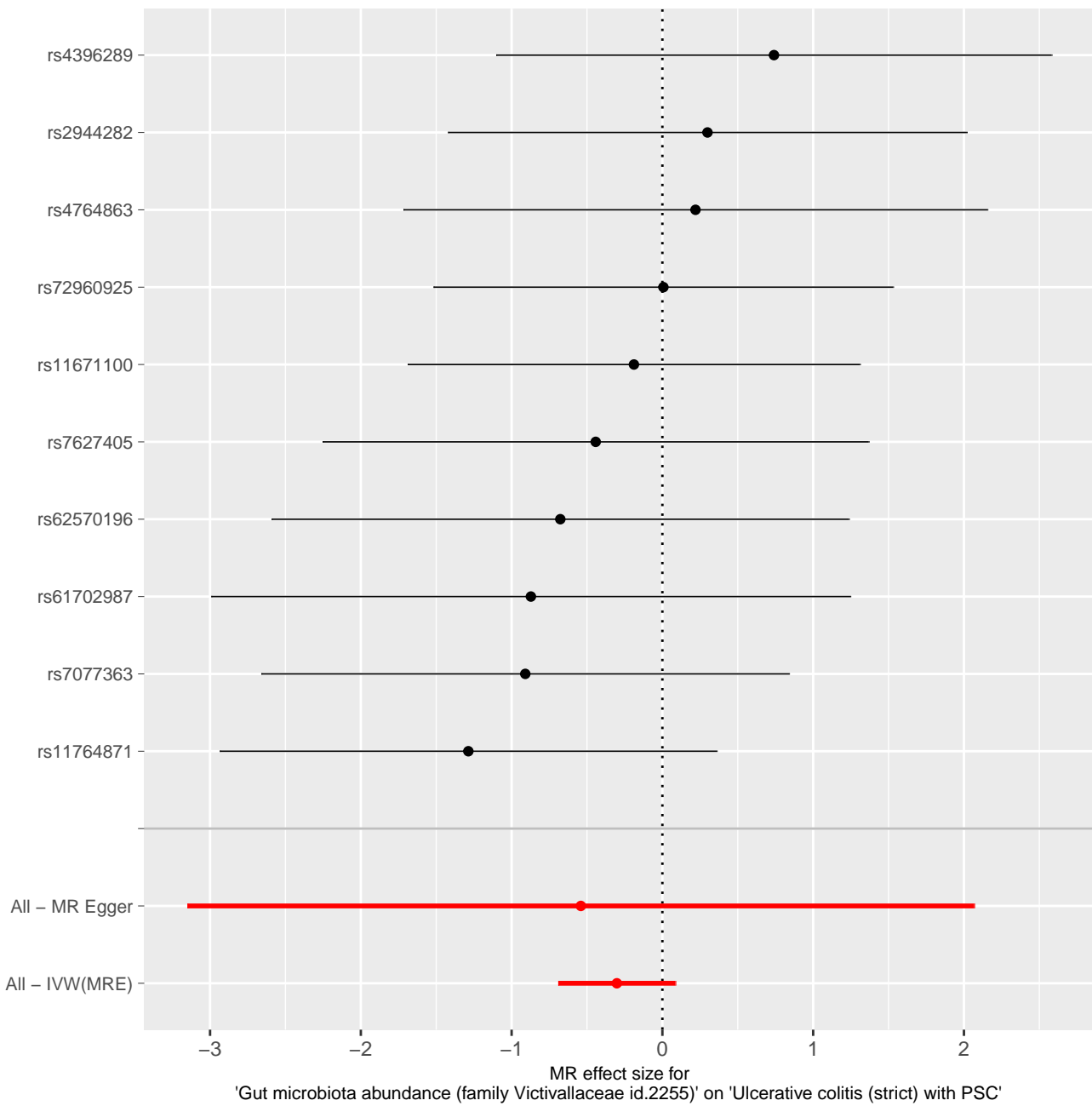


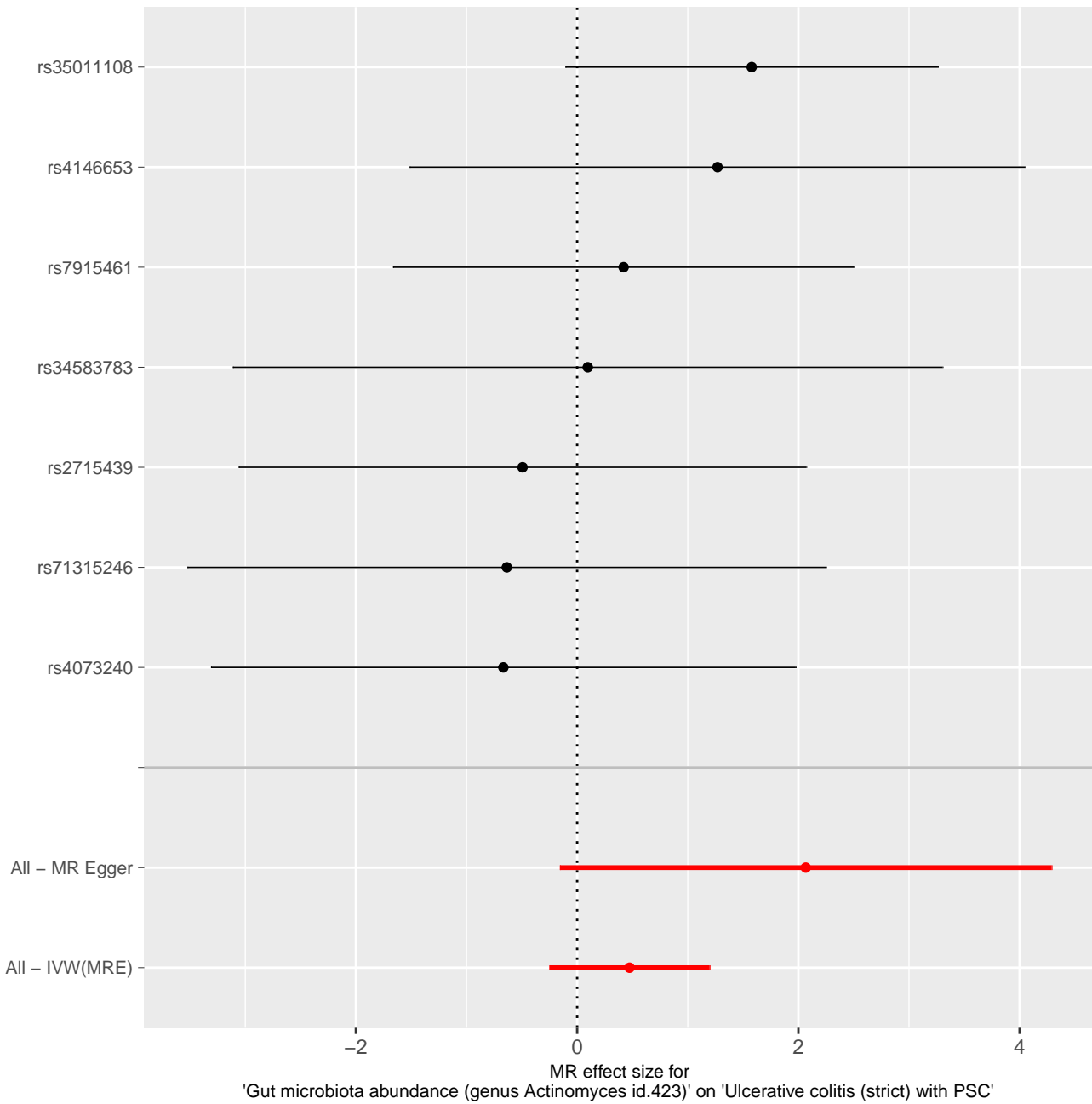


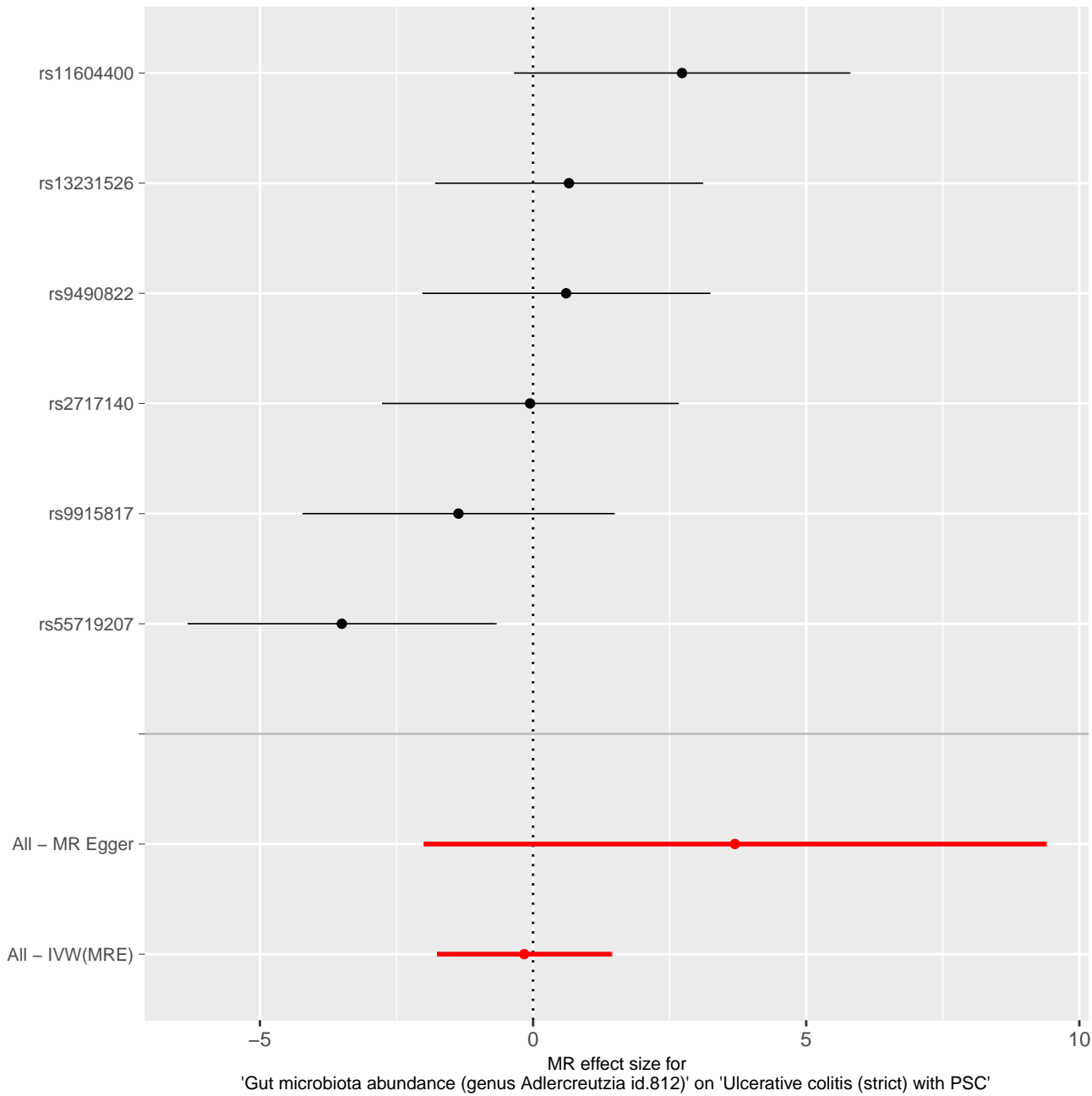




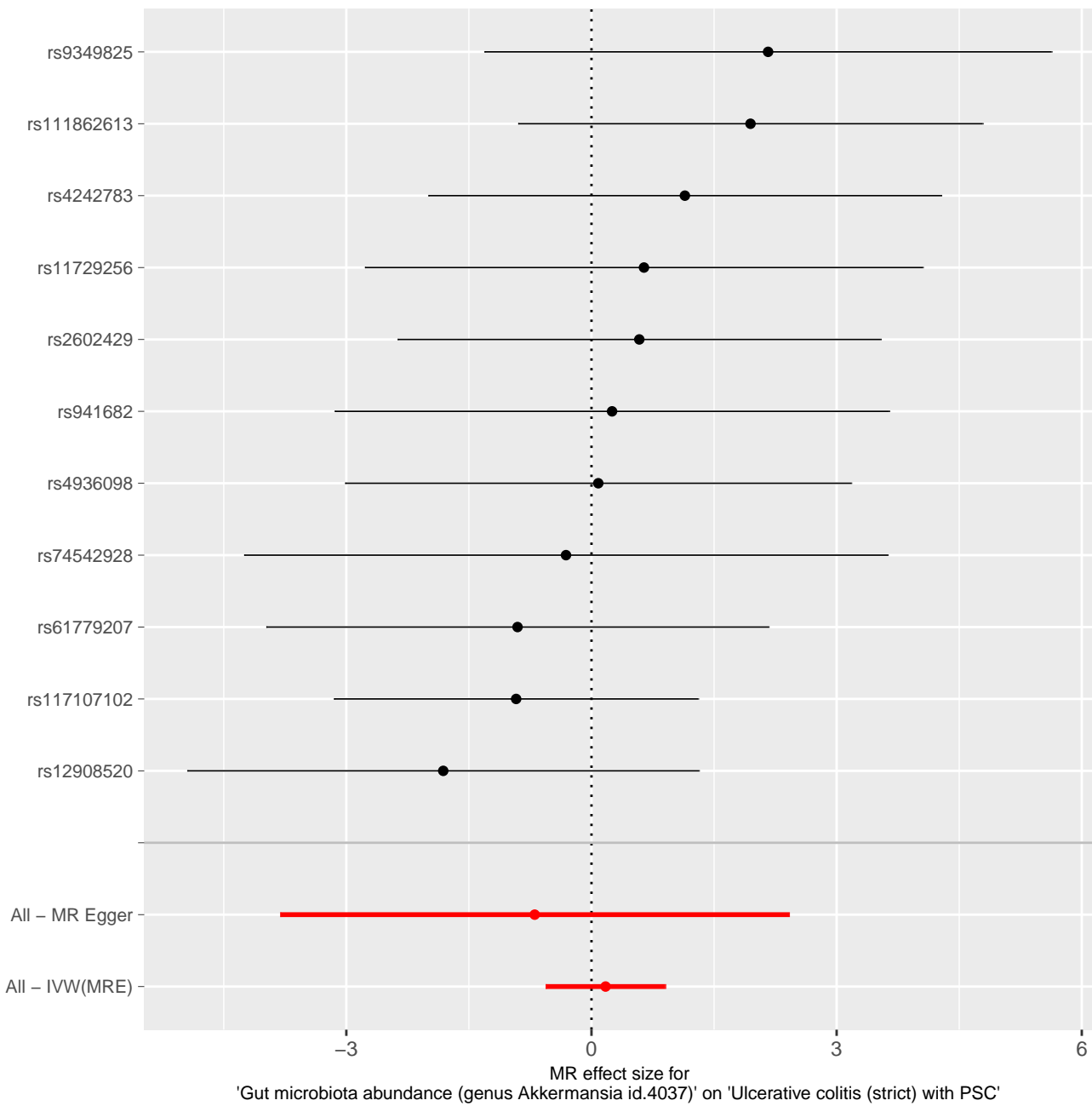


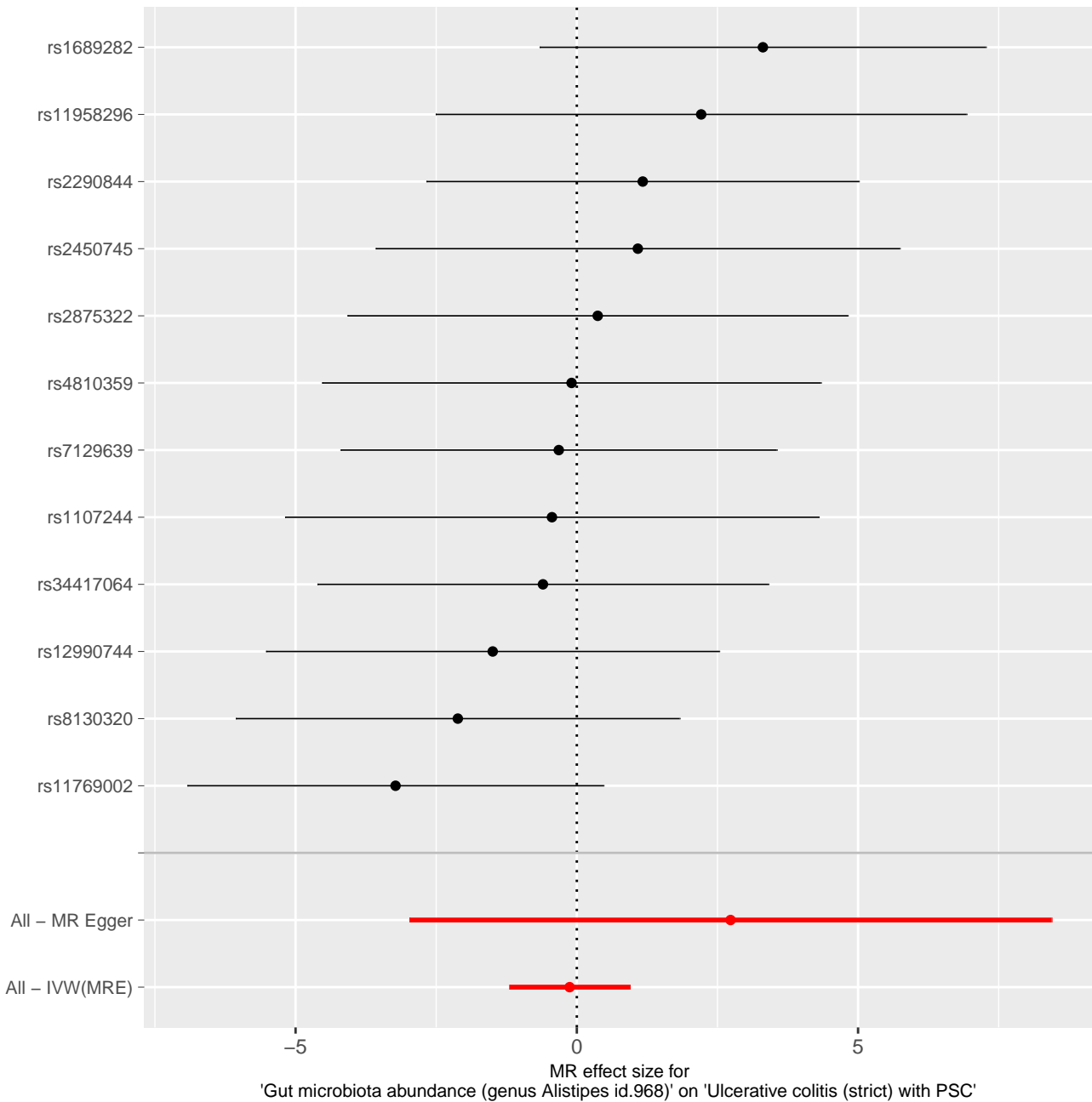


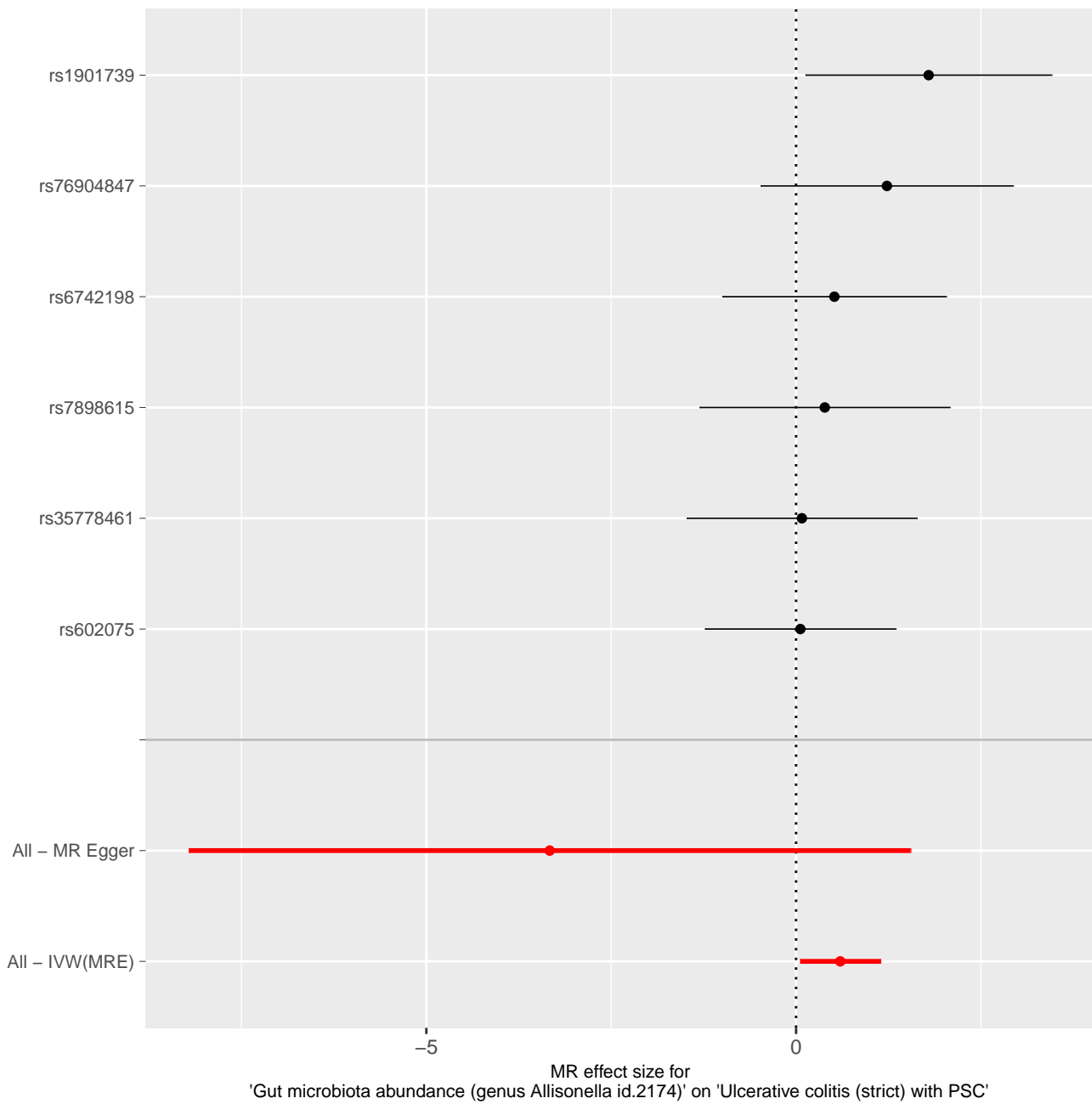


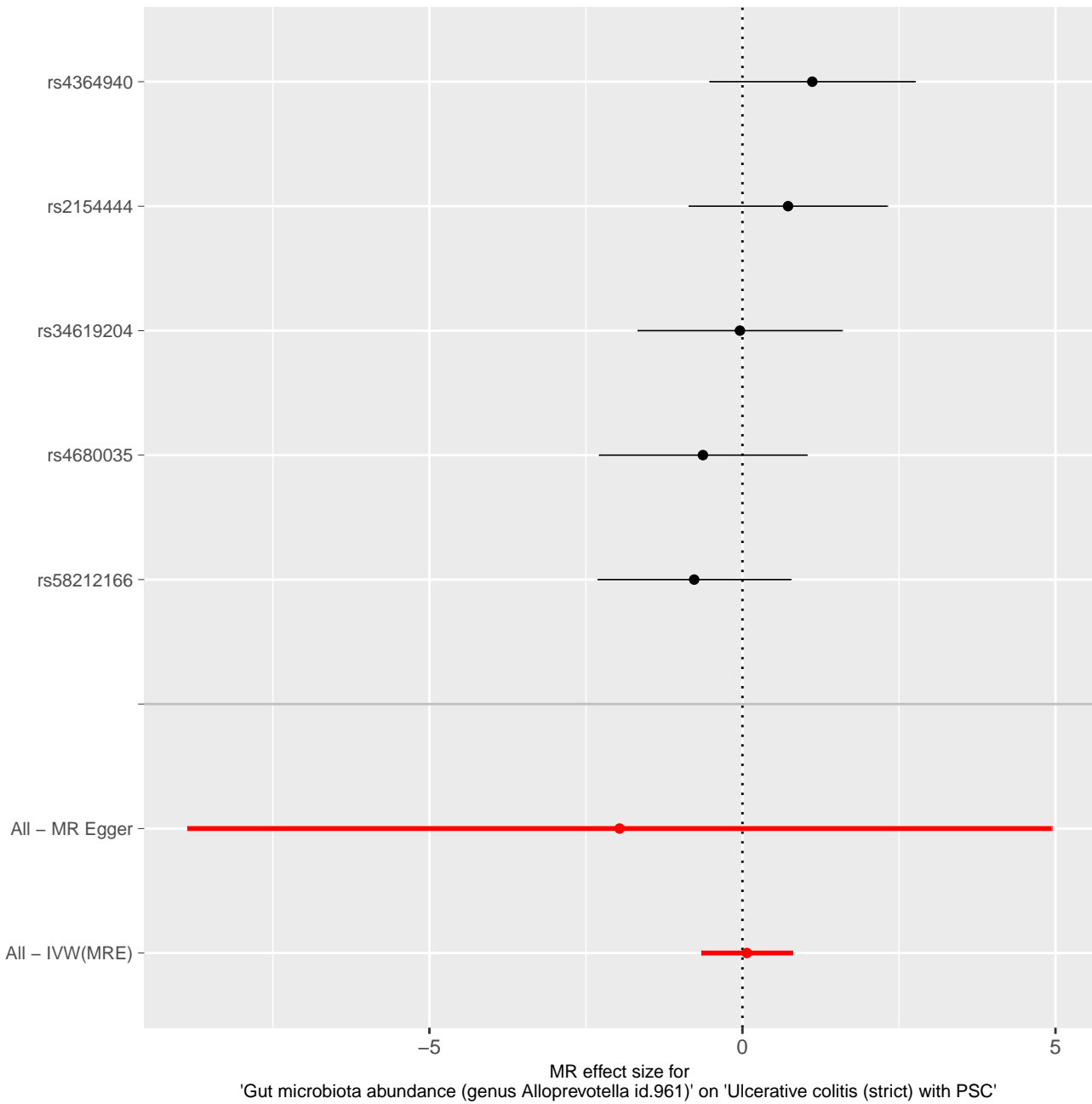


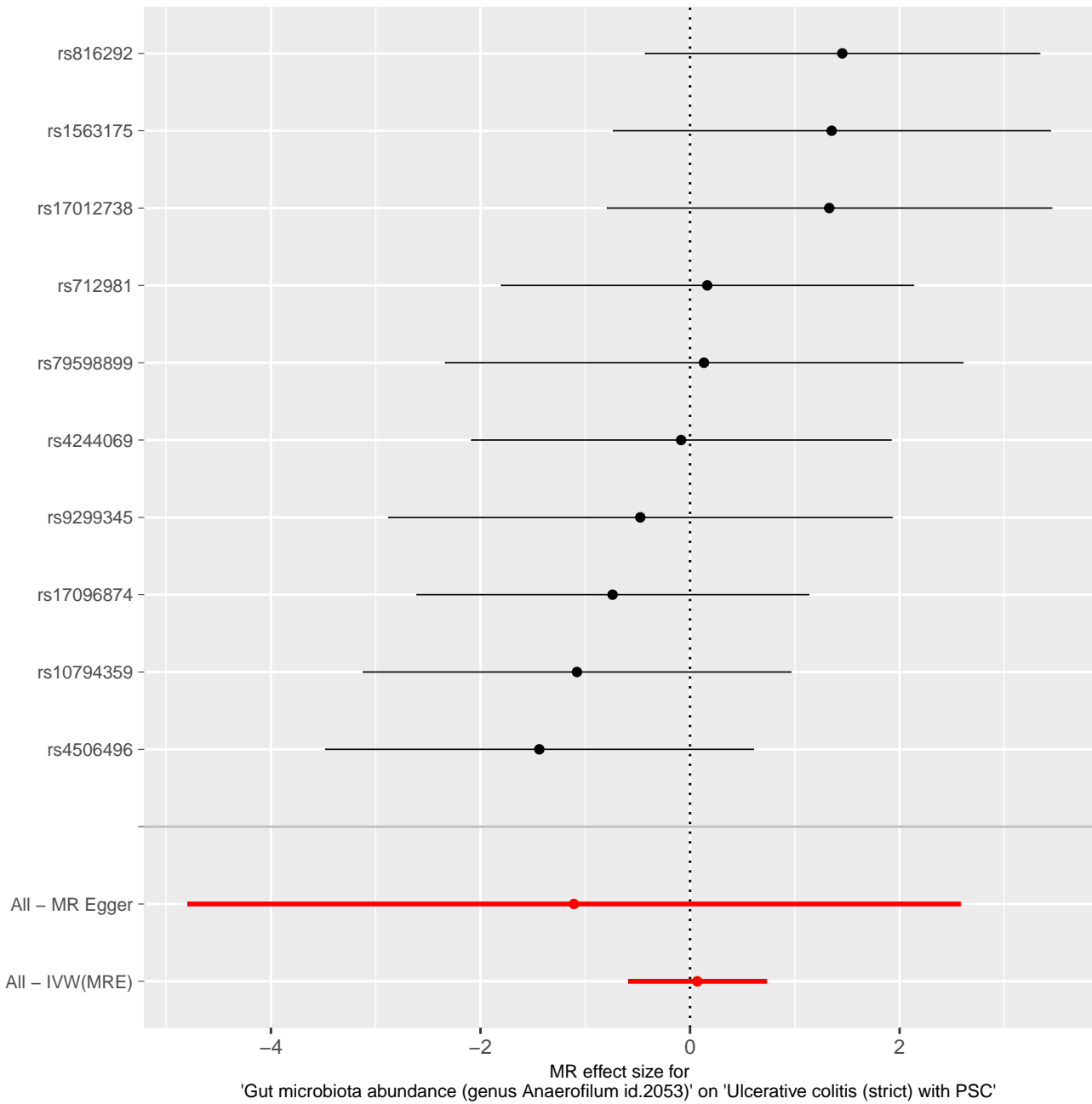
Batch 635 : Gut microbiota abundance (genus Akkermansia id.4037) on Ulcerative colitis (strict) with PSC

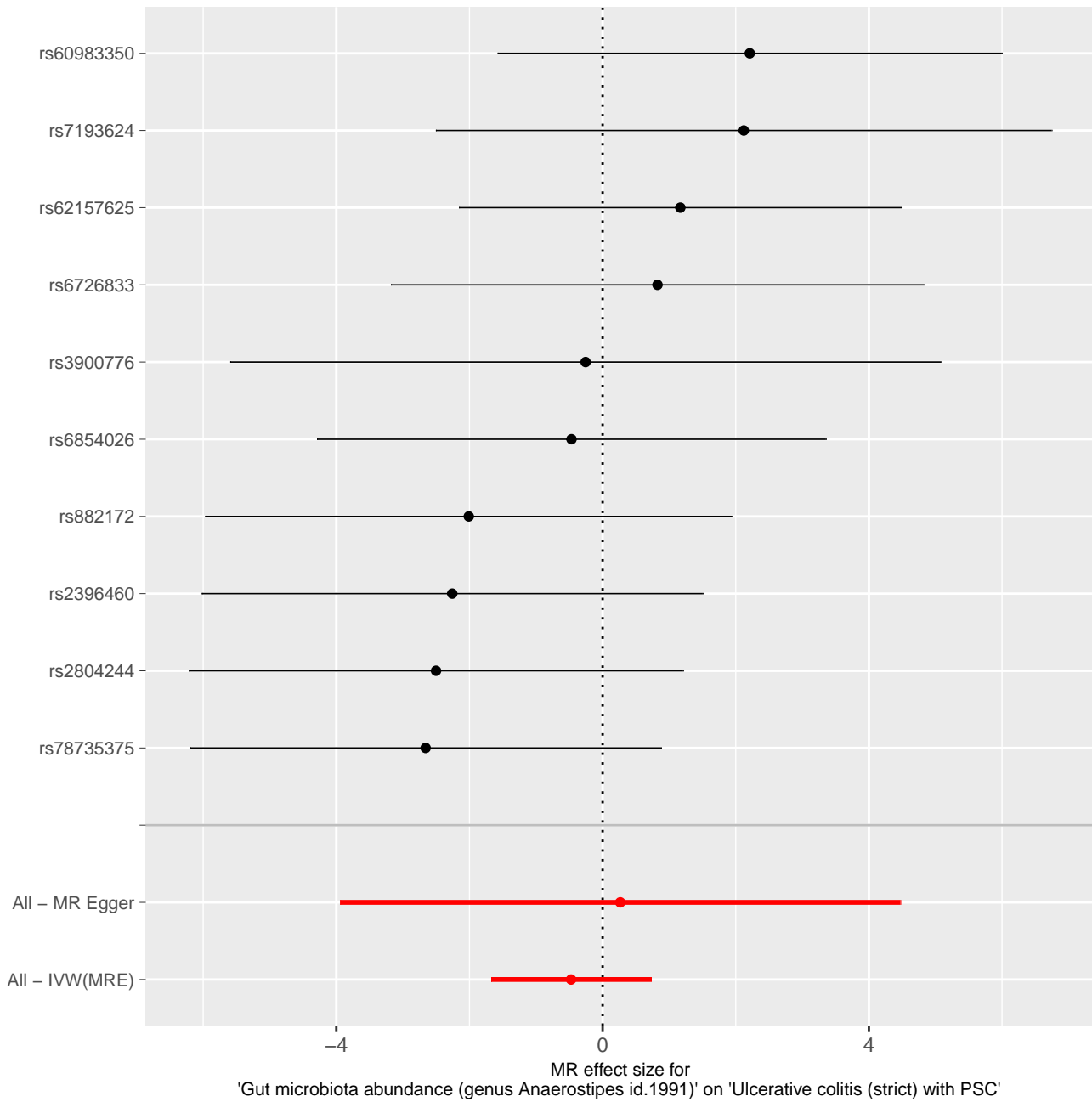


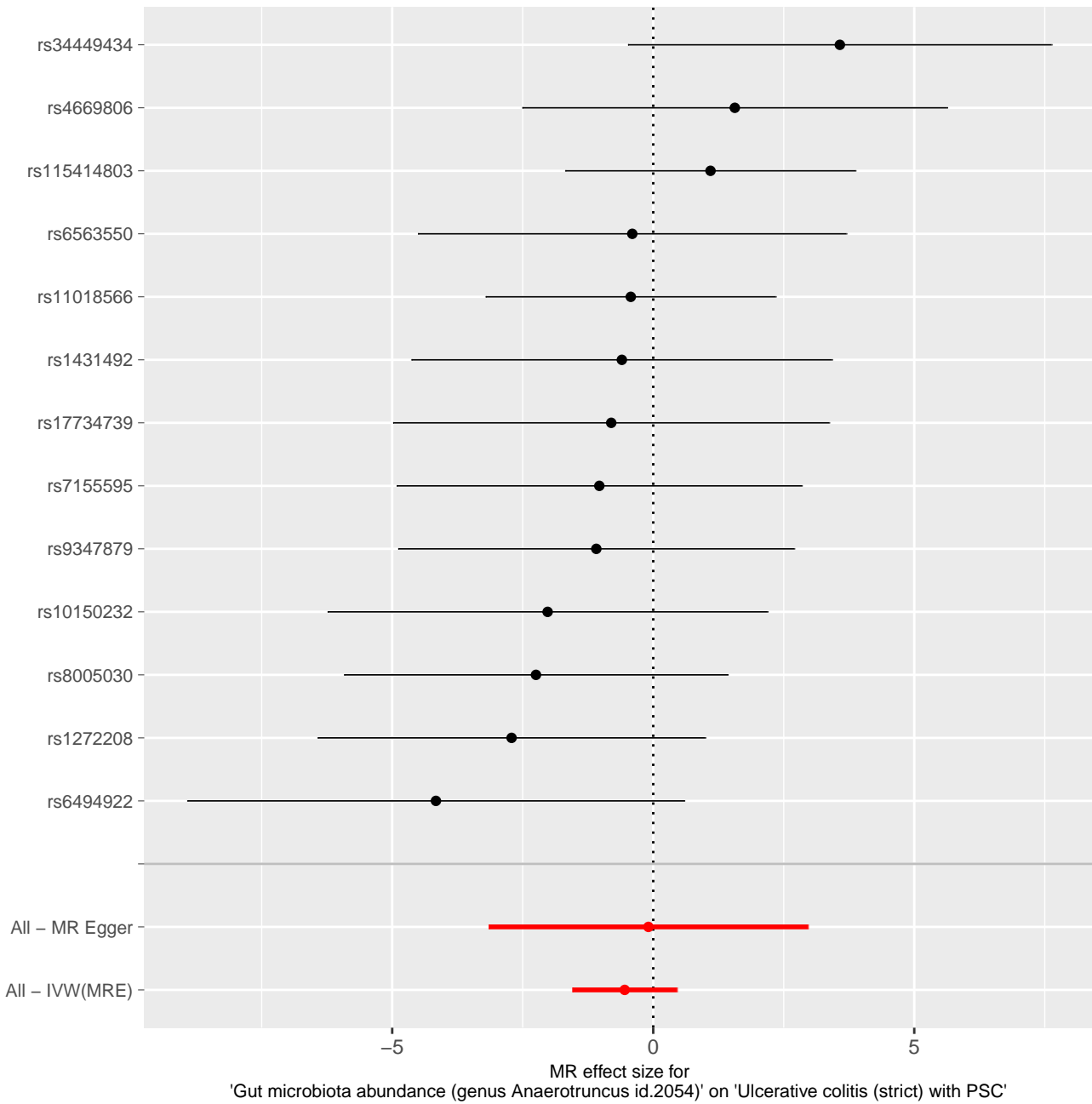


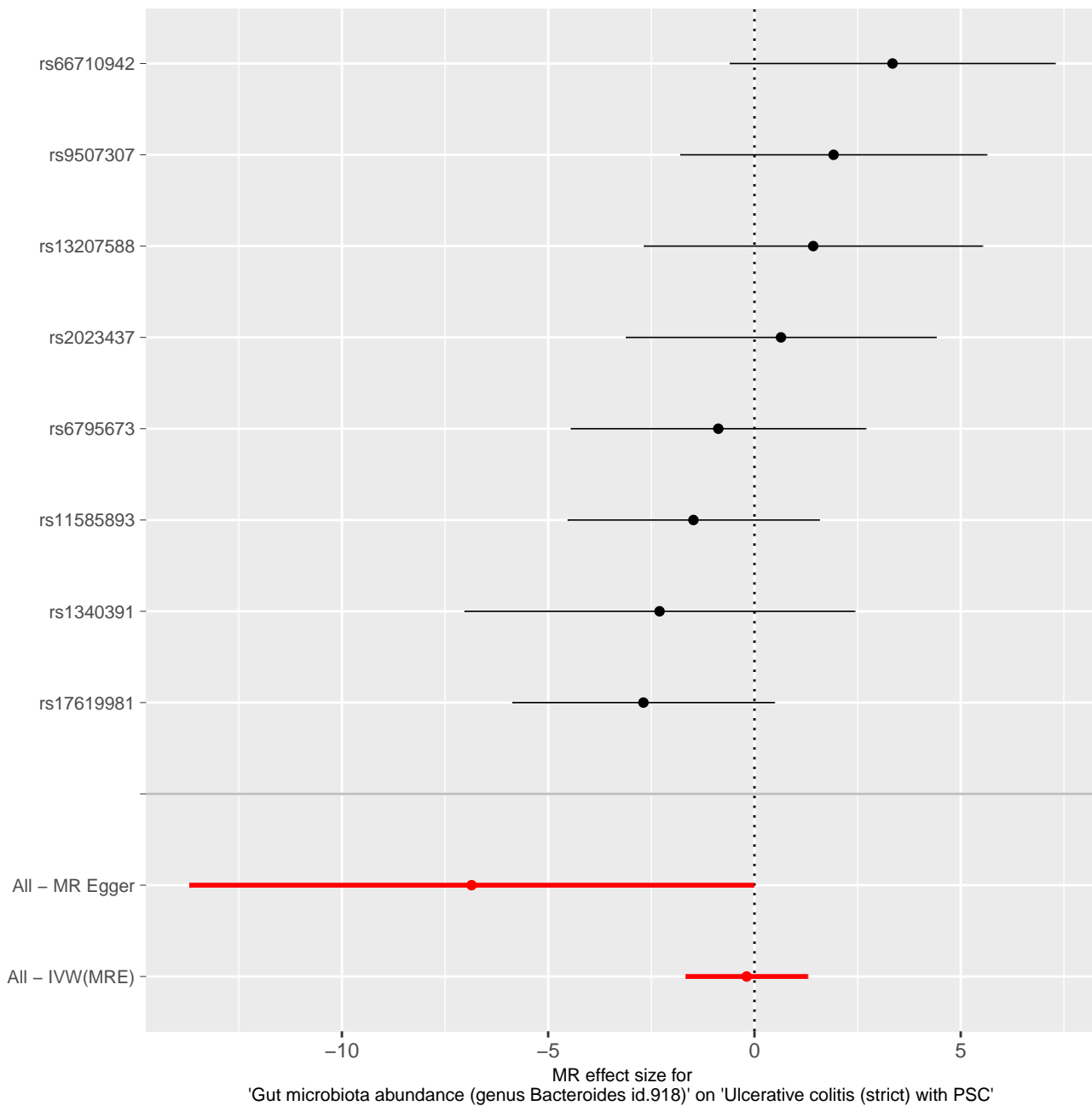


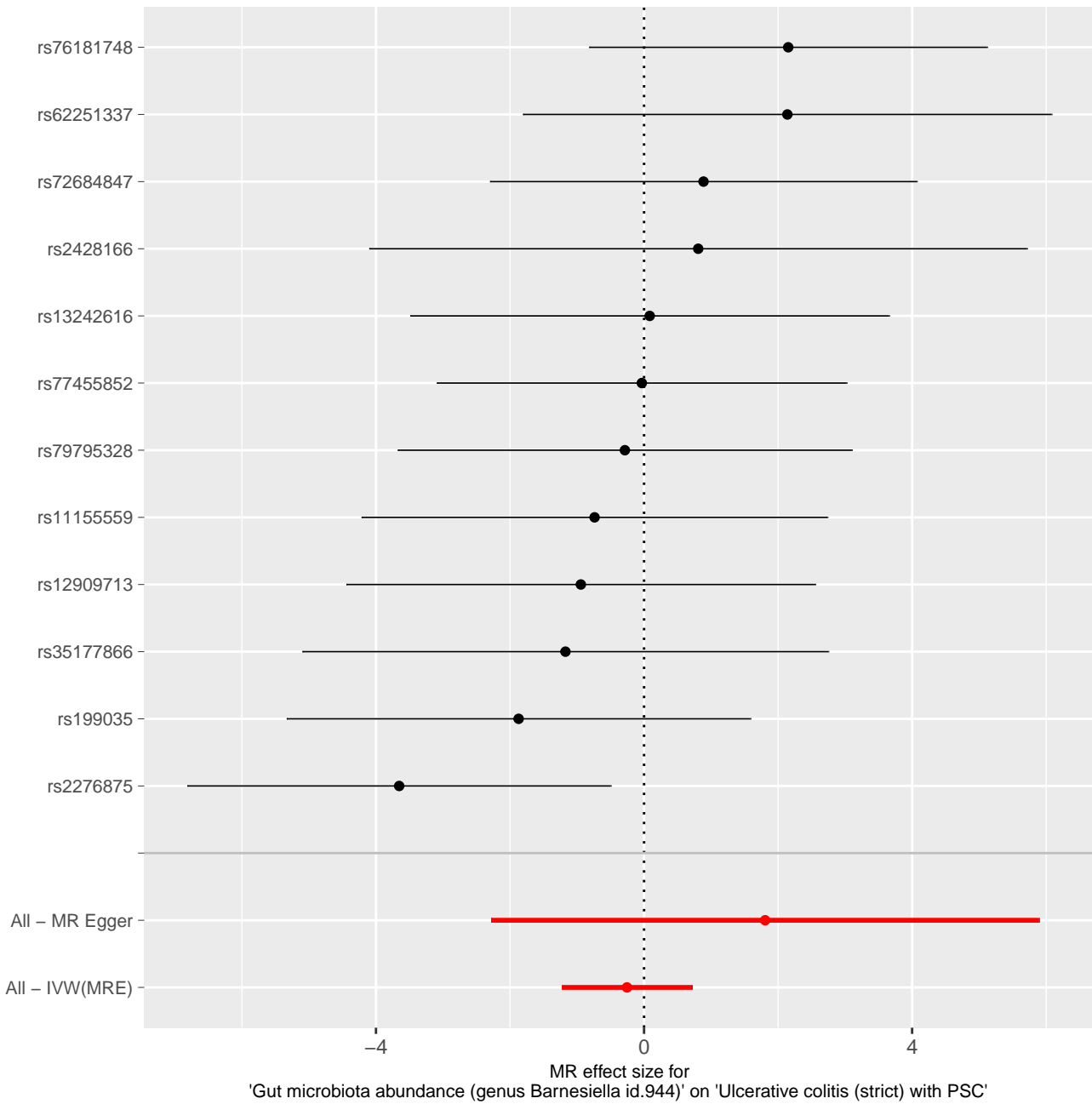


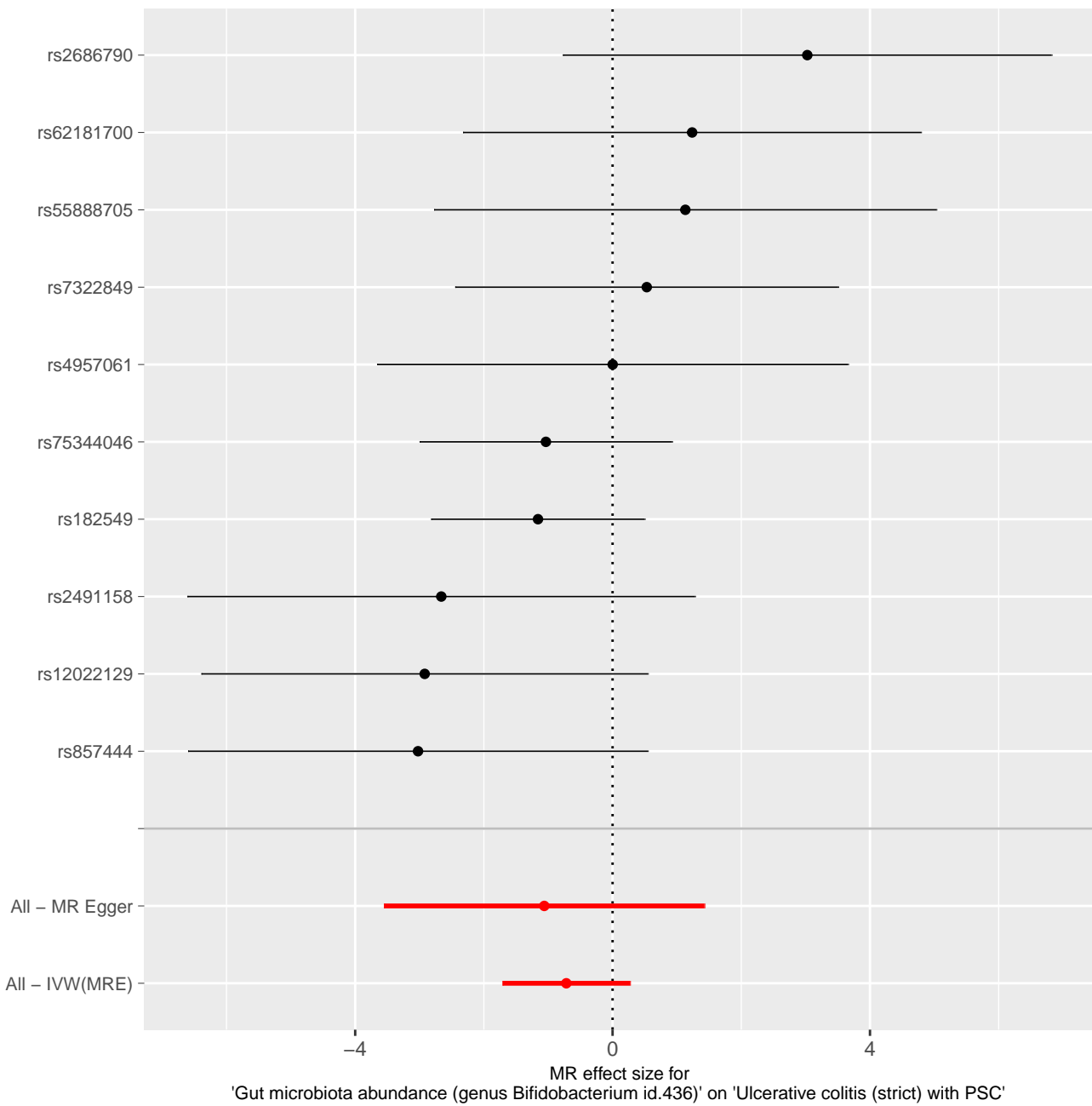


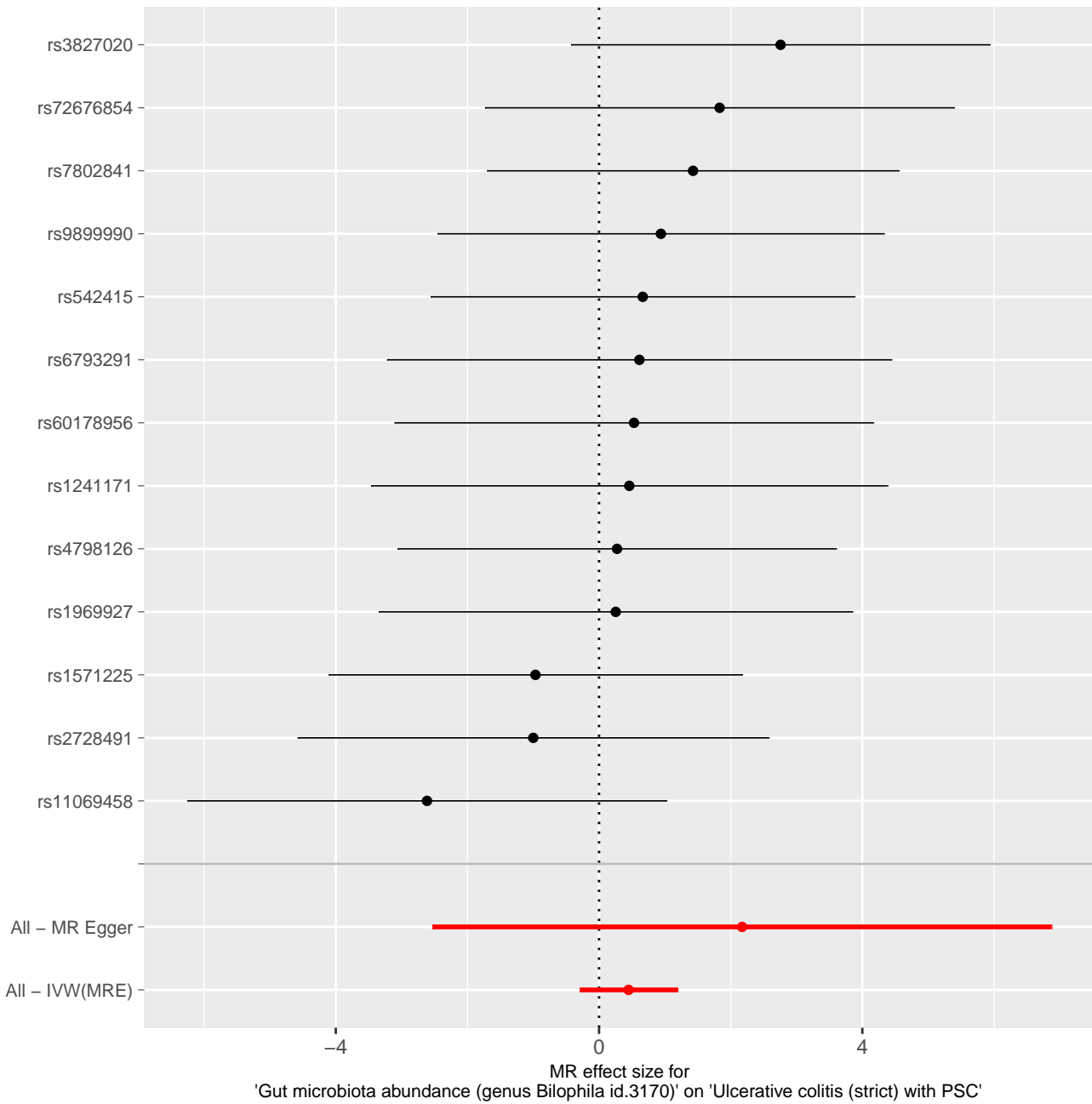


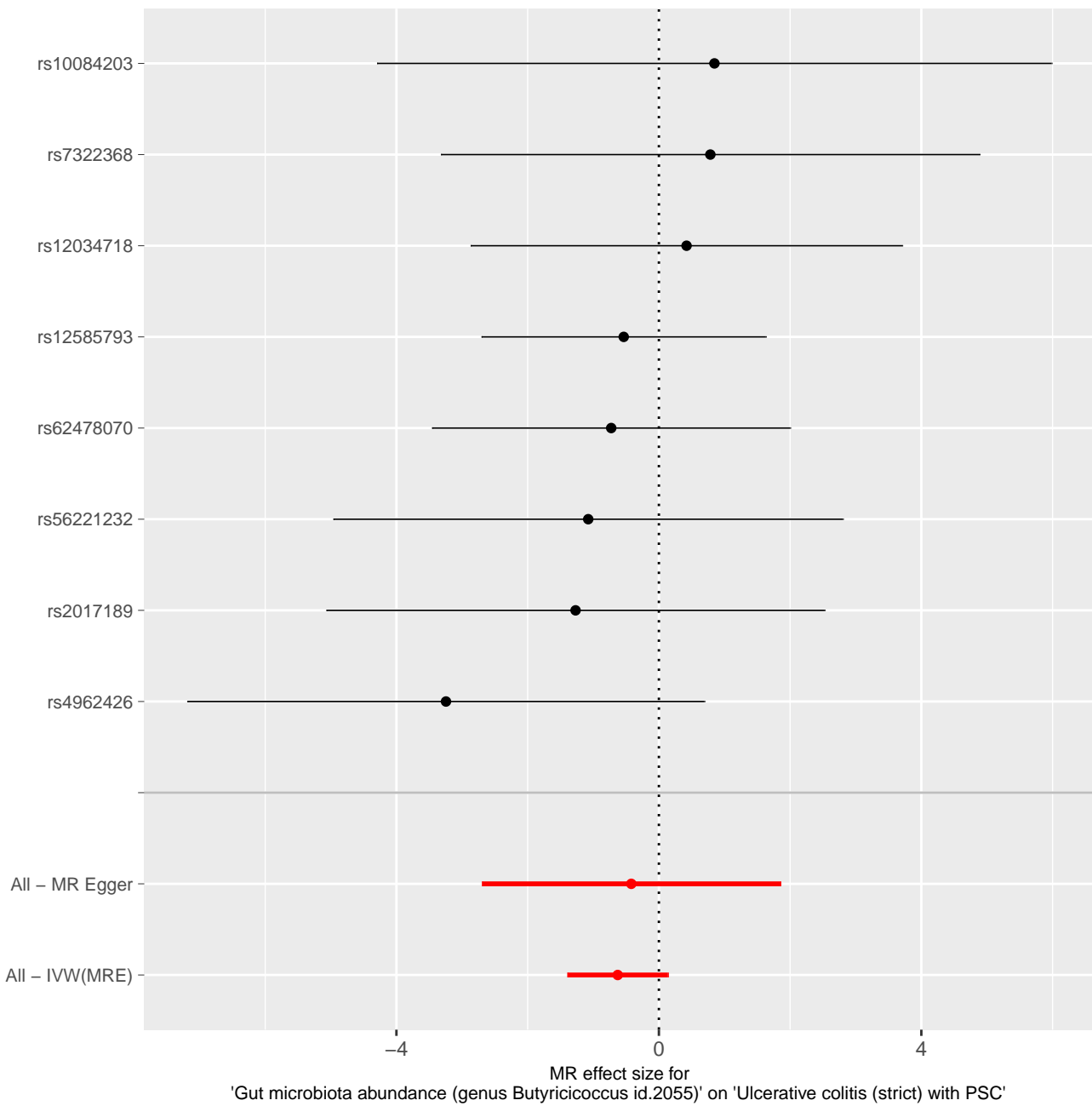


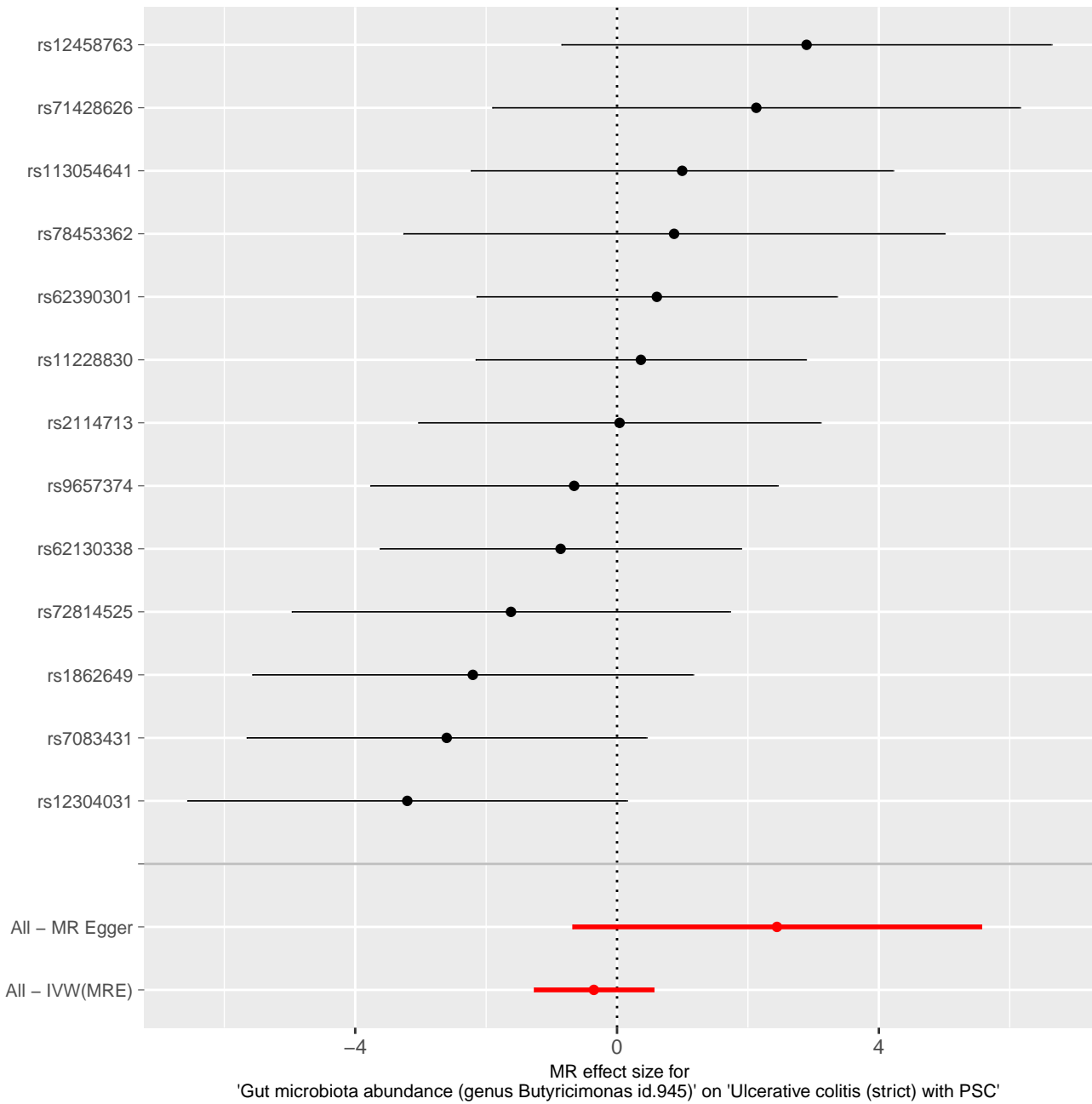




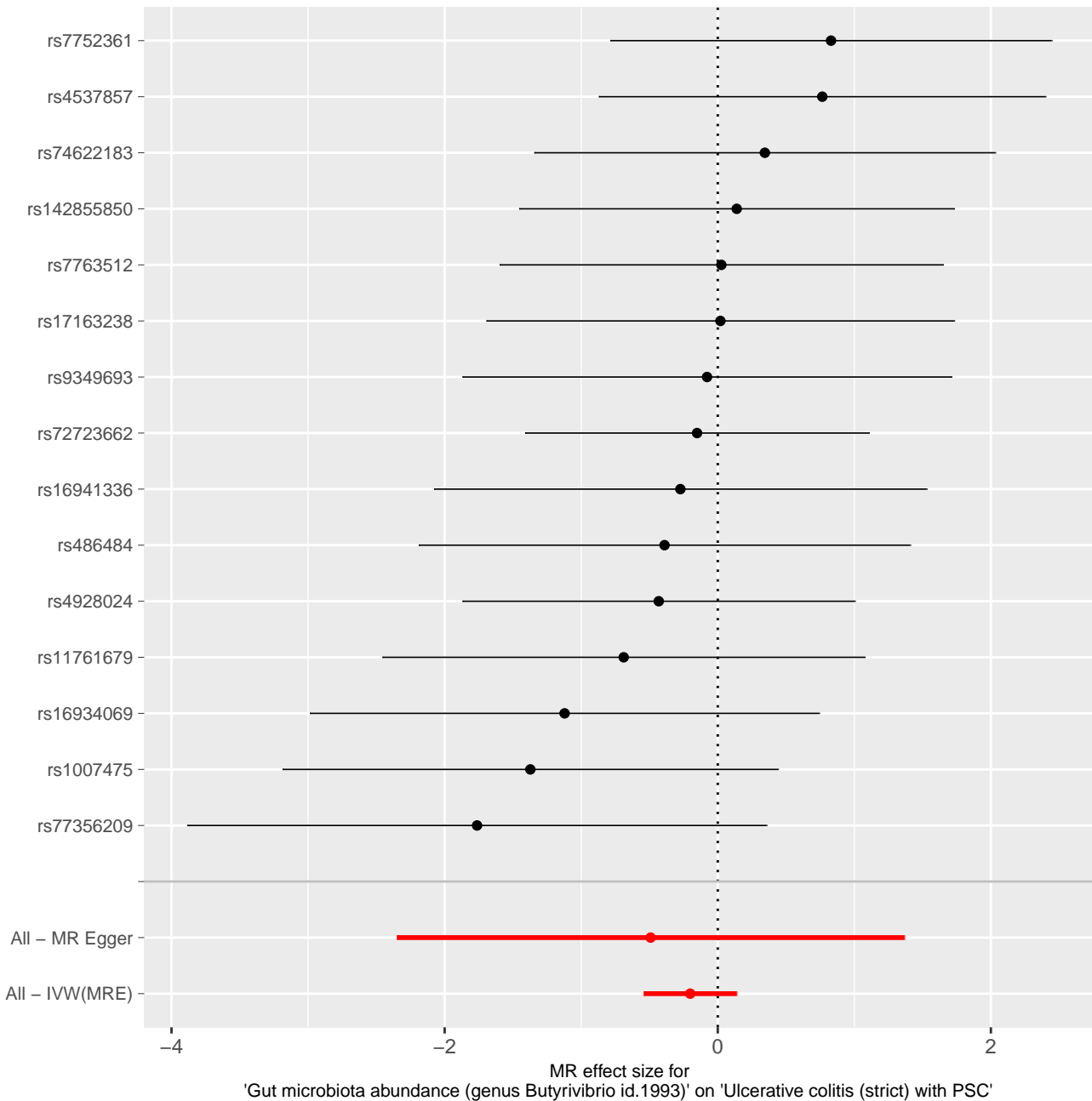


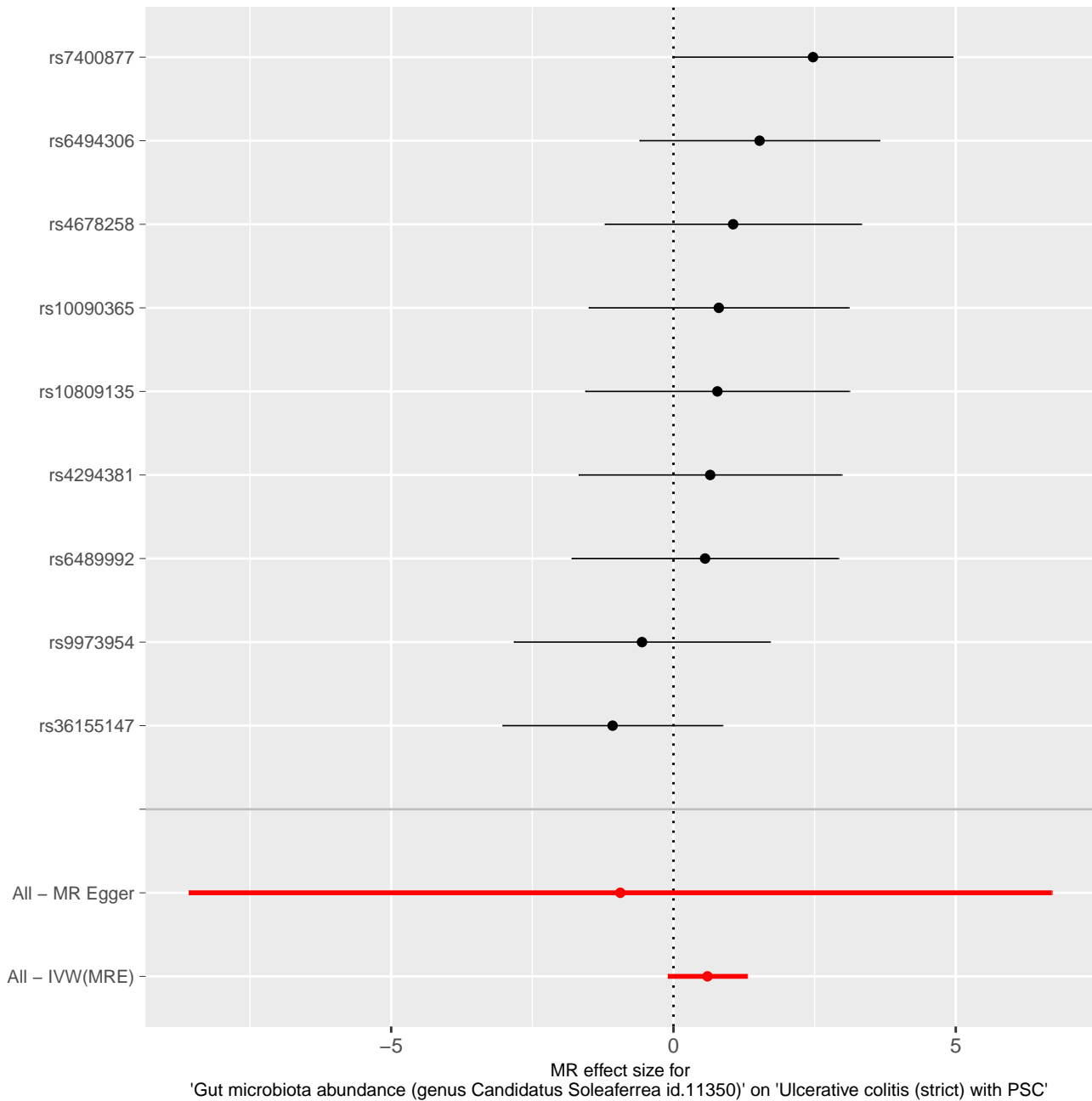


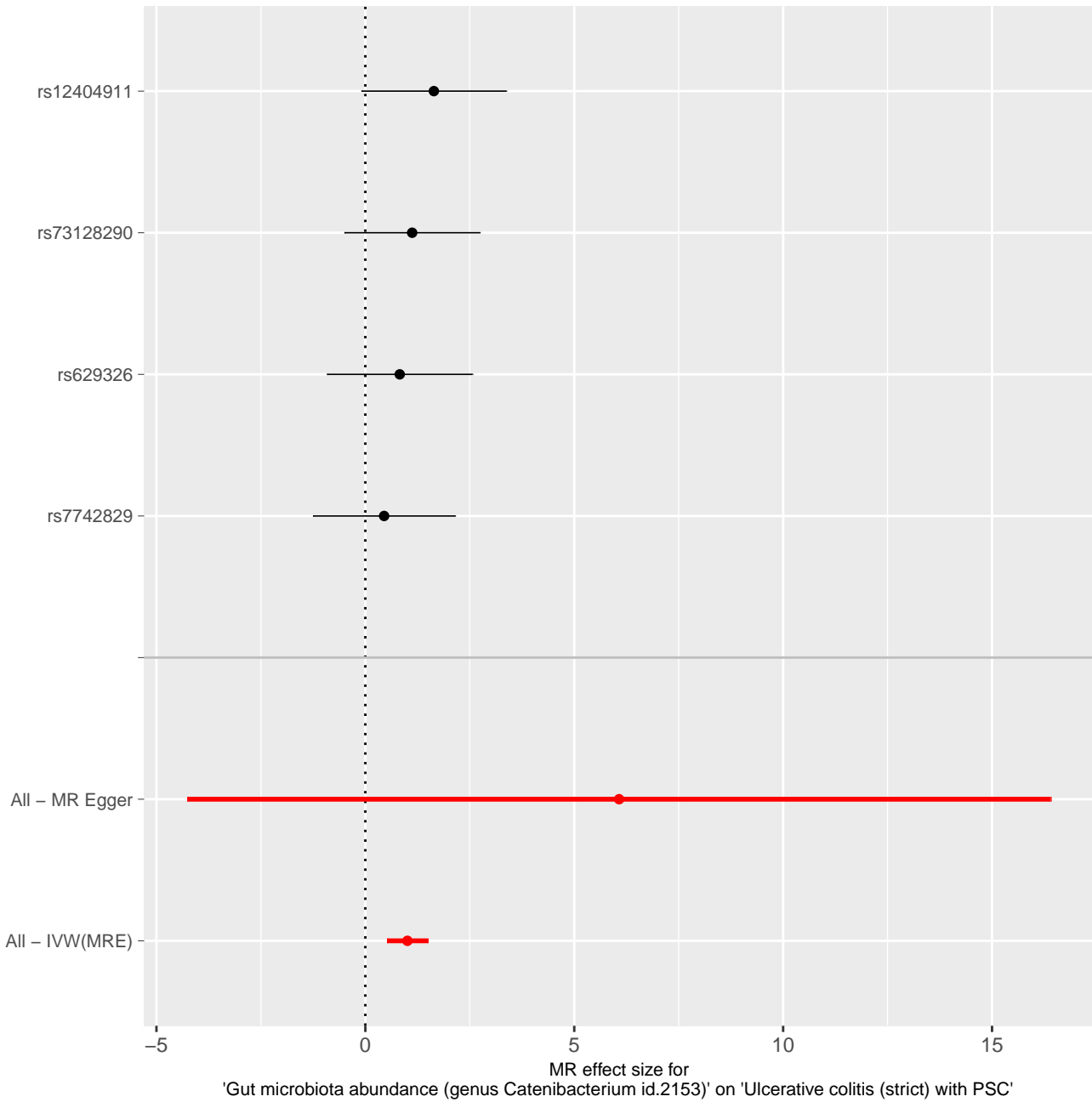


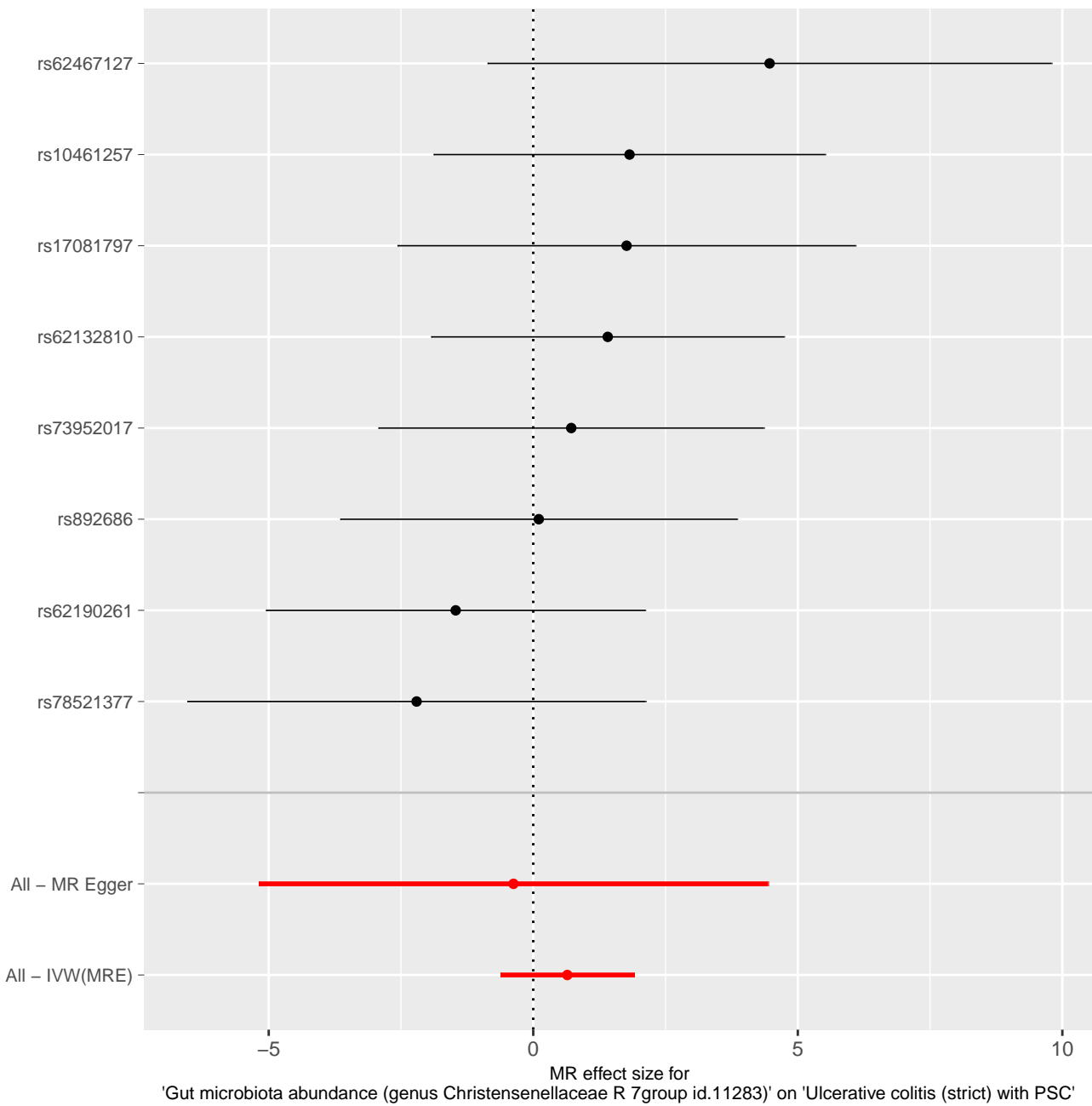


Batch 649 : Gut microbiota abundance (genus Butyrivibrio id.1993) on Ulcerative colitis (strict) with PSC

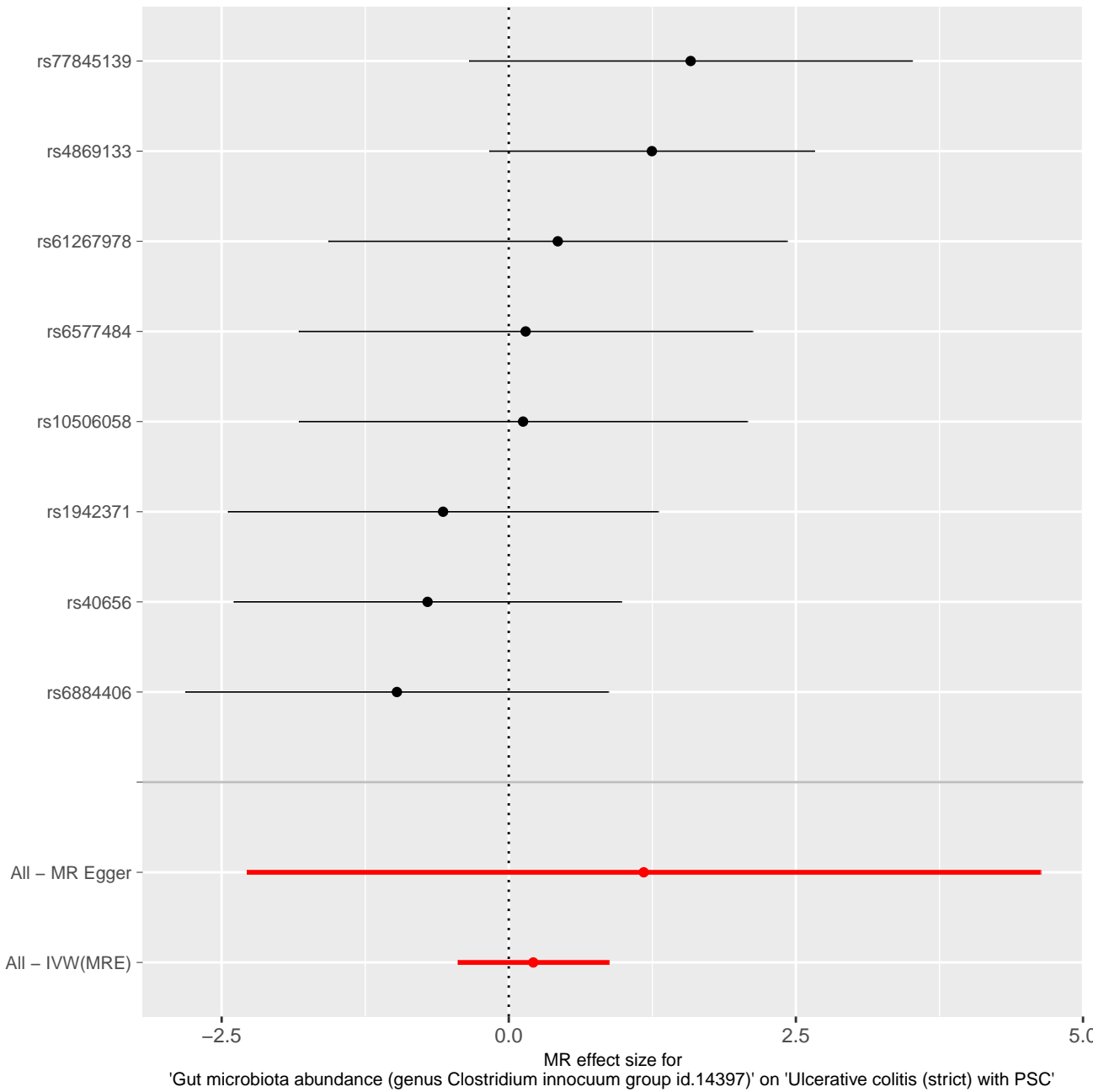




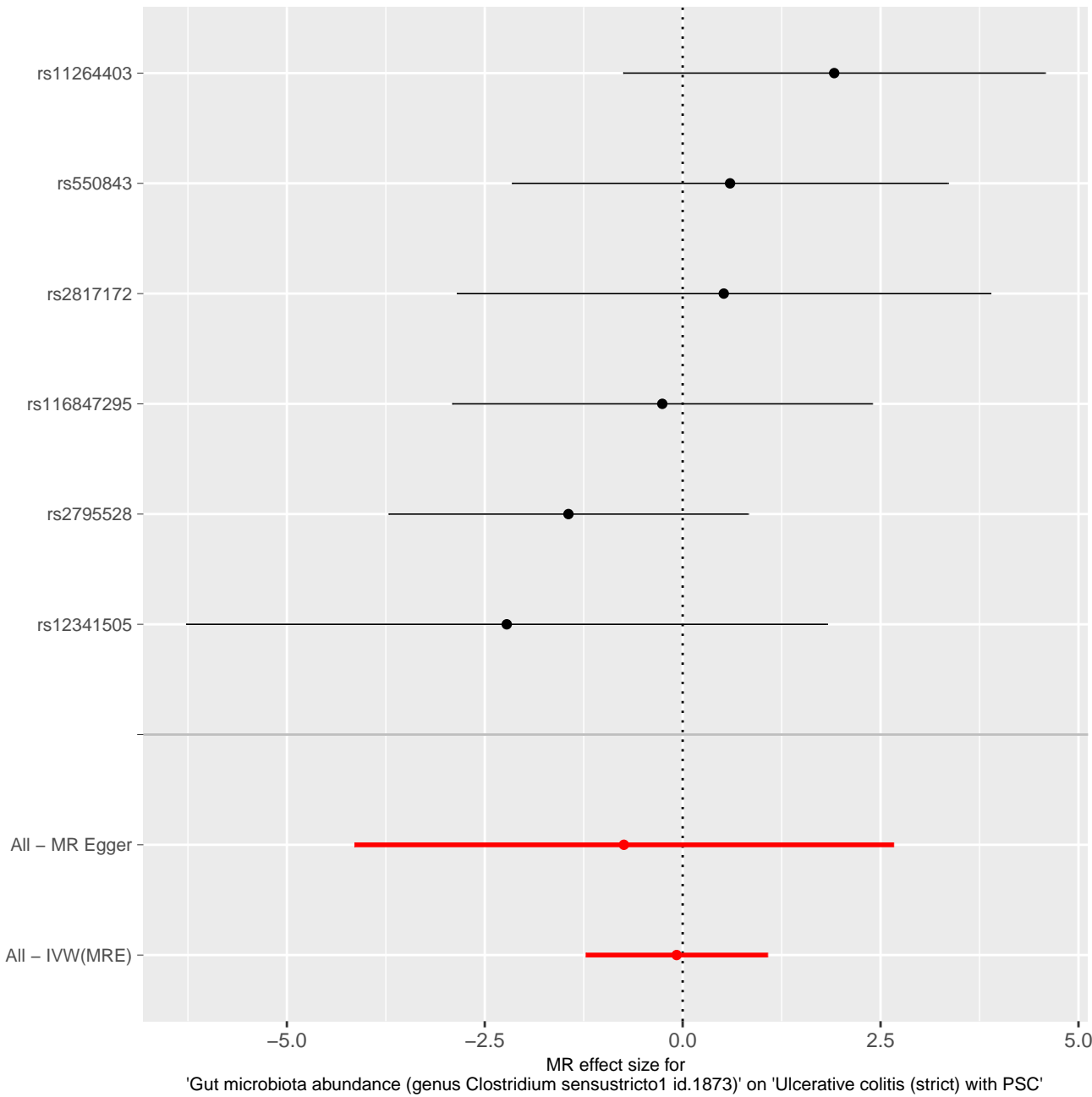




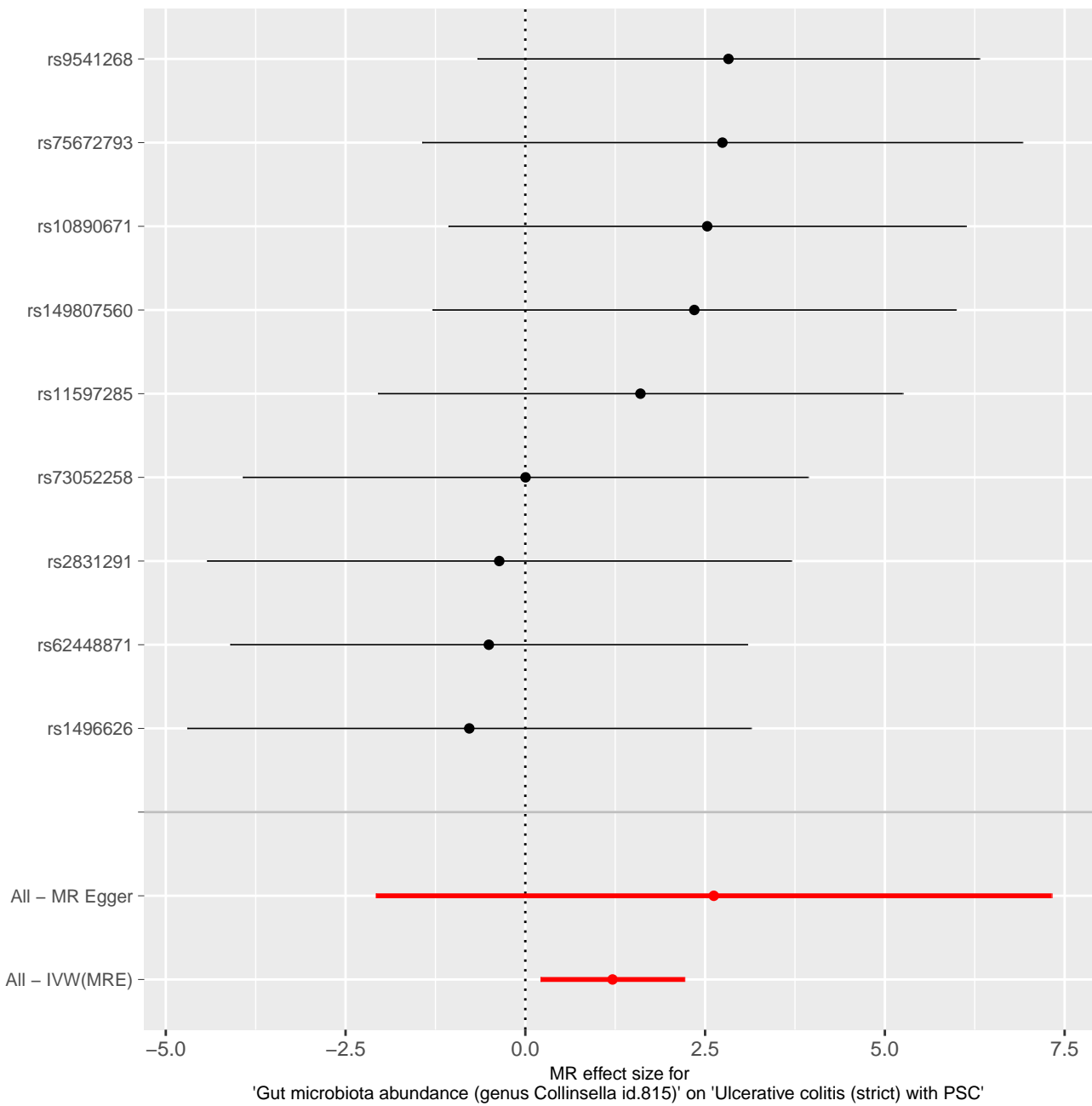
'Gut microbiota abundance (genus Christensenellaceae R 7group id.11283)' on 'Ulcerative colitis (strict) with PSC'

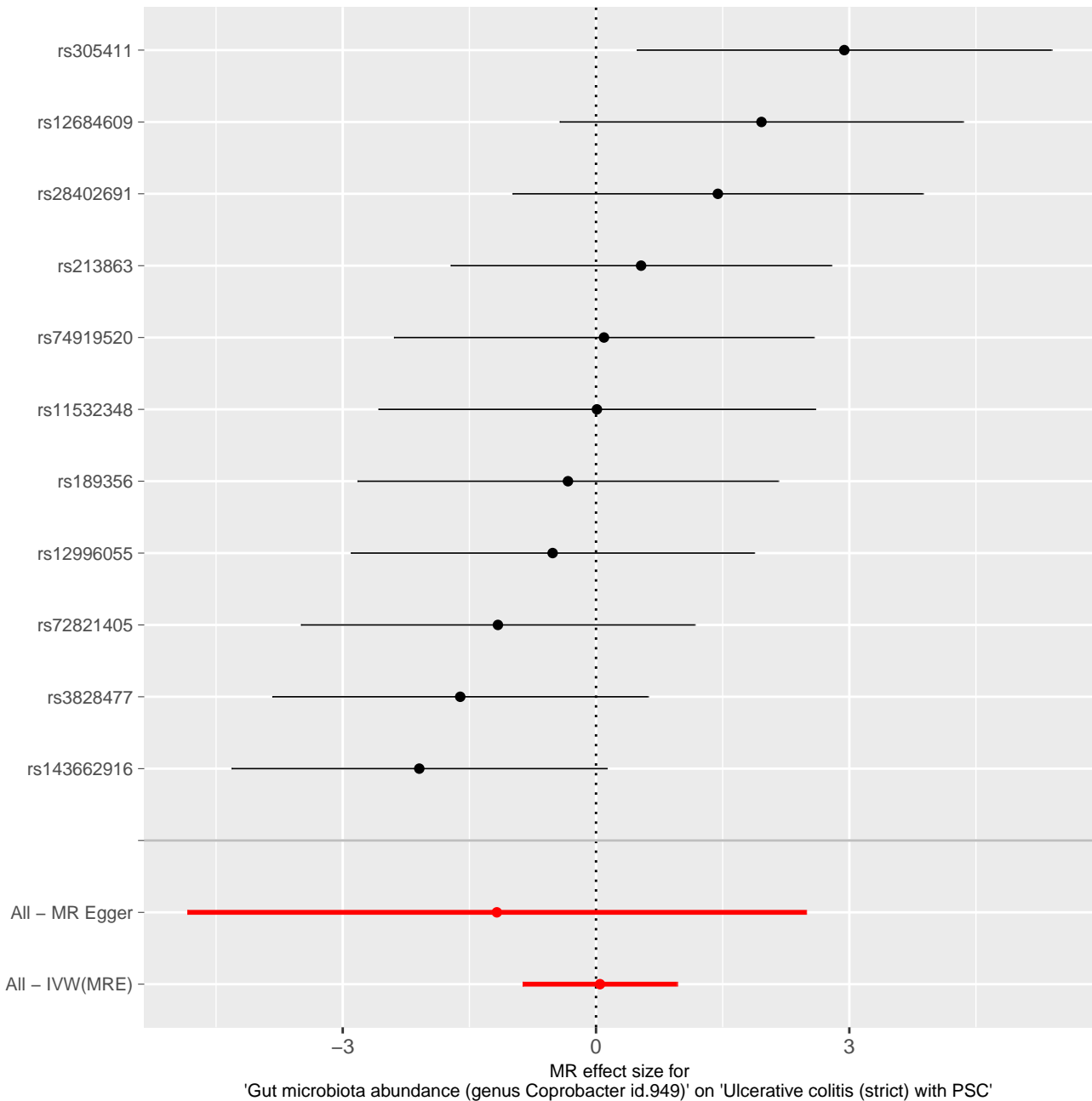


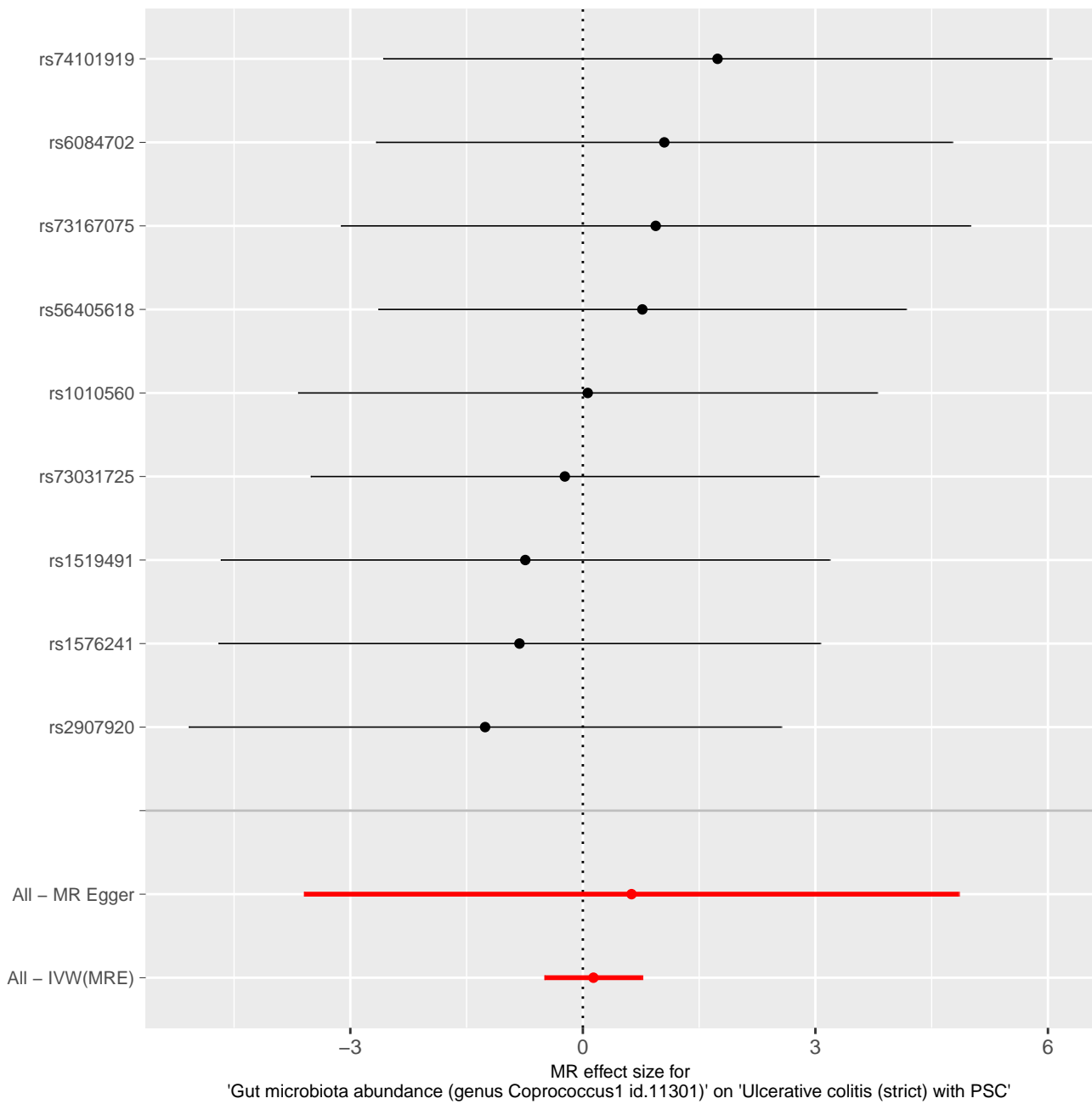
'Gut microbiota abundance (genus Clostridium innocuum group id.14397) on 'Ulcerative colitis (strict) with PSC'

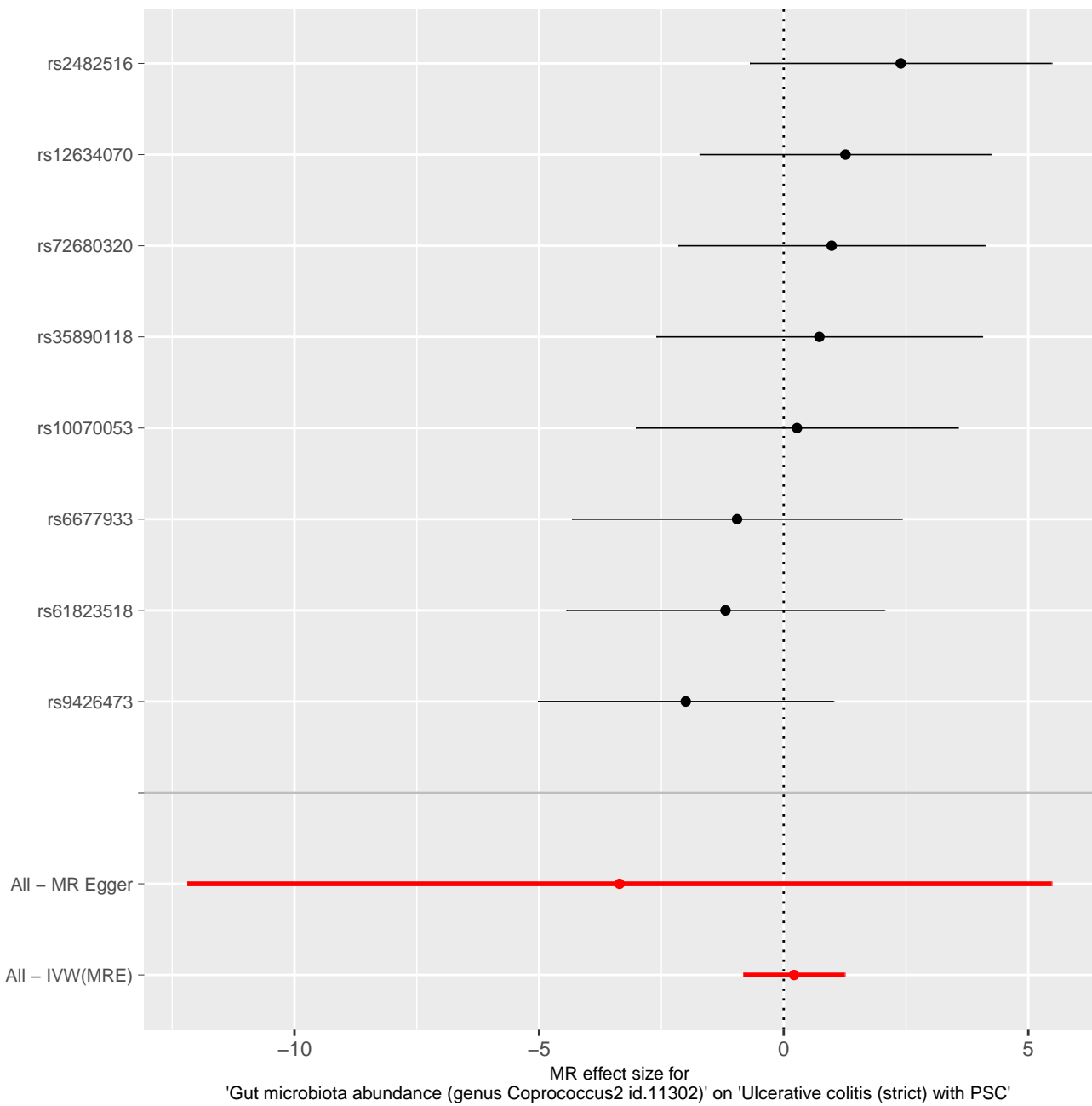


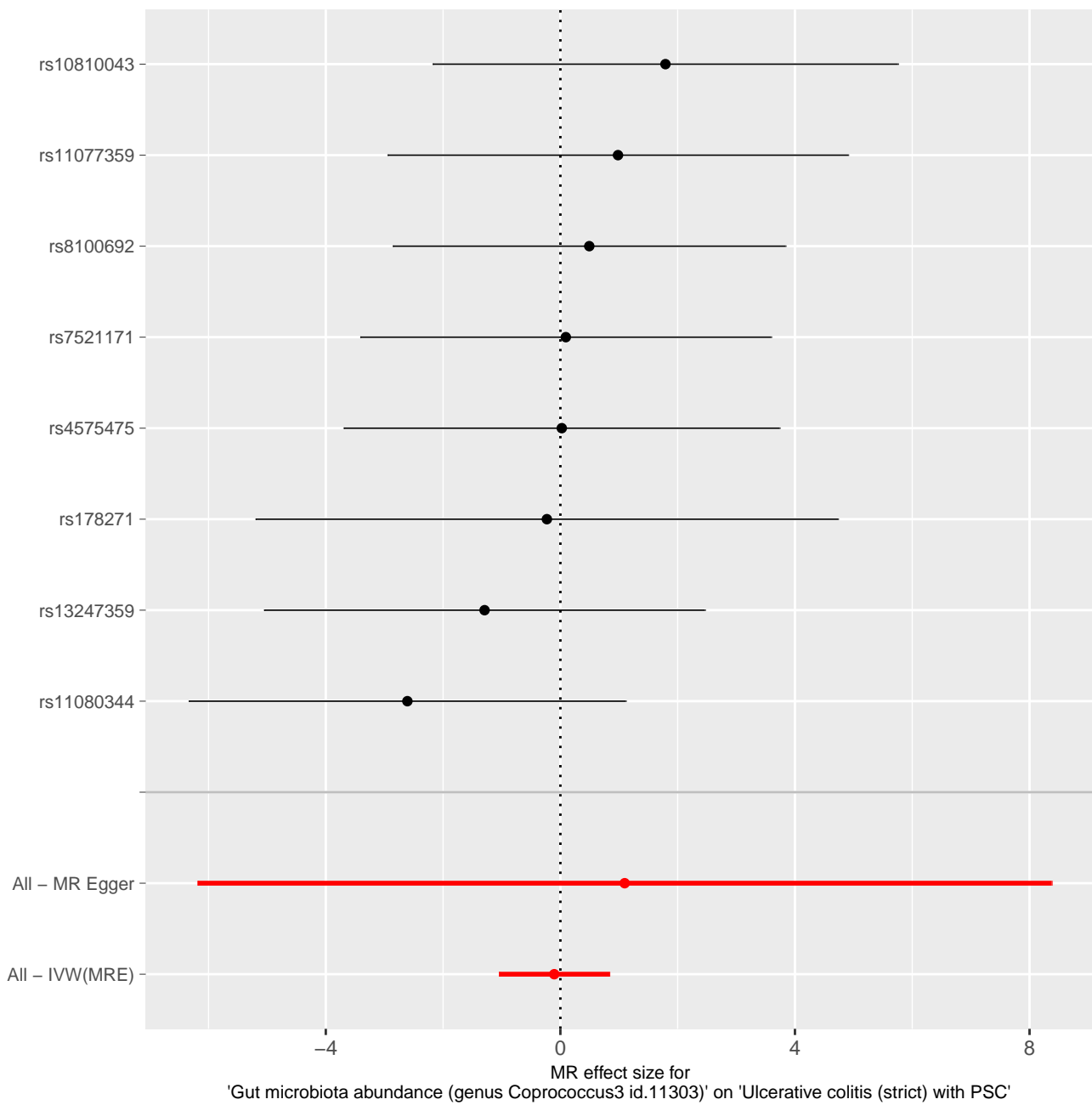
MR effect size for
'Gut microbiota abundance (genus Clostridium sensustricto1 id.1873)' on 'Ulcerative colitis (strict) with PSC'

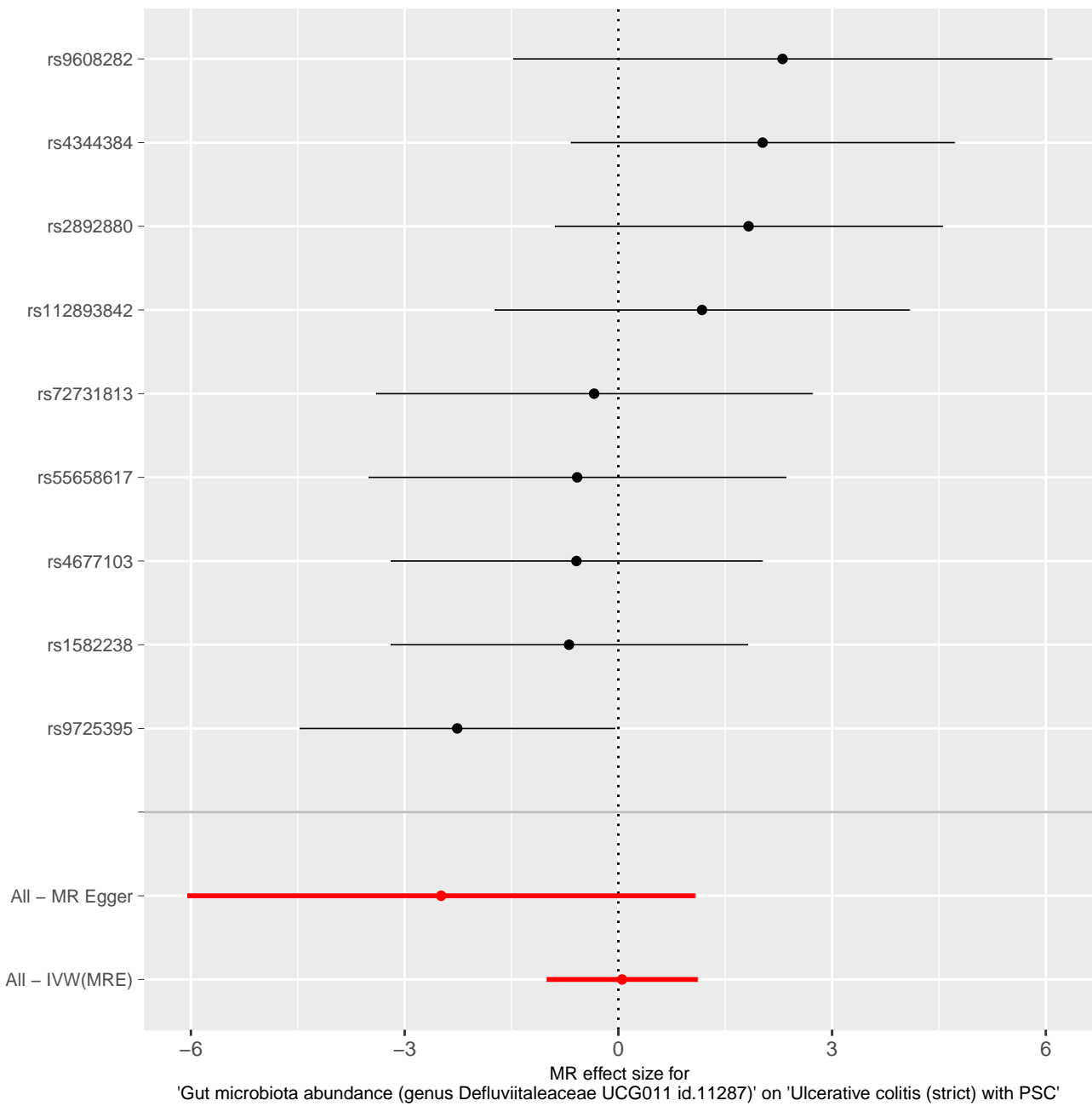


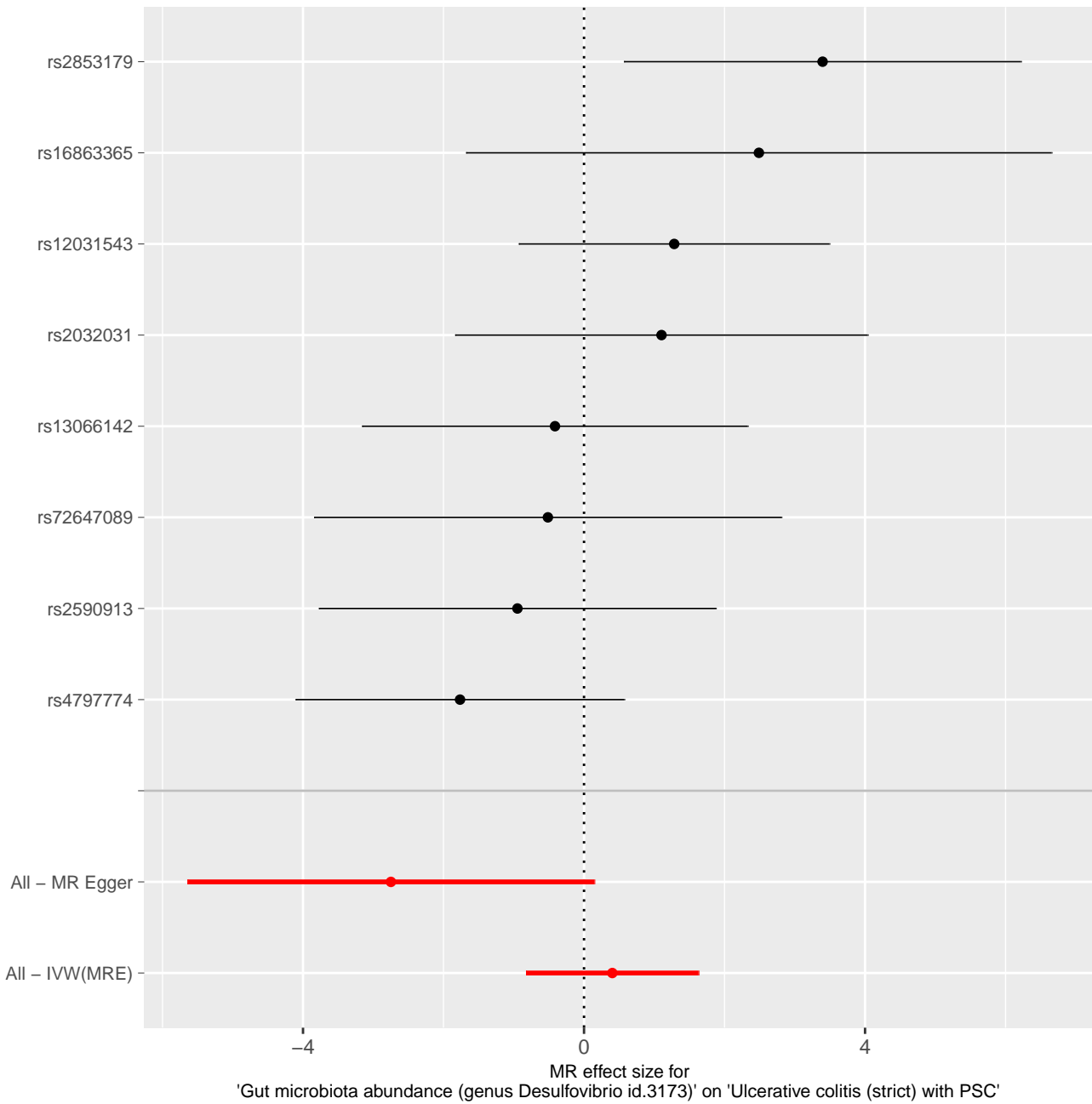




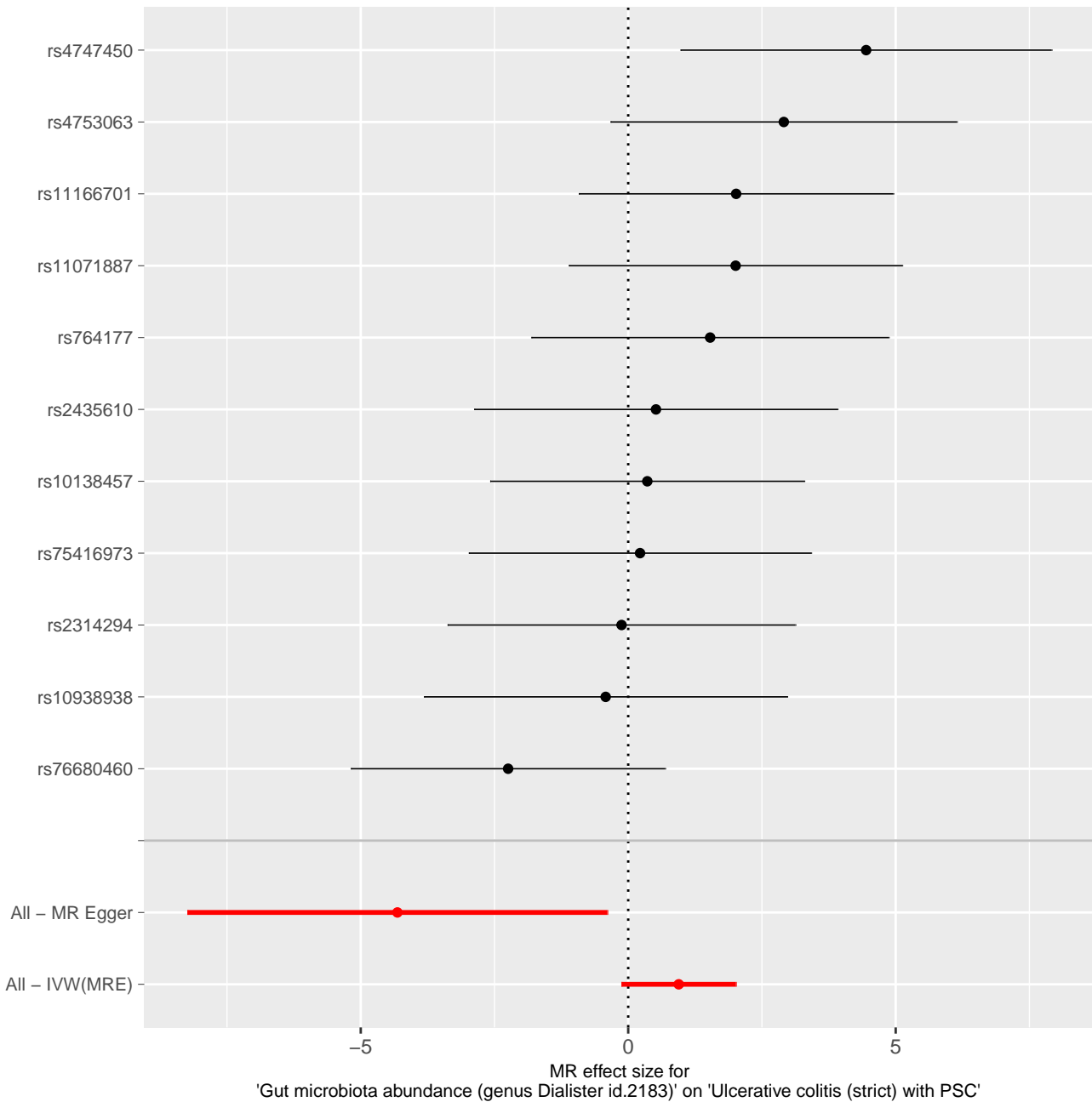


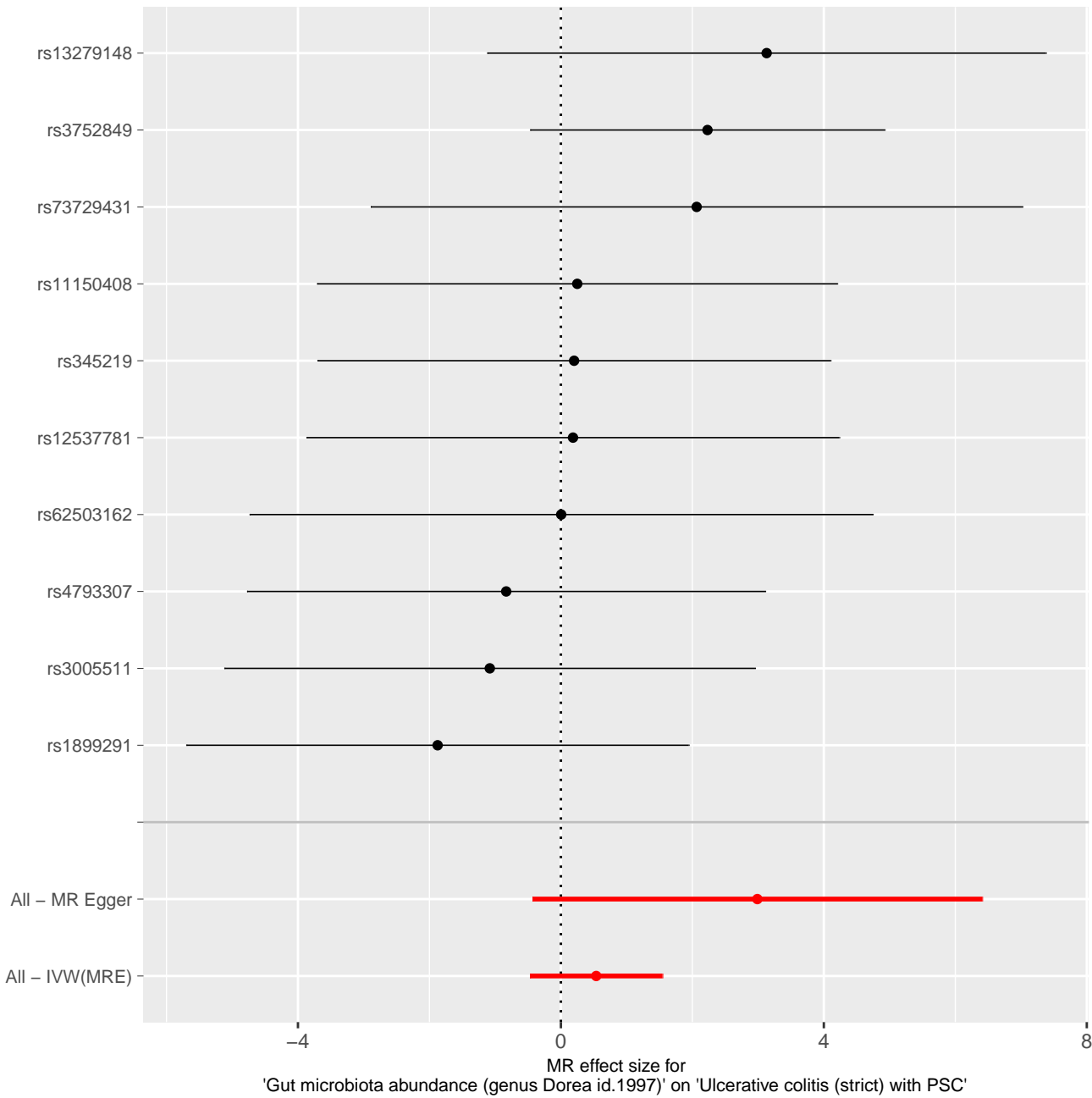


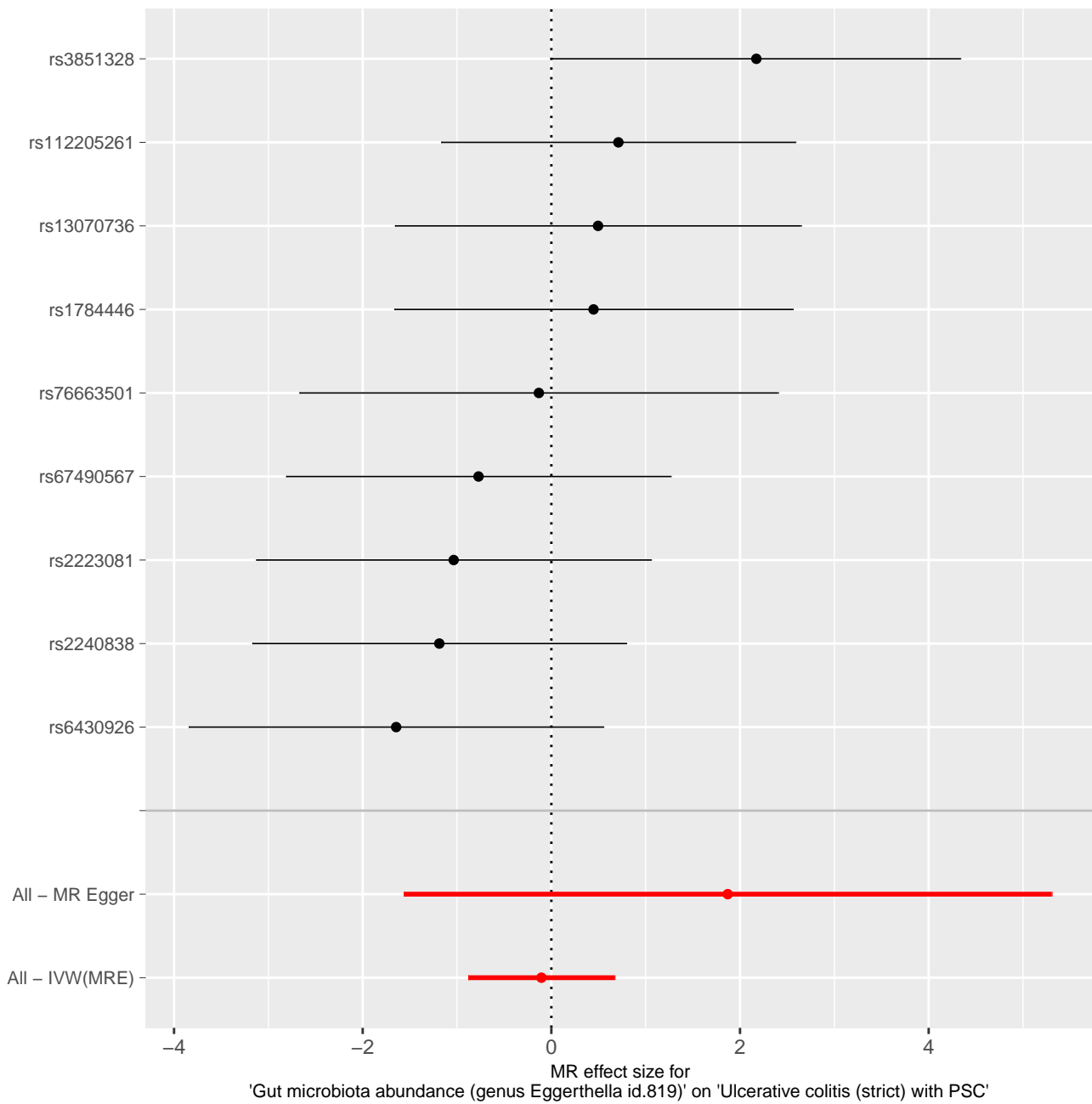


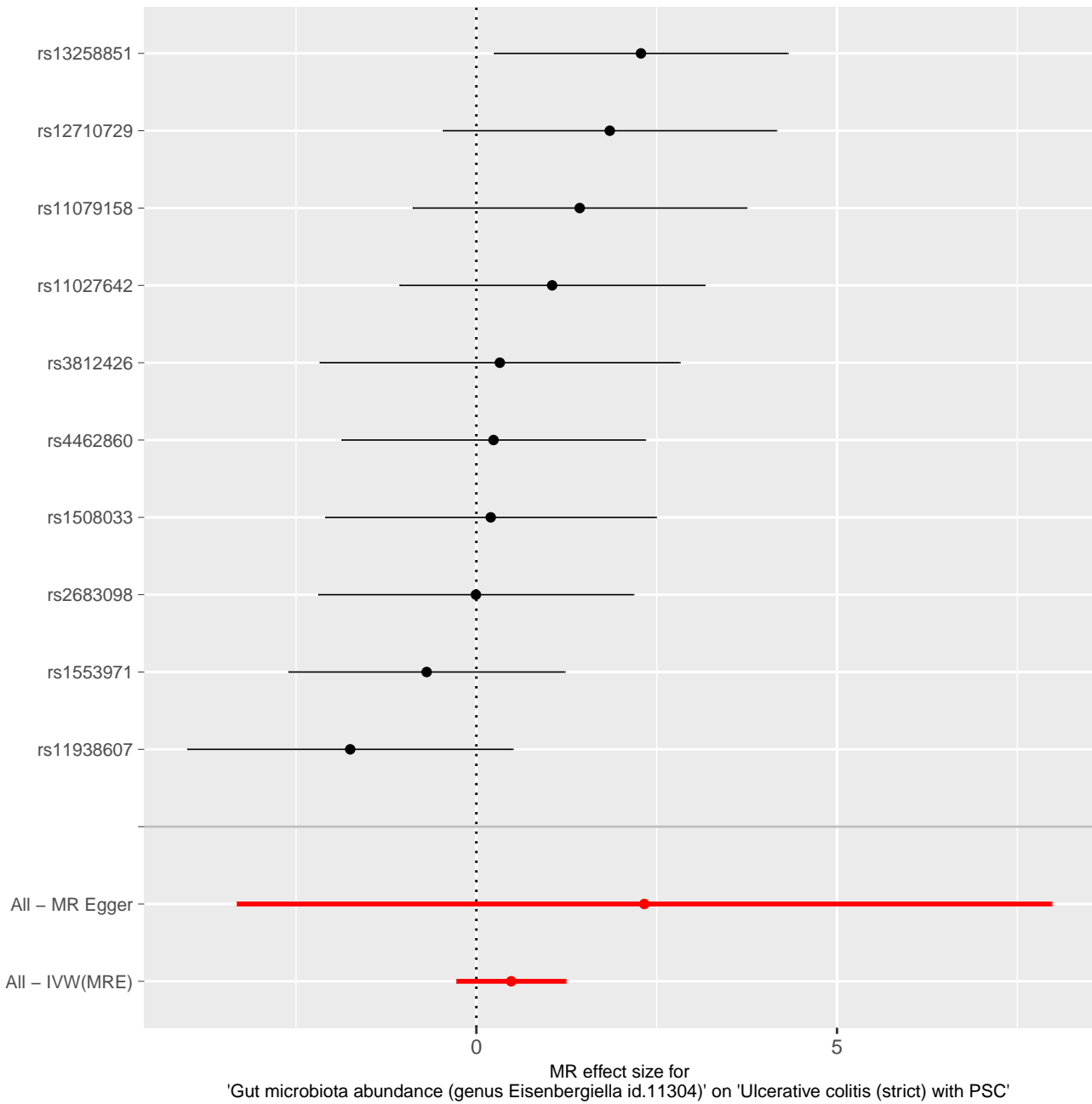


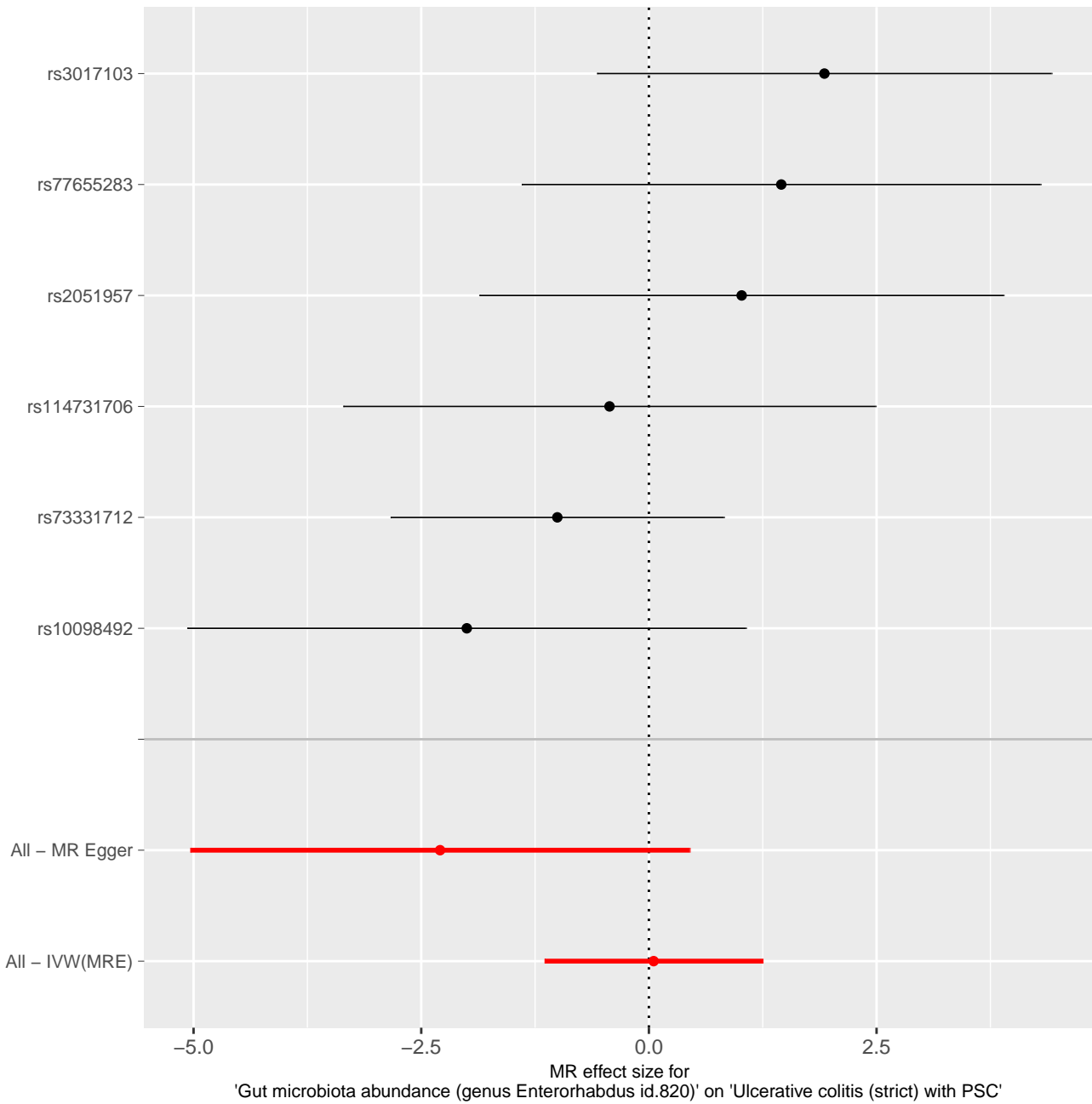
Batch 662 : Gut microbiota abundance (genus Dialister id.2183) on Ulcerative colitis (strict) with PSC

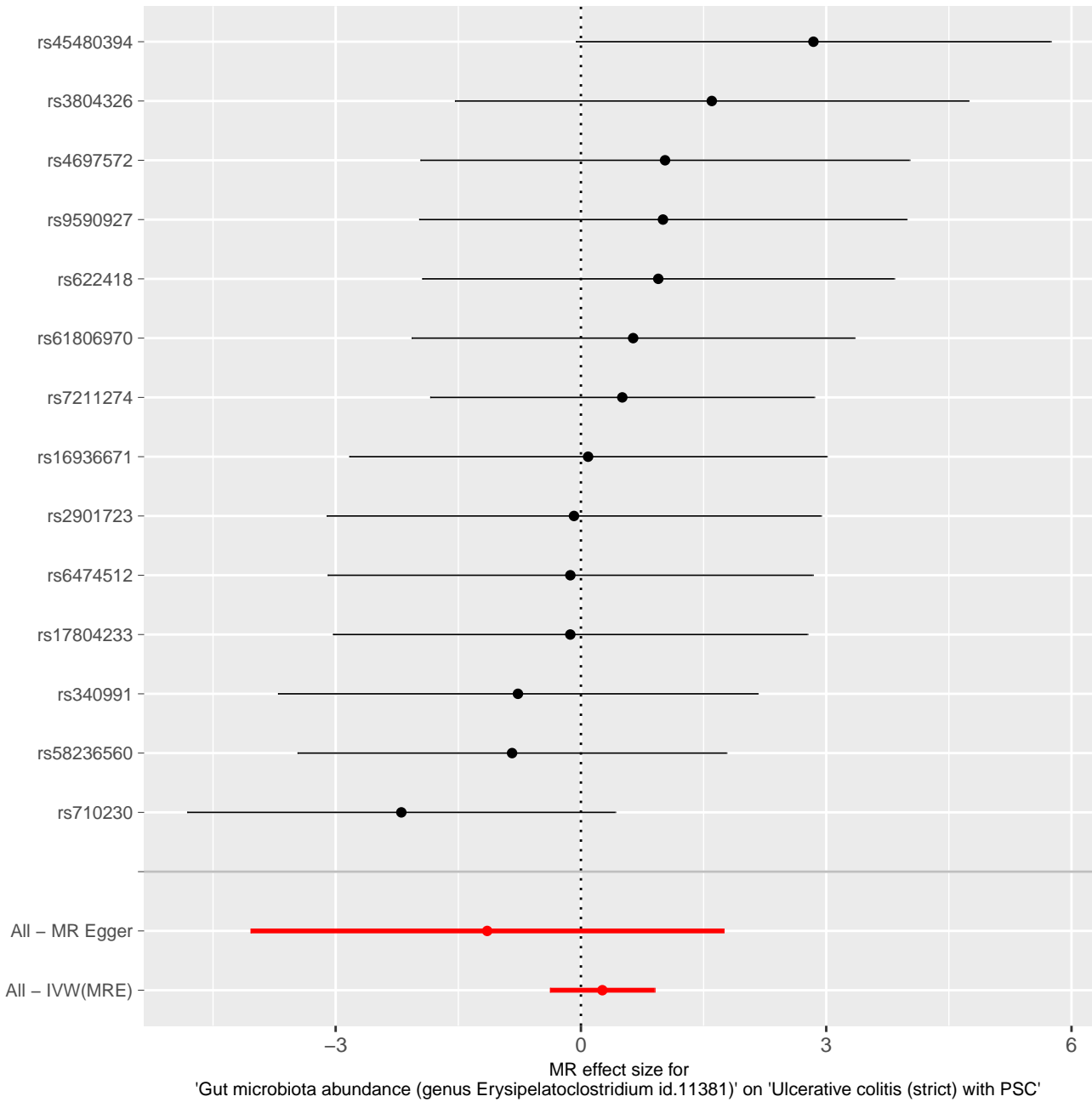




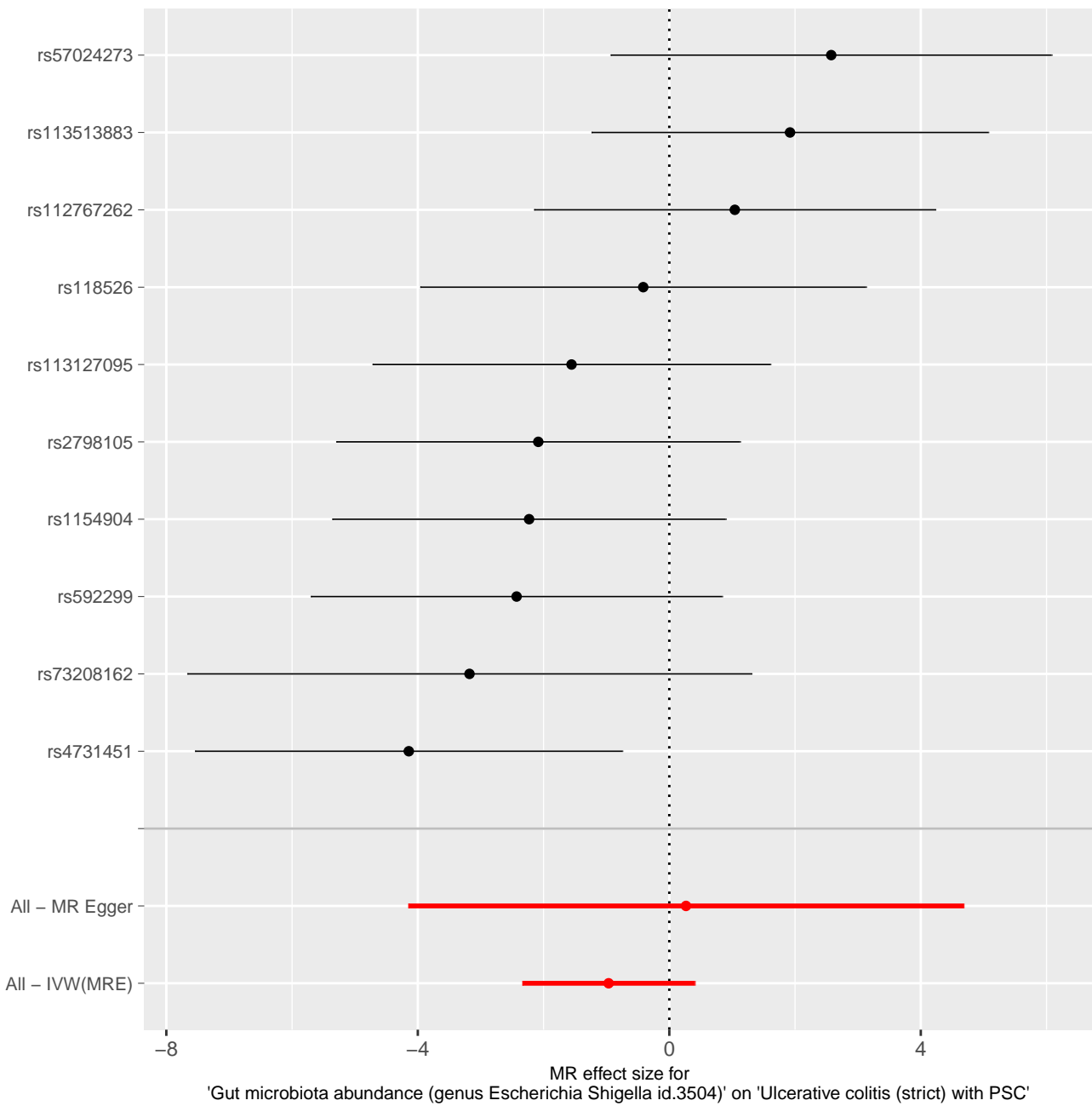


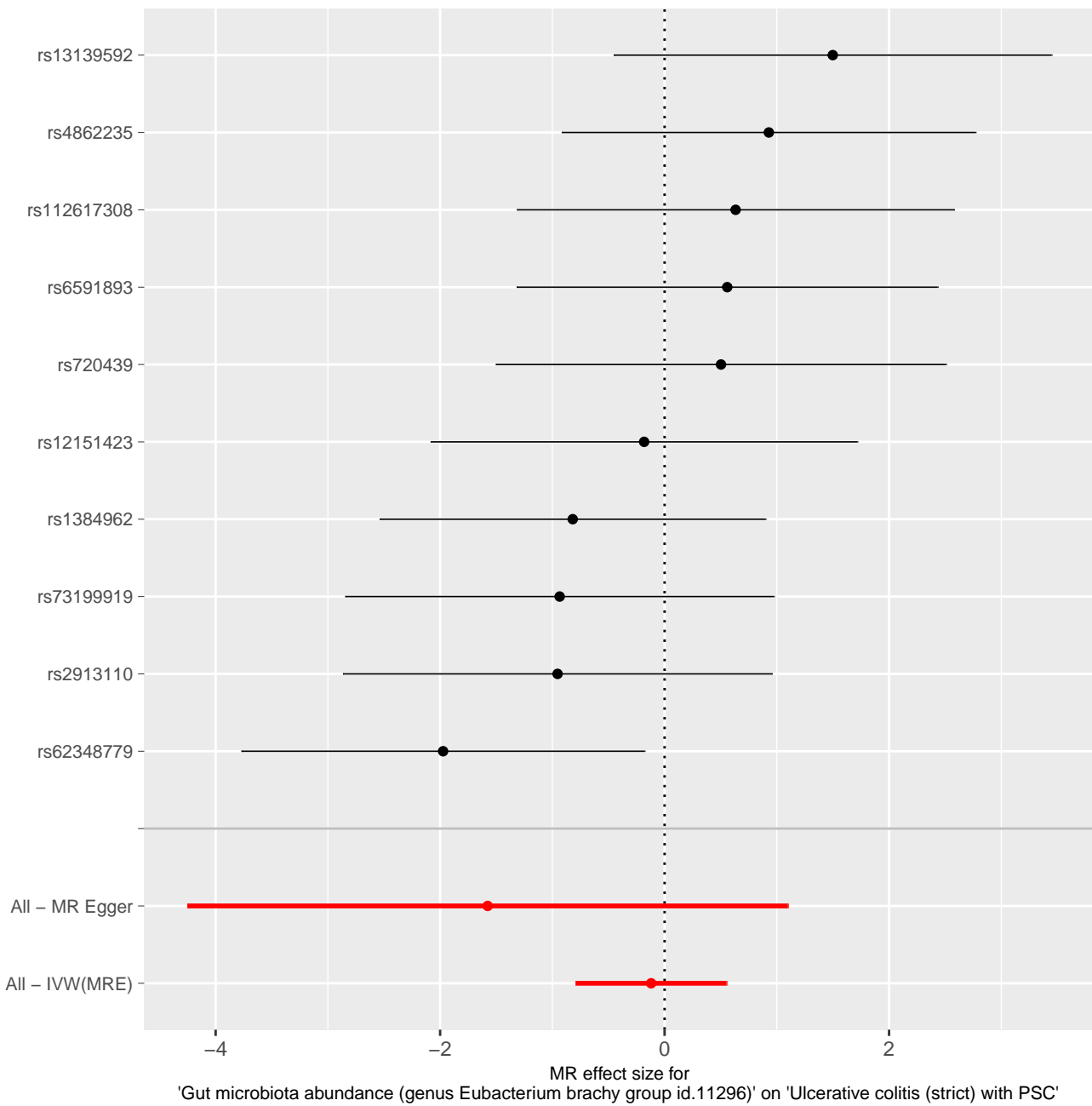


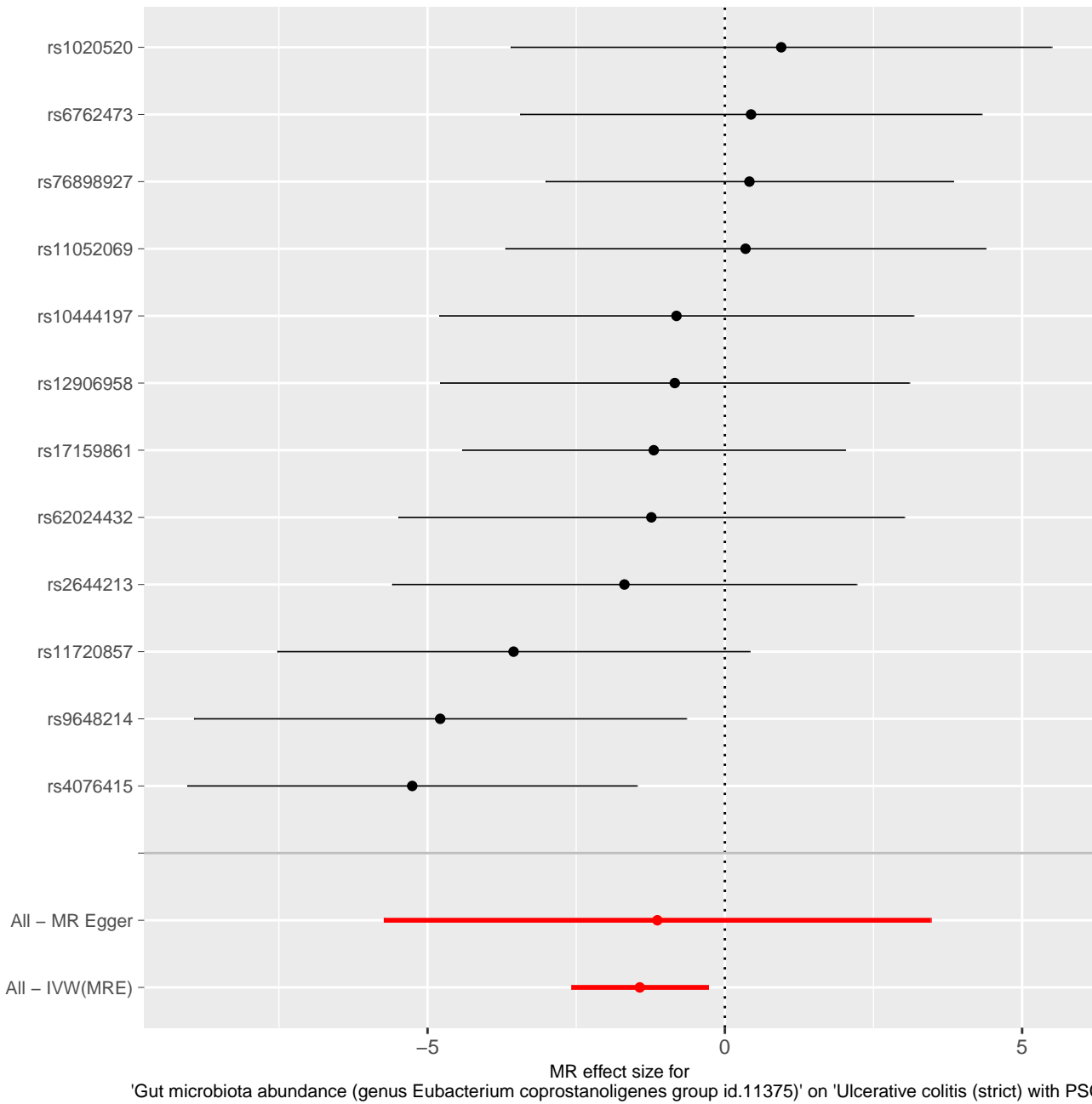


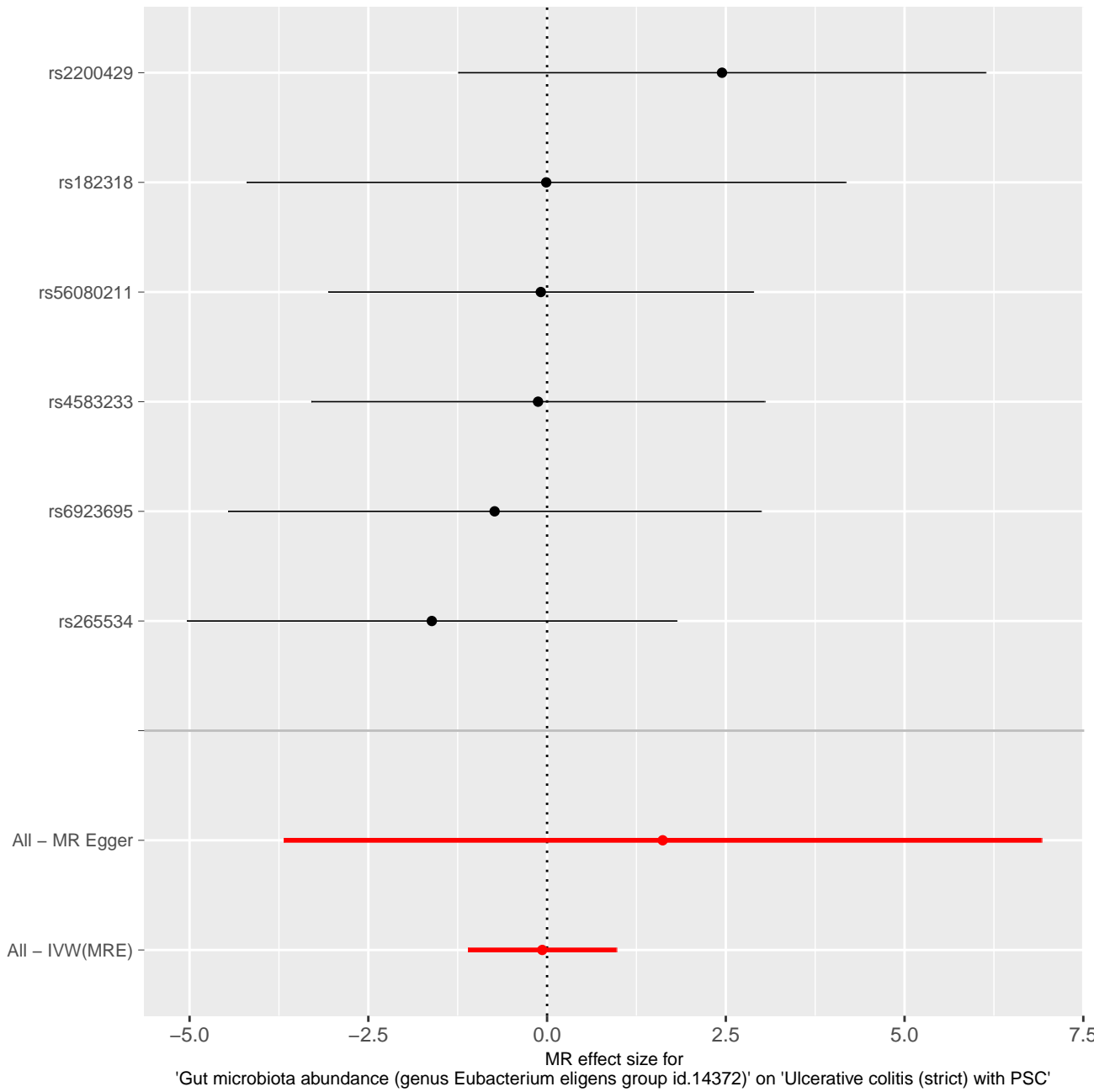


MR effect size for 'Gut microbiota abundance (genus Erysipelatoclostridium id.11381)' on 'Ulcerative colitis (strict) with PSC'

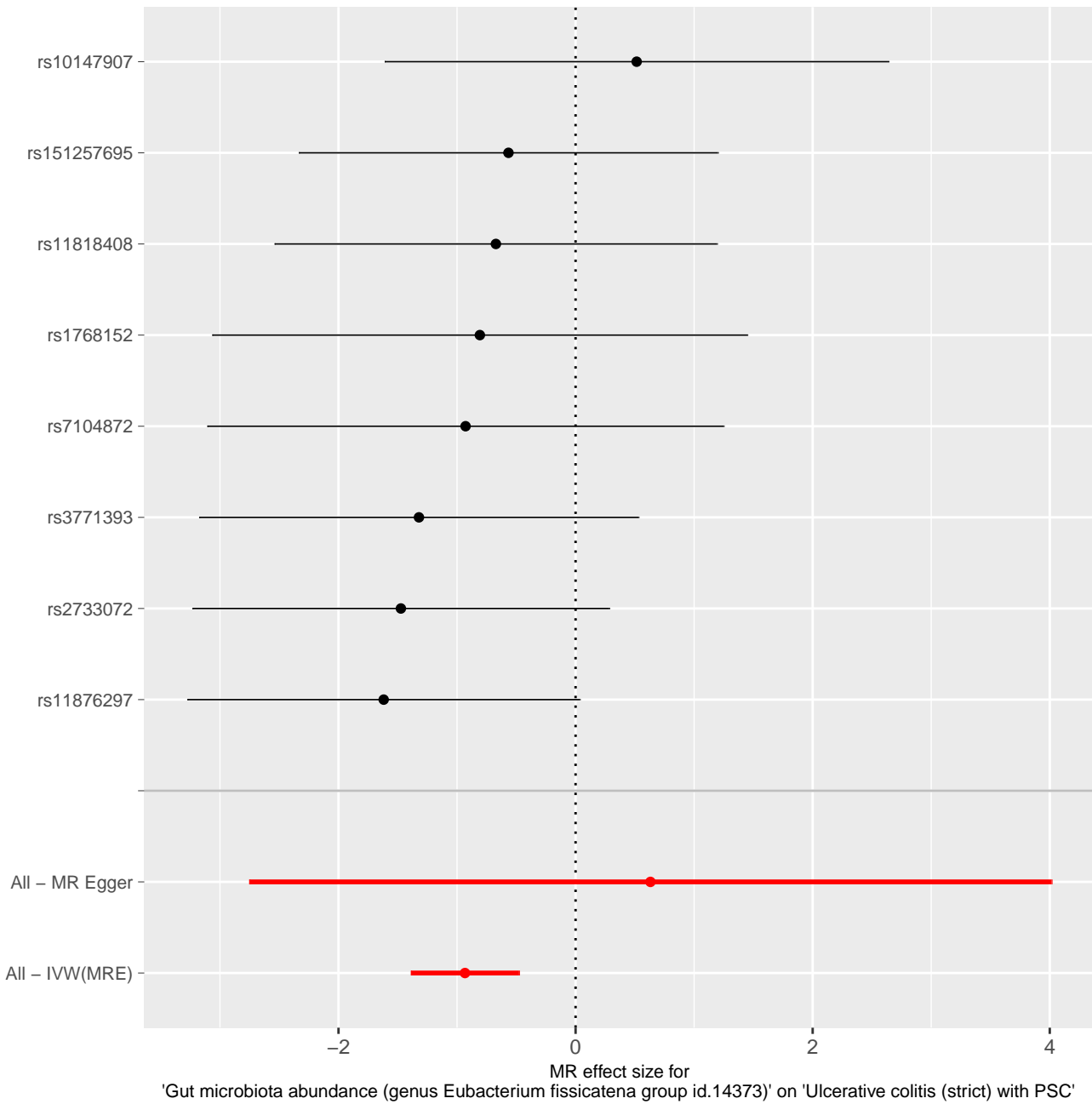


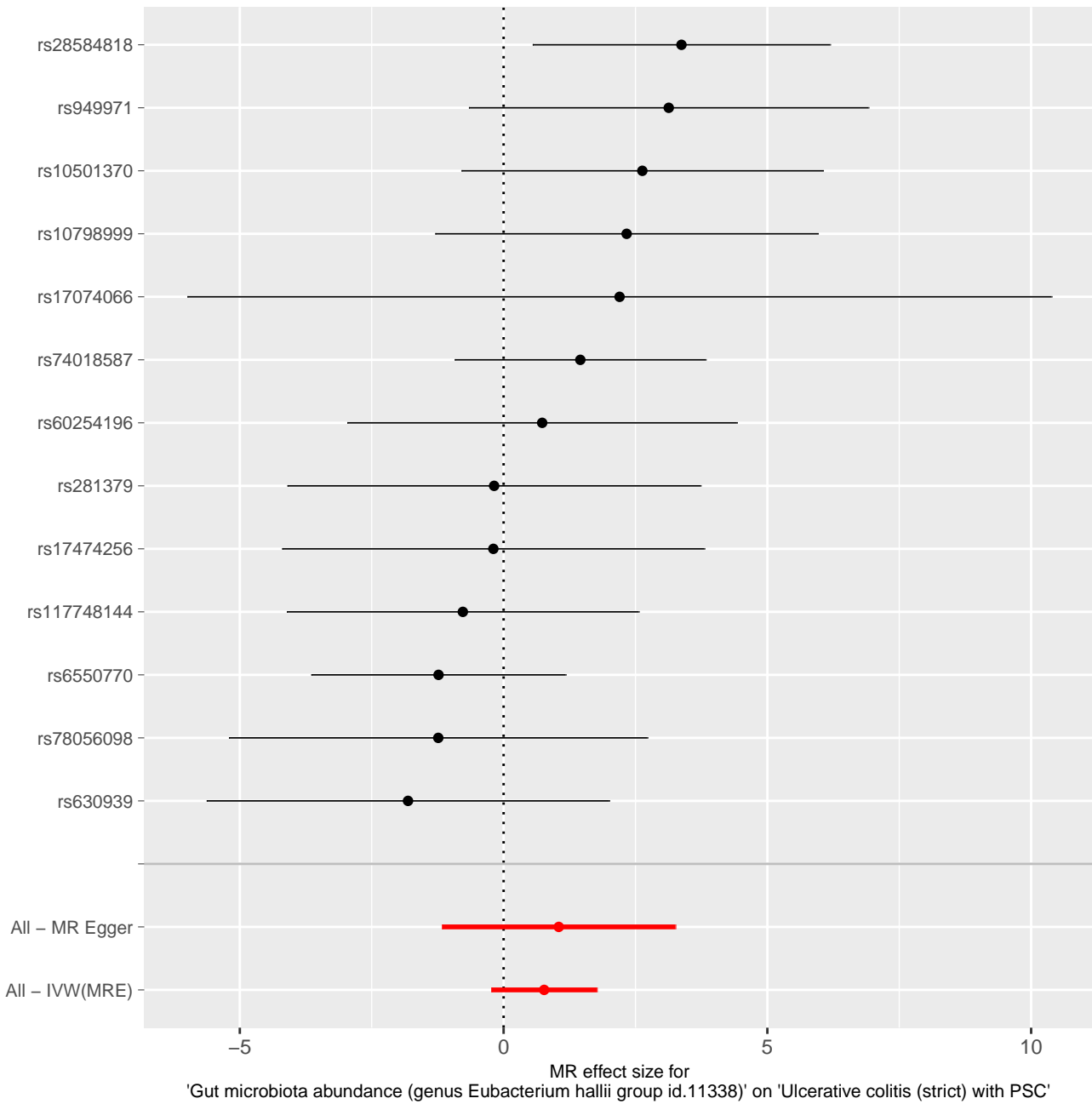


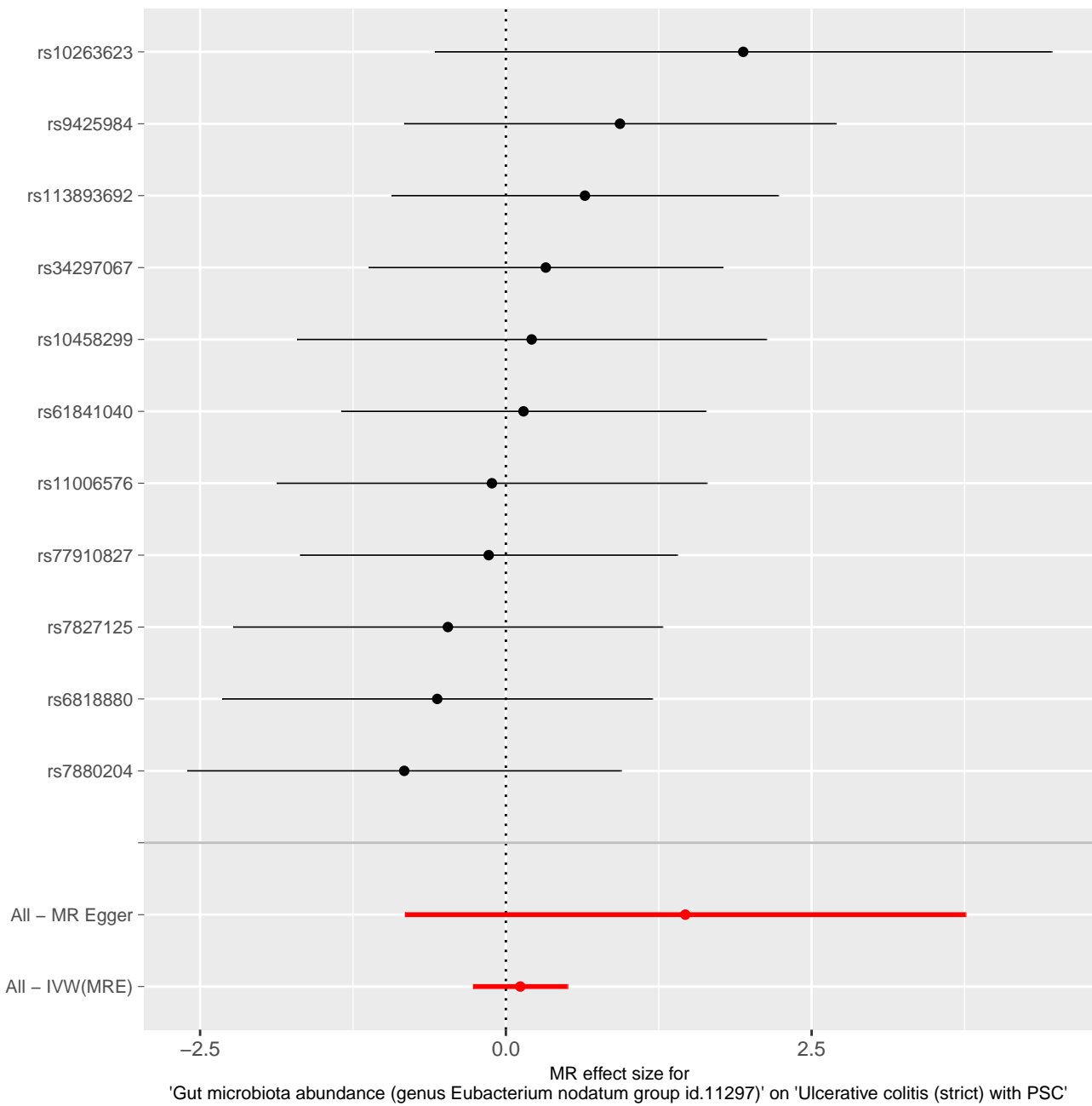


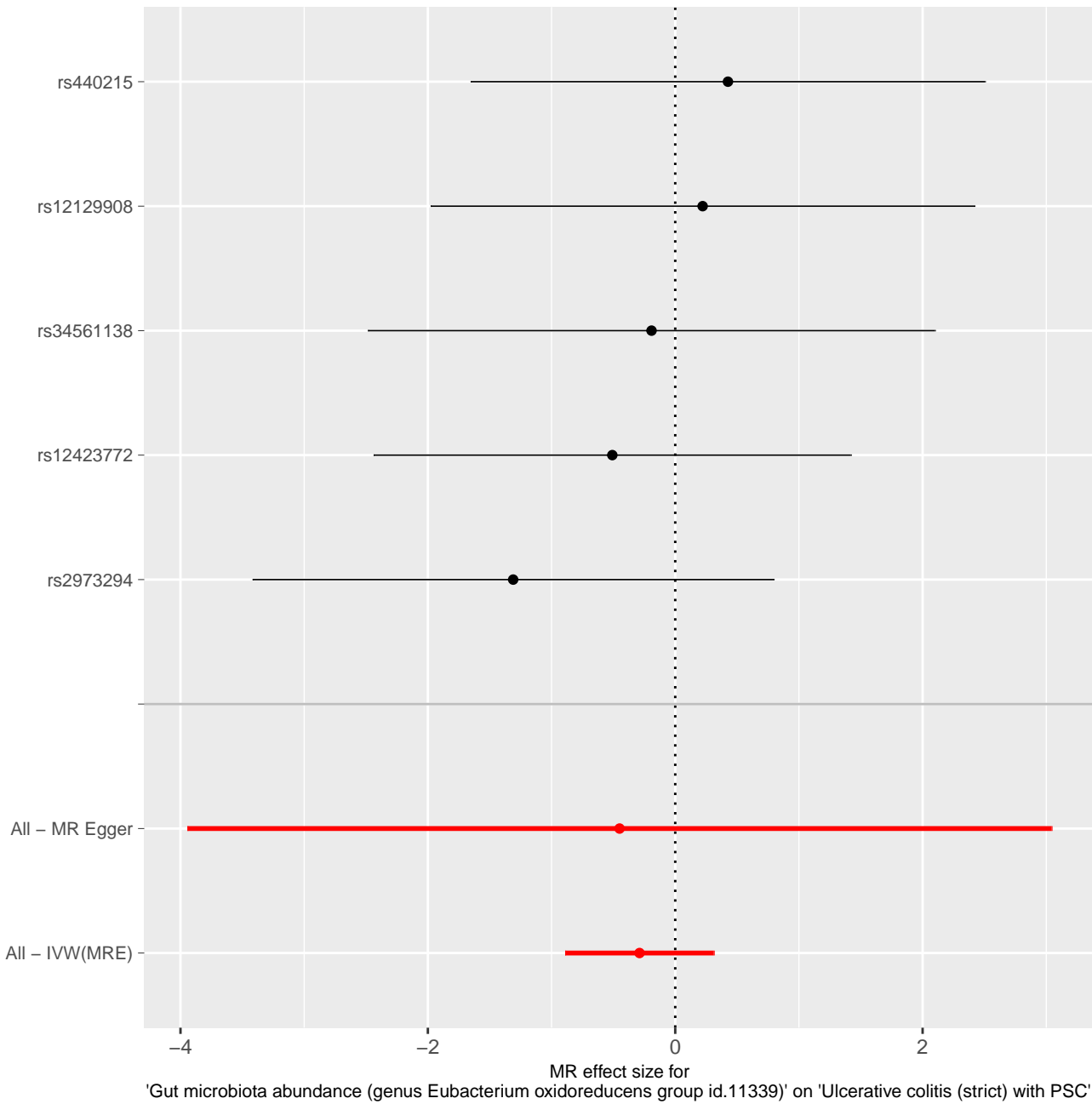


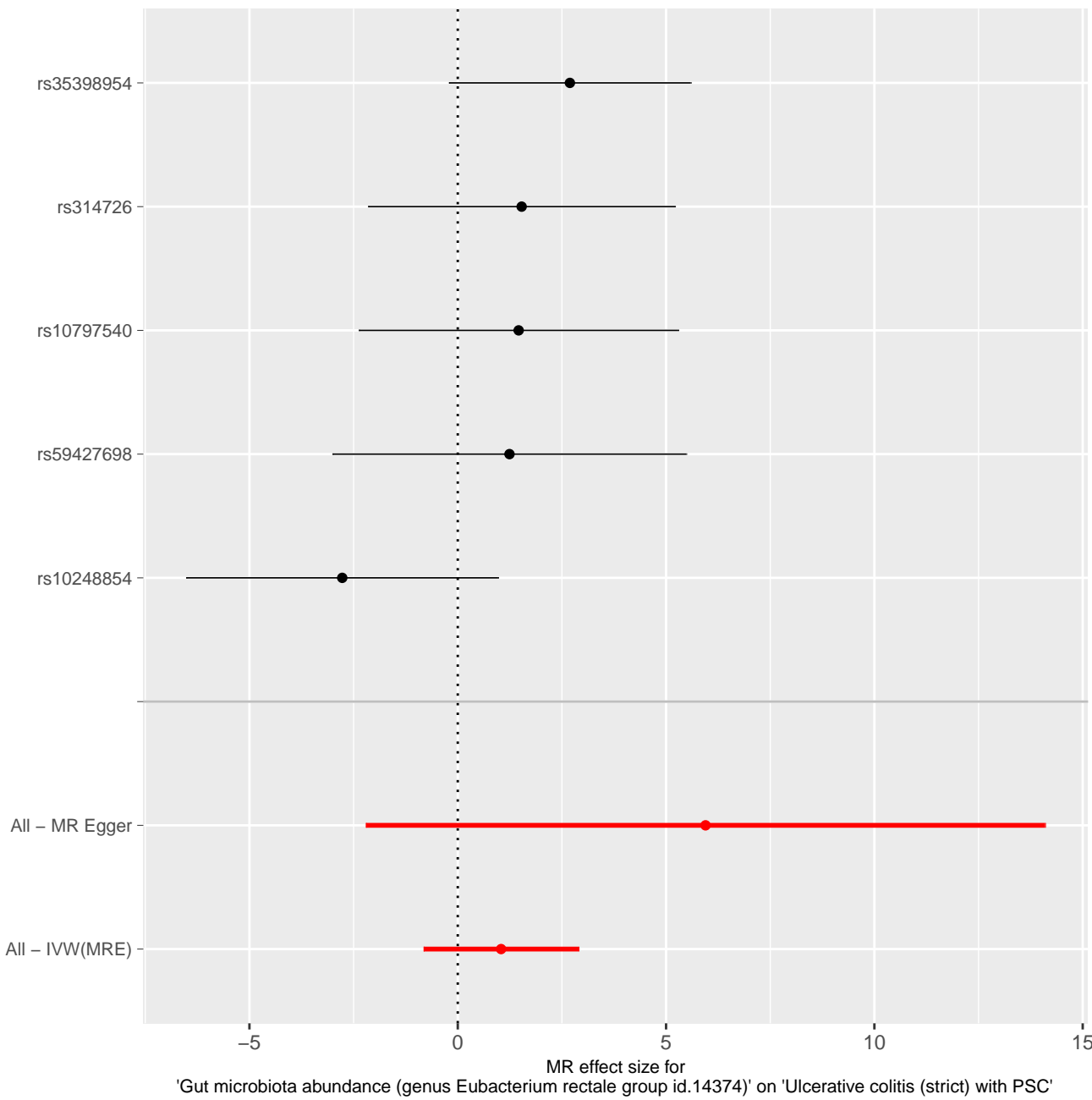
MR effect size for 'Gut microbiota abundance (genus Eubacterium eligens group id.14372)' on 'Ulcerative colitis (strict) with PSC'

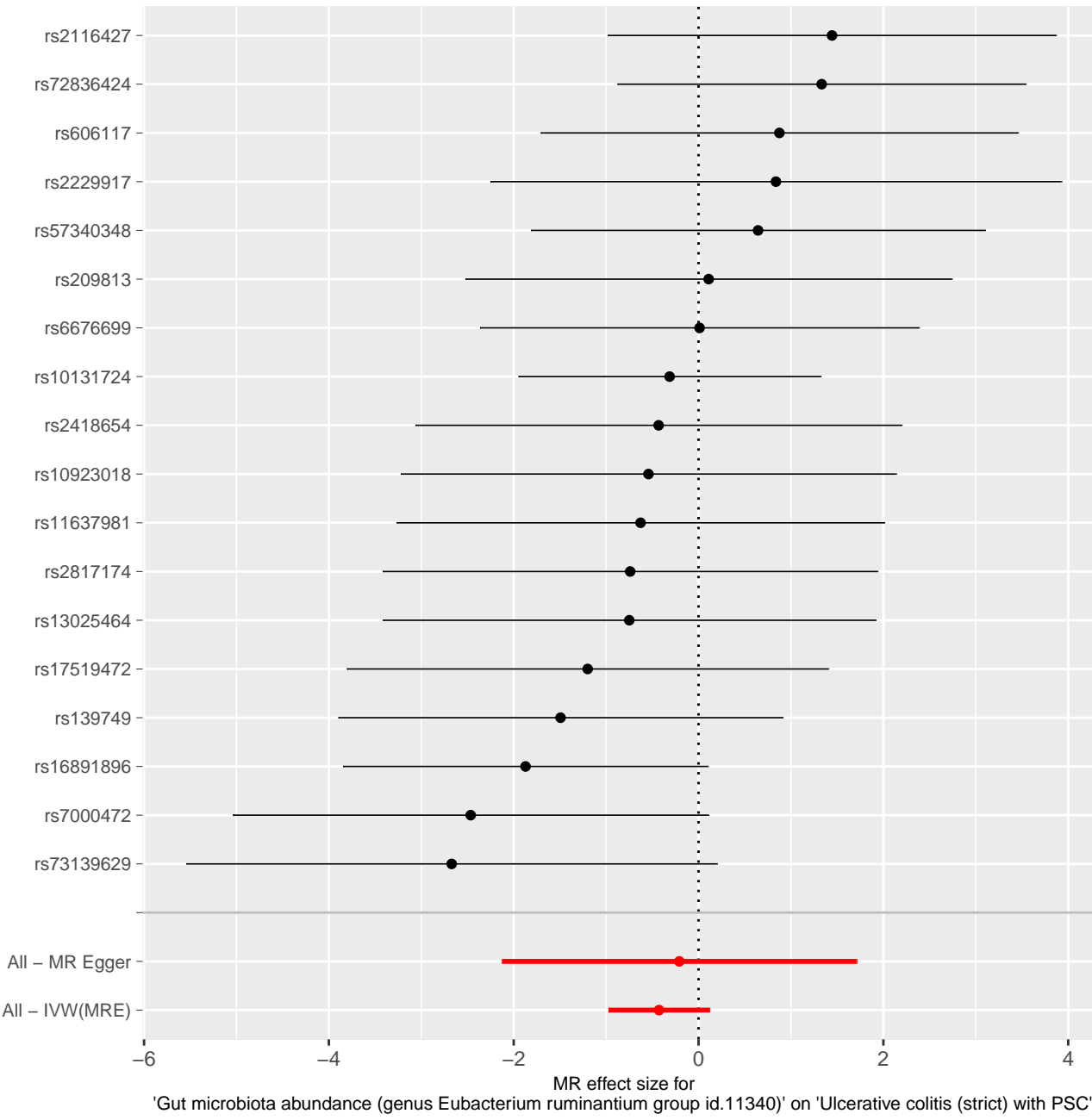


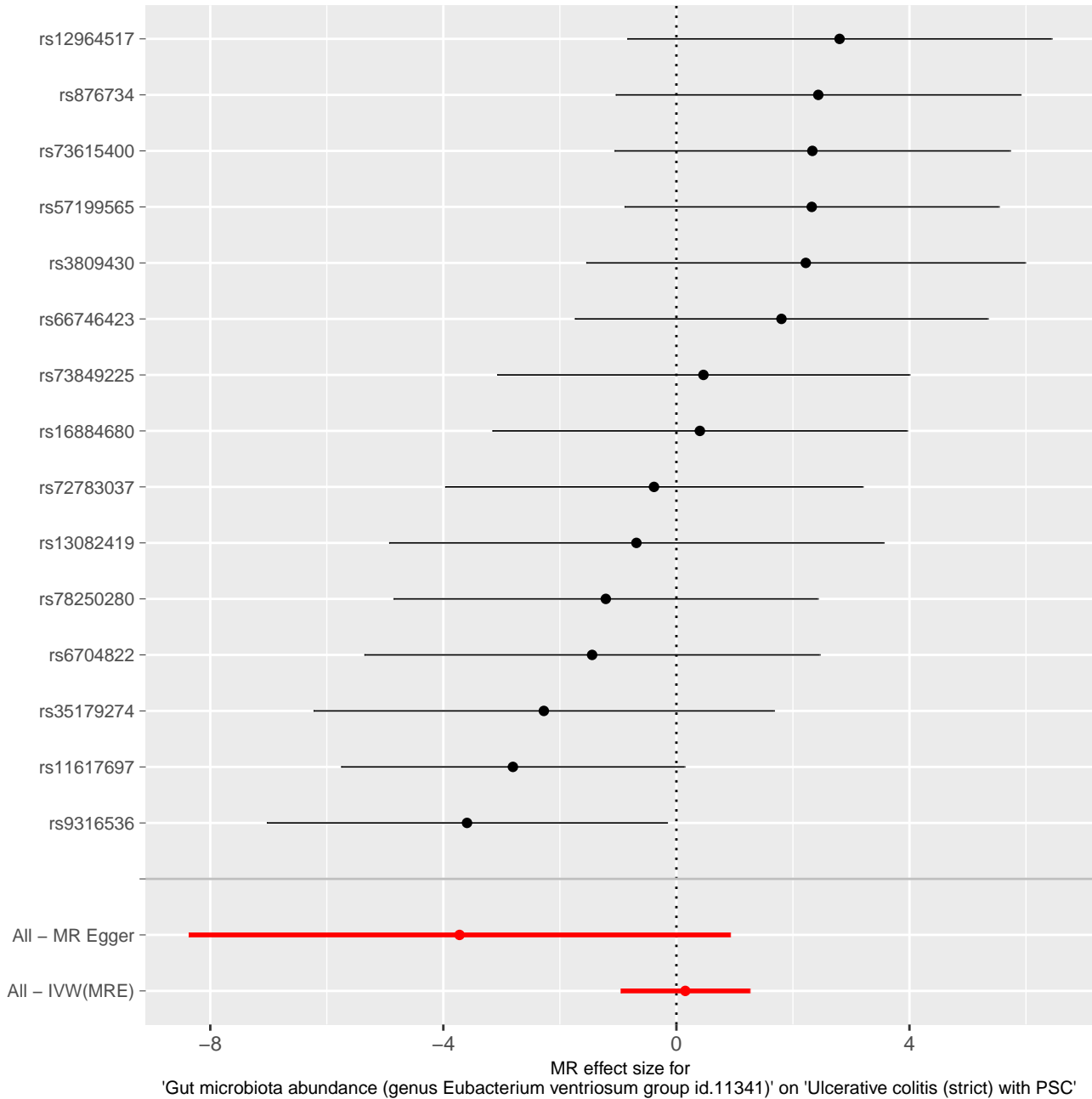


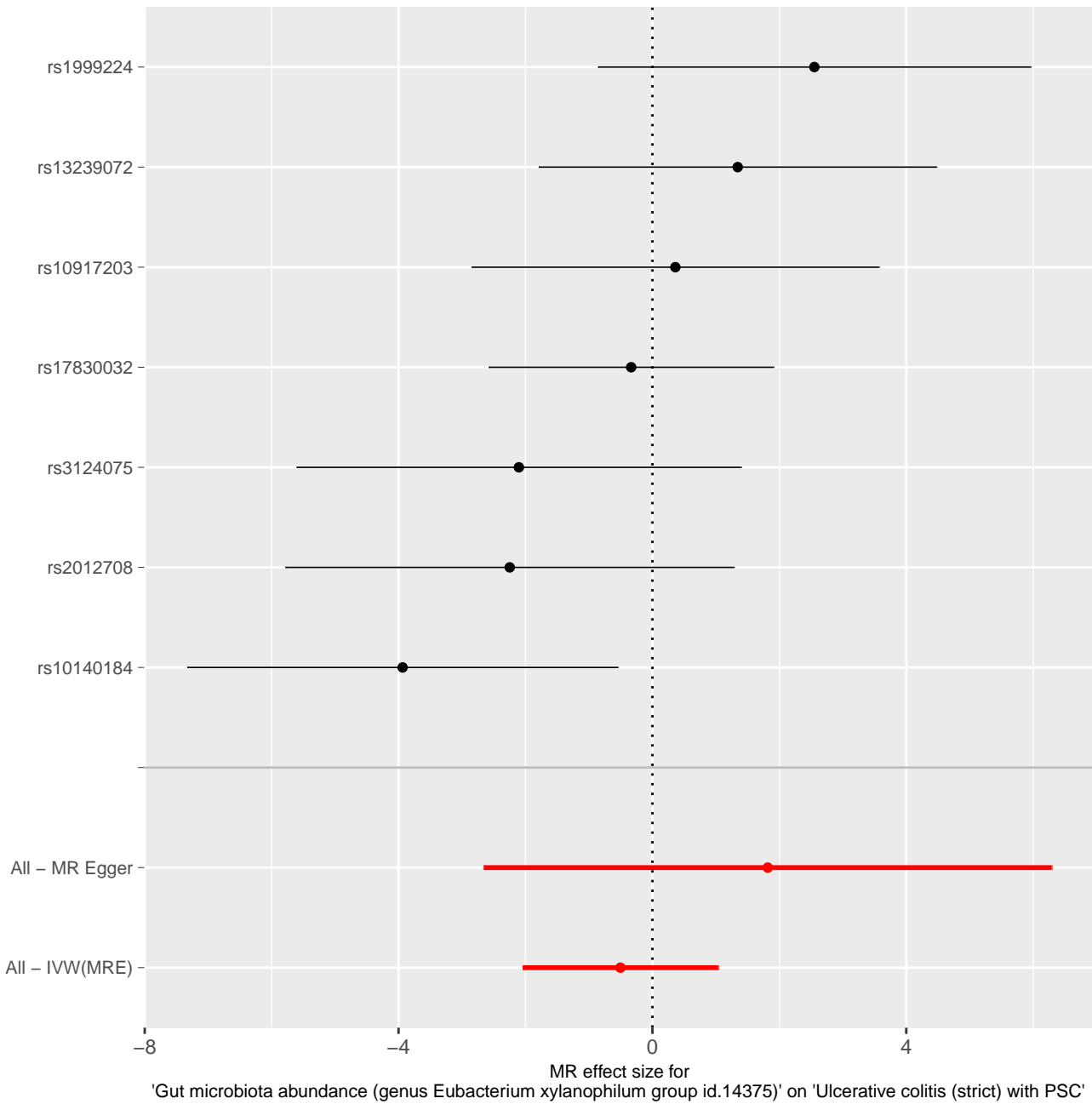


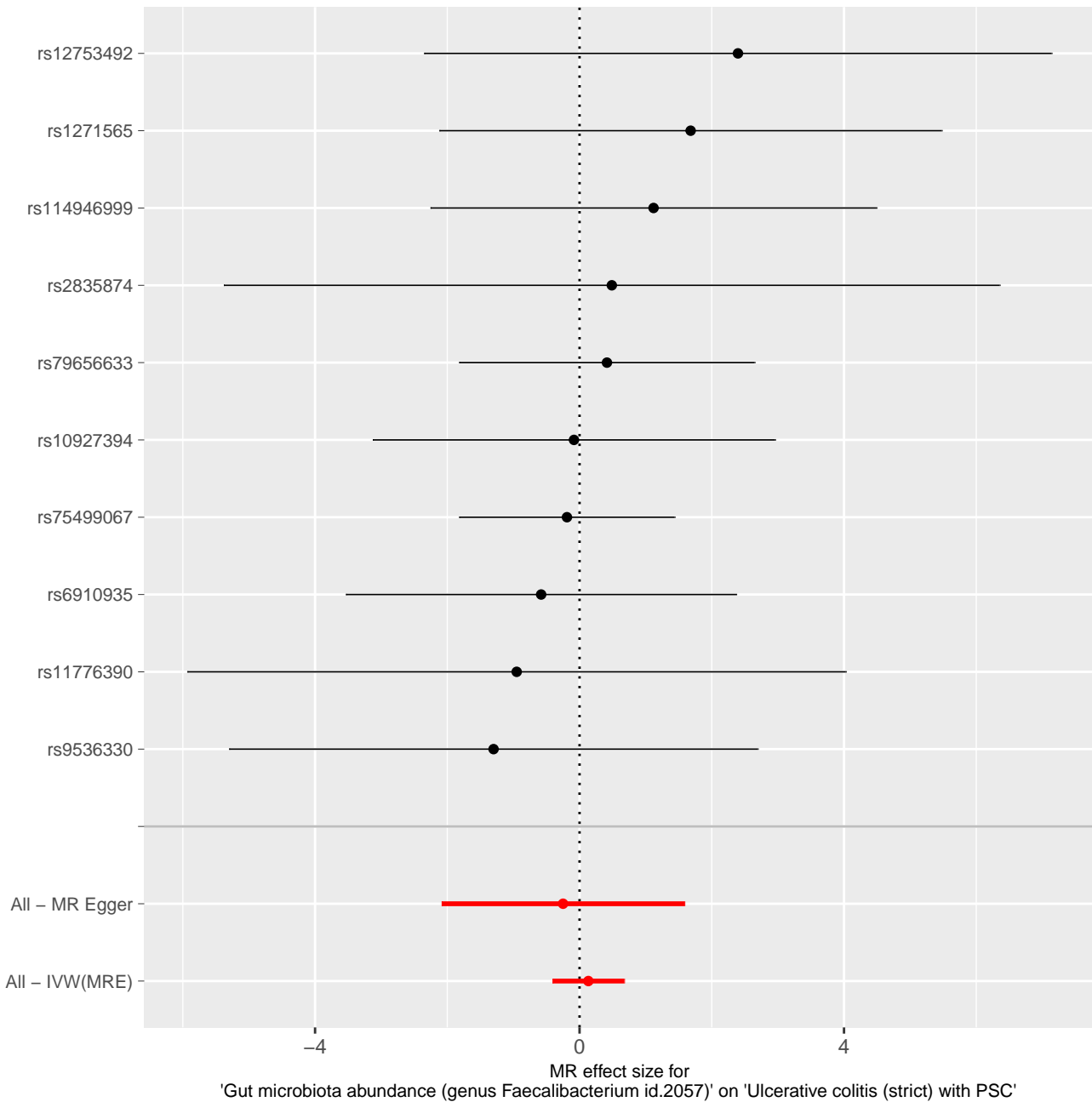


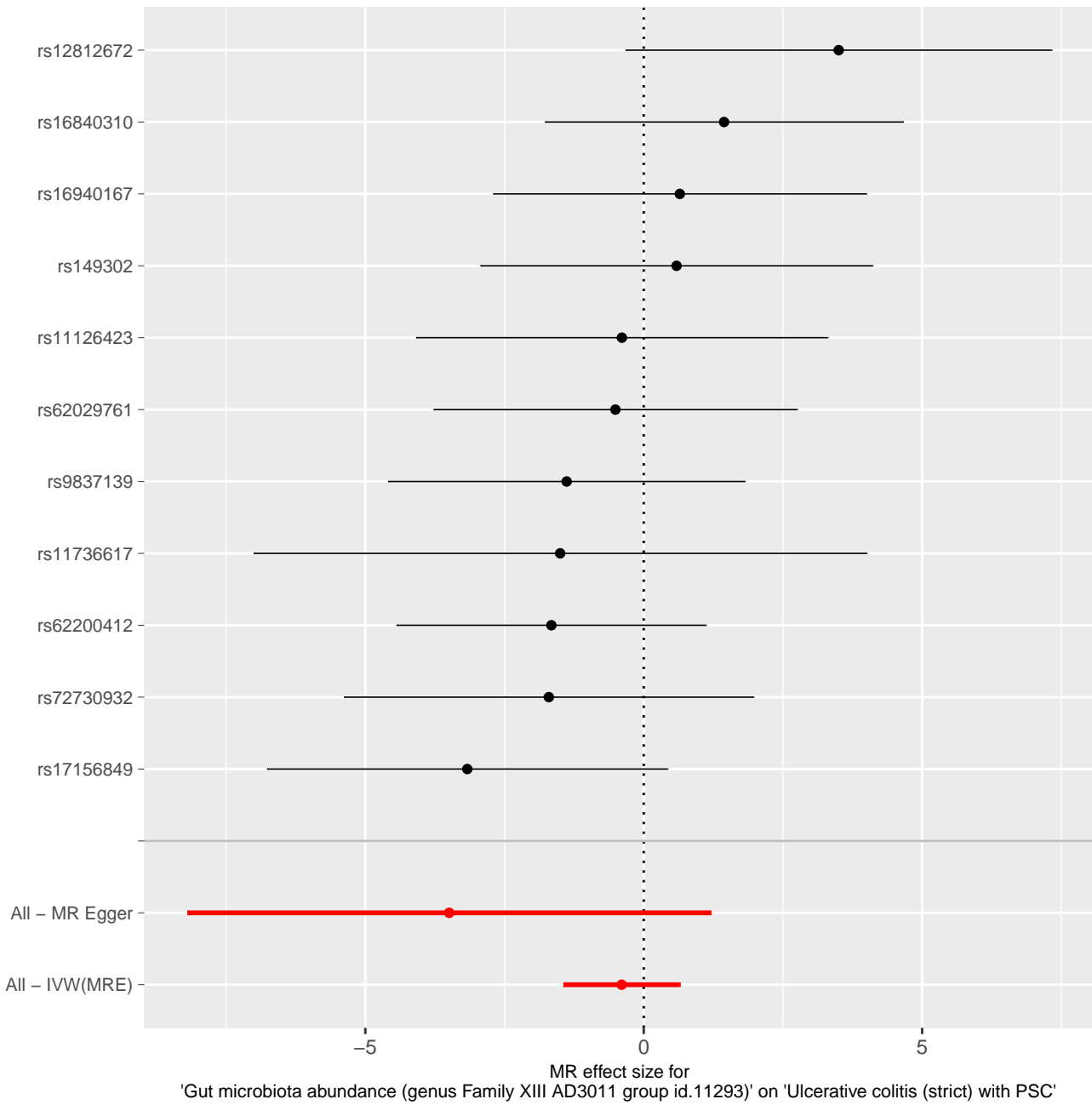


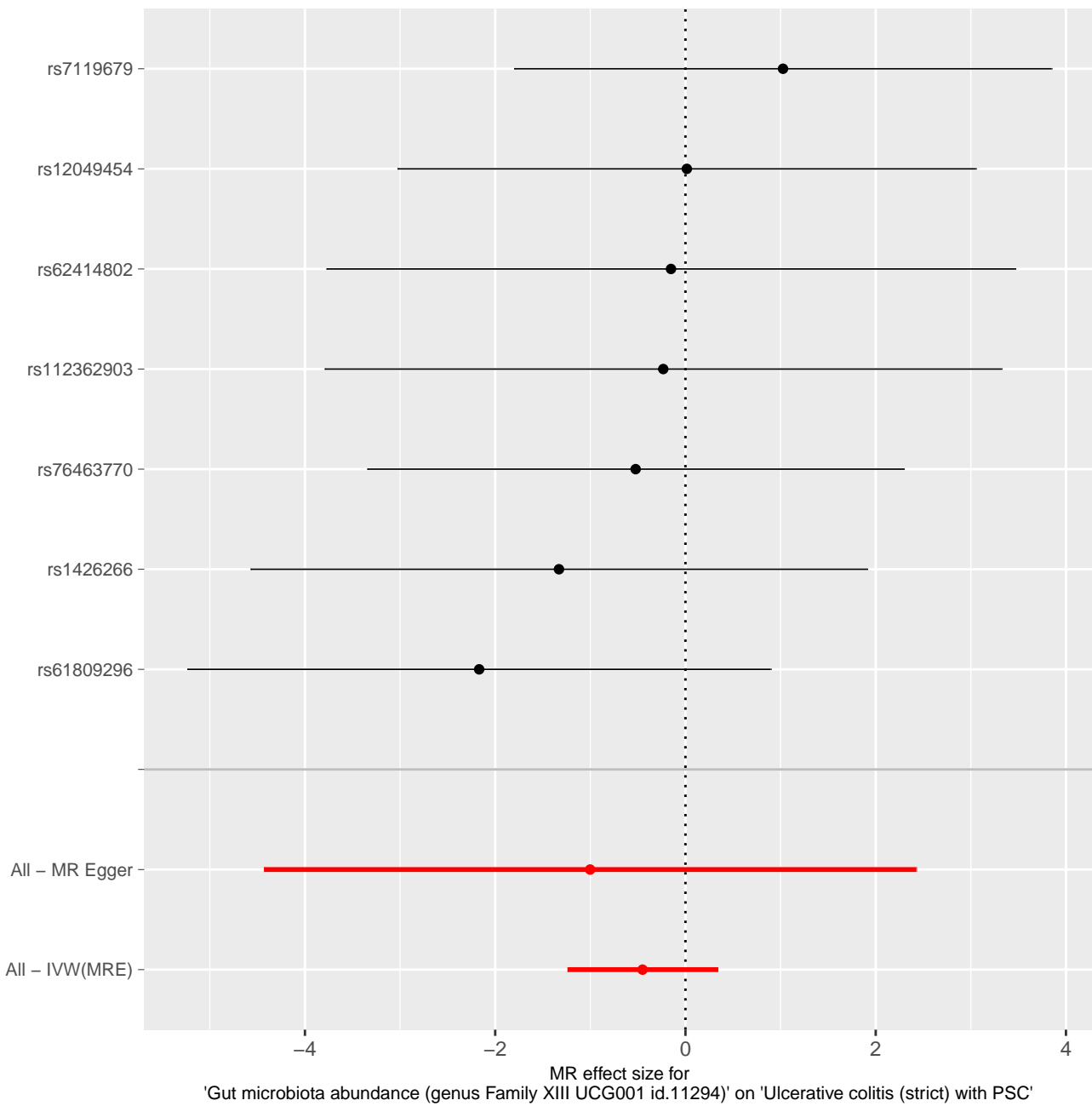


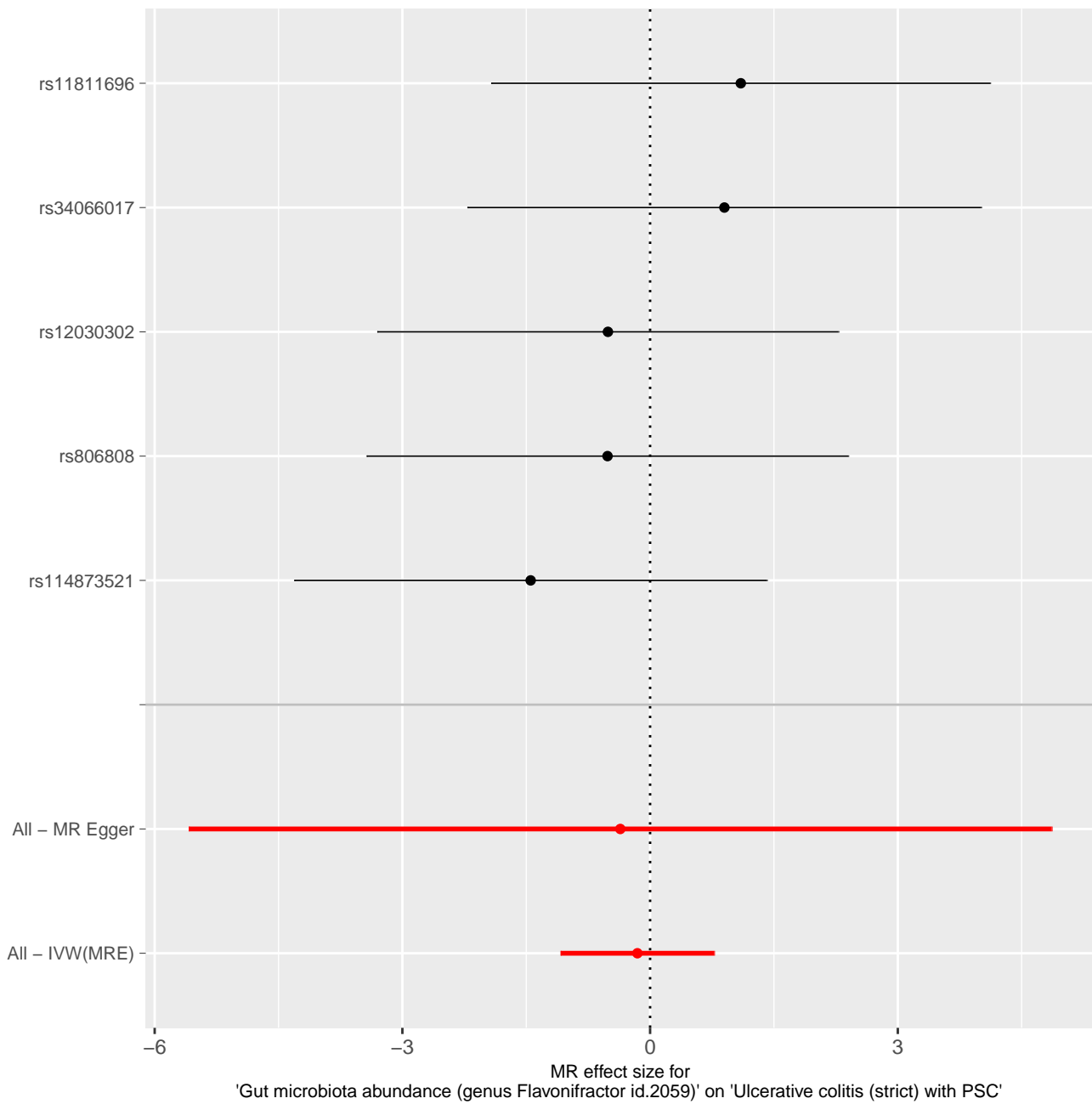


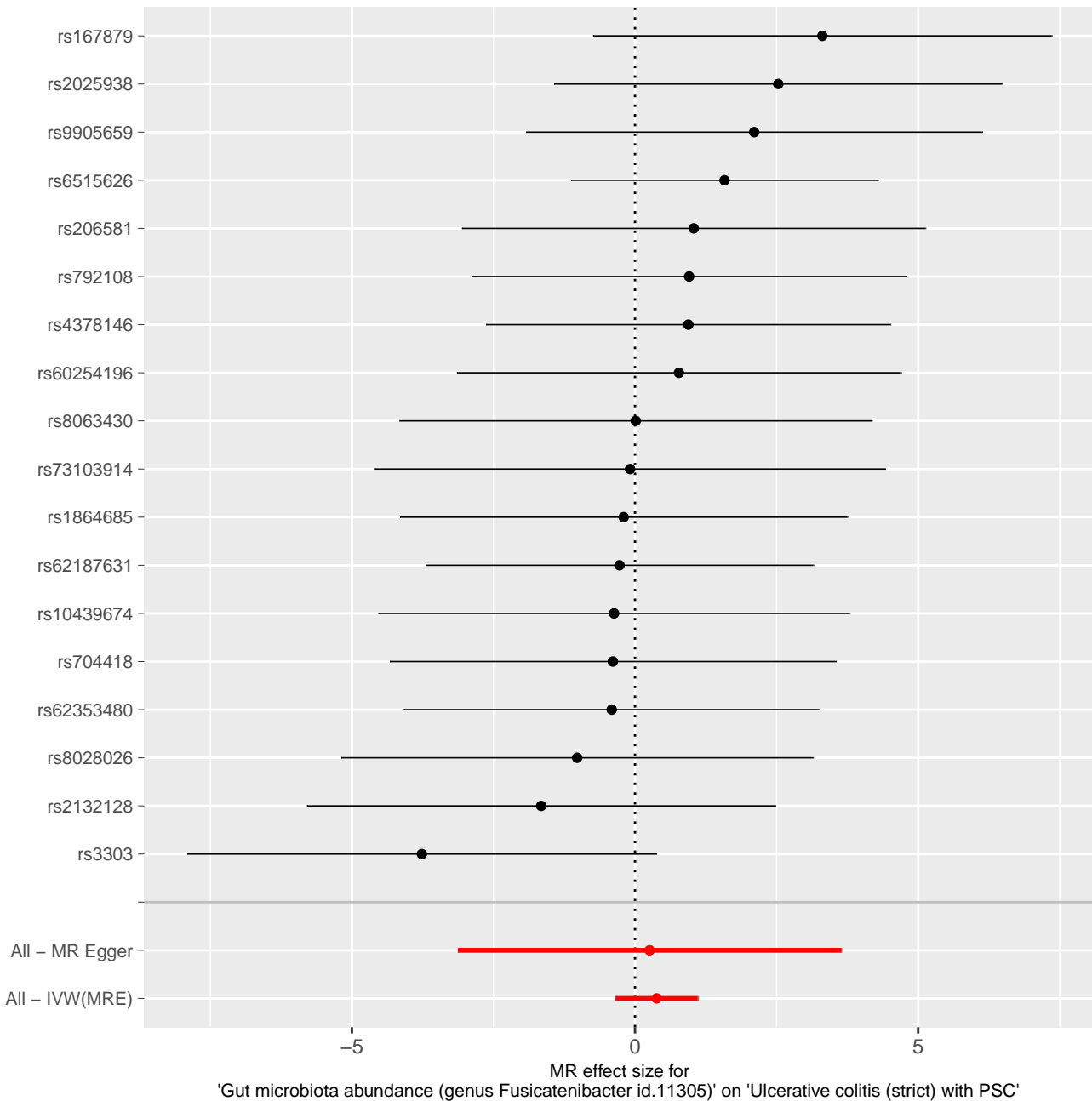


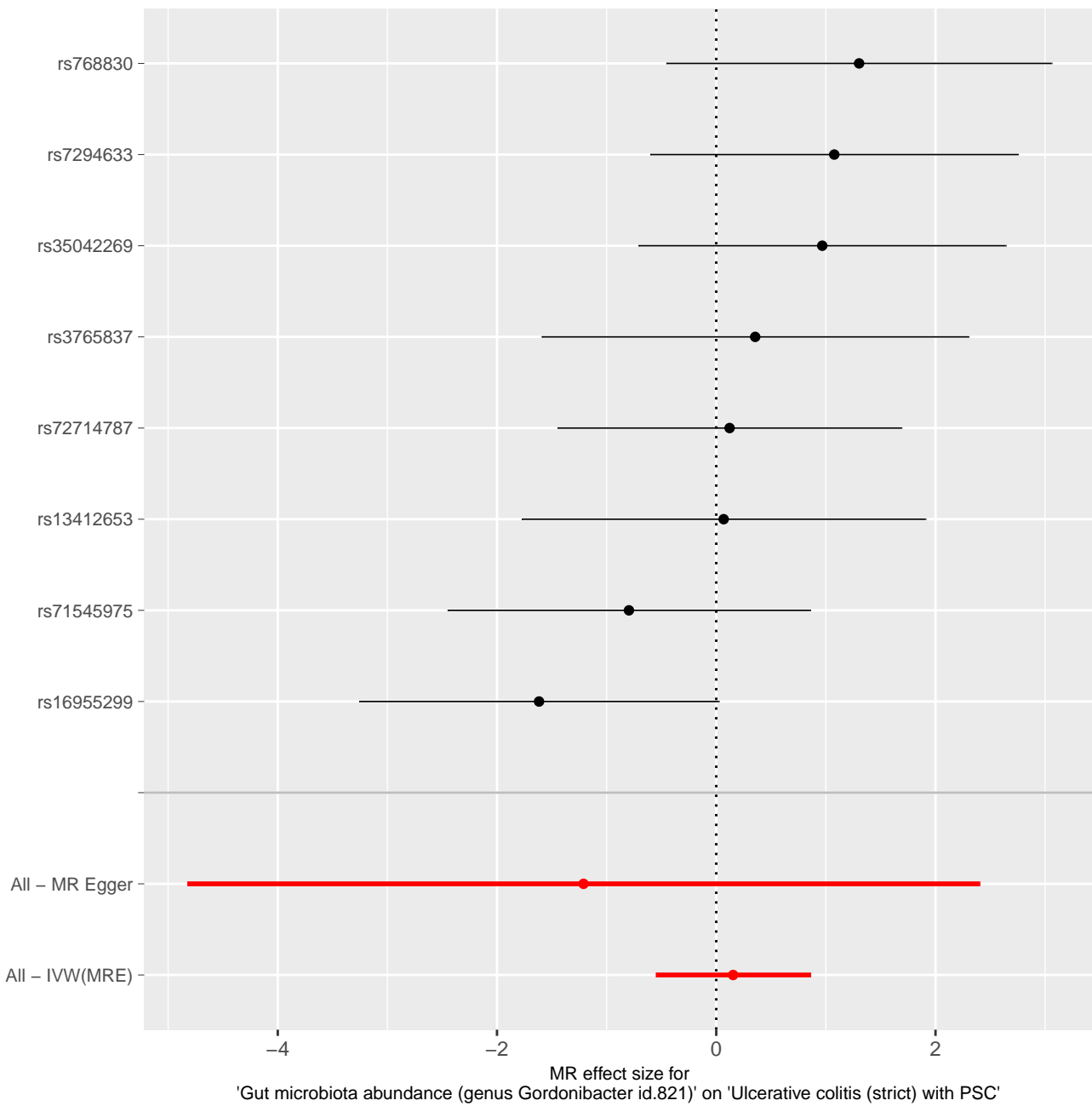


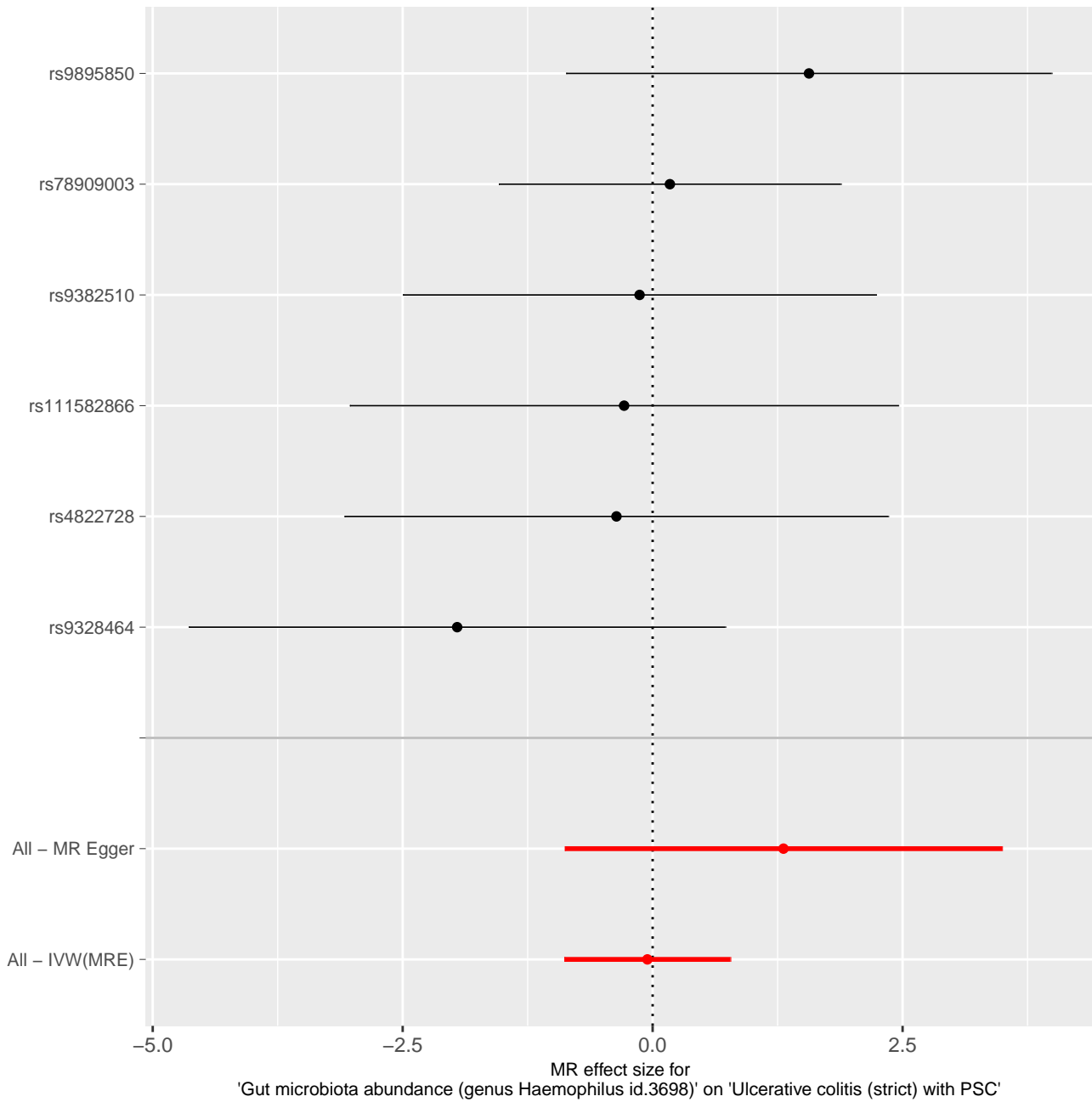




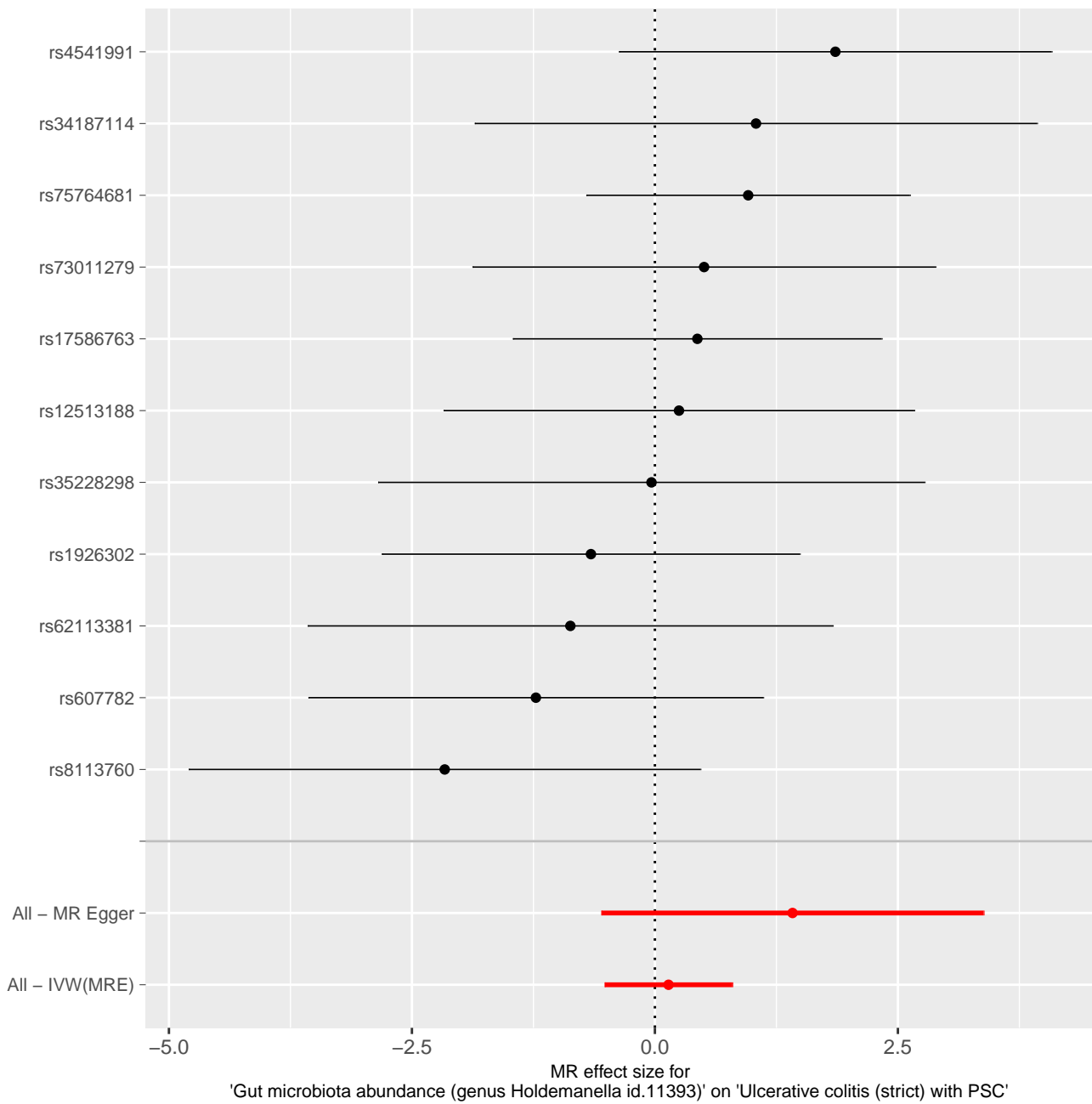




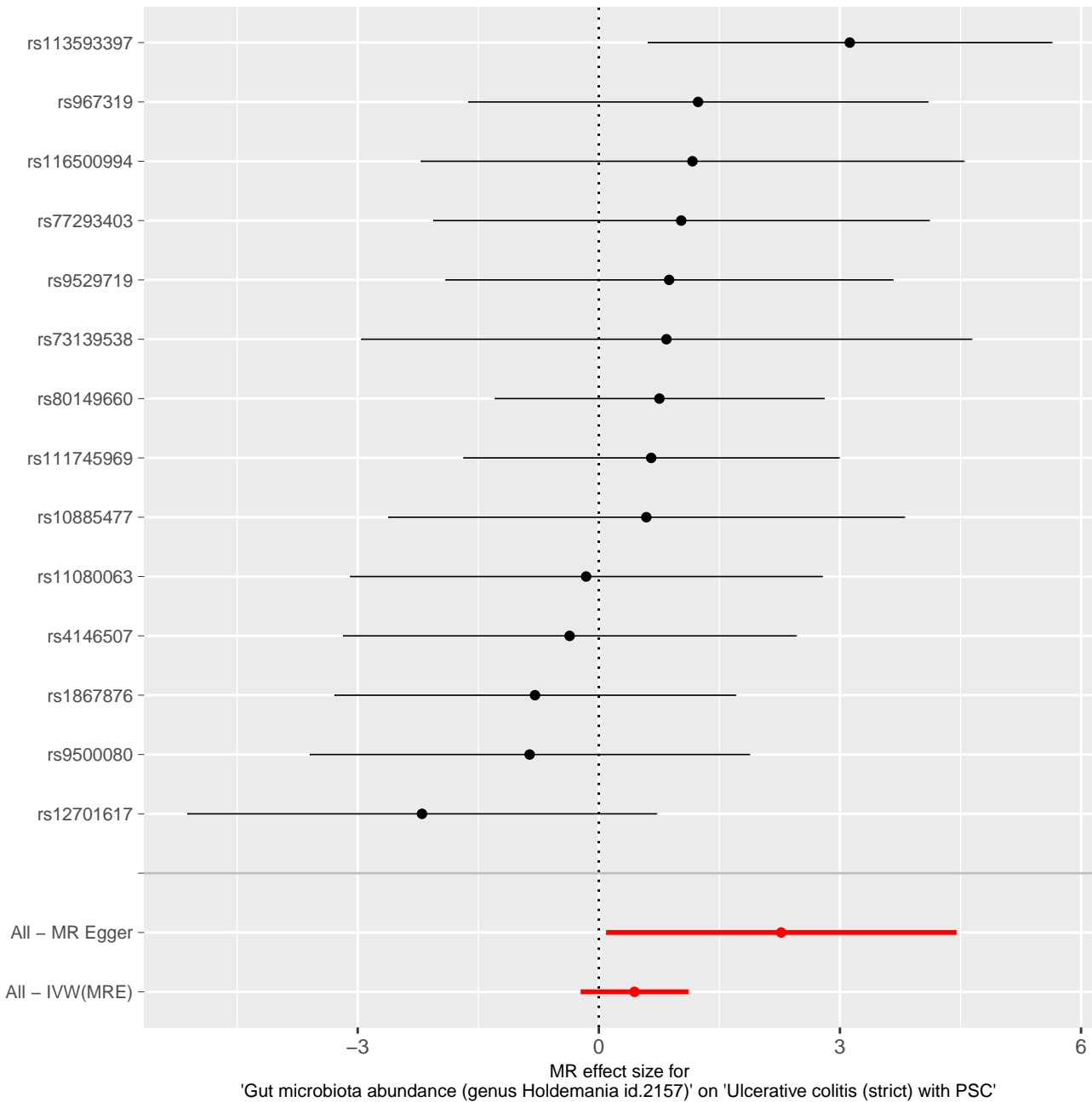


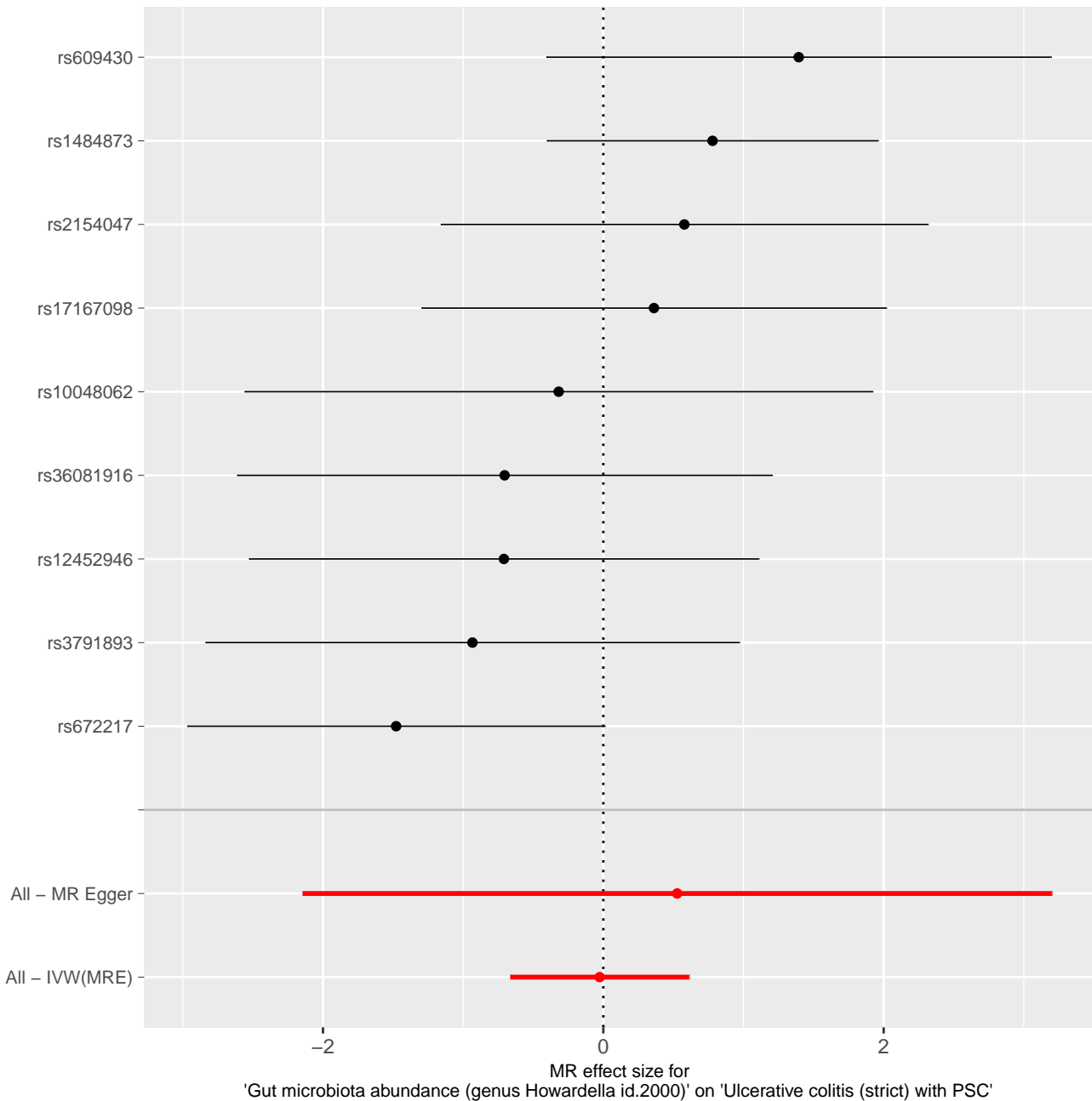


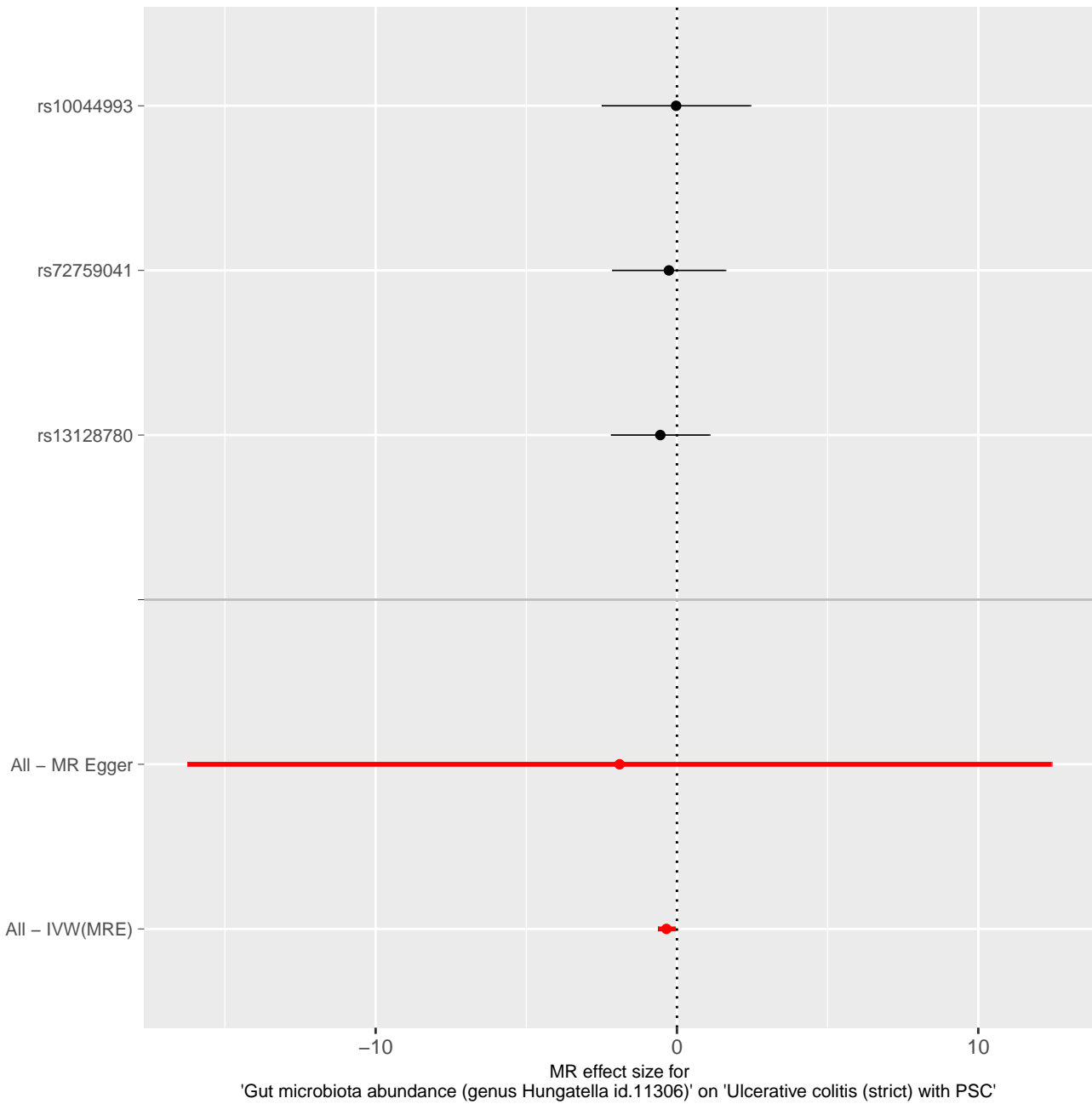
Batch 688 : Gut microbiota abundance (genus Holdemanella id.11393) on Ulcerative colitis (strict) with PSC



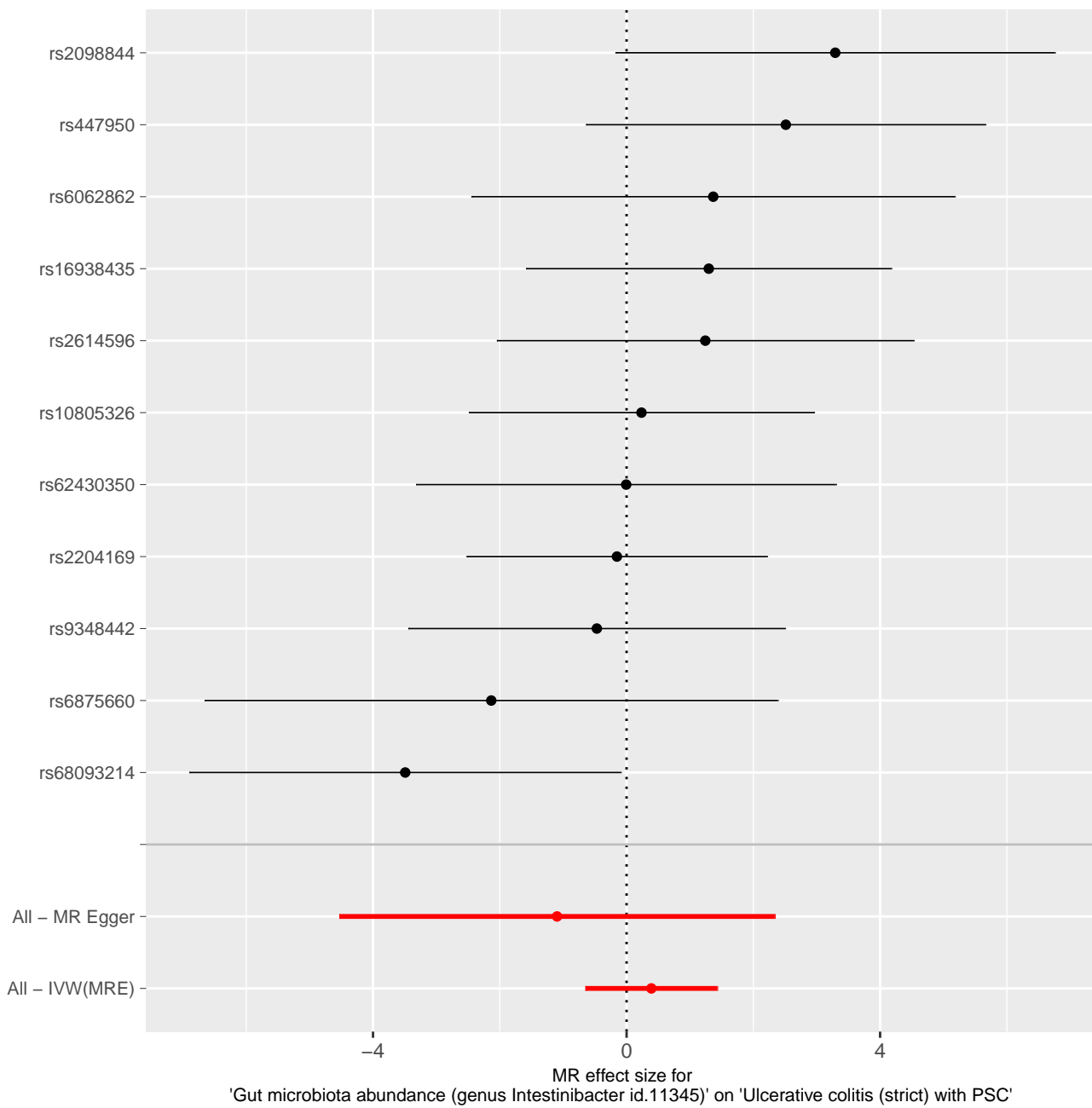
Batch 689 : Gut microbiota abundance (genus Holdemania id.2157) on Ulcerative colitis (strict) with PSC



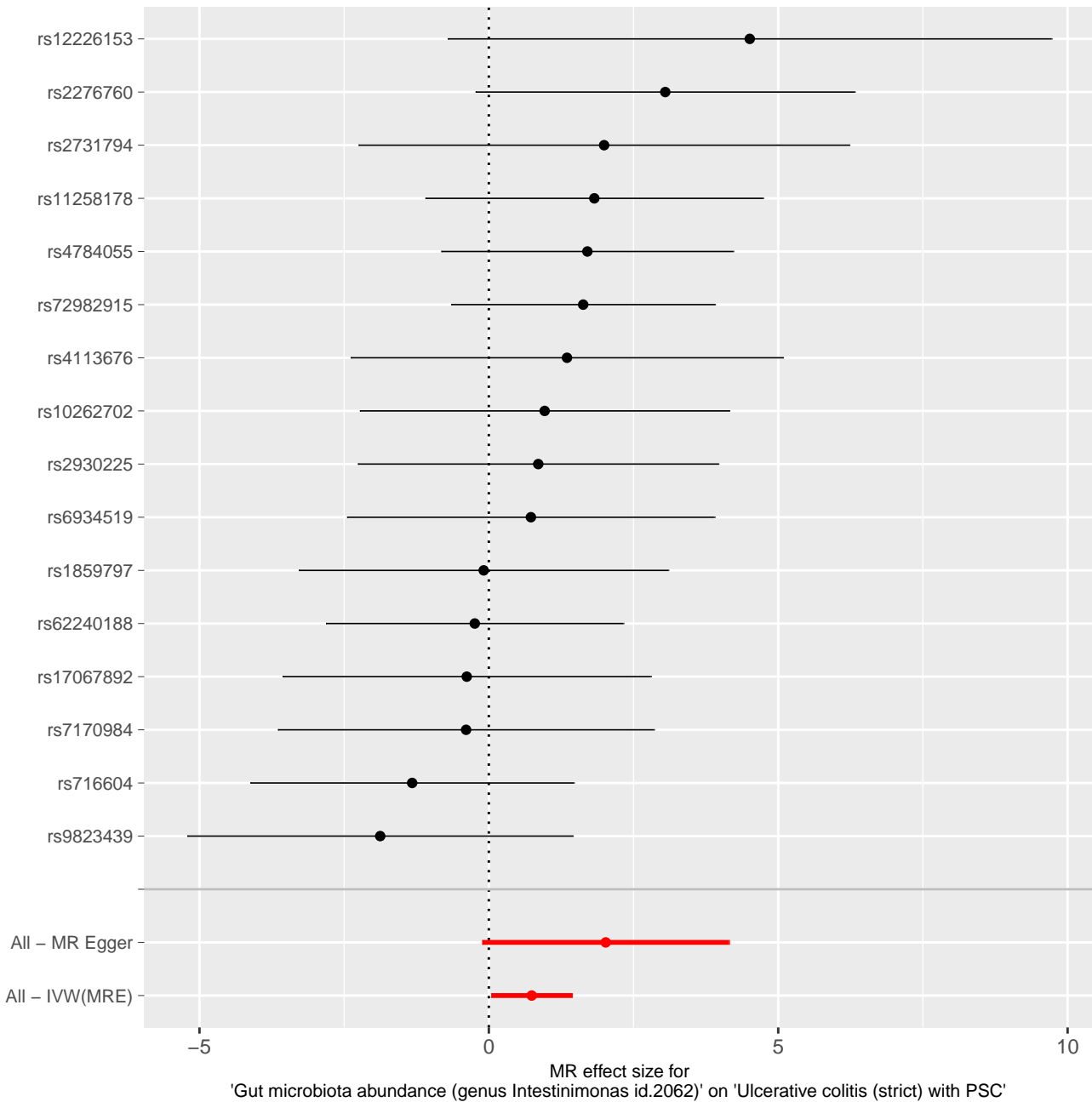


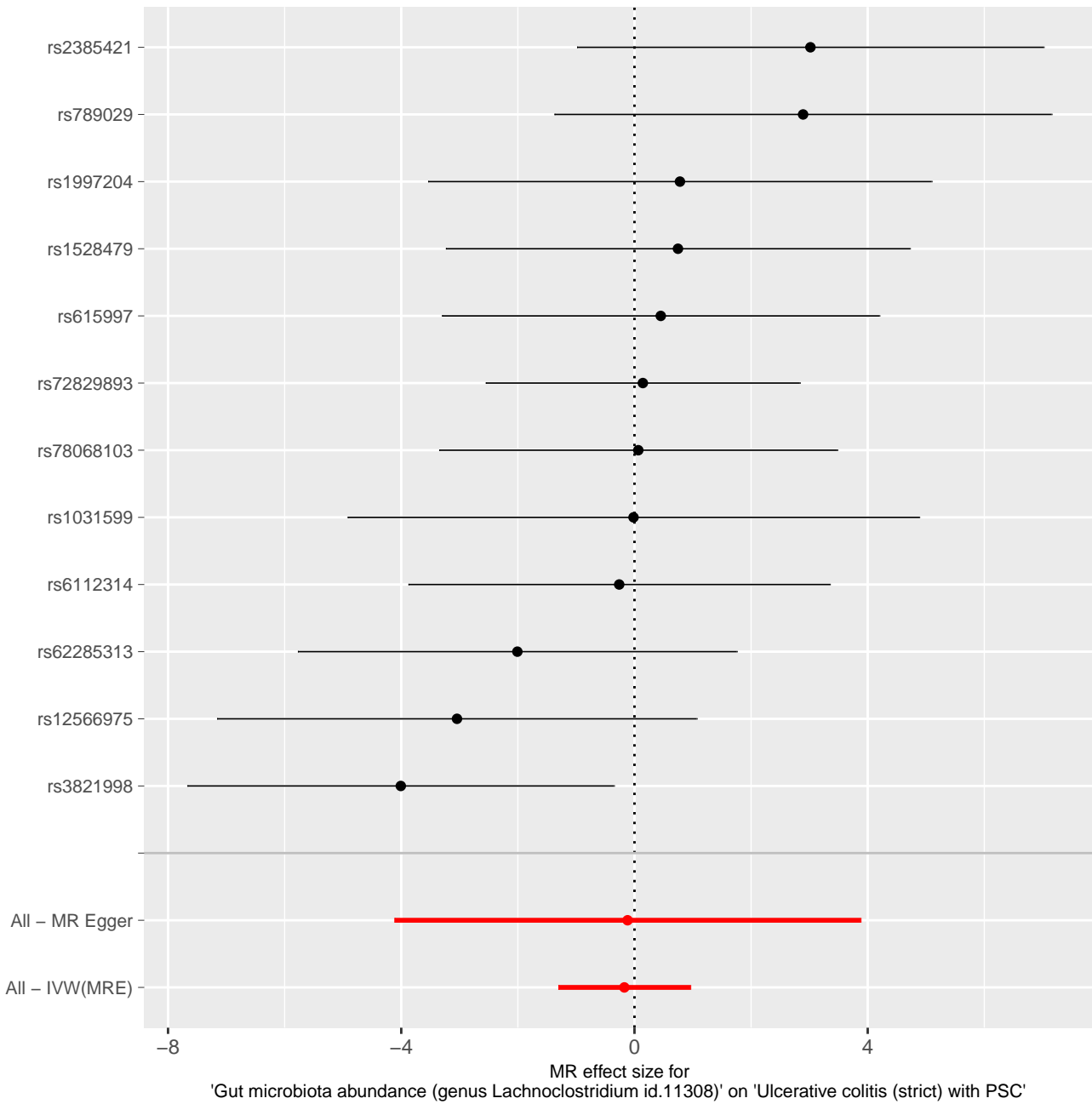


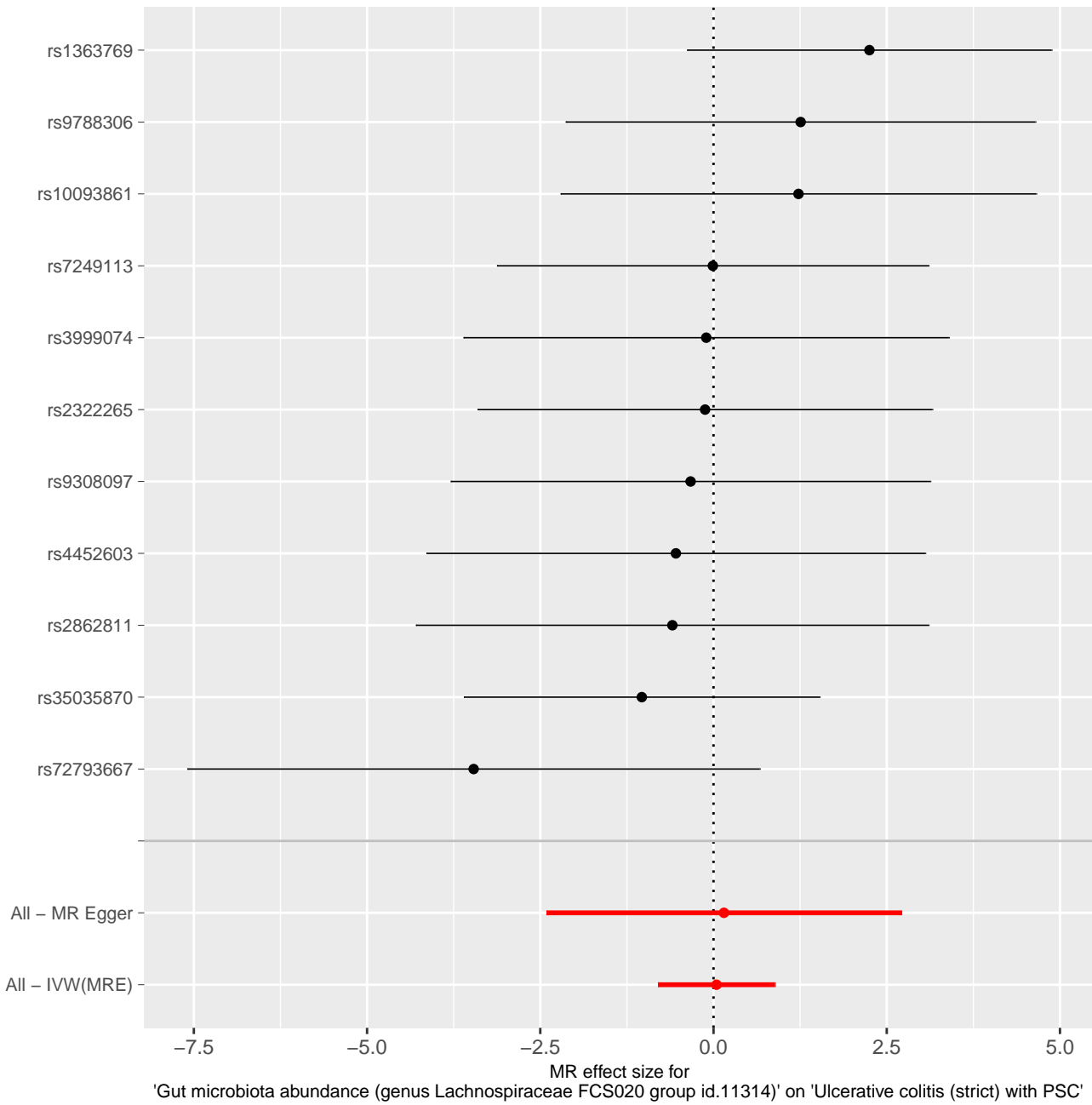
Batch 692 : Gut microbiota abundance (genus Intestinibacter id.11345) on Ulcerative colitis (strict) with PSC

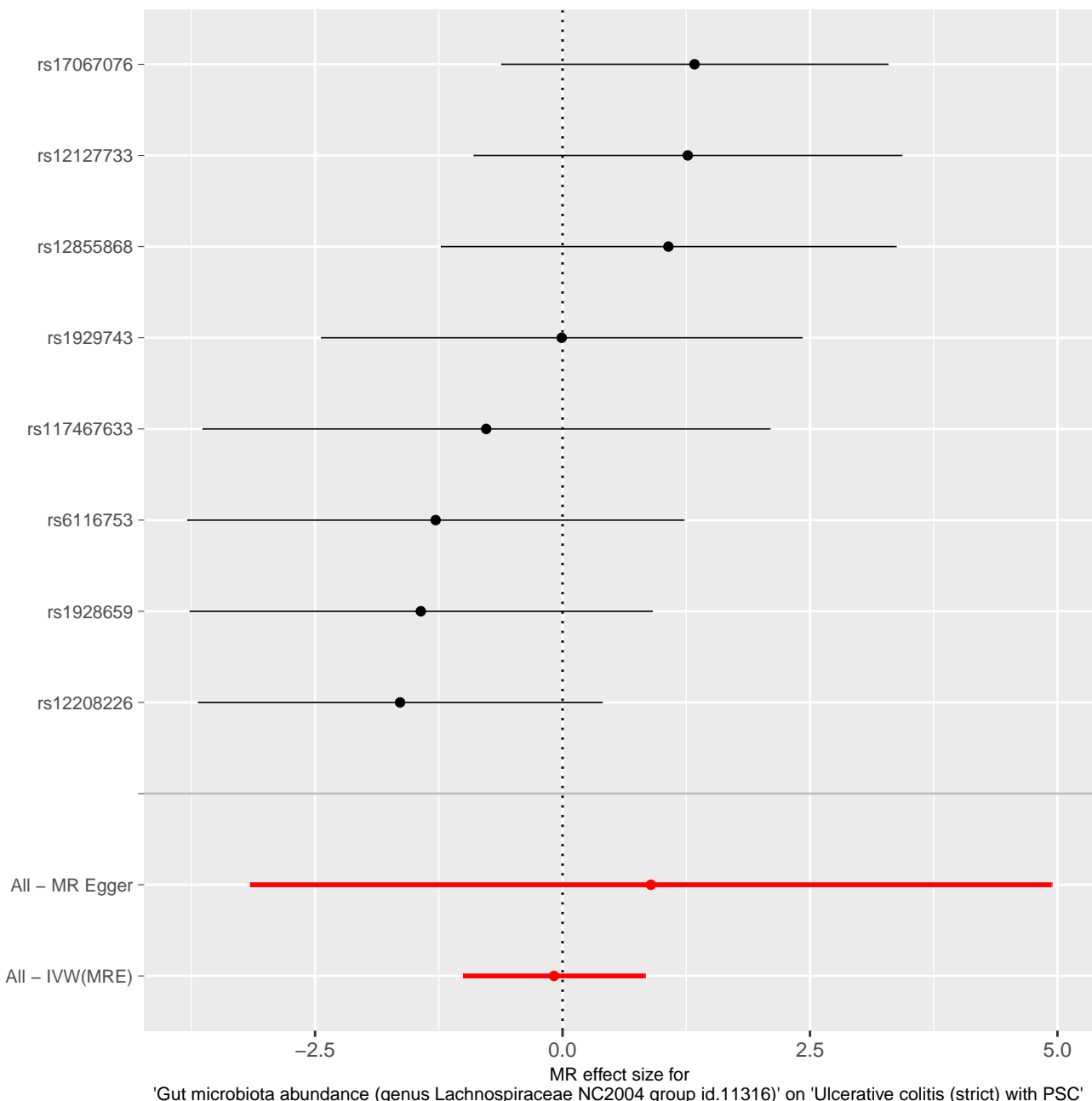


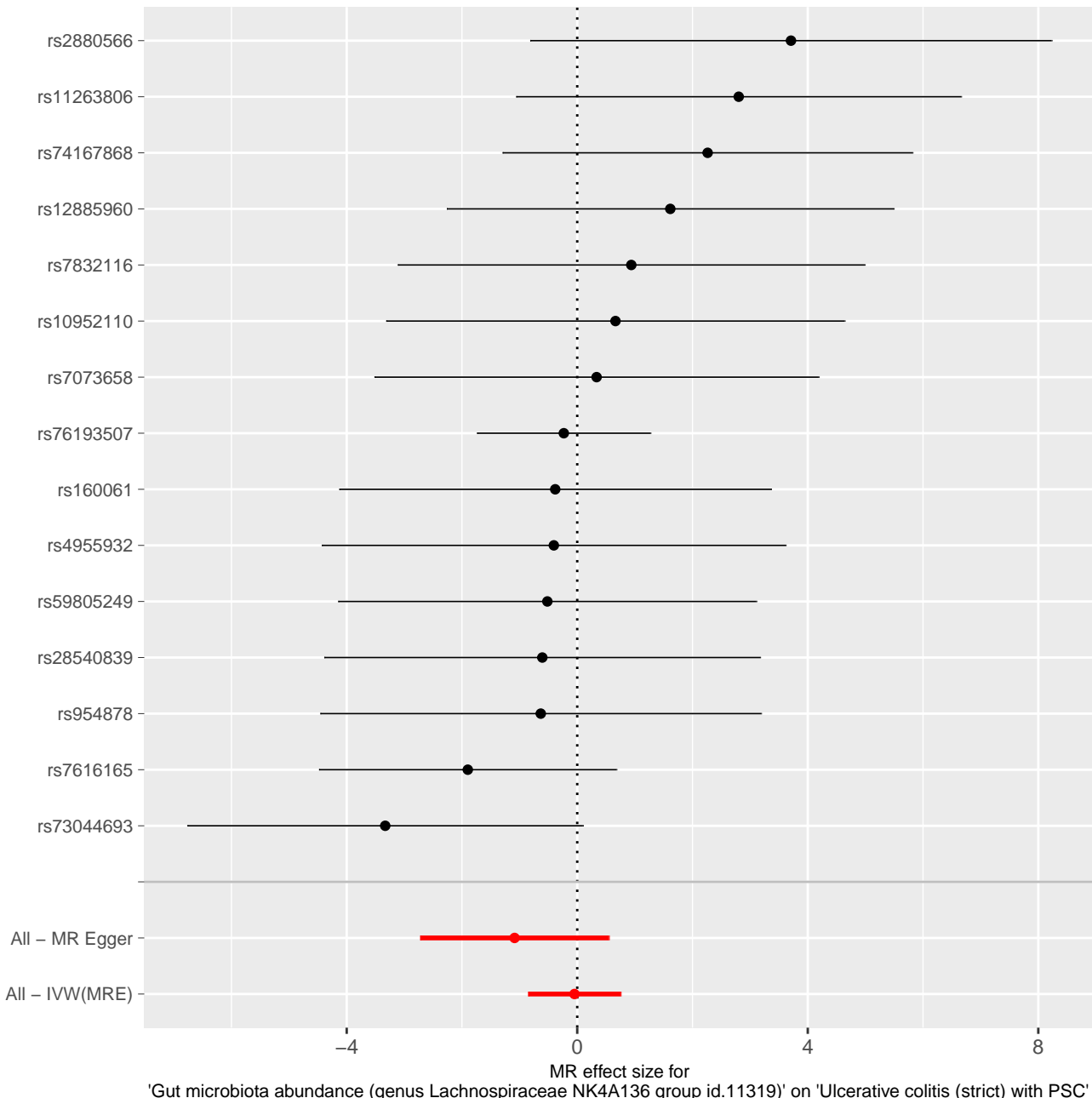
Batch 693 : Gut microbiota abundance (genus Intestinimonas id.2062) on Ulcerative colitis (strict) with PSC

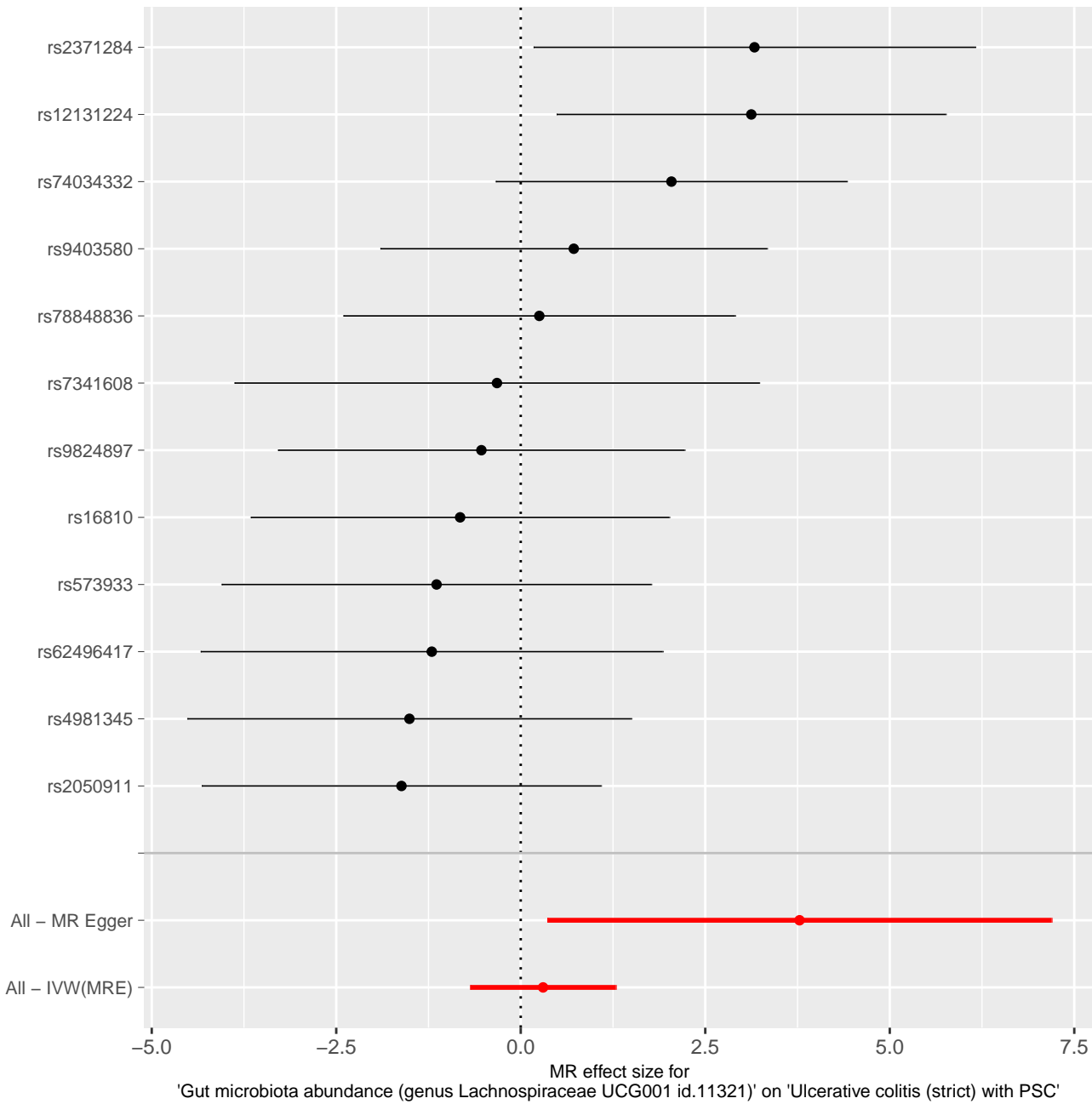


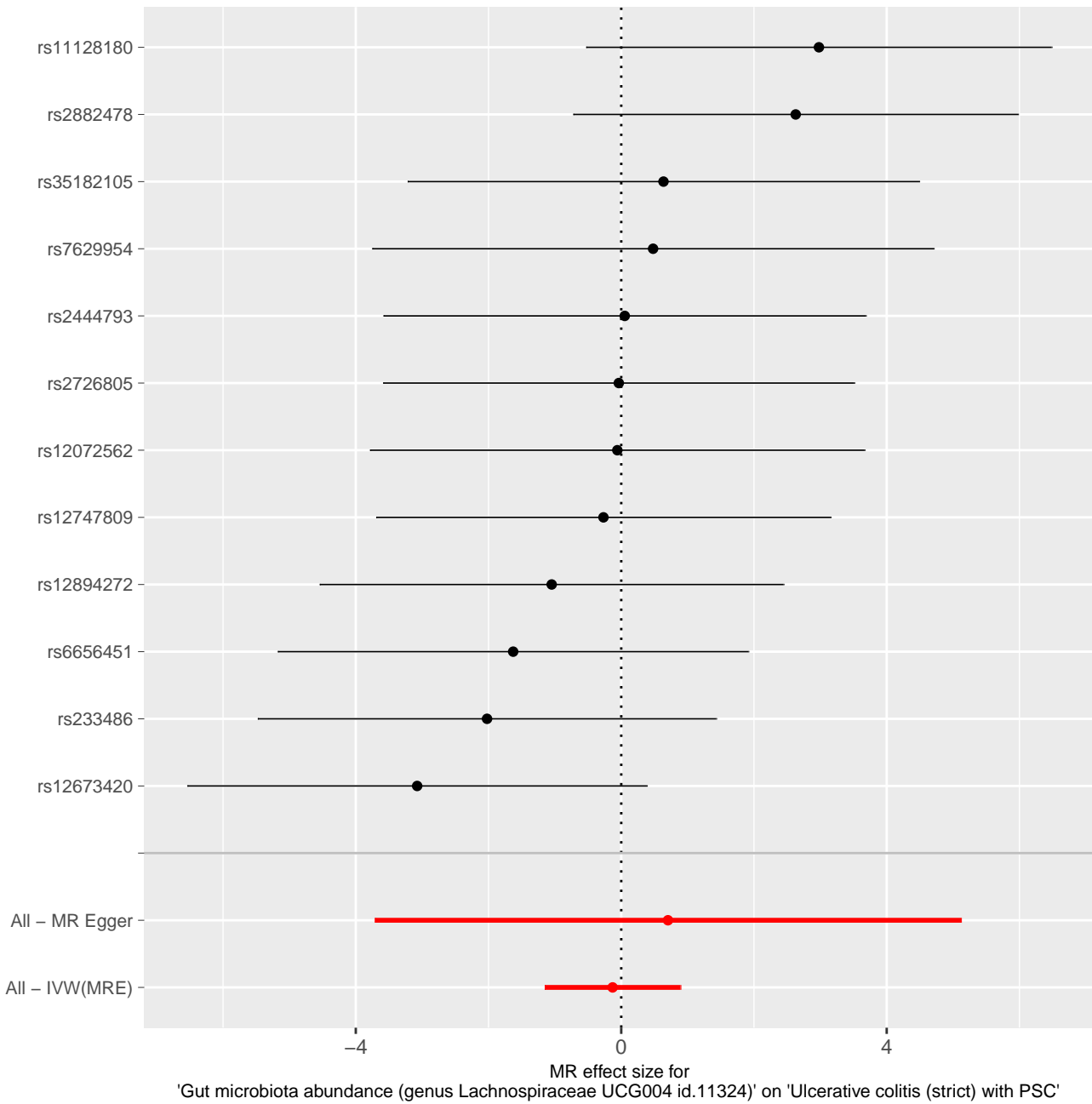


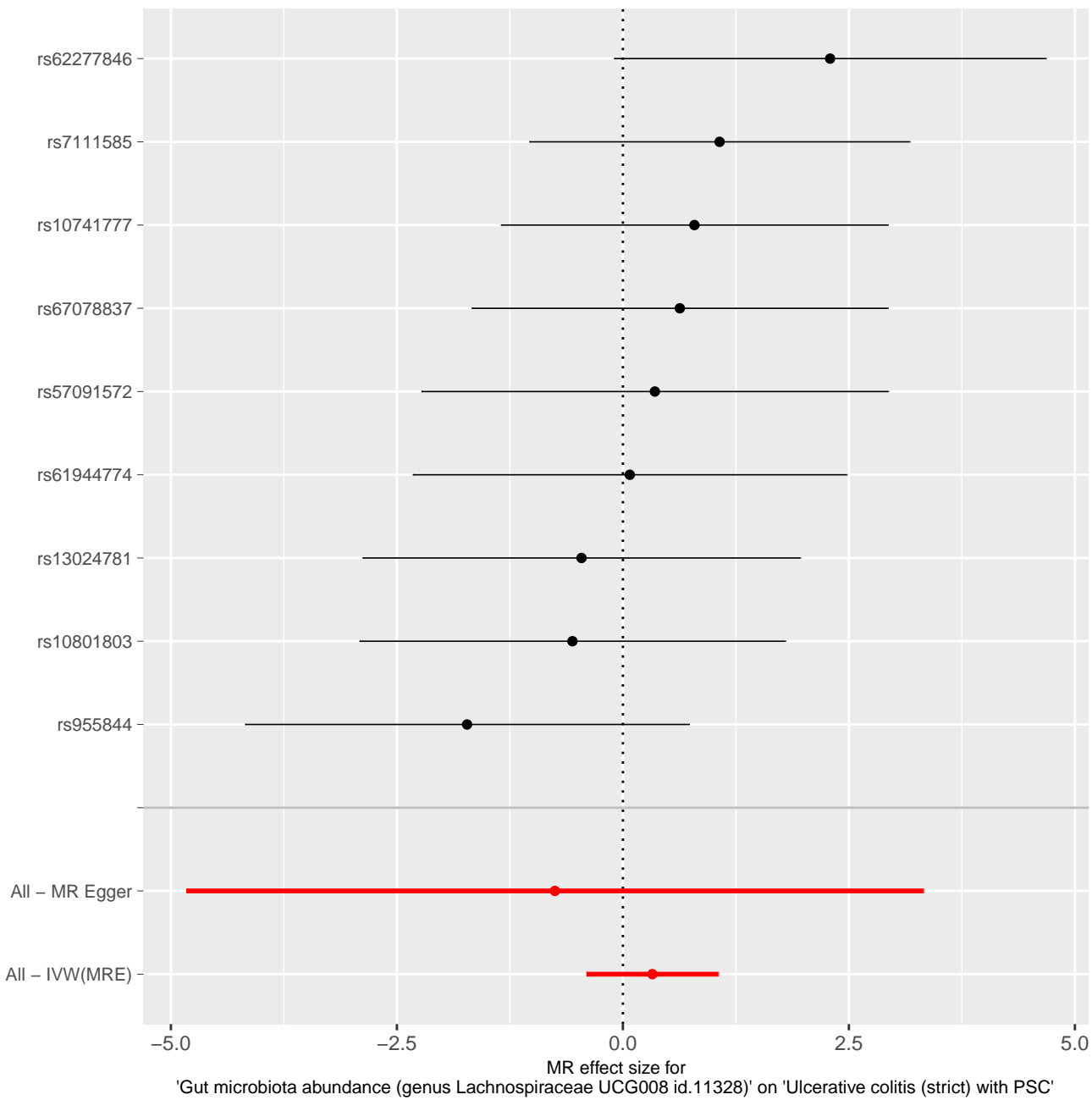


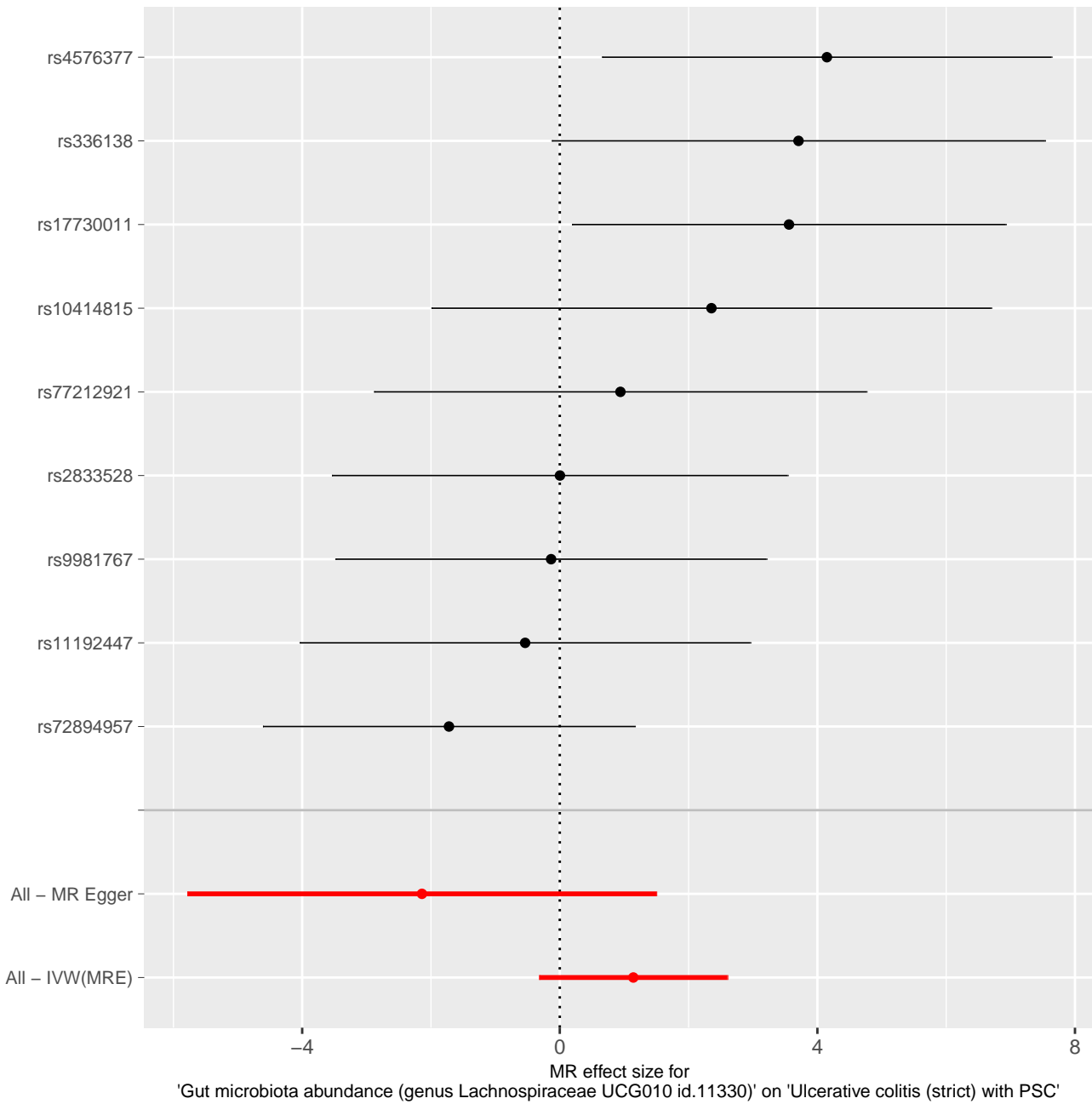






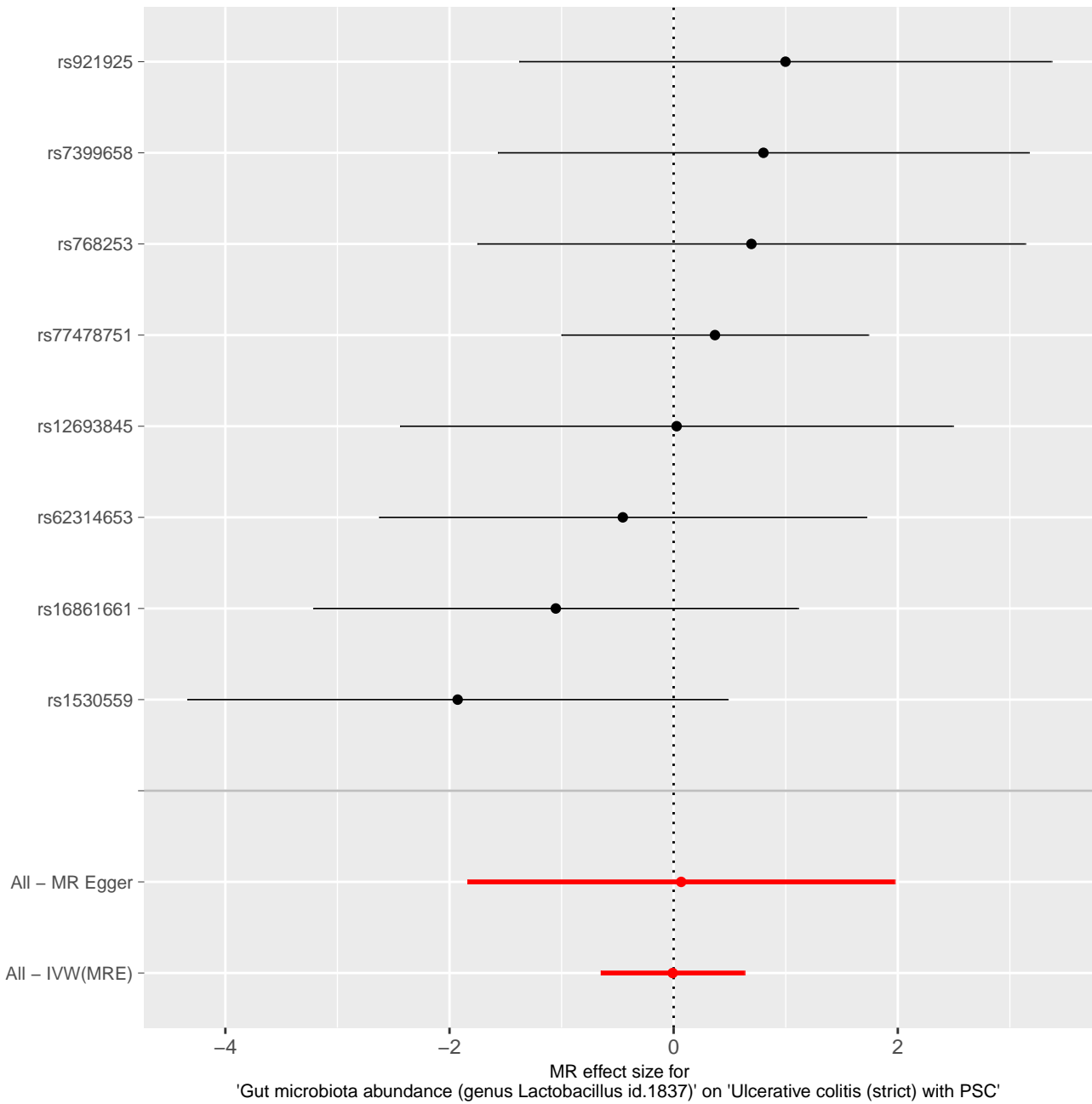


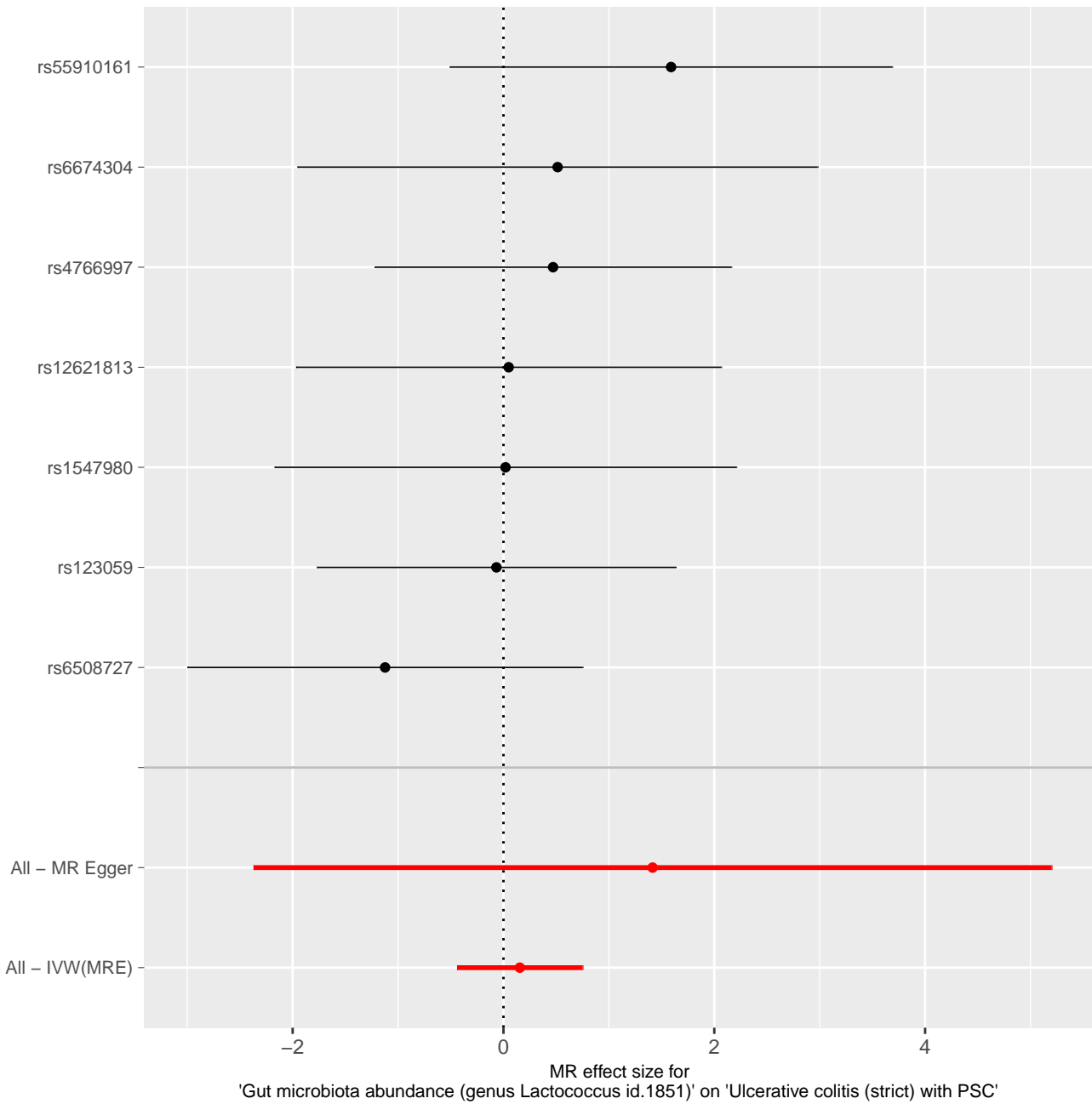


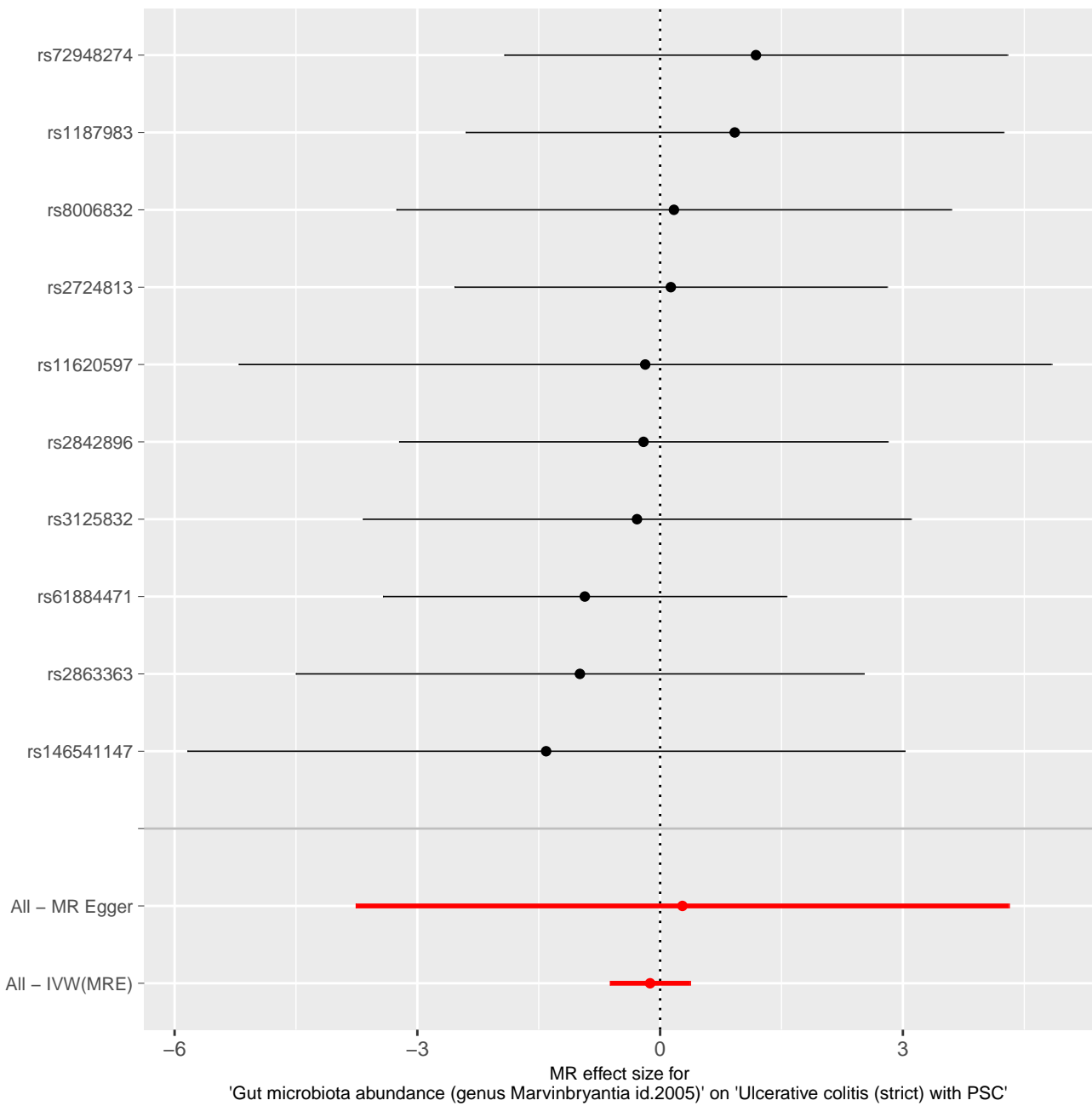


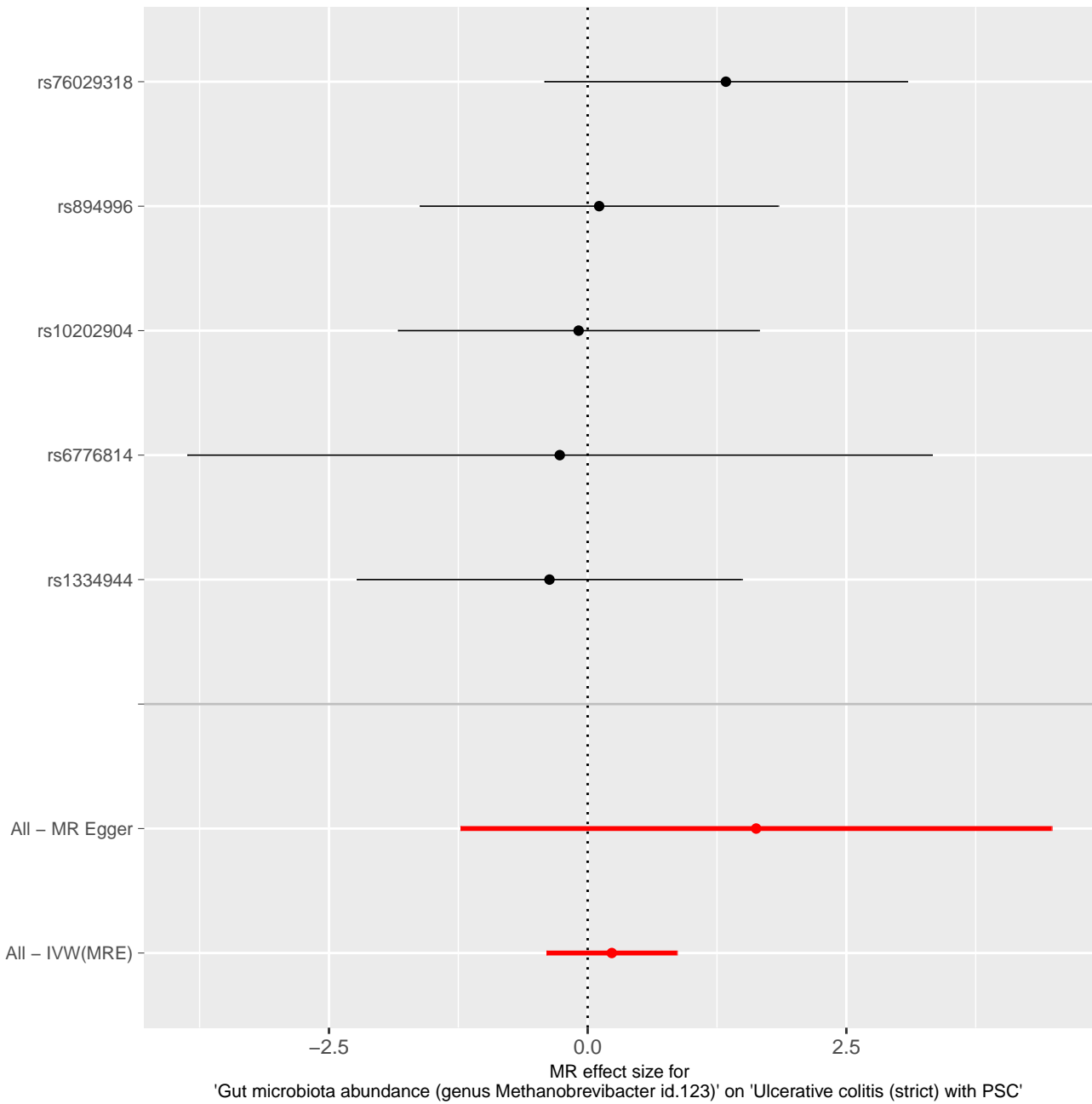
'Gut microbiota abundance (genus Lachnospiraceae UCG010 id.11330)' on 'Ulcerative colitis (strict) with PSC'

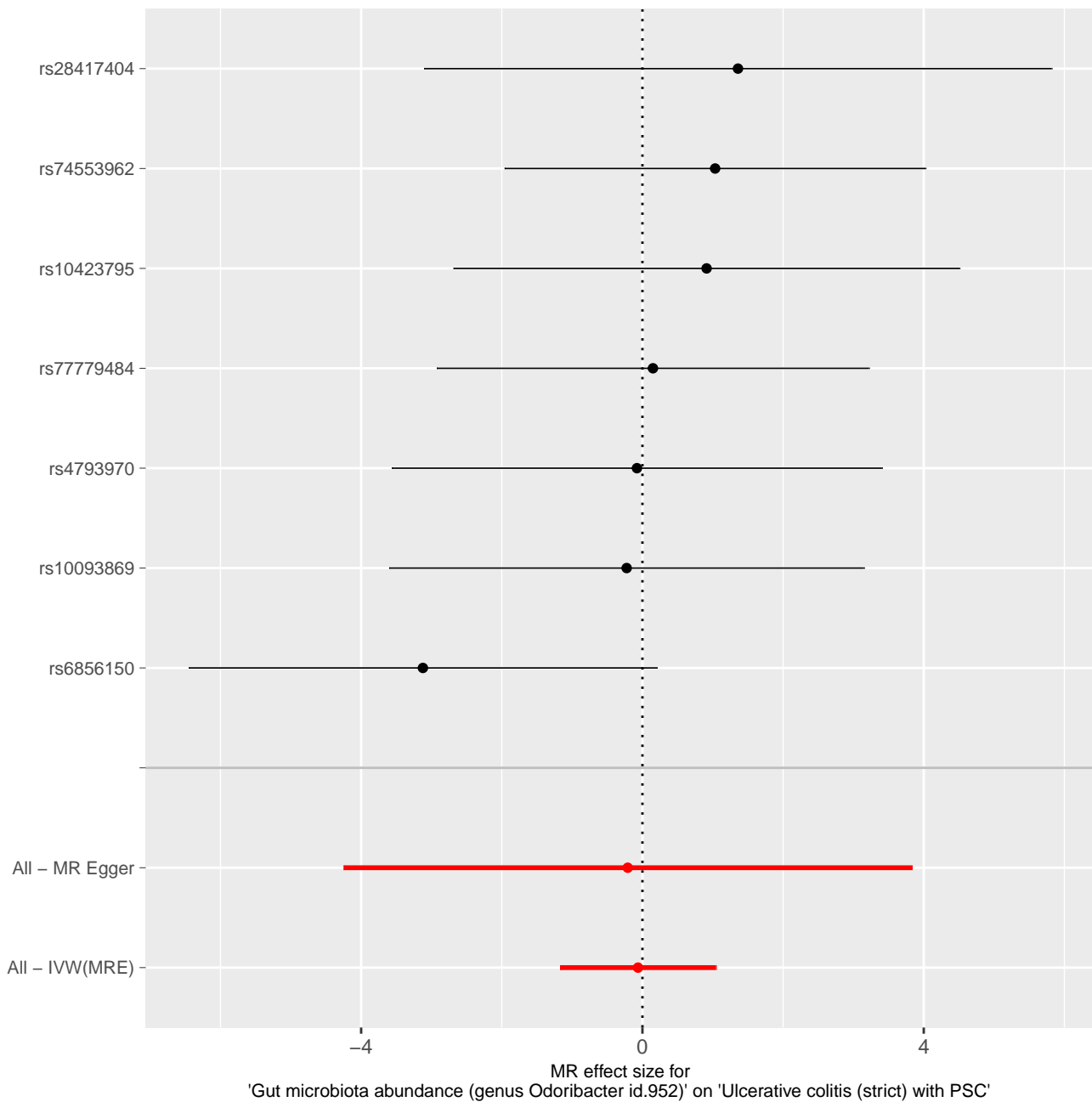
Batch 704 : Gut microbiota abundance (genus Lactobacillus id.1837) on Ulcerative colitis (strict) with PSC

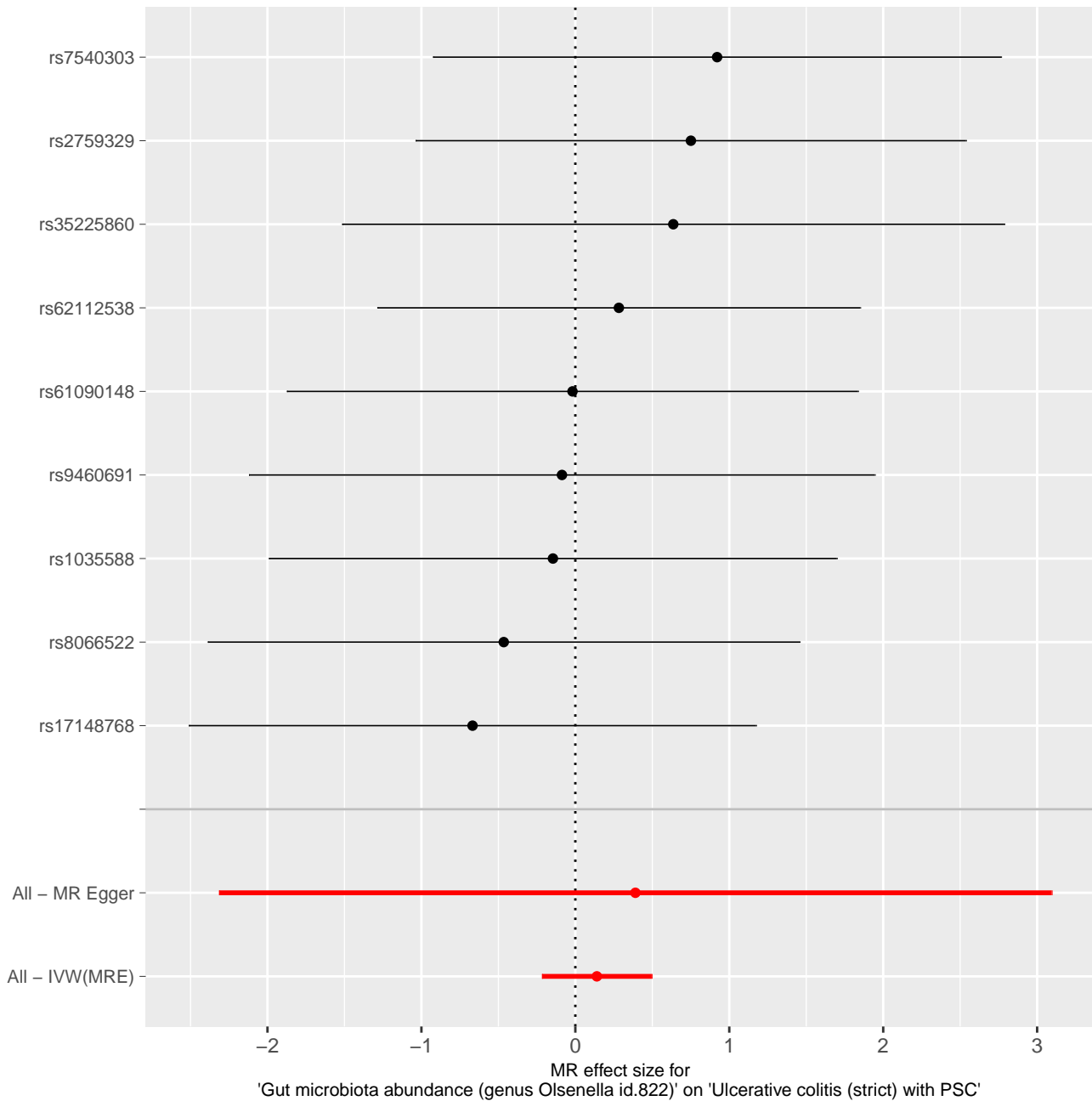


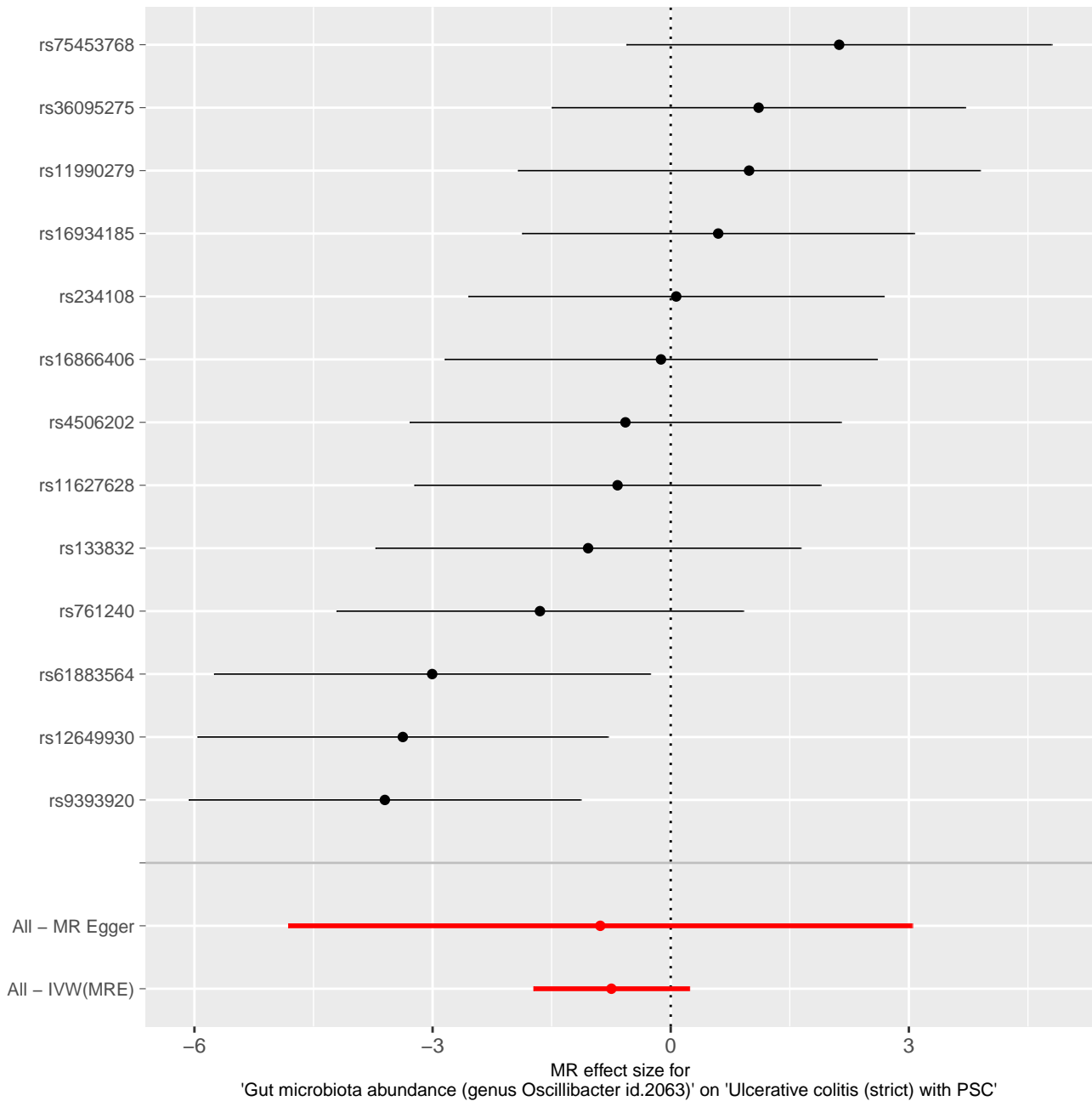


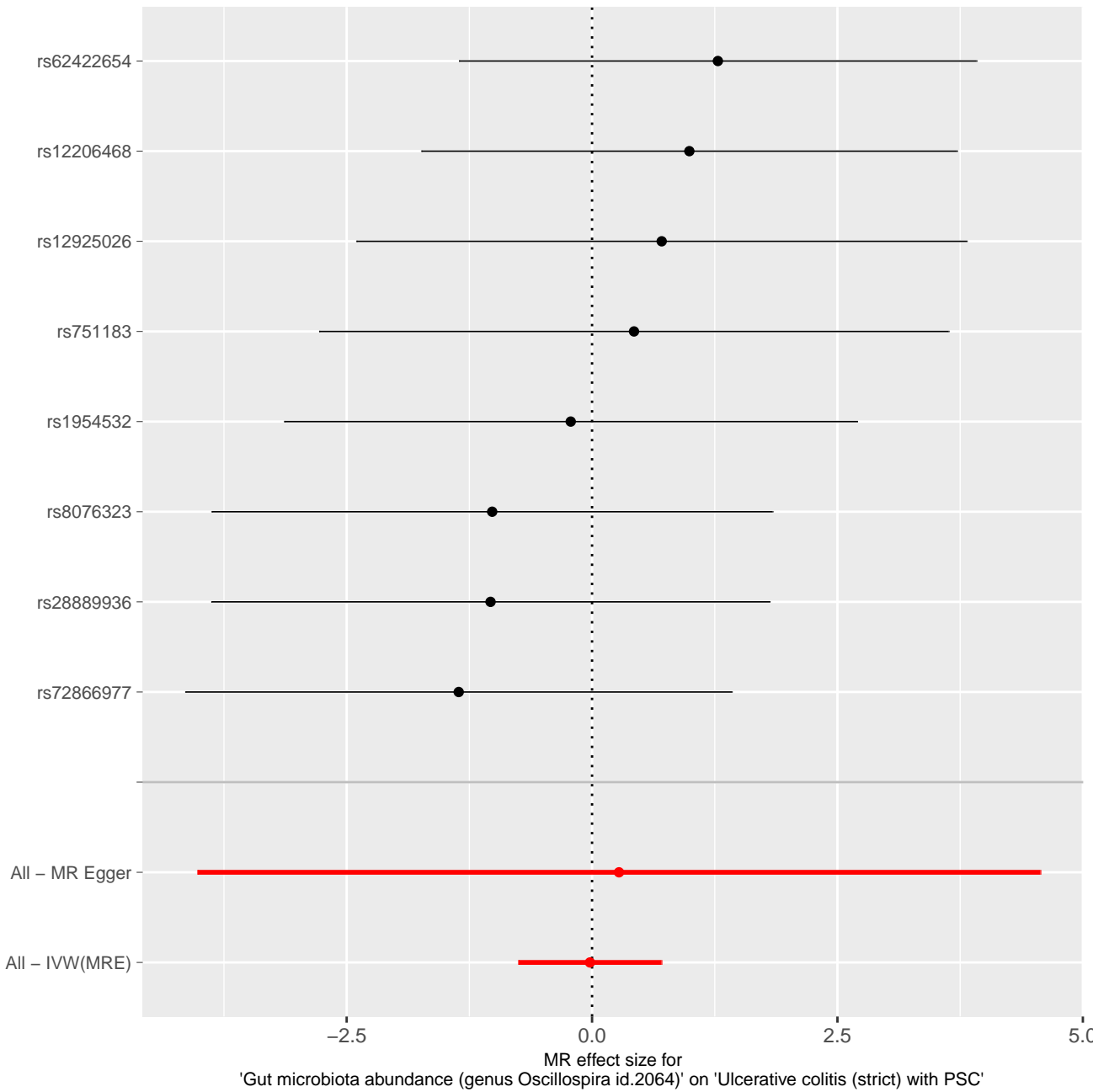


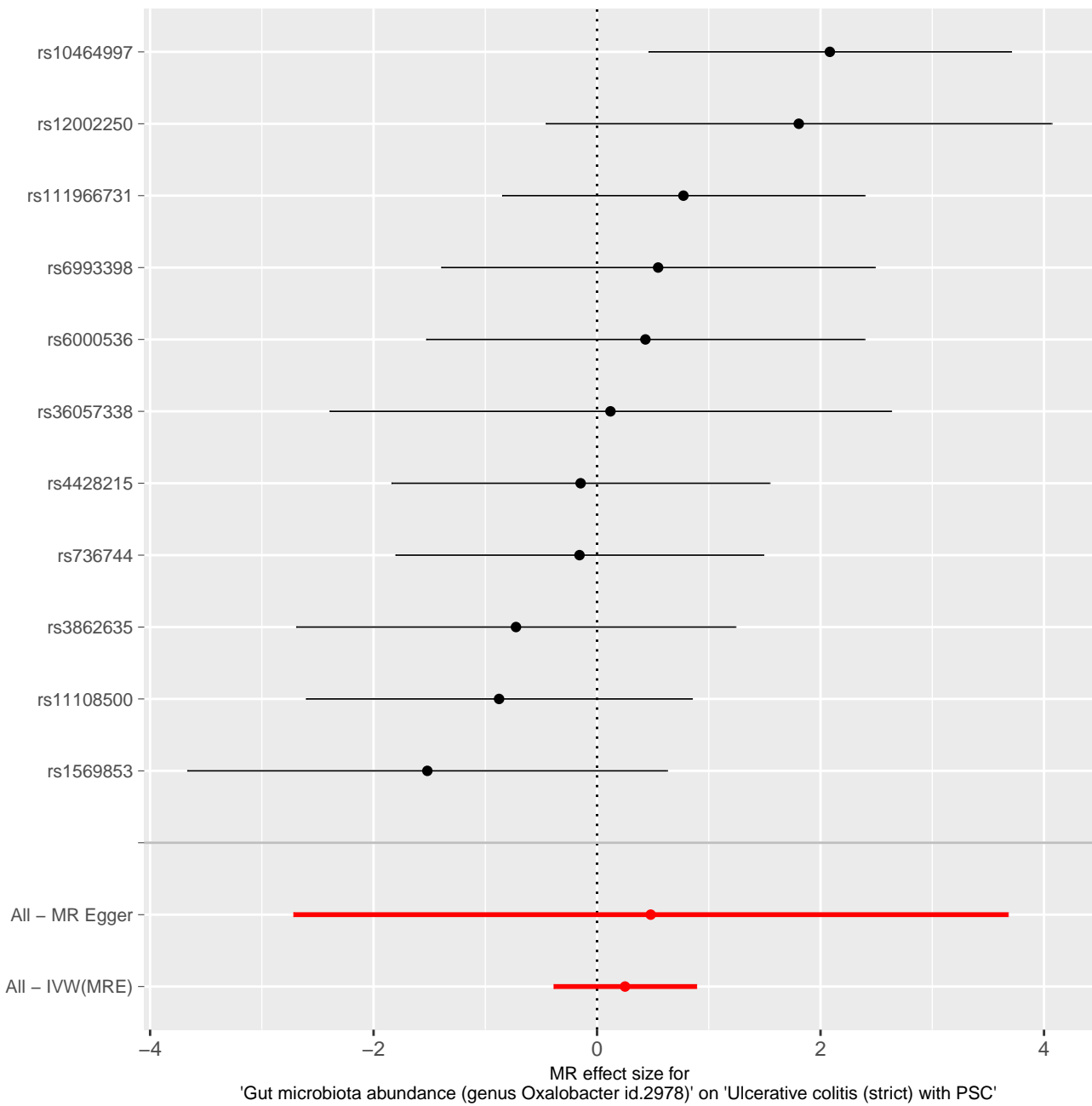


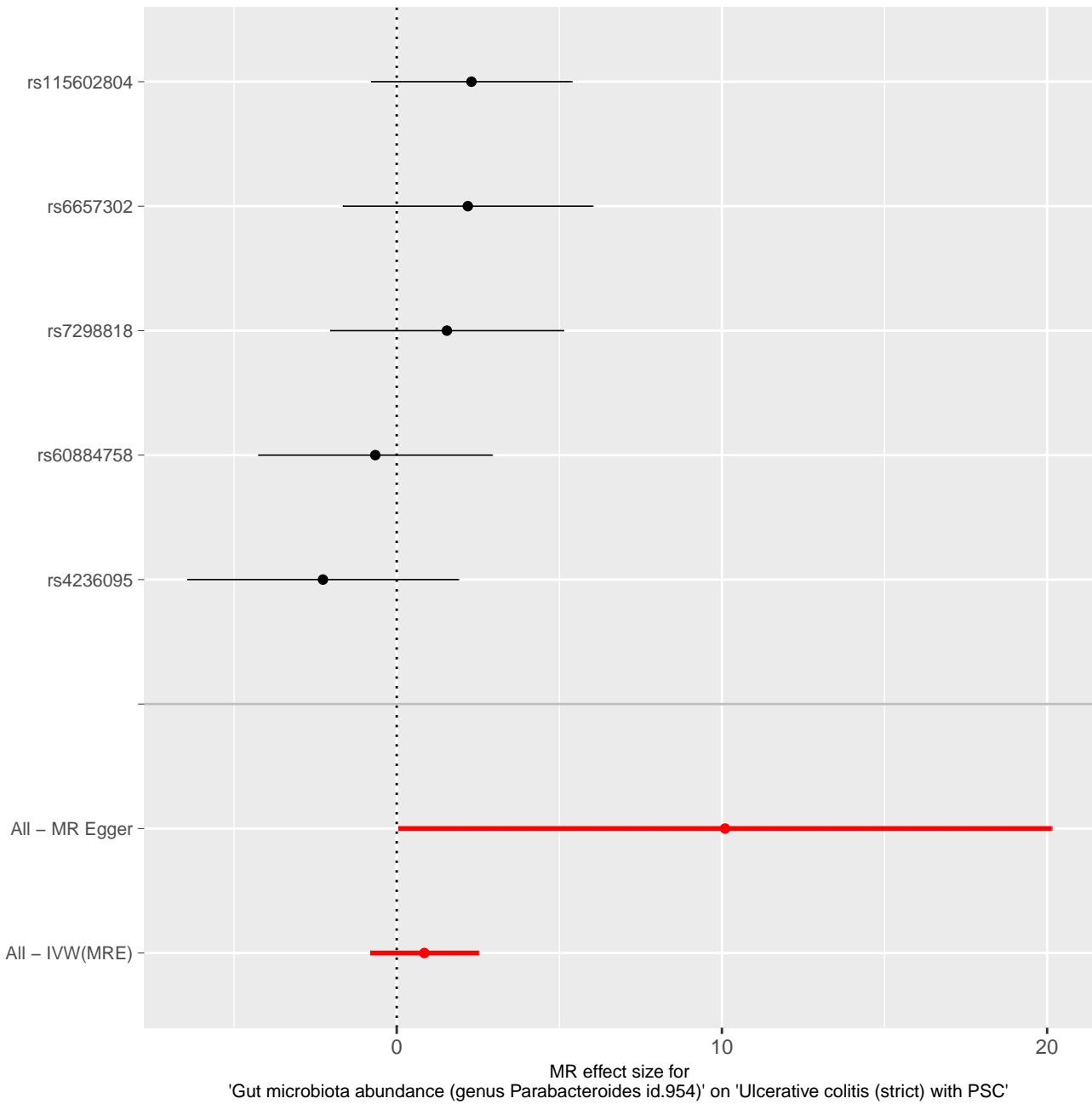




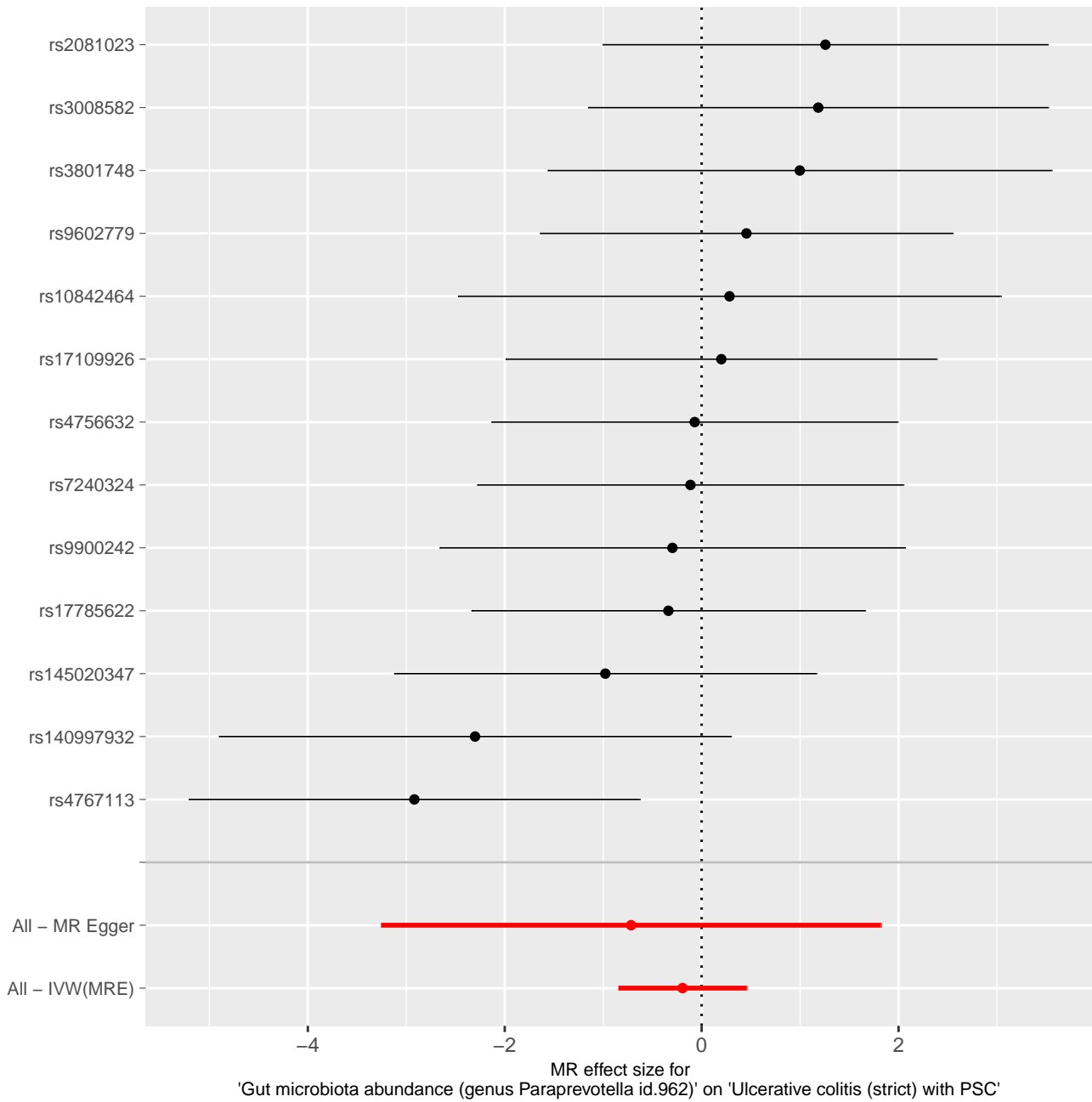




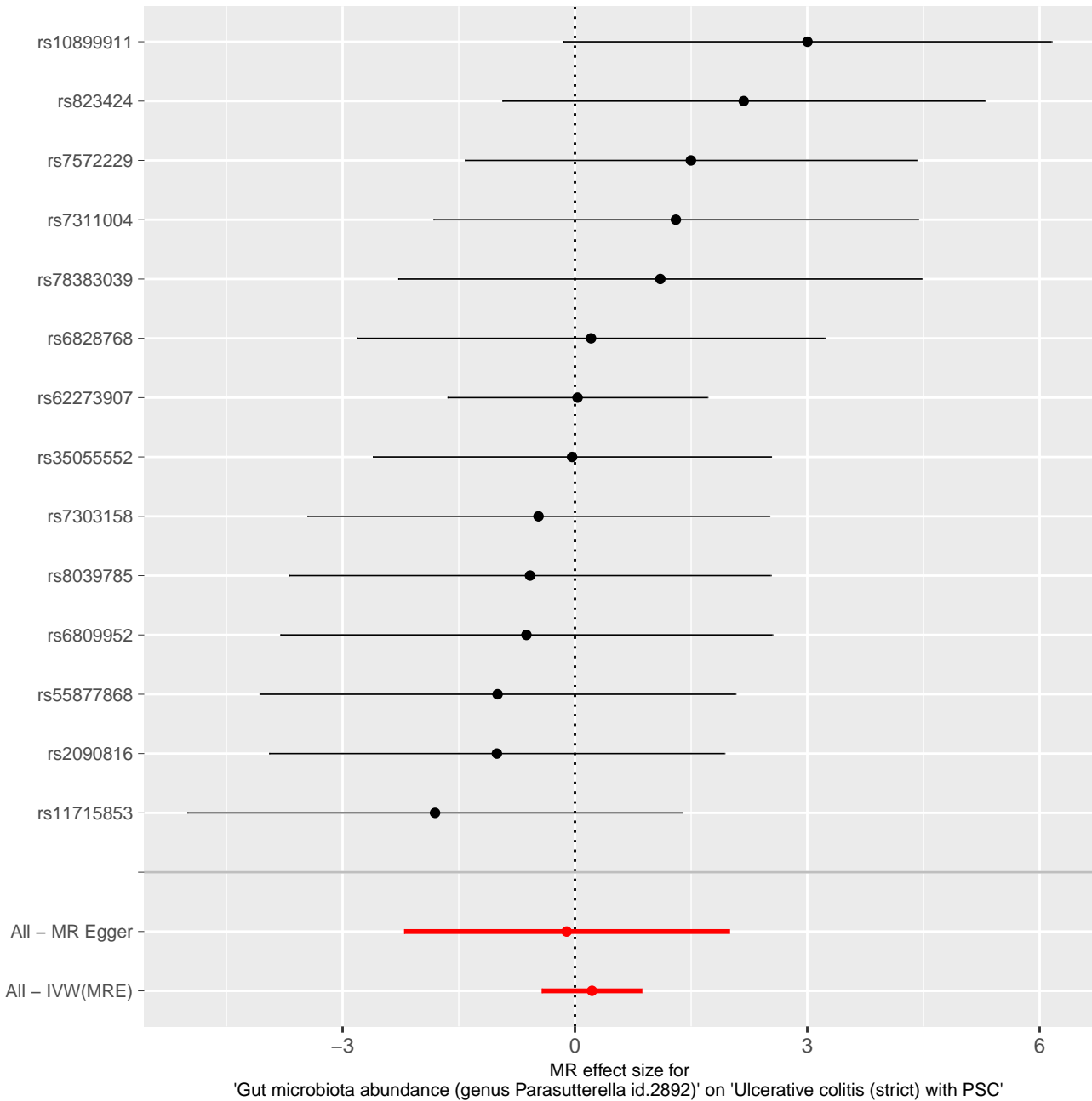


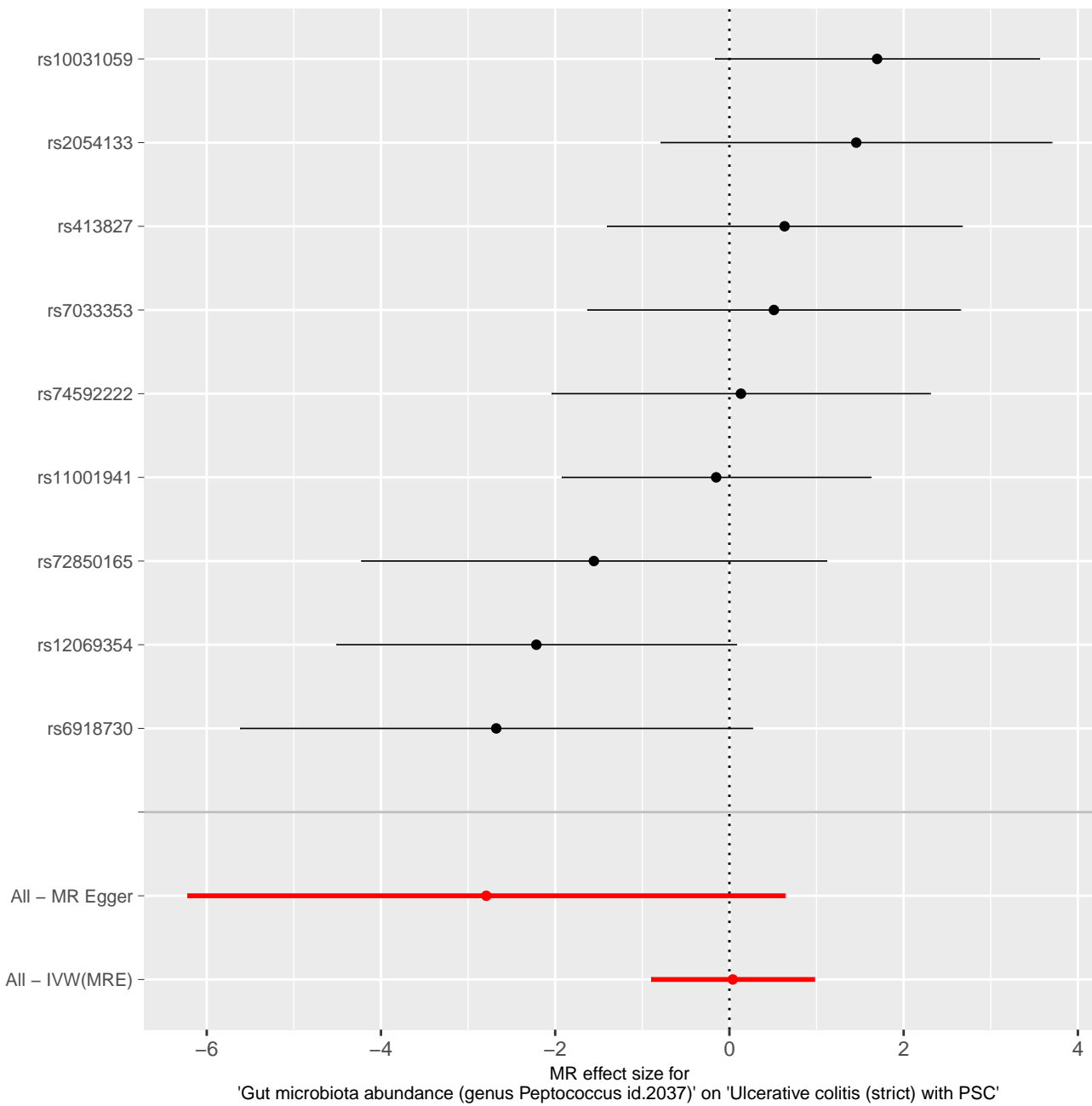


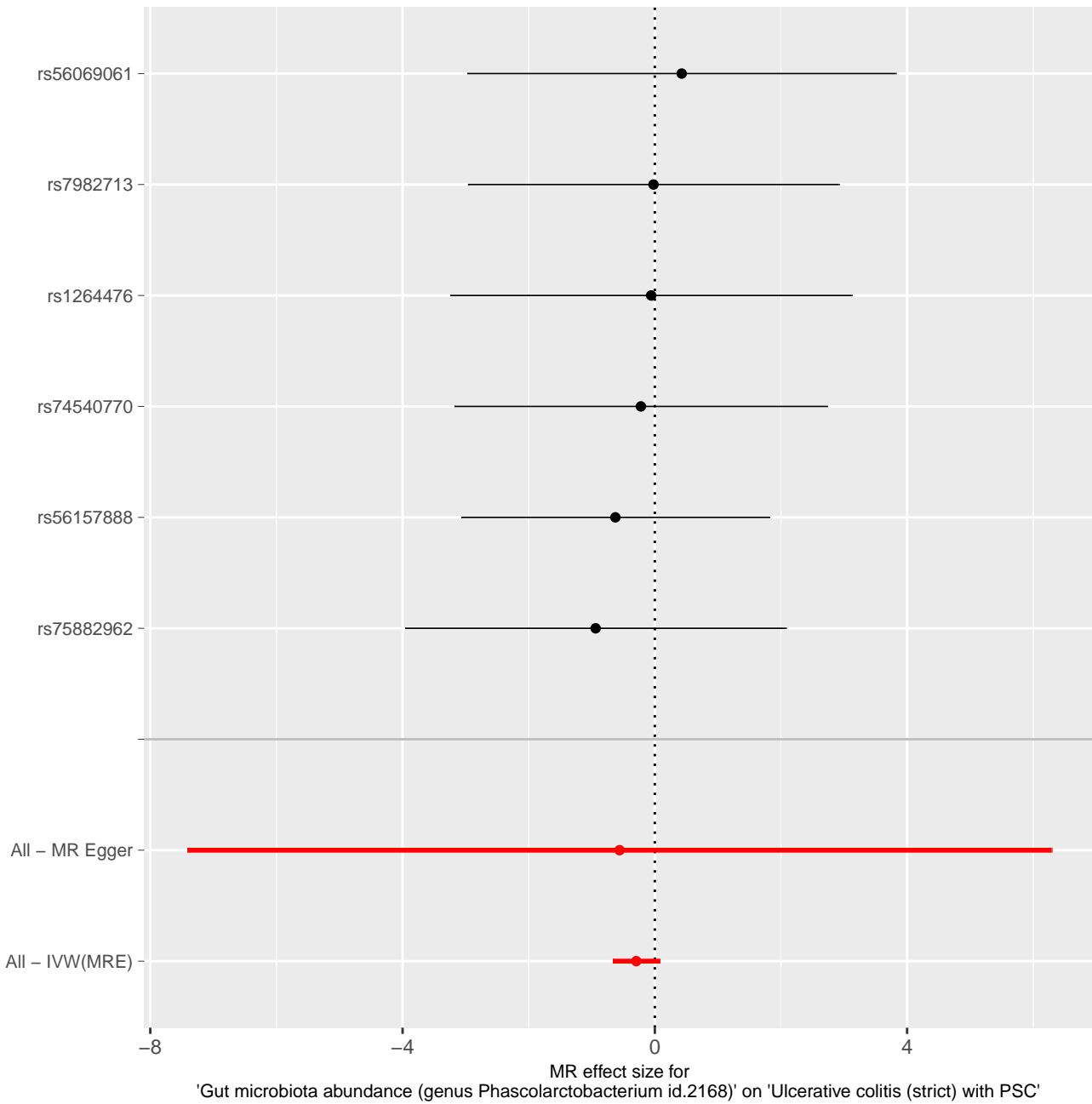
Batch 714 : Gut microbiota abundance (genus Paraprevotella id.962) on Ulcerative colitis (strict) with PSC



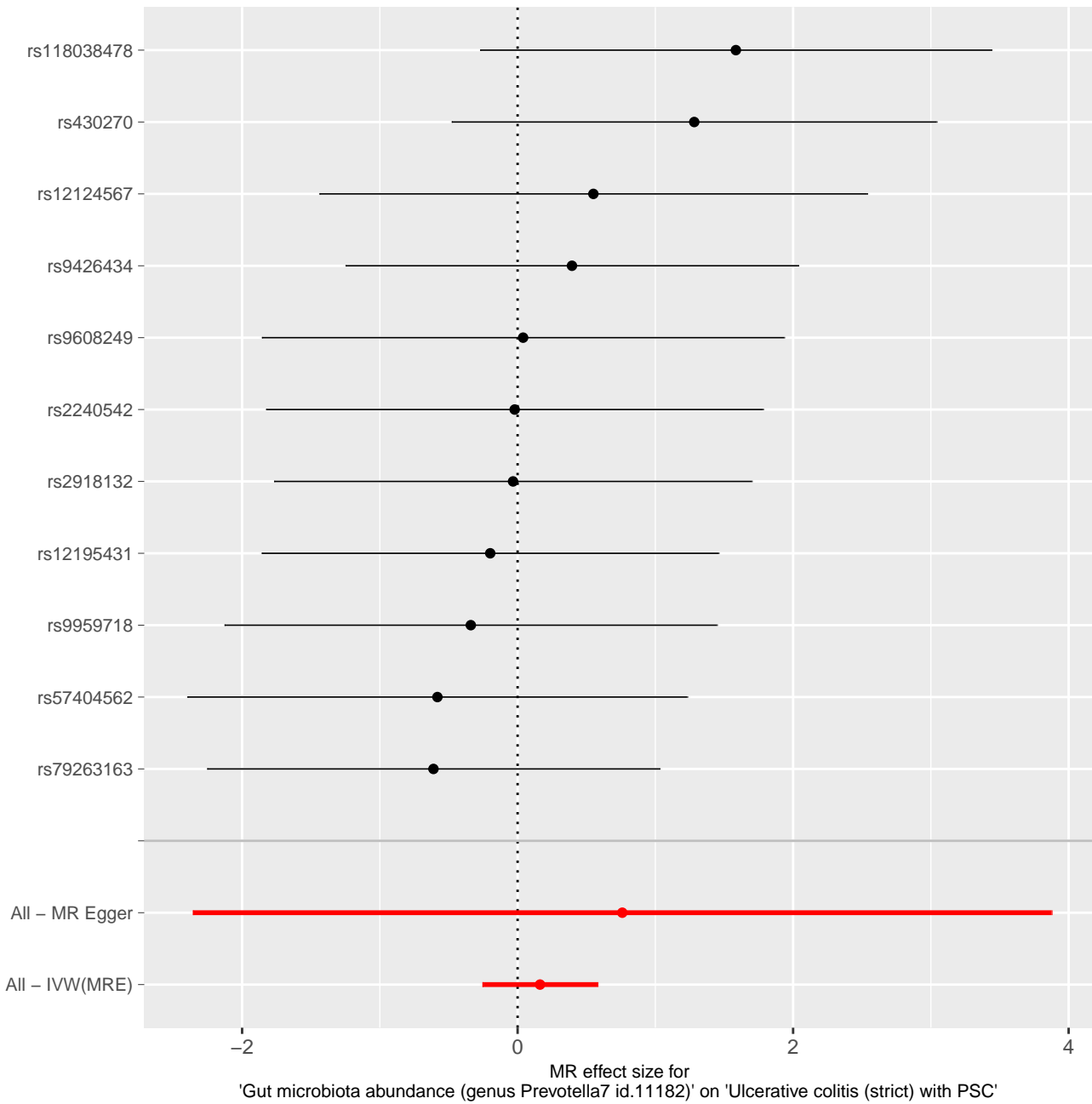
Batch 715 : Gut microbiota abundance (genus Parasutterella id.2892) on Ulcerative colitis (strict) with PSC



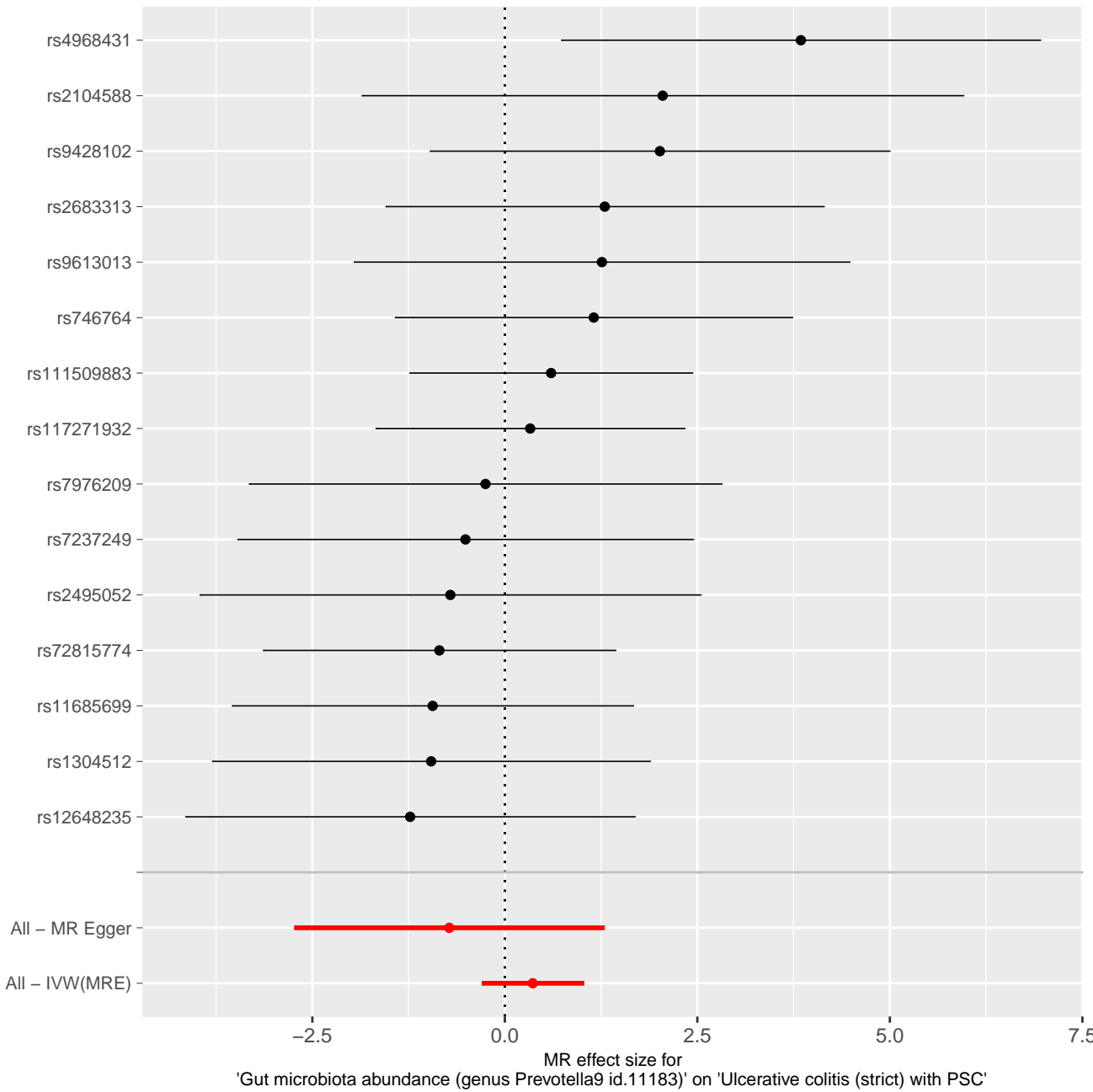


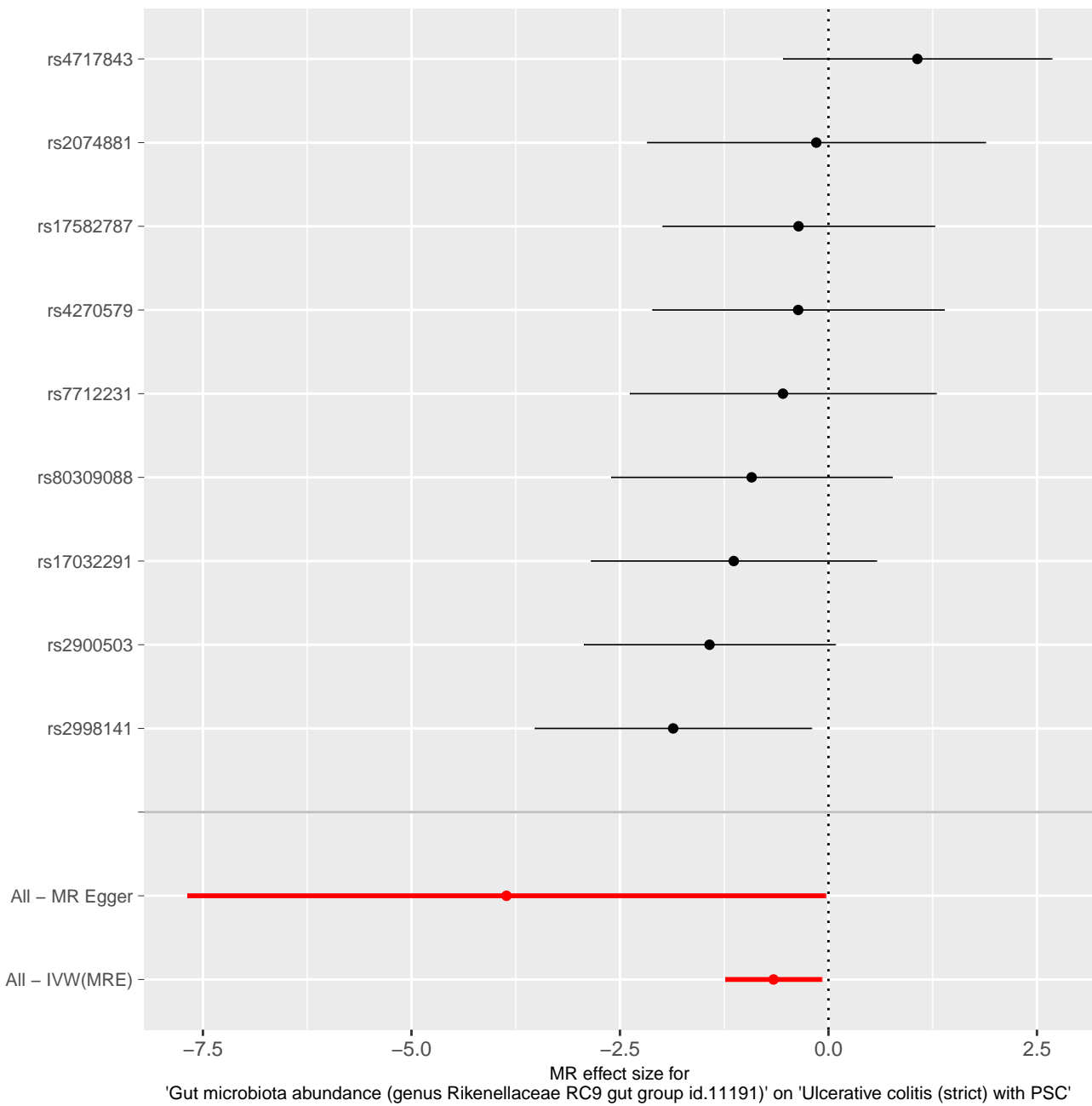


Batch 718 : Gut microbiota abundance (genus Prevotella7 id.11182) on Ulcerative colitis (strict) with PSC

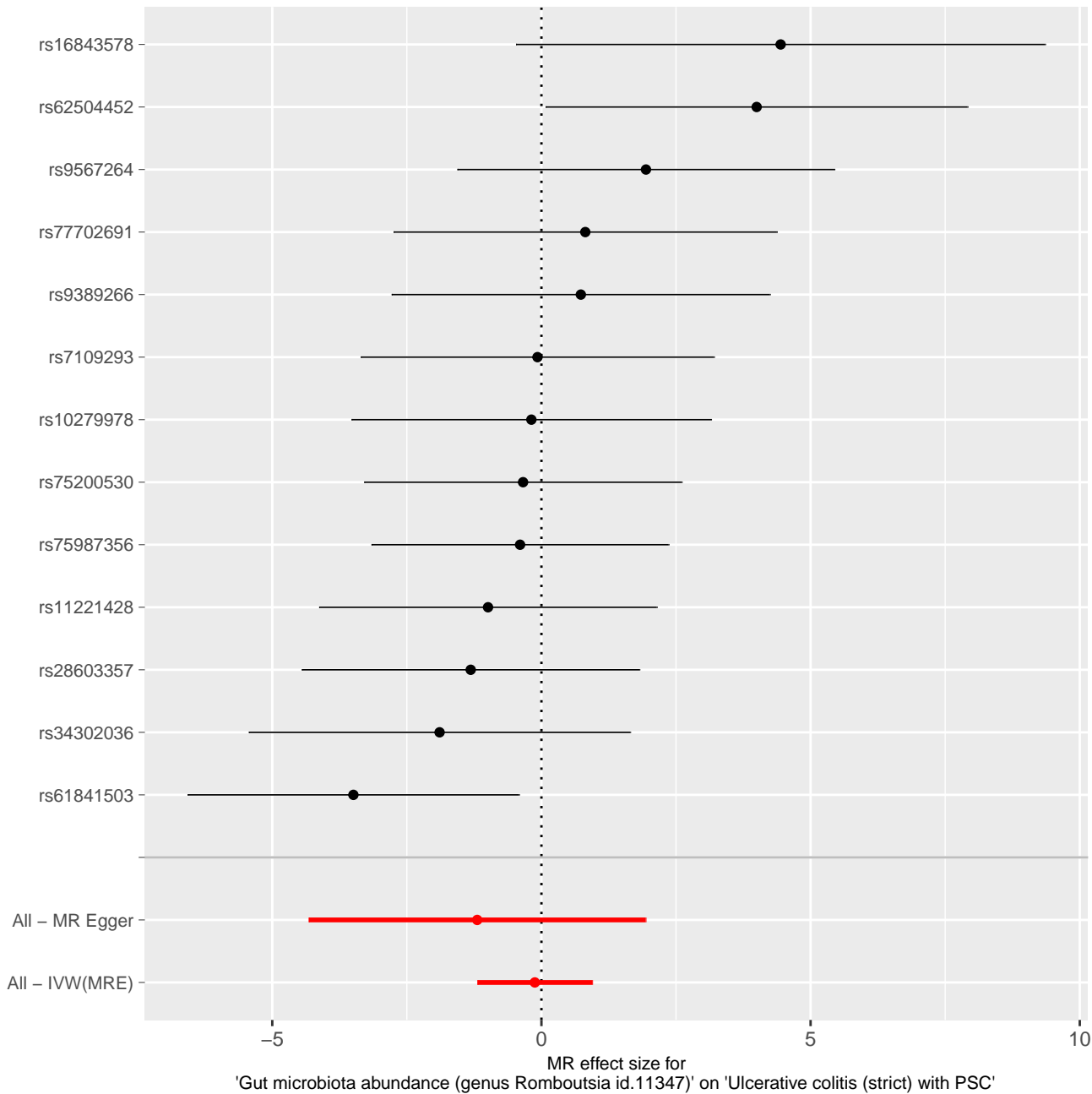


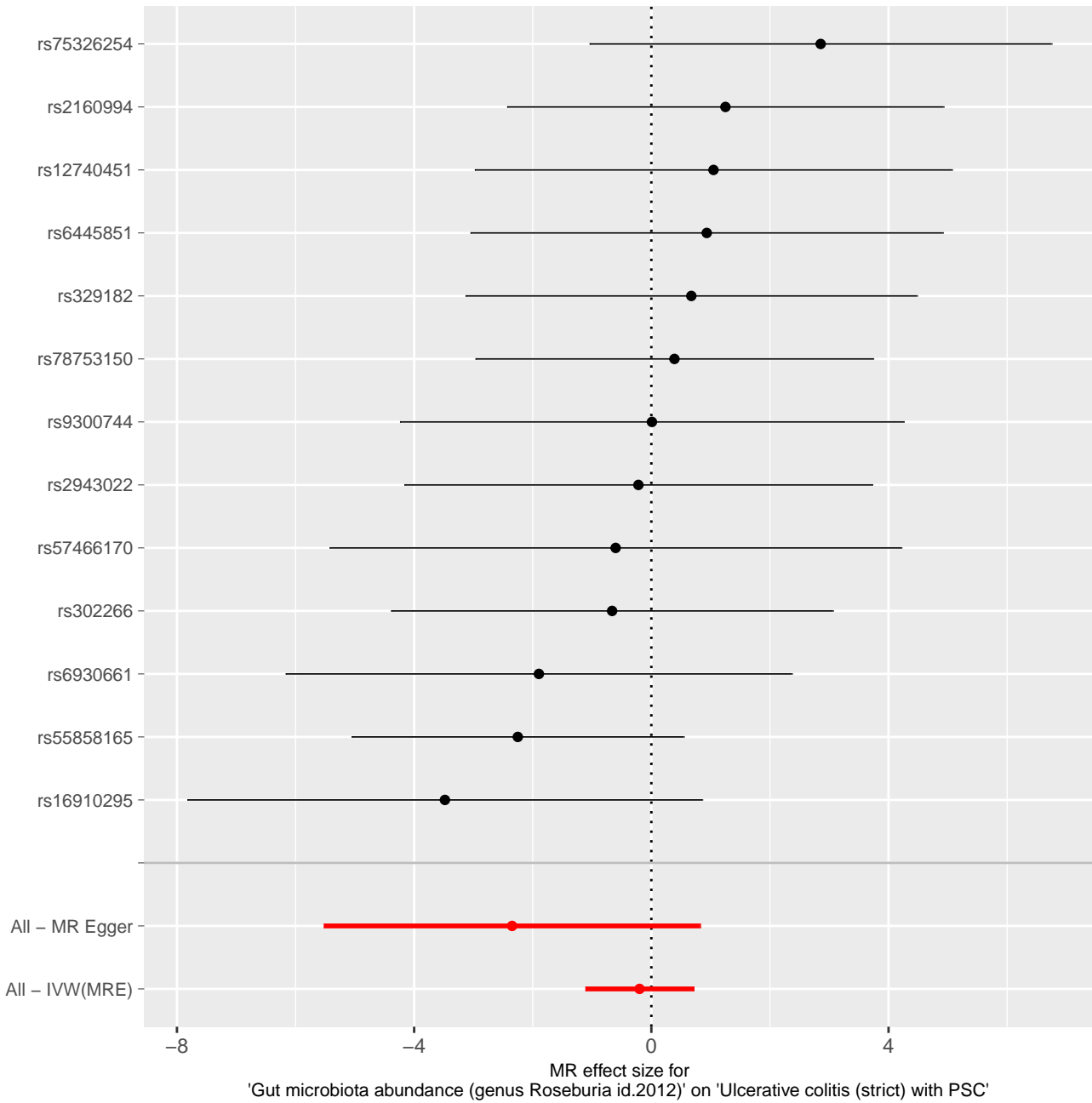
Batch 719 : Gut microbiota abundance (genus Prevotella9 id.11183) on Ulcerative colitis (strict) with PSC

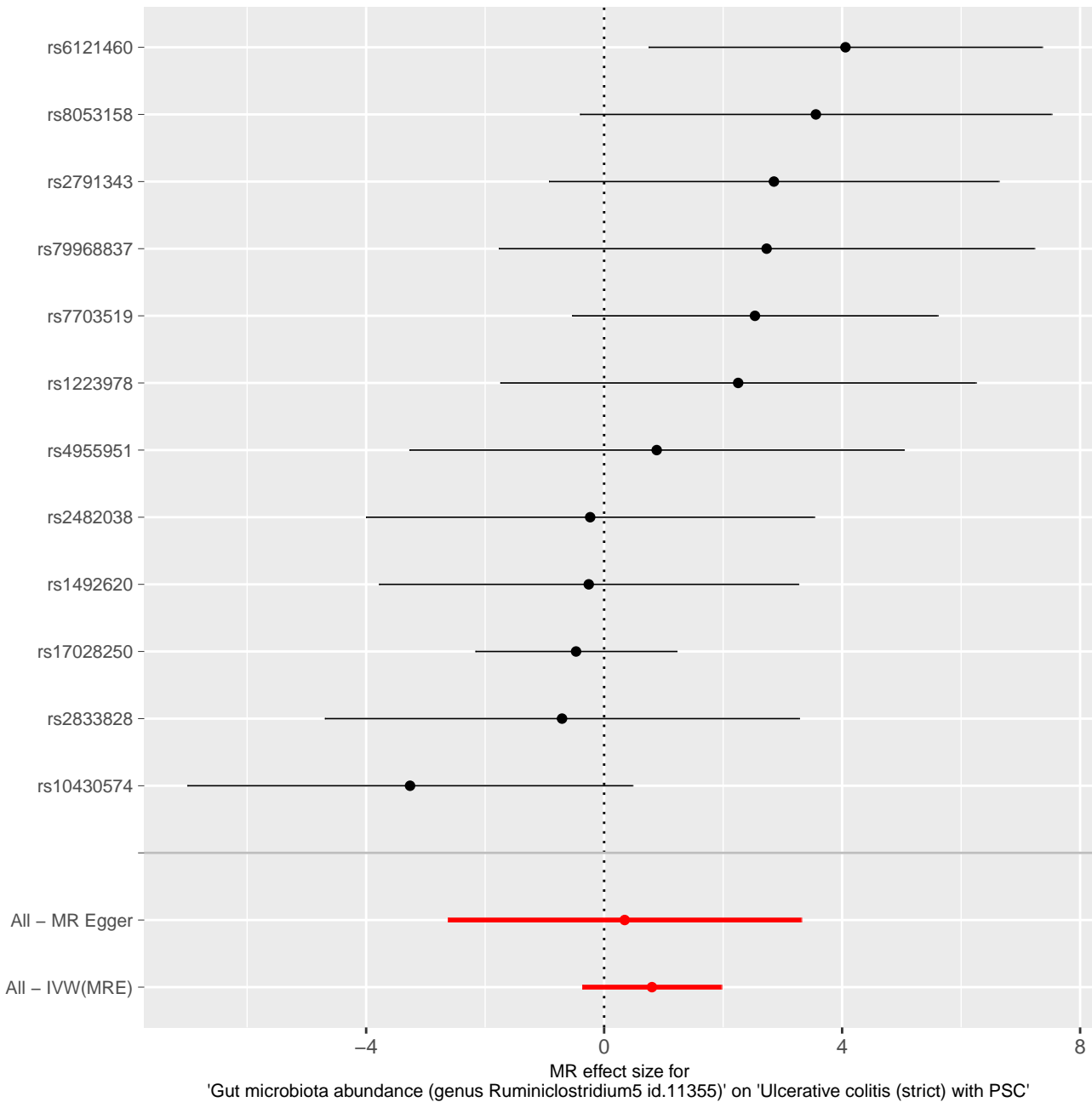


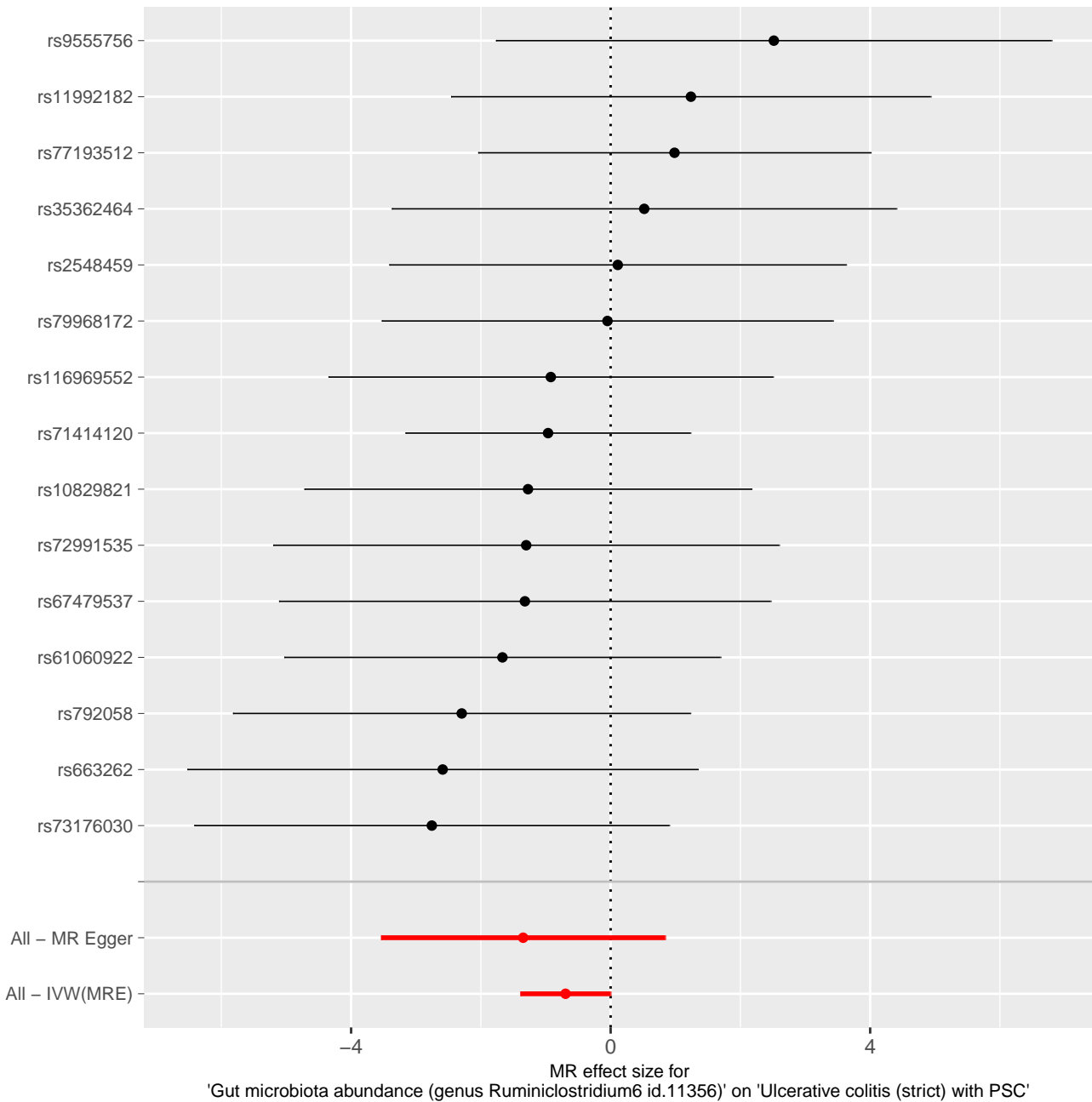


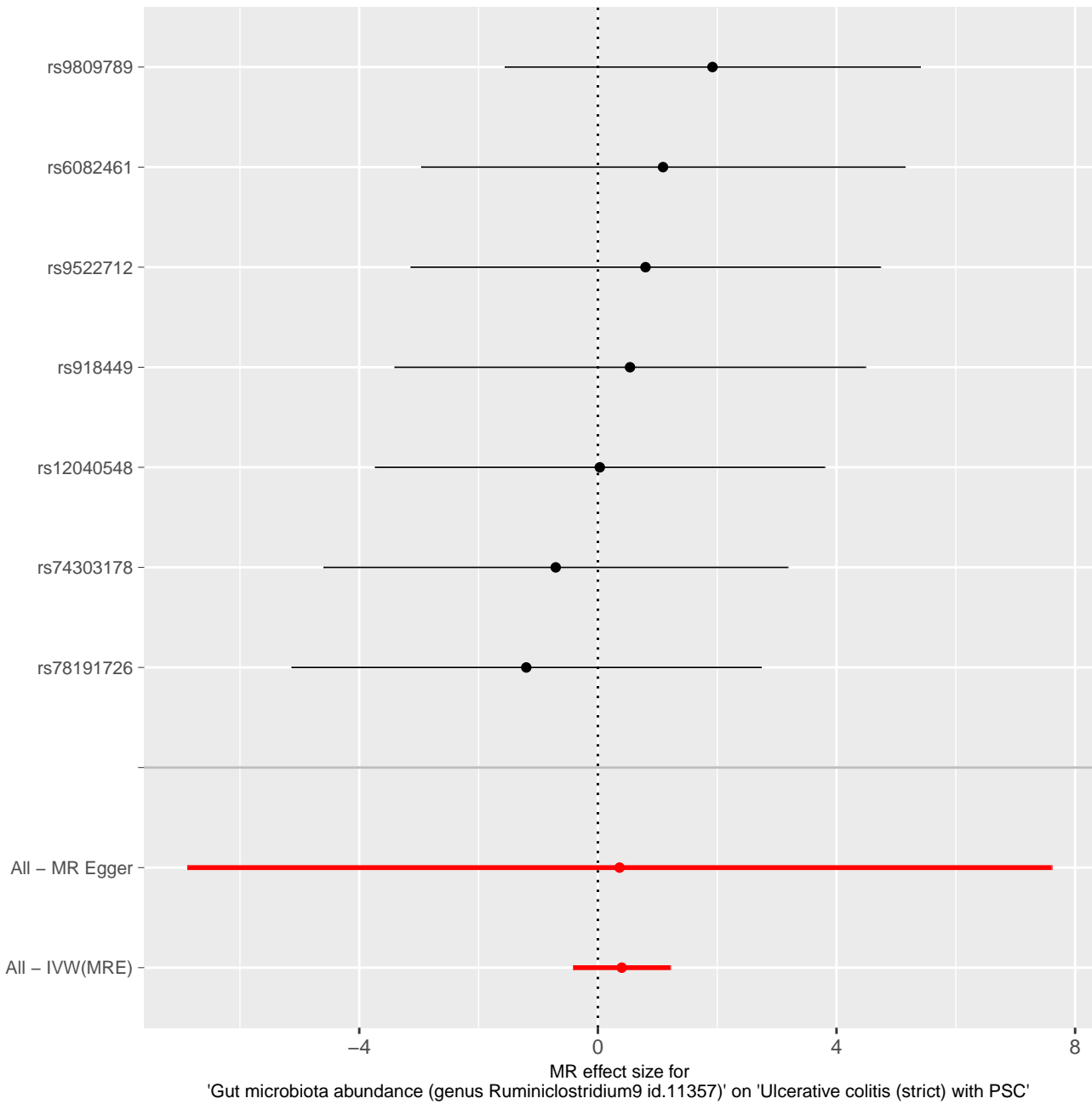
Batch 721 : Gut microbiota abundance (genus Romboutsia id.11347) on Ulcerative colitis (strict) with PSC

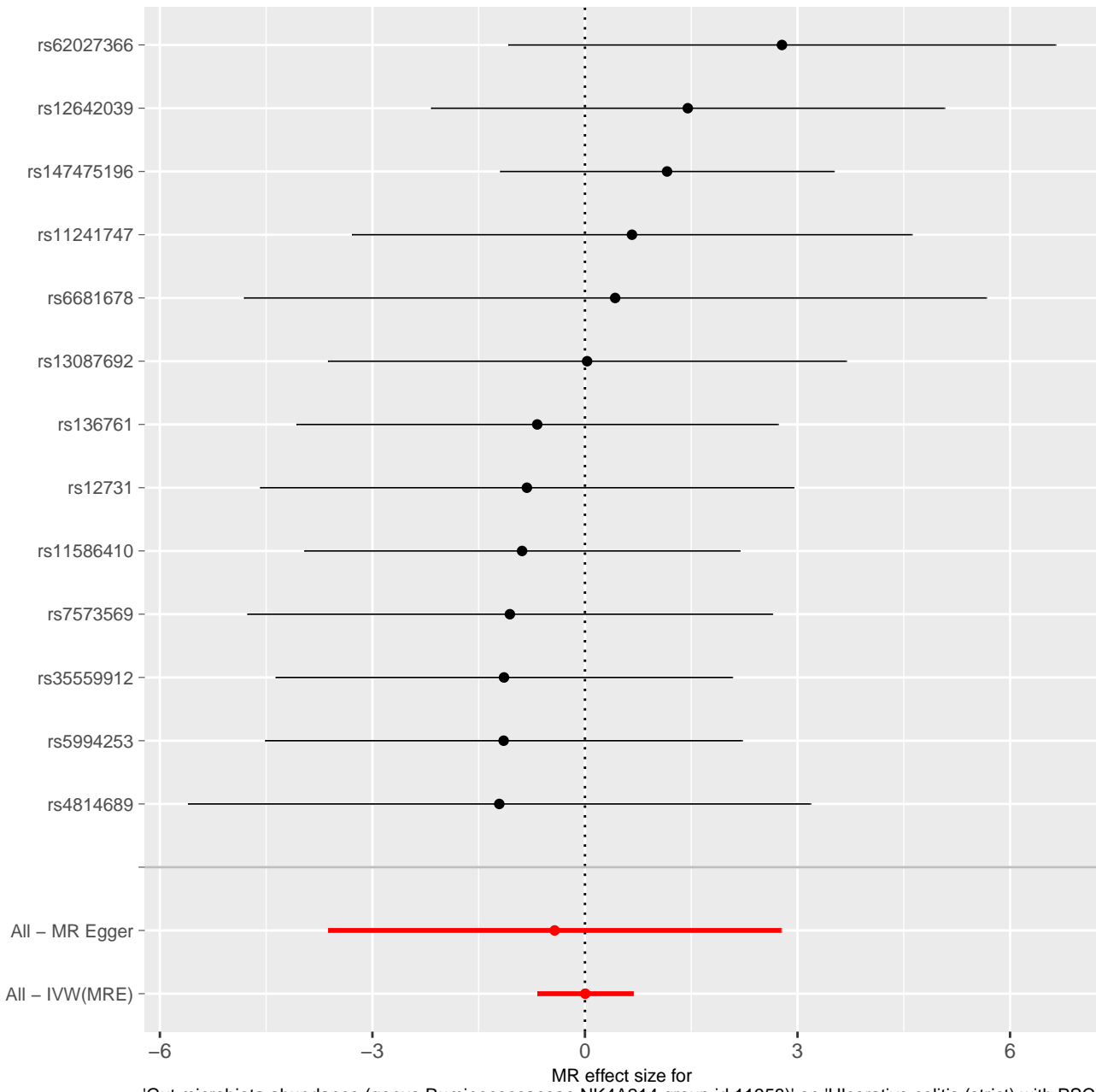




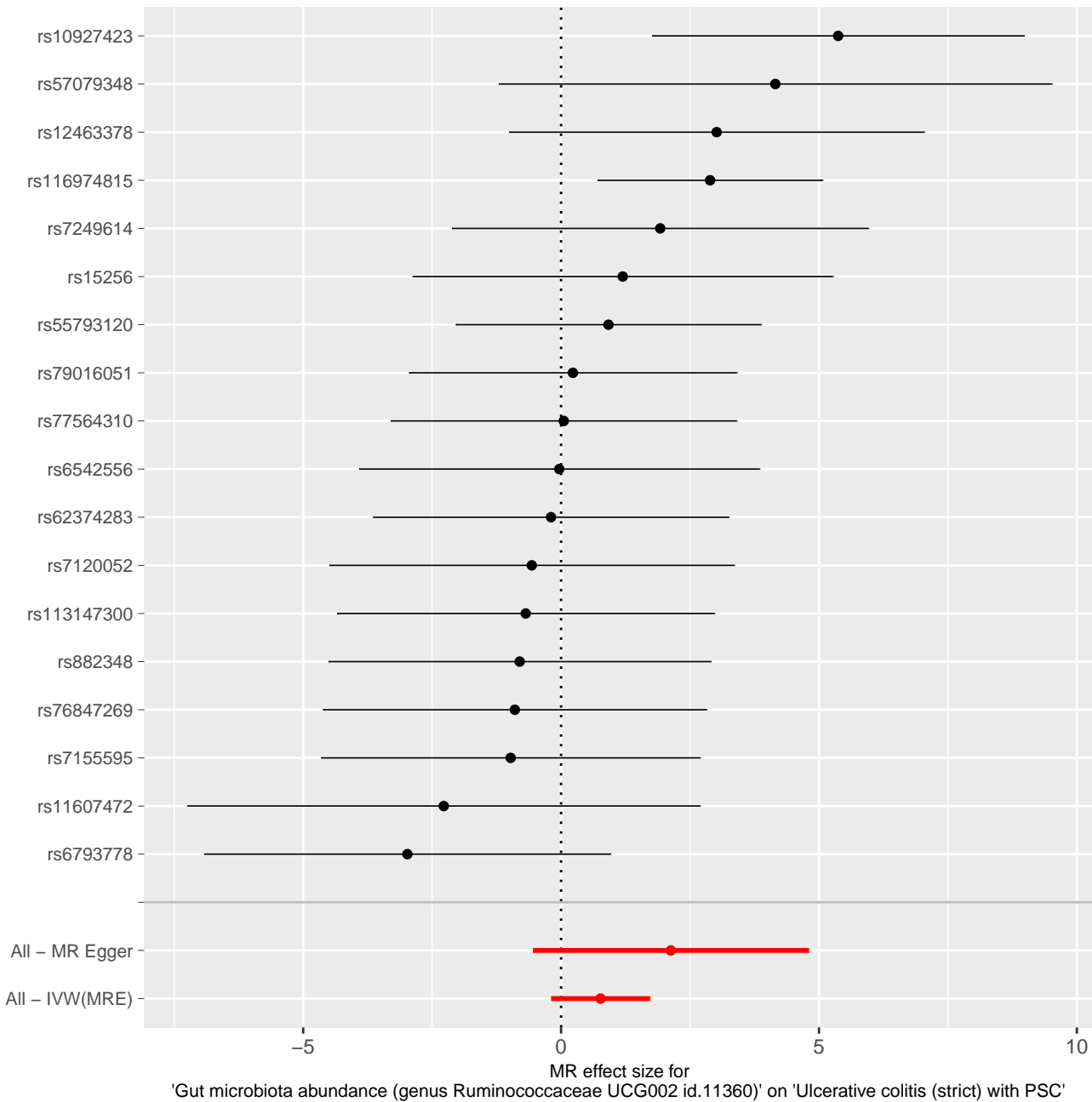


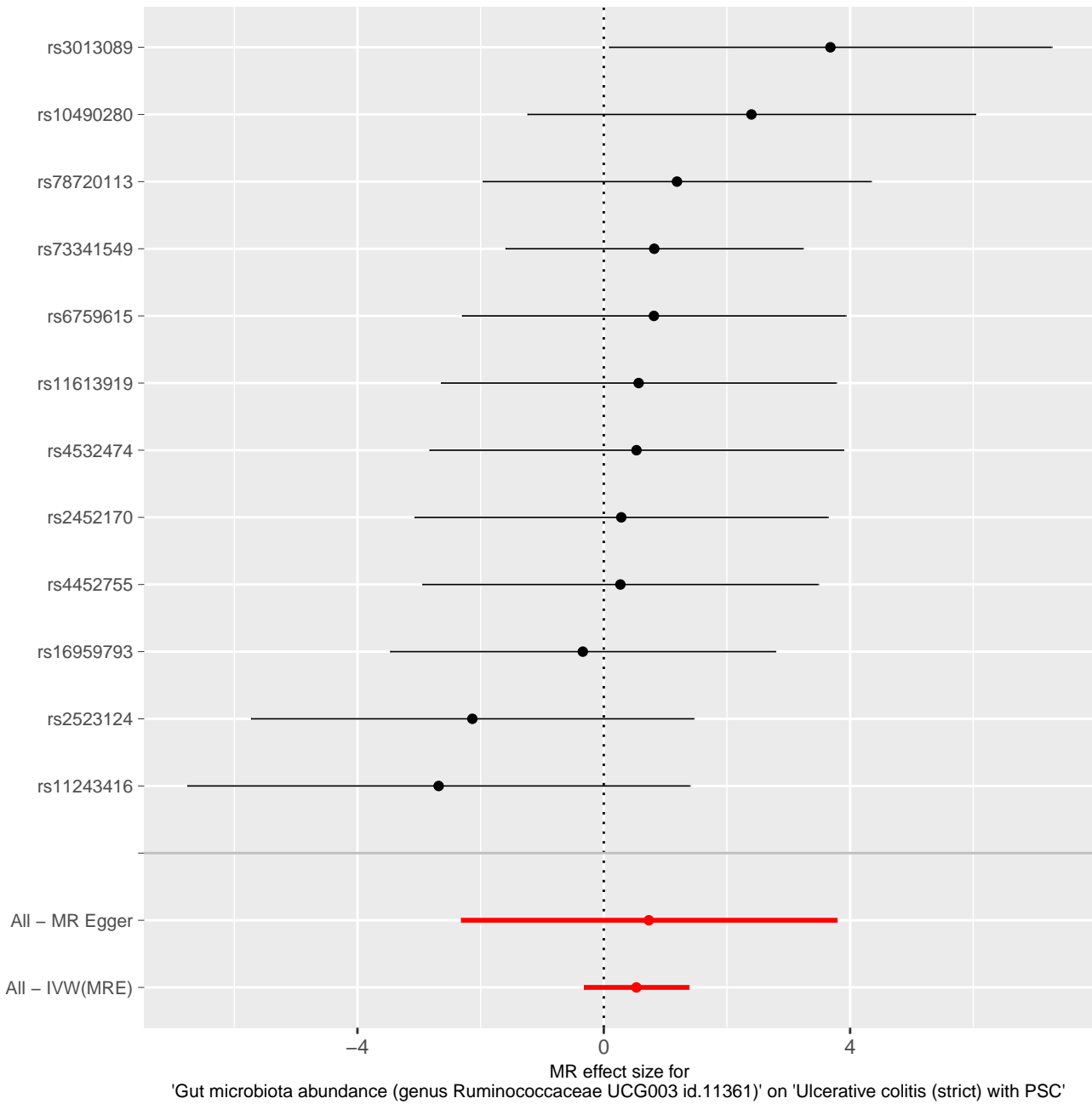


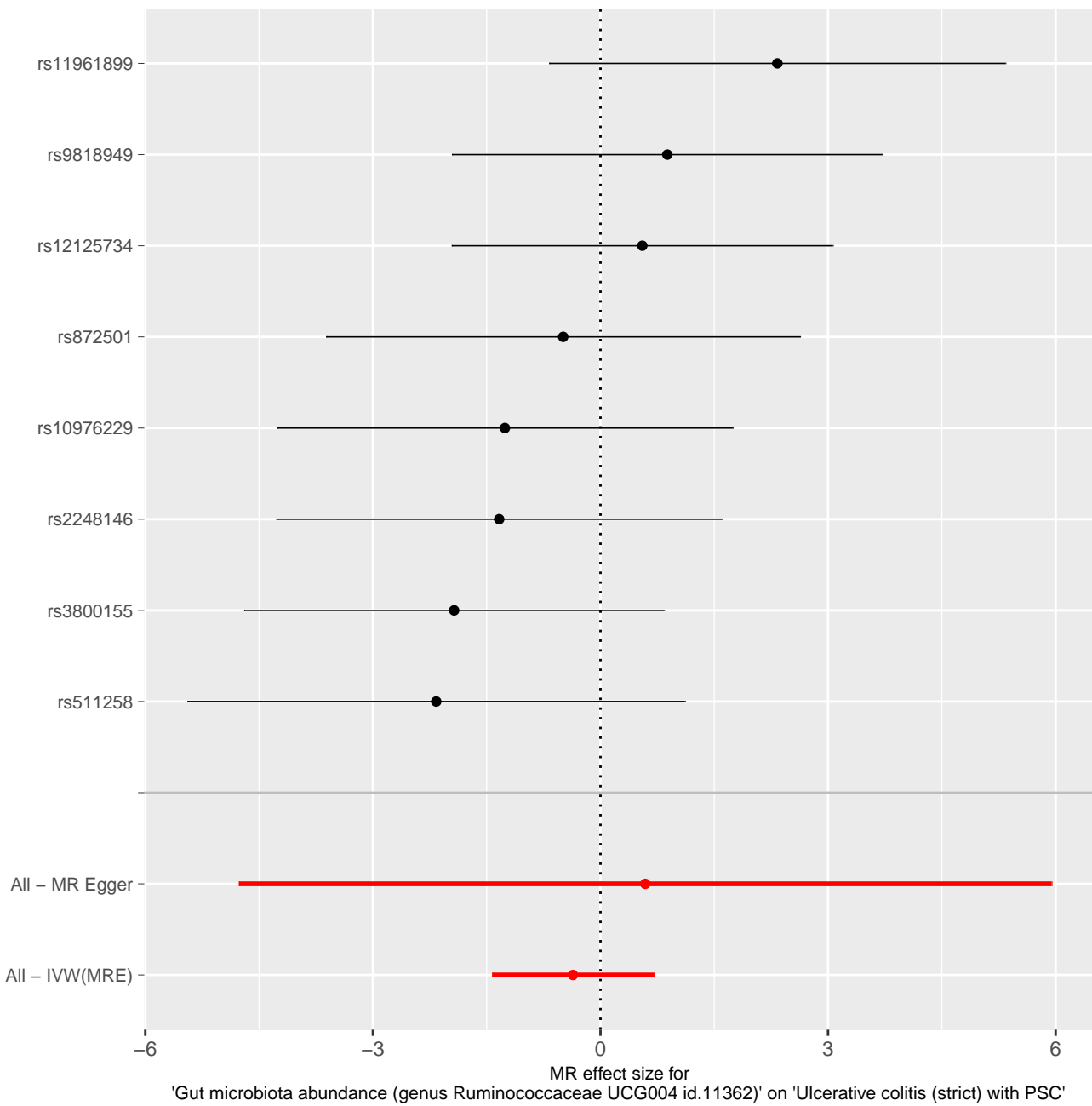


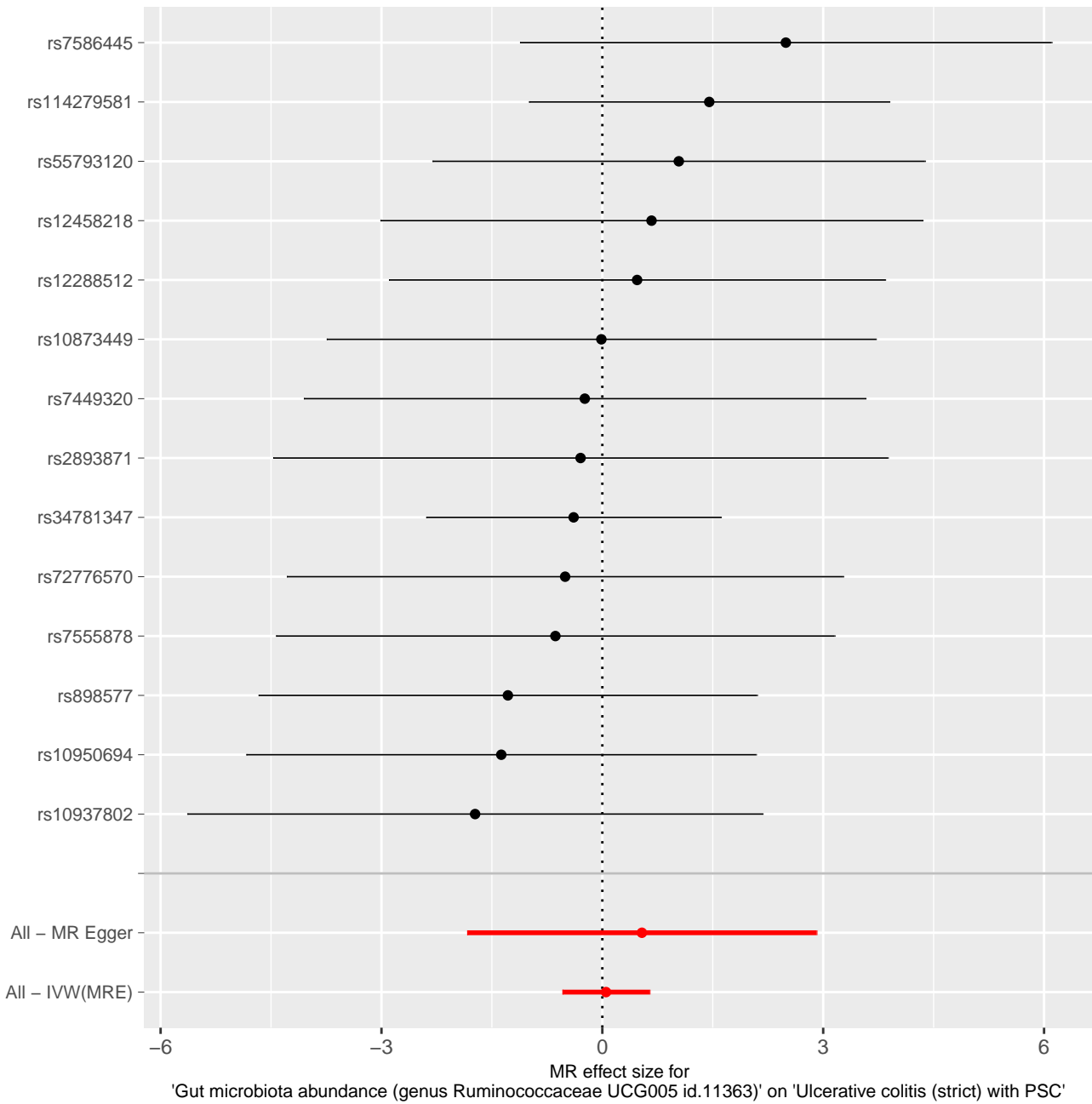


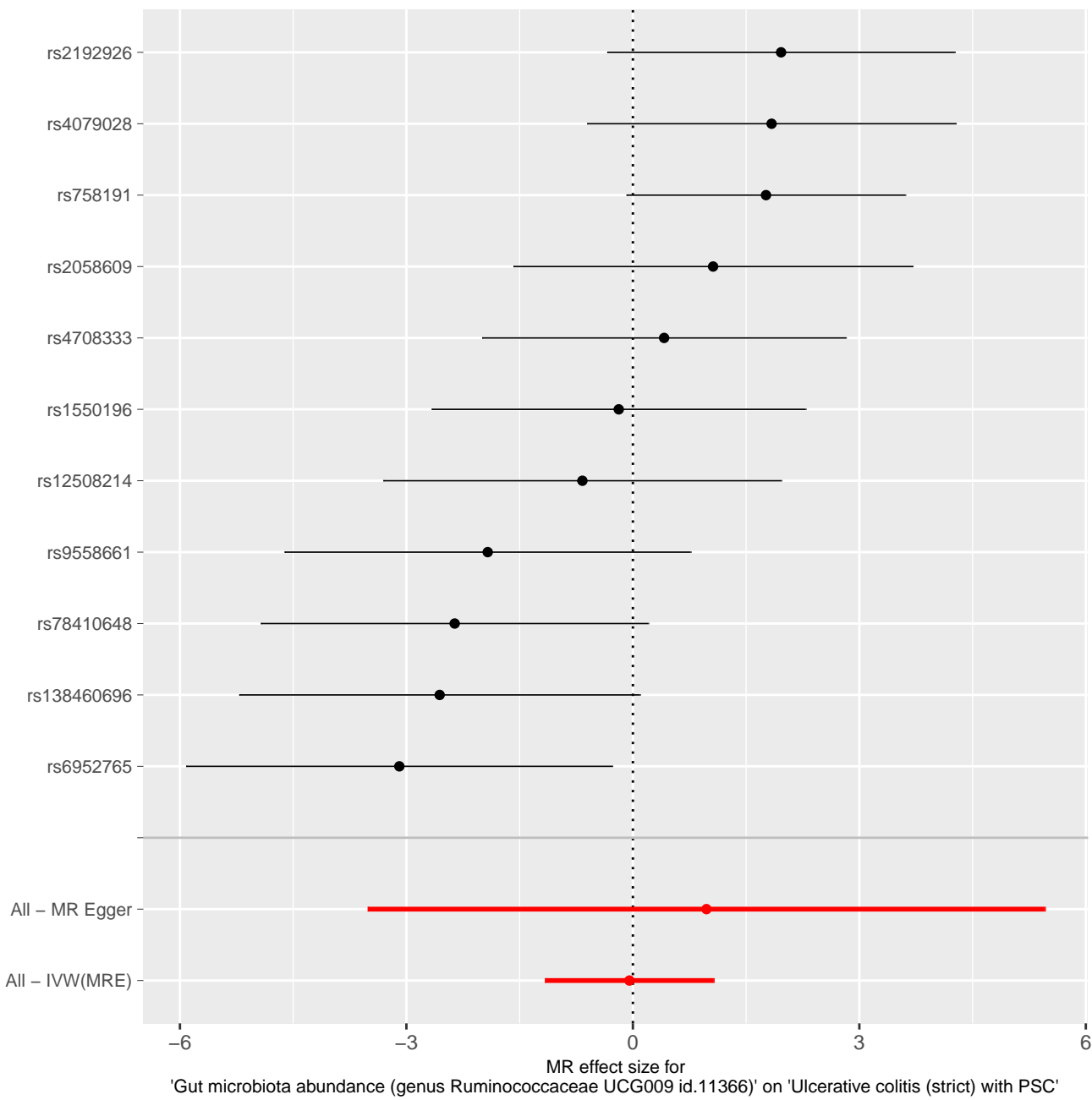
MR effect size for 'Gut microbiota abundance (genus Ruminococcaceae NK4A214 group id.11358)' on 'Ulcerative colitis (strict) with PSC'

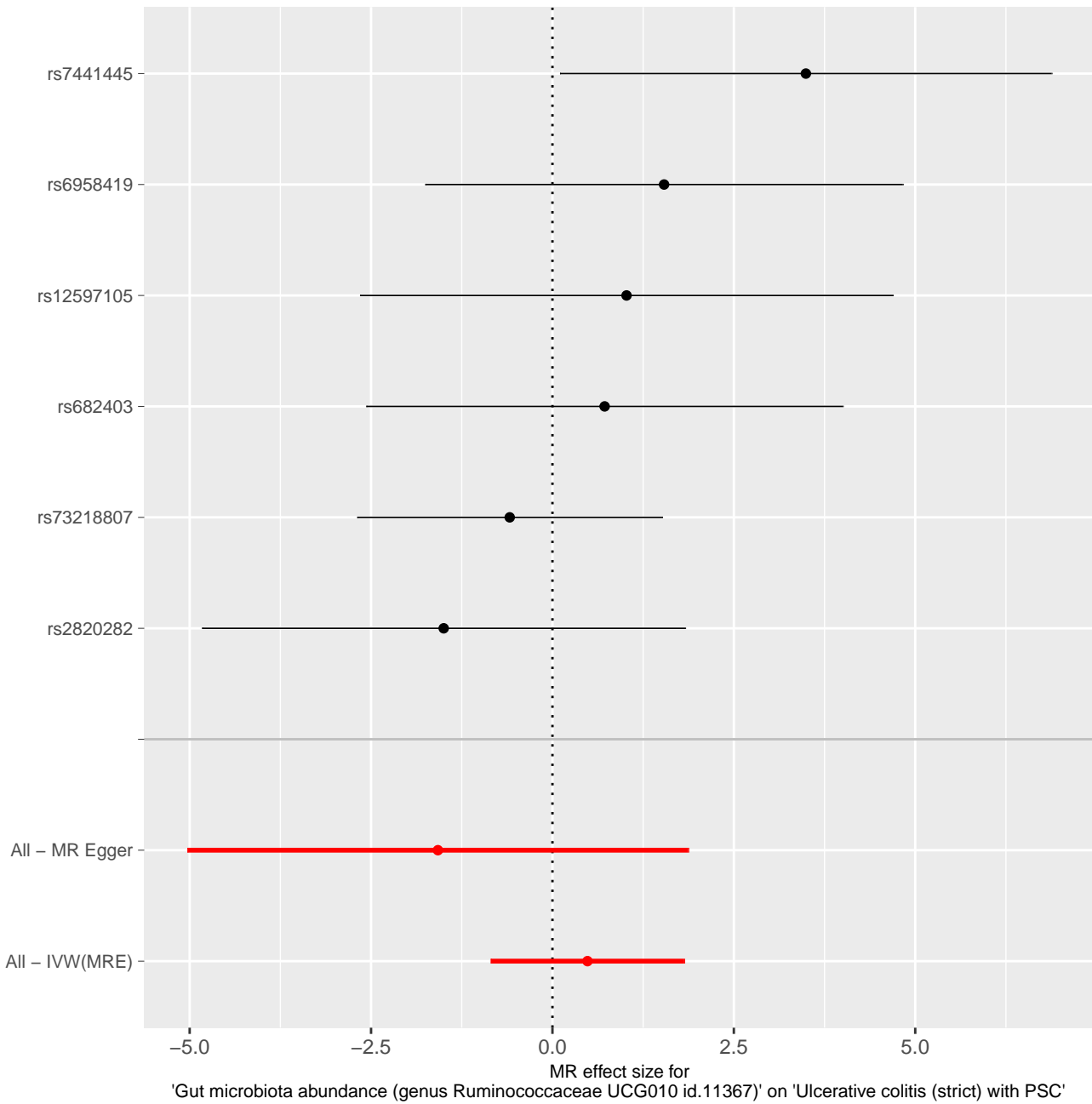


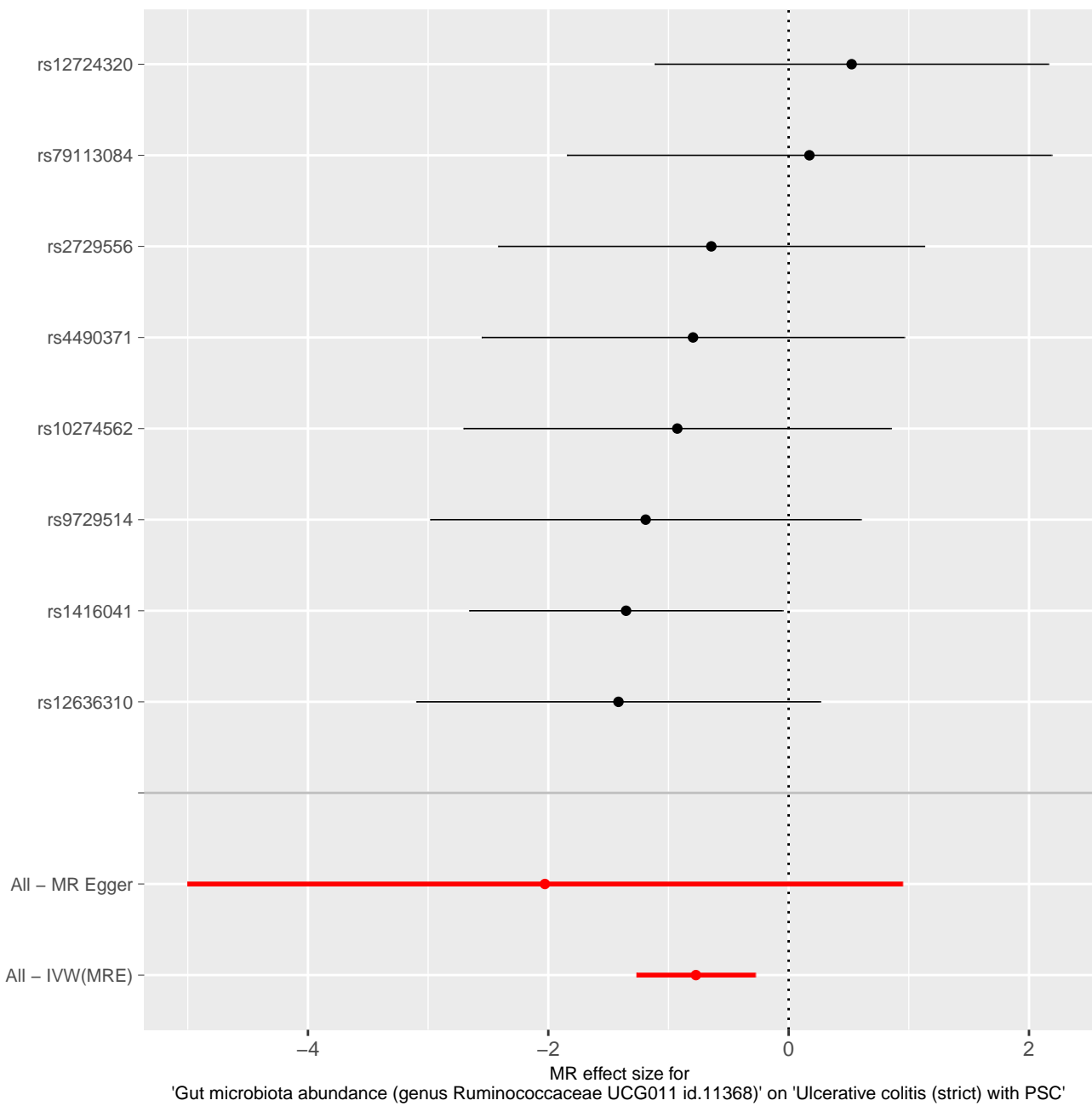




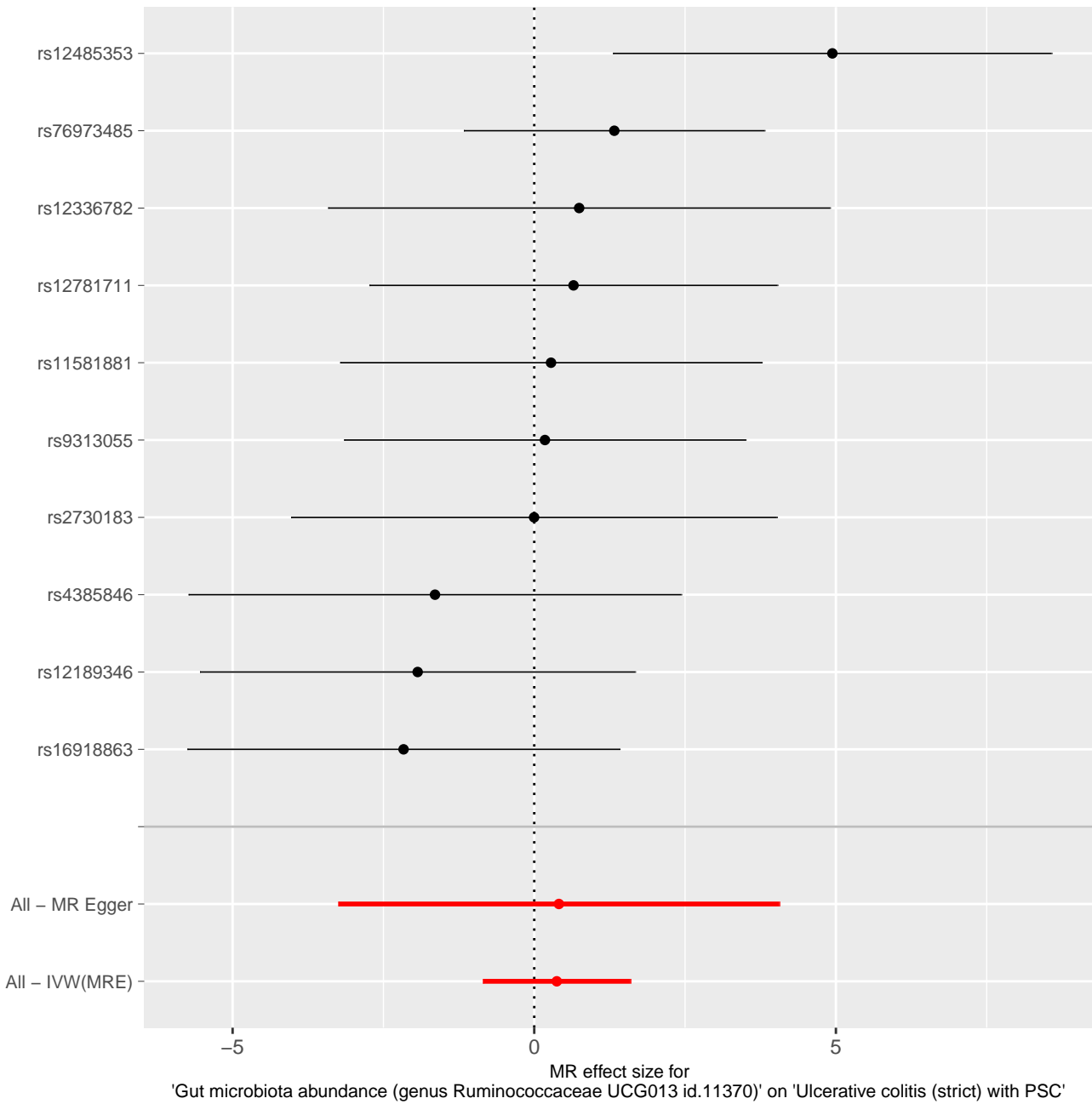


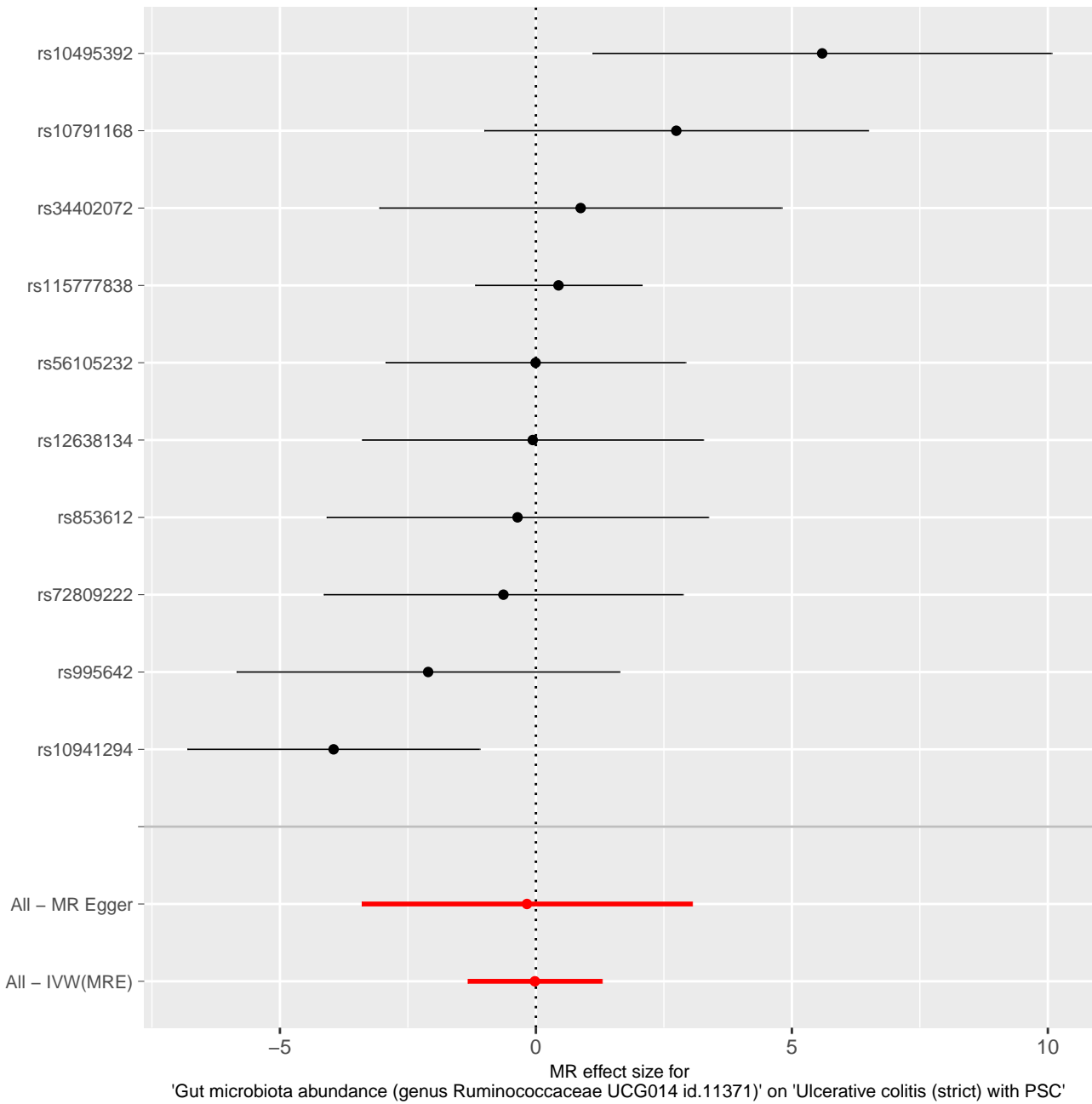


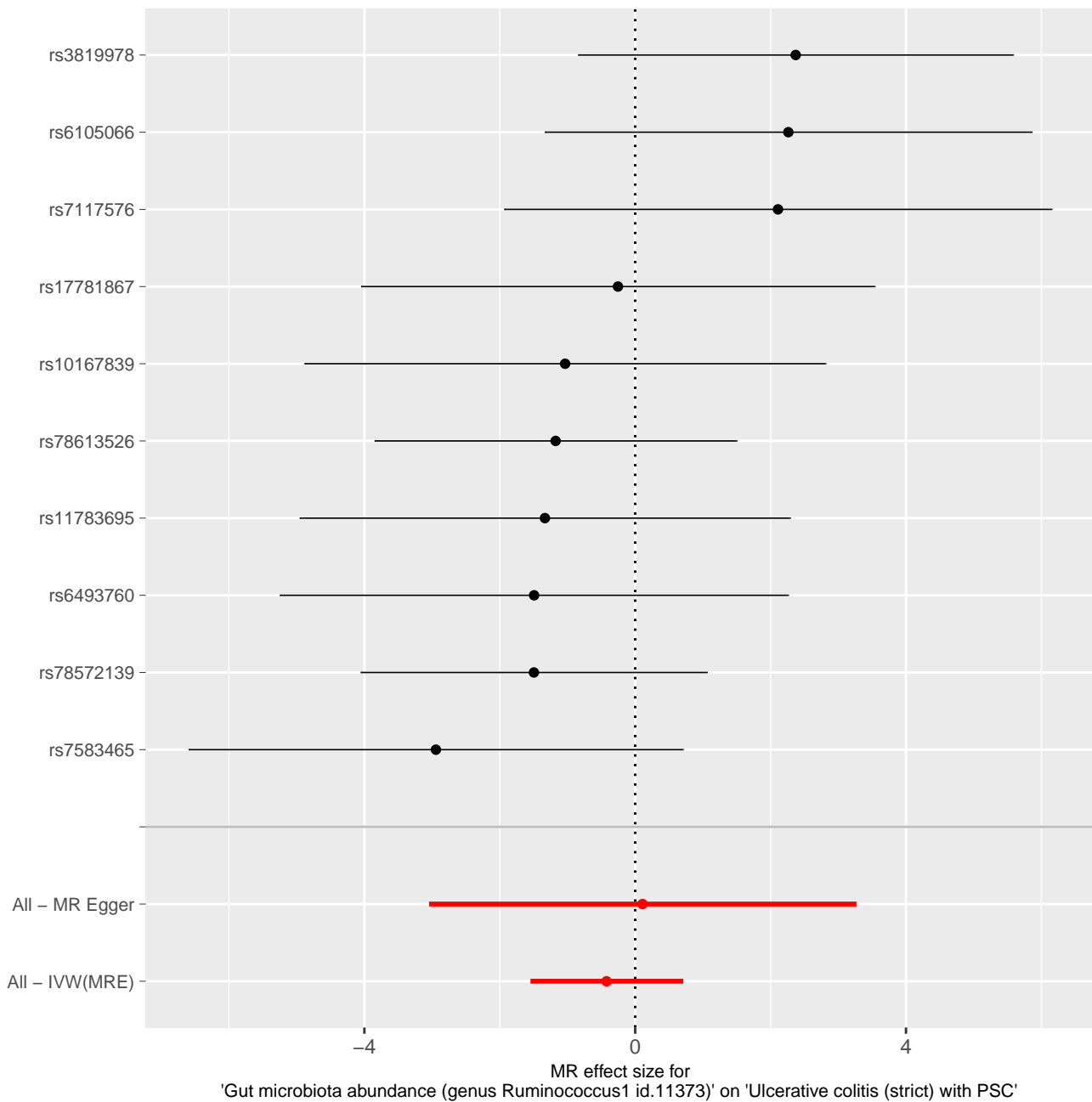


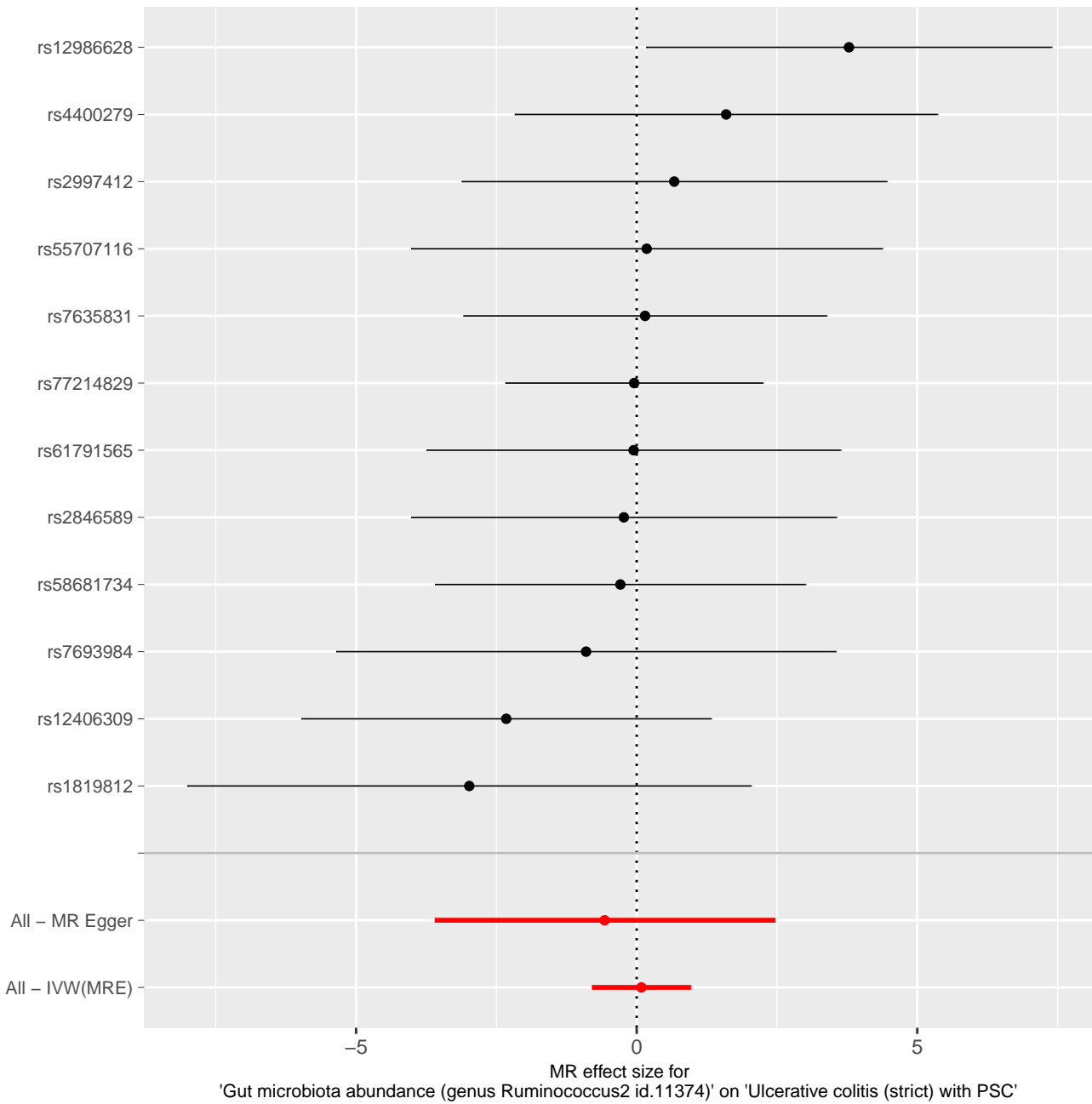


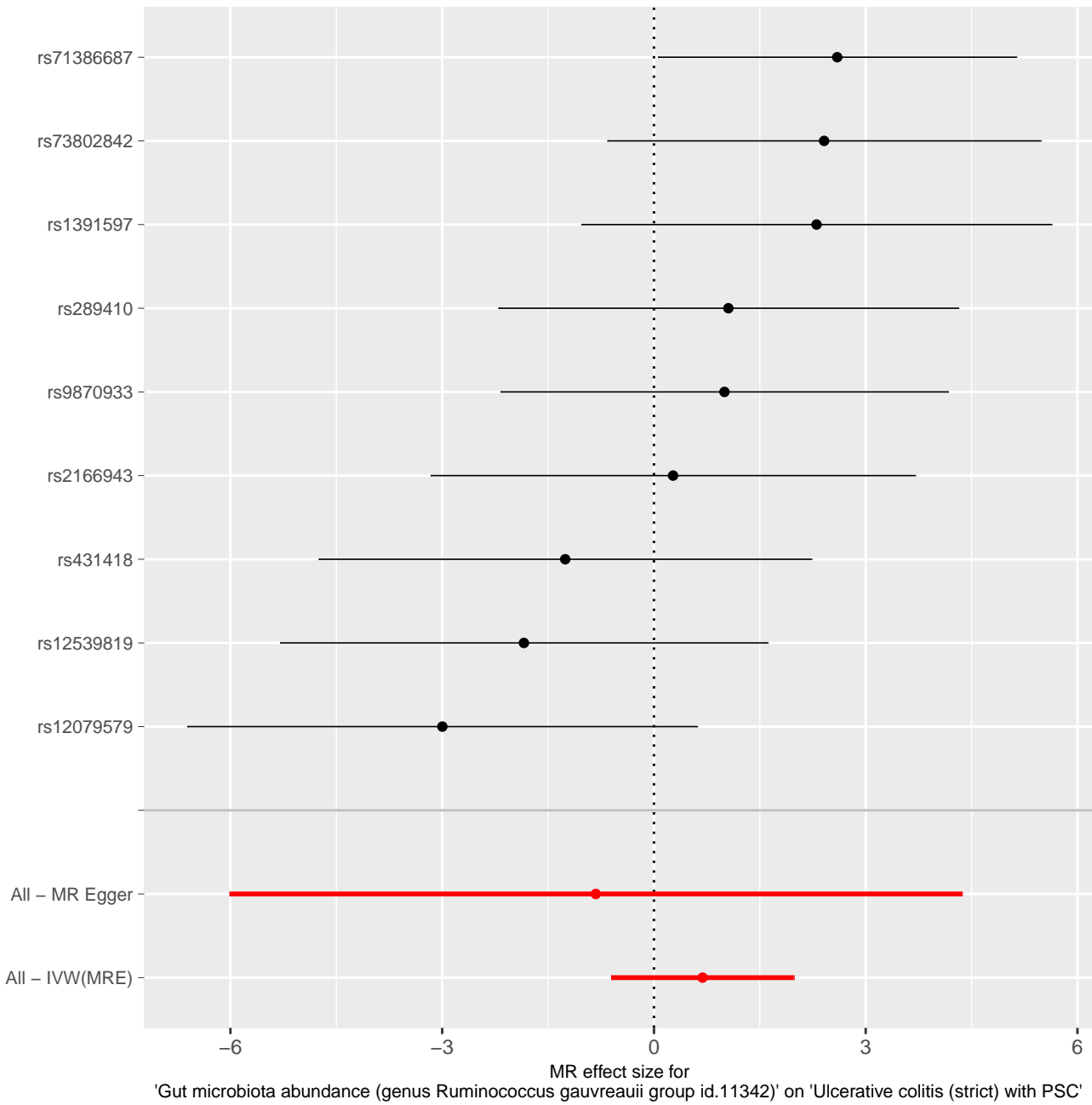
'Gut microbiota abundance (genus Ruminococcaceae UCG011 id.11368)' on 'Ulcerative colitis (strict) with PSC'



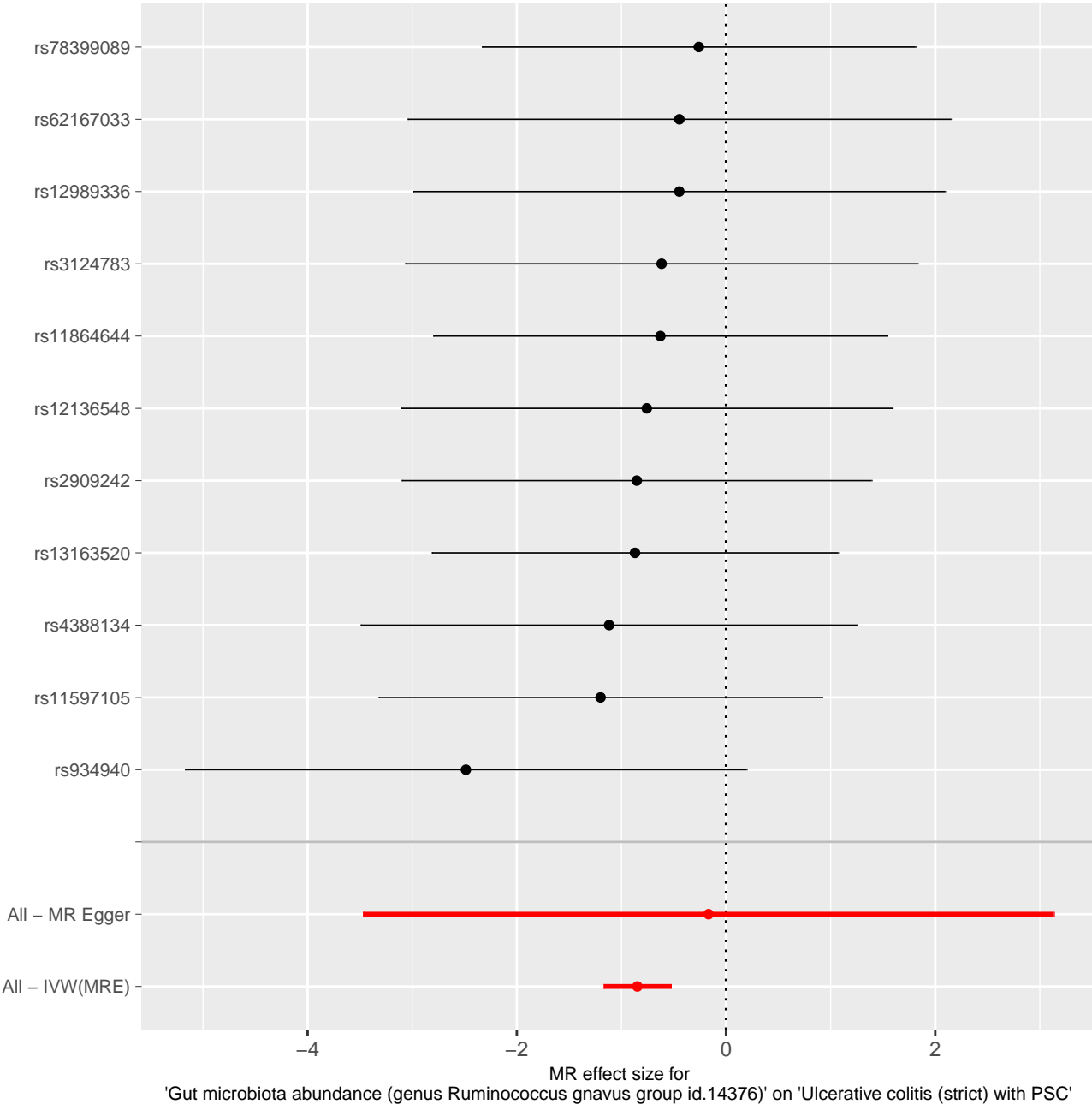


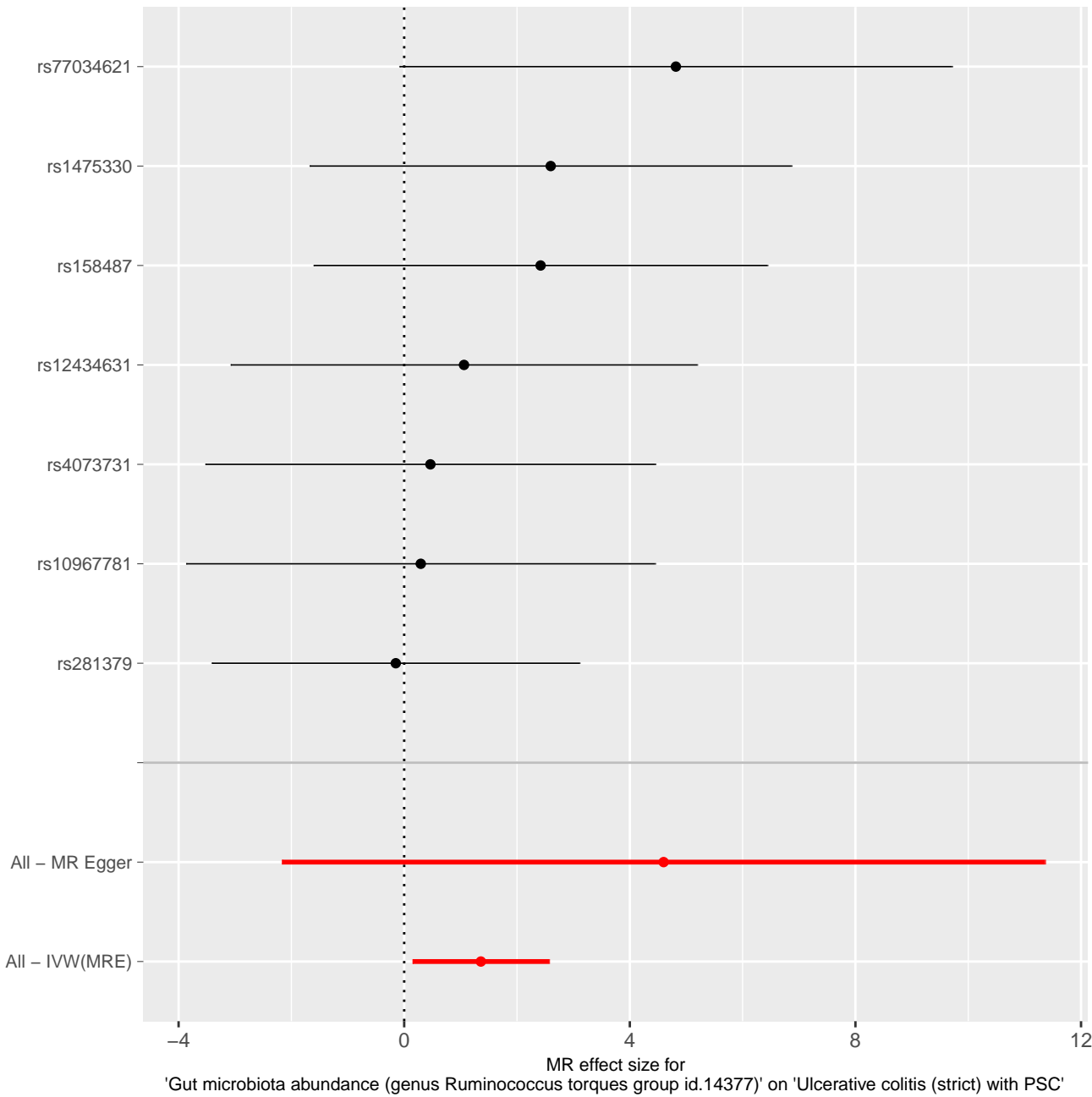




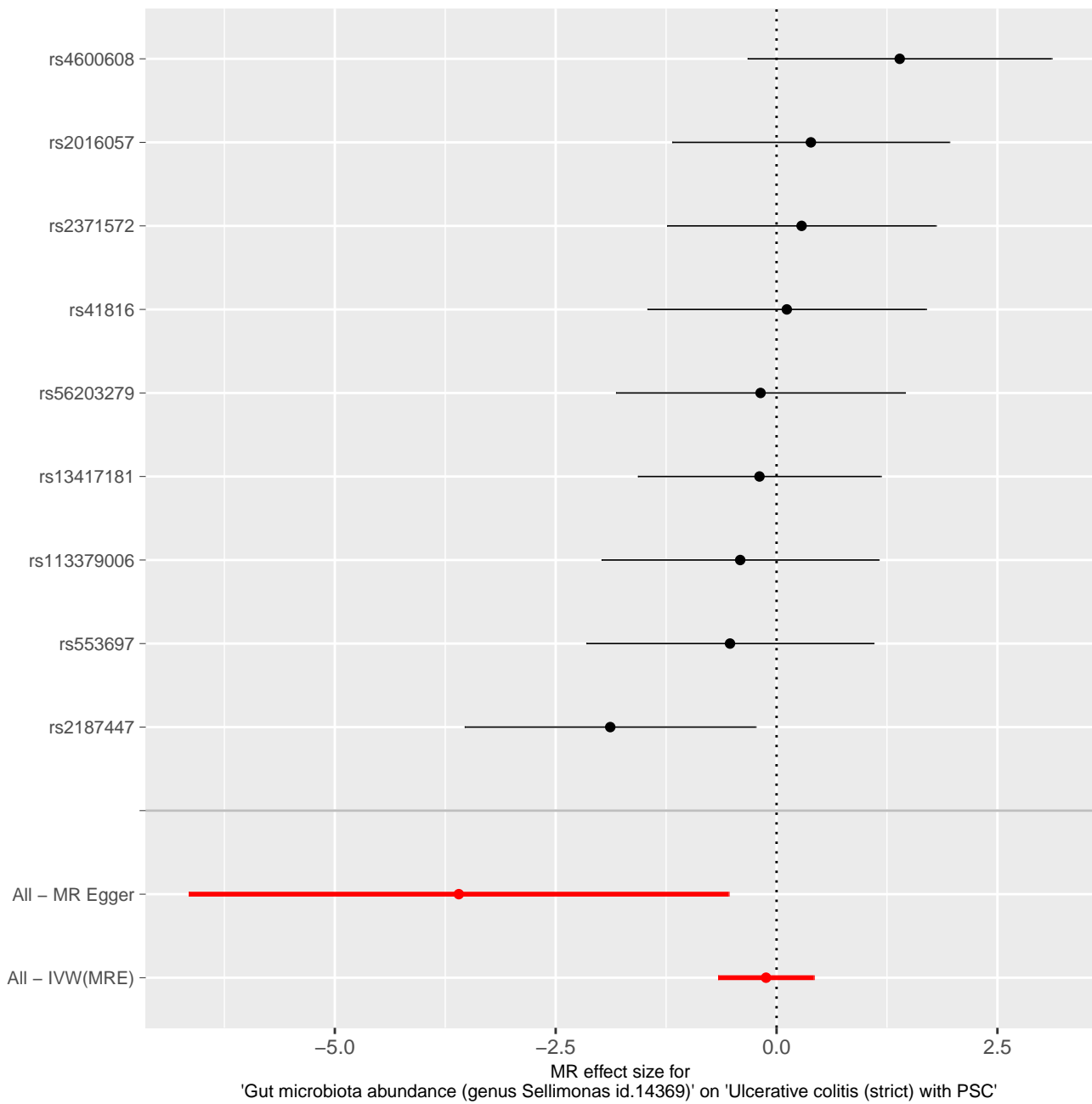


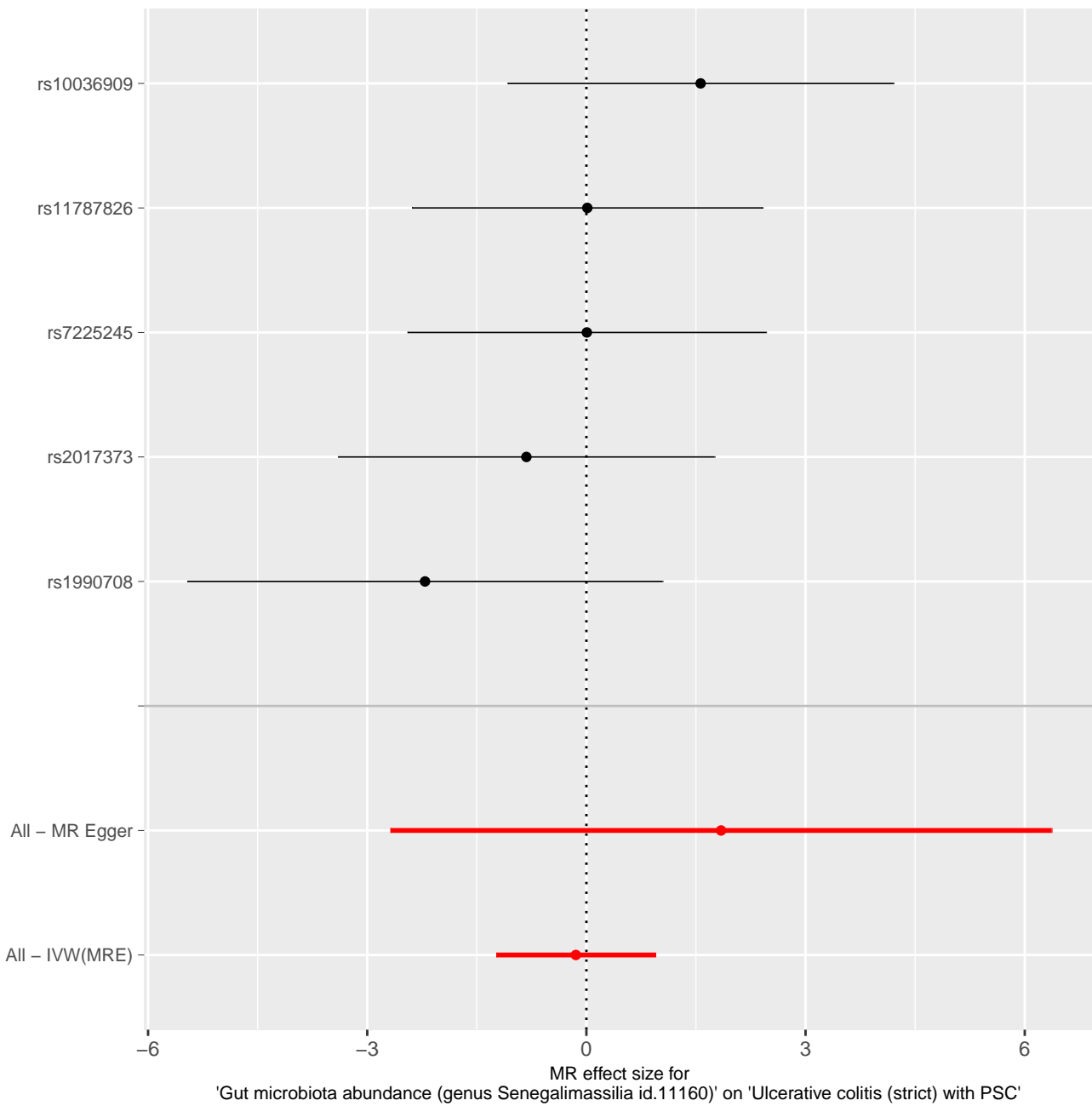
MR effect size for
'Gut microbiota abundance (genus Ruminococcus gauvreauii group id.11342)' on 'Ulcerative colitis (strict) with PSC'



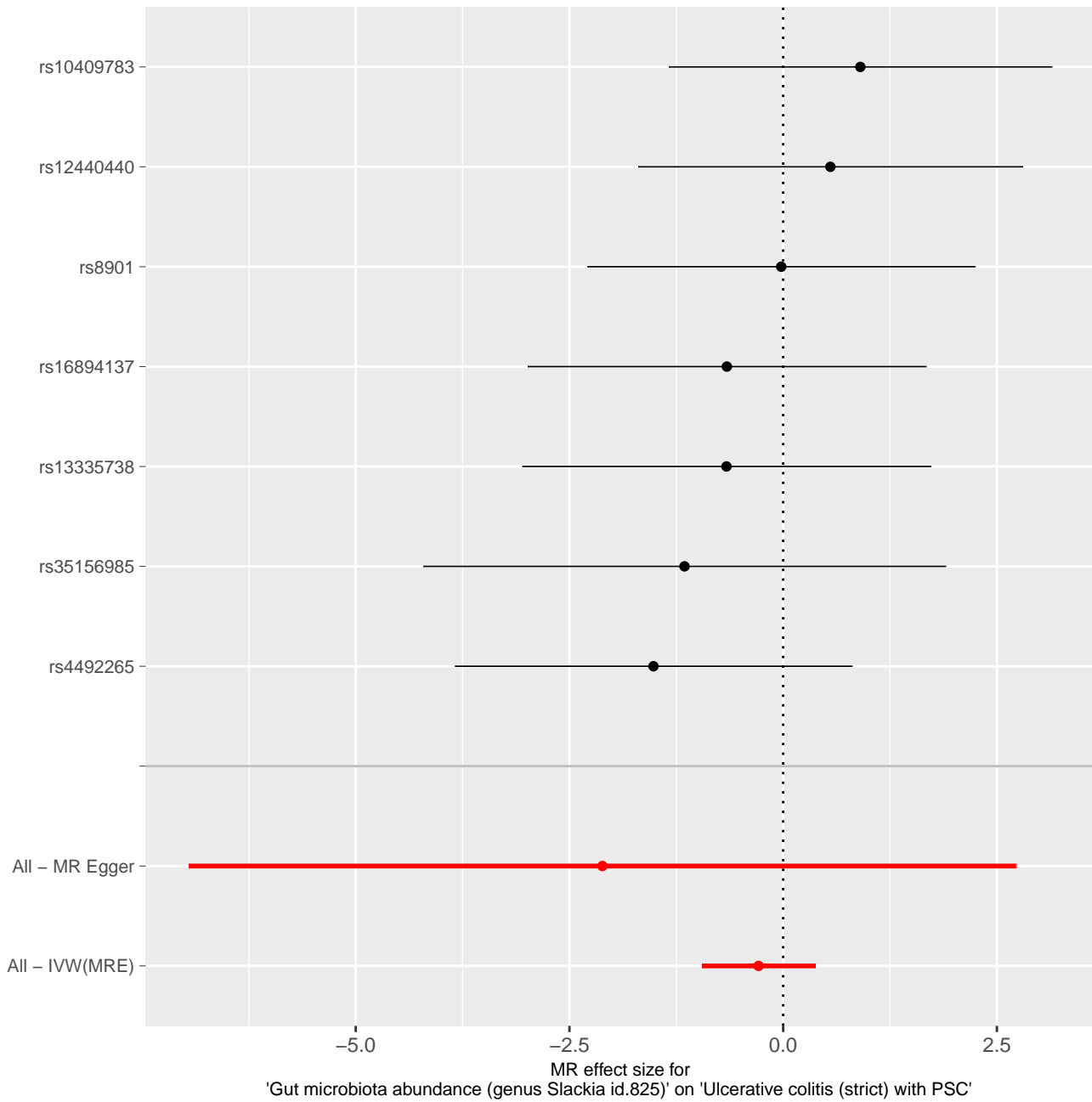


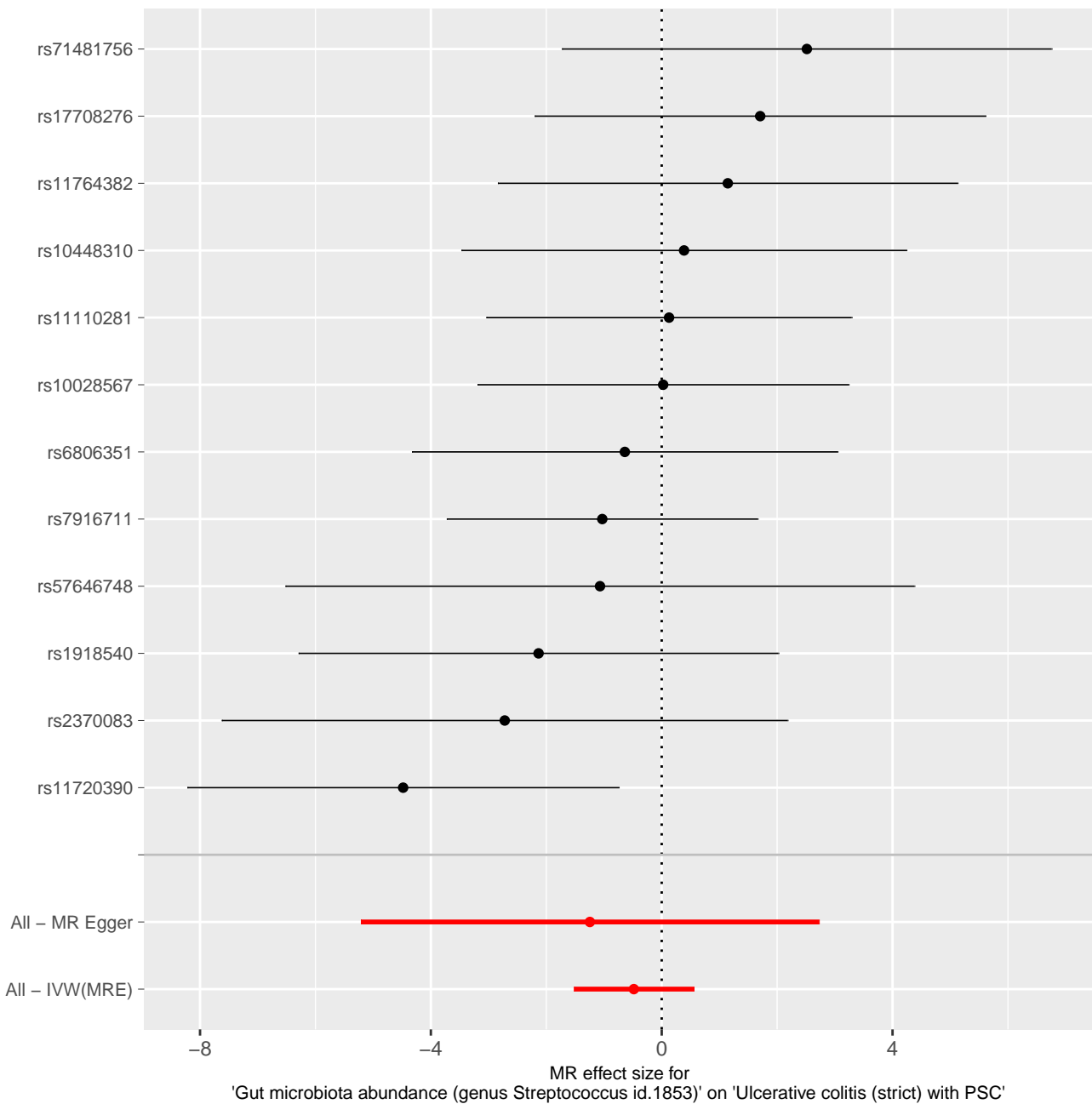
MR effect size for
'Gut microbiota abundance (genus Ruminococcus torques group id.14377)' on 'Ulcerative colitis (strict) with PSC'

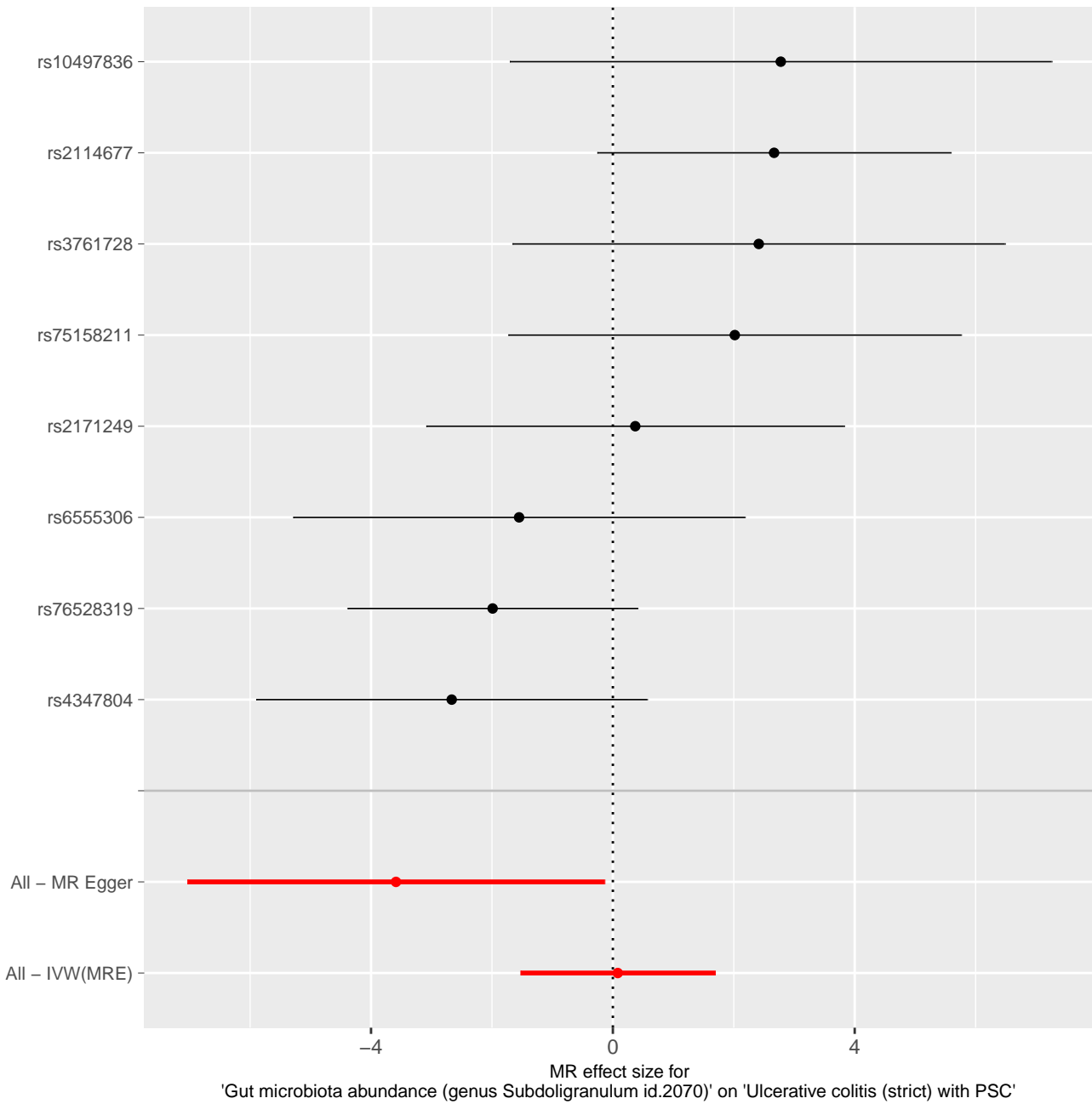


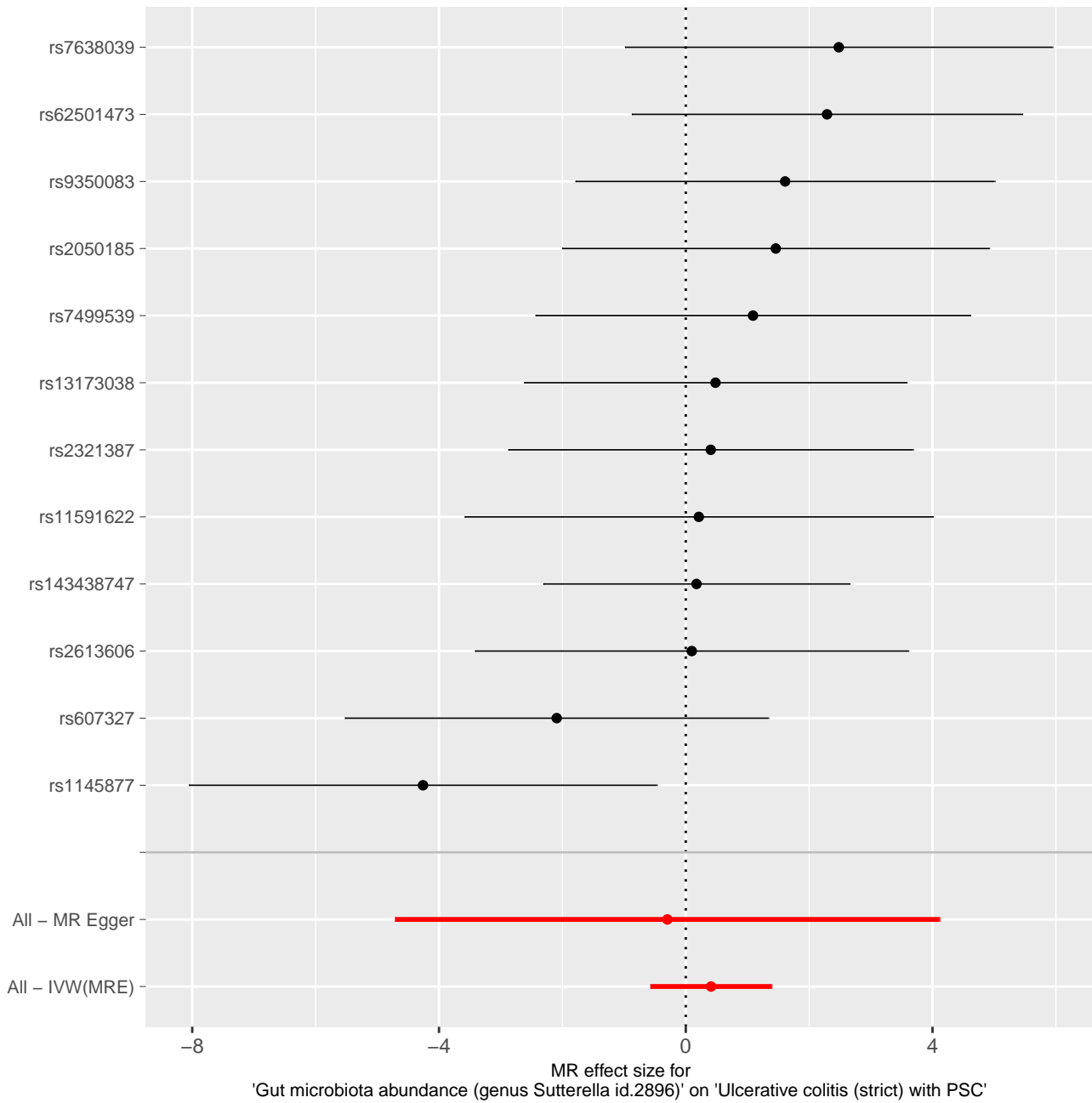


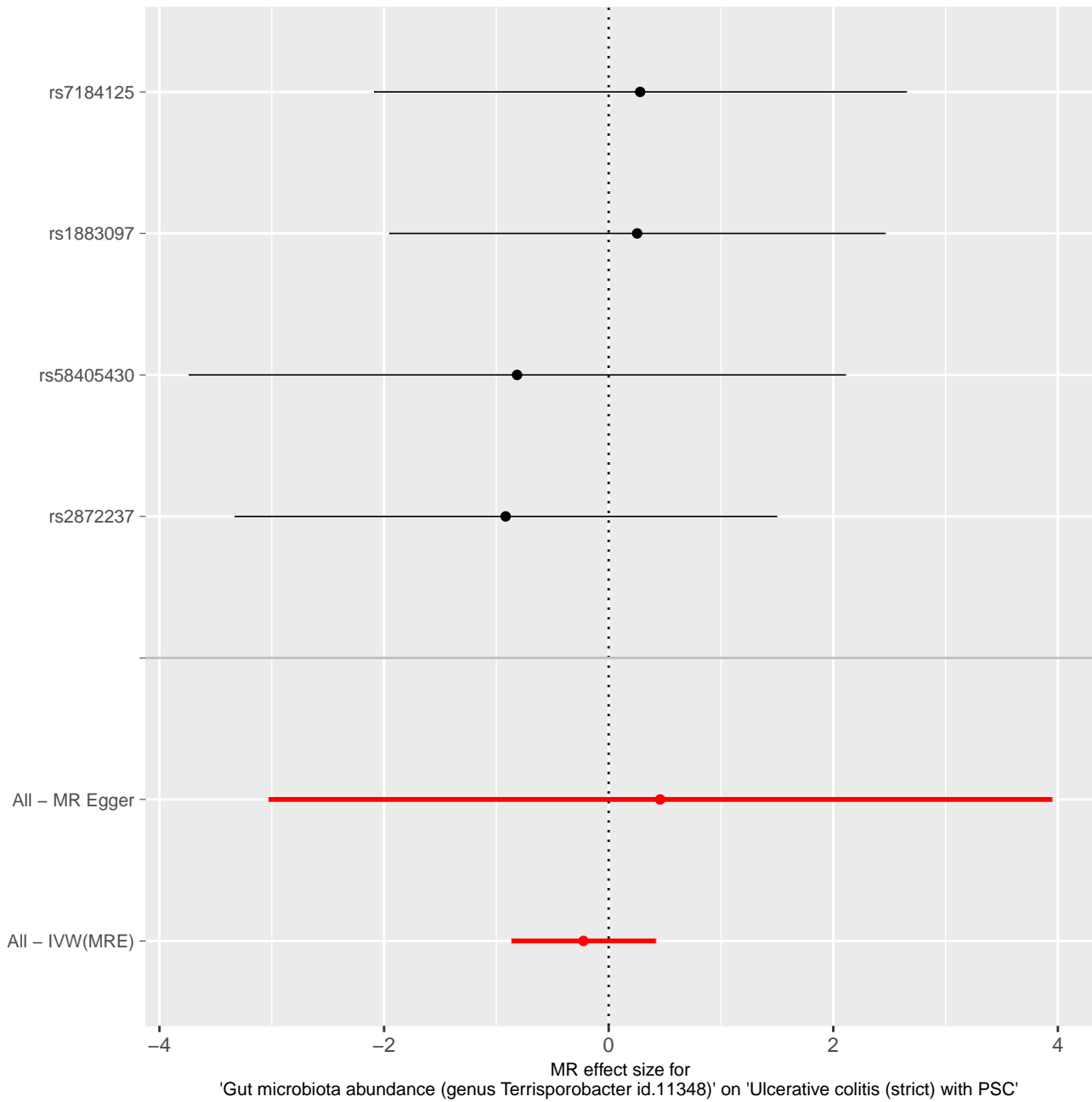
Batch 743 : Gut microbiota abundance (genus Slackia id.825) on Ulcerative colitis (strict) with PSC

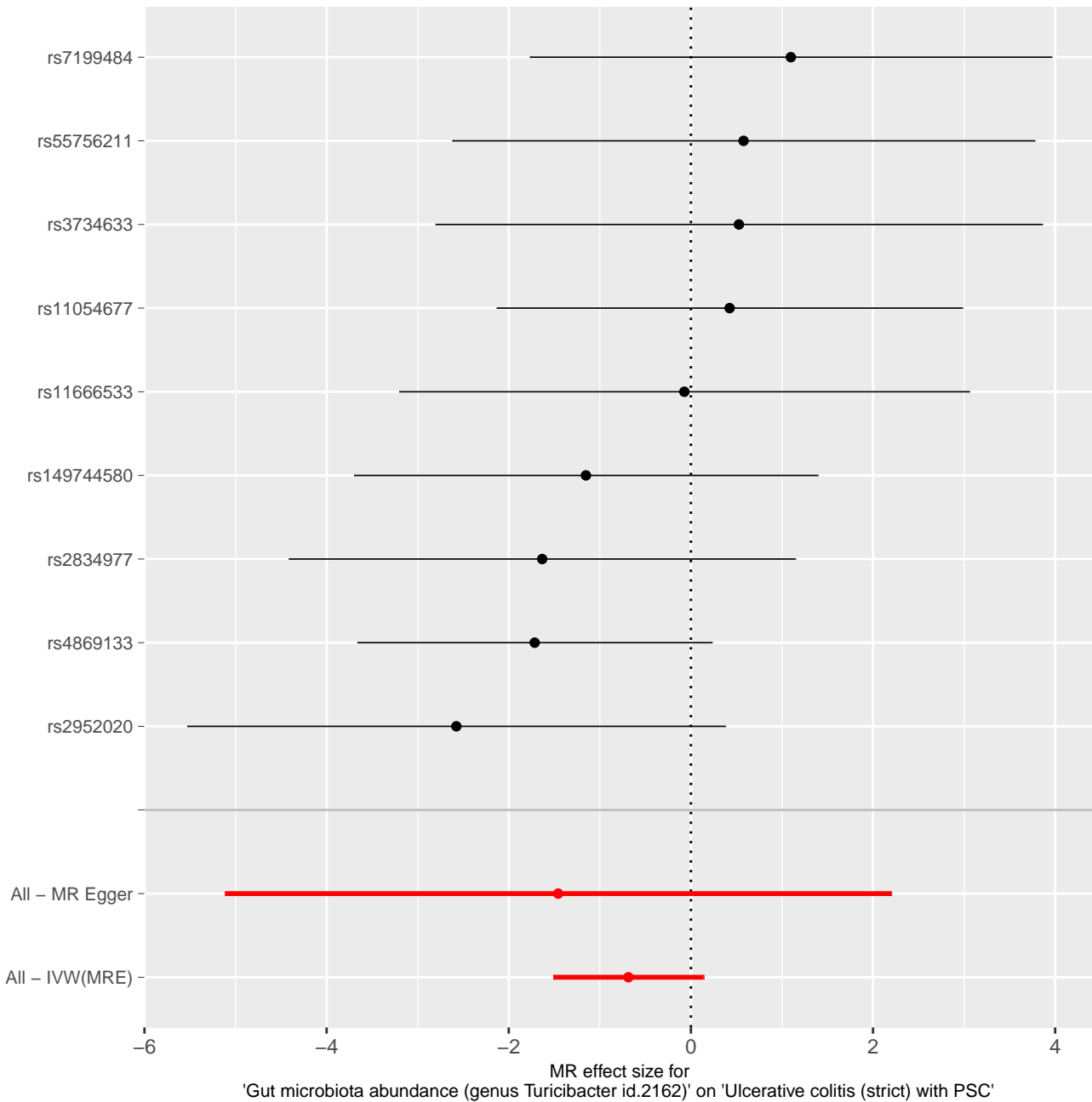


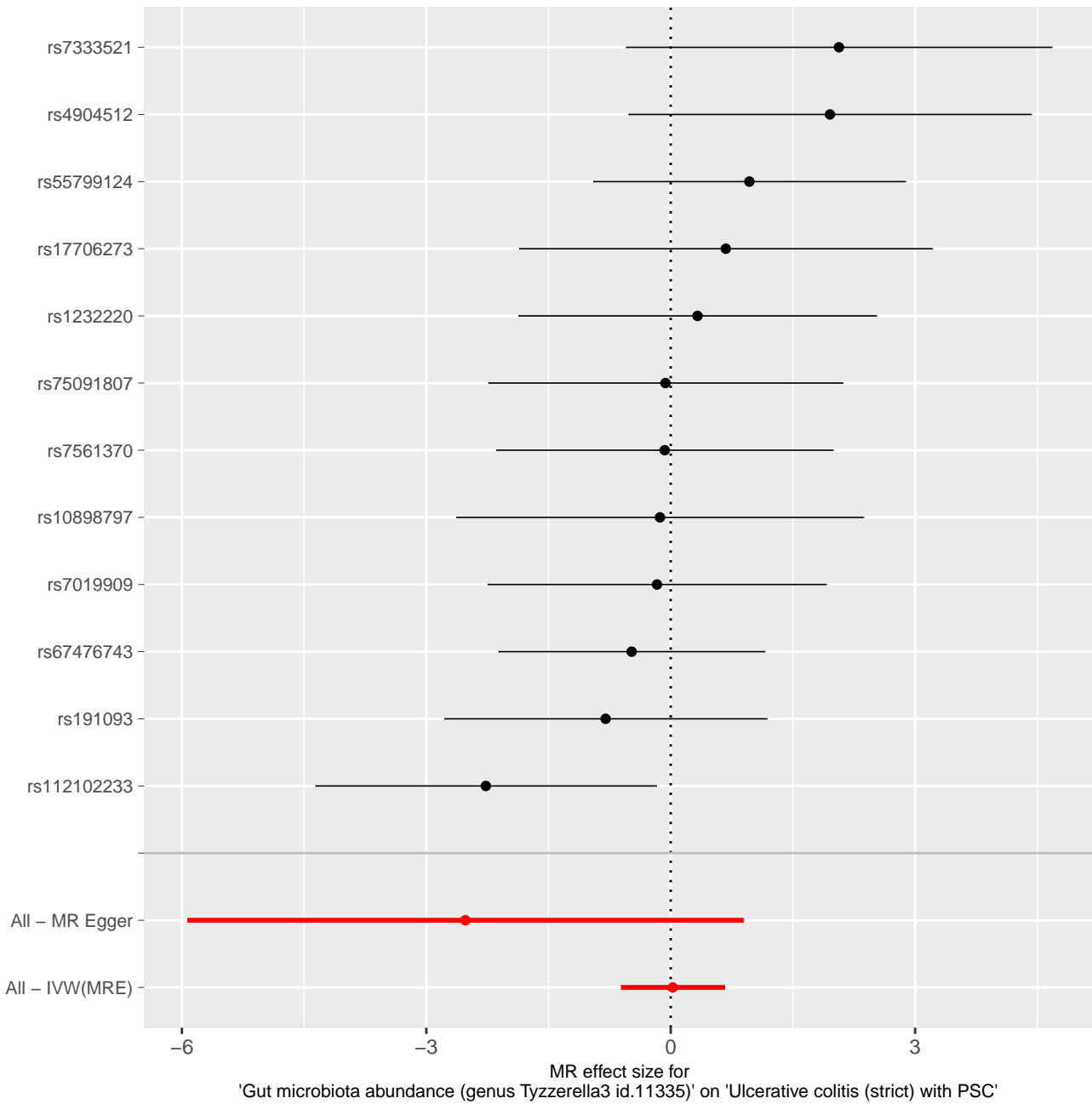


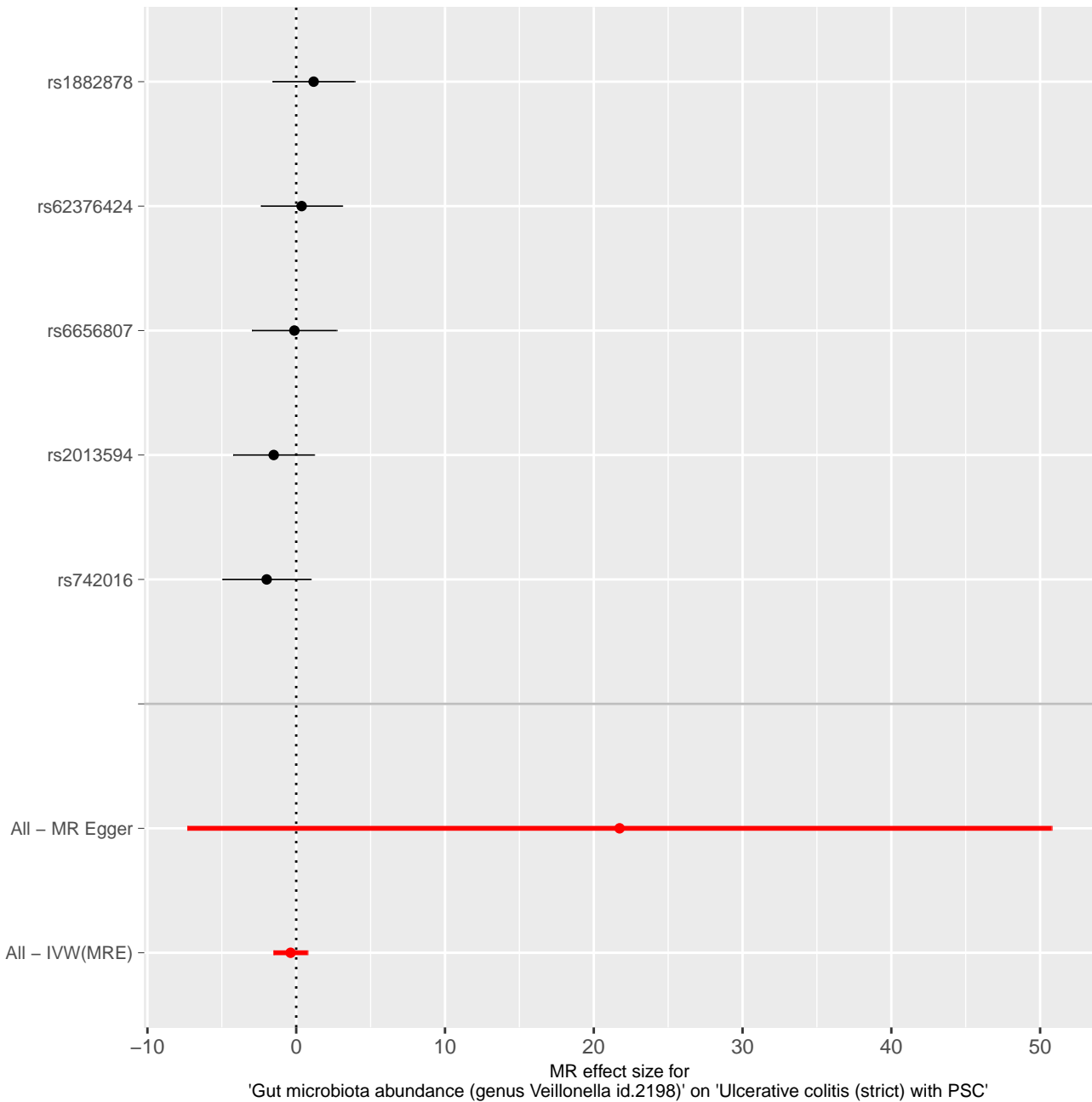


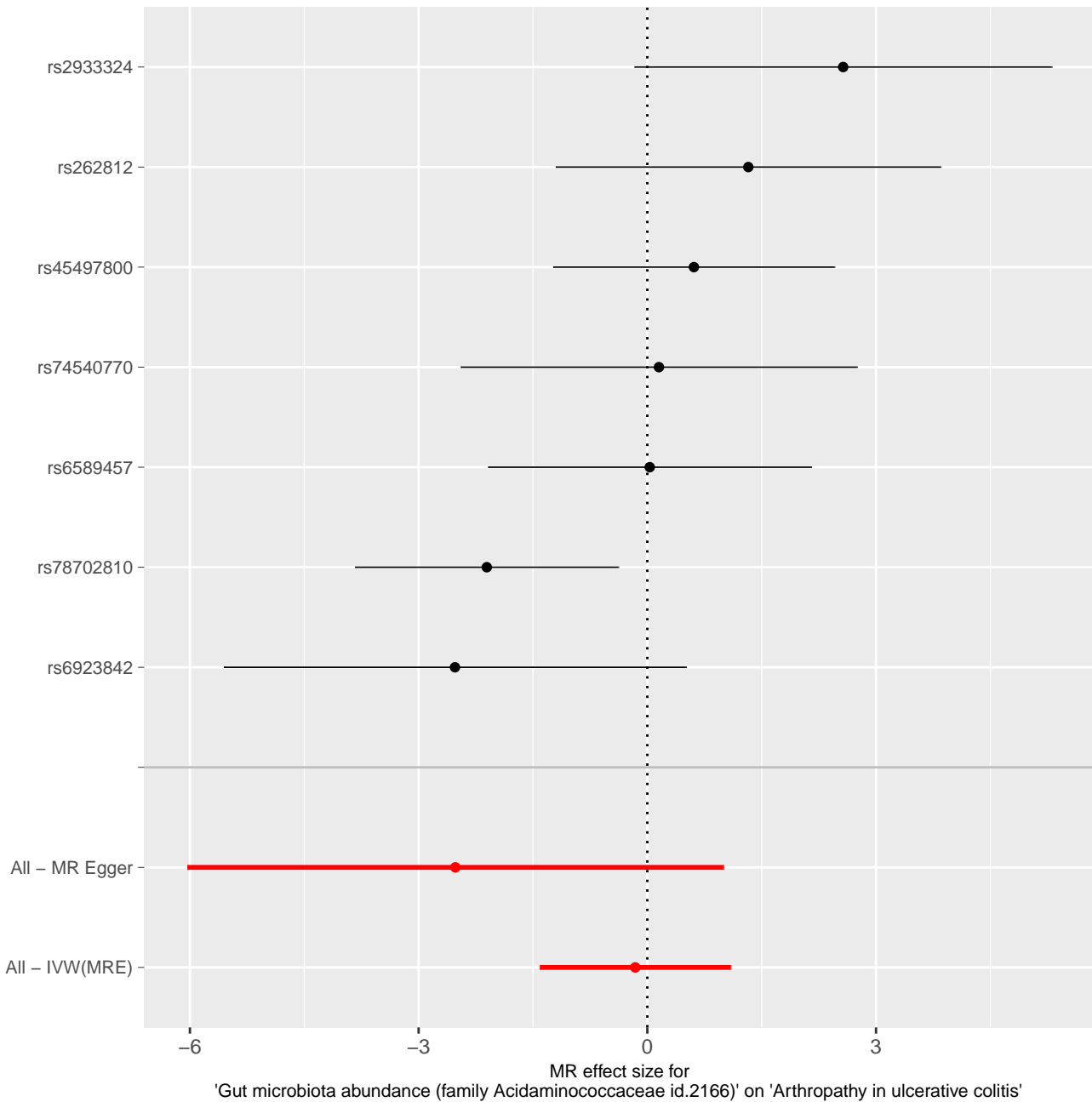


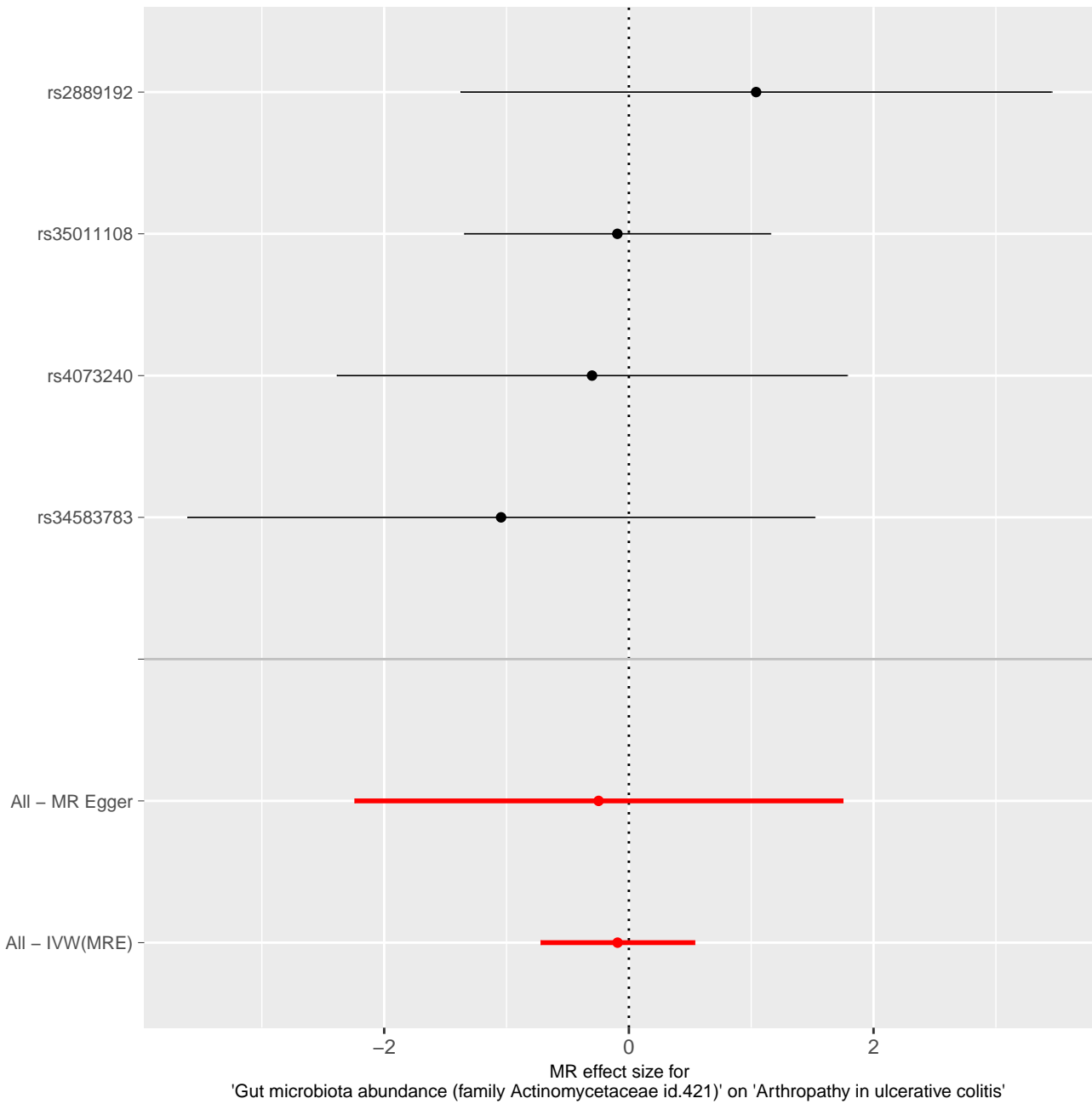


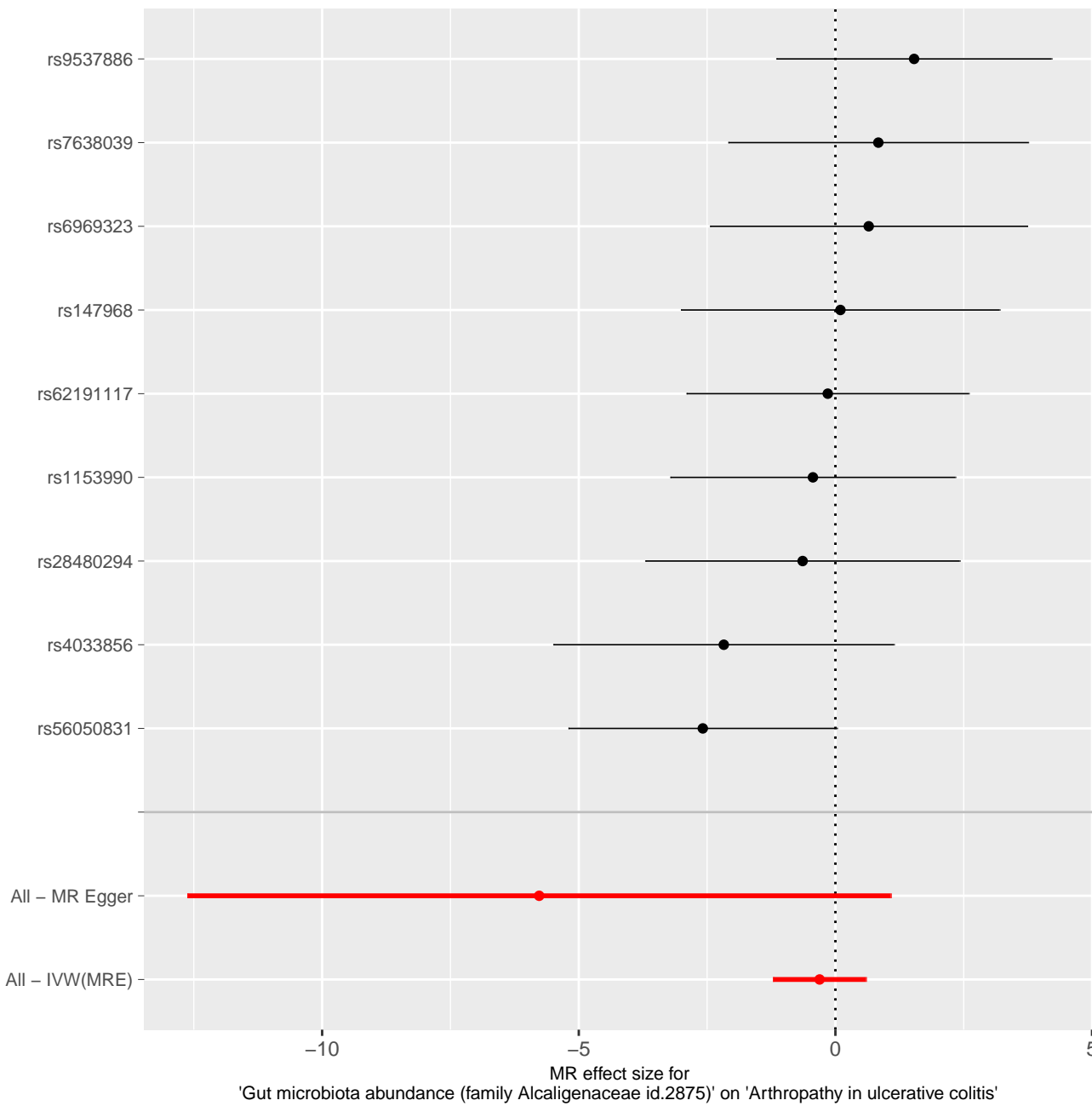


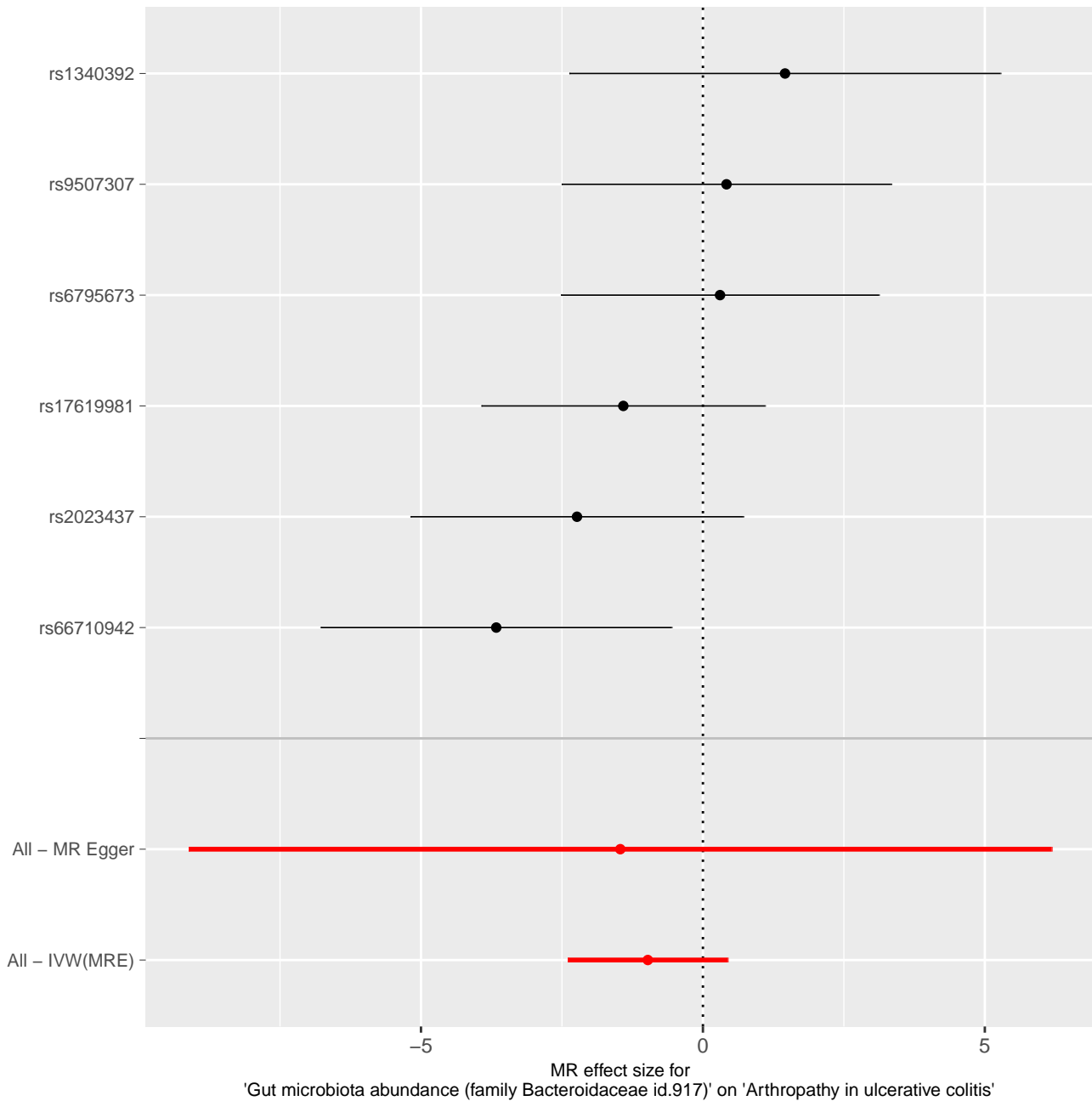


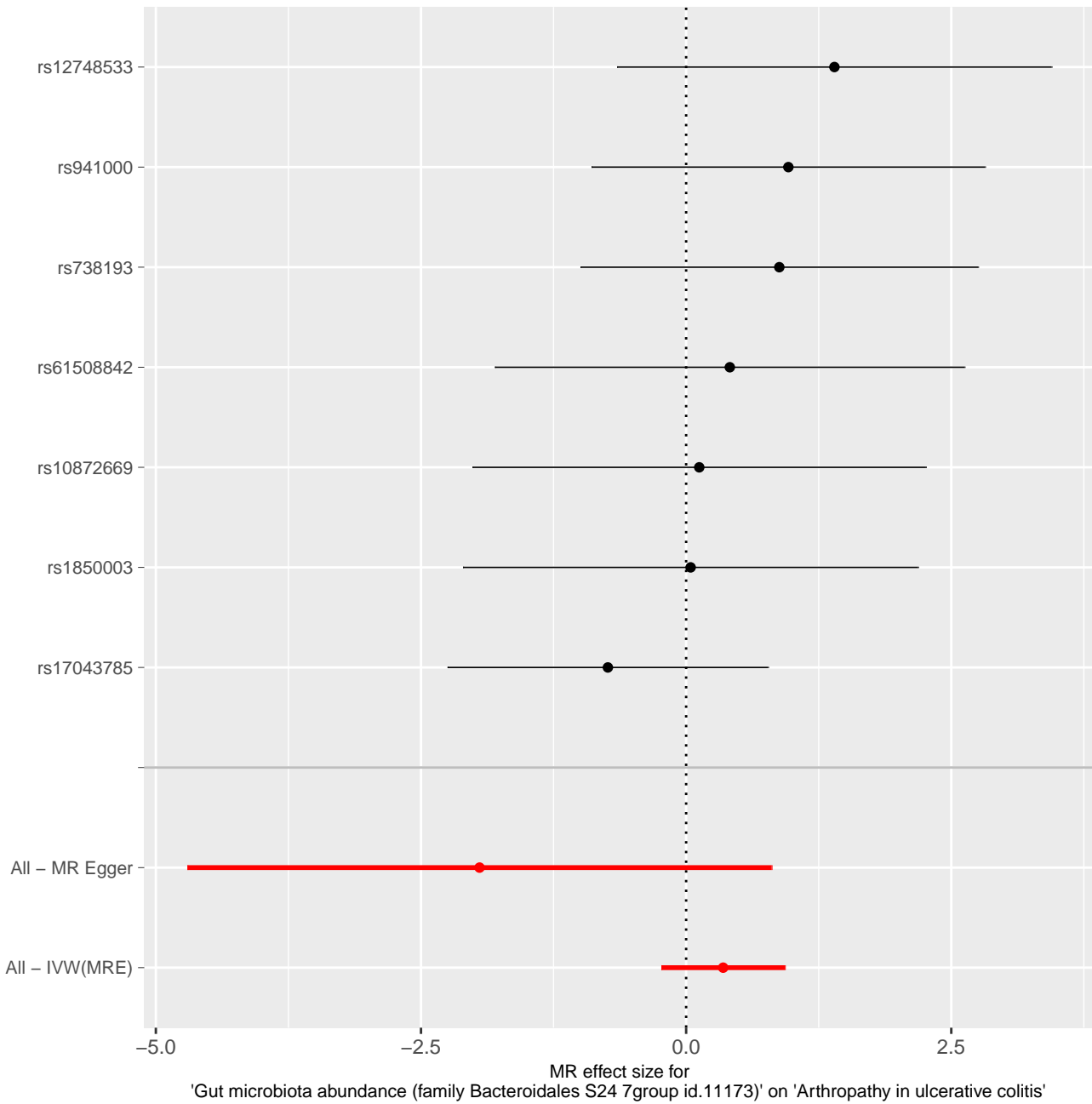




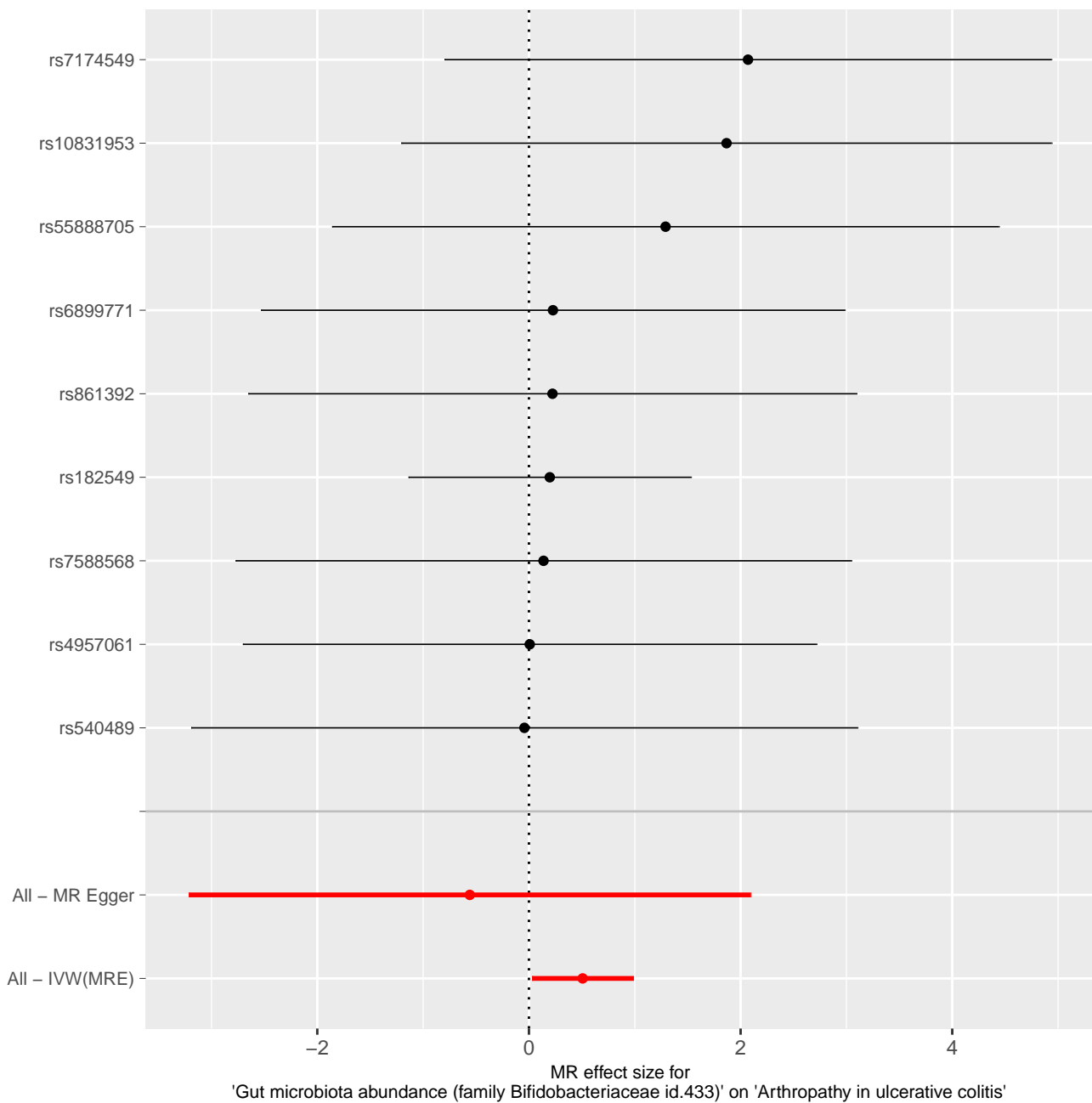




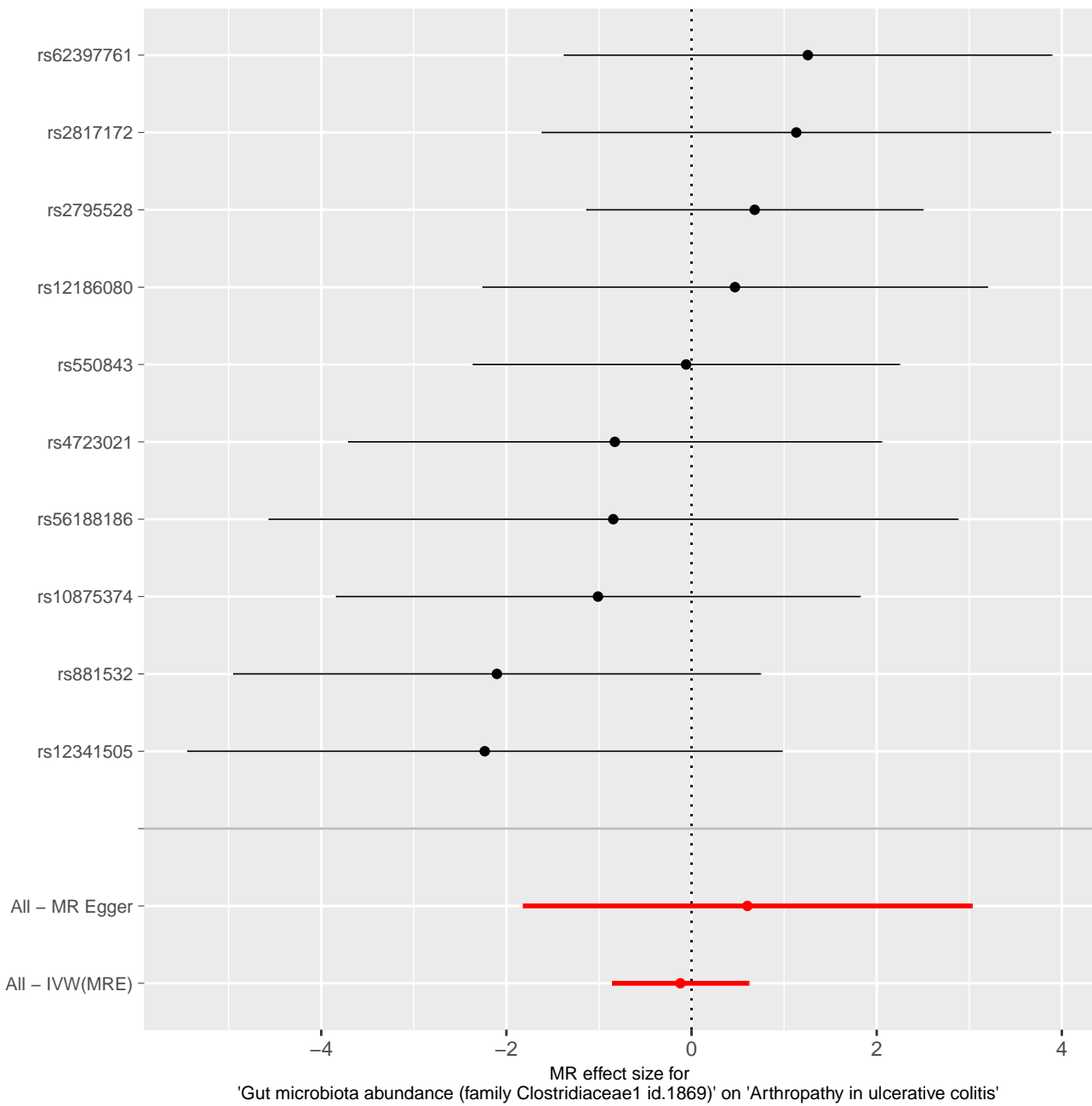


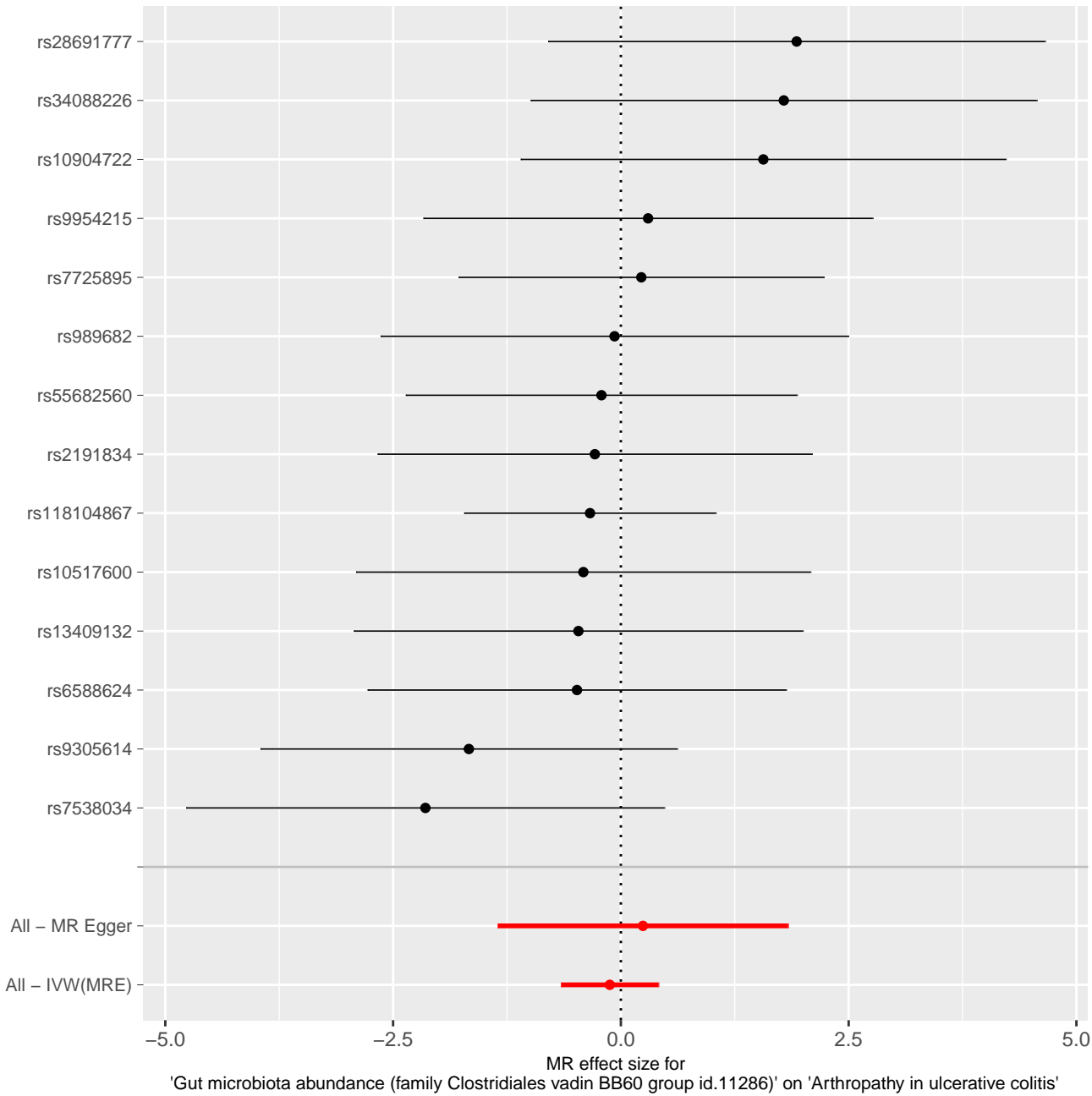


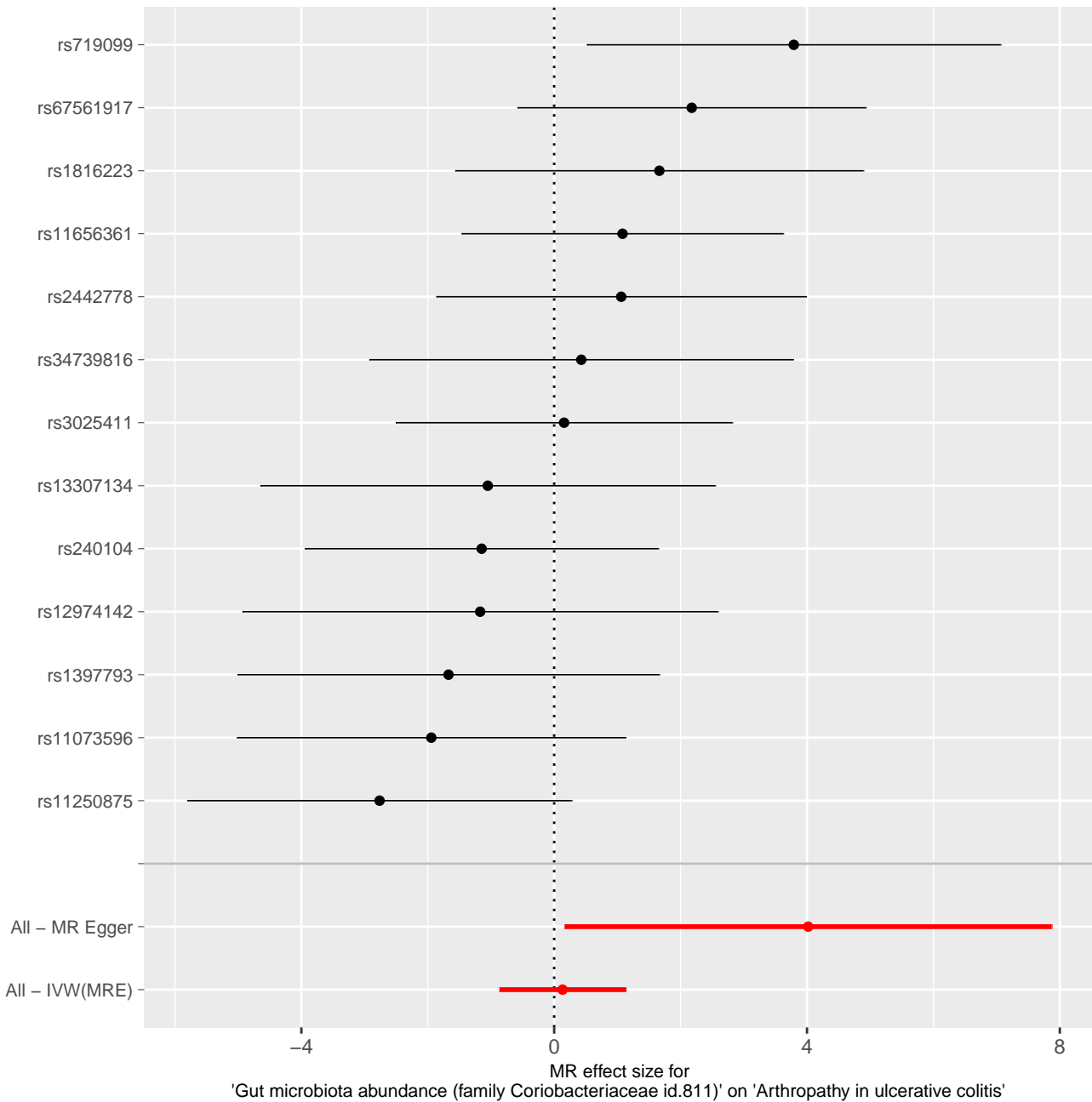
MR effect size for
'Gut microbiota abundance (family Bacteroidales S24 7group id.11173)' on 'Arthropathy in ulcerative colitis'

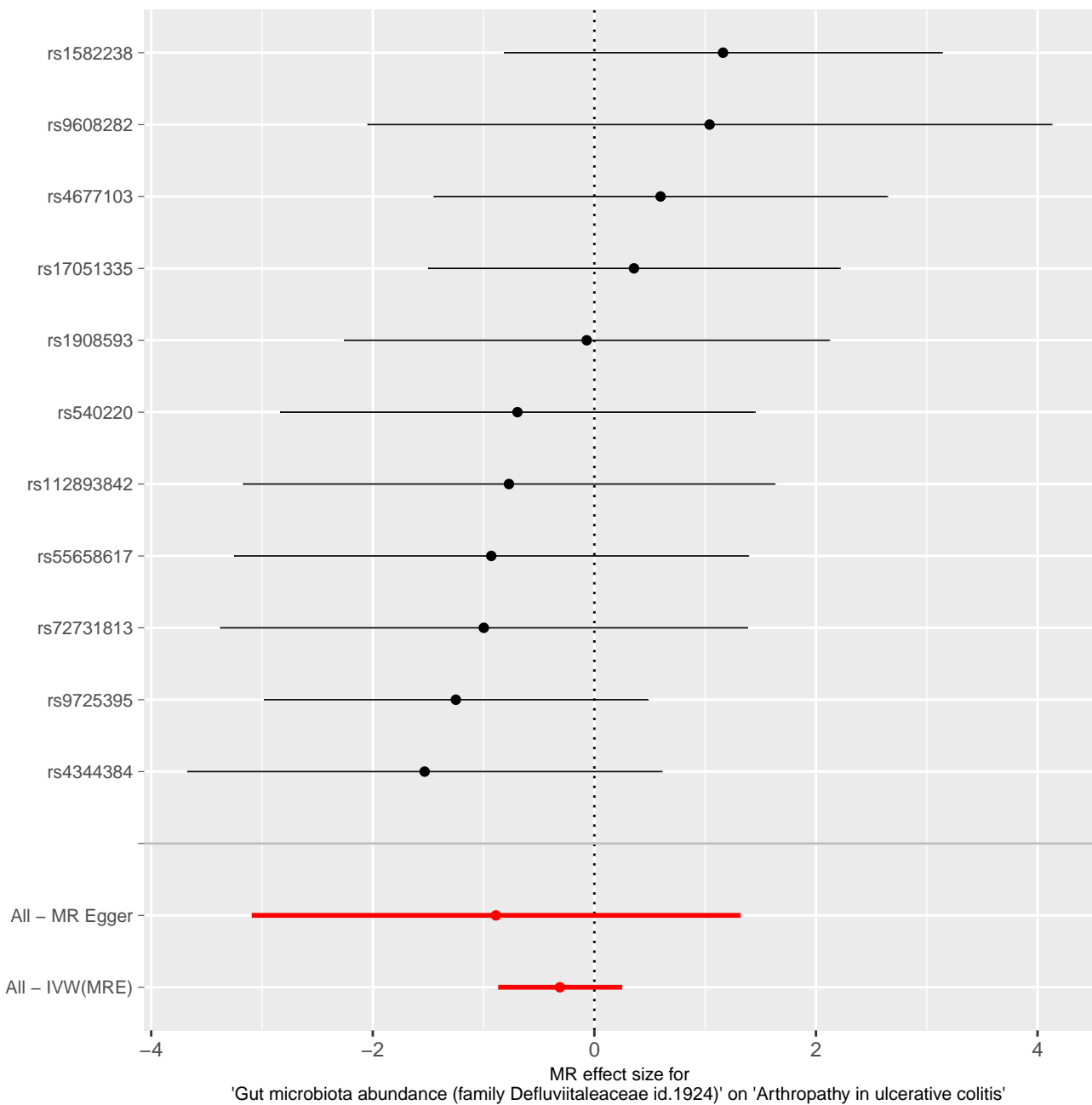


Batch 758 : Gut microbiota abundance (family Clostridiaceae1 id.1869) on Arthropathy in ulcerative colitis

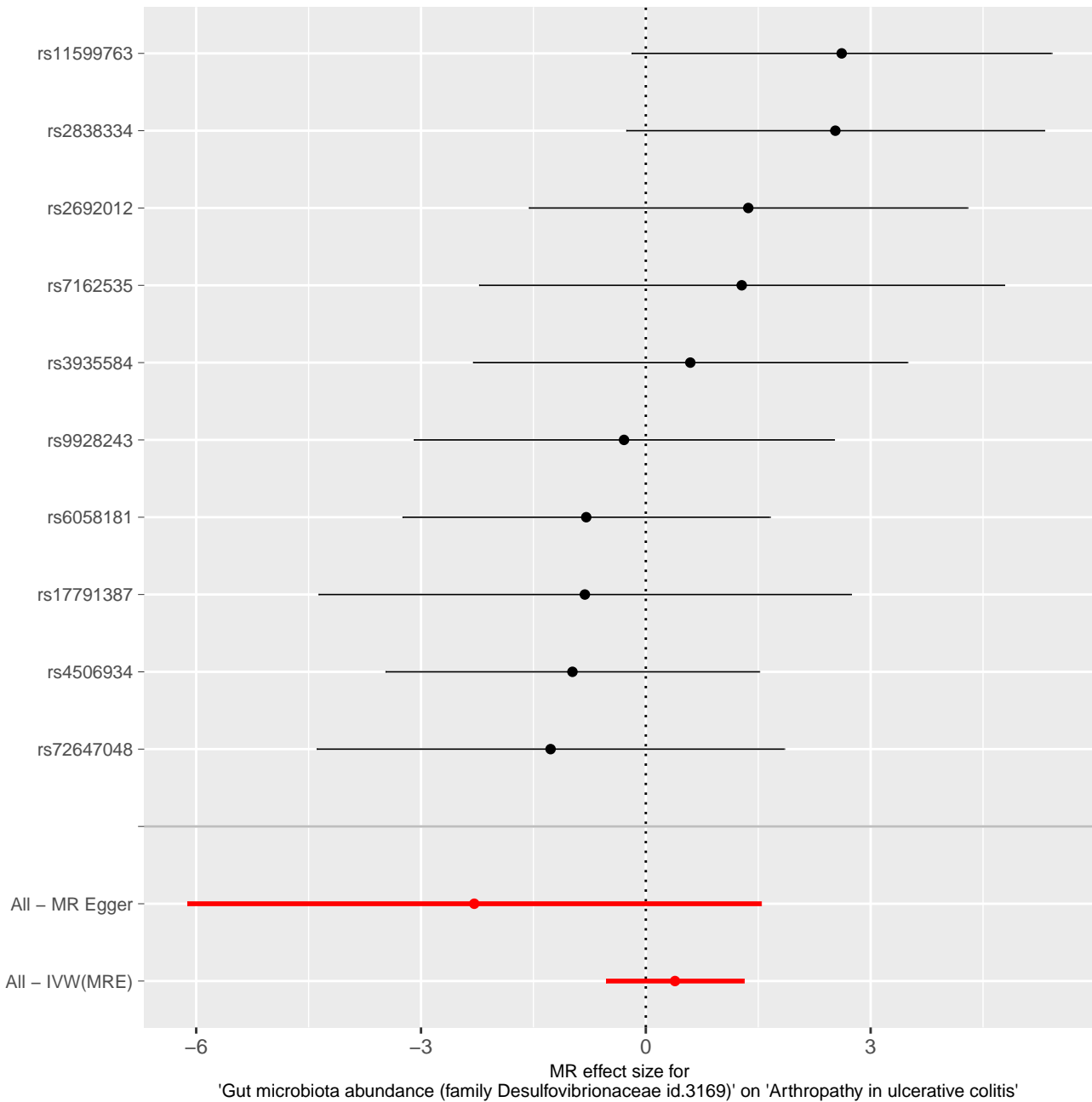


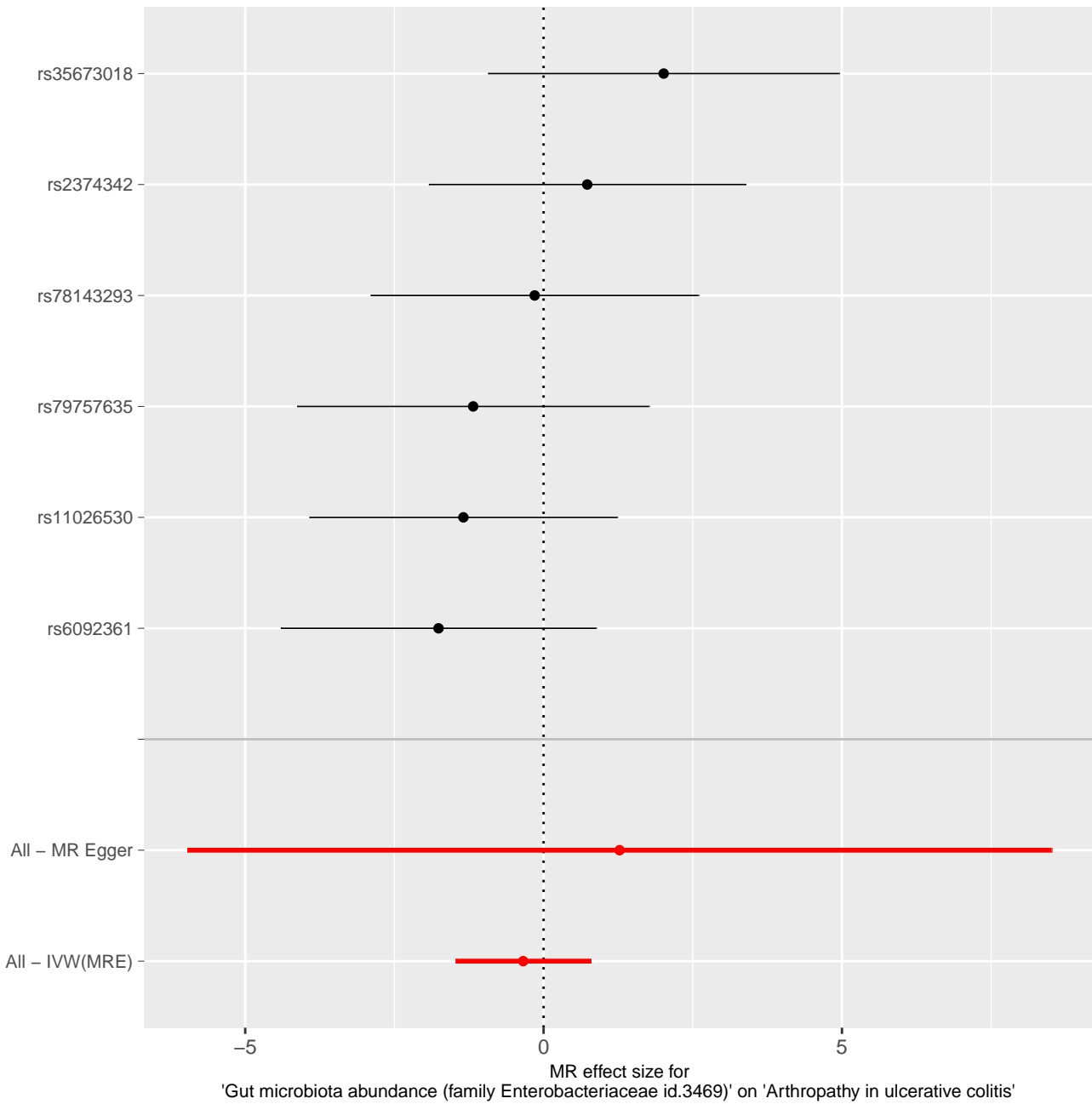




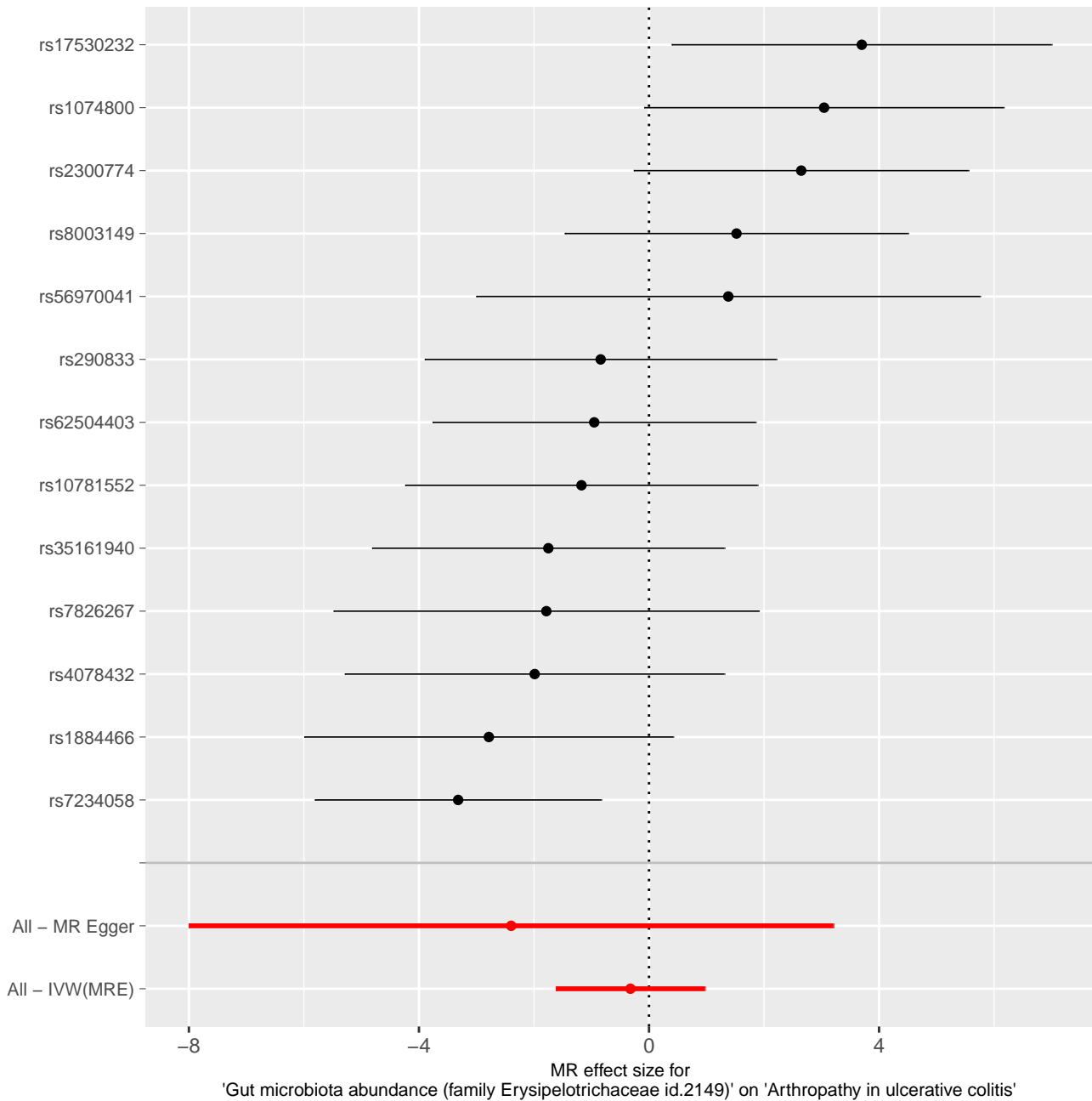


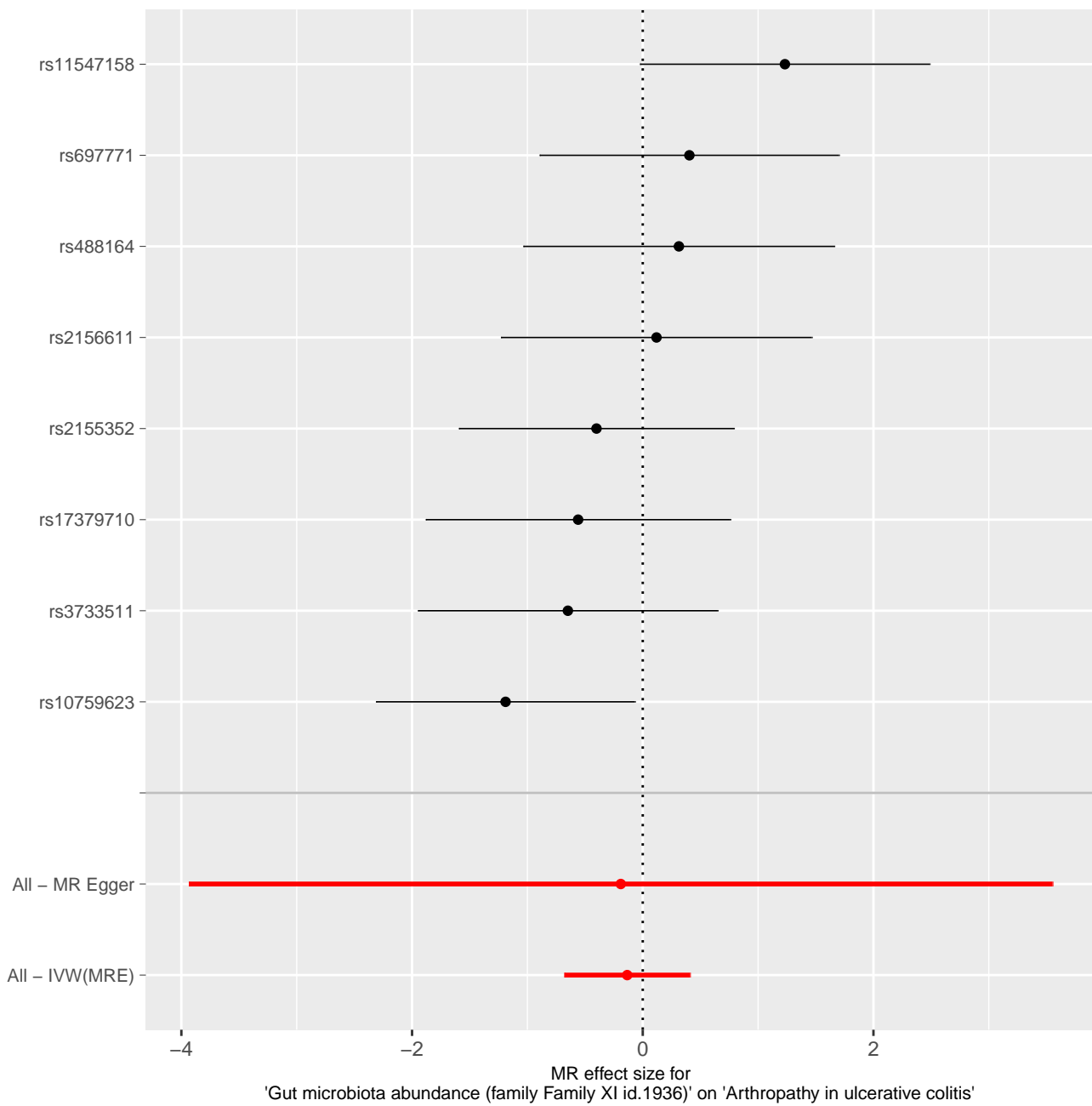
Batch 762 : Gut microbiota abundance (family Desulfovibrionaceae id.3169) on Arthropathy in ulcerative colitis

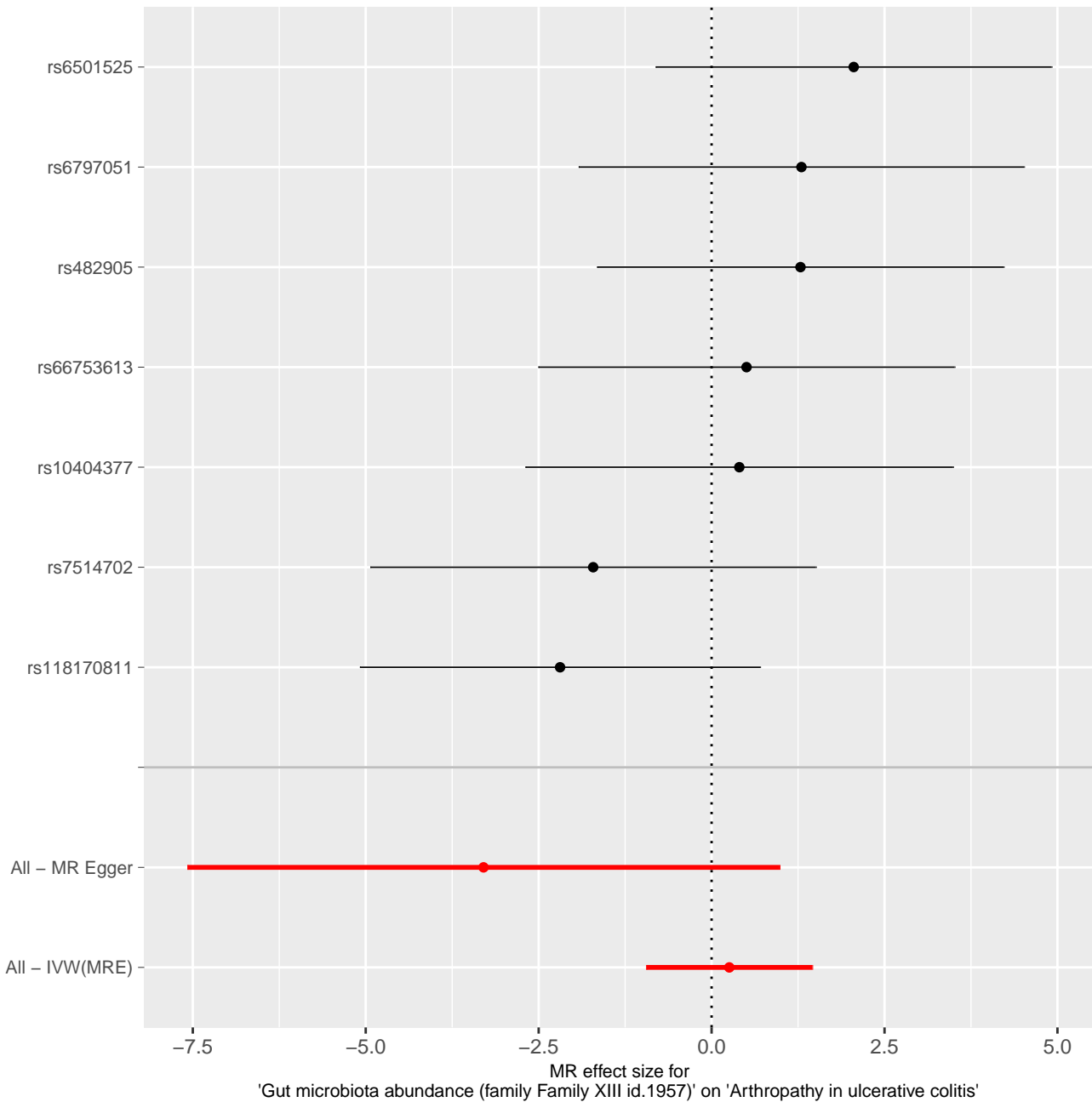




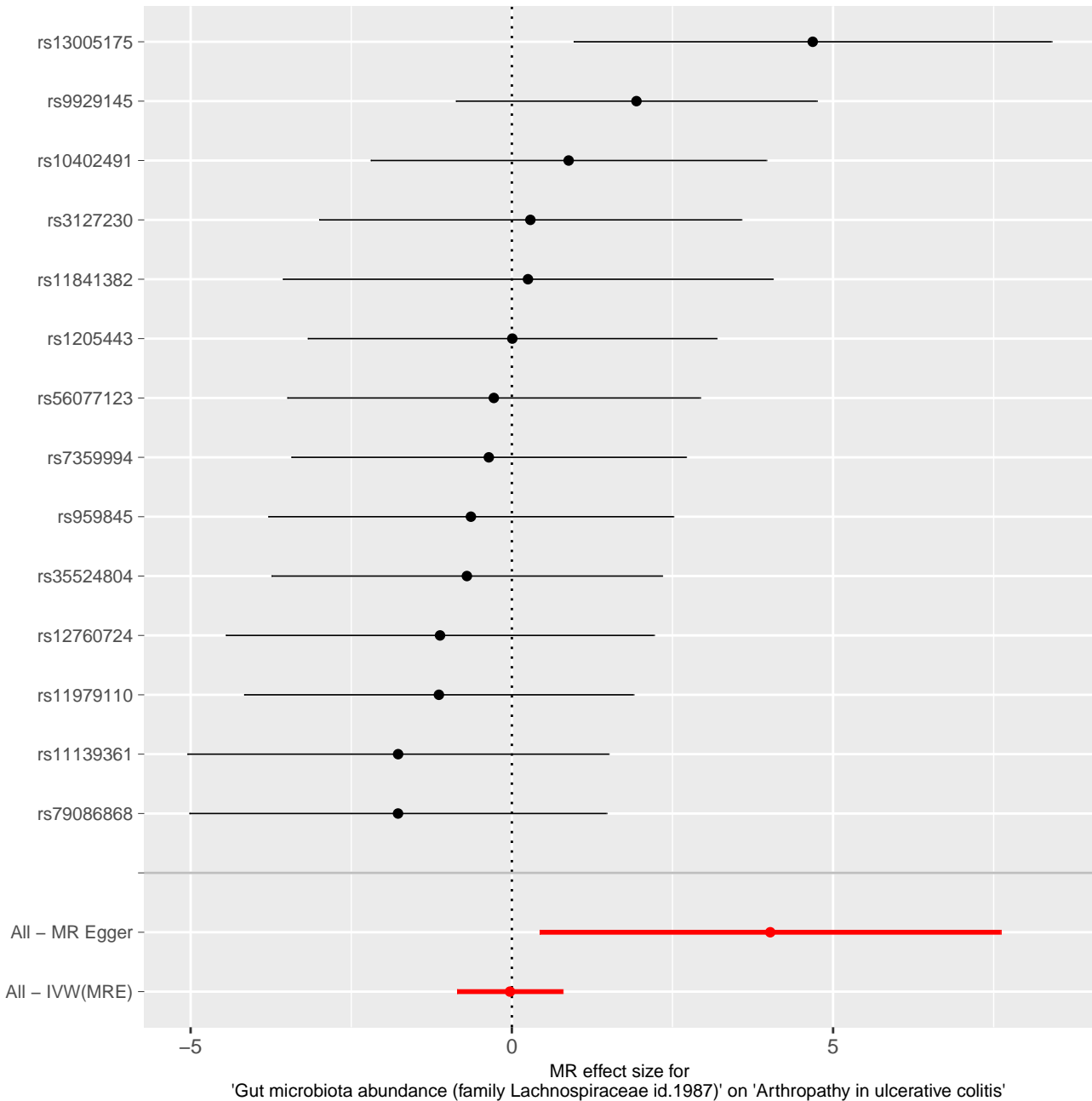
Batch 764 : Gut microbiota abundance (family Erysipelotrichaceae id.2149) on Arthropathy in ulcerative colitis

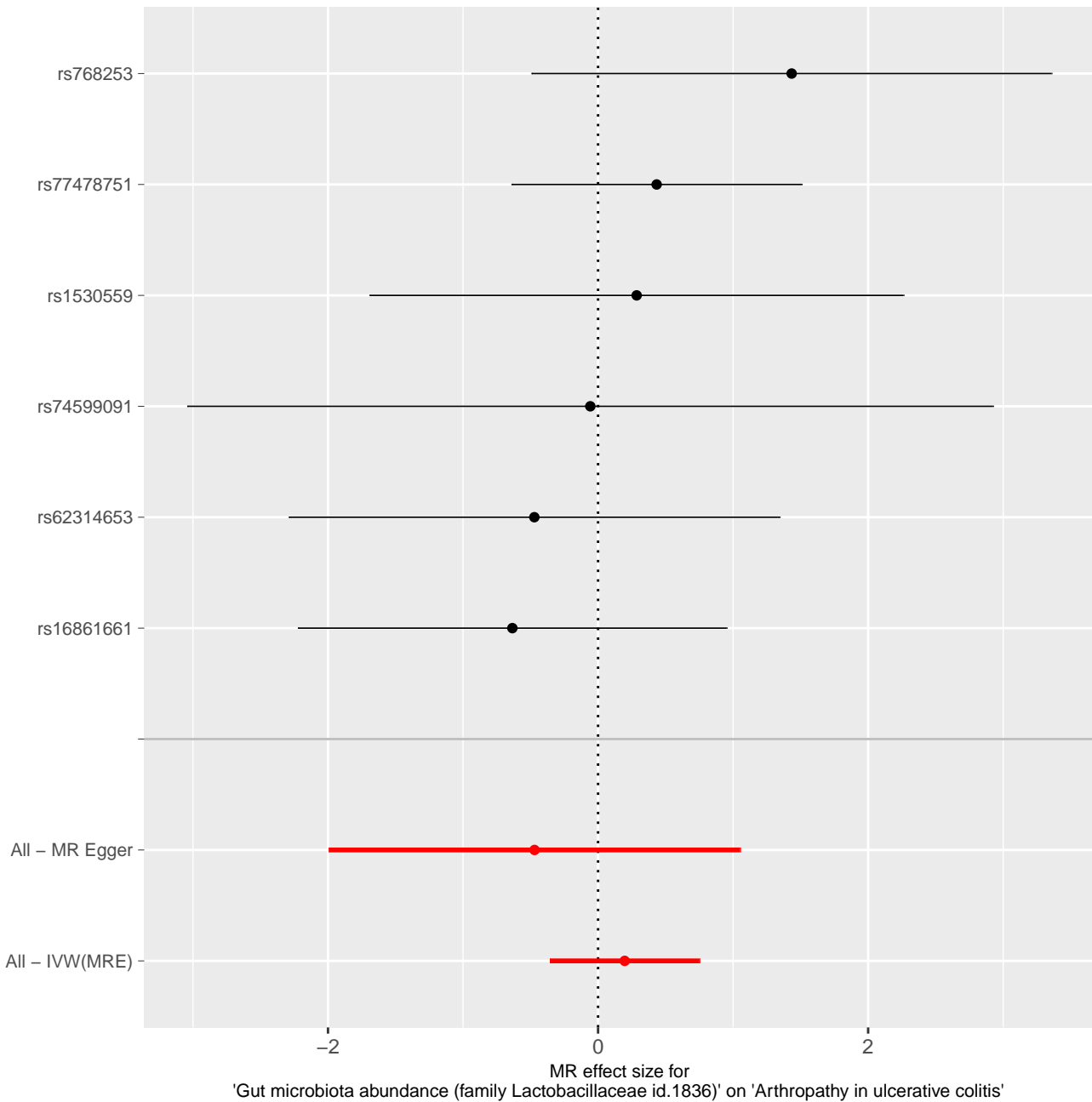


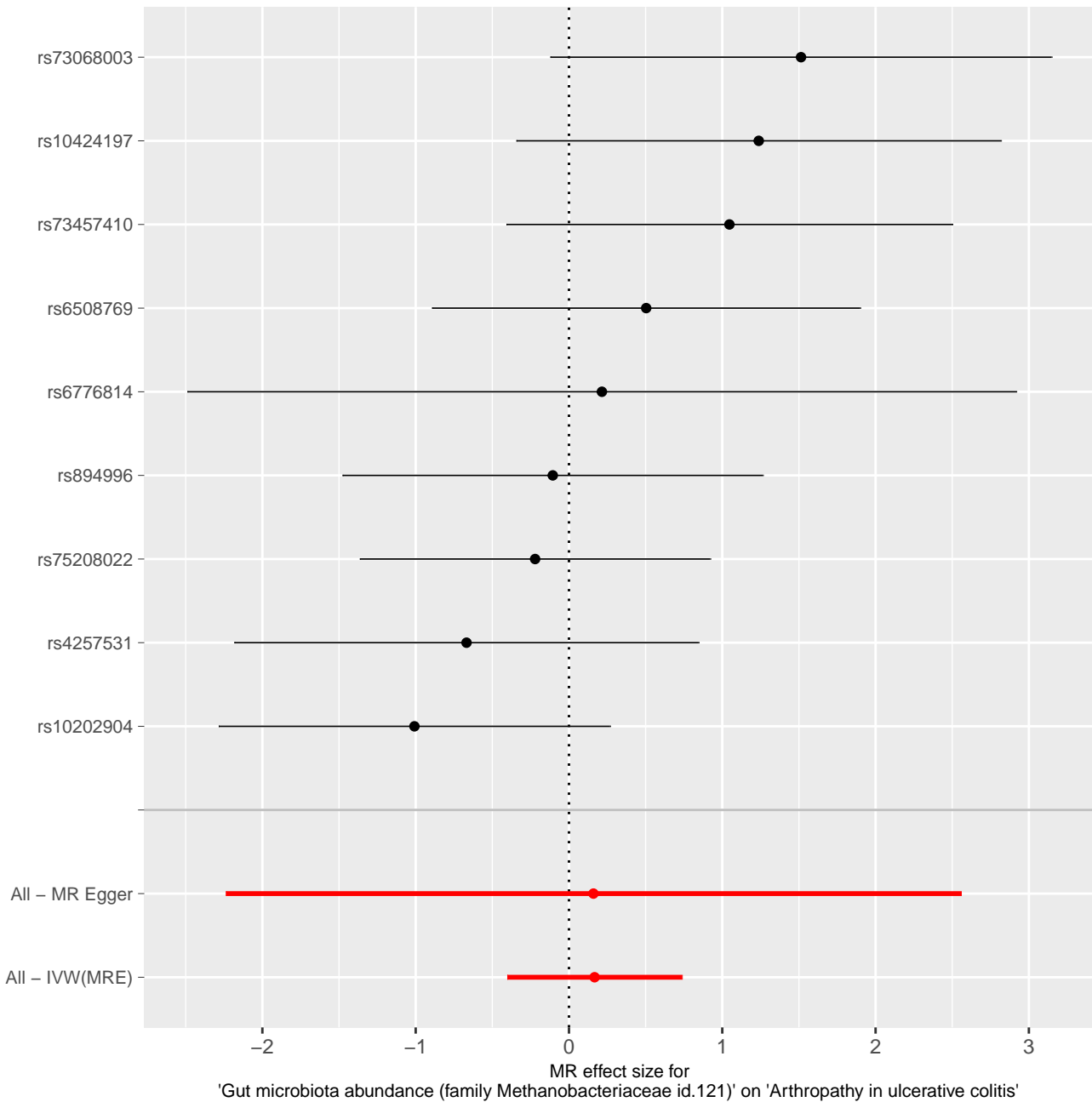




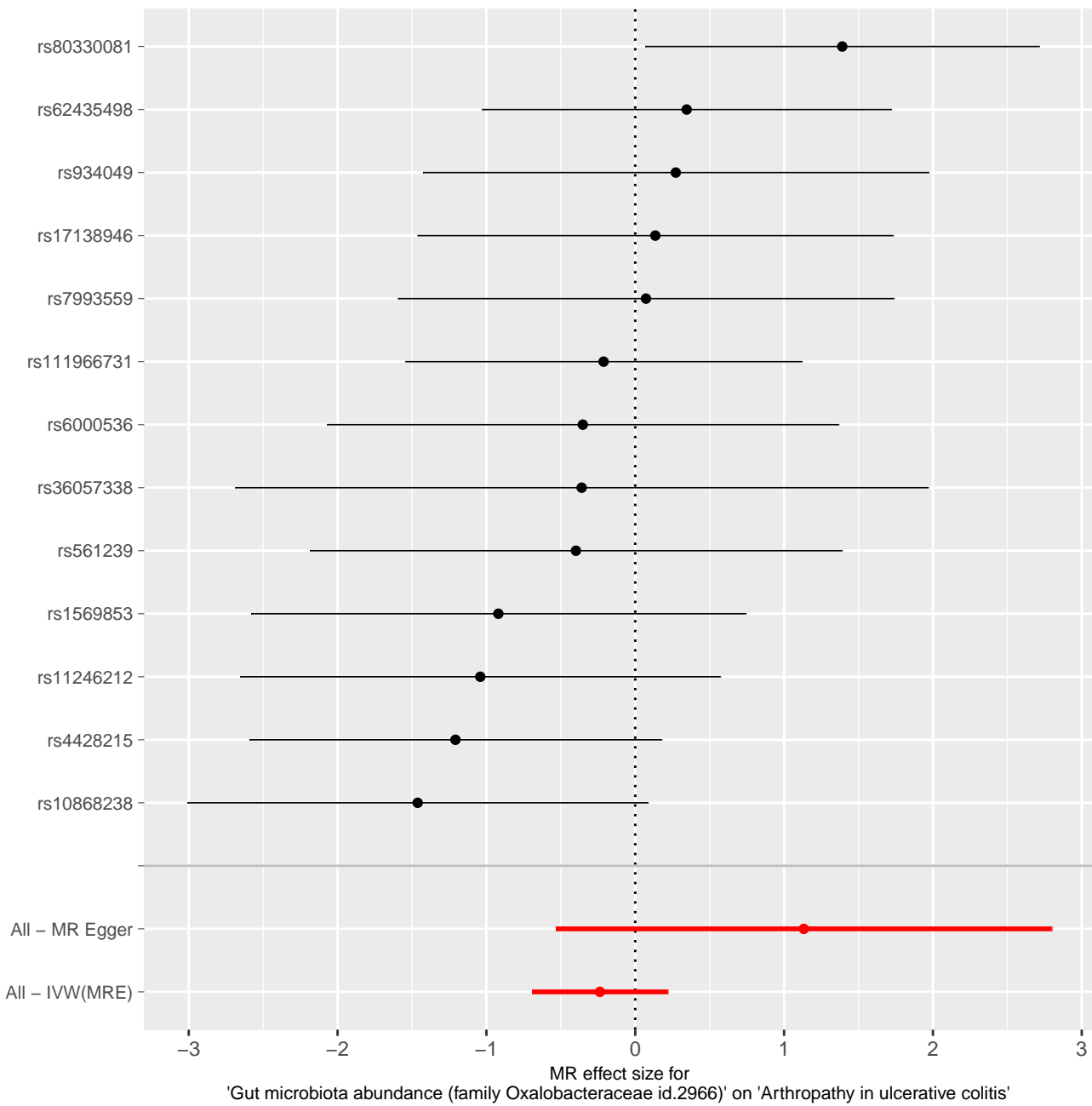
Batch 767 : Gut microbiota abundance (family Lachnospiraceae id.1987) on Arthropathy in ulcerative colitis



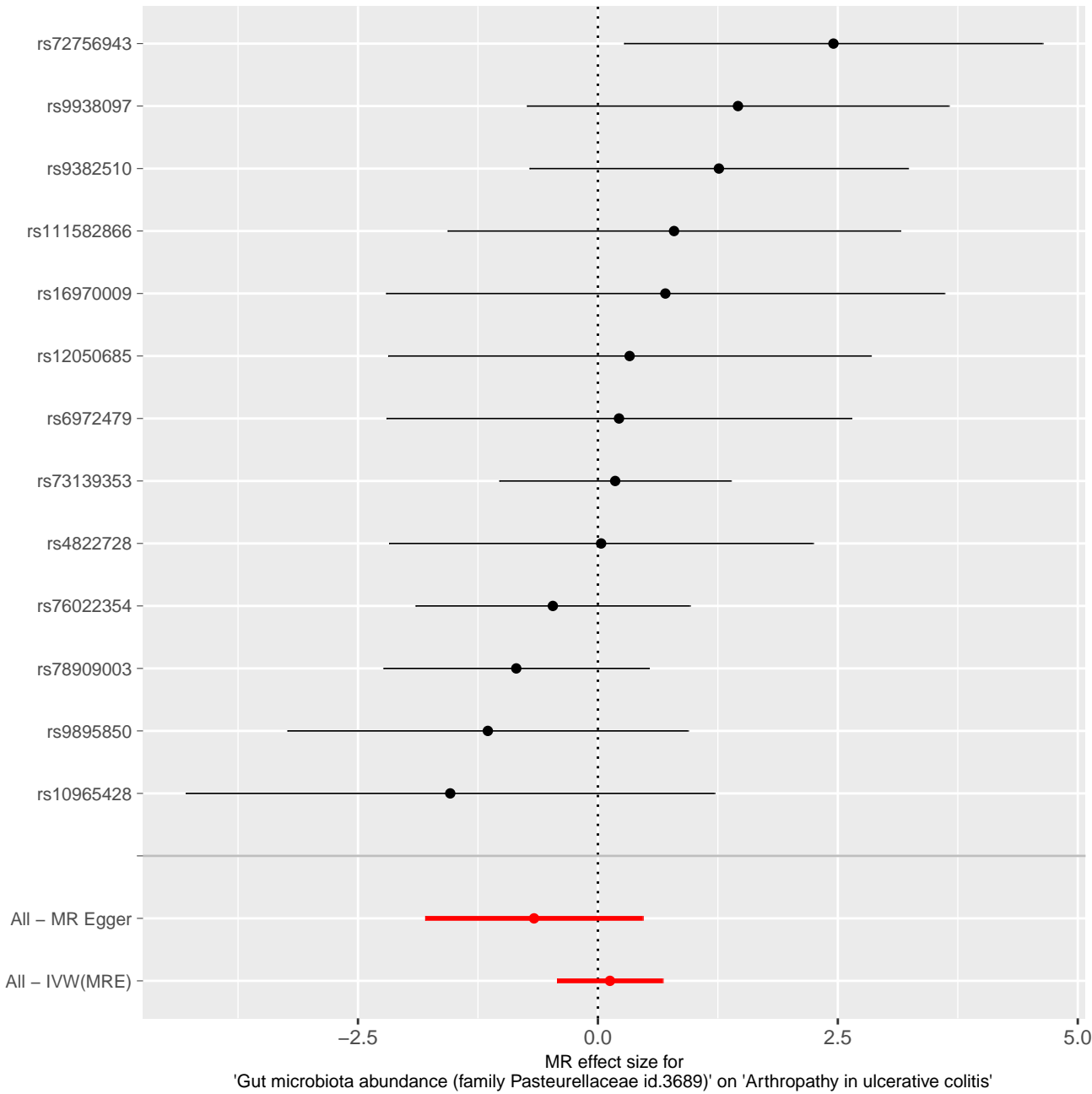


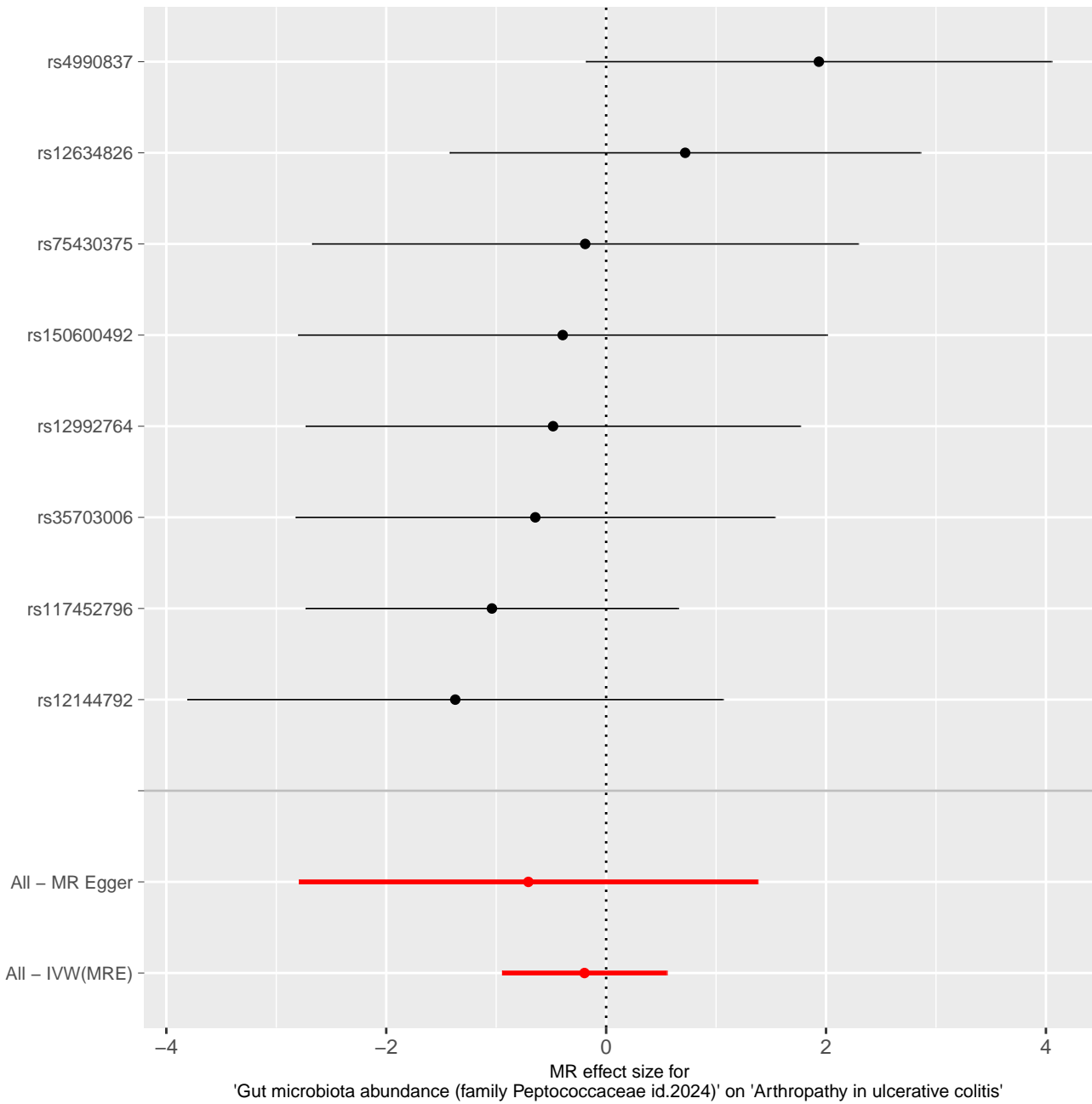


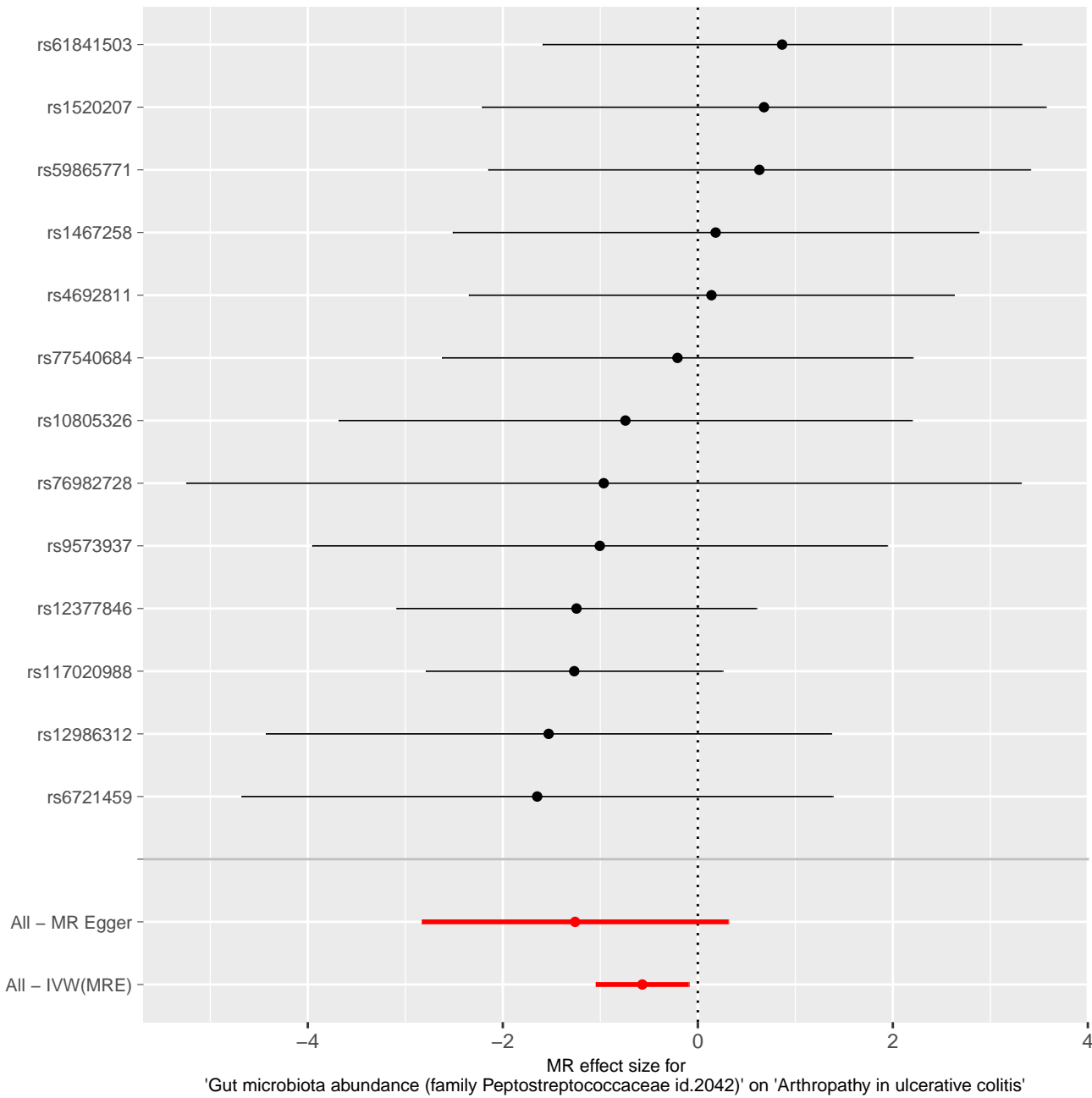
Batch 770 : Gut microbiota abundance (family Oxalobacteraceae id.2966) on Arthropathy in ulcerative colitis

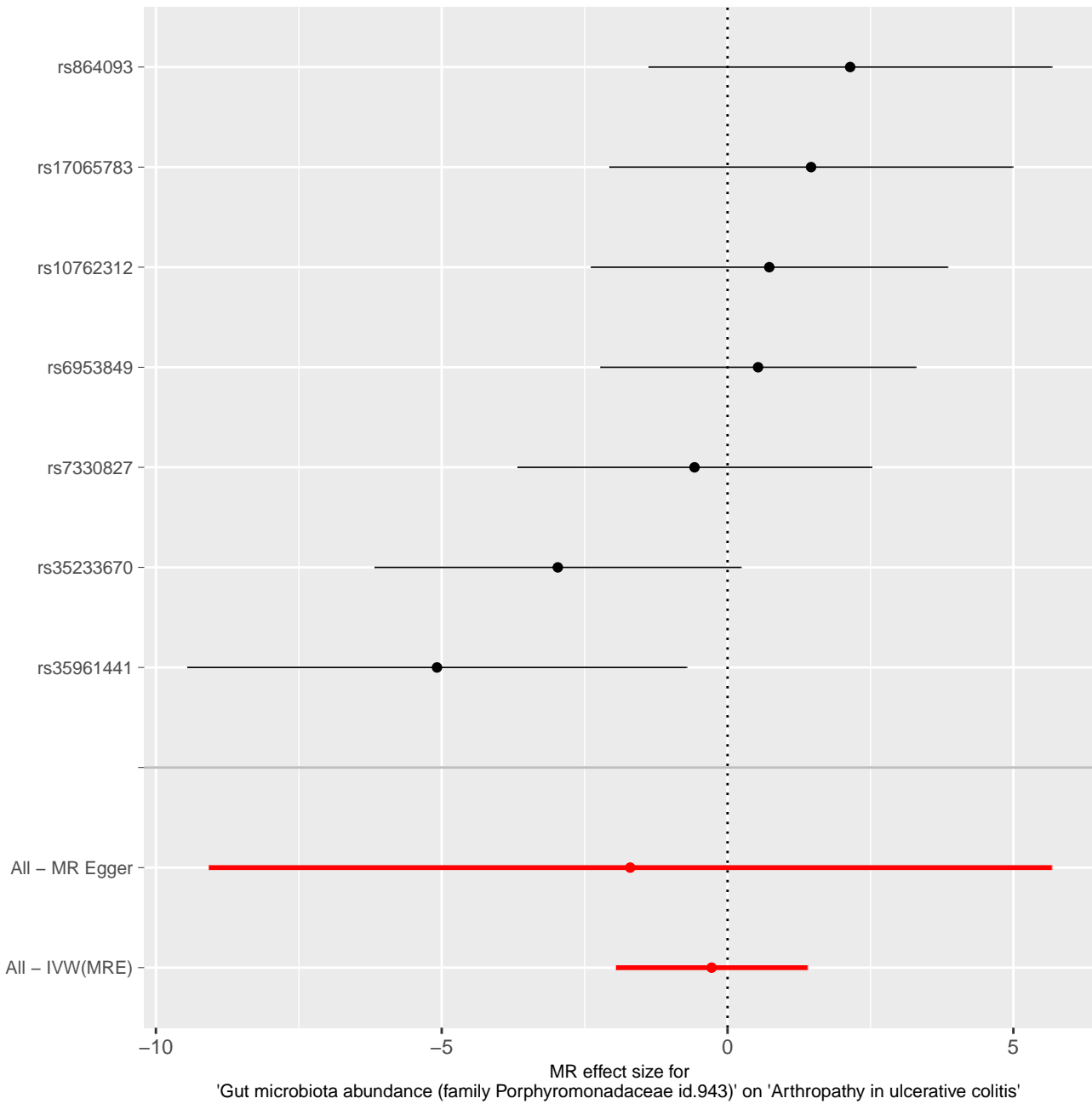


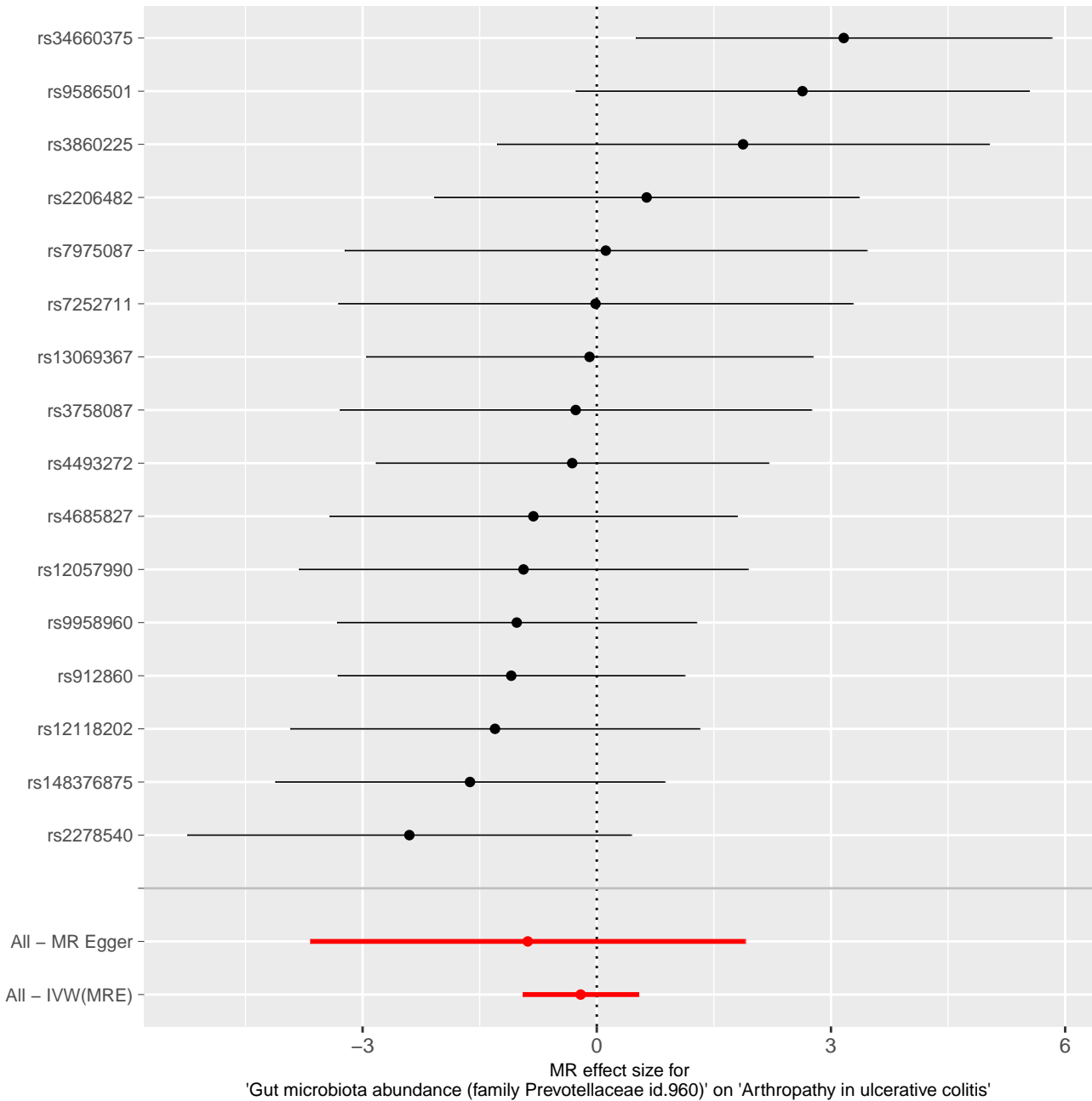
Batch 771 : Gut microbiota abundance (family Pasteurellaceae id.3689) on Arthropathy in ulcerative colitis

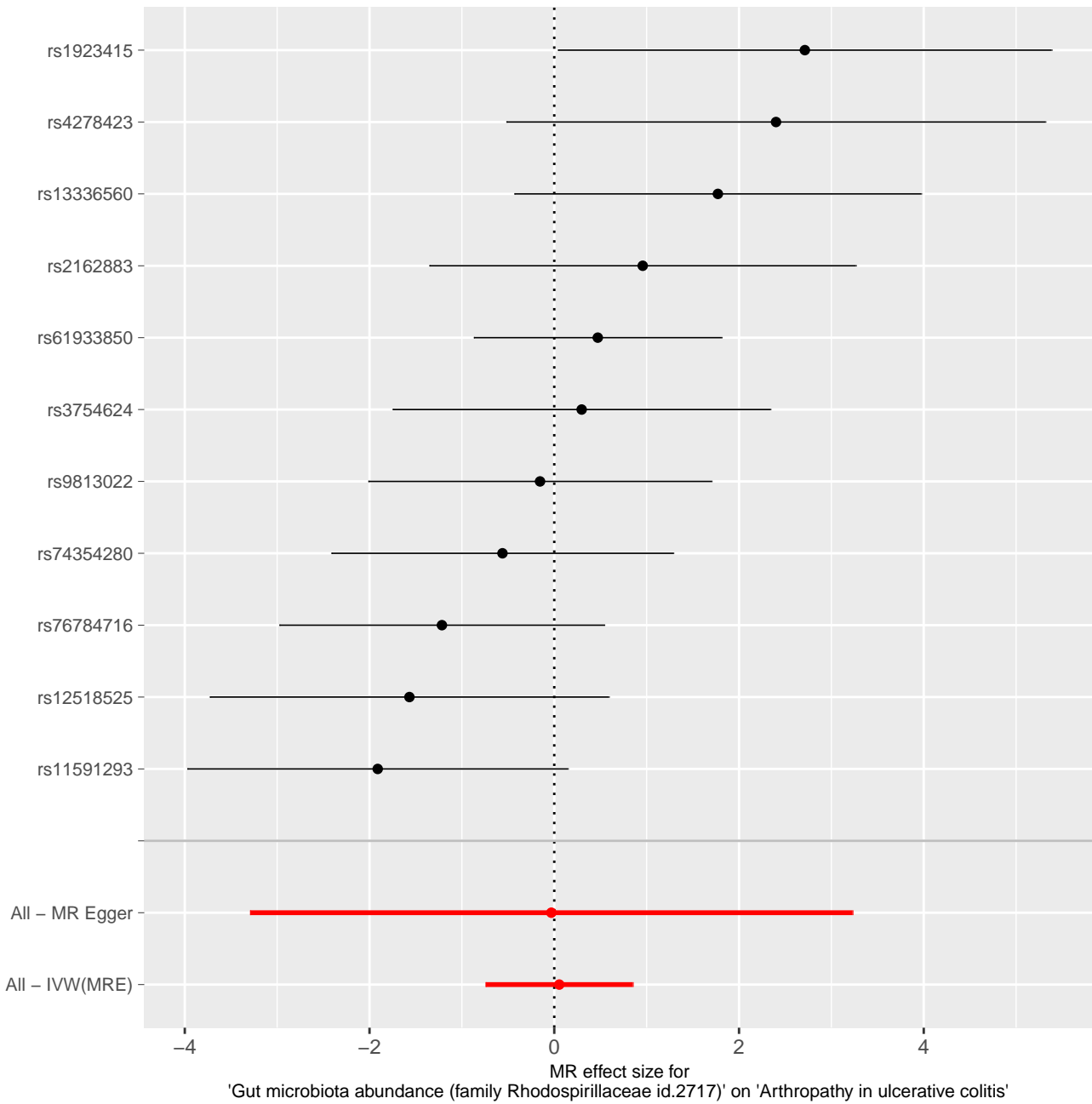


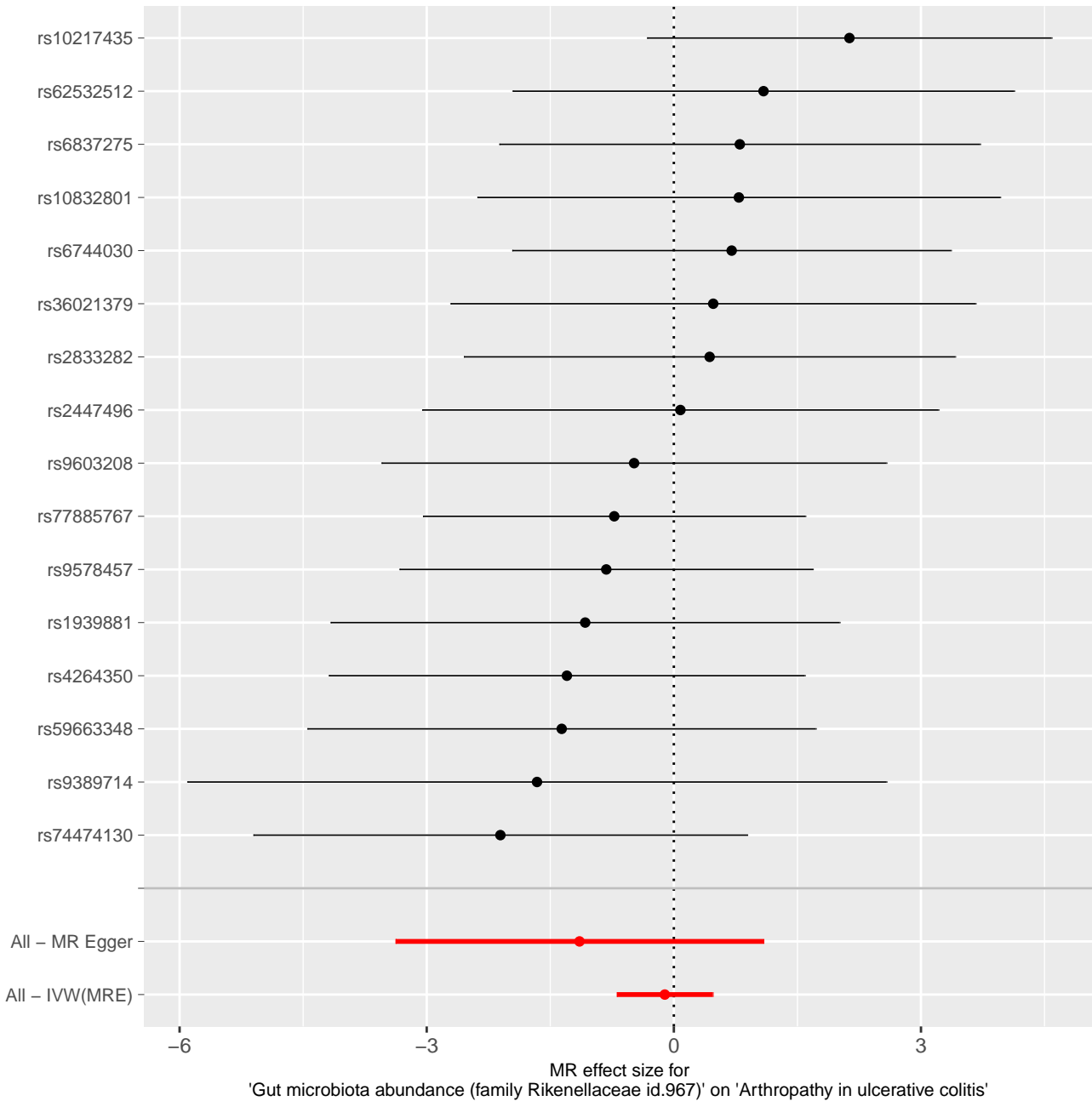




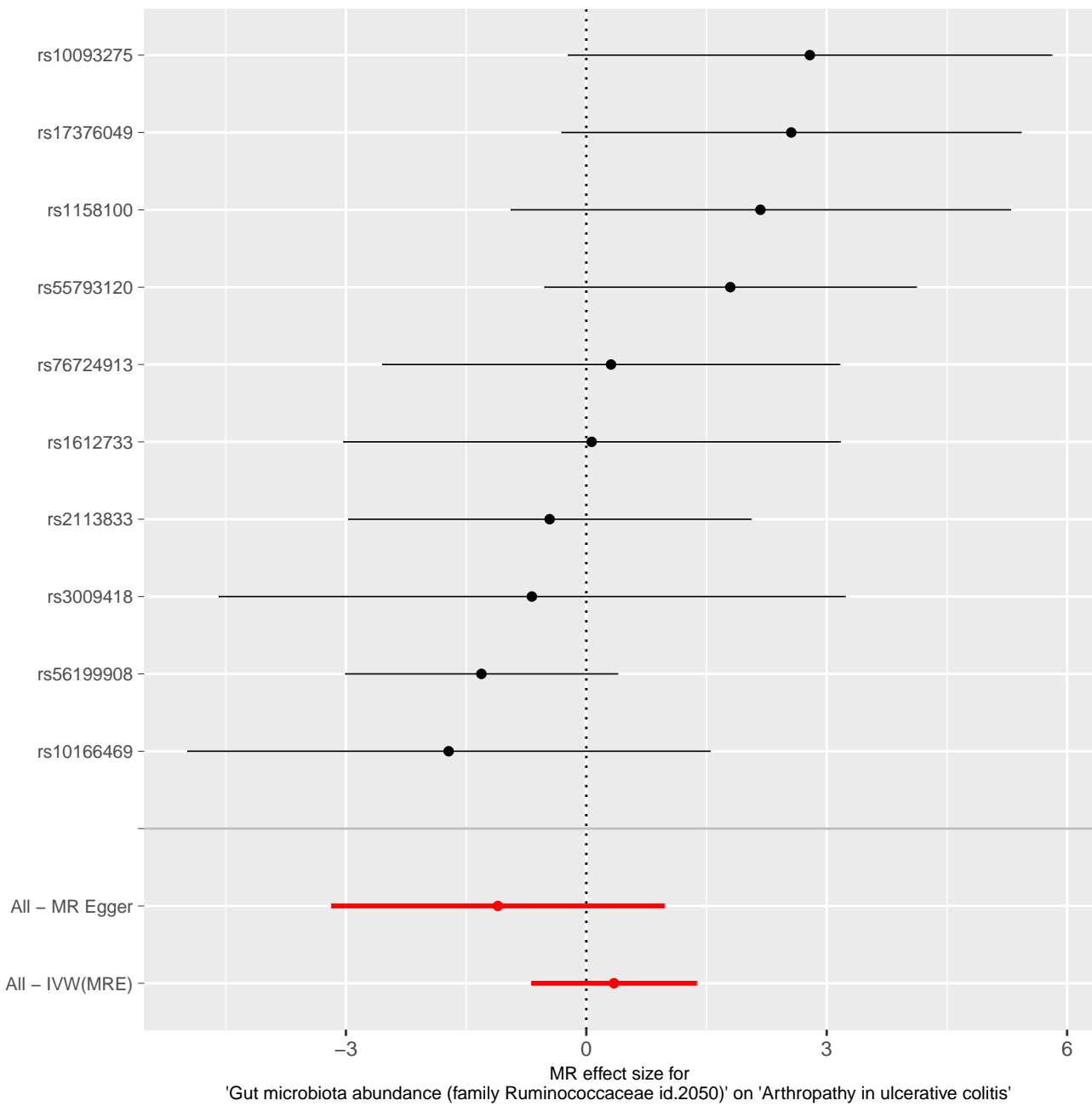




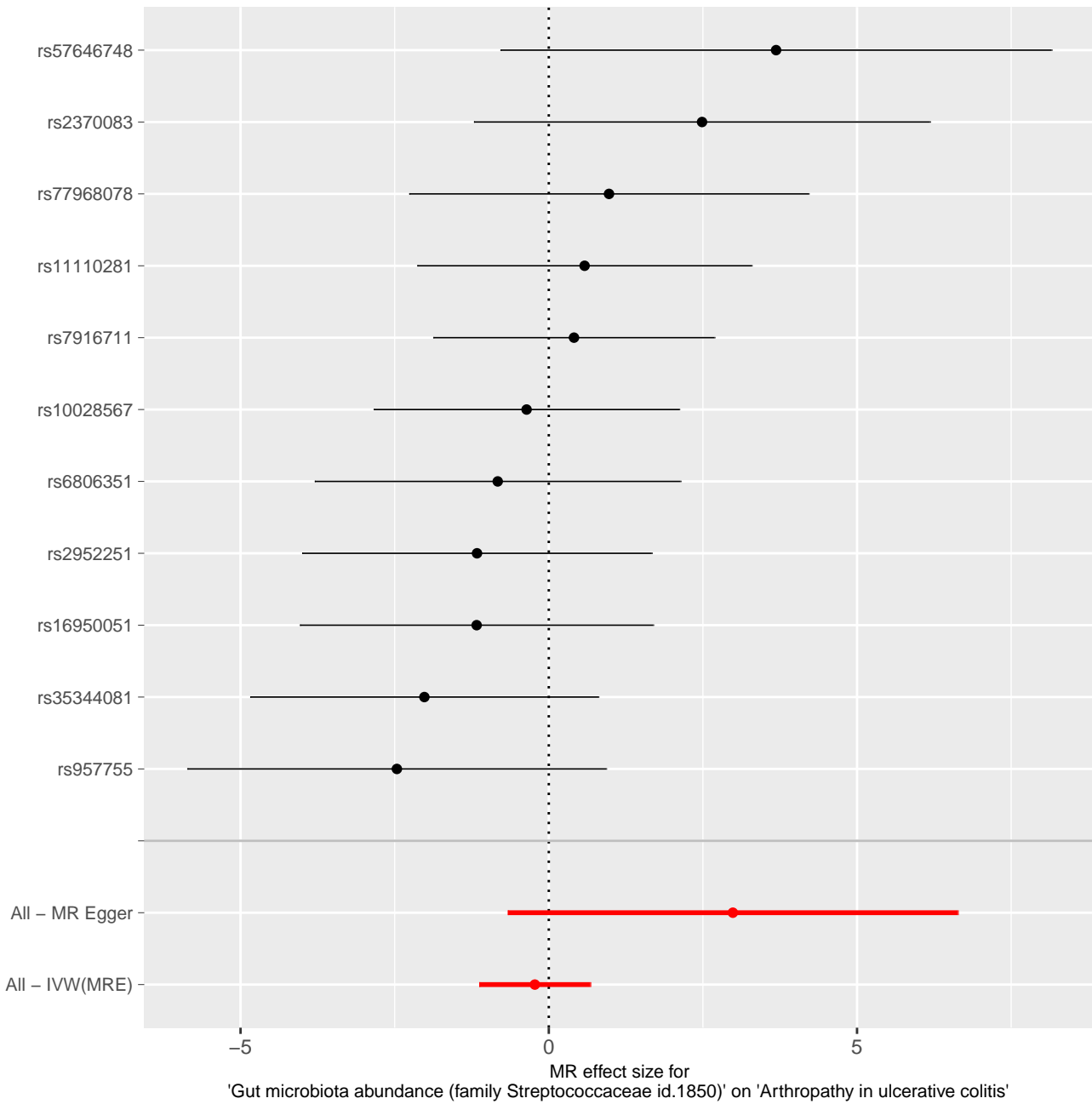




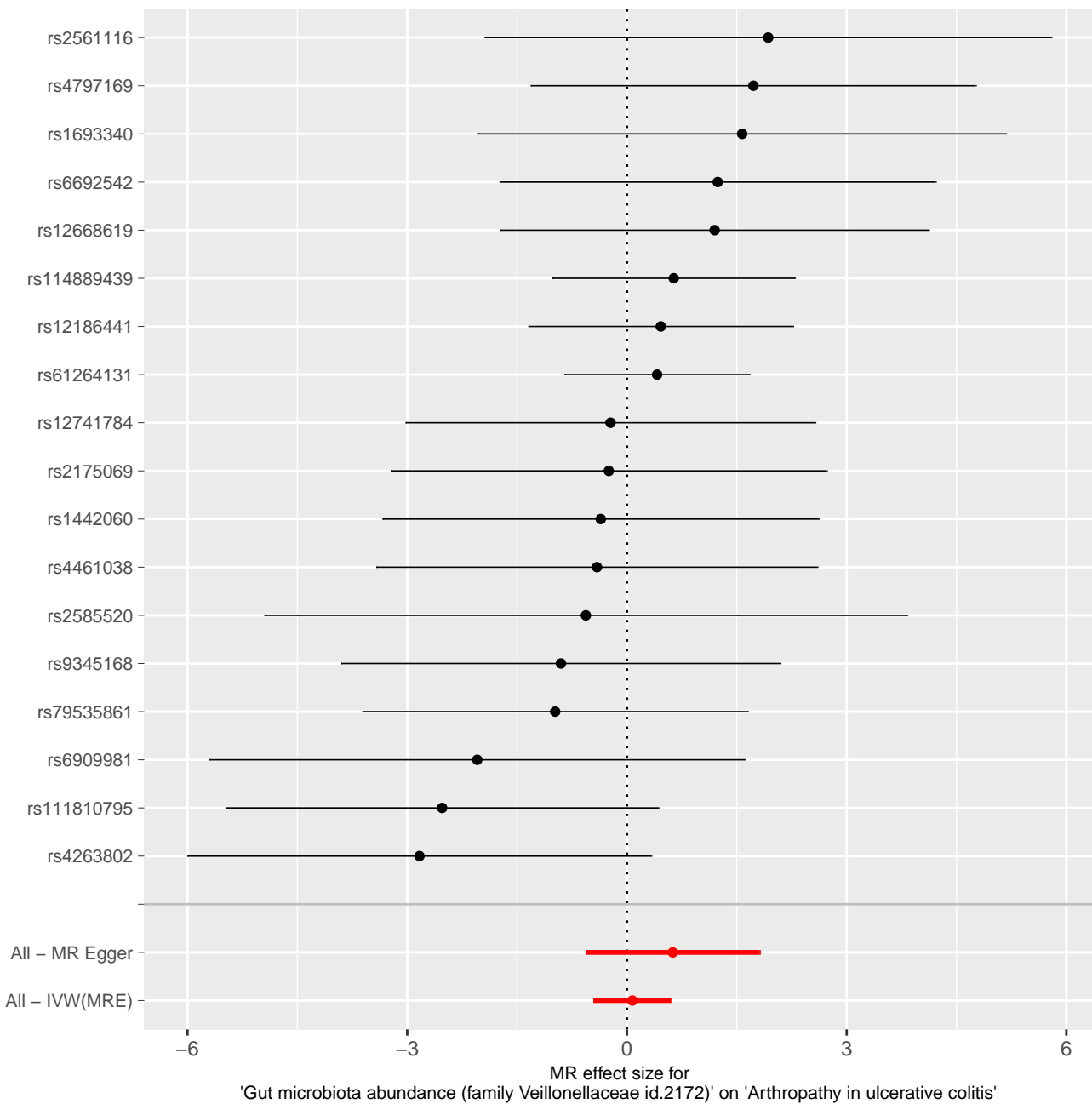
Batch 778 : Gut microbiota abundance (family Ruminococcaceae id.2050) on Arthropathy in ulcerative colitis



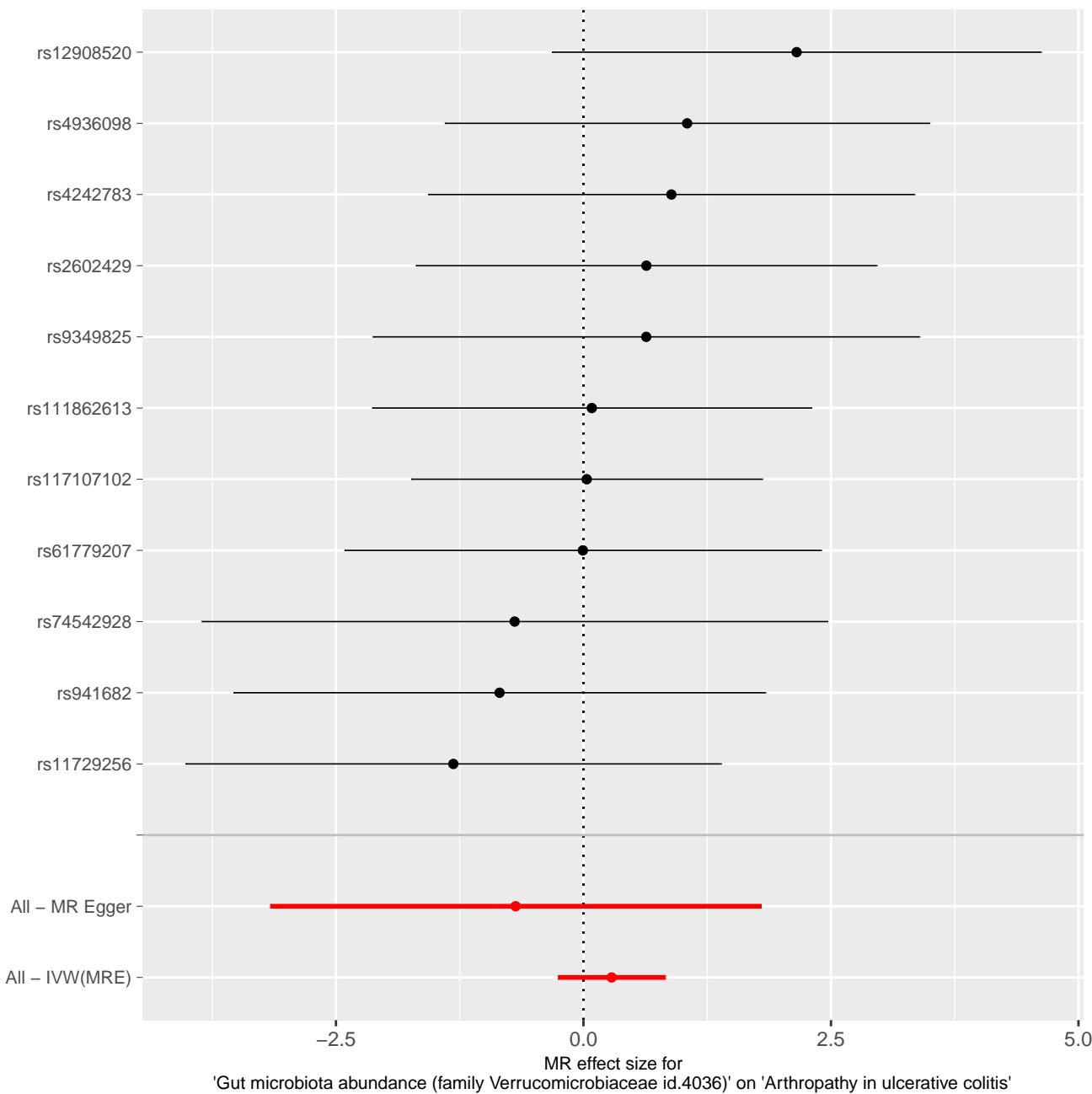
Batch 779 : Gut microbiota abundance (family Streptococcaceae id.1850) on Arthropathy in ulcerative colitis

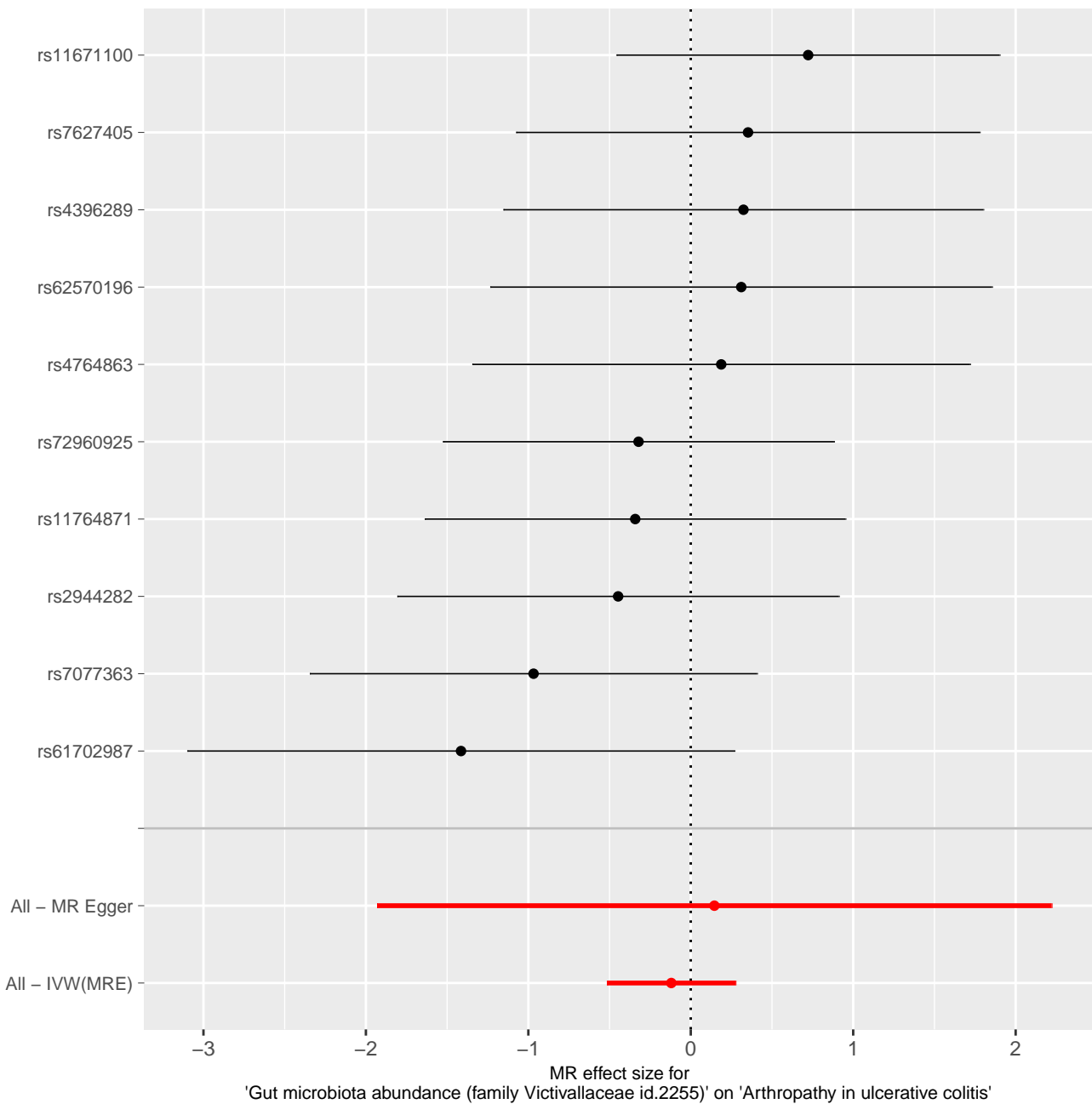


Batch 780 : Gut microbiota abundance (family Veillonellaceae id.2172) on Arthropathy in ulcerative colitis

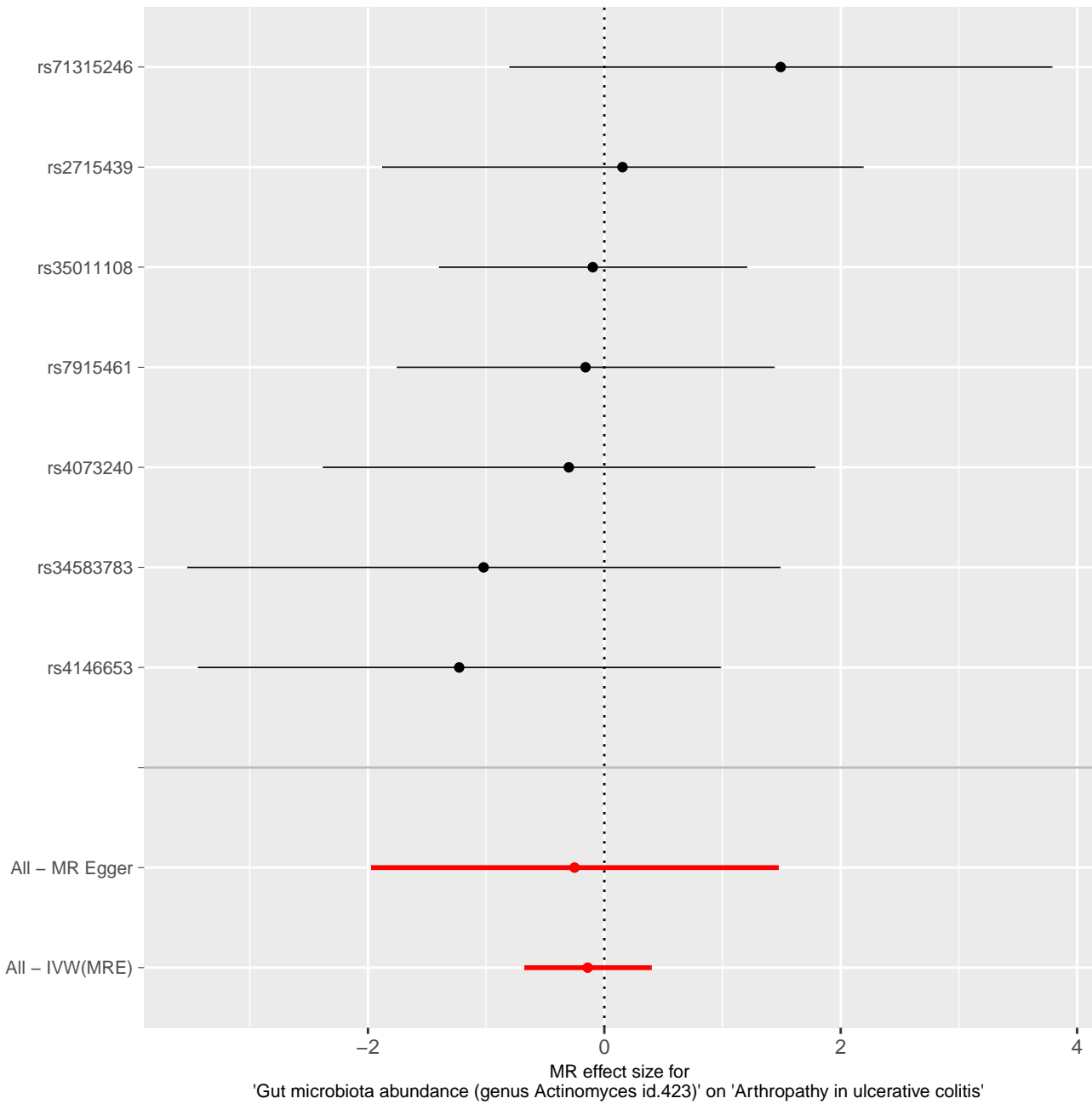


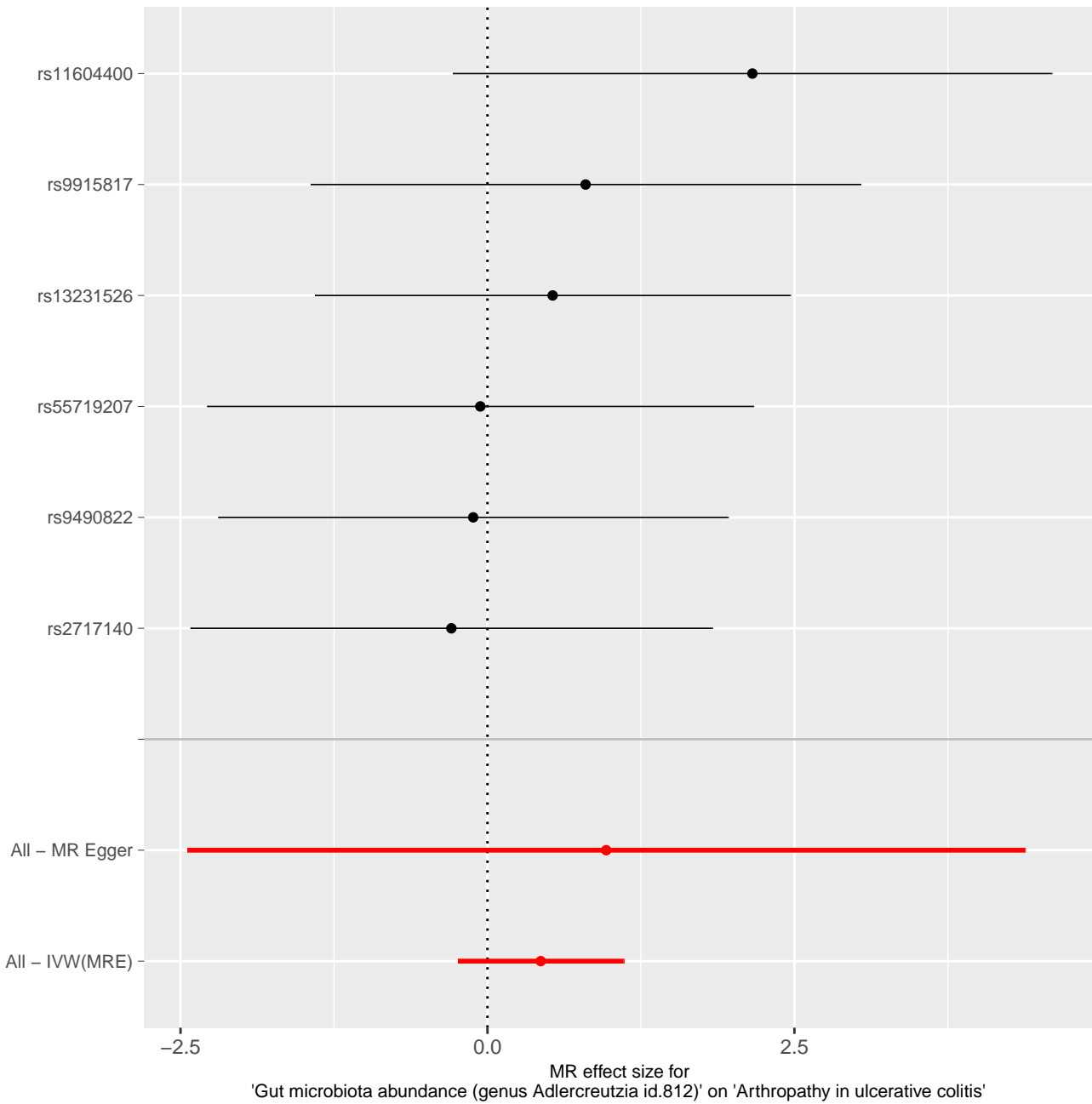
Batch 781 : Gut microbiota abundance (family Verrucomicrobiaceae id.4036) on Arthropathy in ulcerative colitis



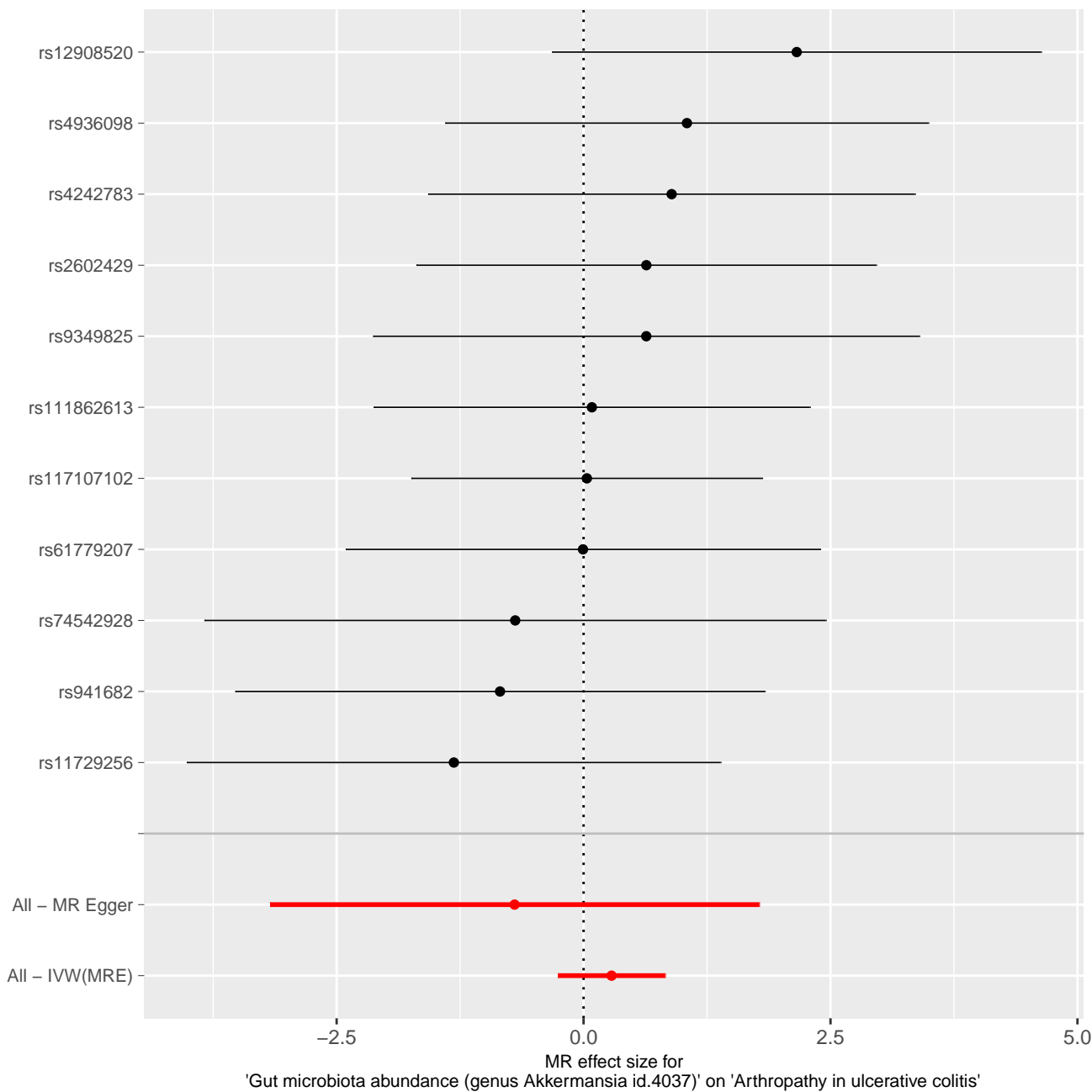


Batch 783 : Gut microbiota abundance (genus Actinomyces id.423) on Arthropathy in ulcerative colitis

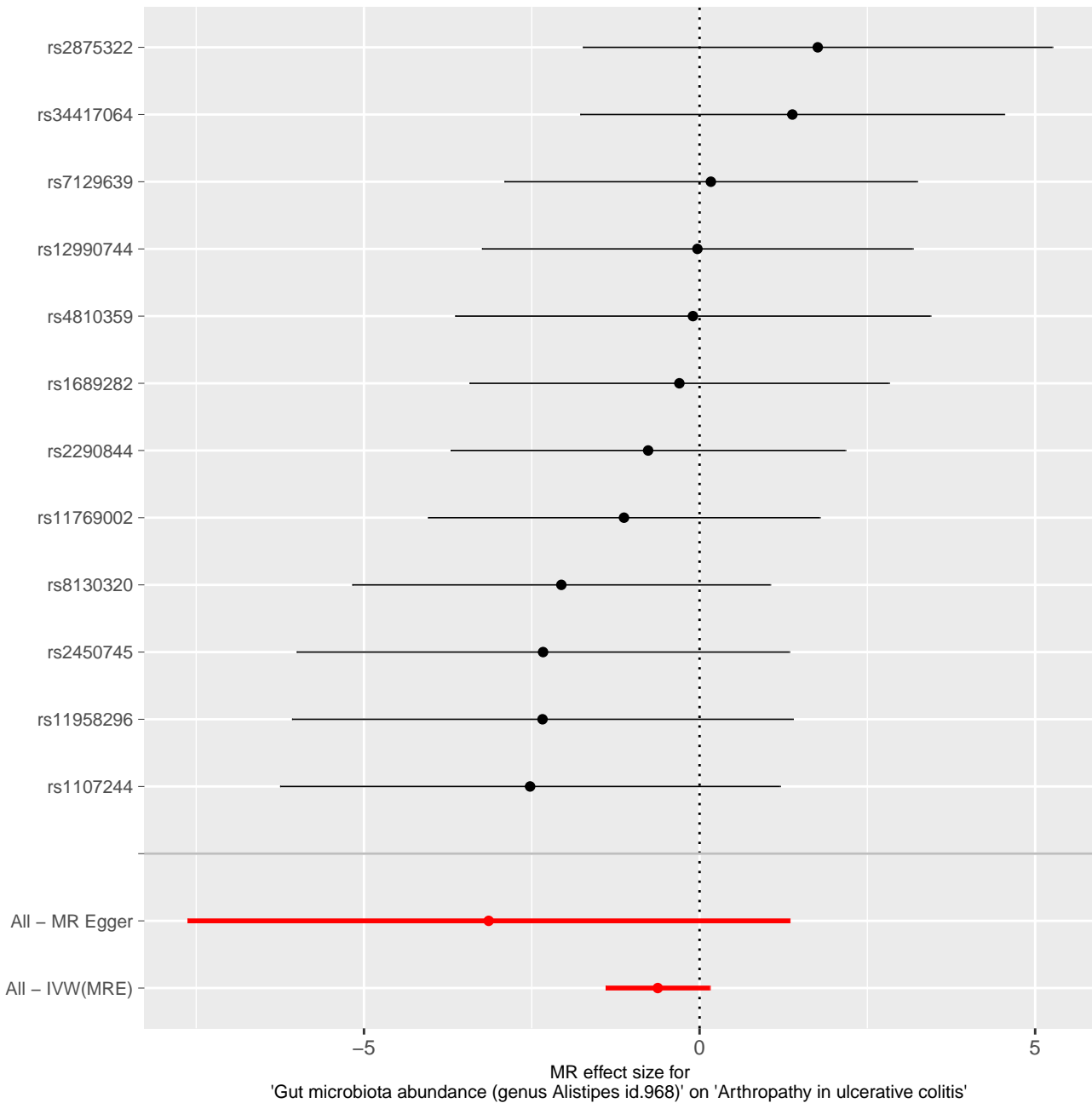


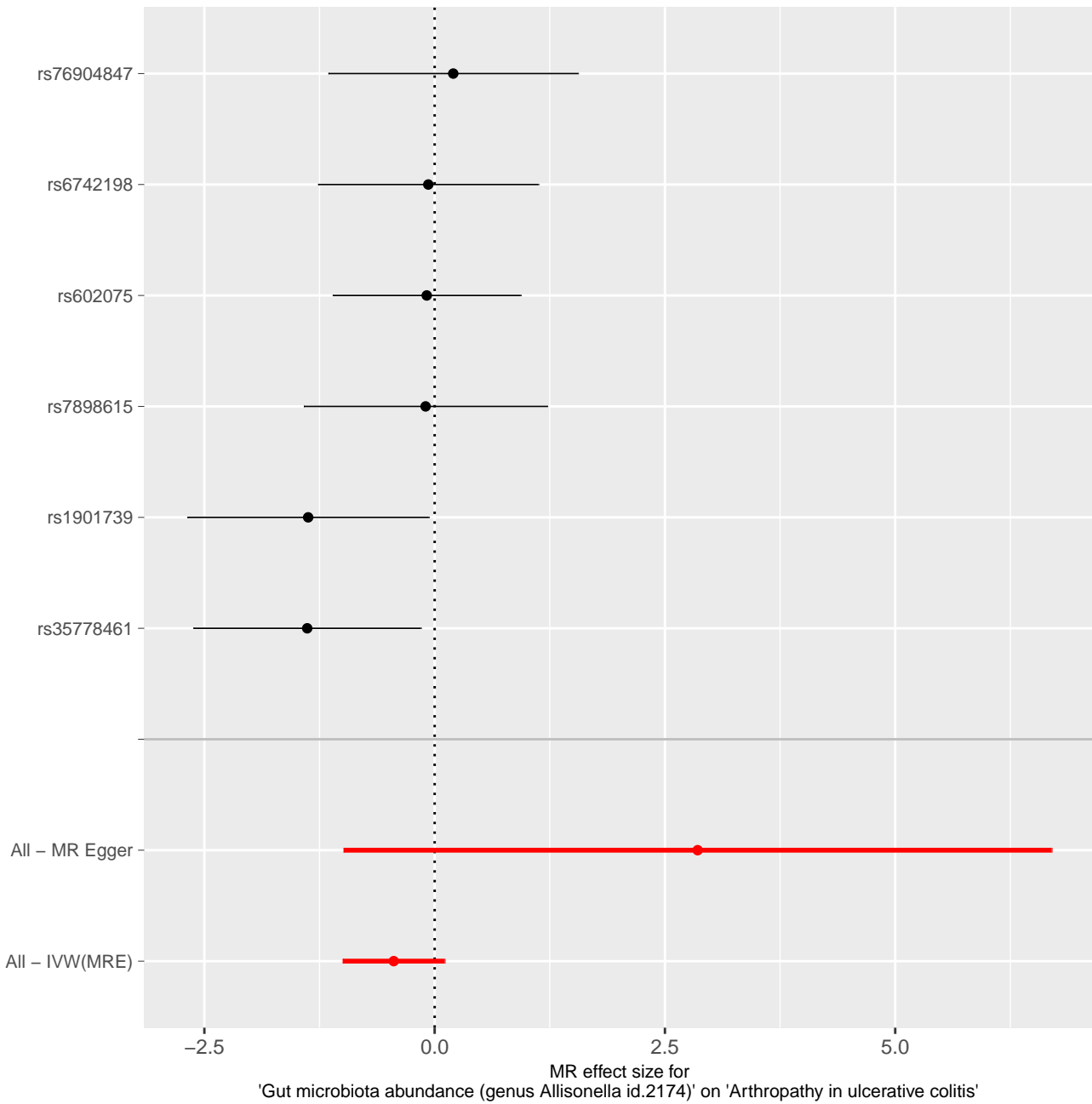


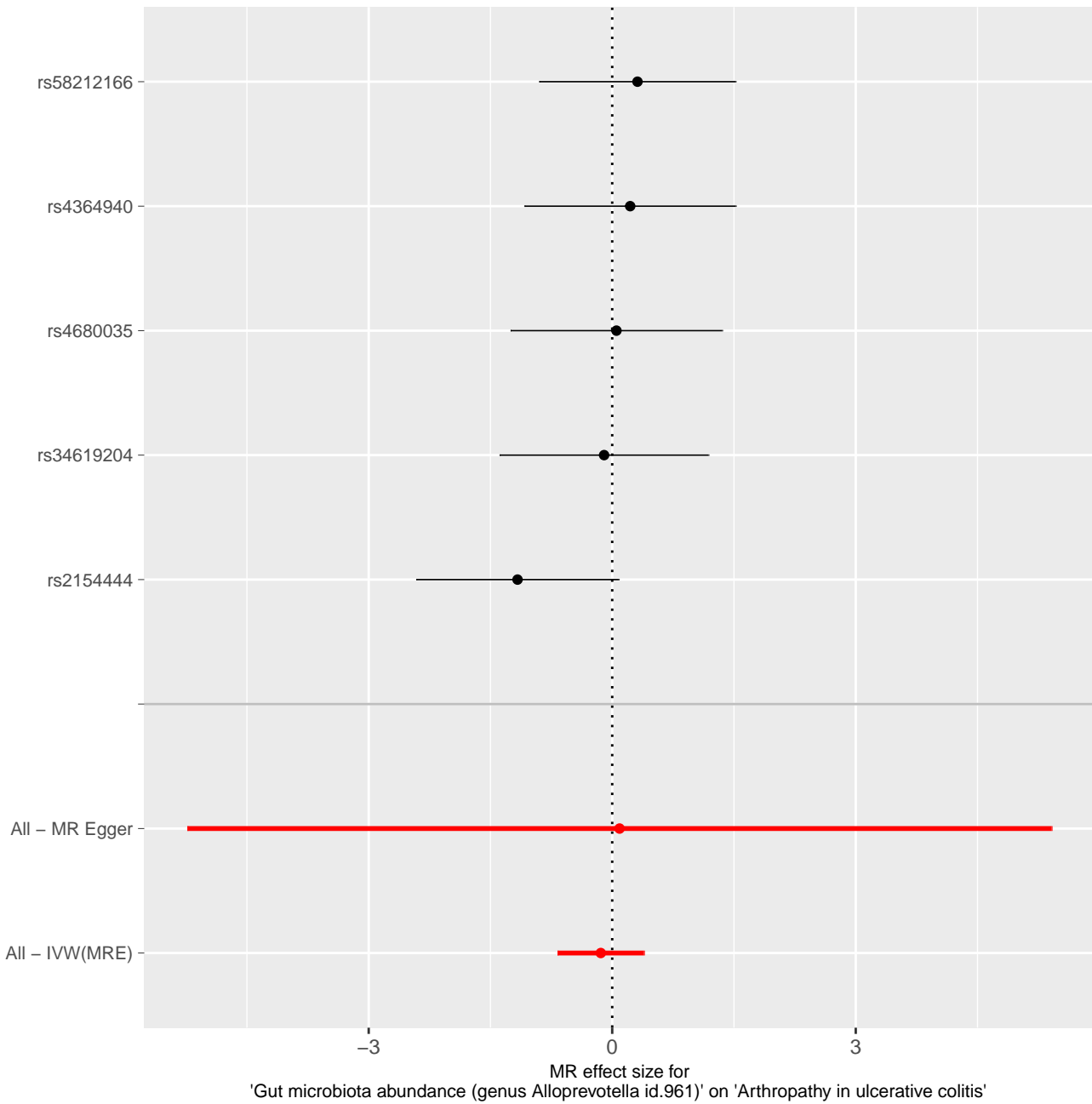
Batch 785 : Gut microbiota abundance (genus Akkermansia id.4037) on Arthropathy in ulcerative colitis



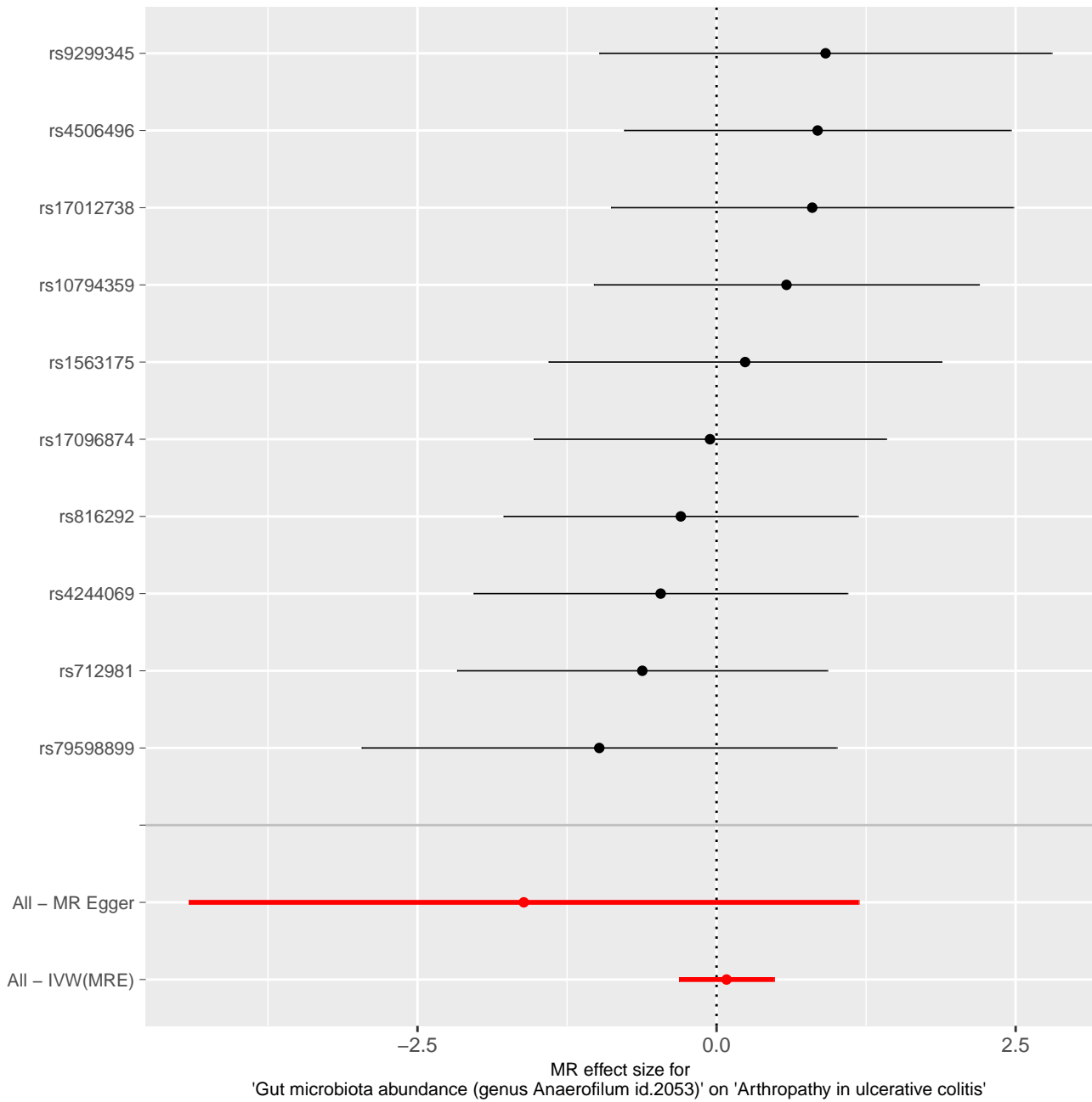
Batch 786 : Gut microbiota abundance (genus Alistipes id.968) on Arthropathy in ulcerative colitis



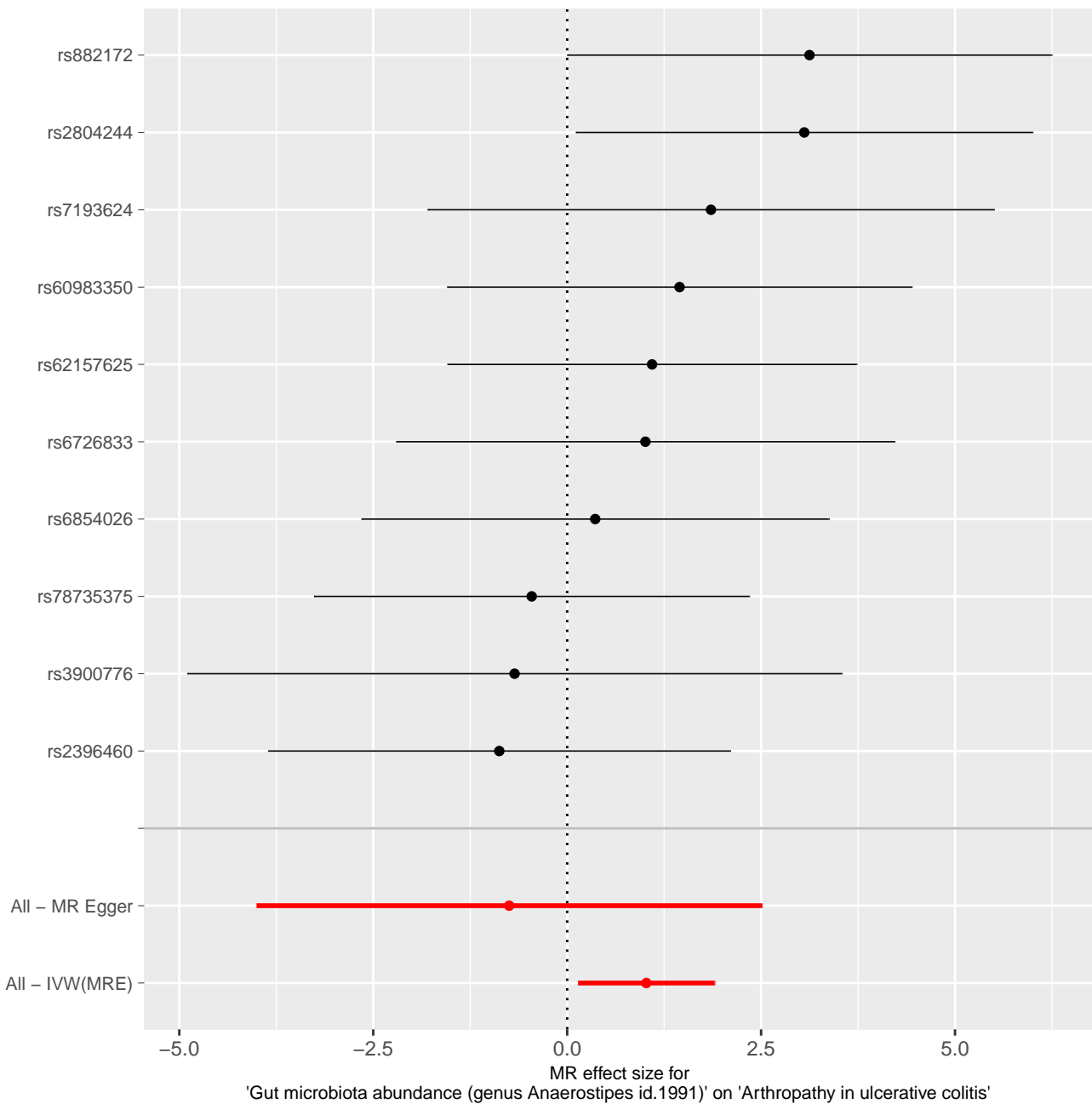




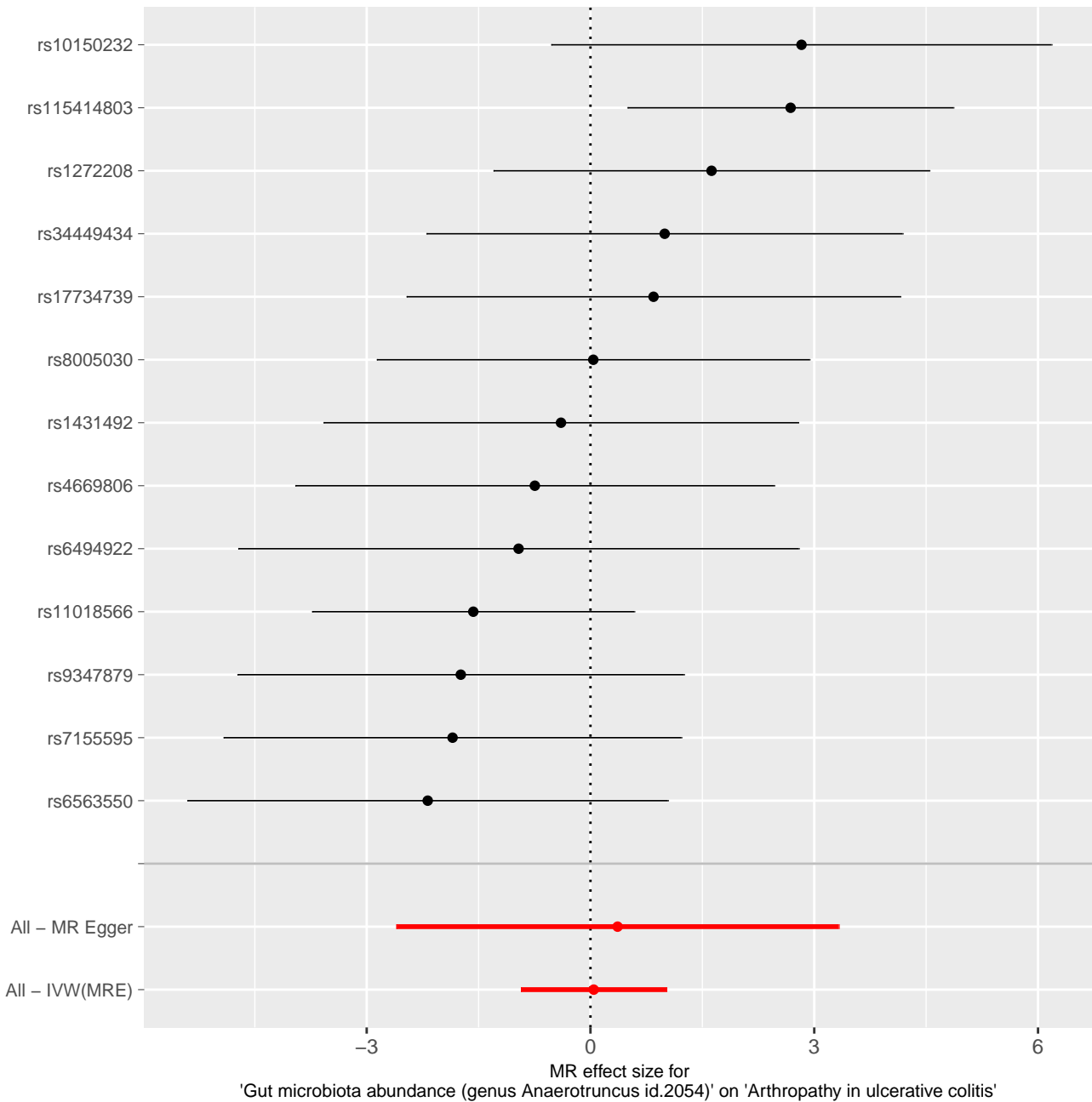
Batch 789 : Gut microbiota abundance (genus Anaerofilum id.2053) on Arthropathy in ulcerative colitis

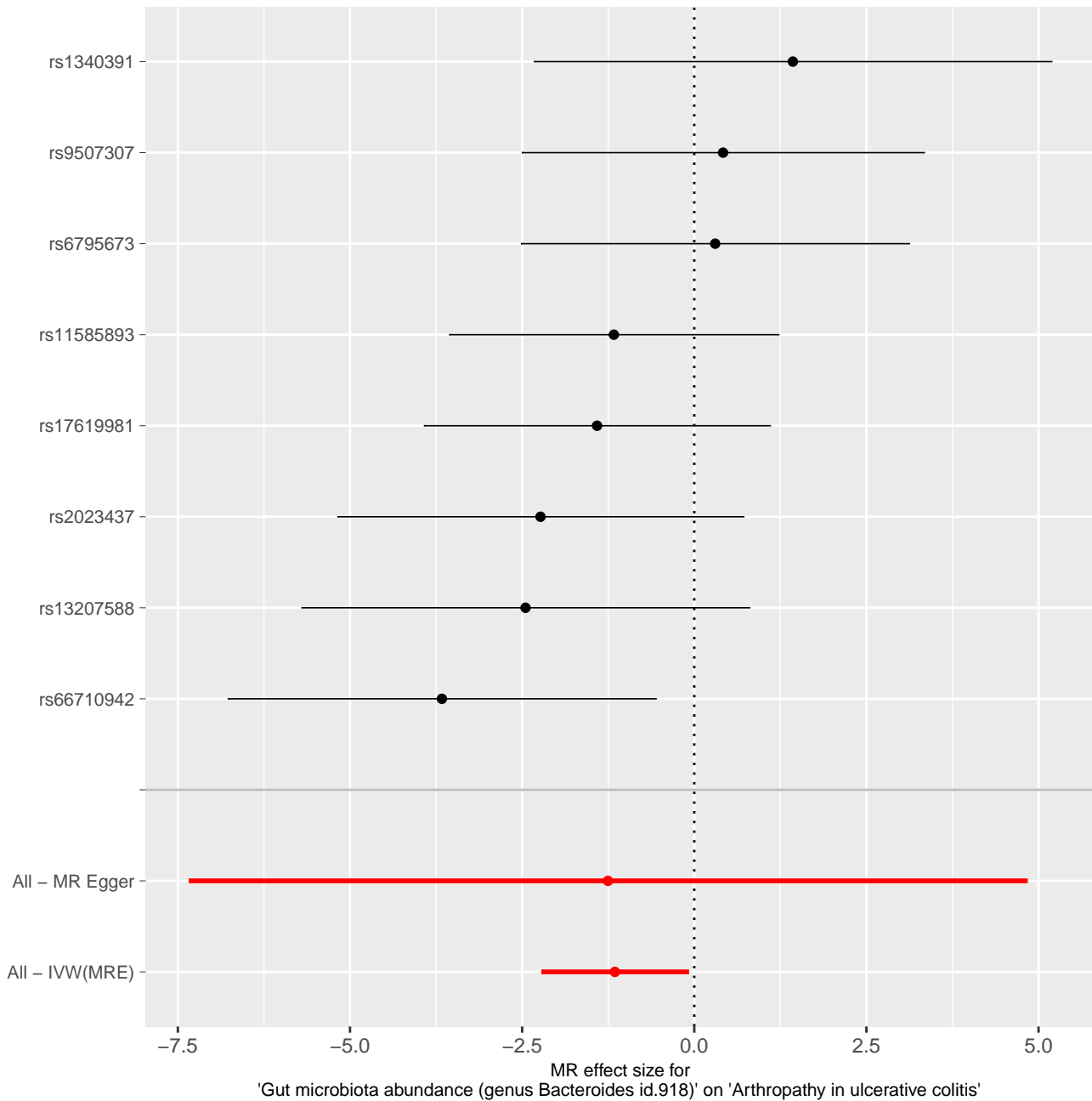


Batch 790 : Gut microbiota abundance (genus Anaerostipes id.1991) on Arthropathy in ulcerative colitis

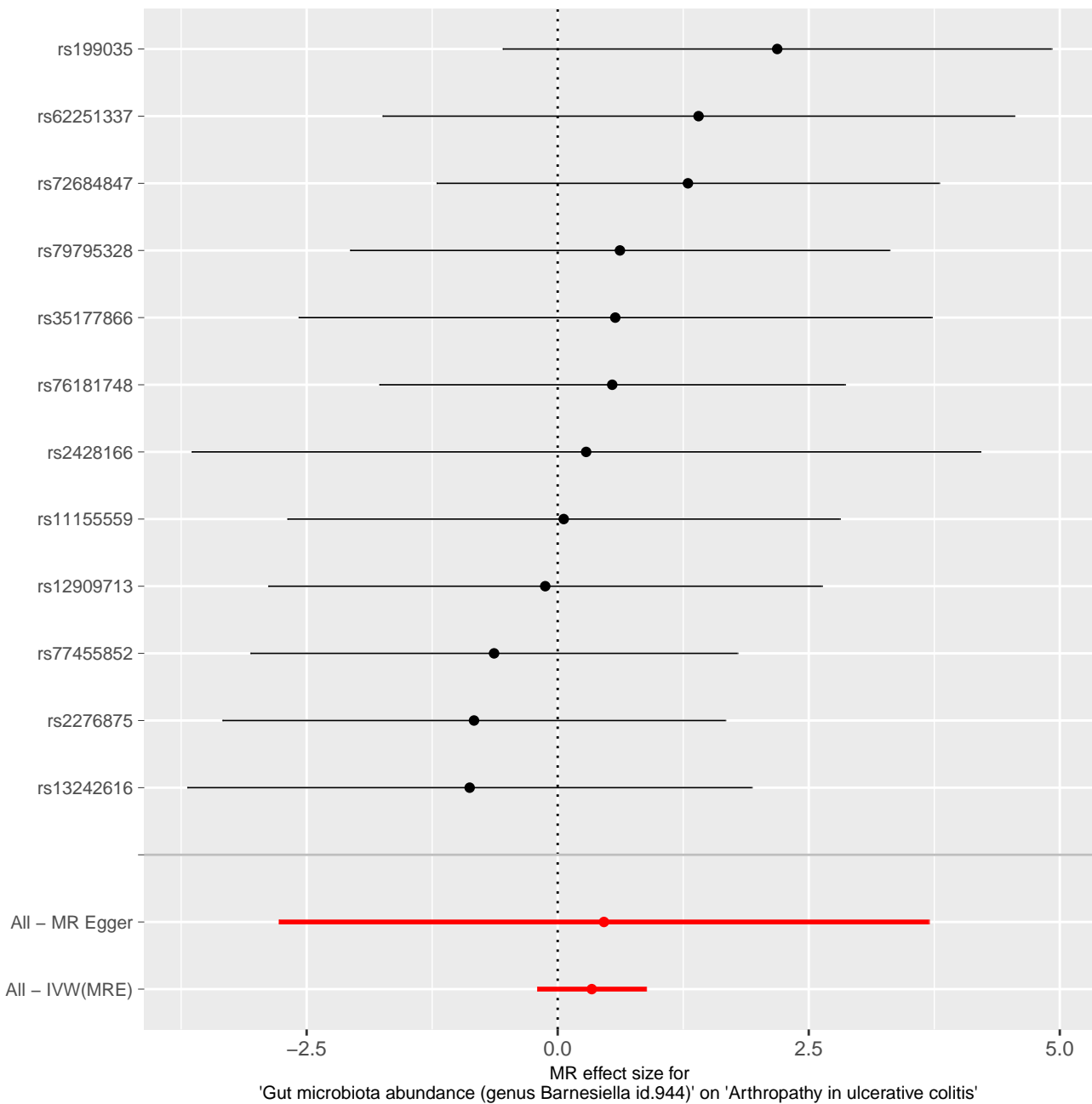


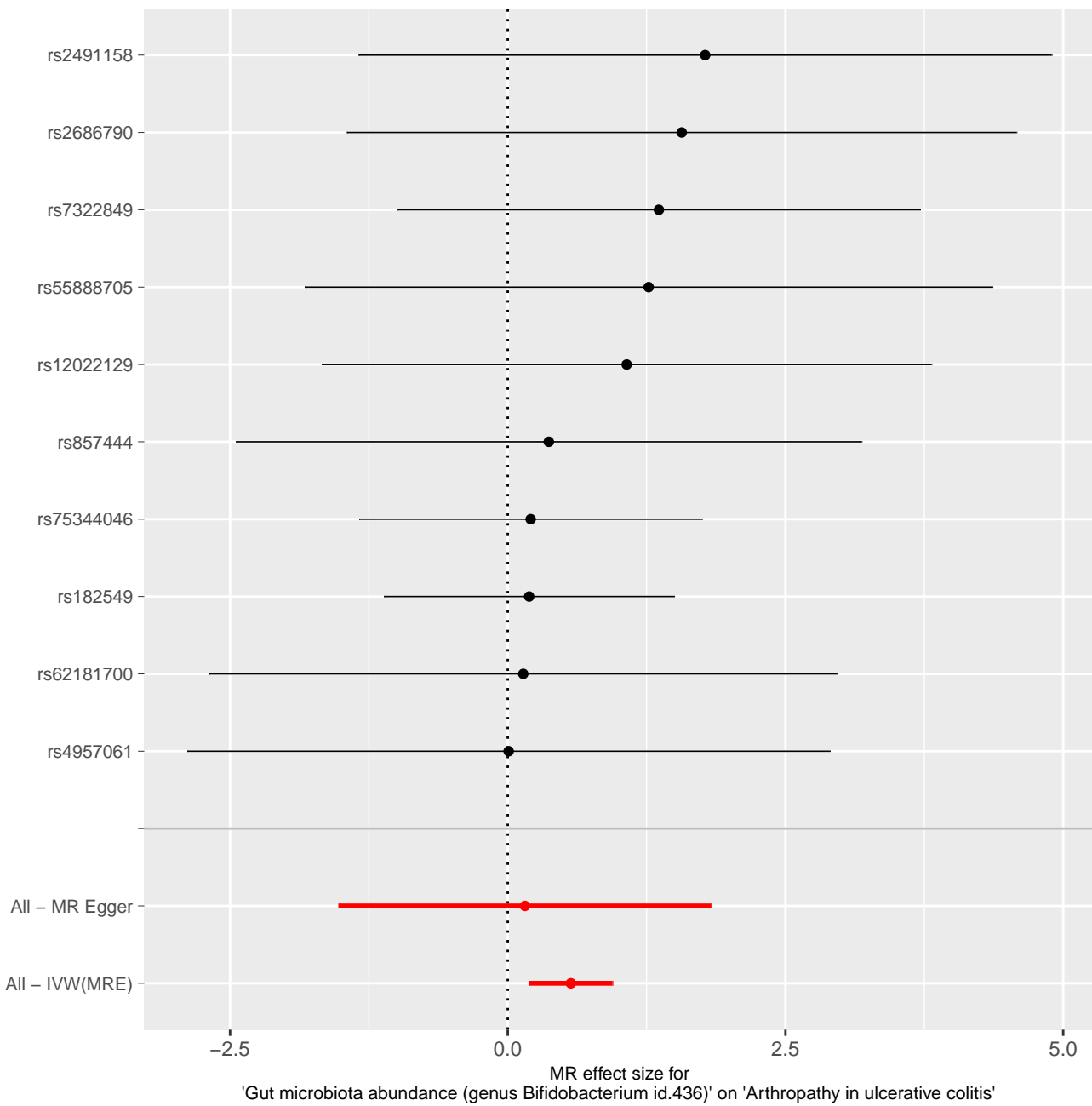
Batch 791 : Gut microbiota abundance (genus Anaerotruncus id.2054) on Arthropathy in ulcerative colitis



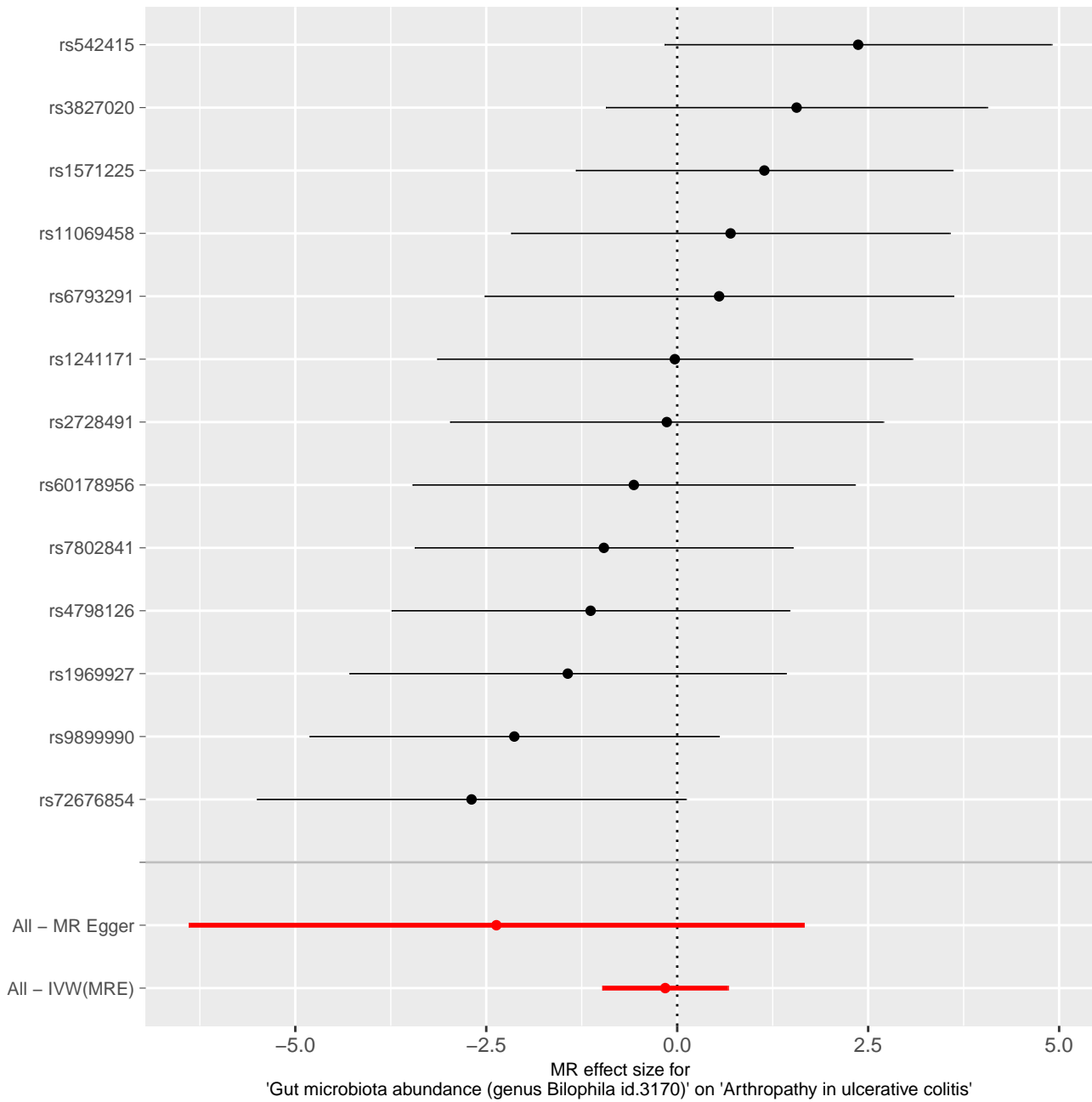


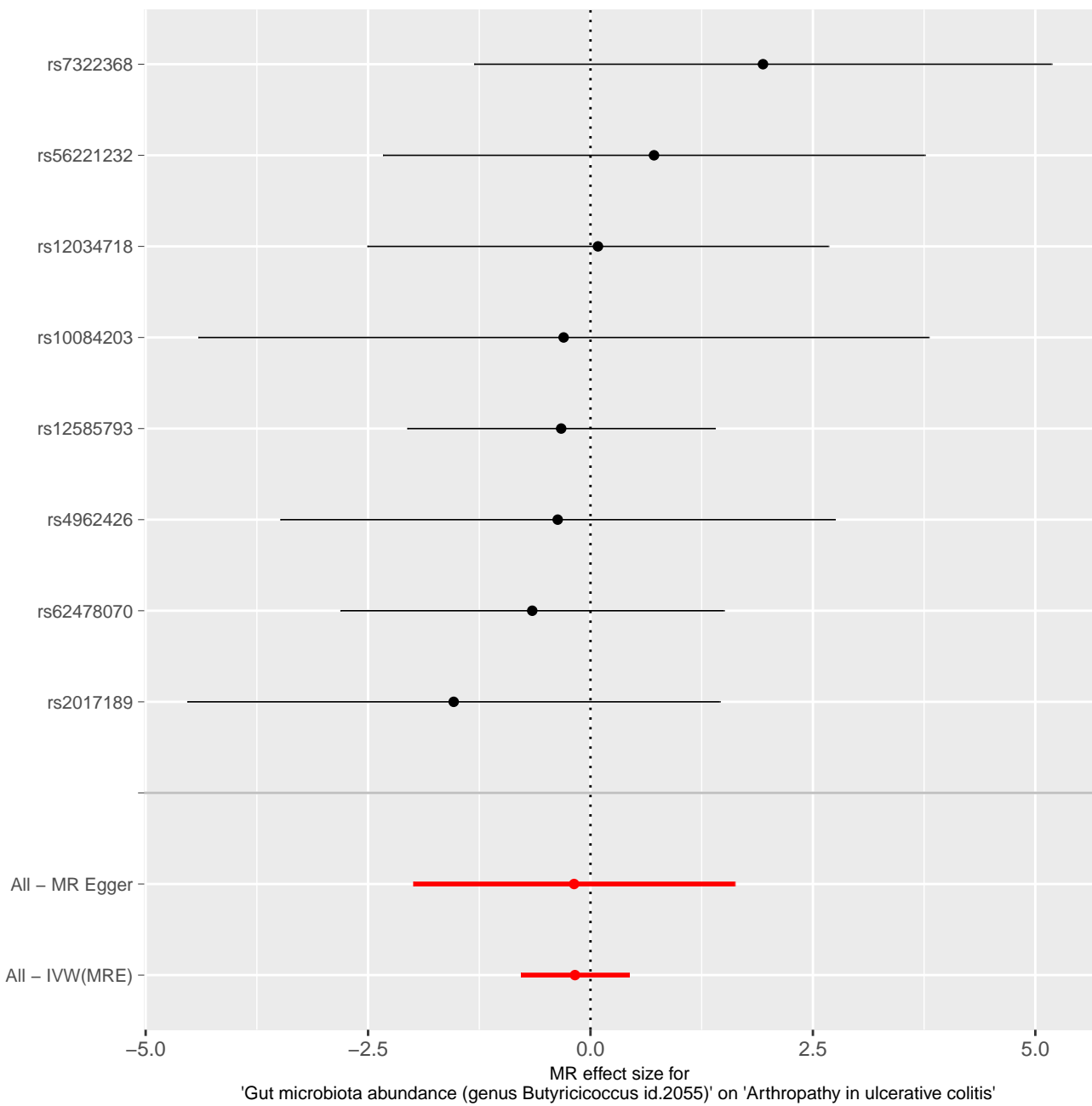
Batch 793 : Gut microbiota abundance (genus *Barnesiella* id.944) on Arthropathy in ulcerative colitis

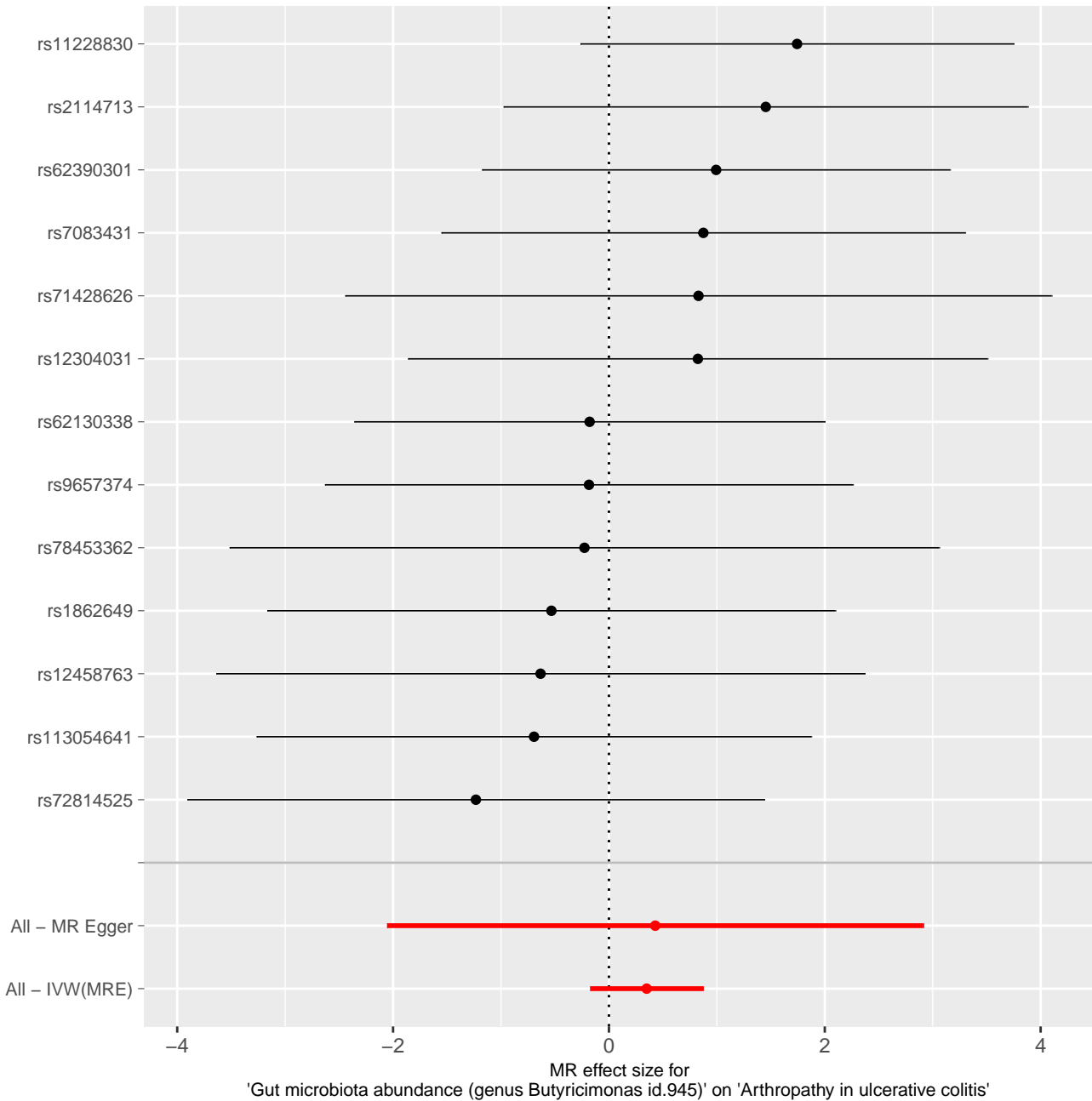




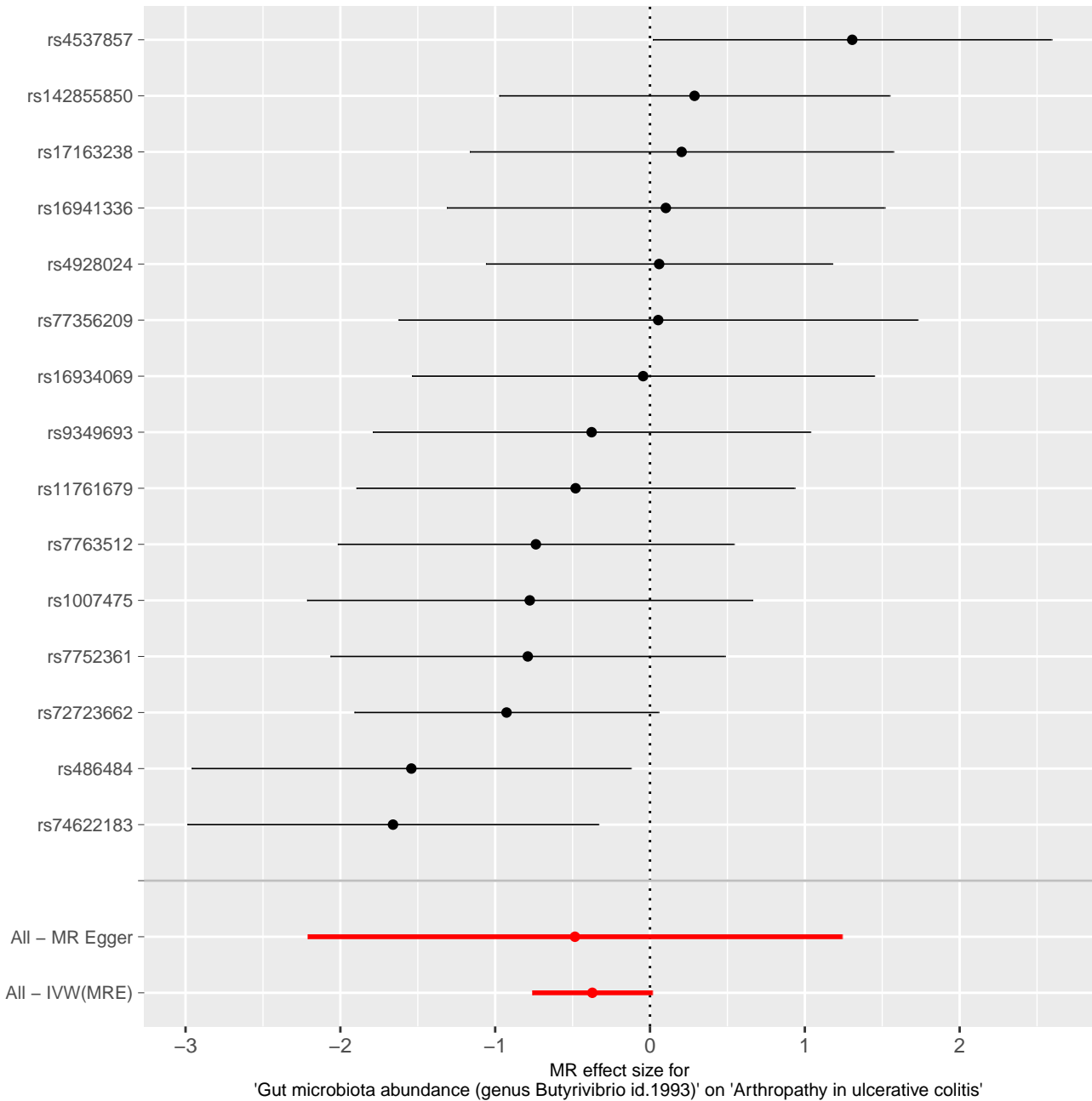
Batch 795 : Gut microbiota abundance (genus Bilophila id.3170) on Arthropathy in ulcerative colitis

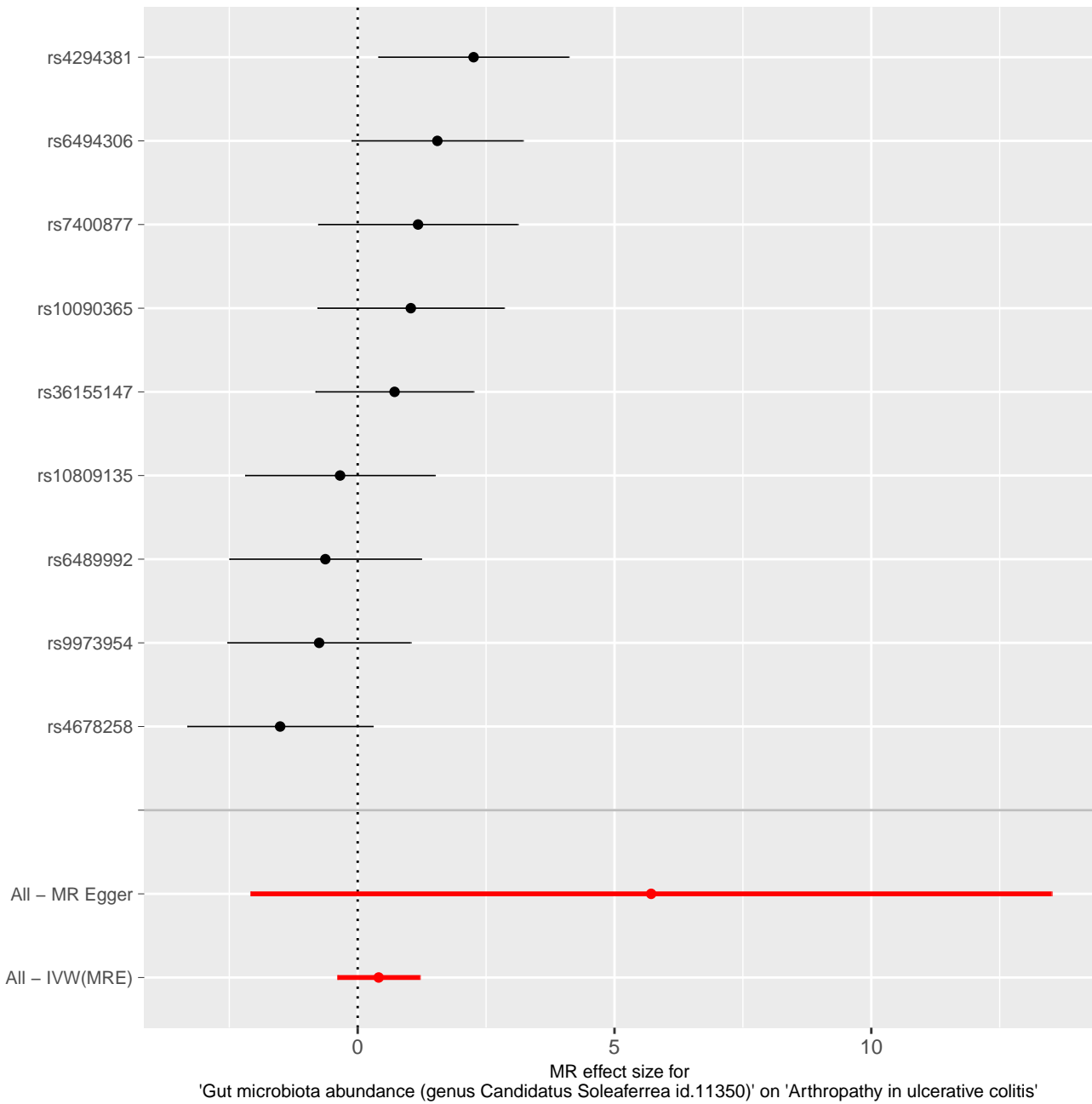


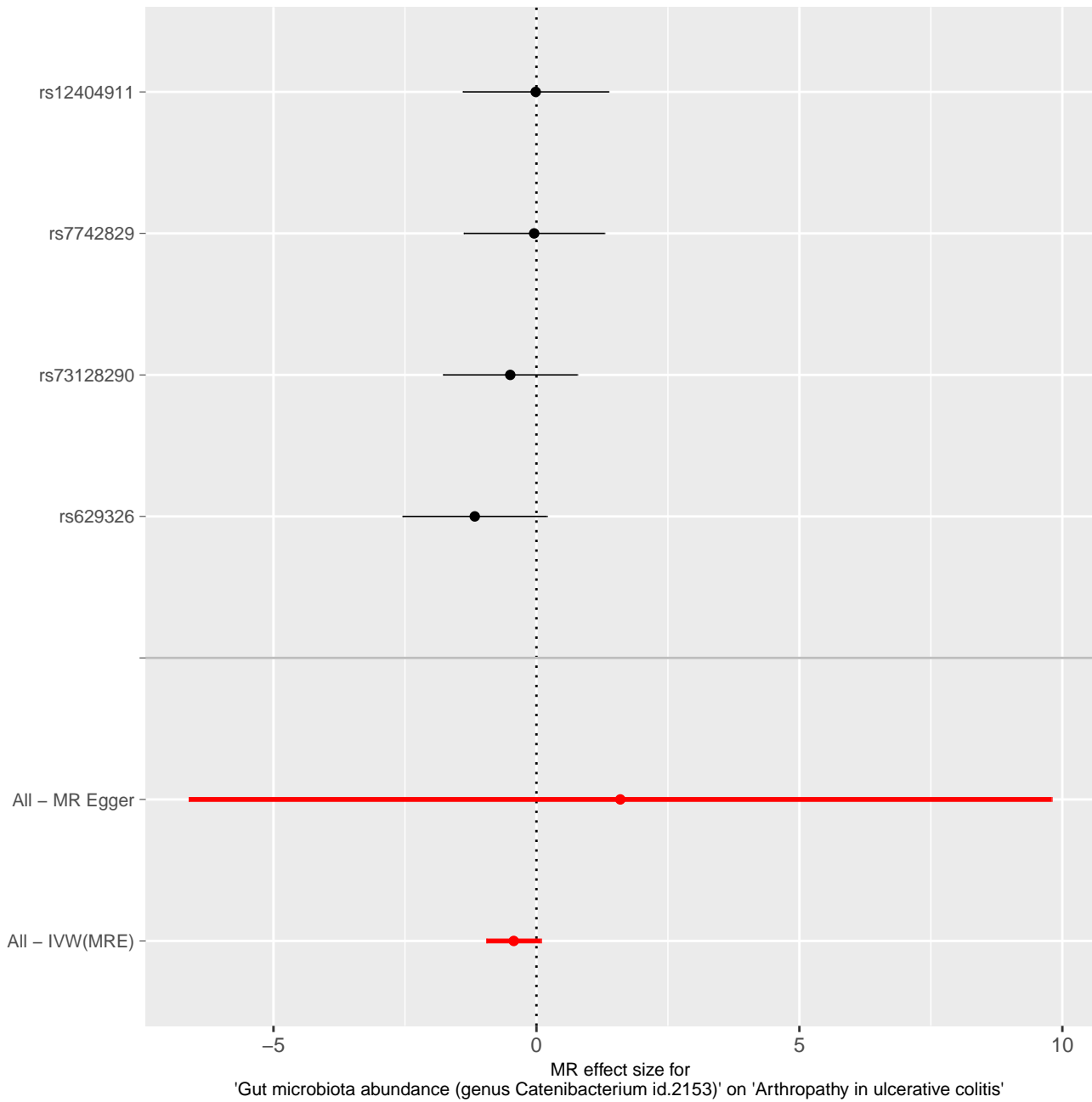


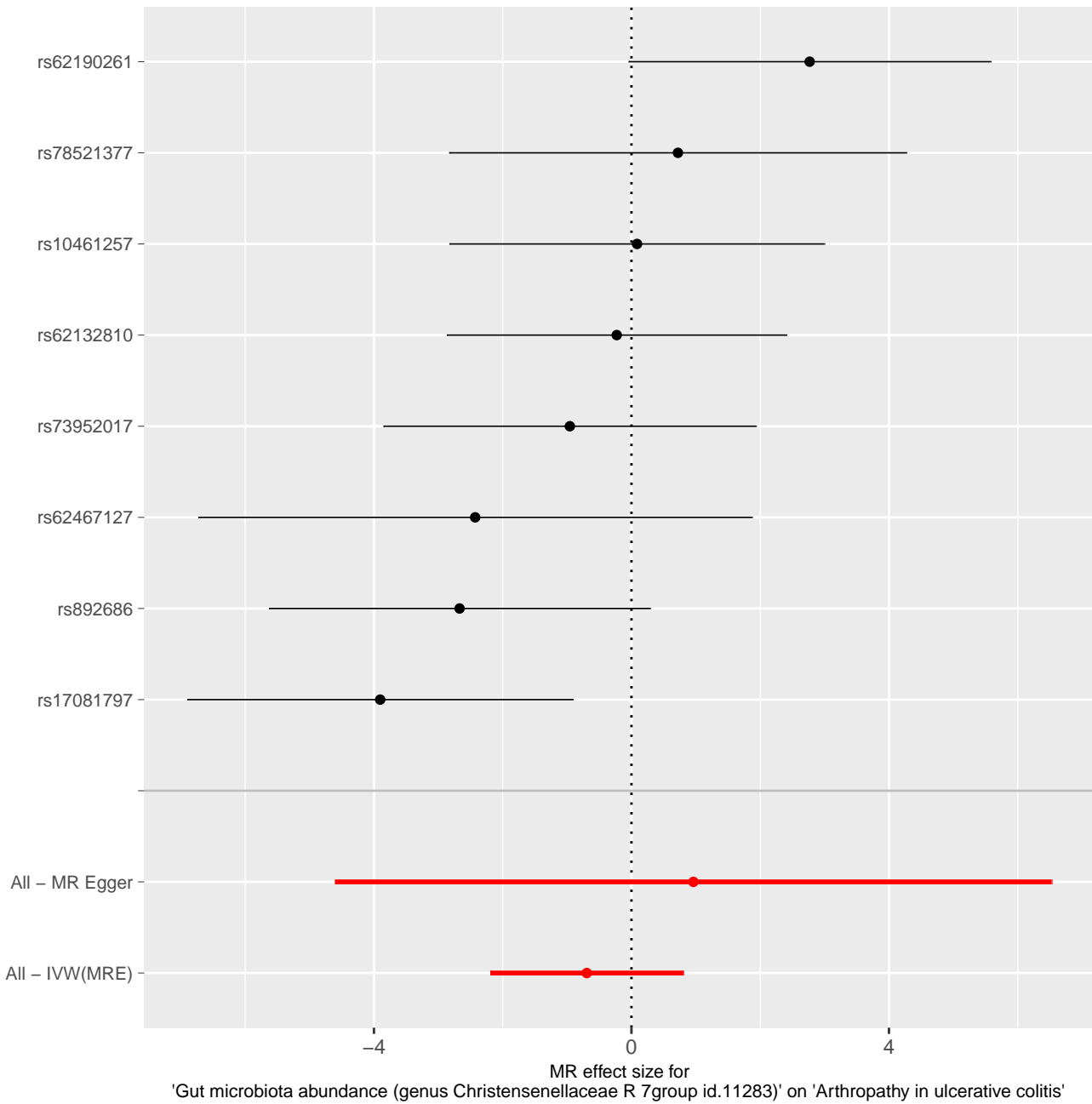


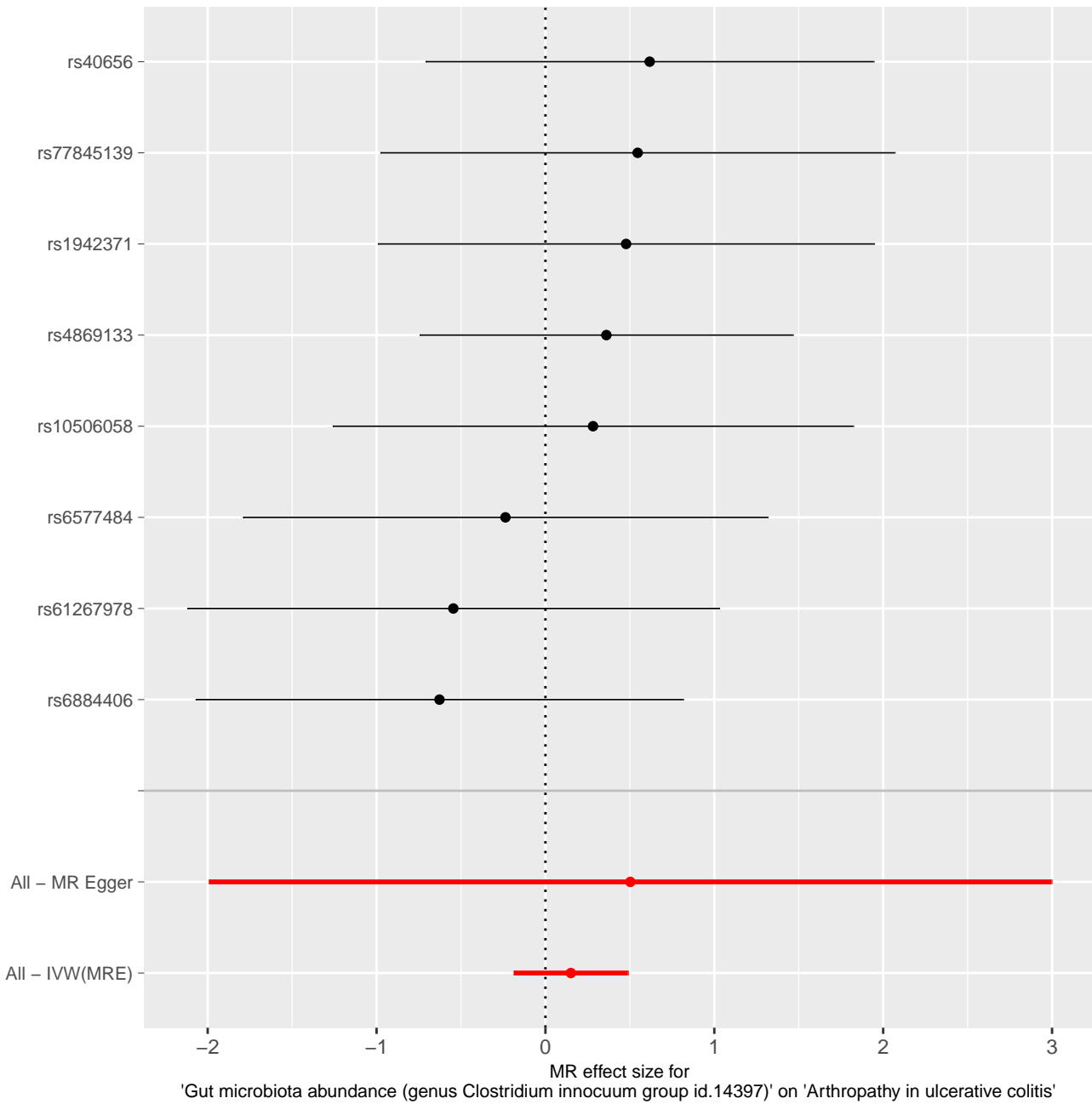
Batch 799 : Gut microbiota abundance (genus Butyrivibrio id.1993) on Arthropathy in ulcerative colitis

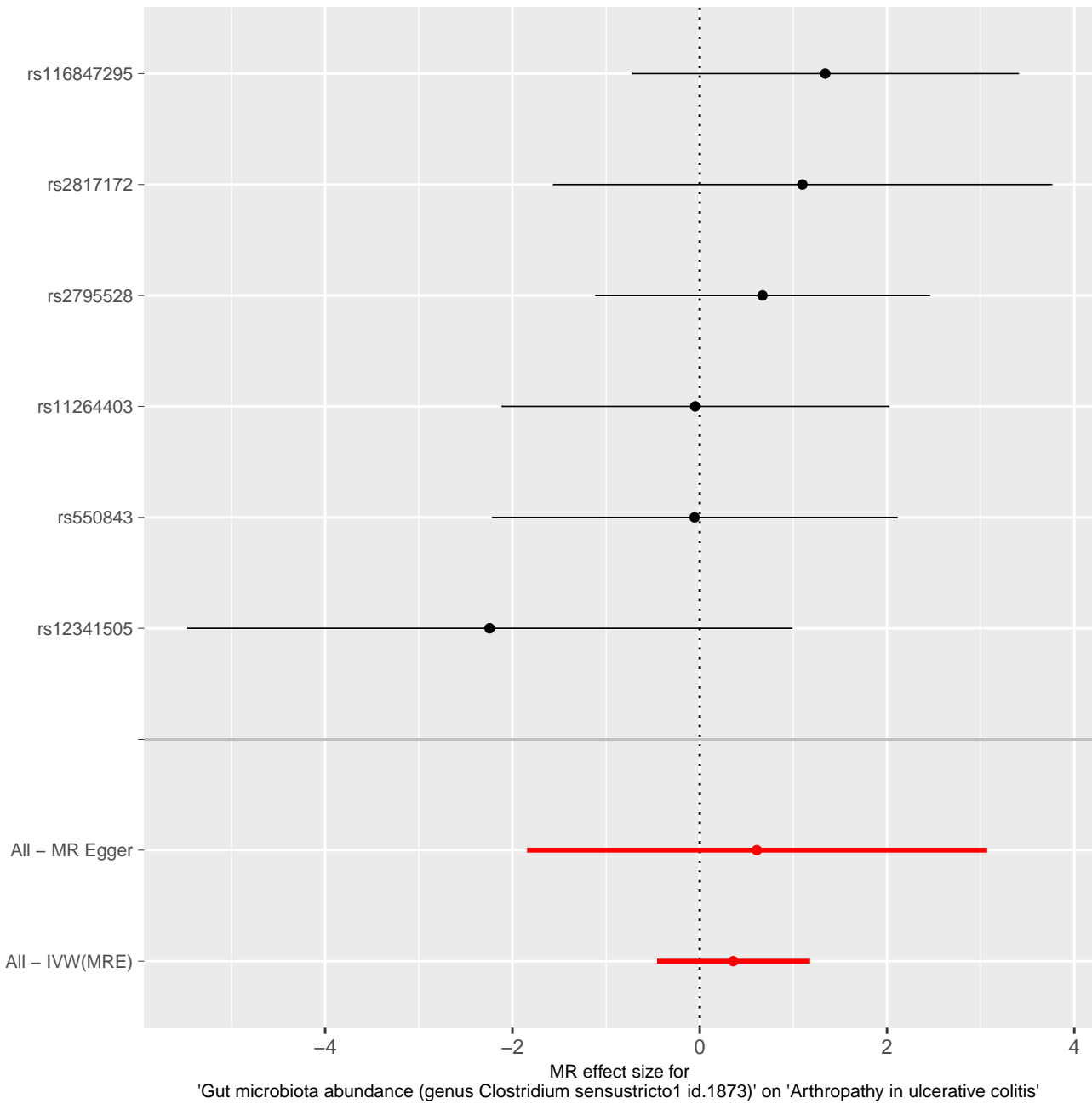


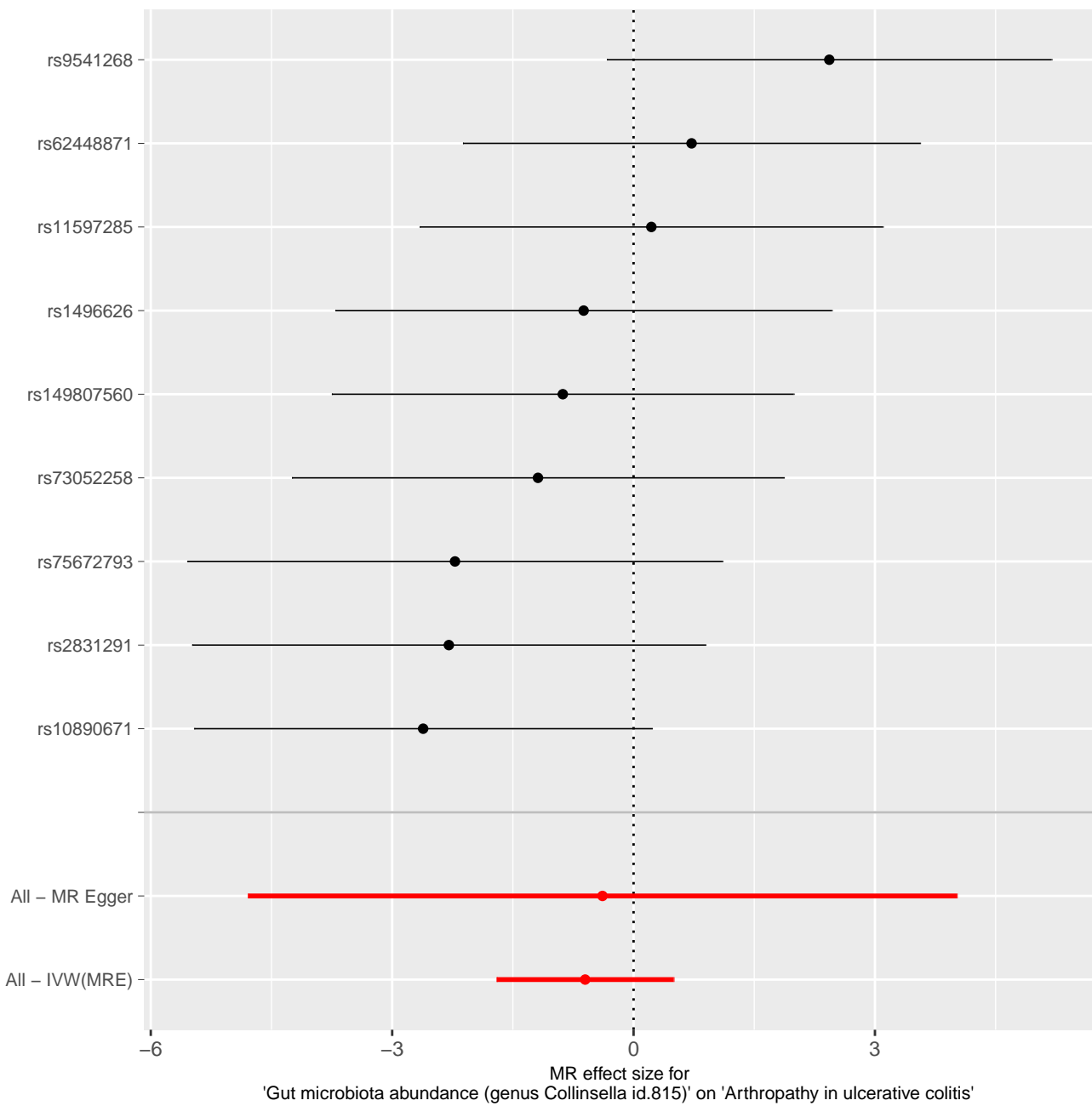


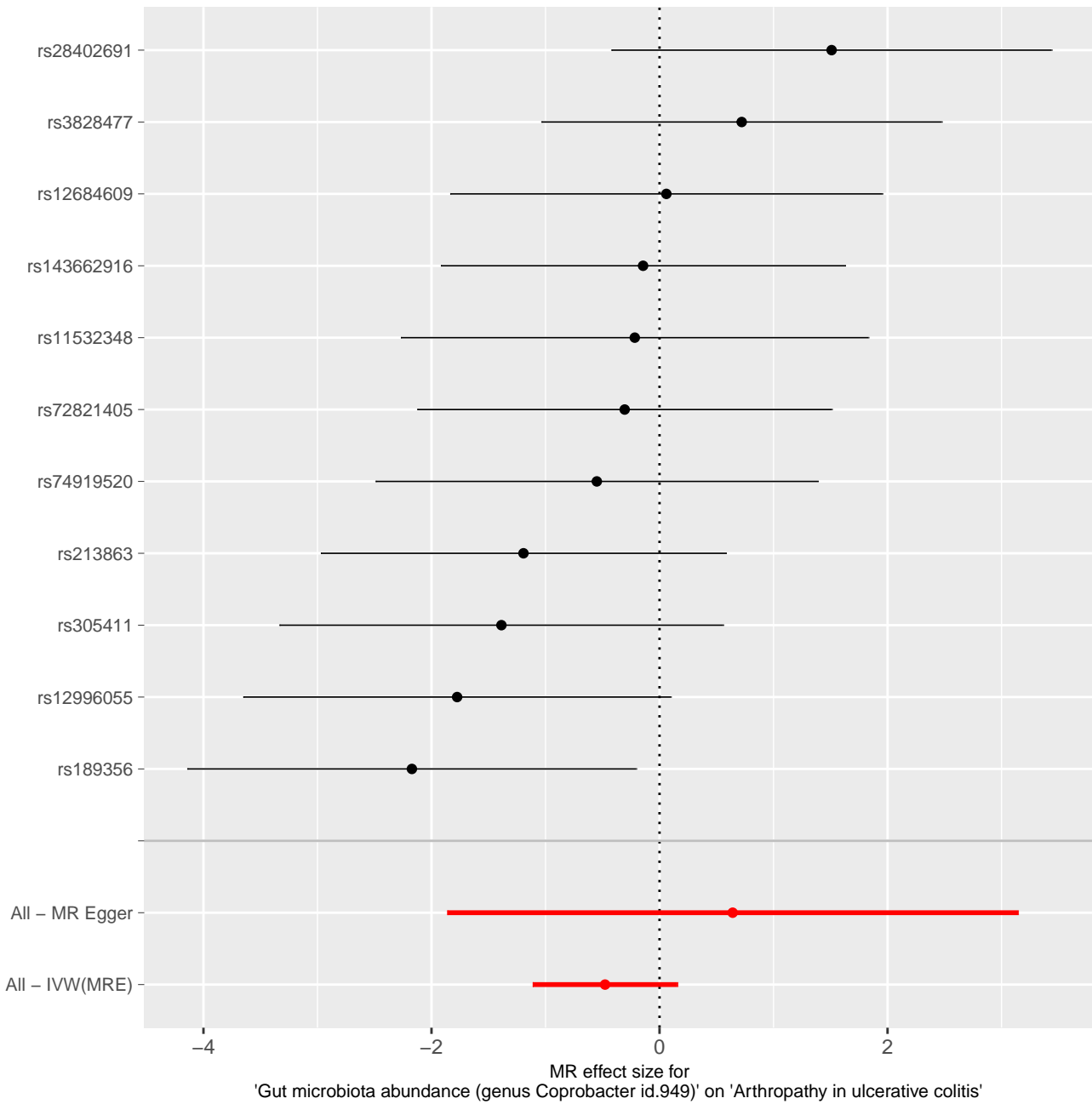


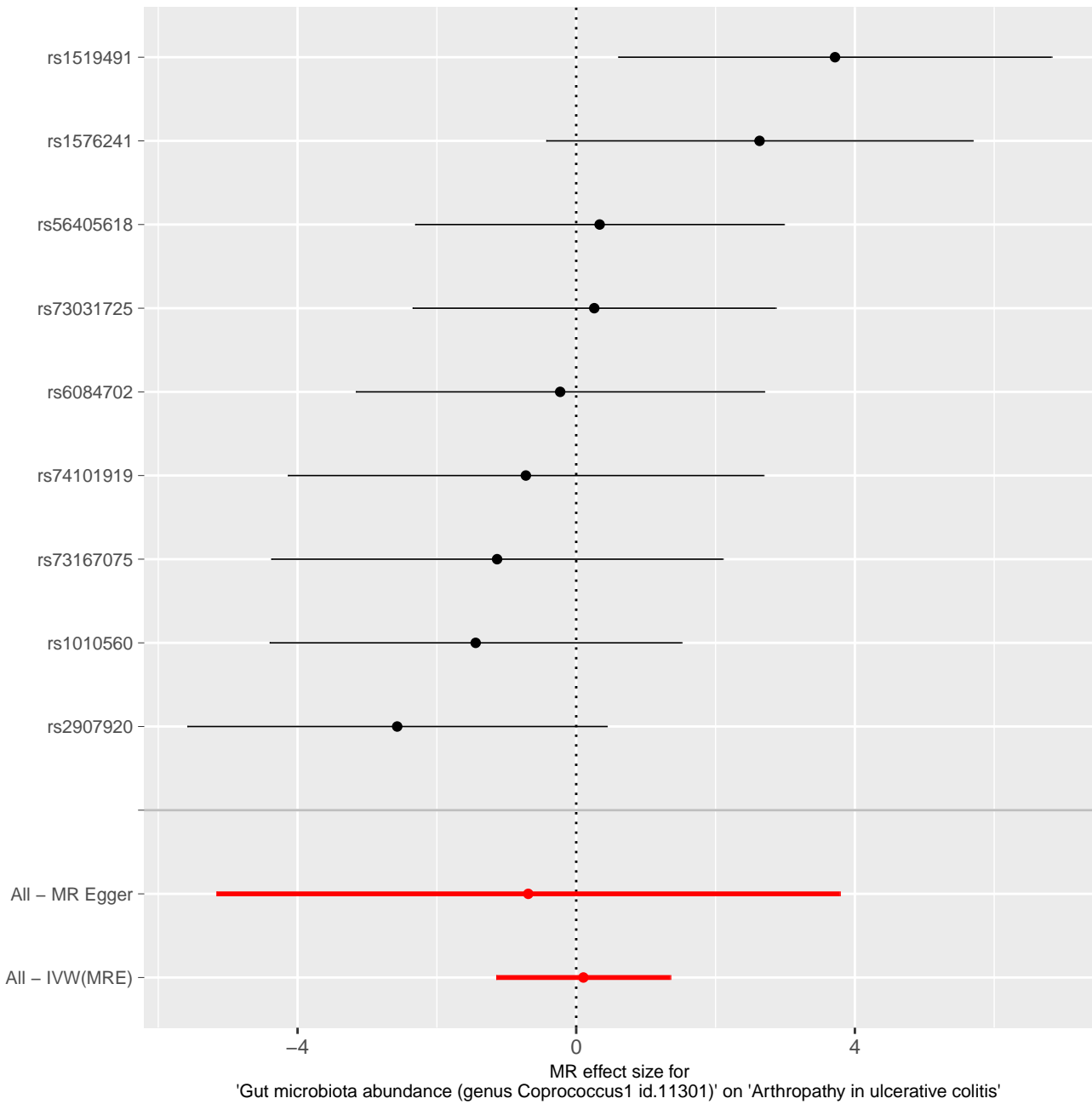


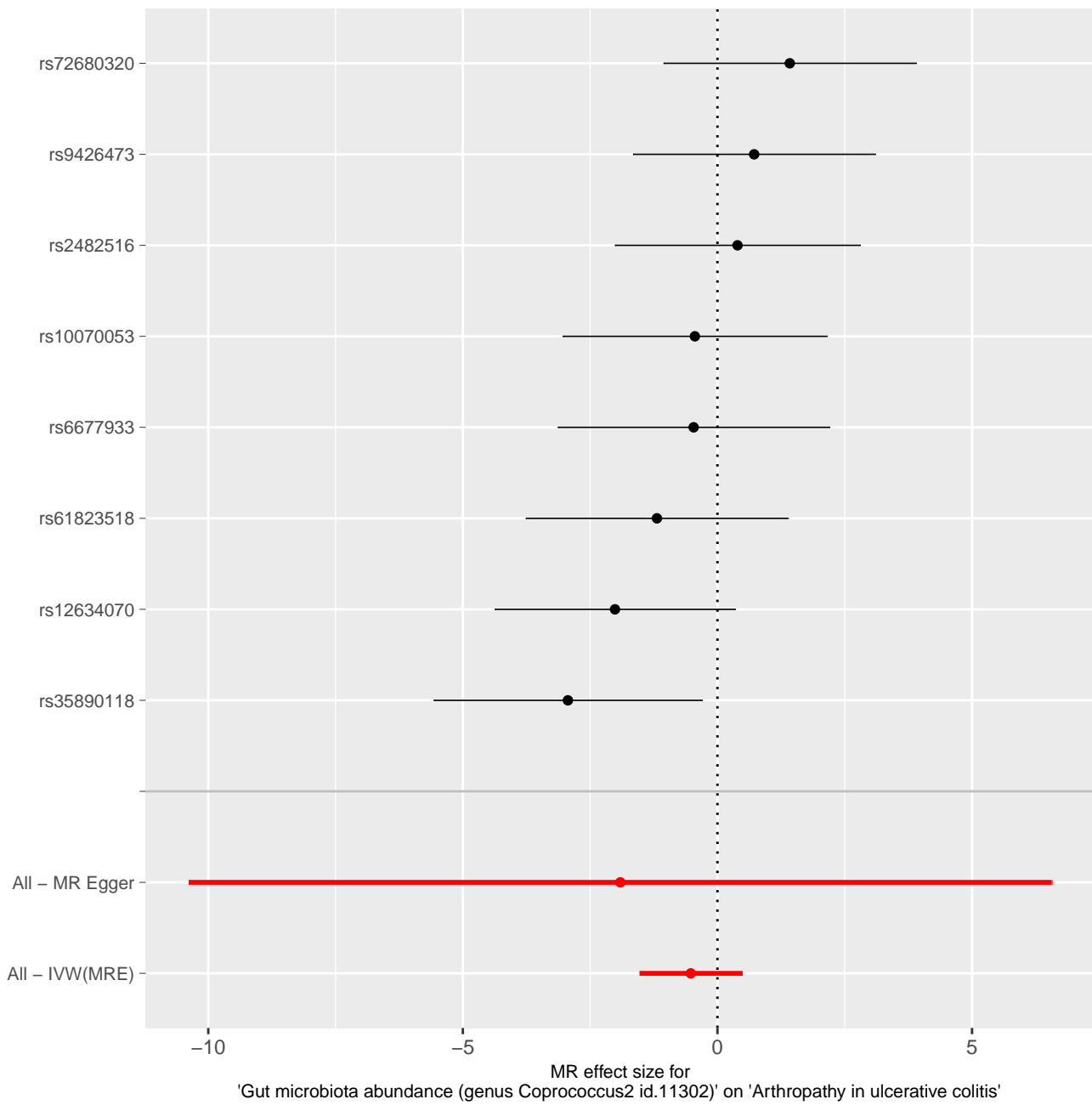


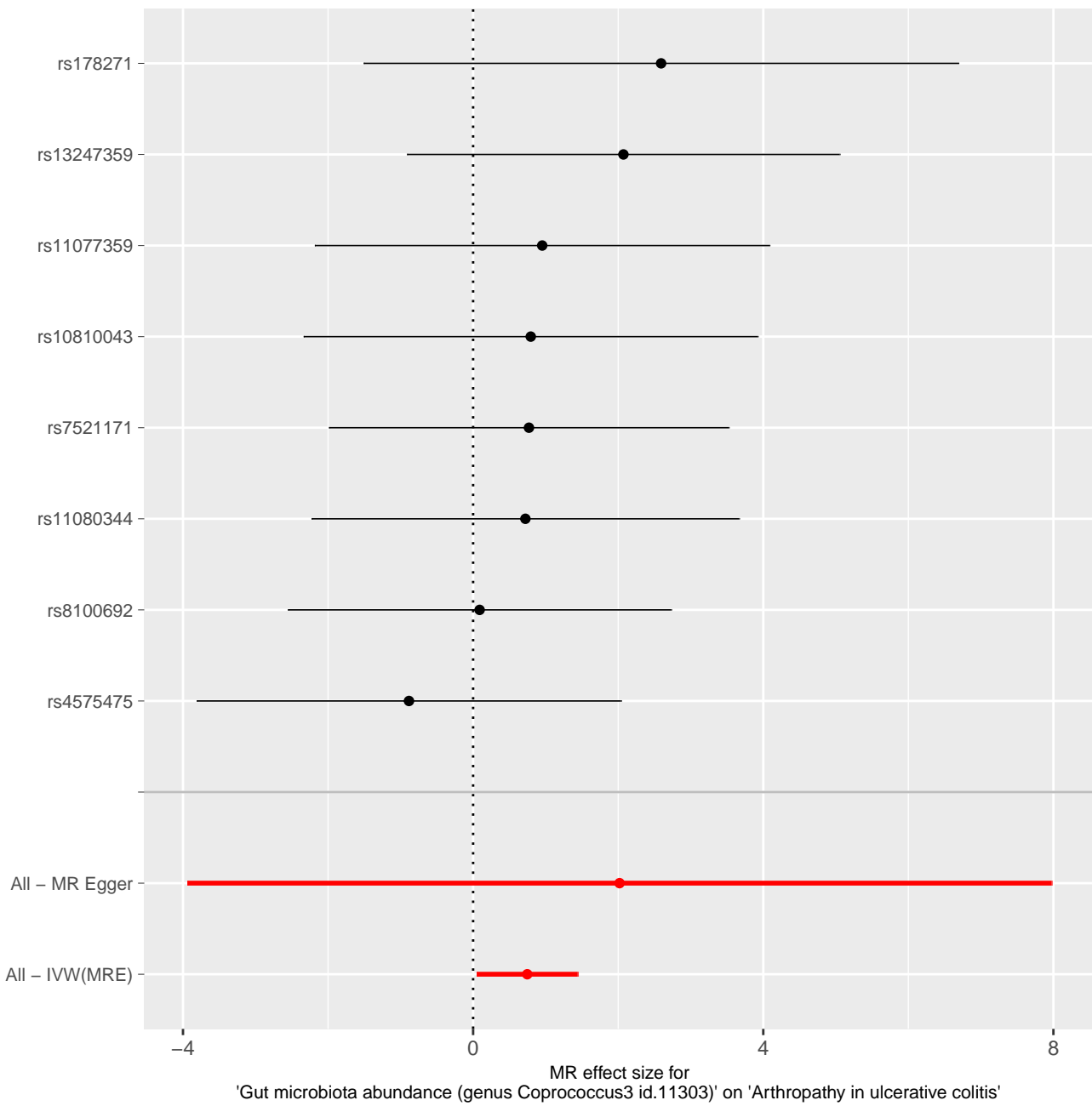


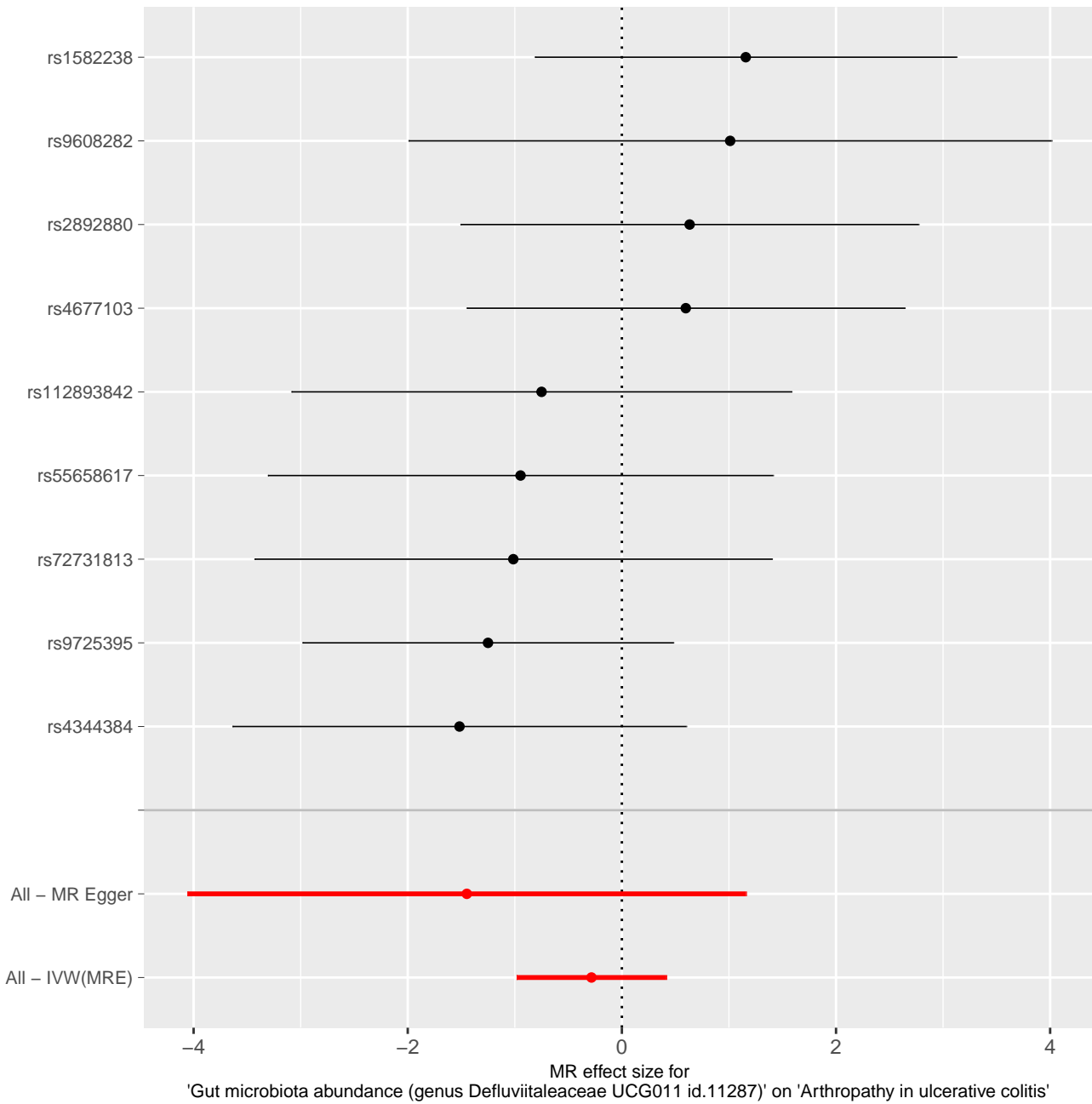


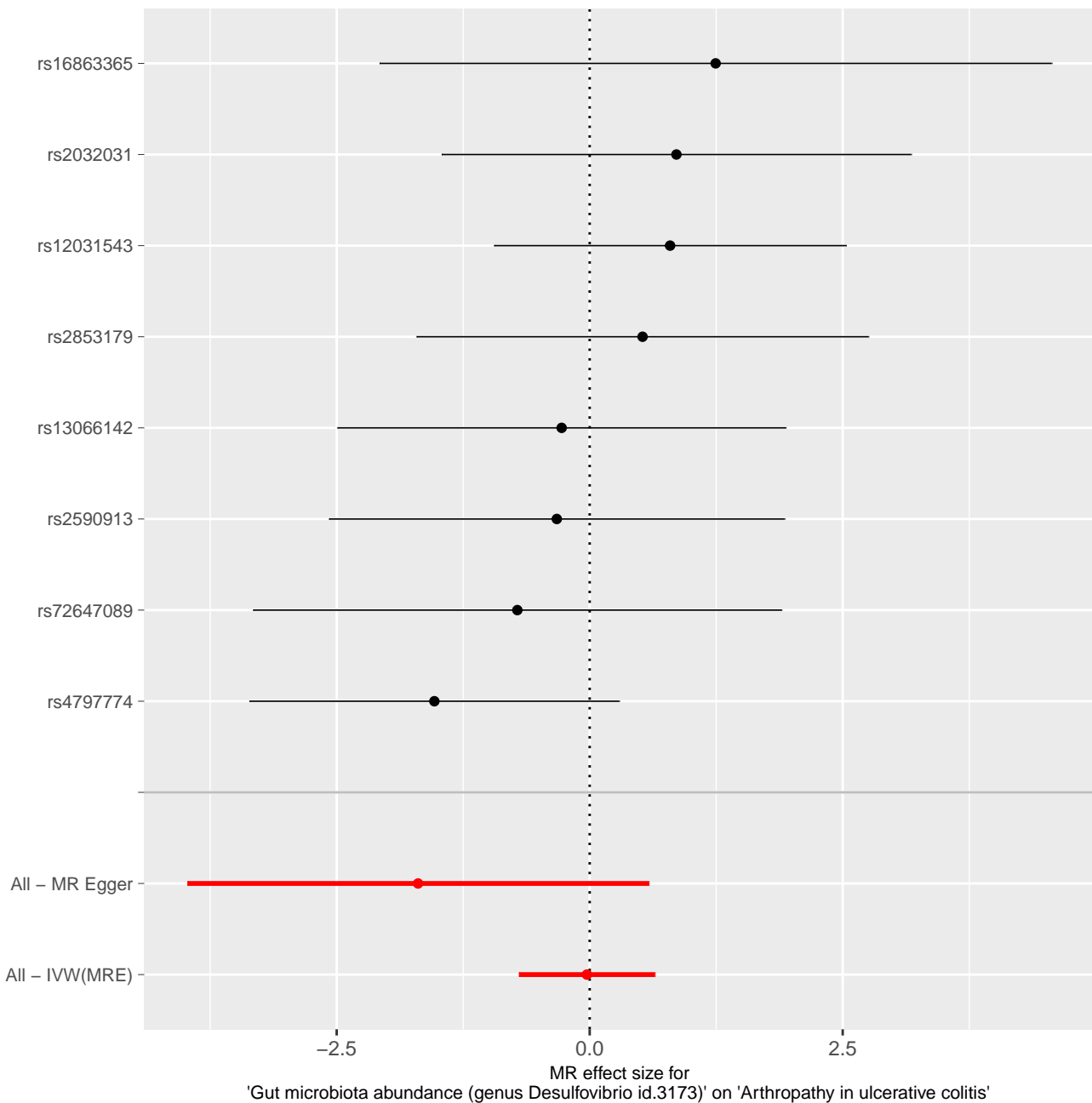


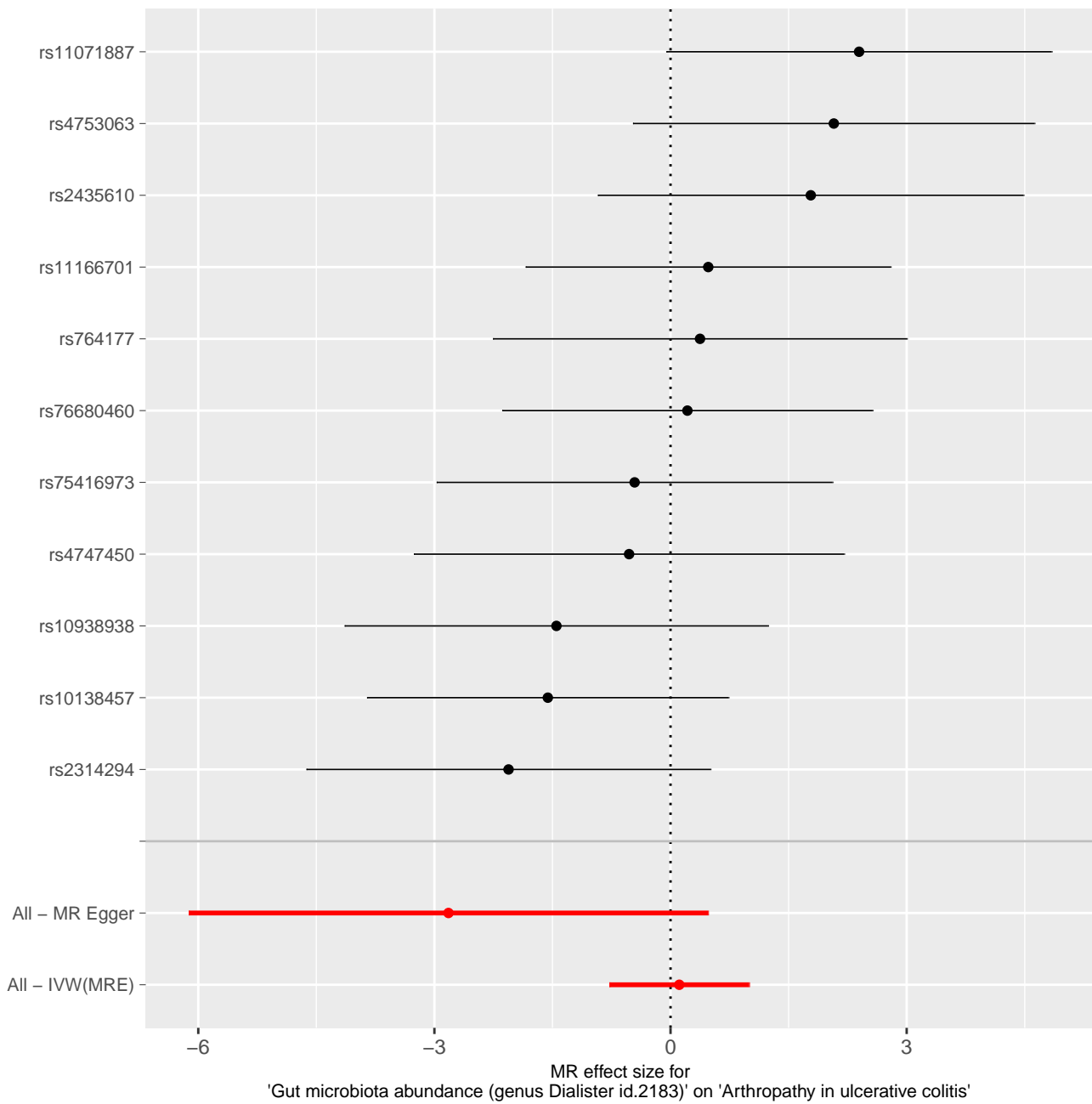


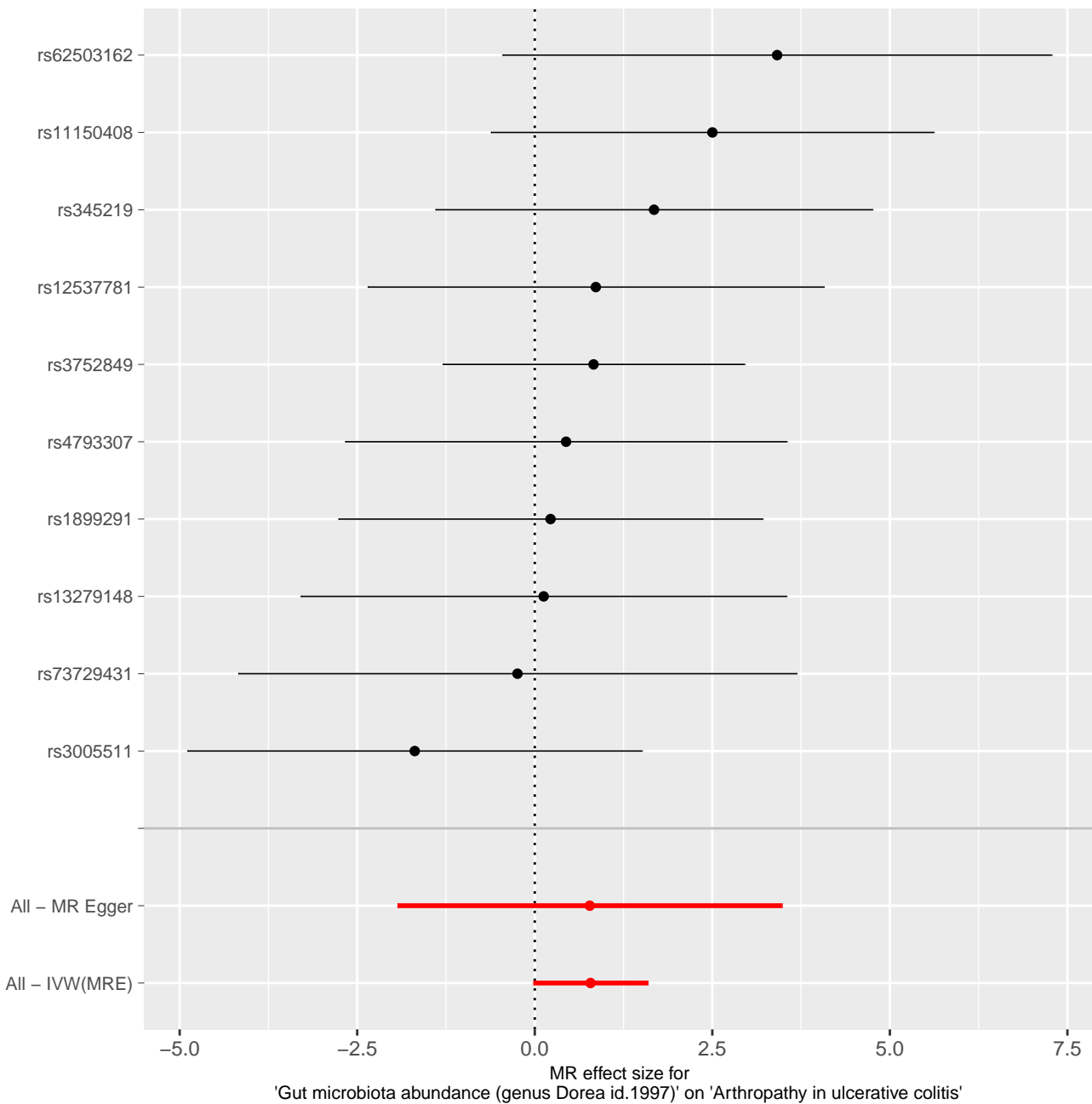




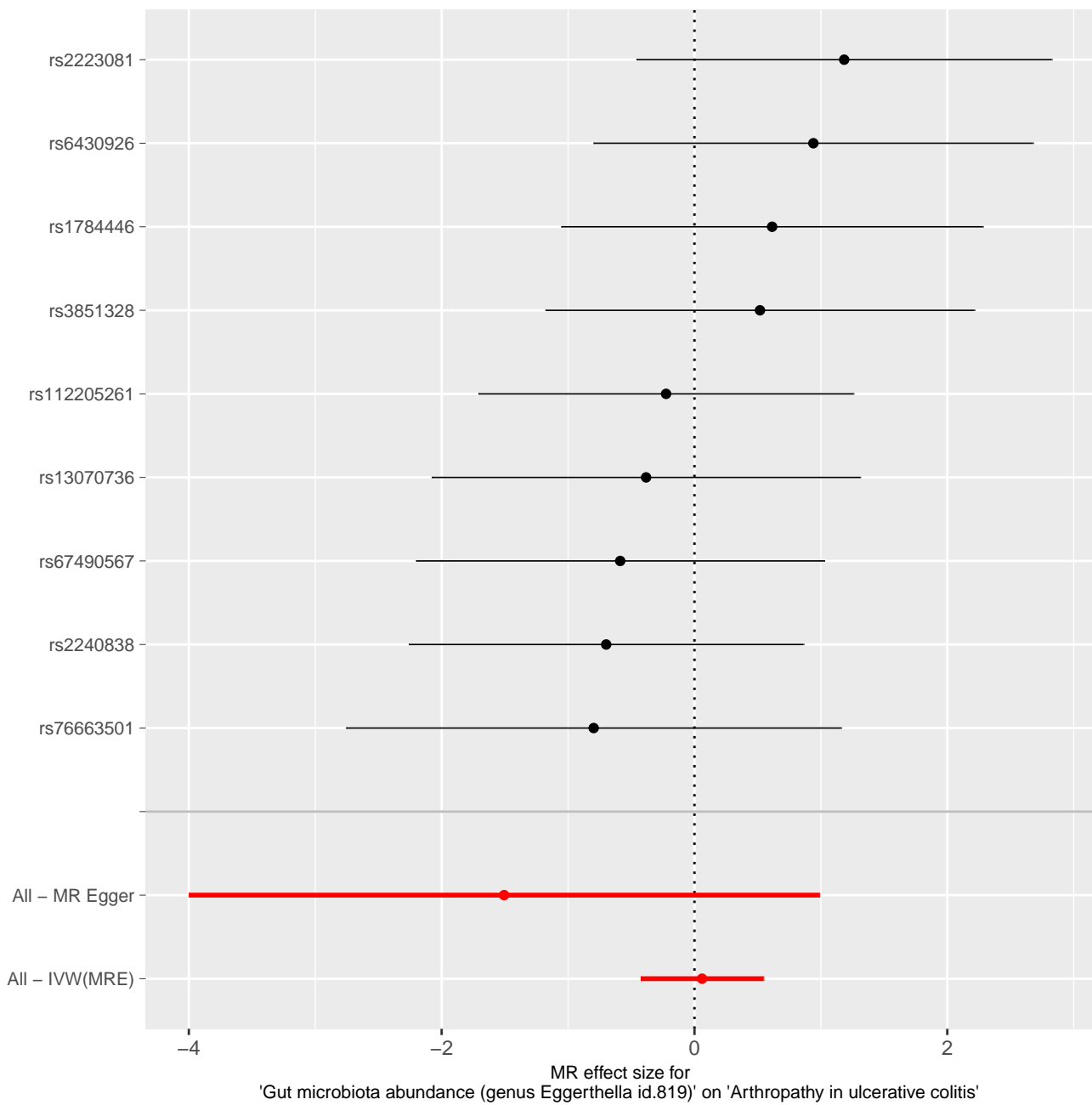


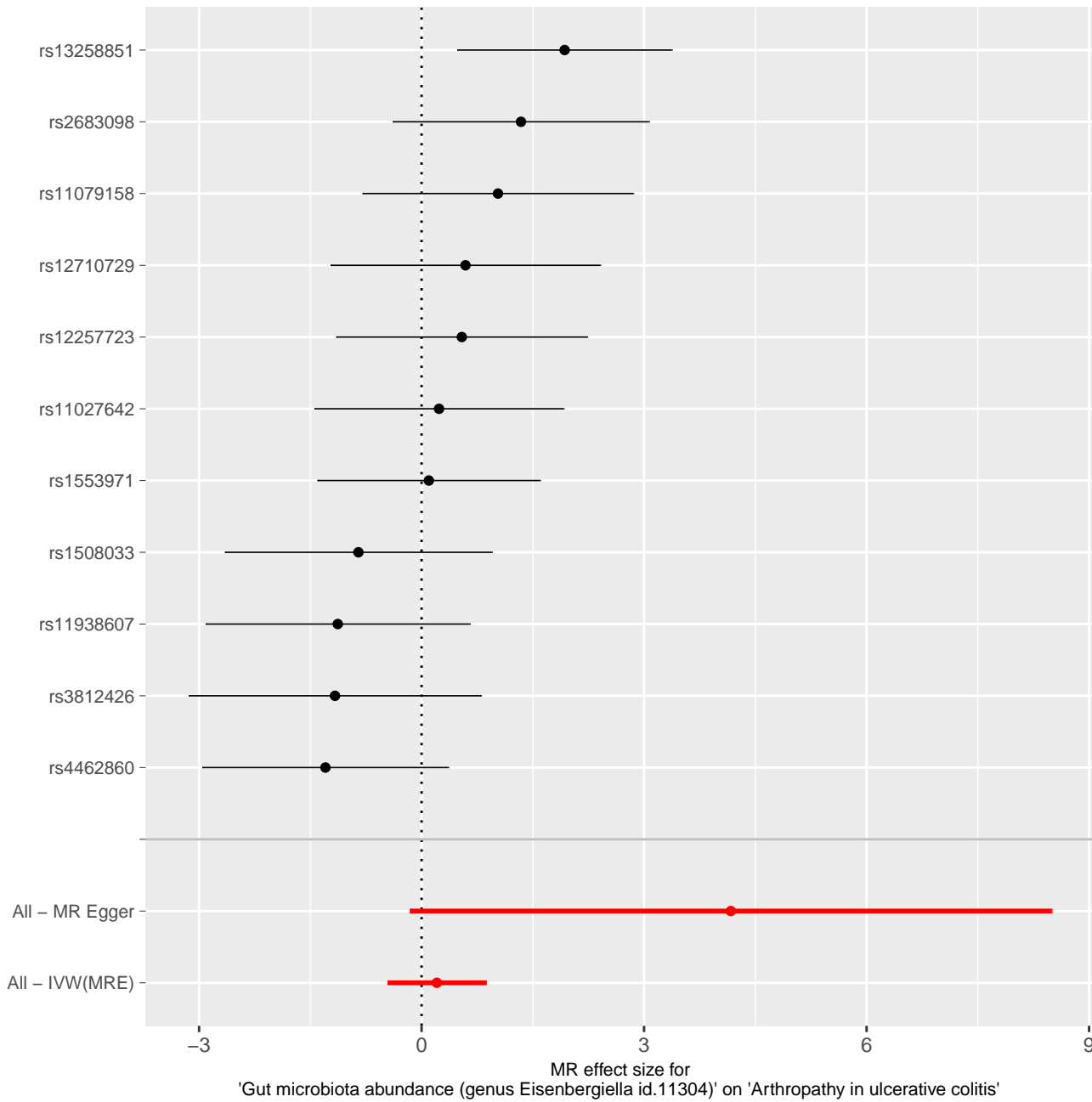


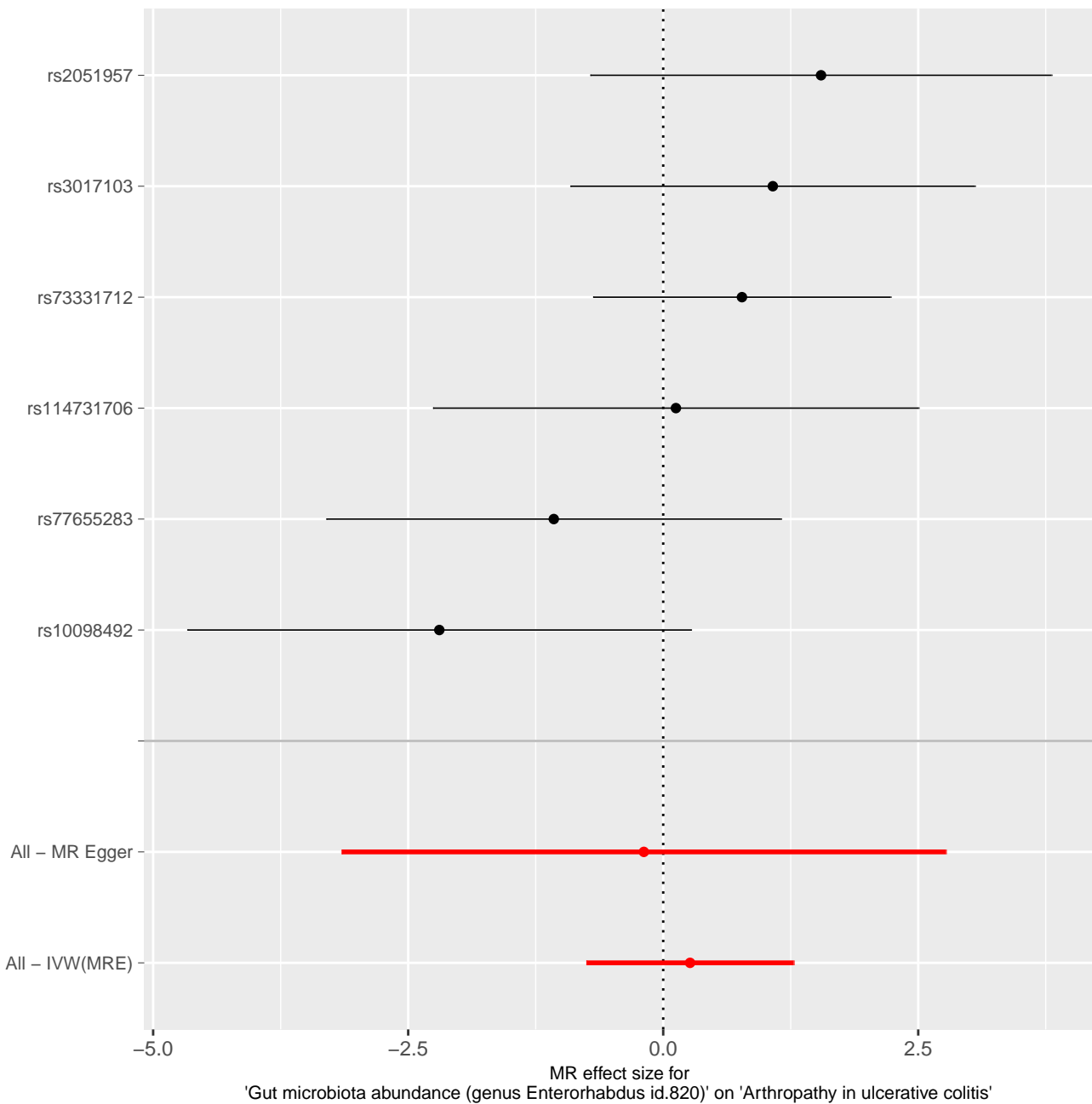




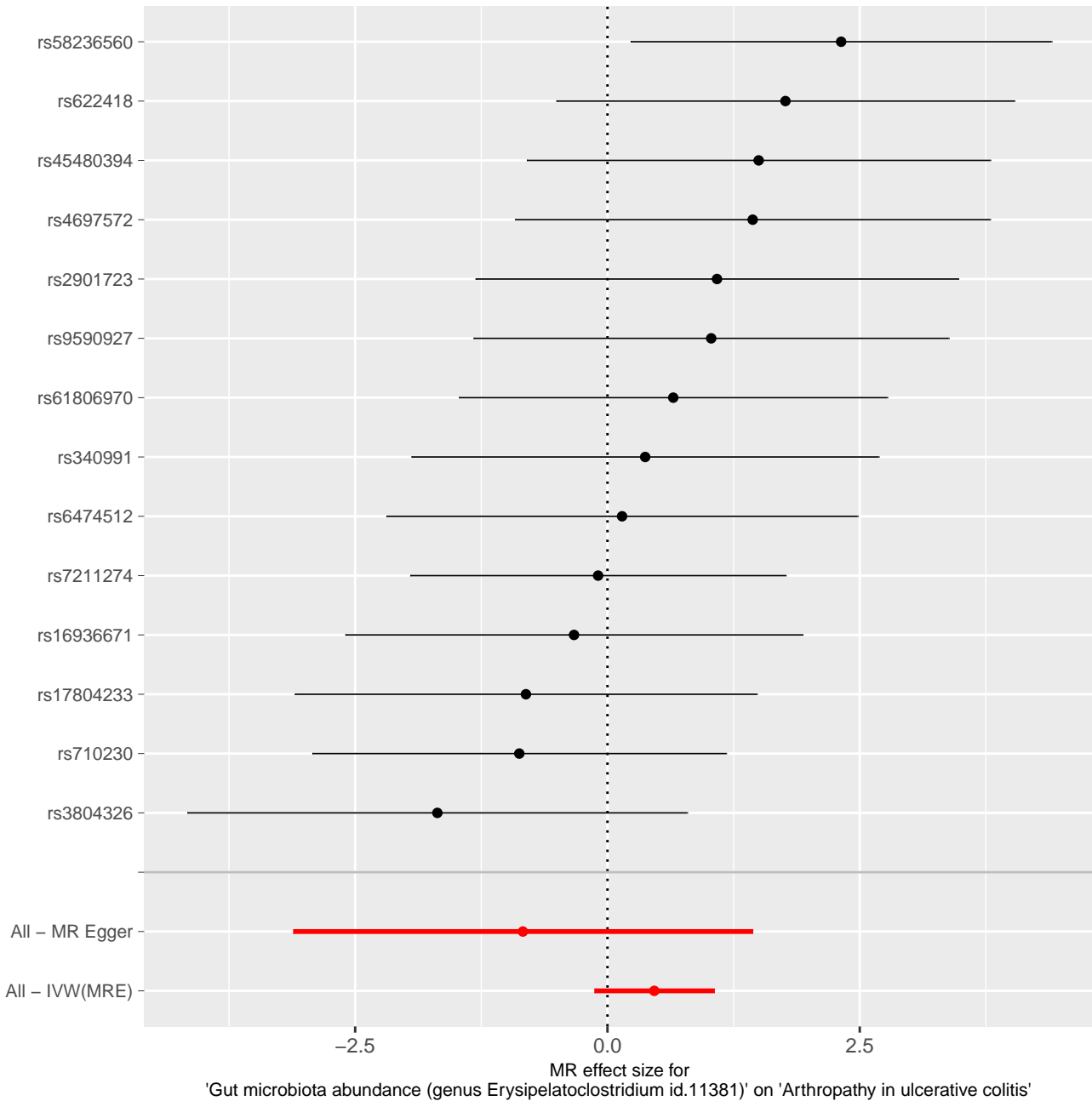
Batch 814 : Gut microbiota abundance (genus Eggerthella id.819) on Arthropathy in ulcerative colitis

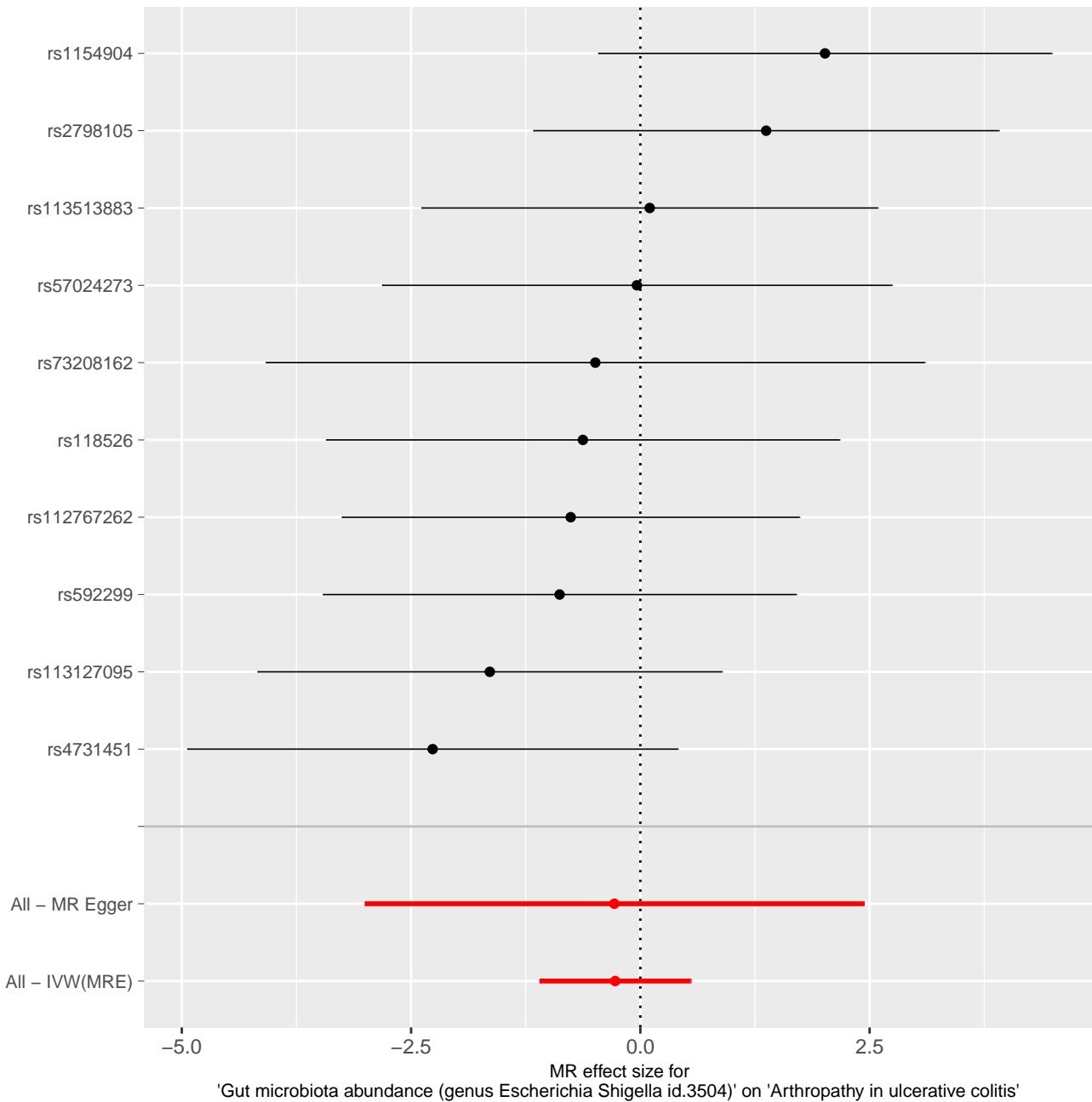


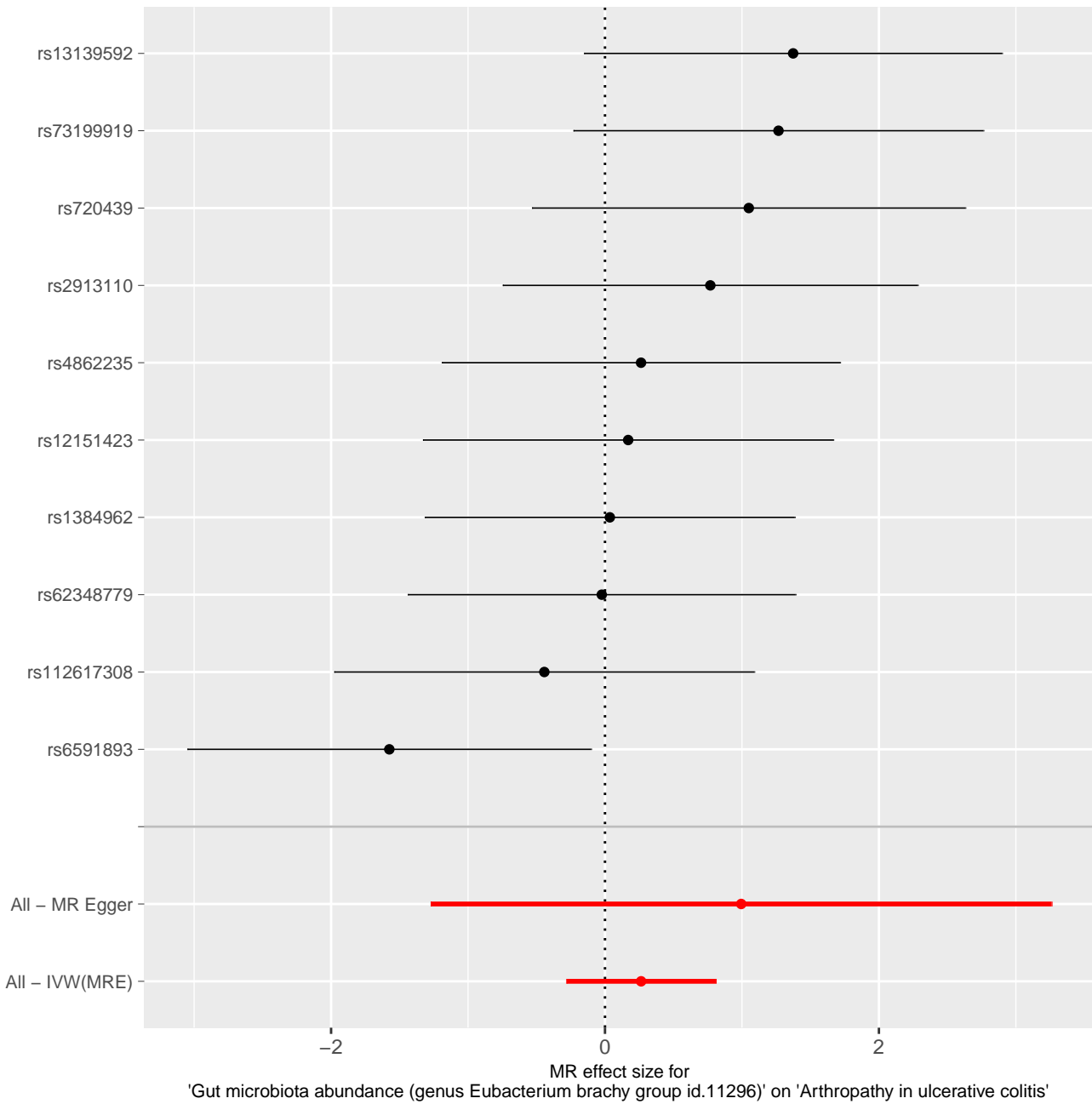


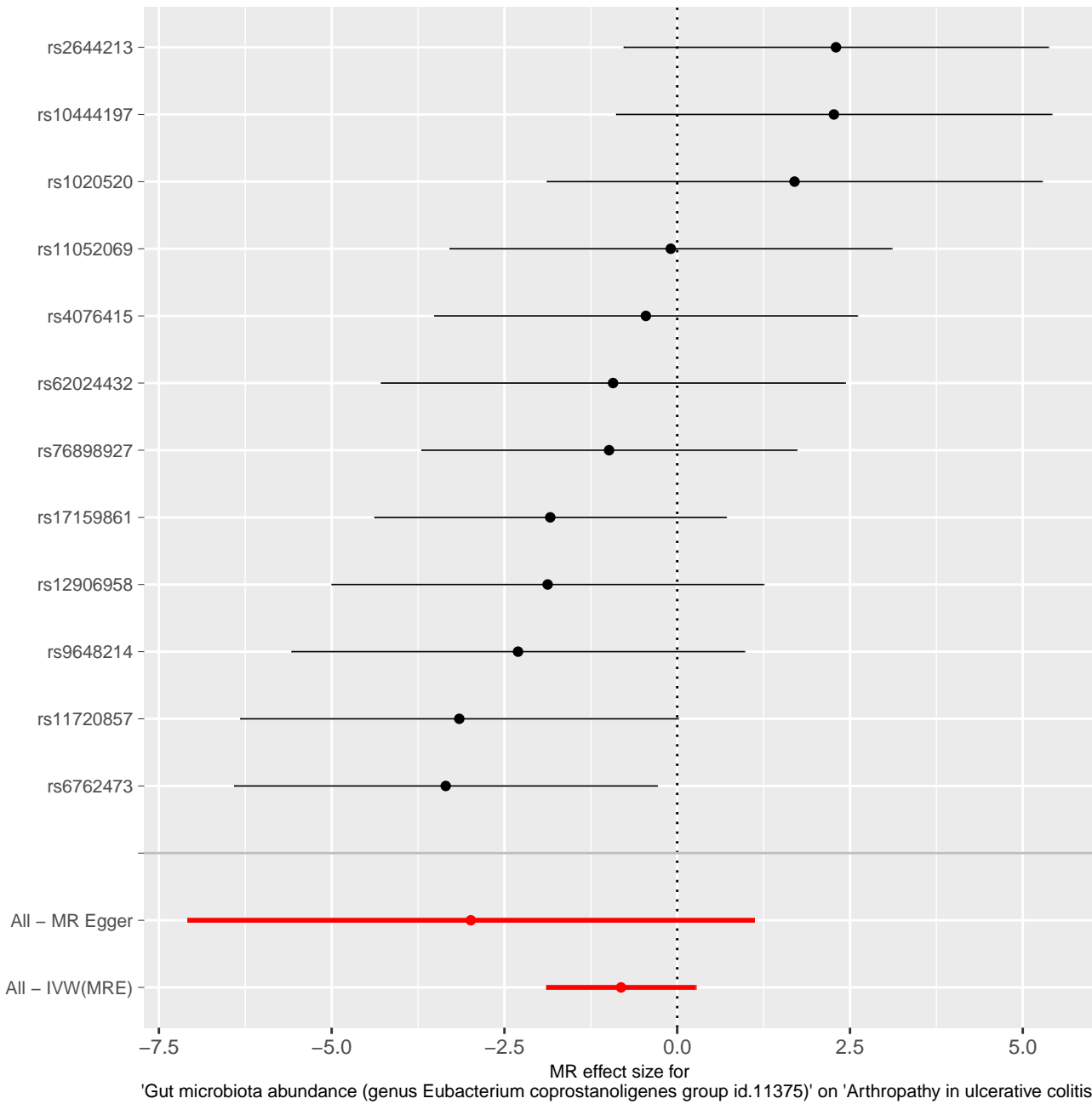


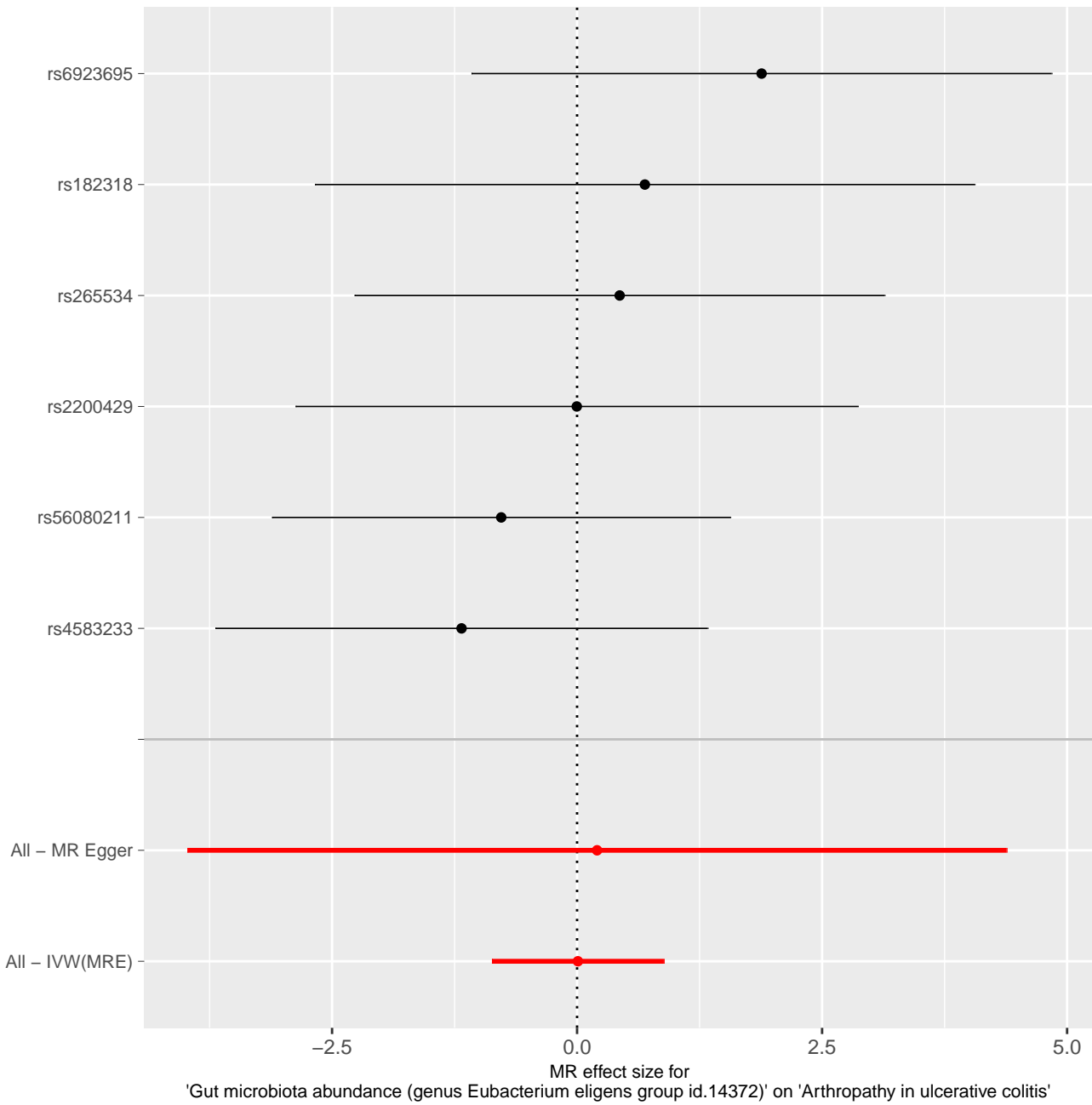
Batch 817 : Gut microbiota abundance (genus Erysipelatoclostridium id.11381) on Arthropathy in ulcerative colitis

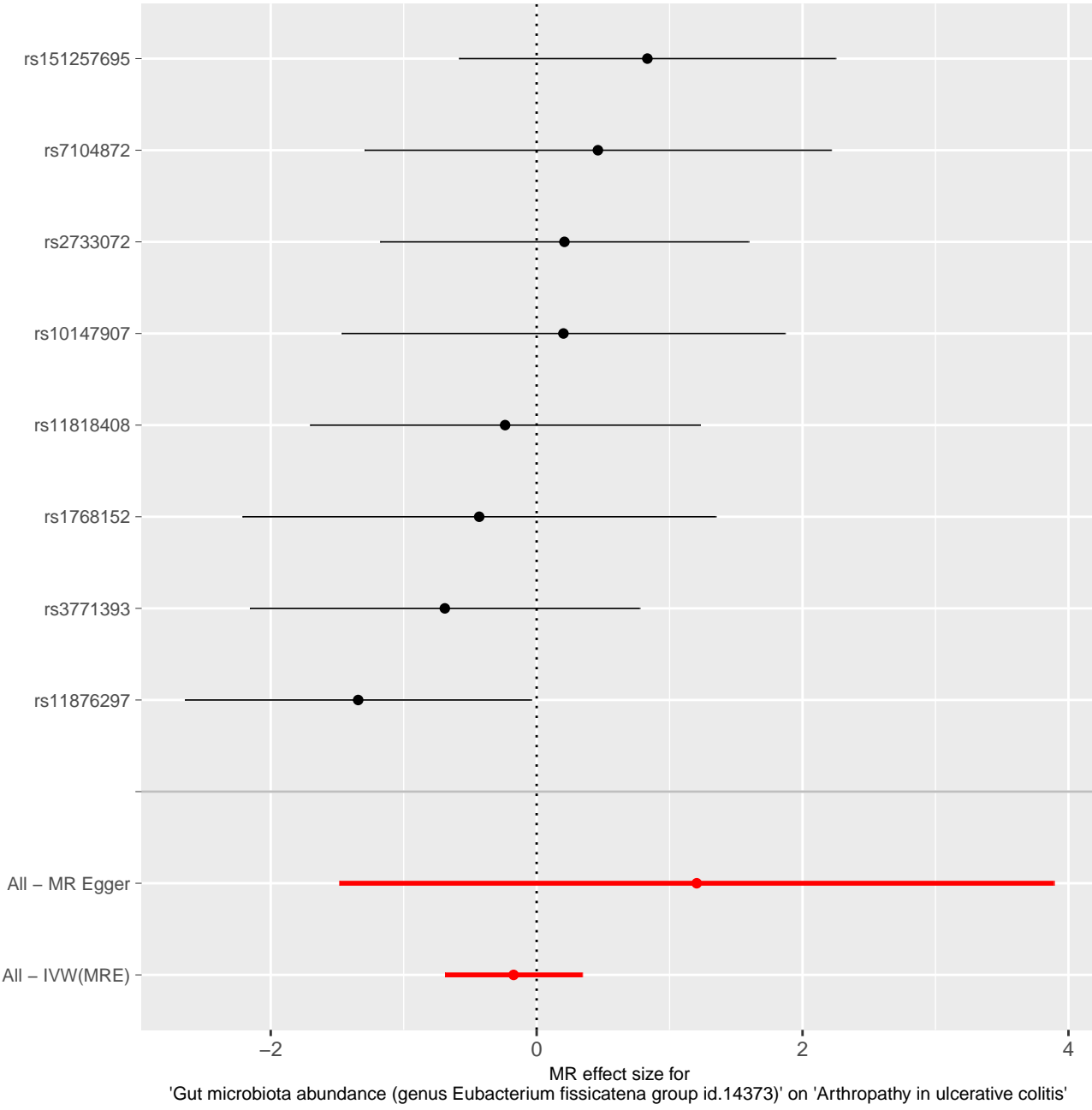


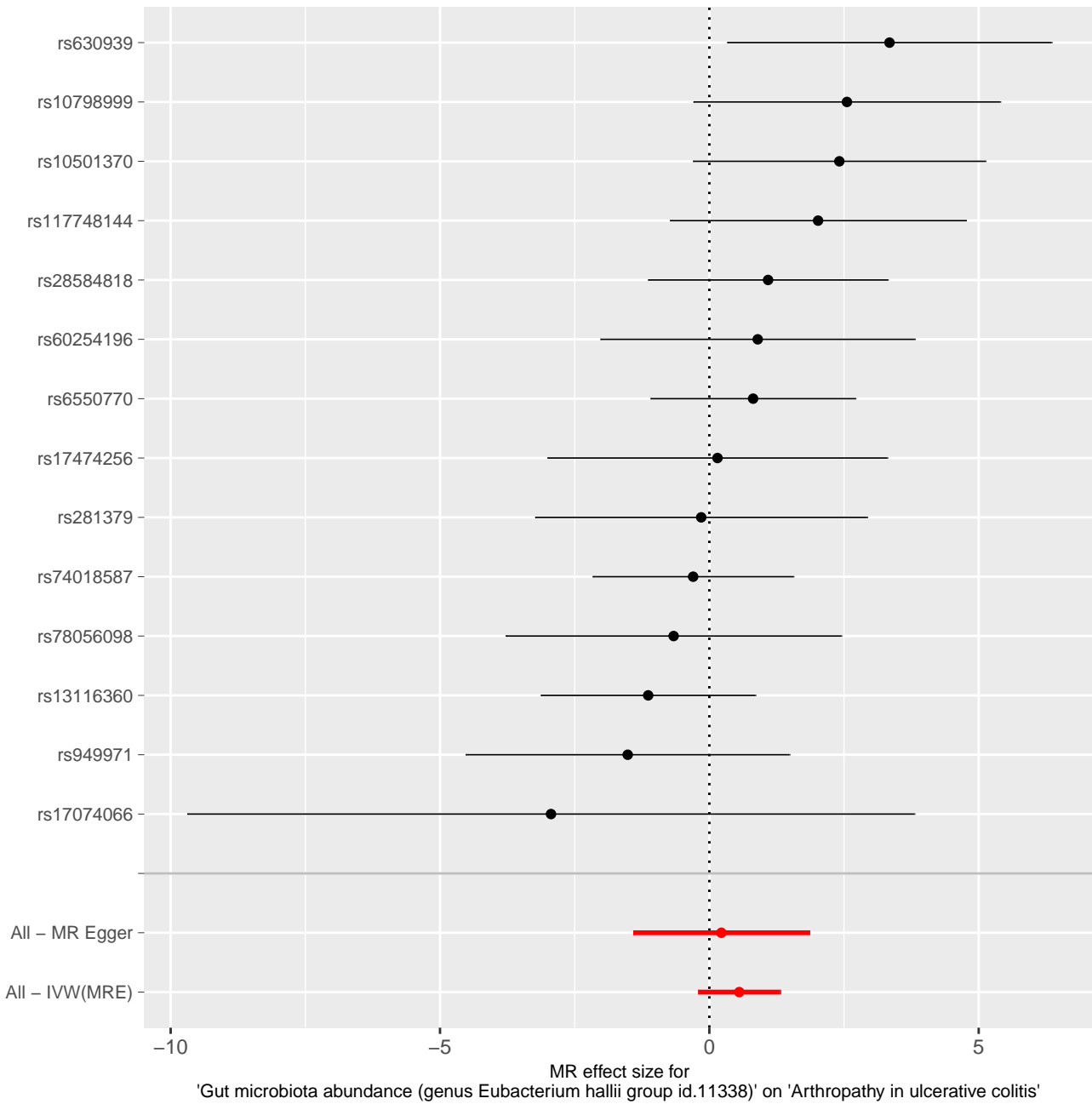




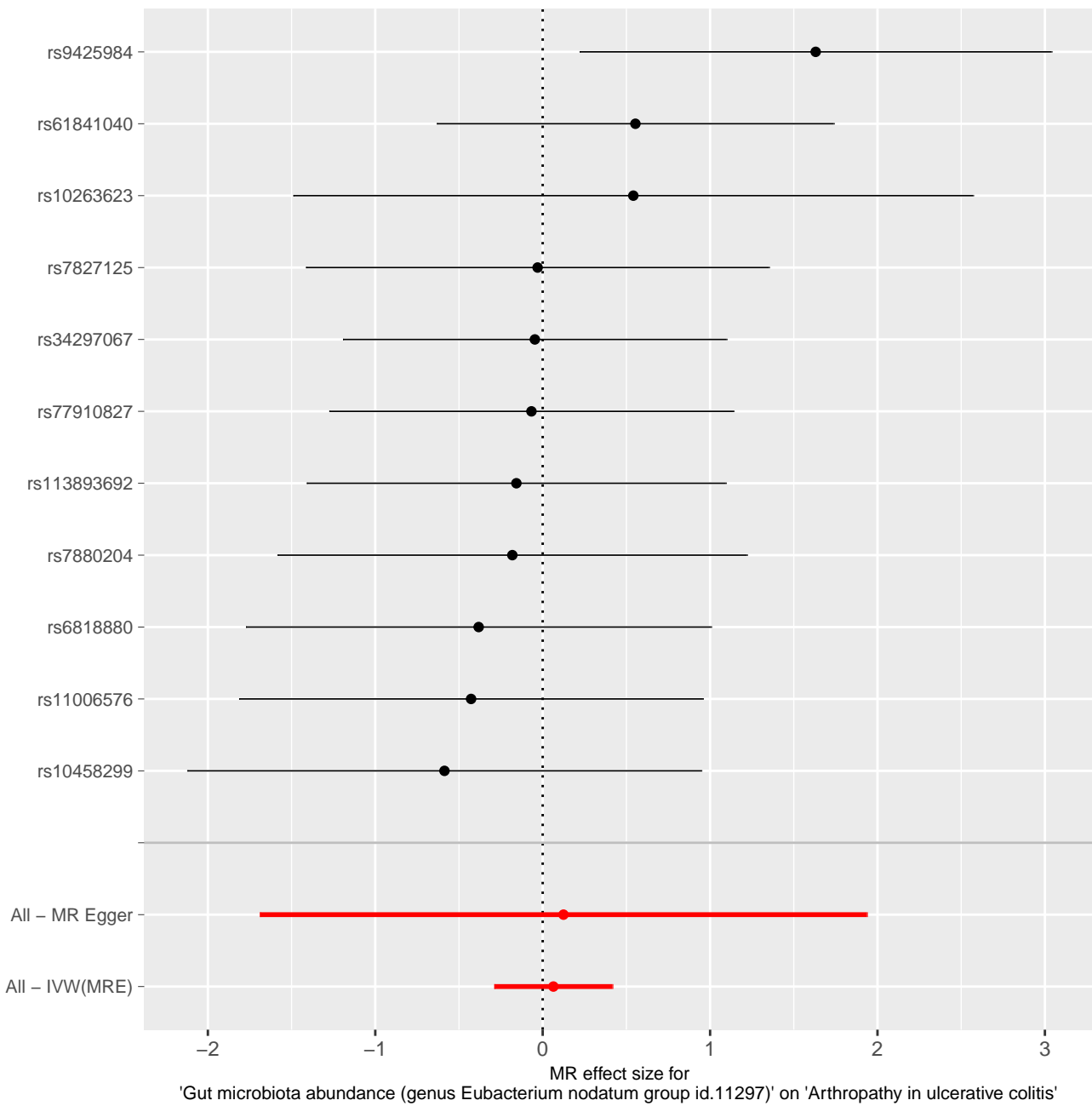


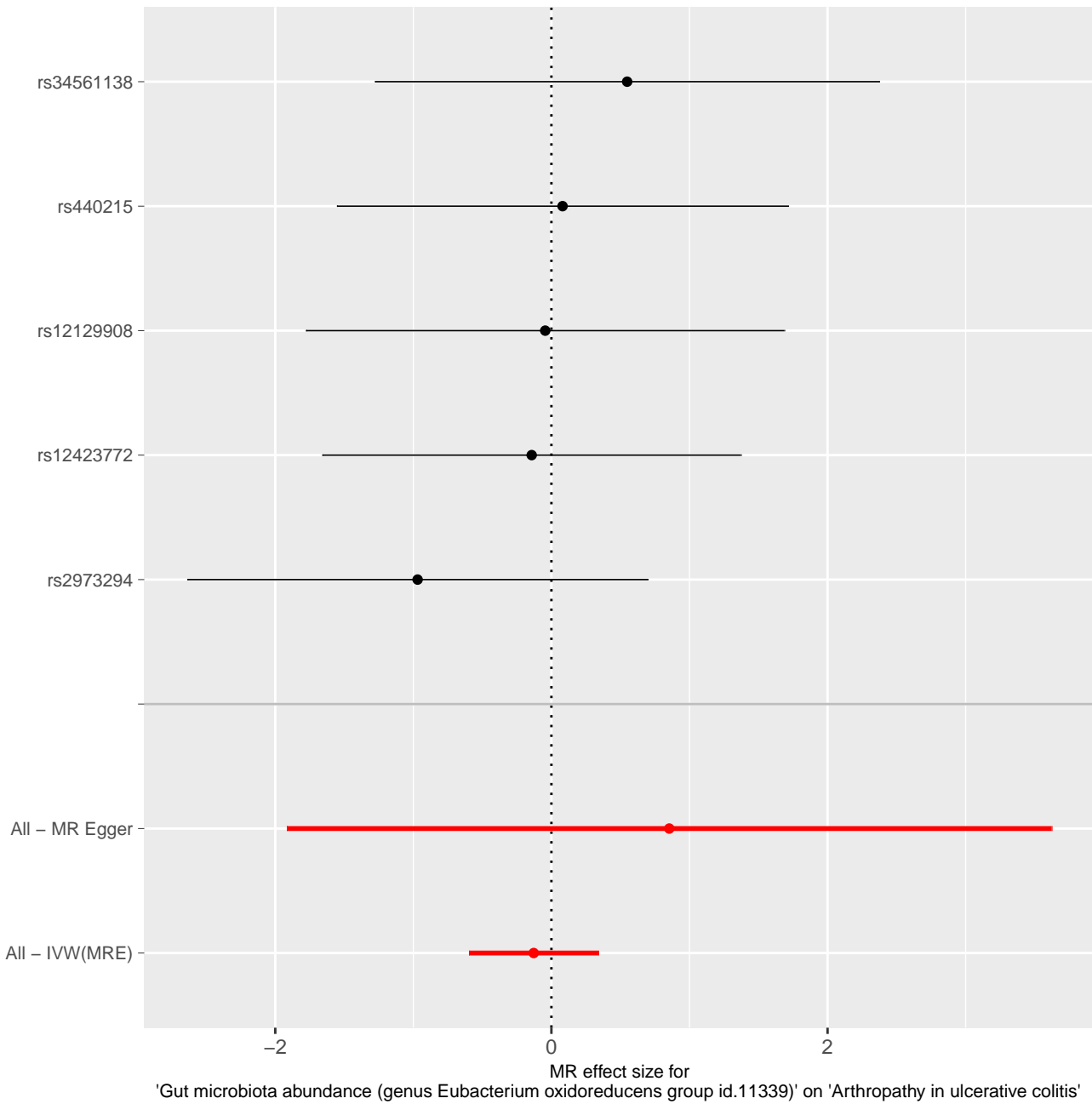


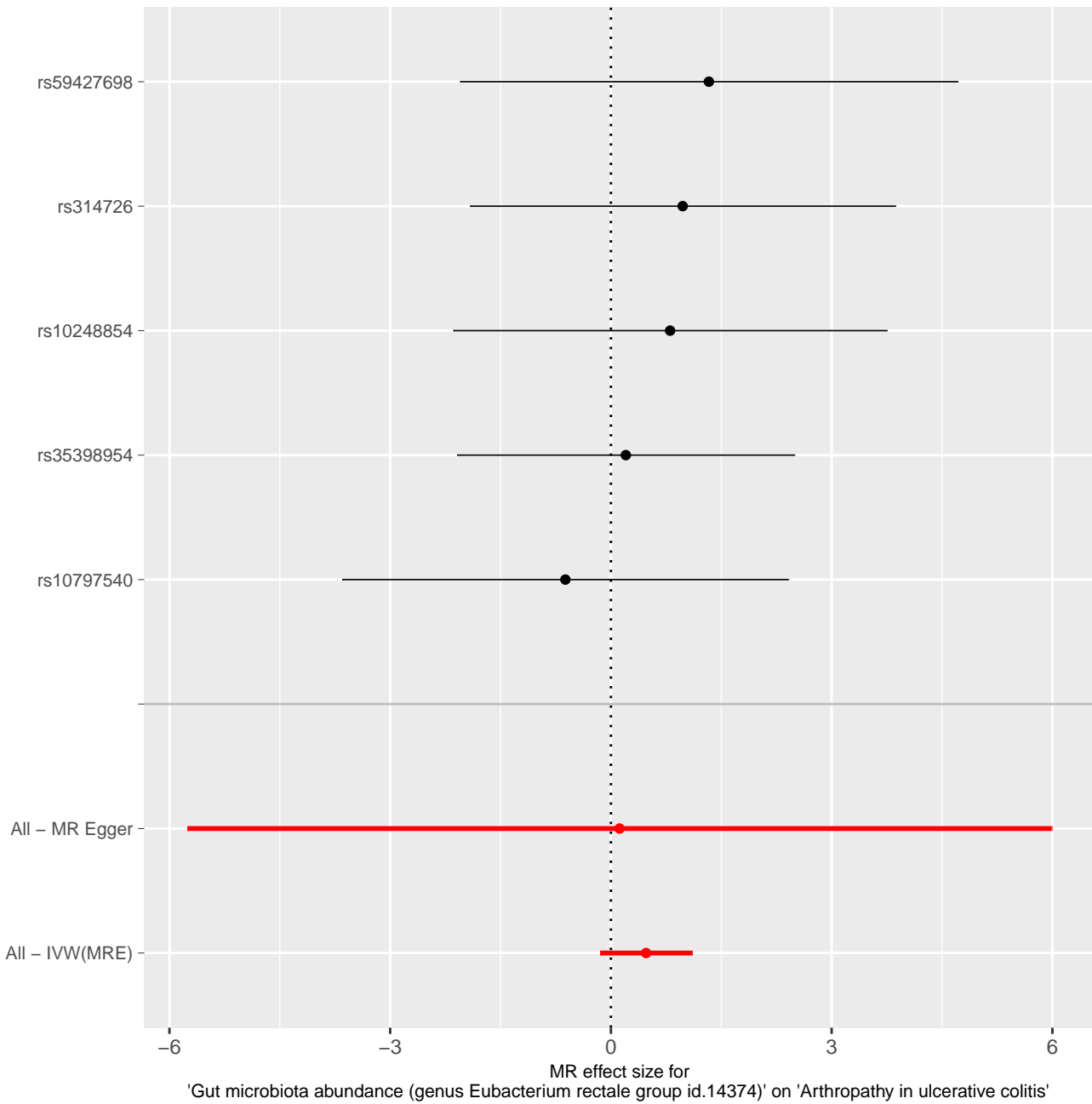


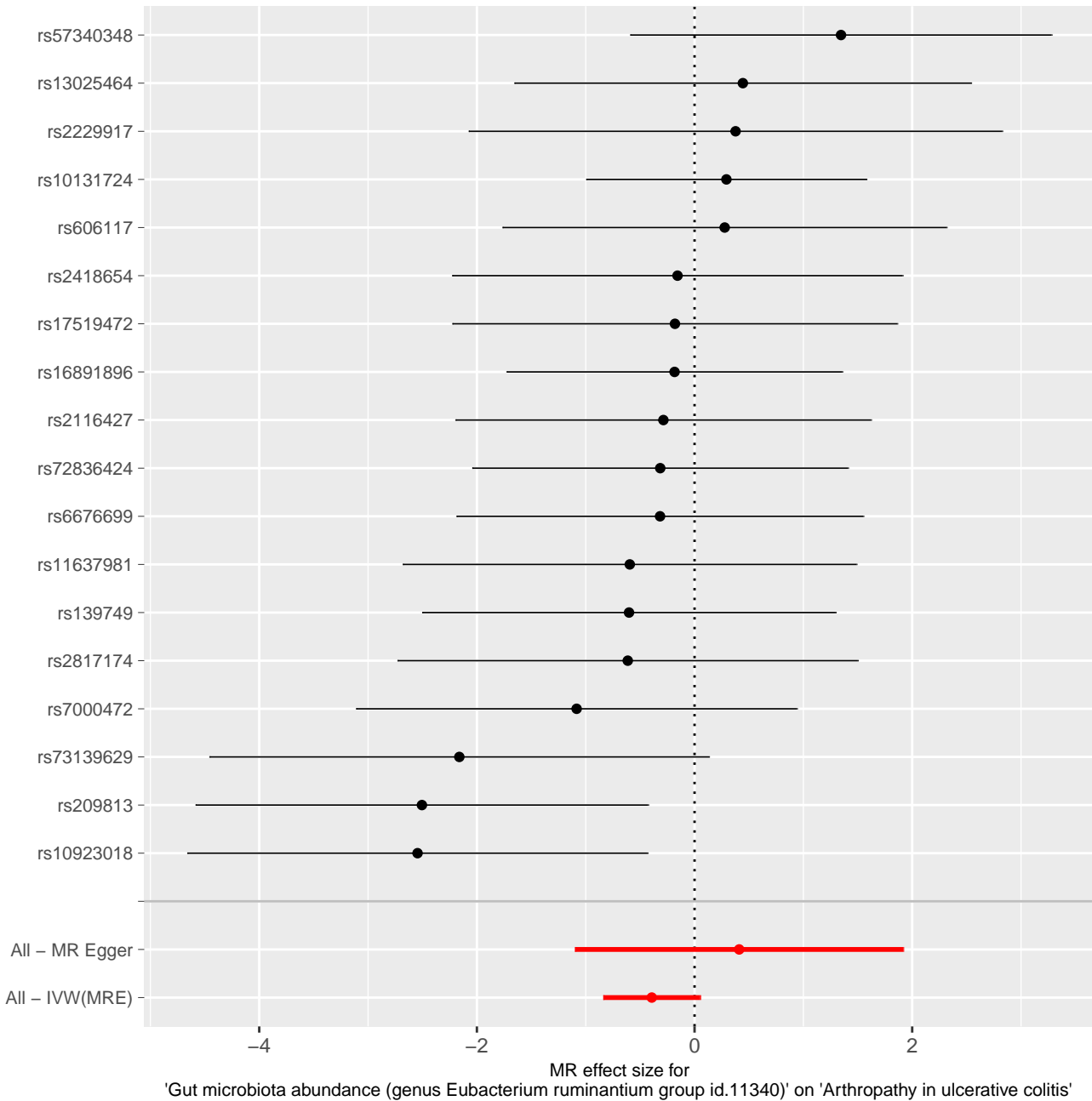


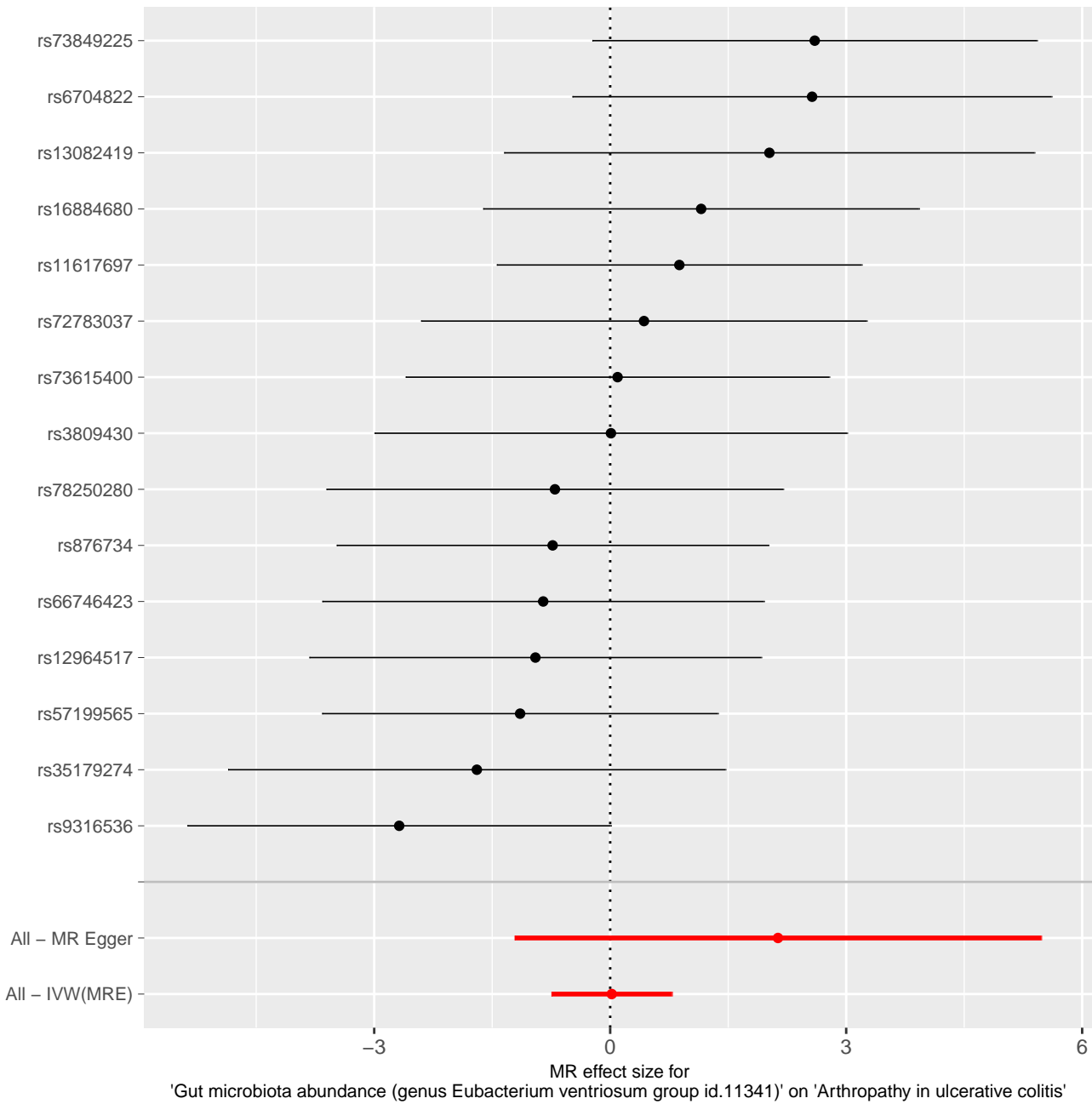
Batch 825 : Gut microbiota abundance (genus Eubacterium nodatum group id.11297) on Arthropathy in ulcerative colitis

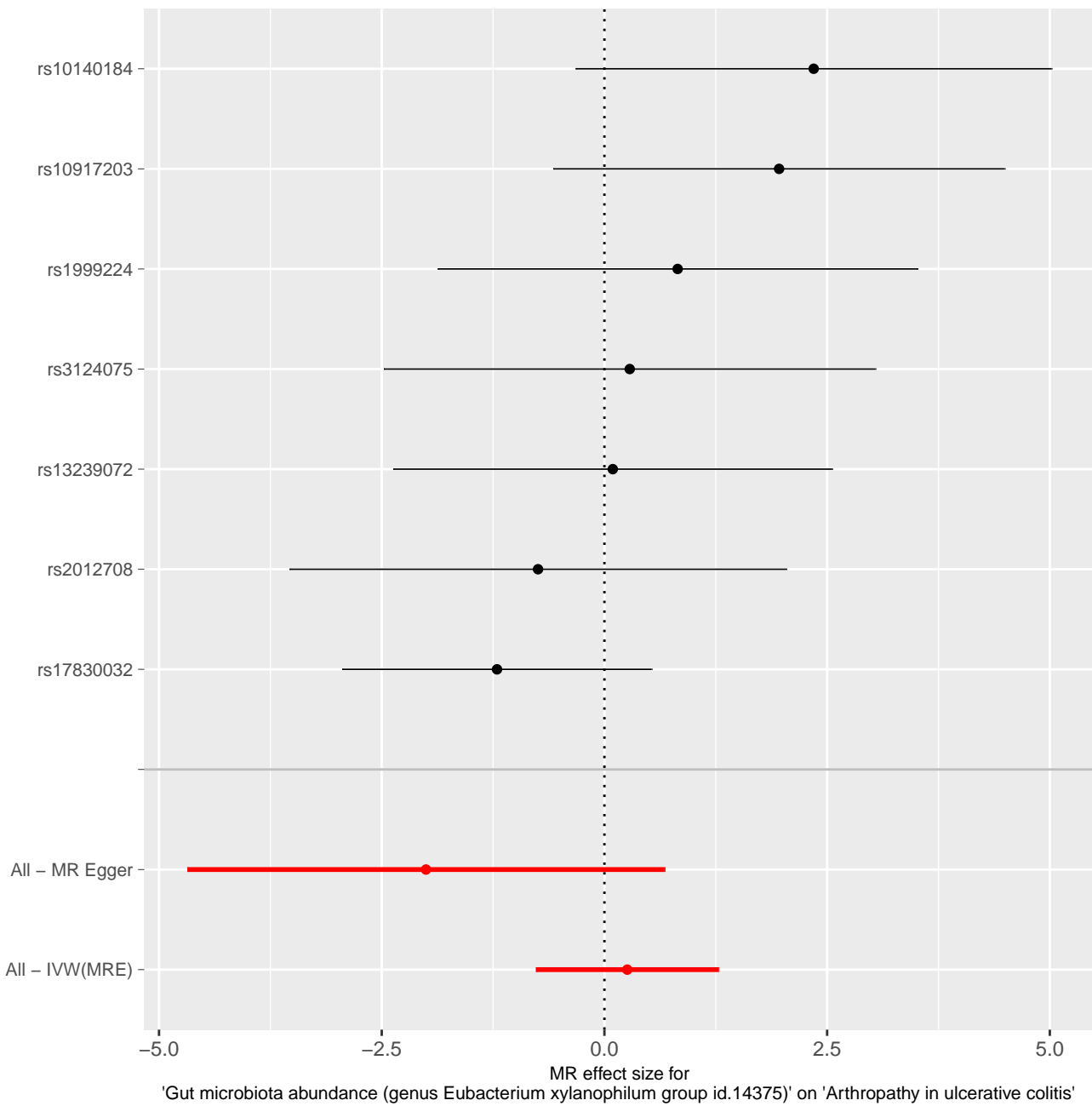


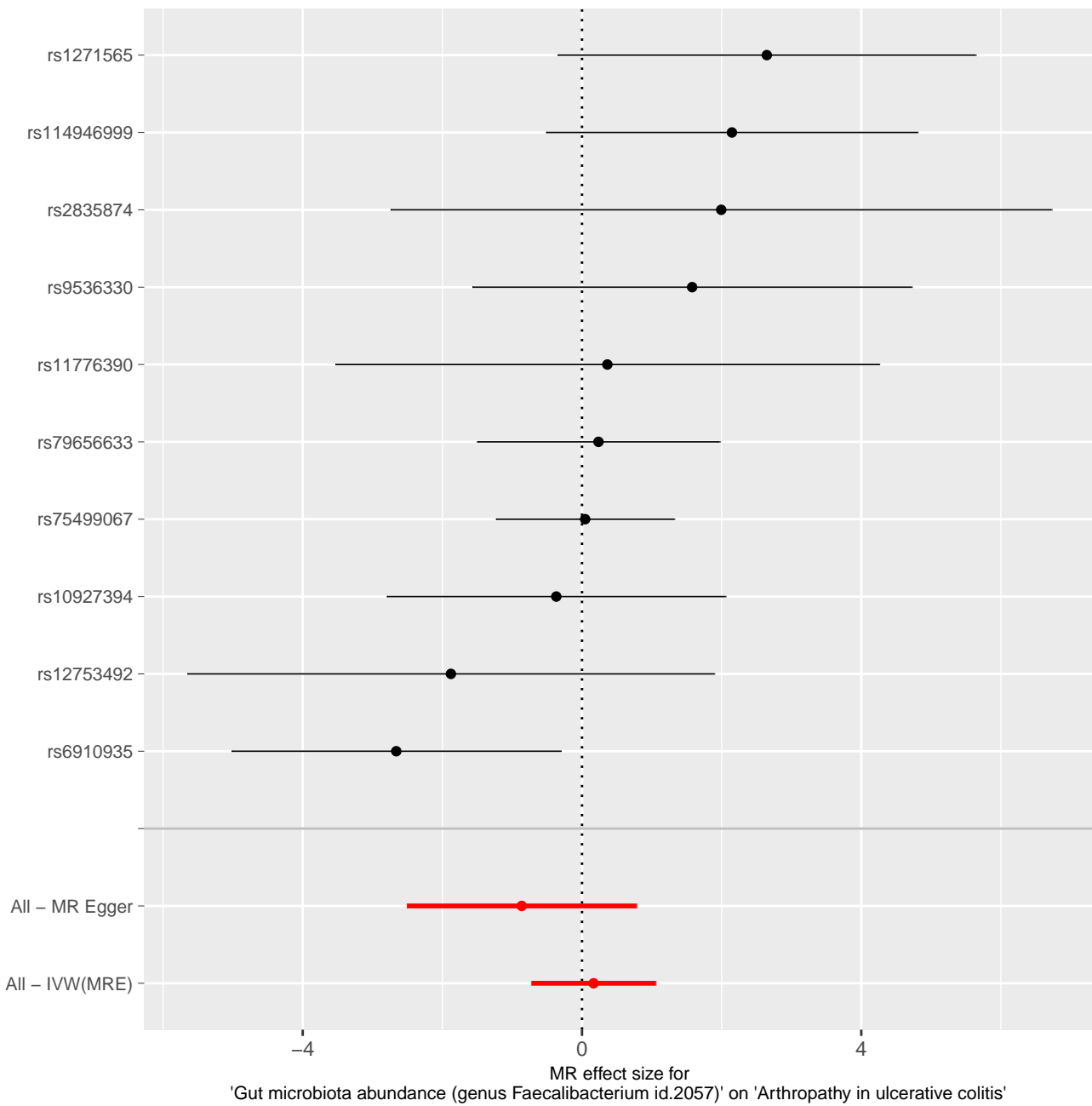


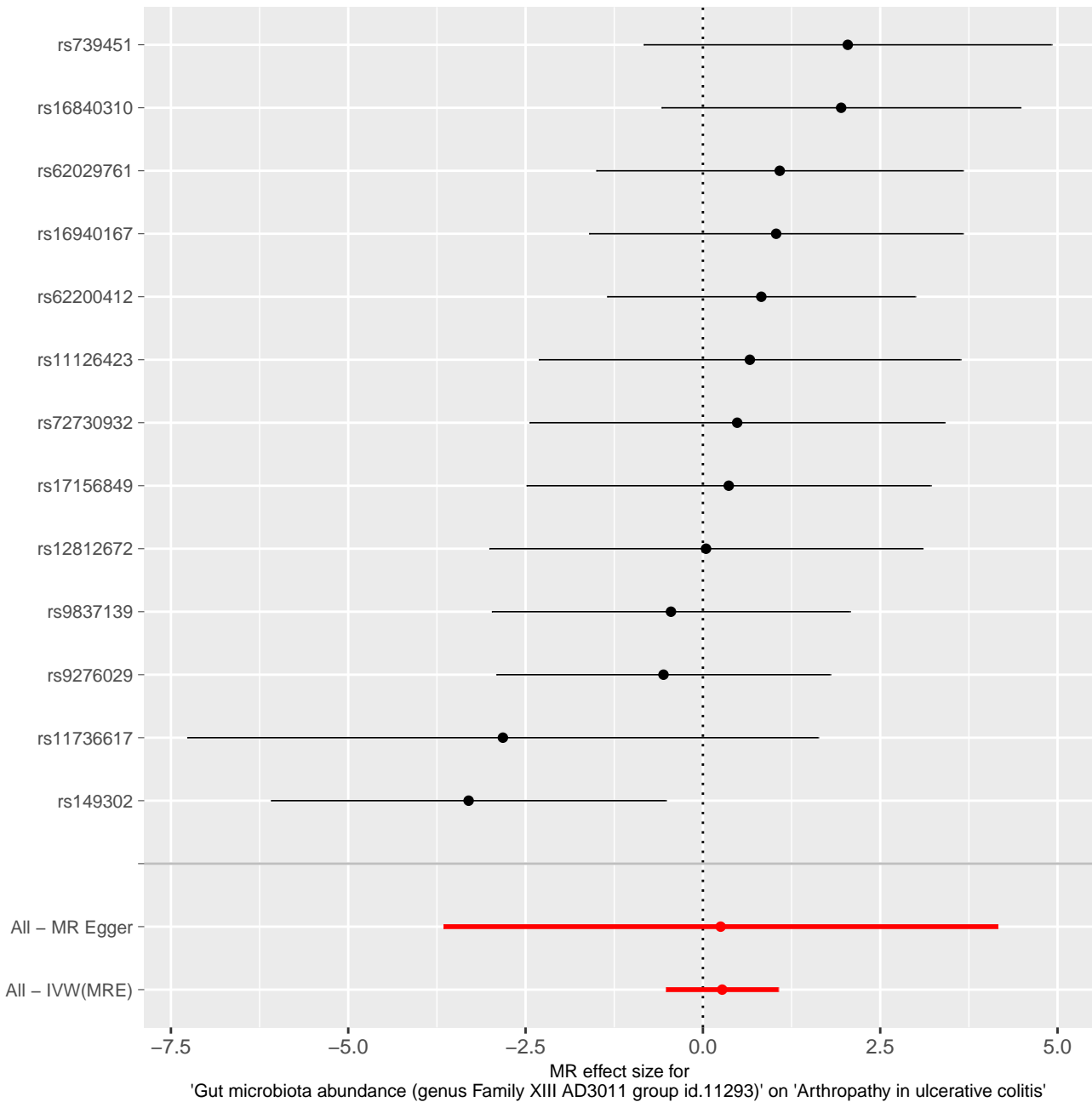


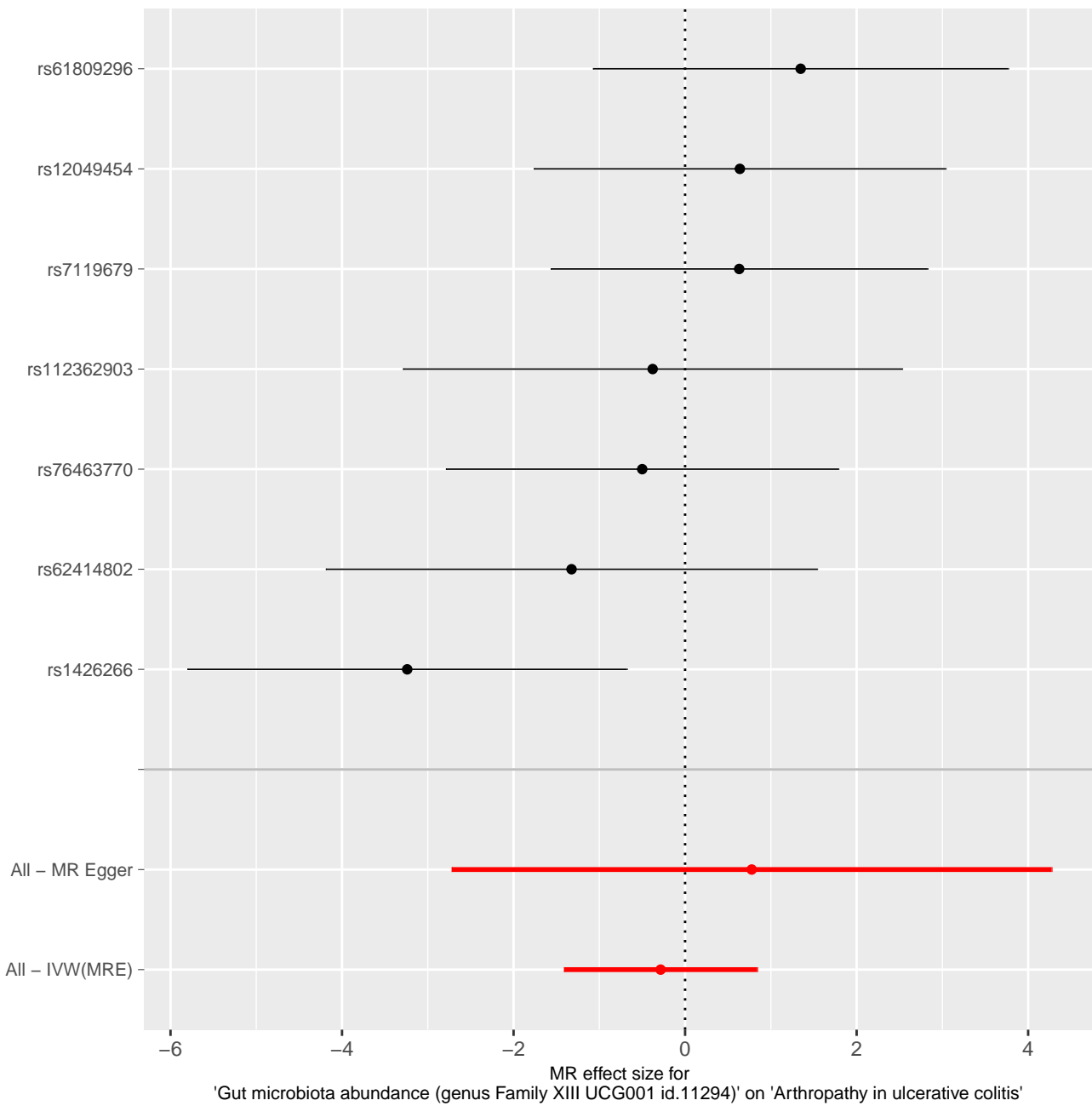


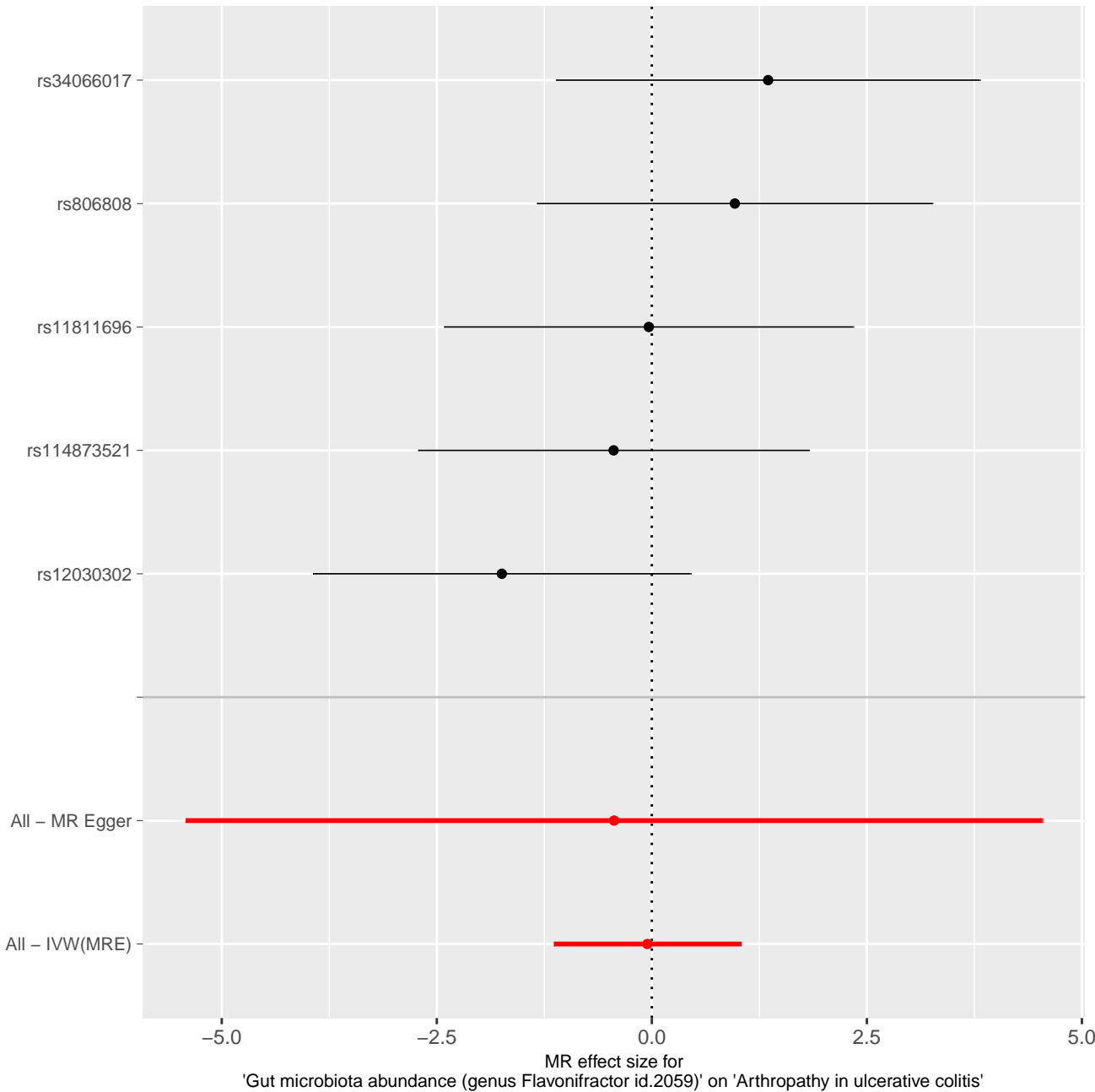


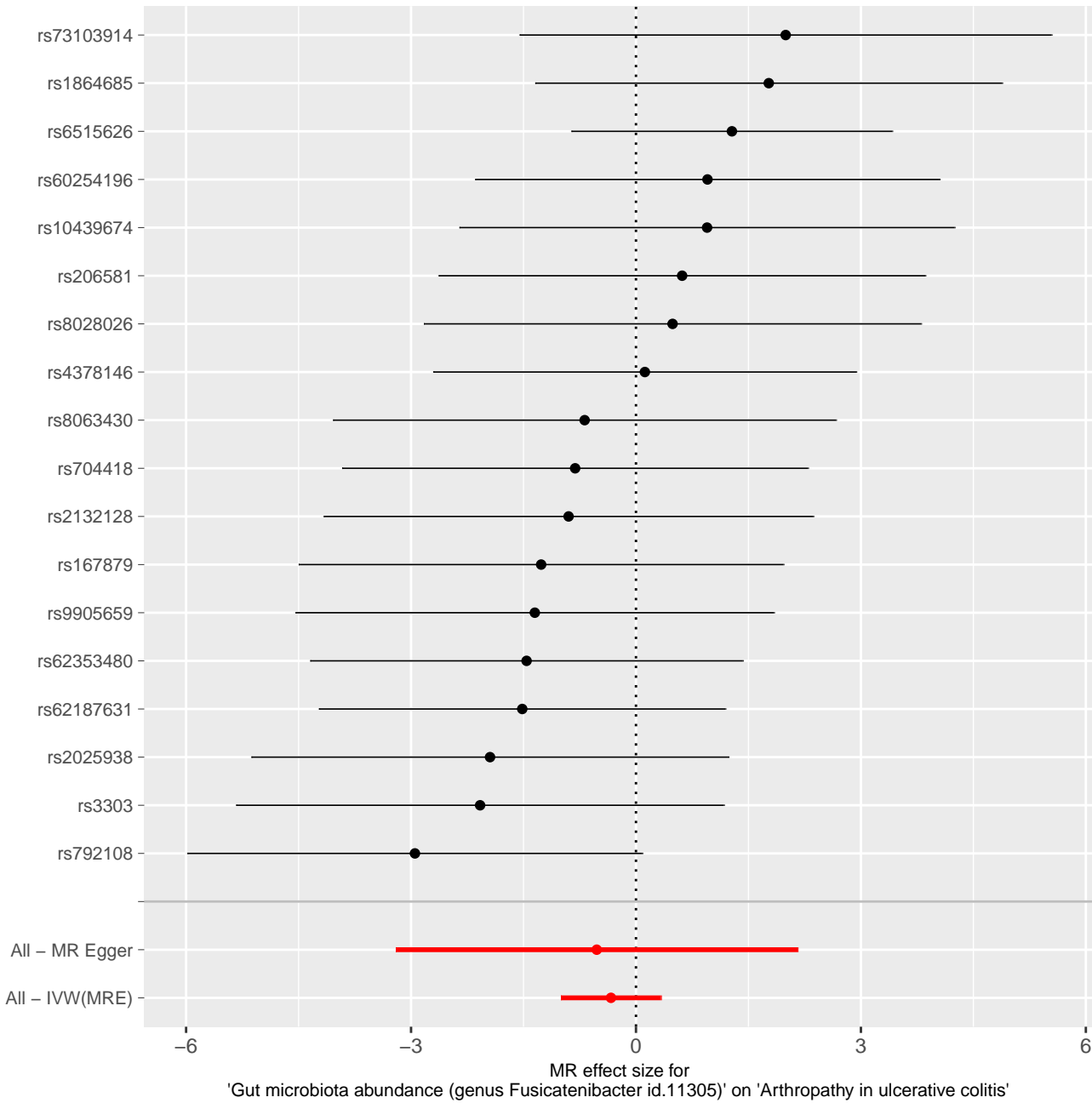


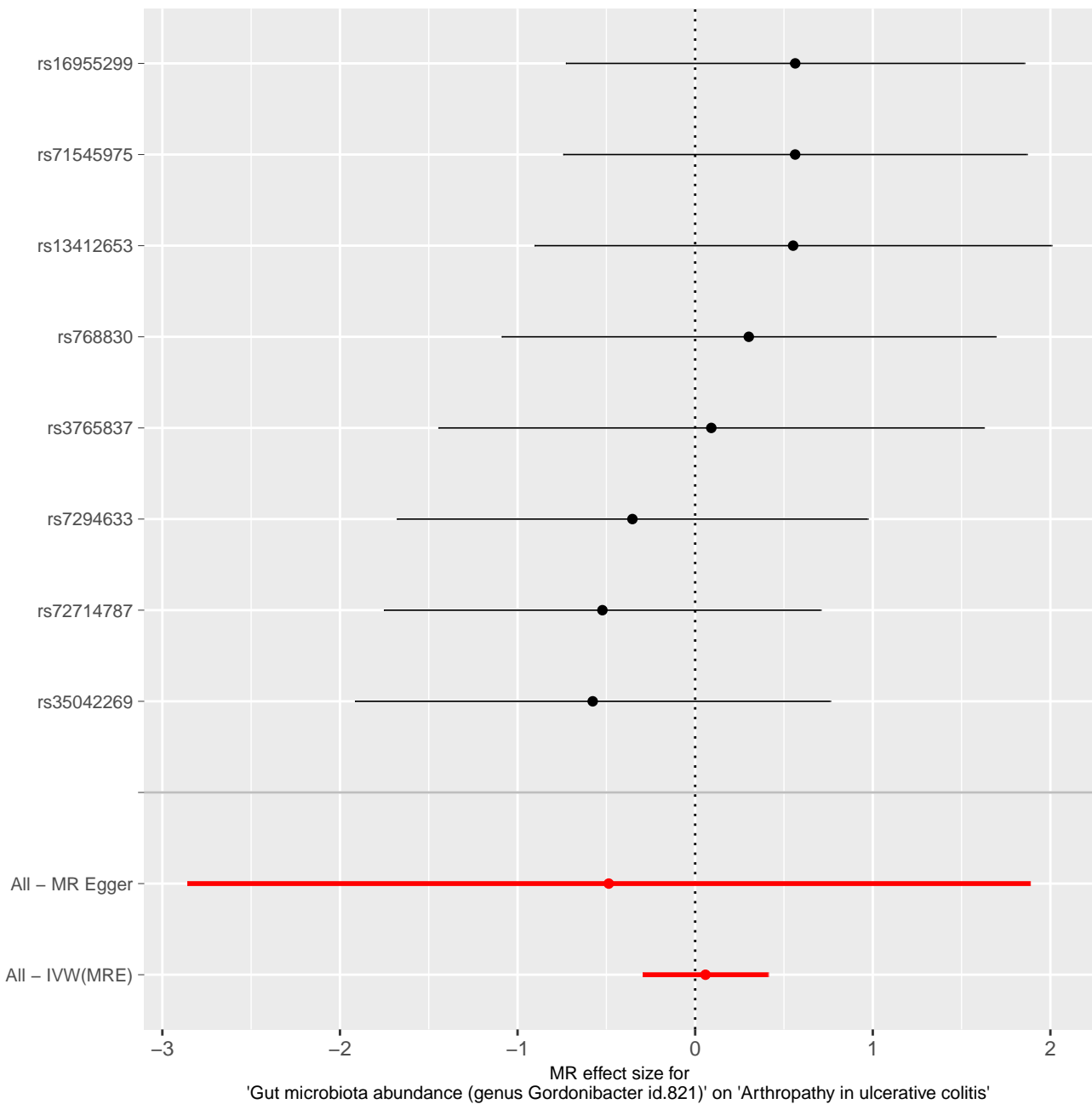


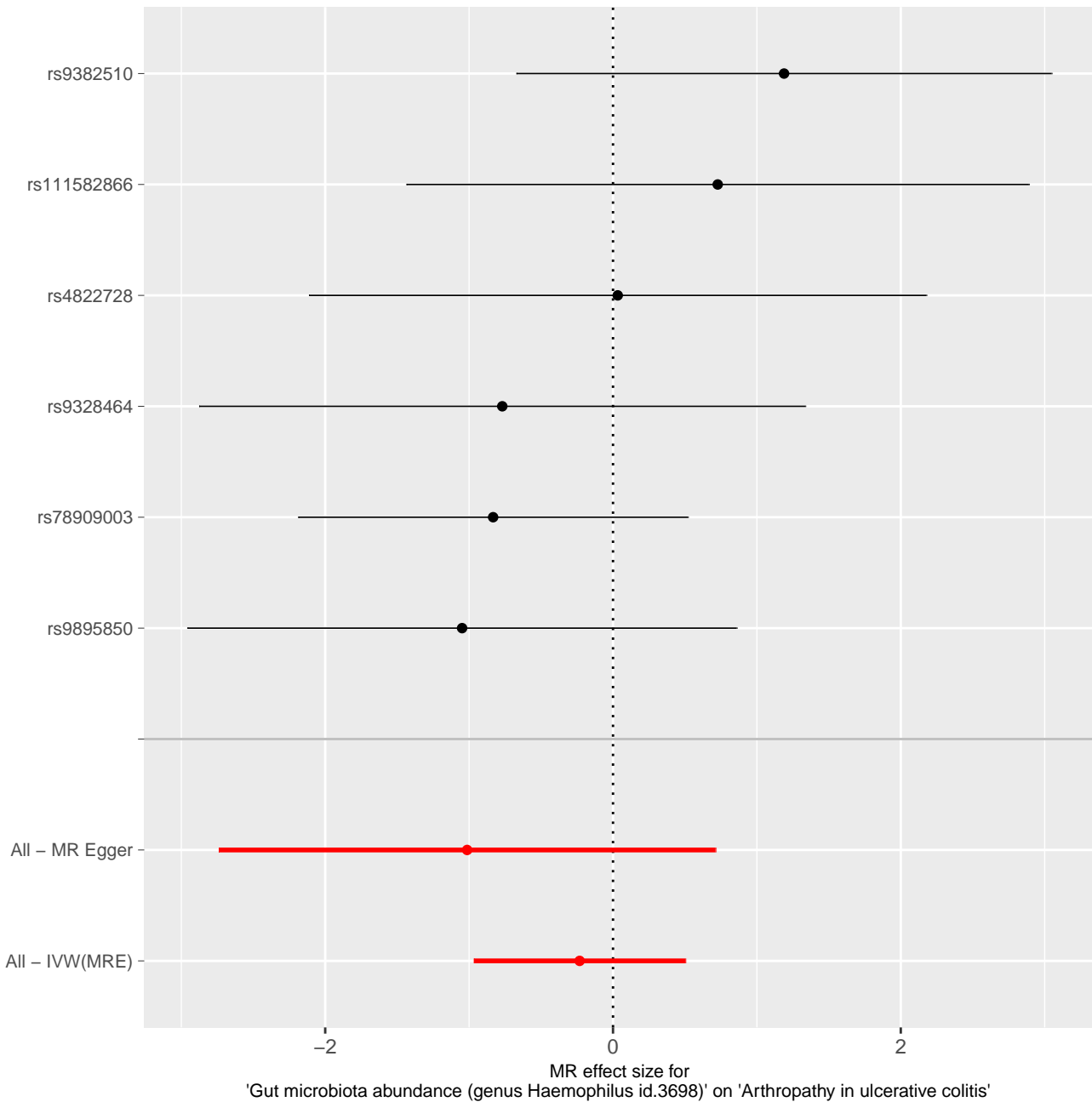


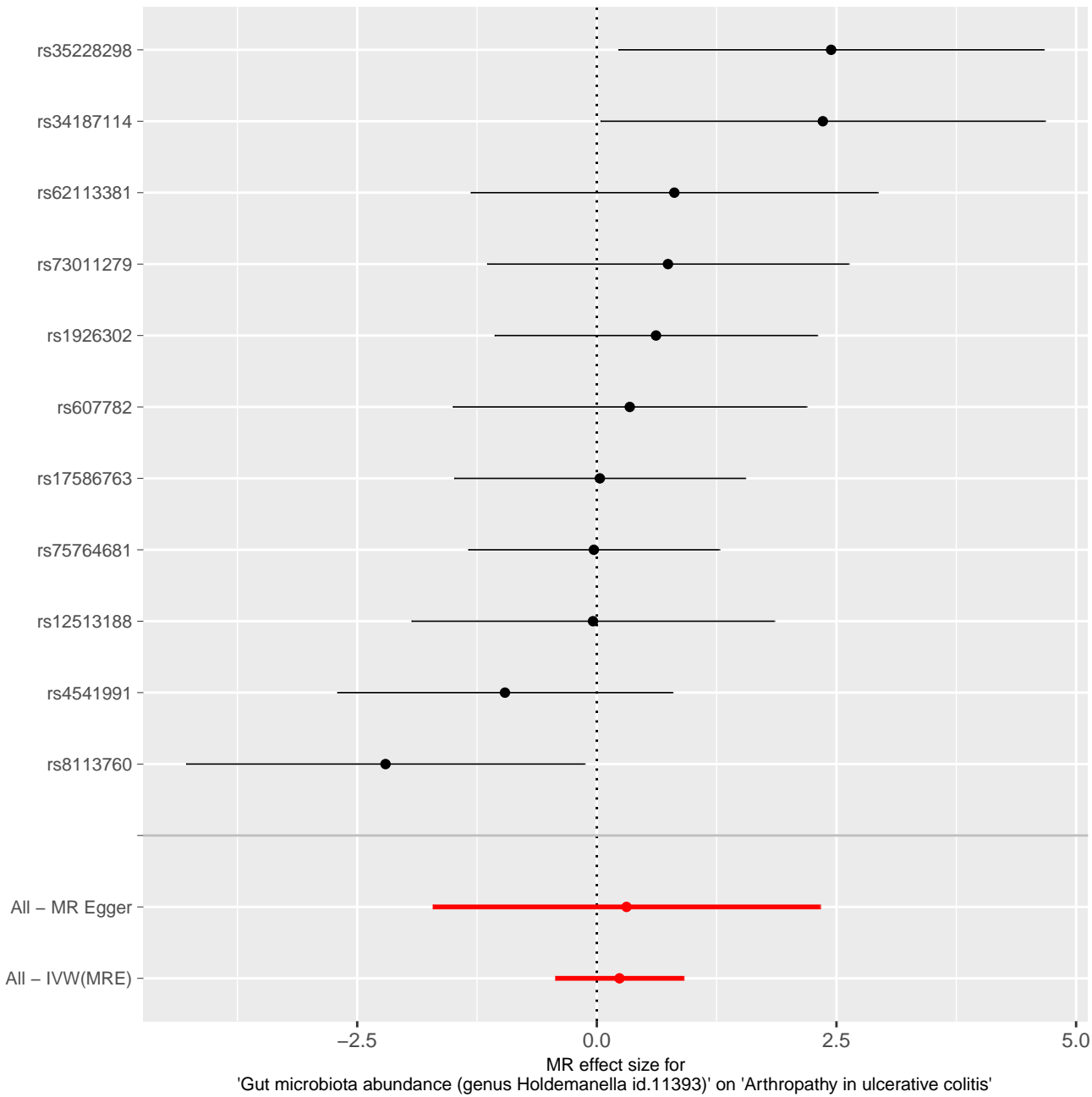




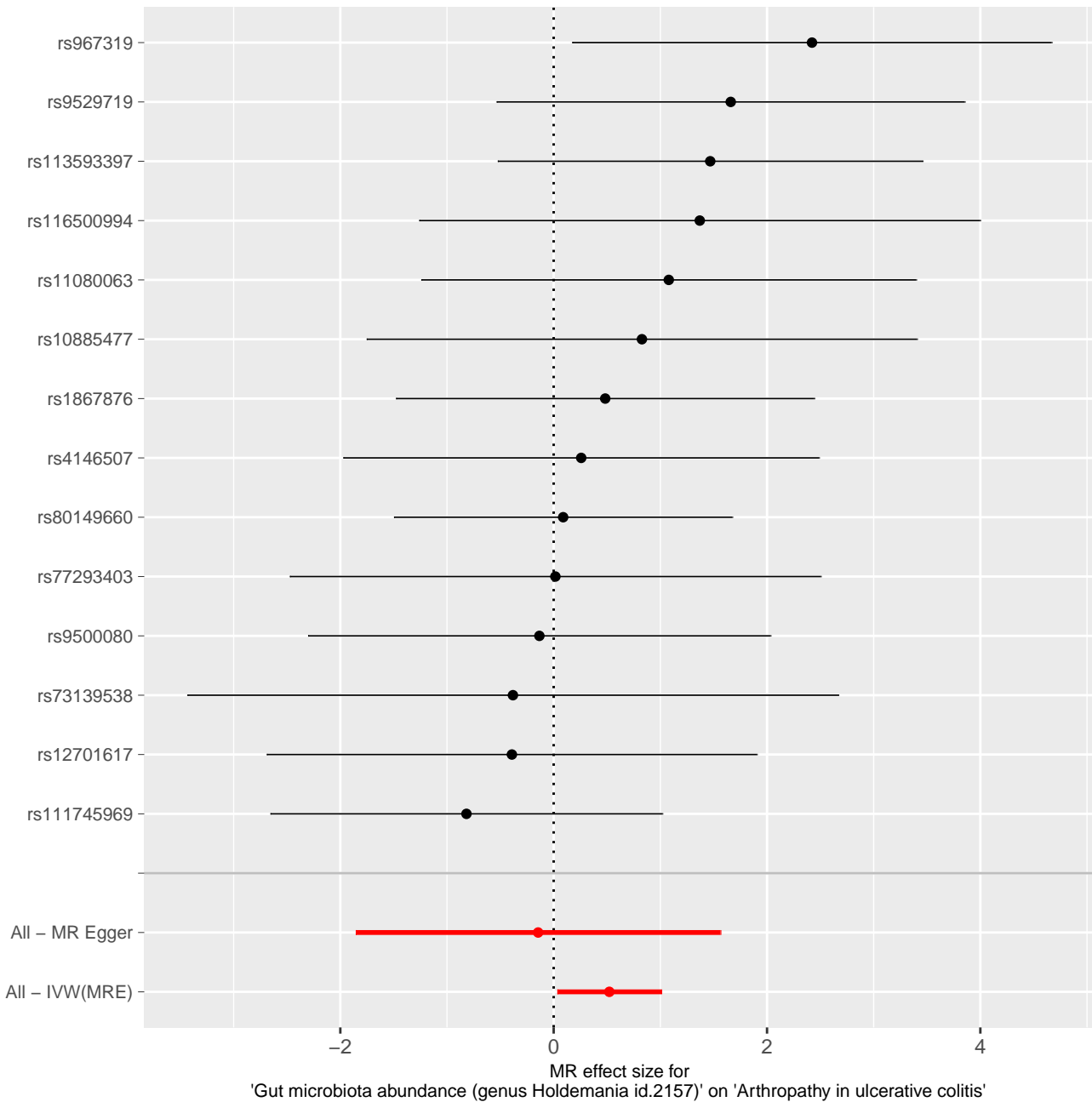


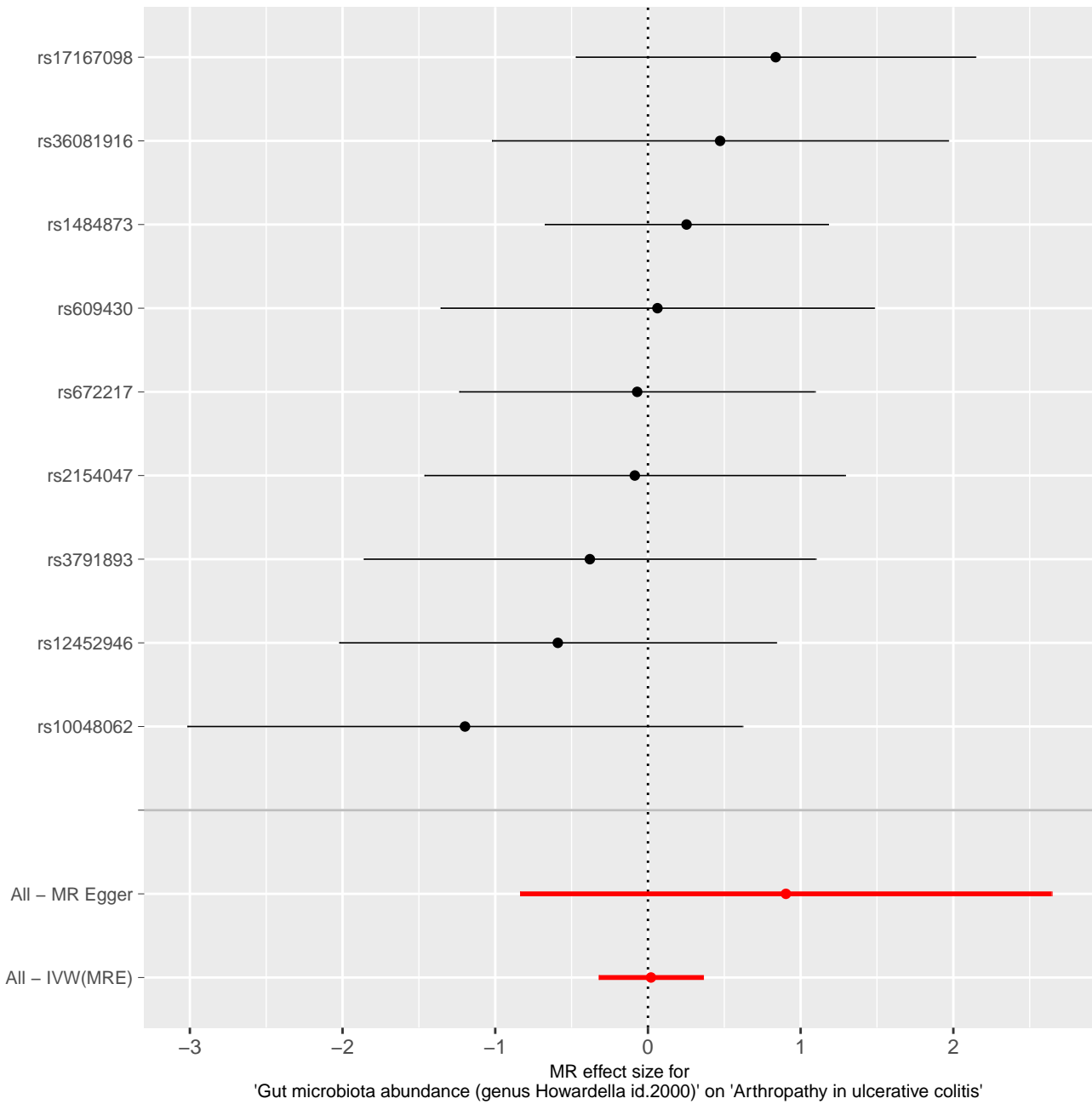


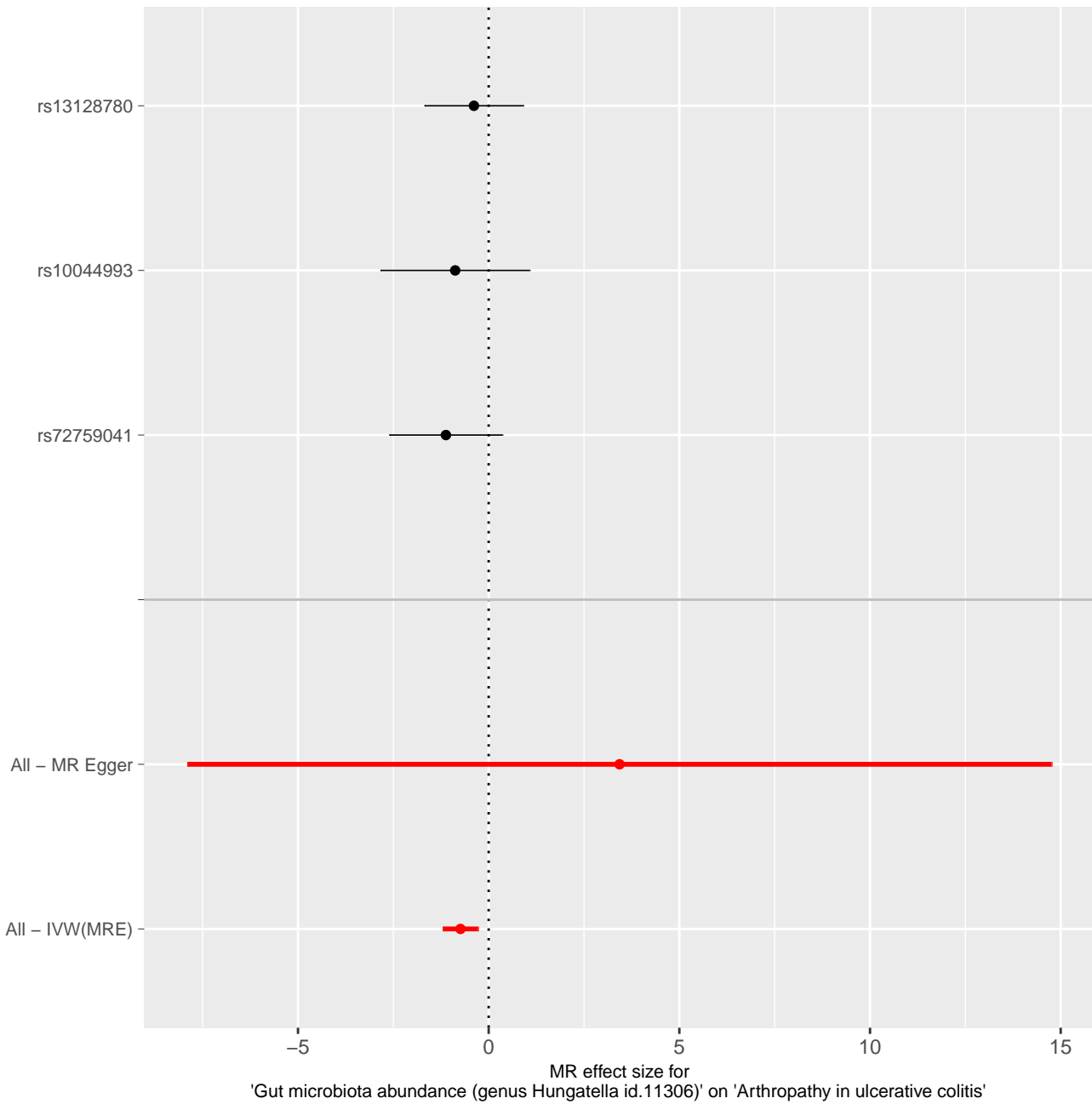


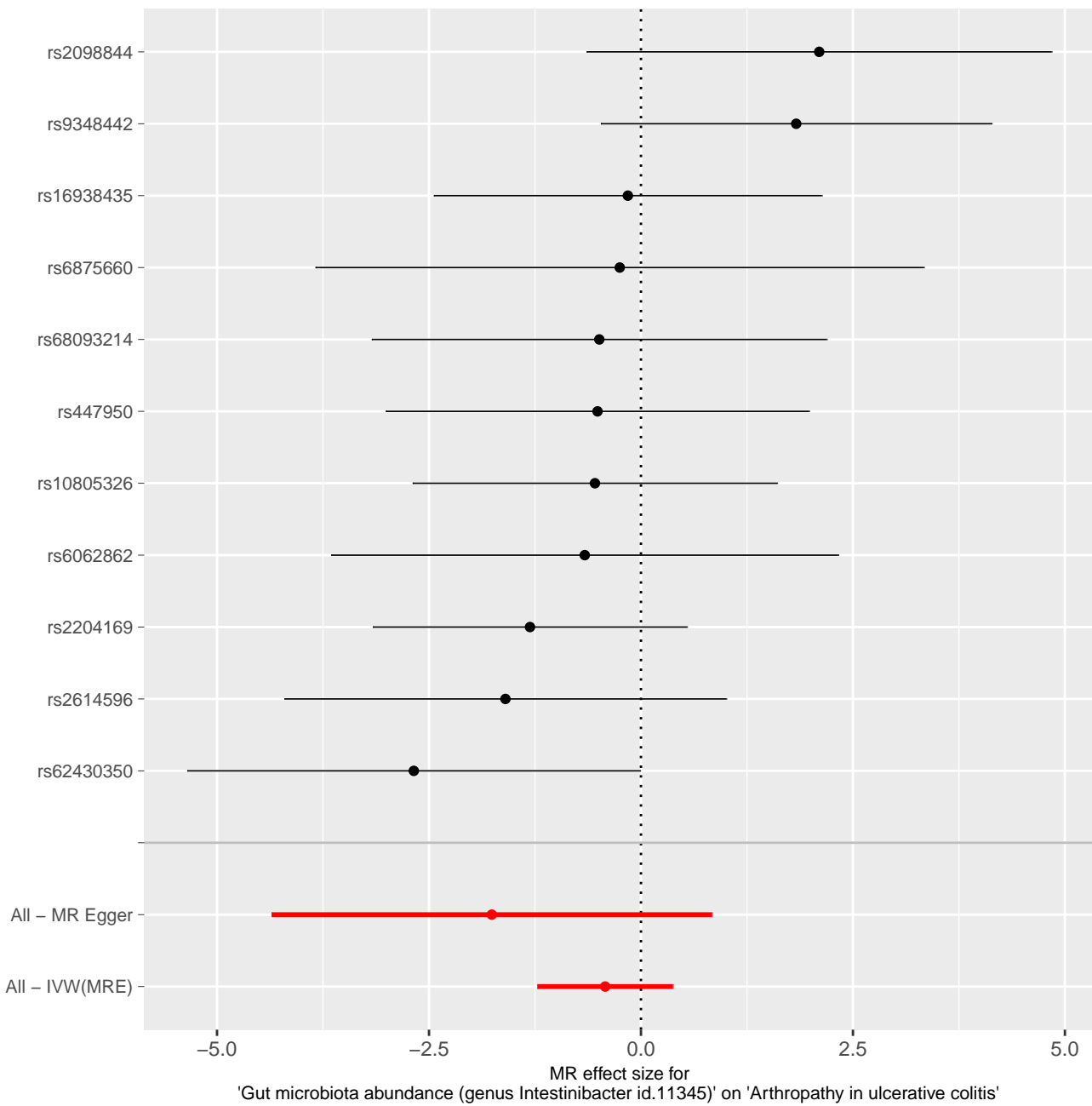


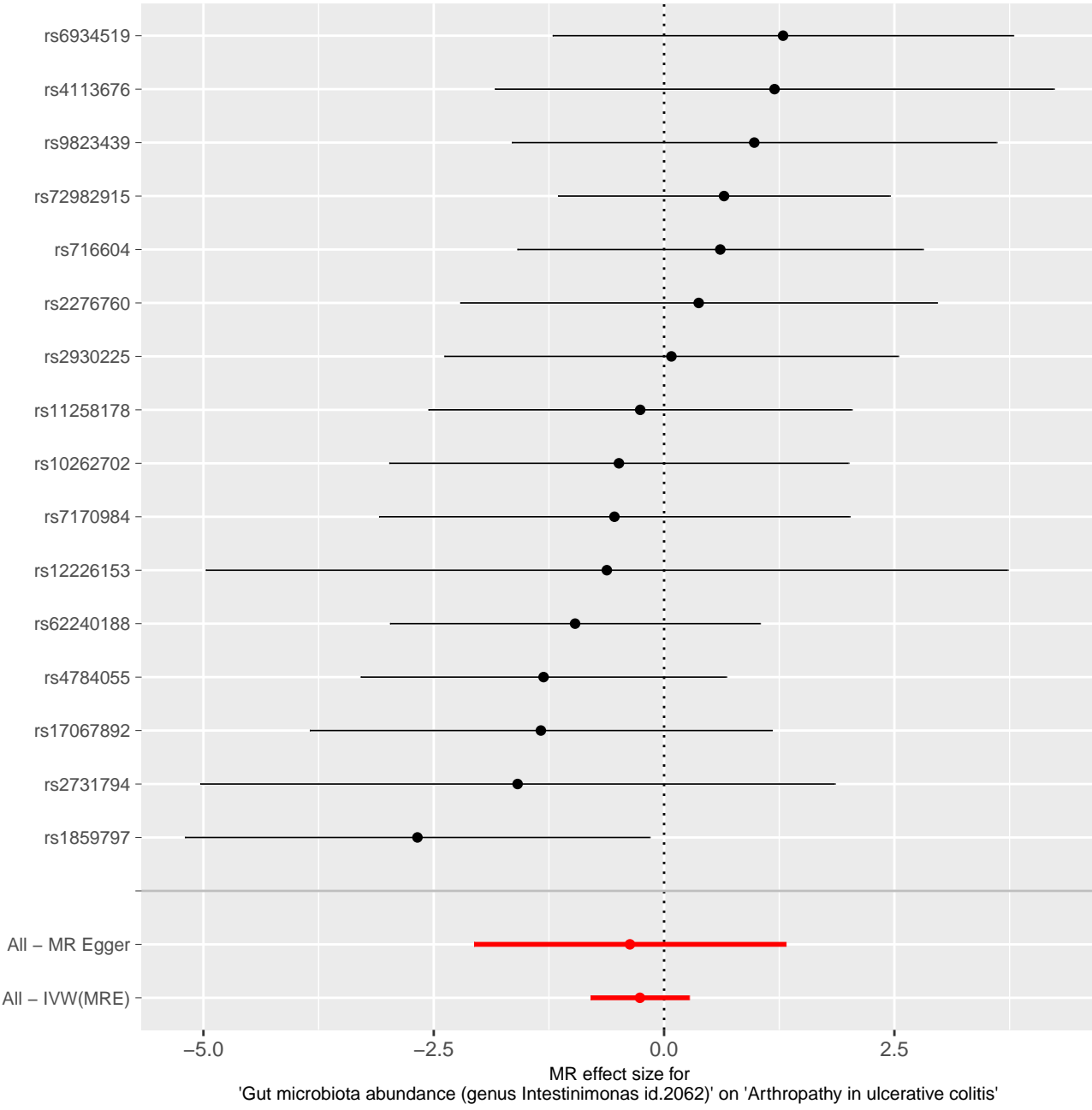
Batch 839 : Gut microbiota abundance (genus Holdemania id.2157) on Arthropathy in ulcerative colitis

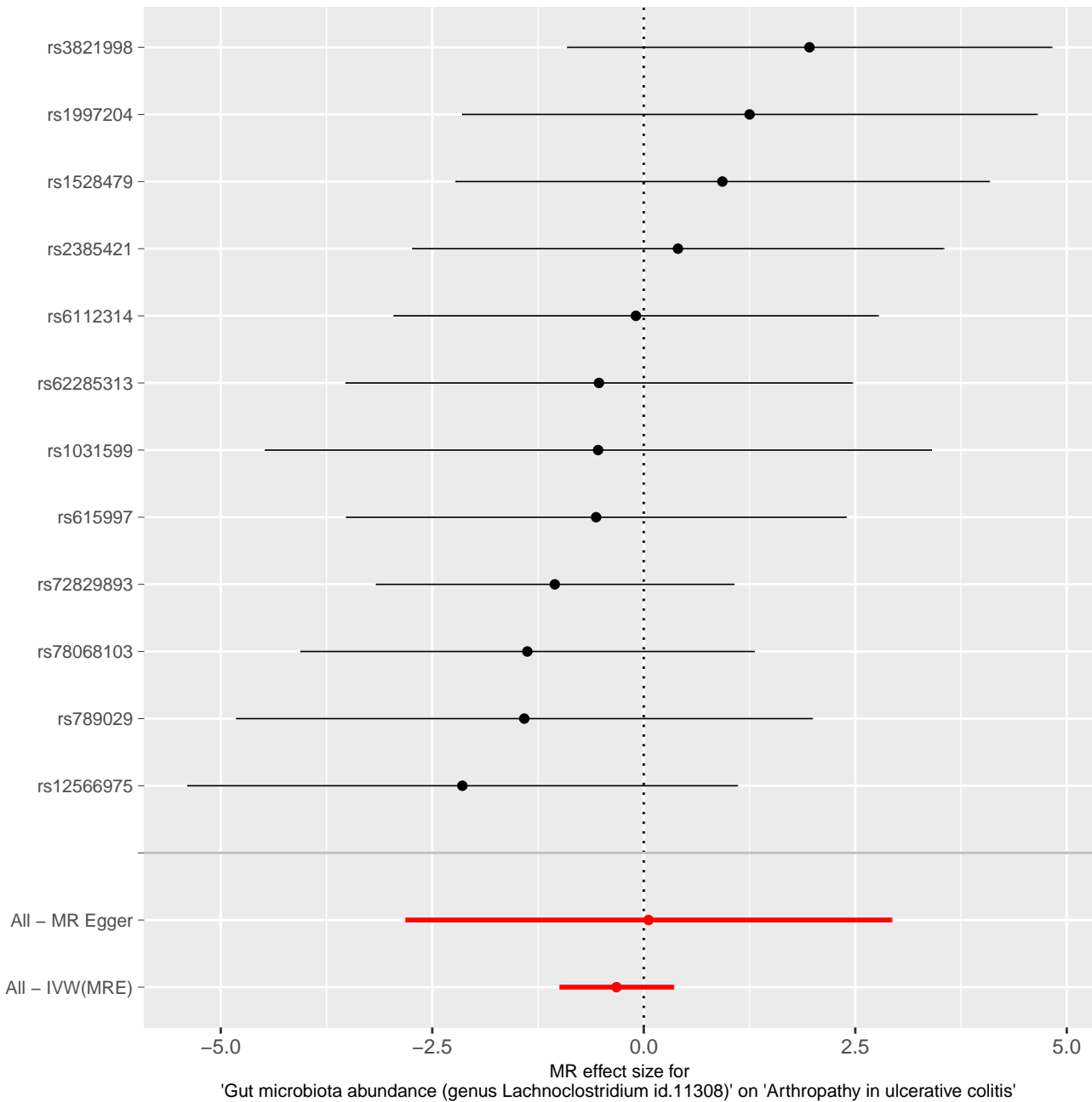


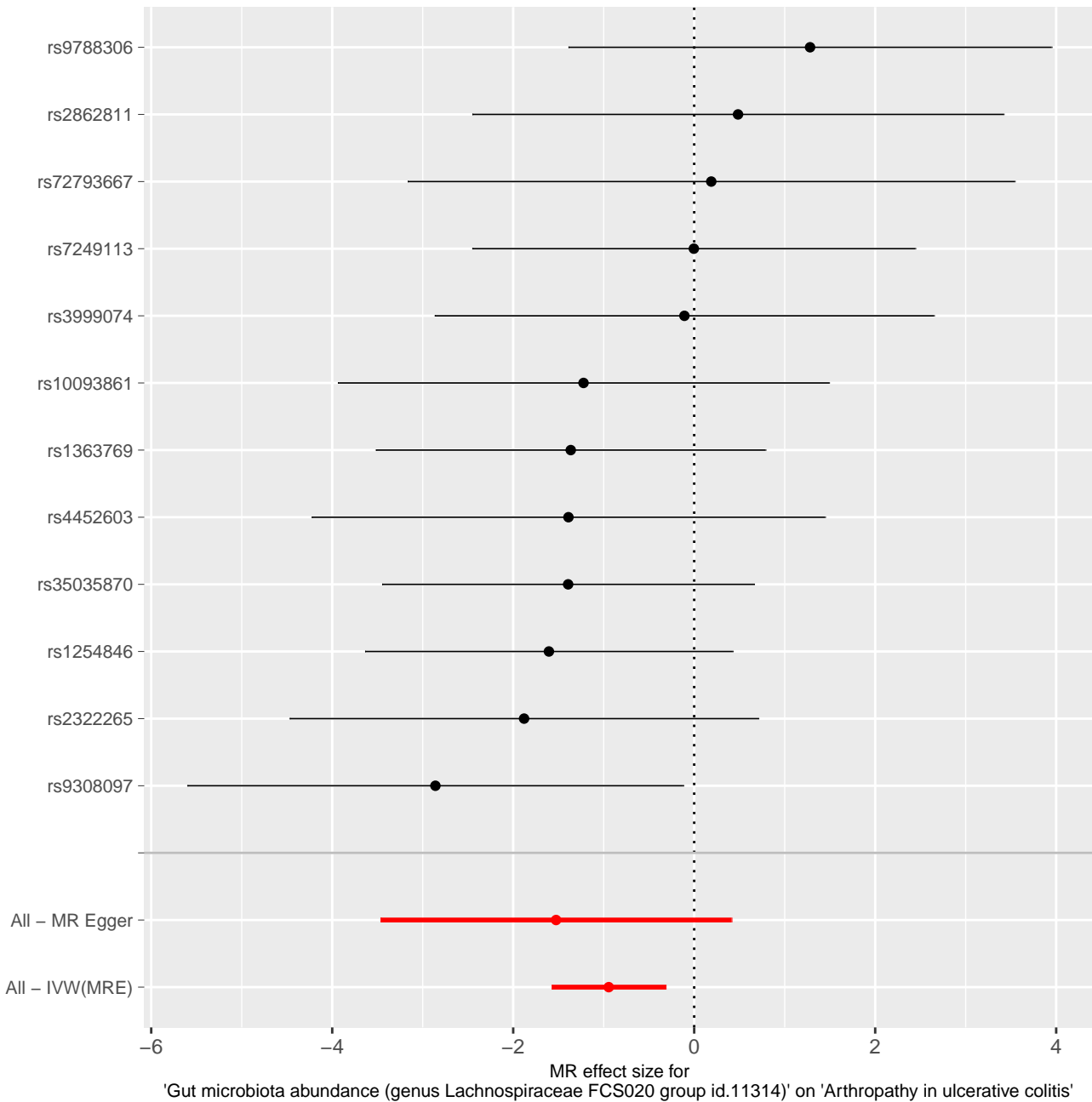


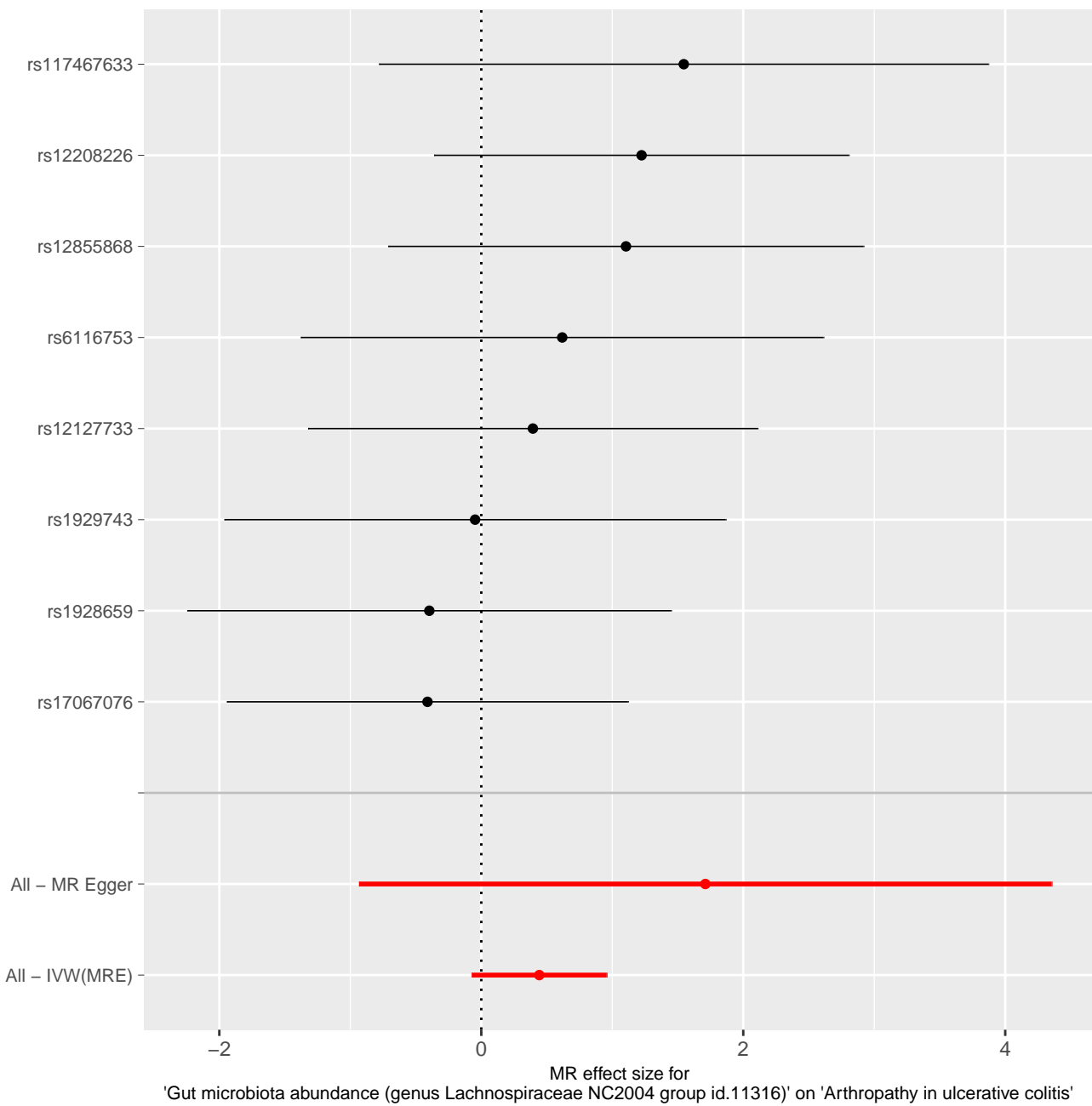


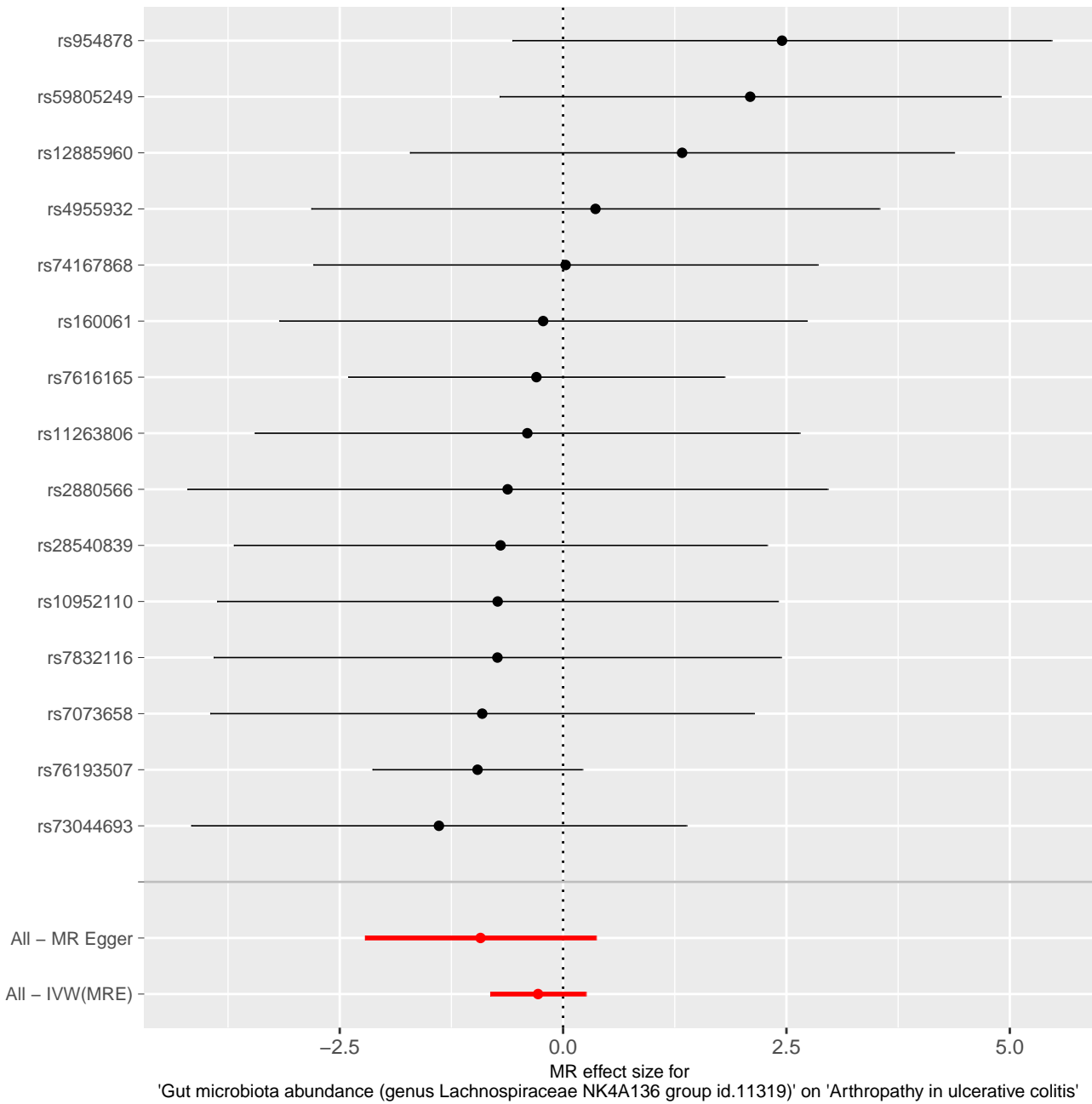


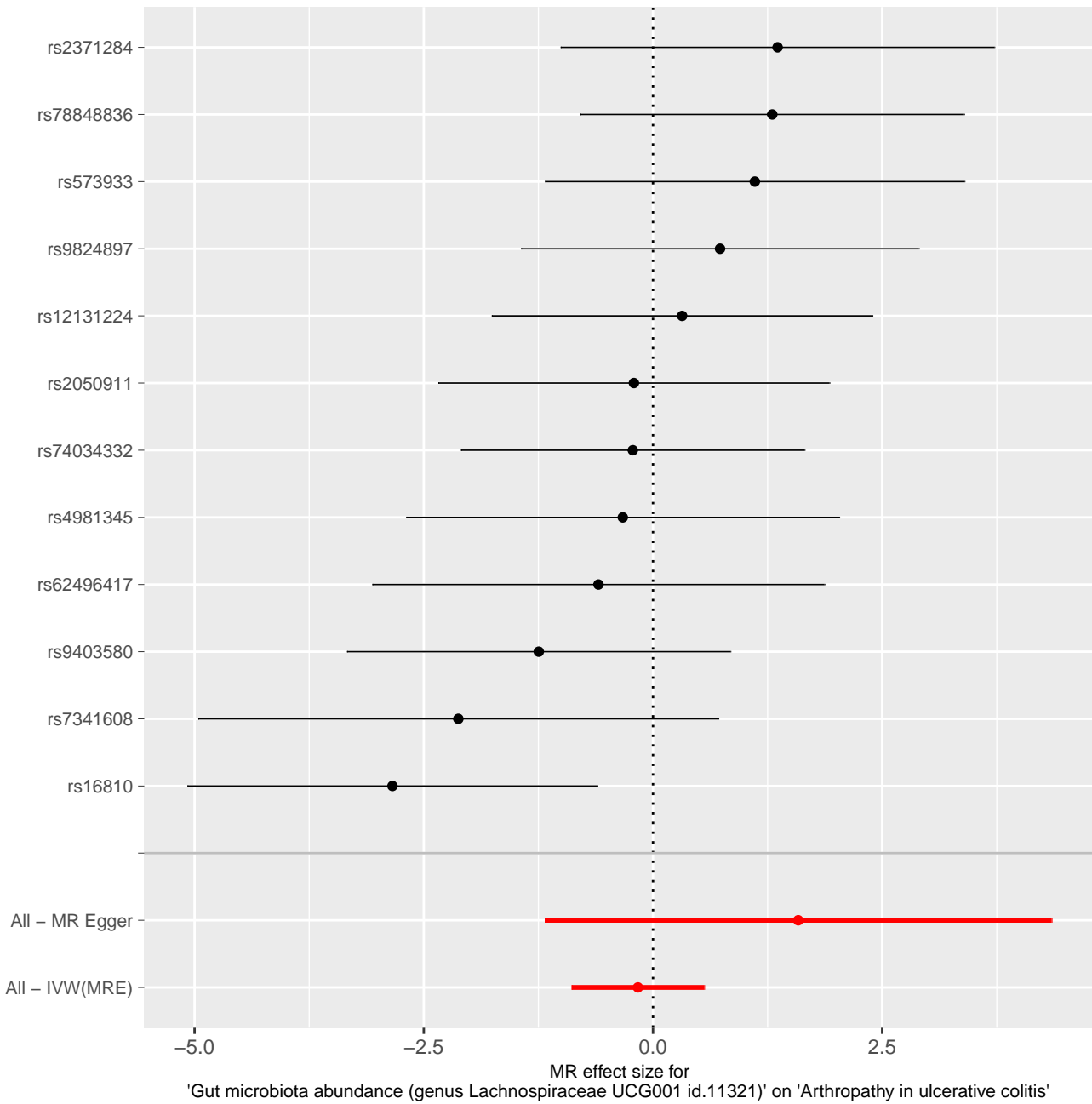


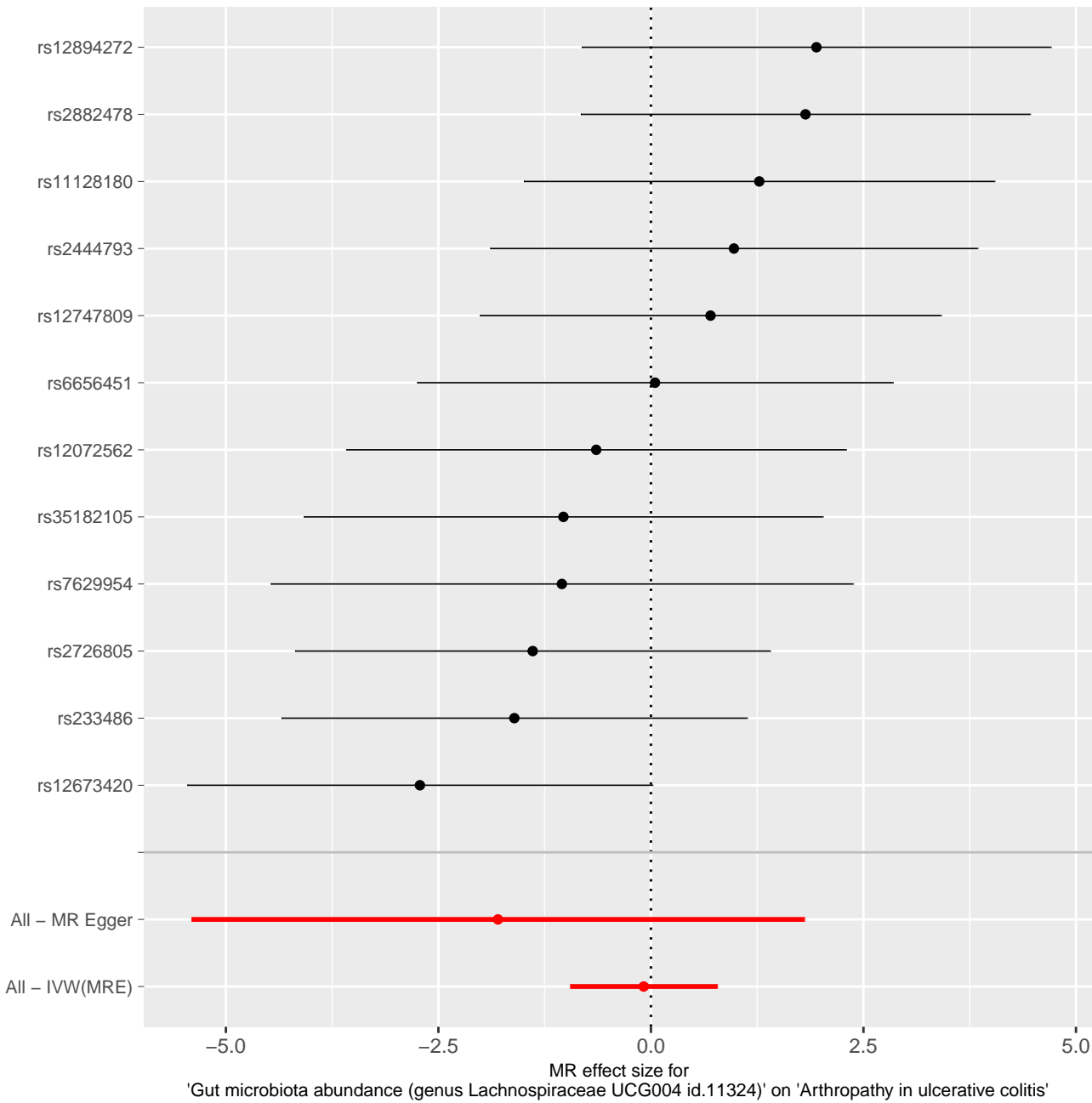


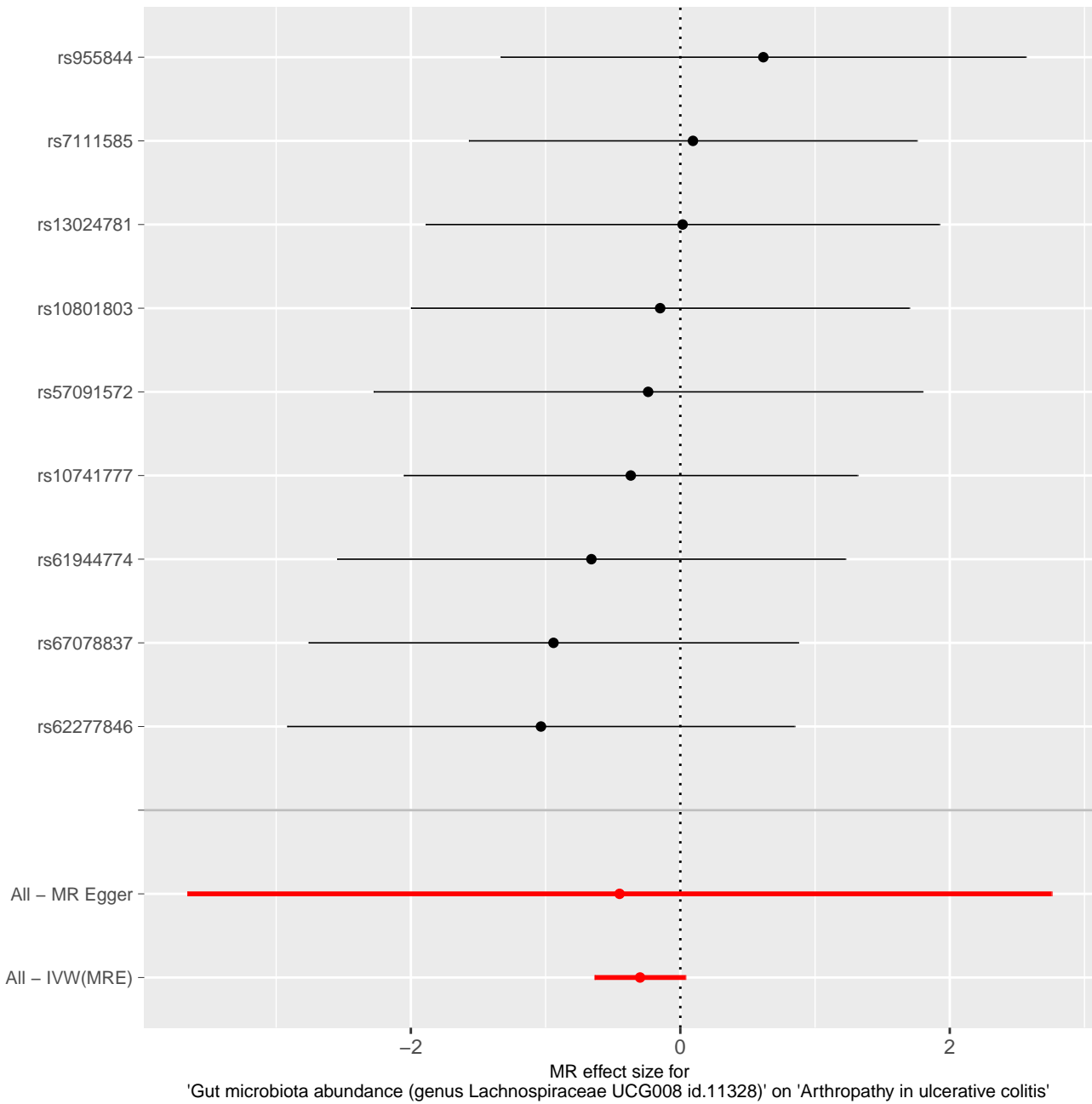


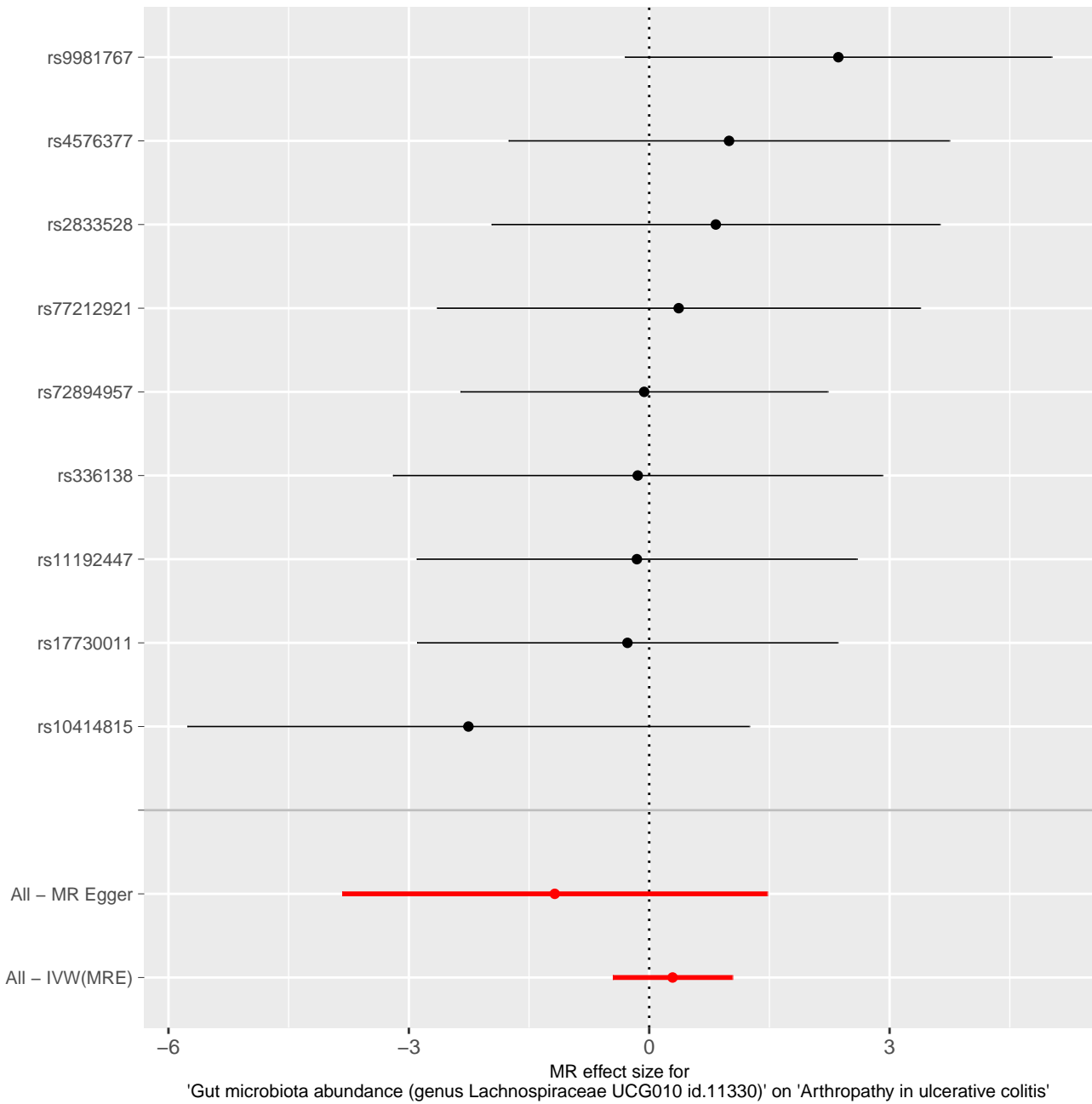


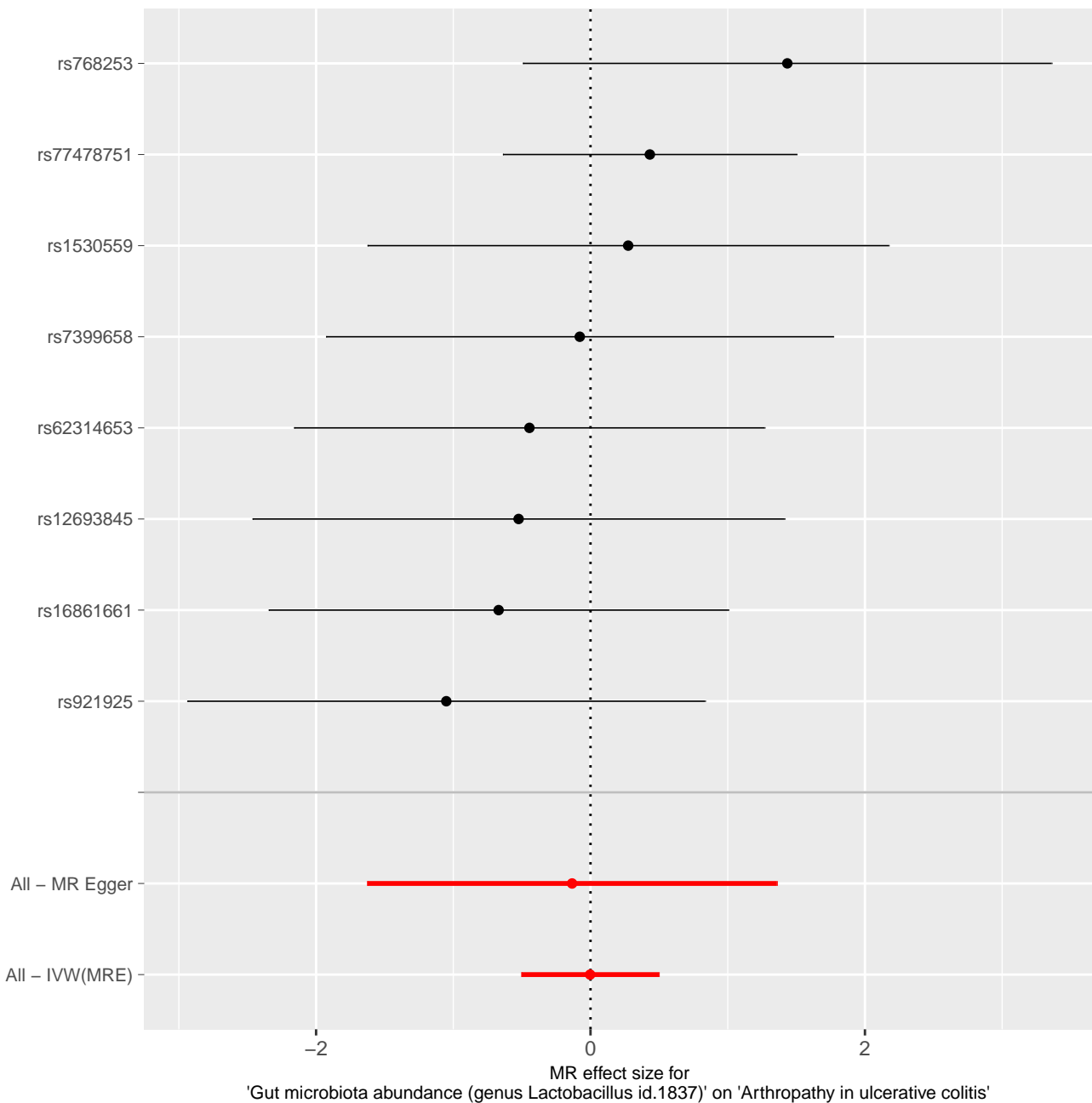


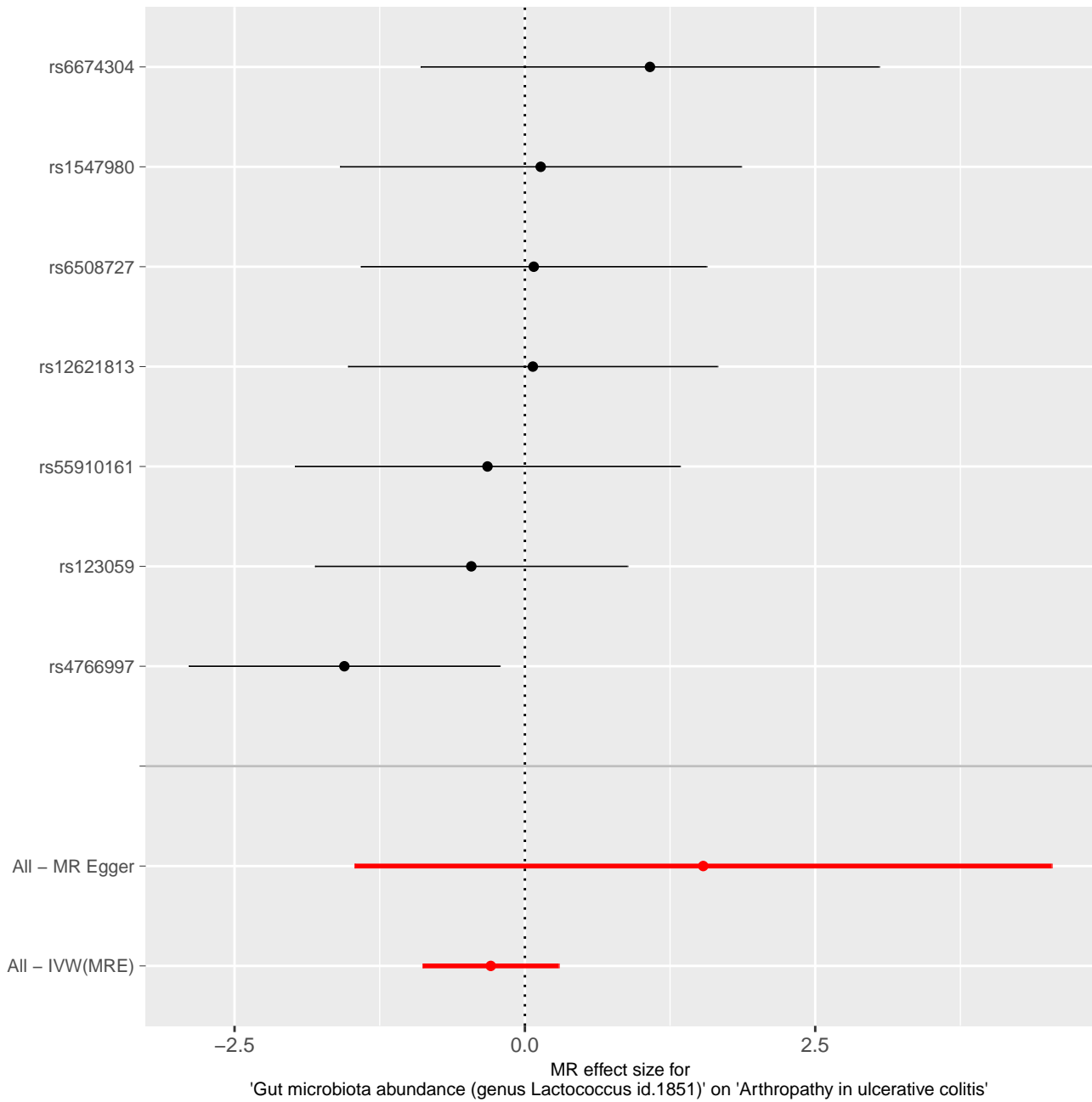


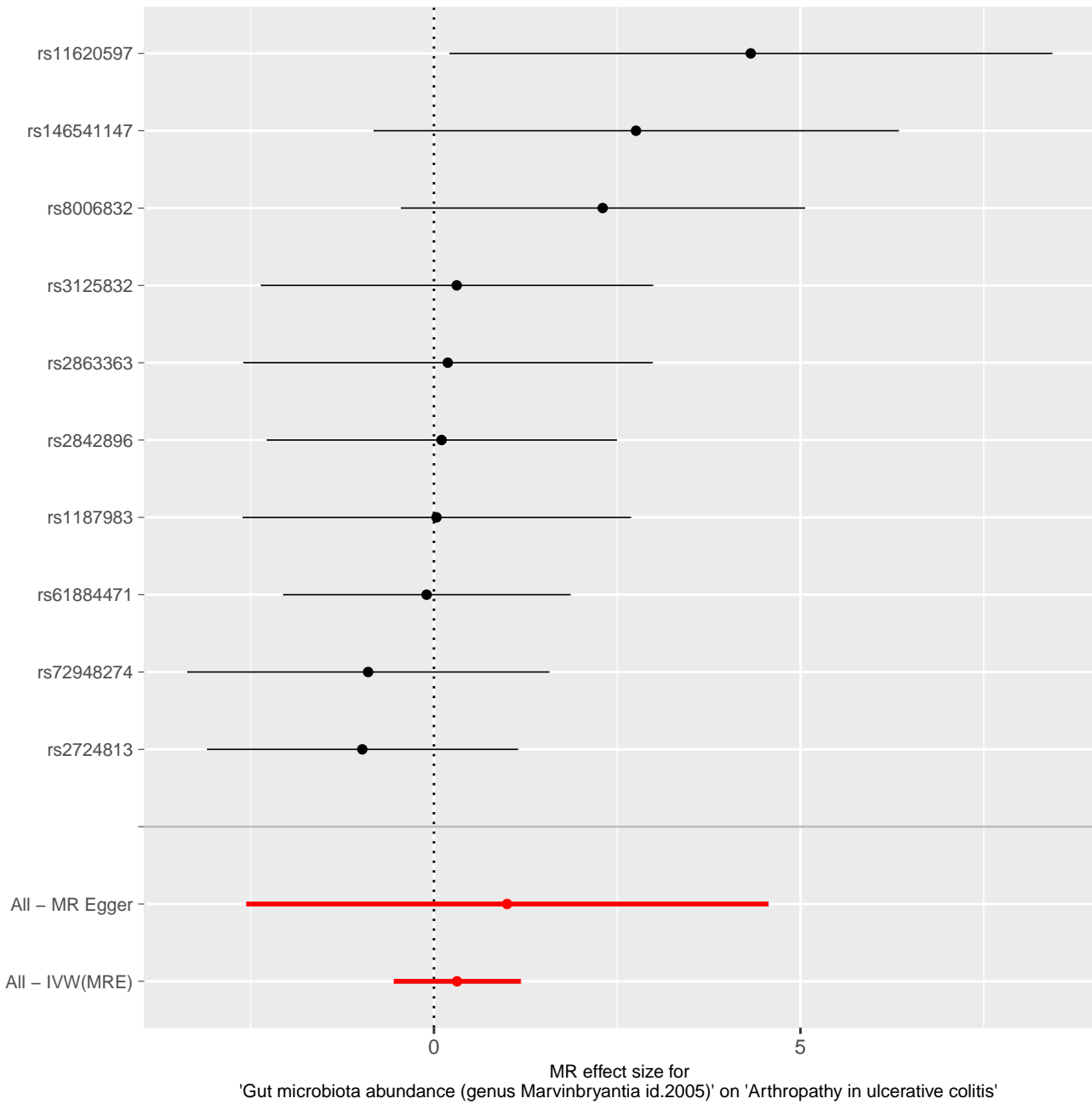


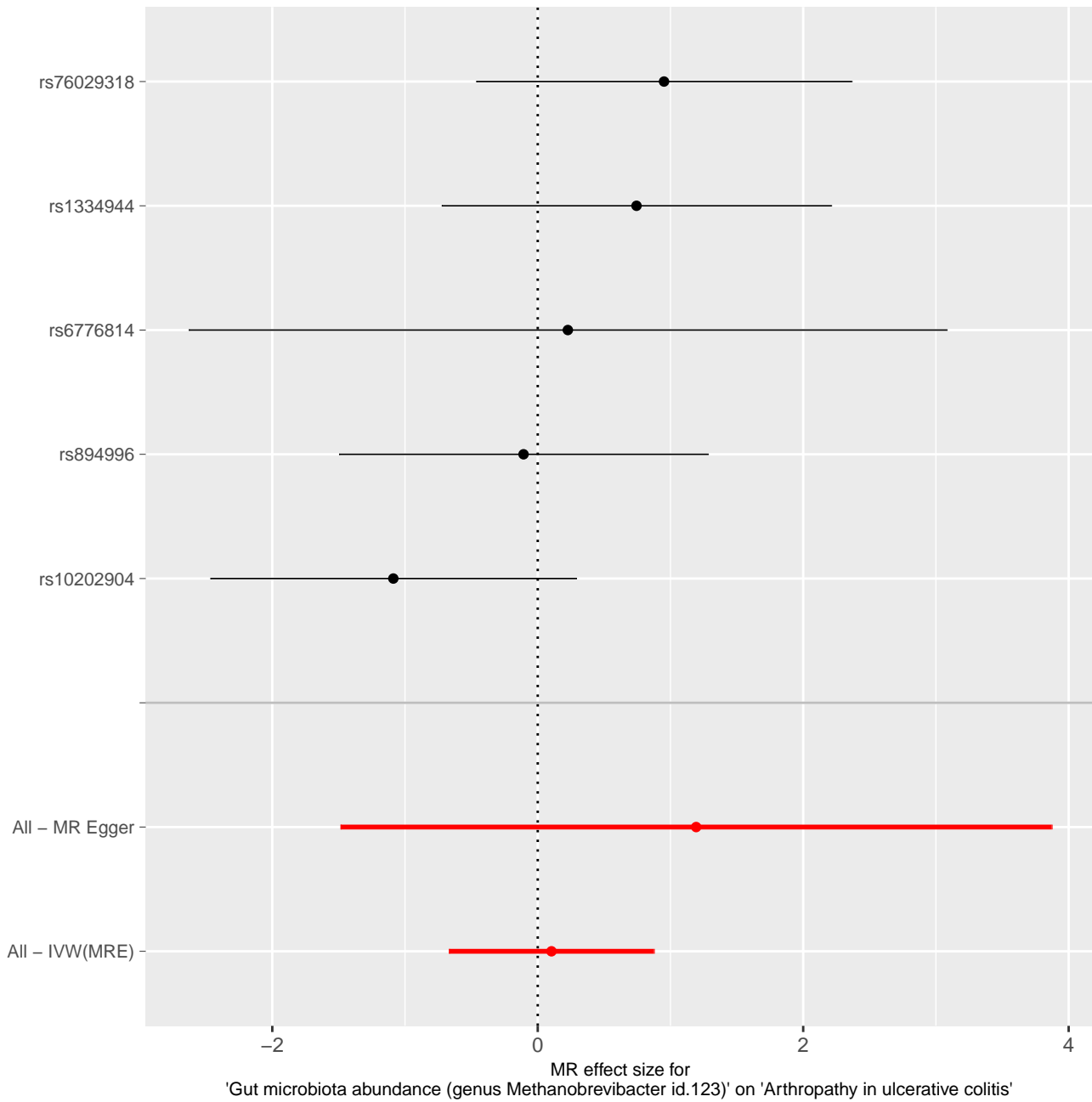


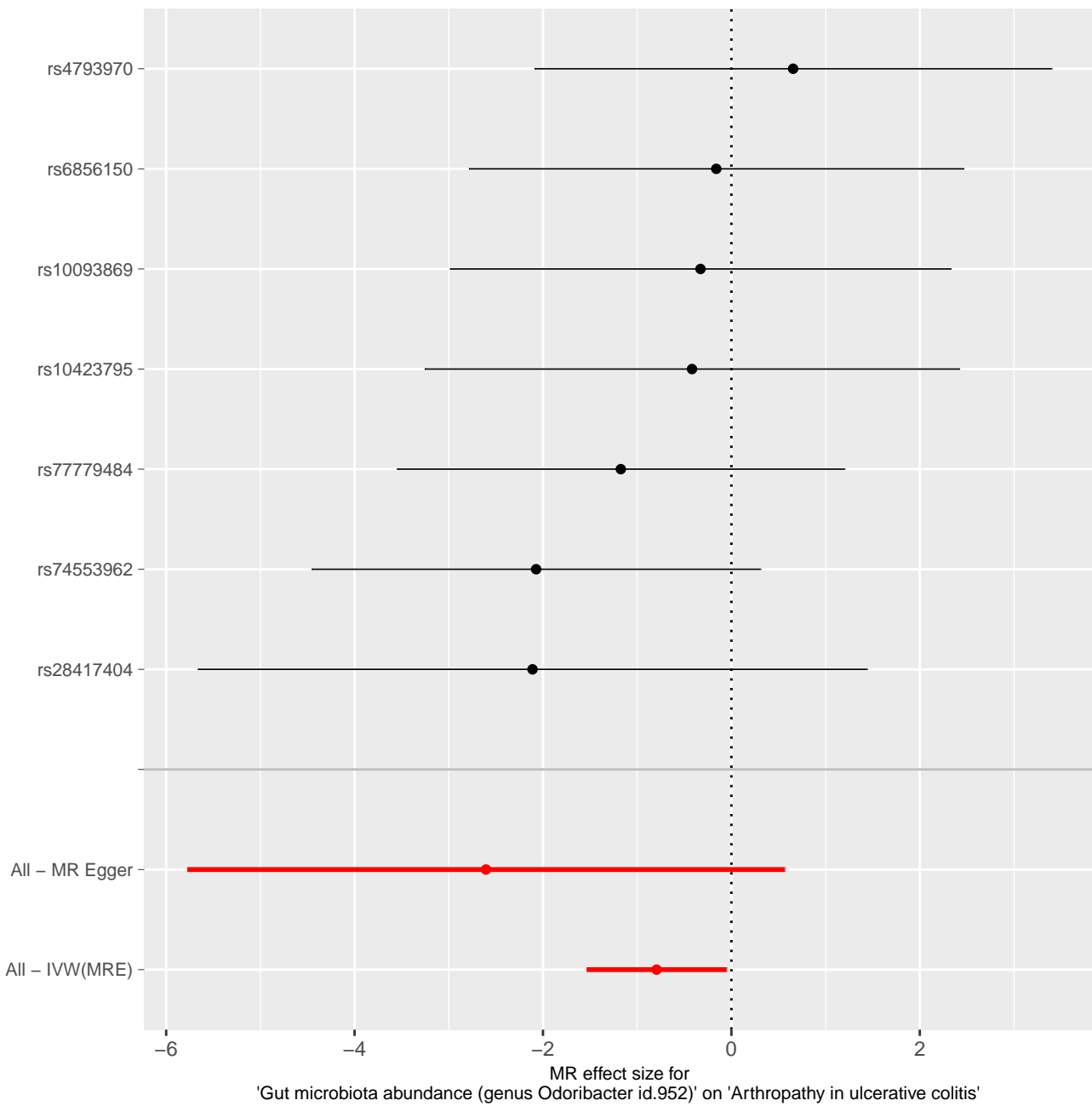


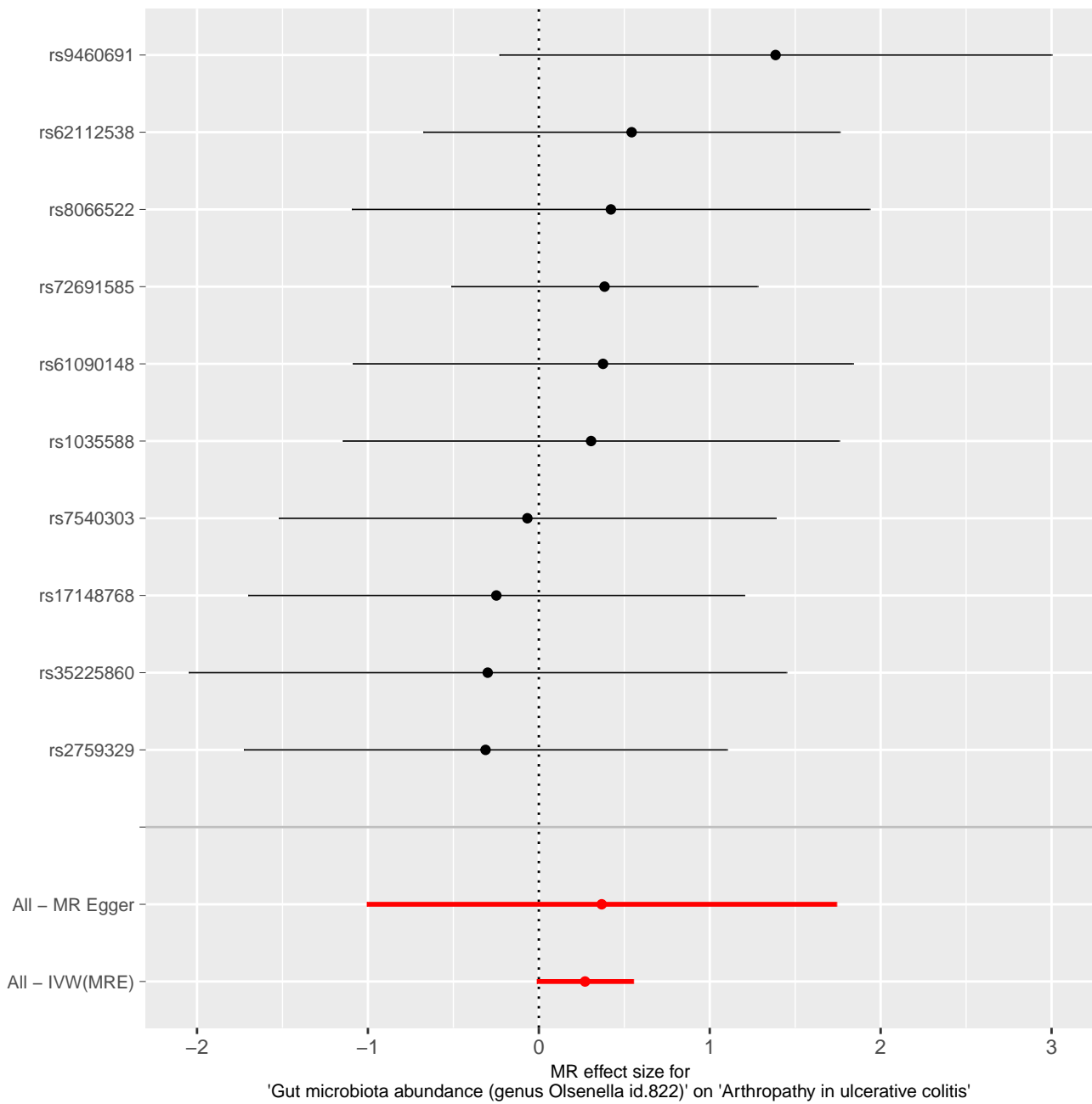


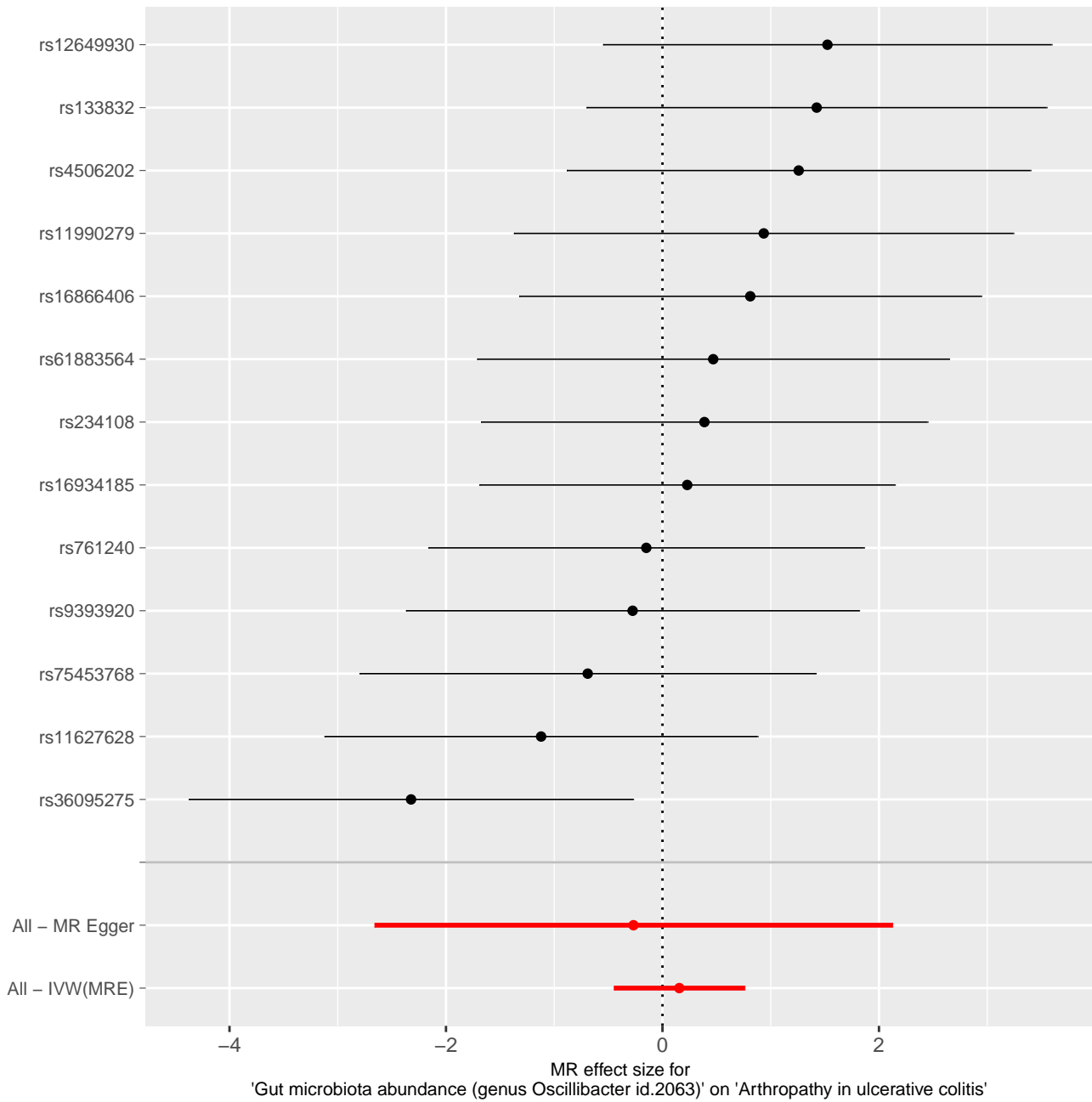


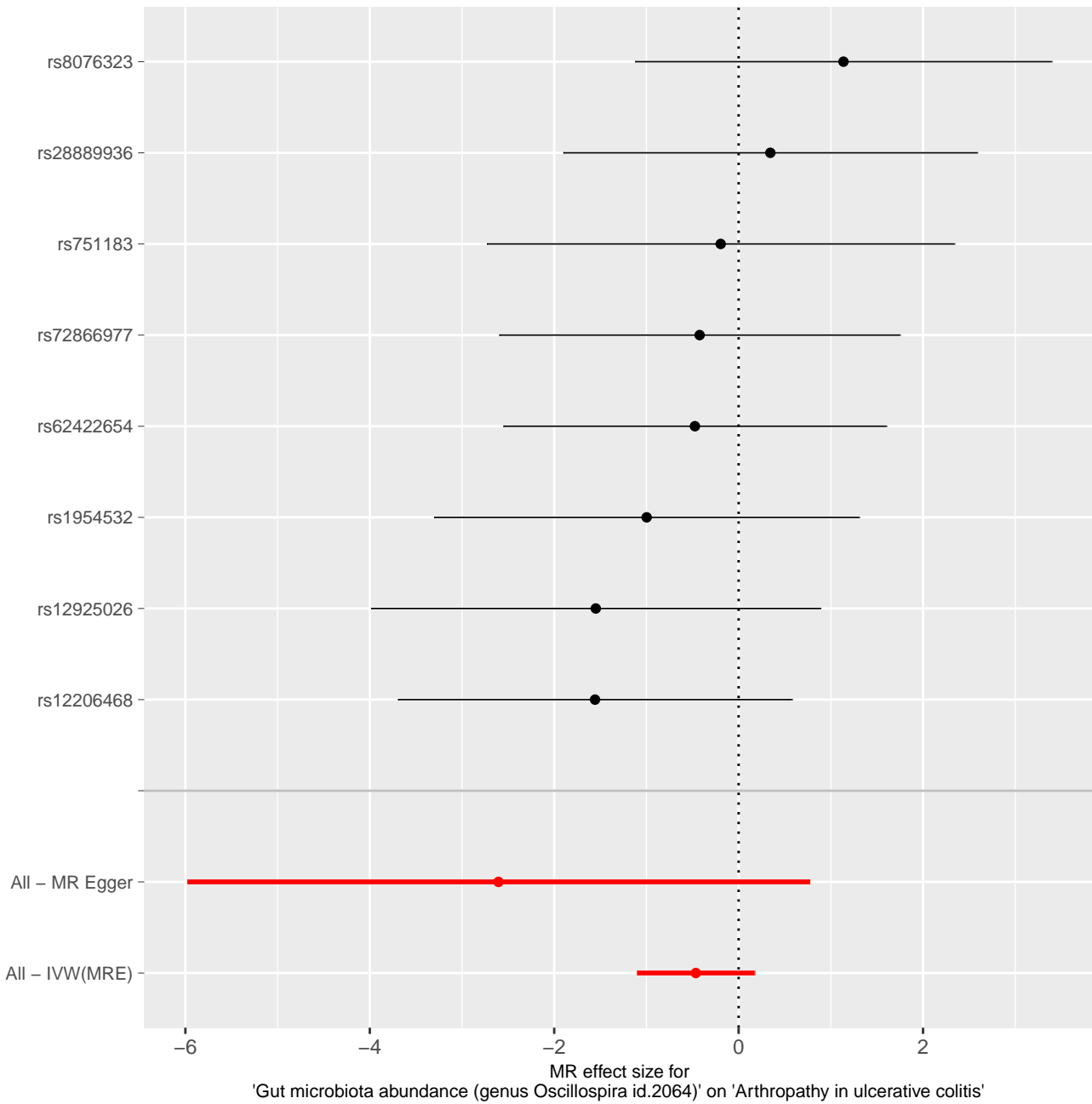




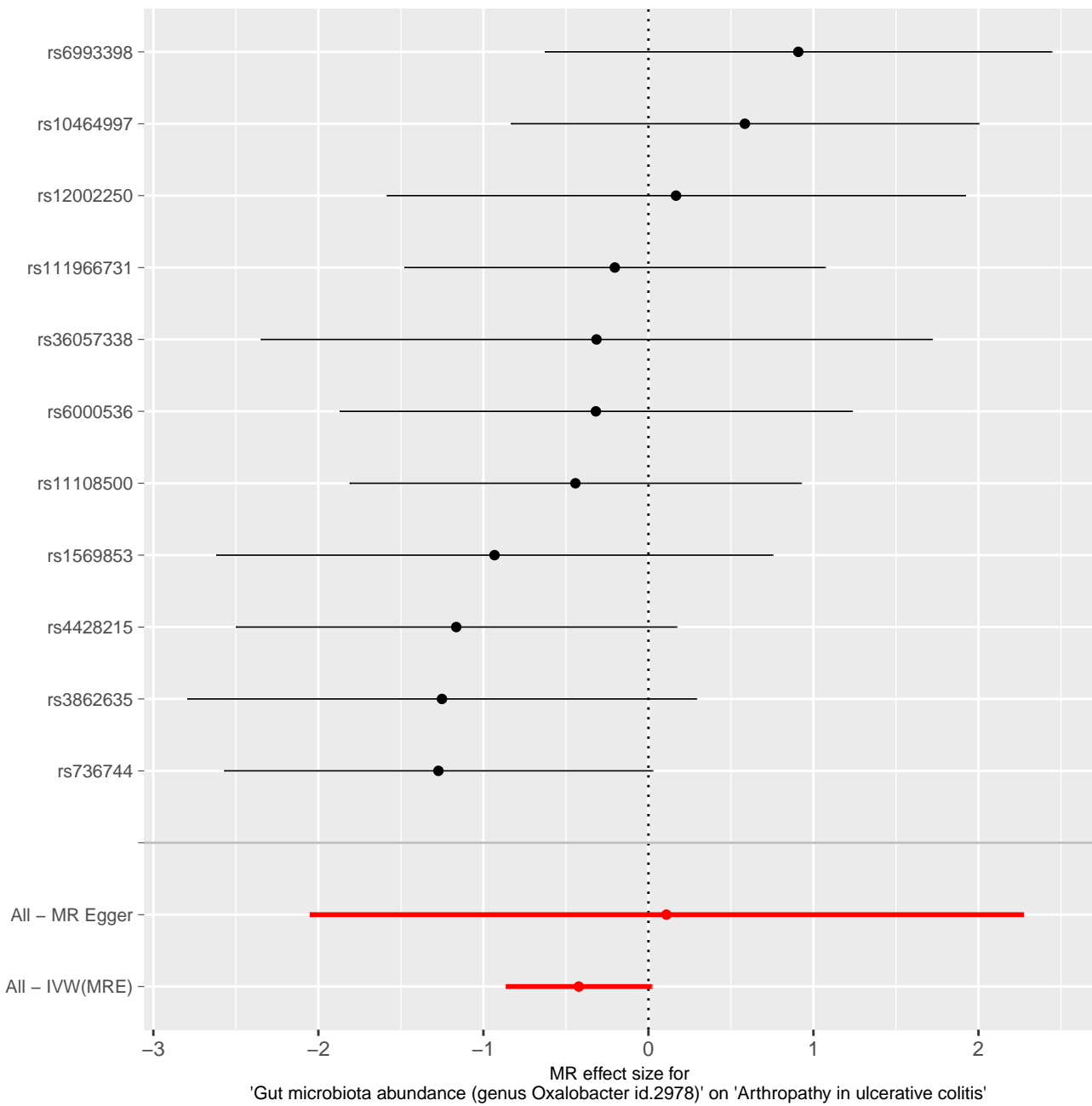


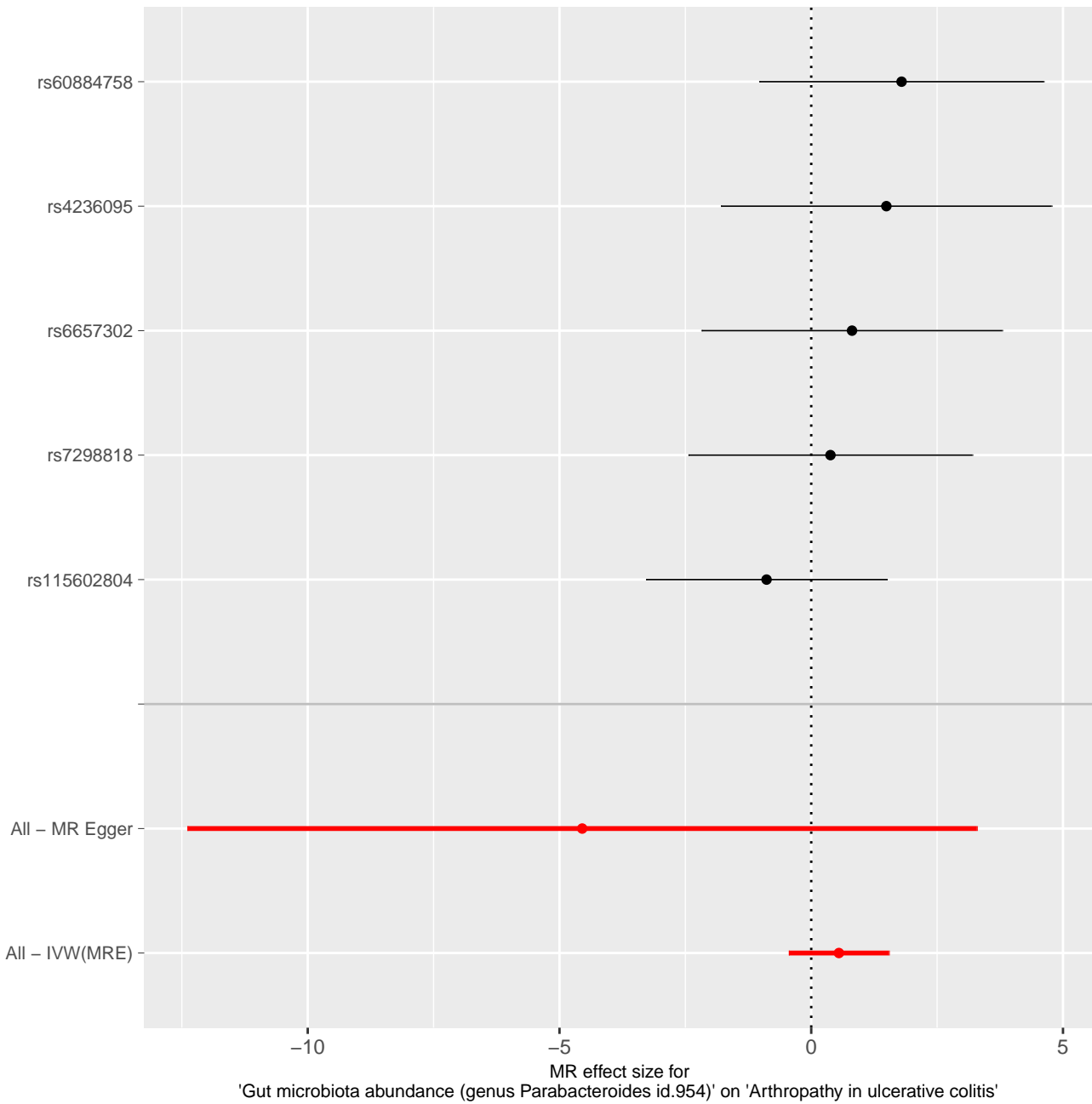


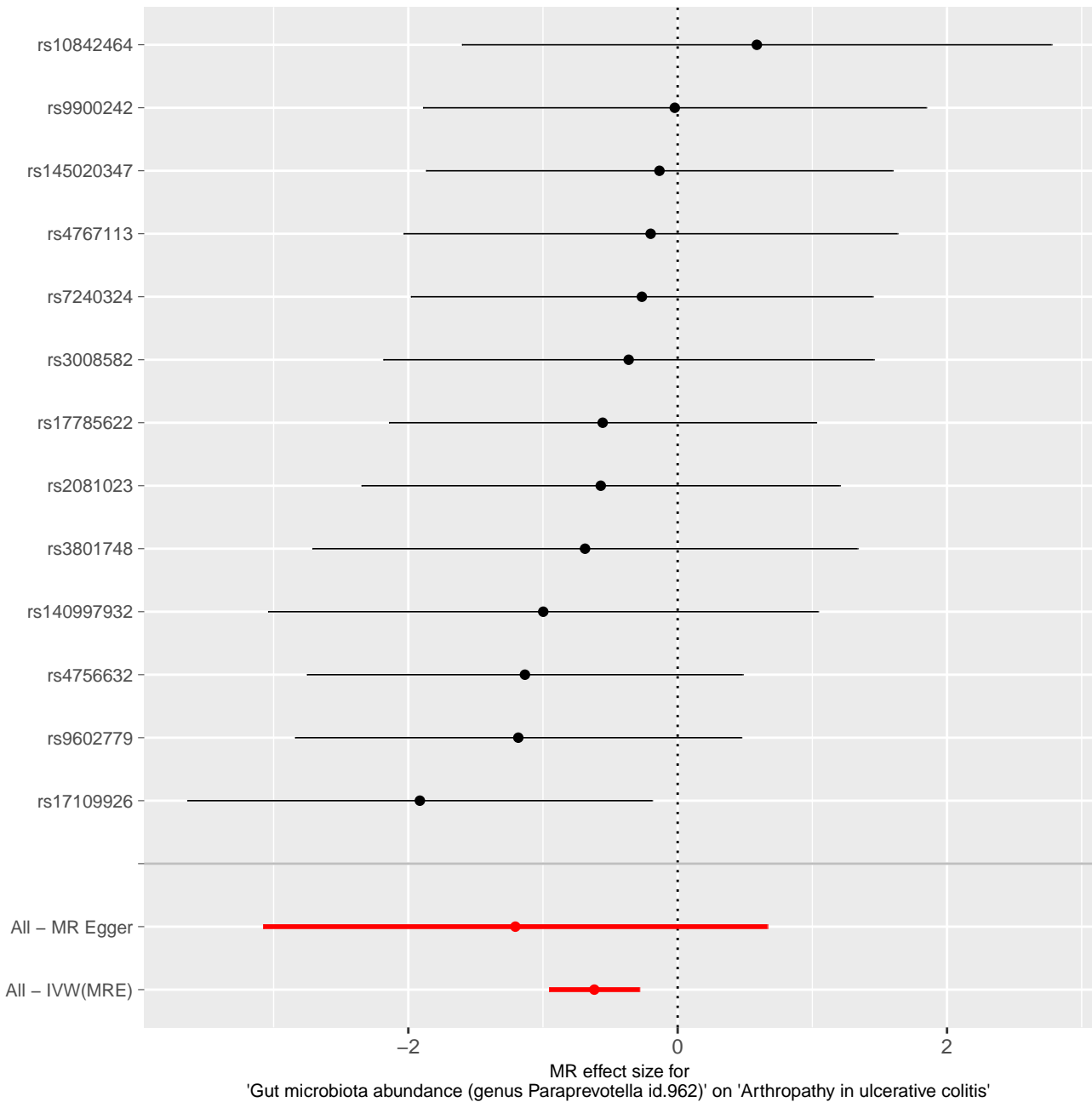




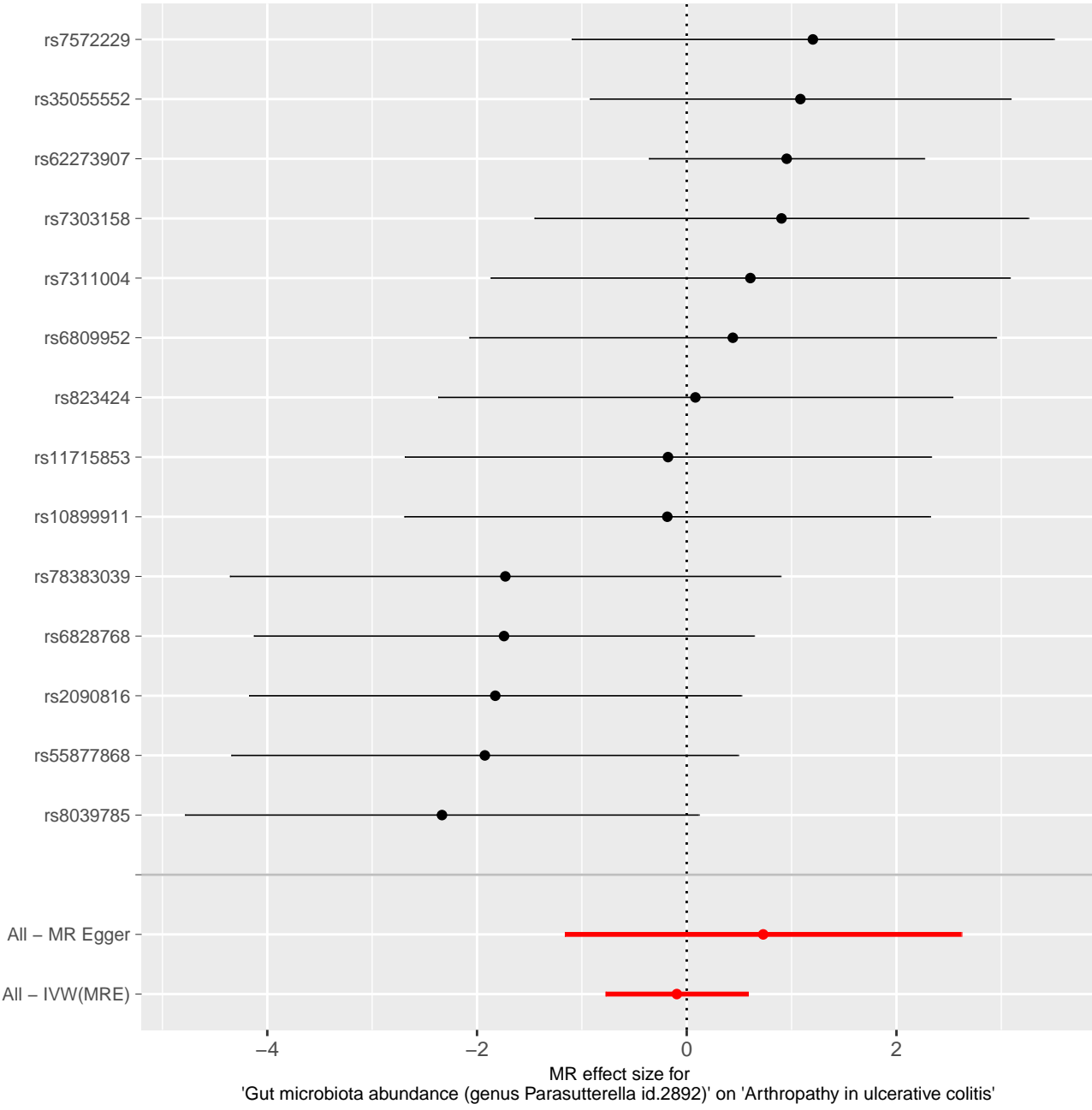
Batch 862 : Gut microbiota abundance (genus Oxalobacter id.2978) on Arthropathy in ulcerative colitis

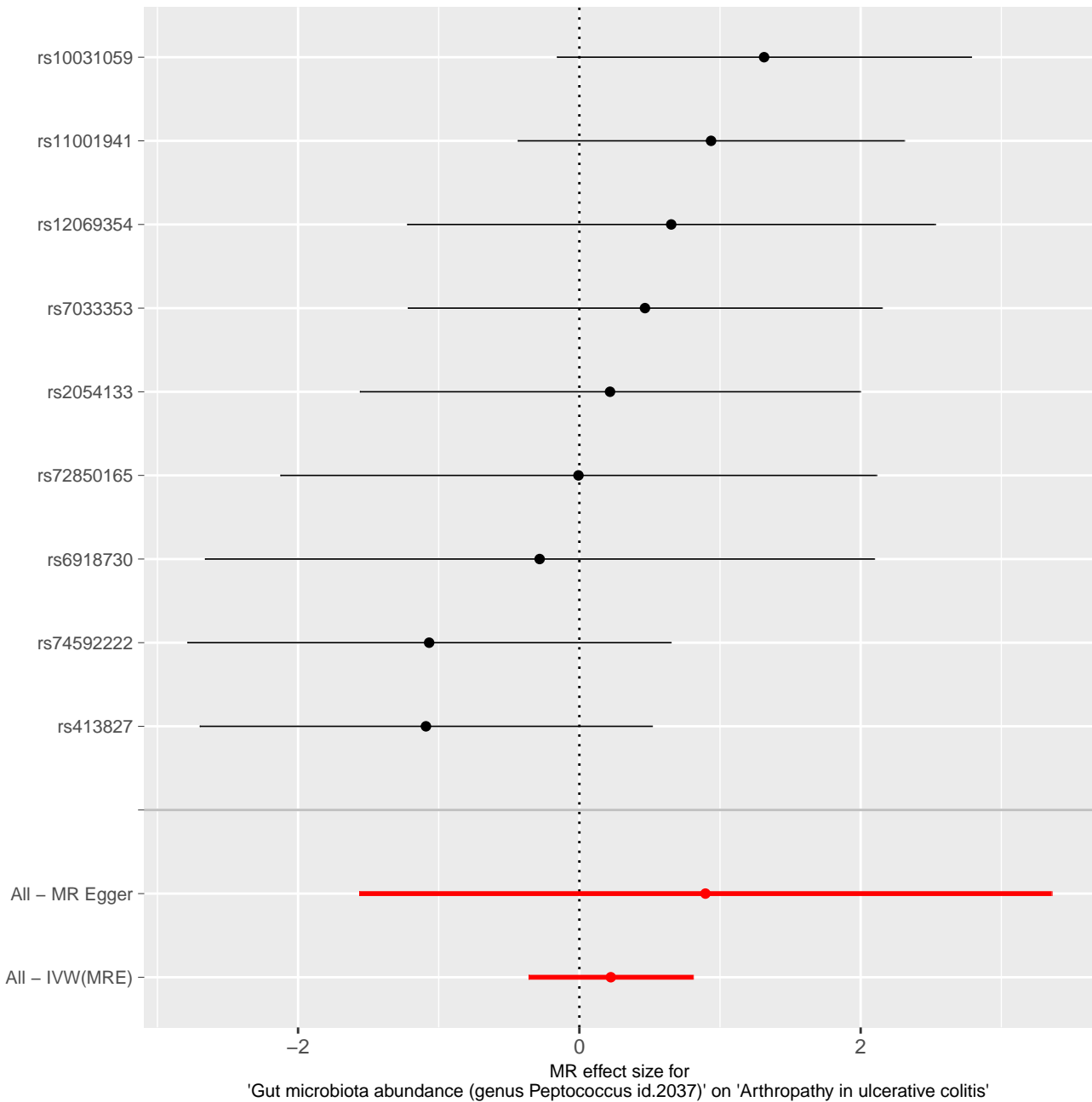


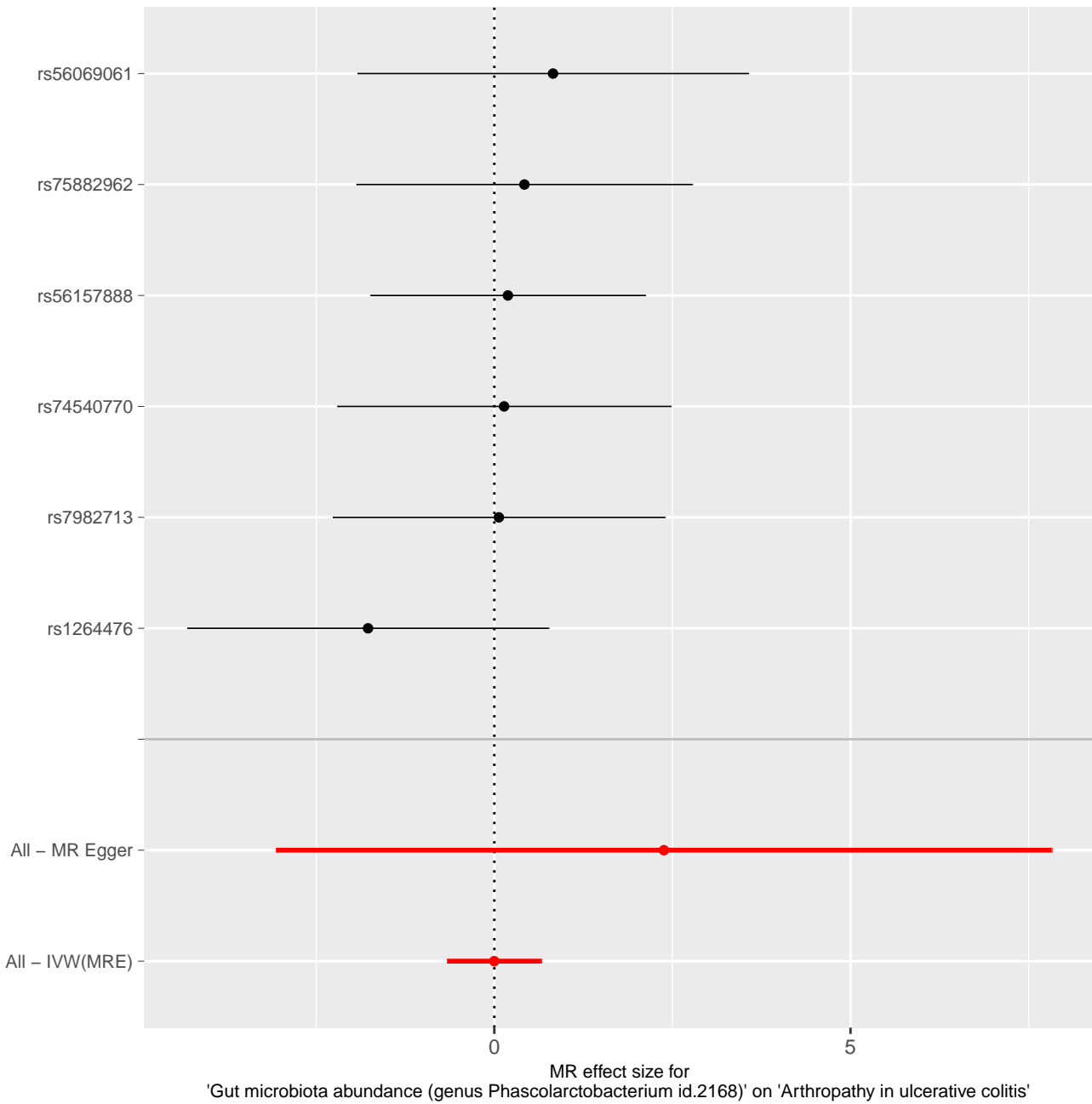




Batch 865 : Gut microbiota abundance (genus Parasutterella id.2892) on Arthropathy in ulcerative colitis

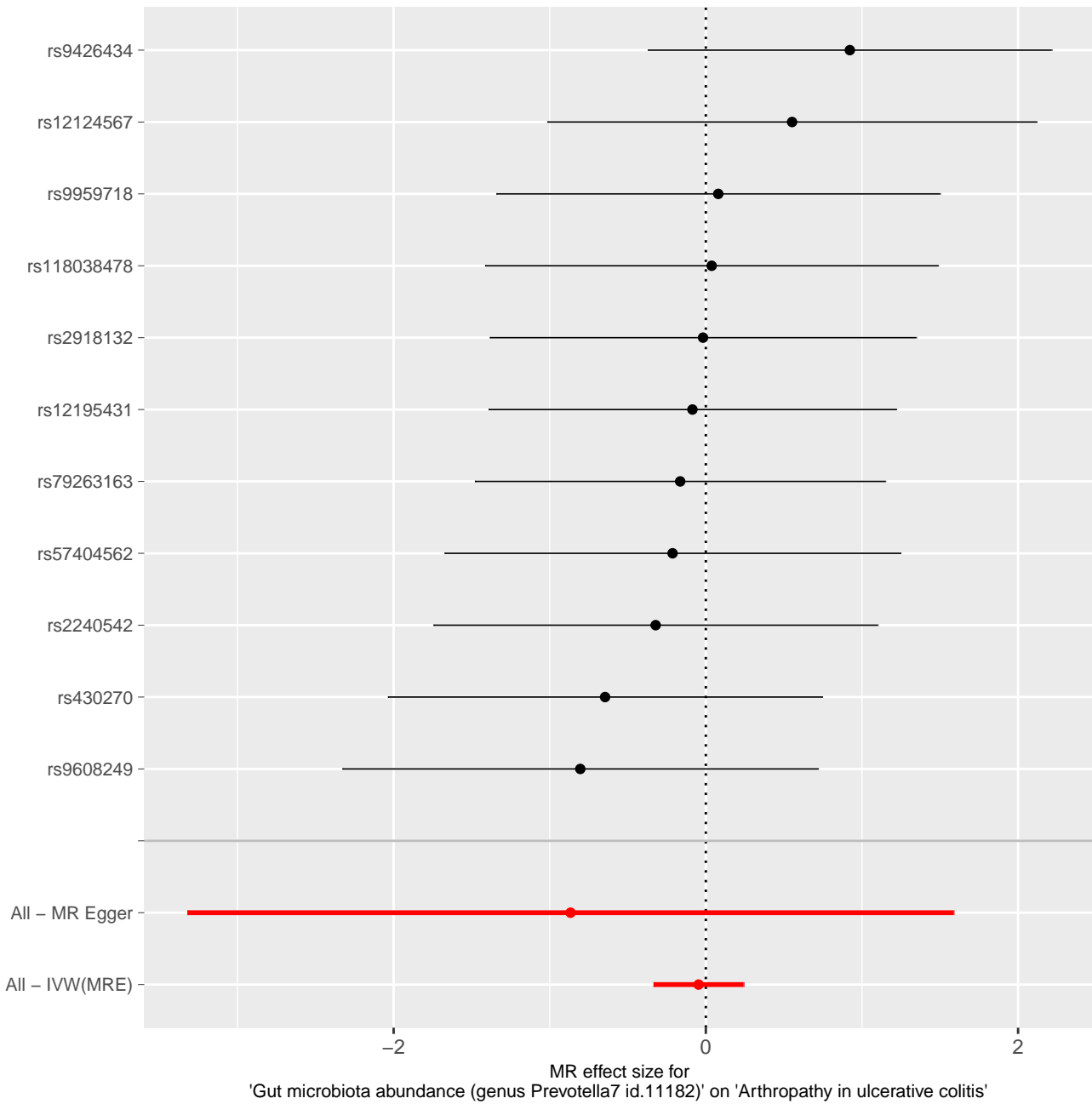




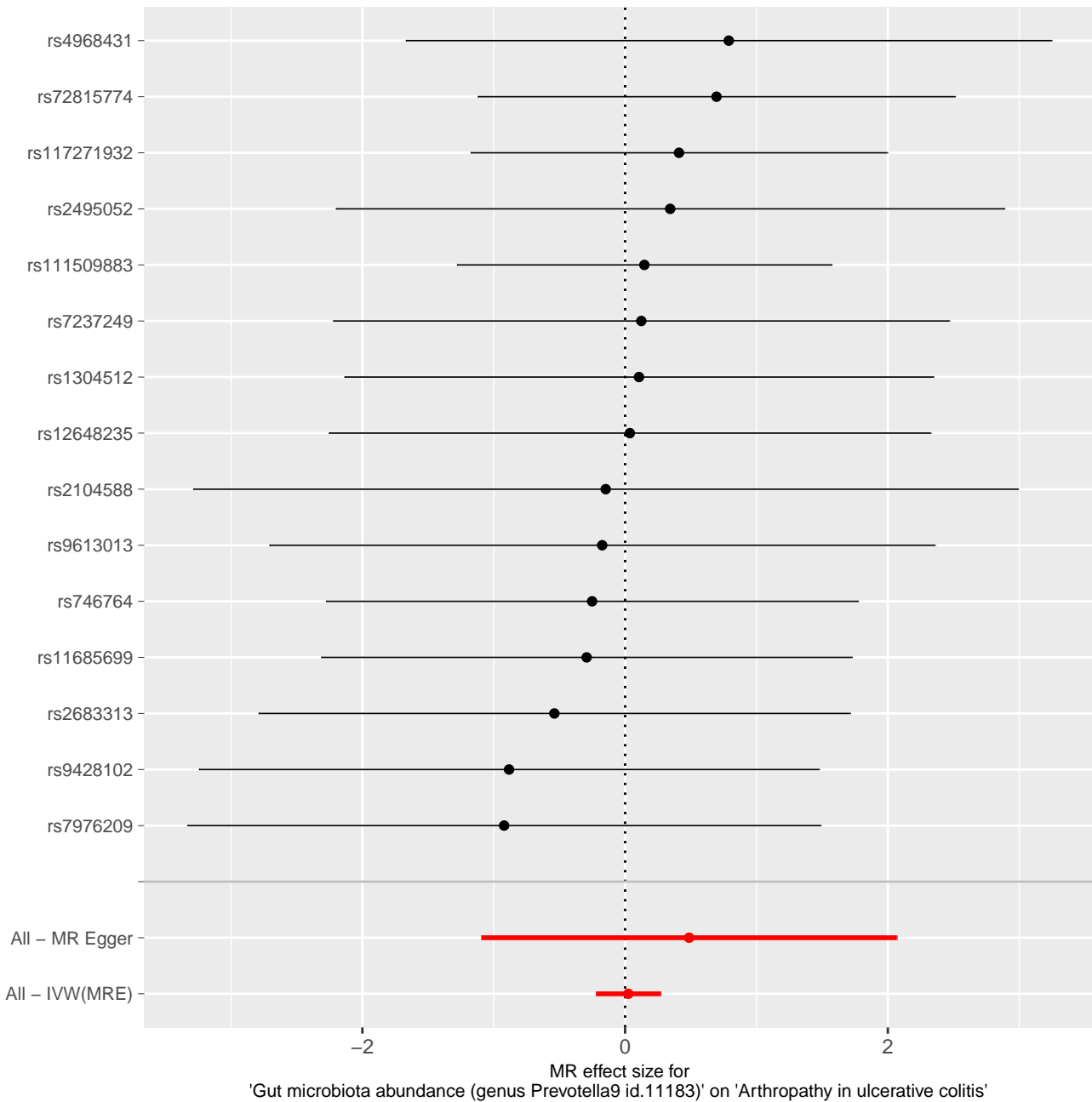


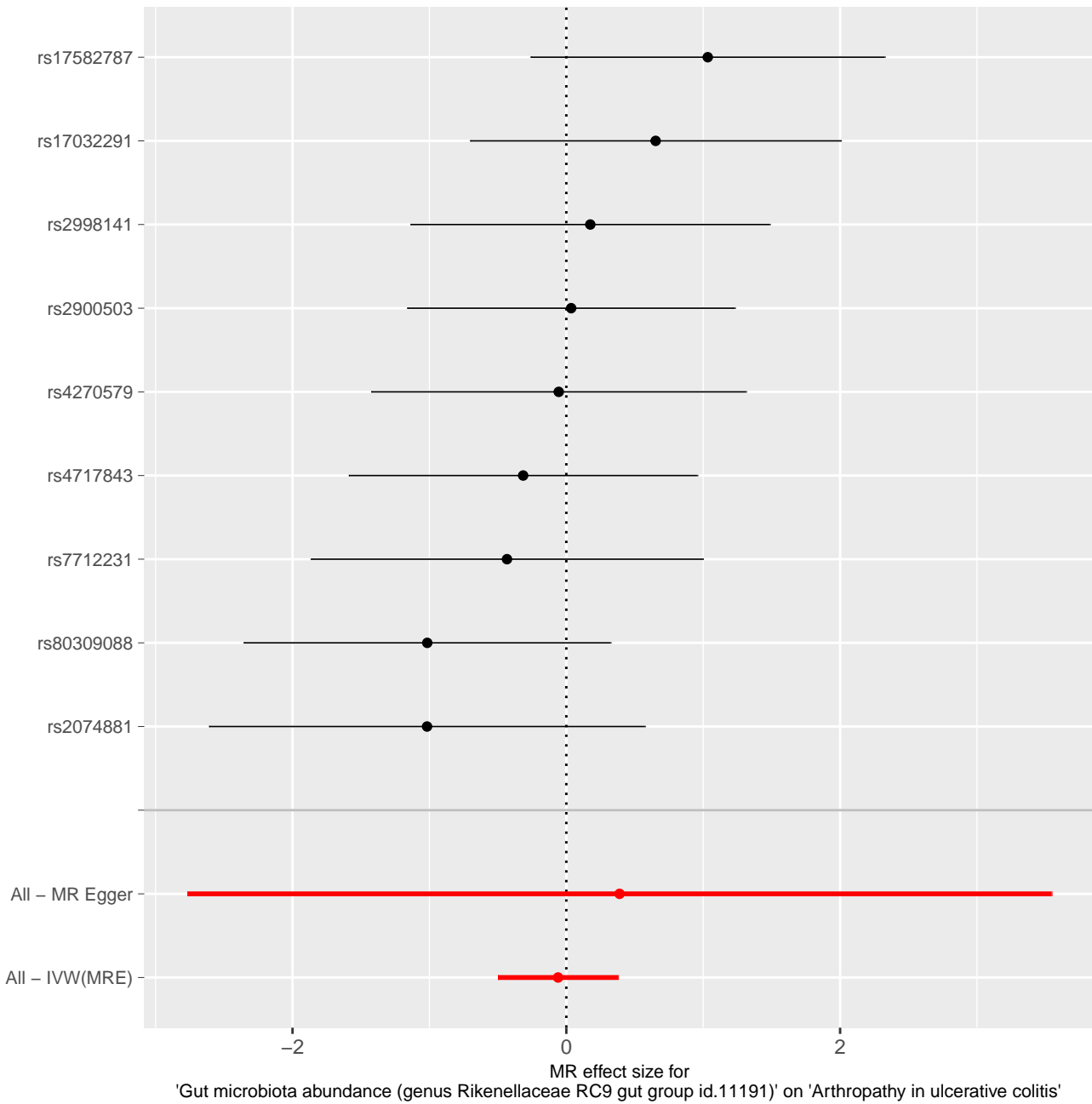
MR effect size for 'Gut microbiota abundance (genus Phascolarctobacterium id.2168)' on 'Arthropathy in ulcerative colitis'

Batch 868 : Gut microbiota abundance (genus Prevotella7 id.11182) on Arthropathy in ulcerative colitis

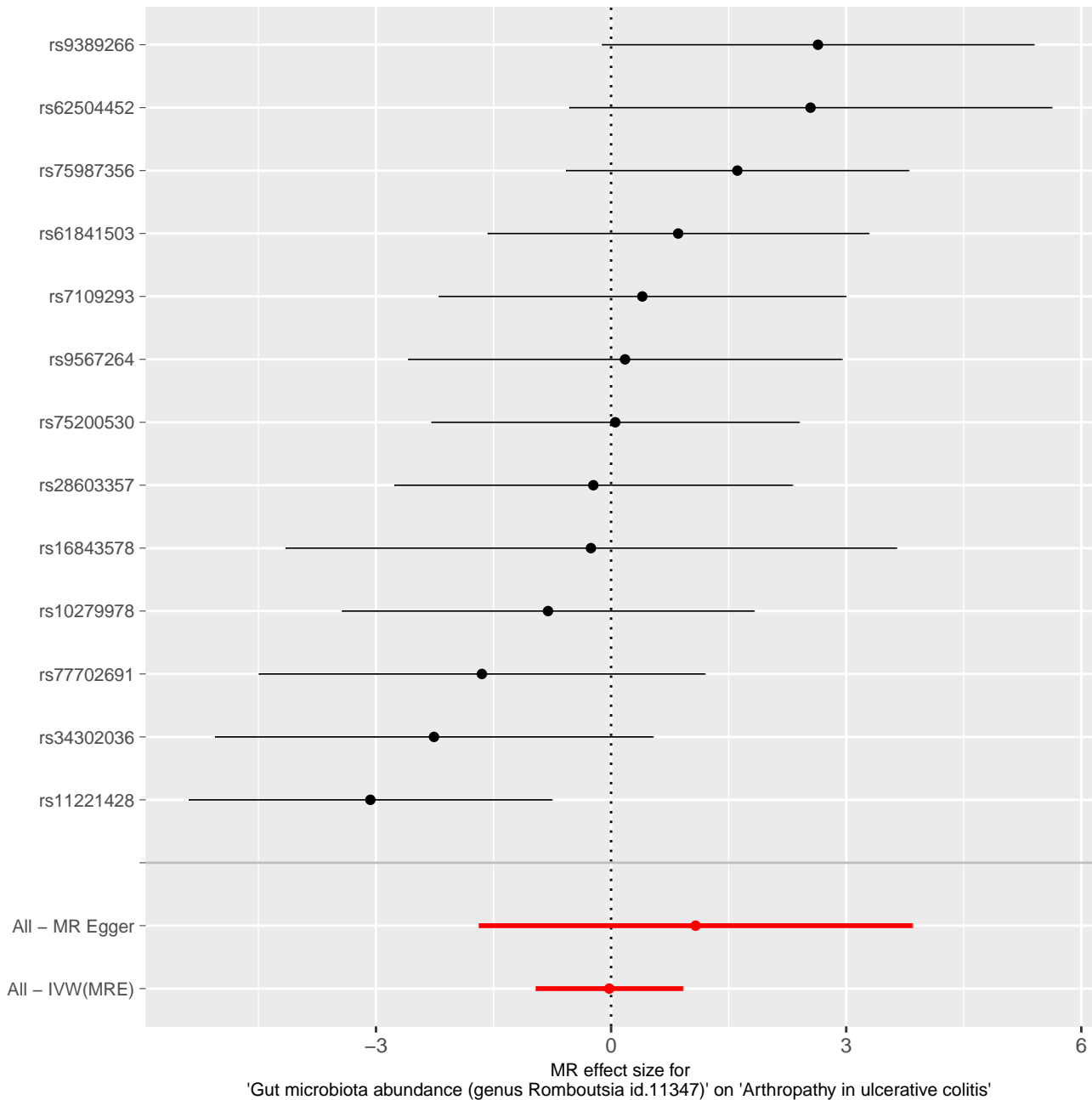


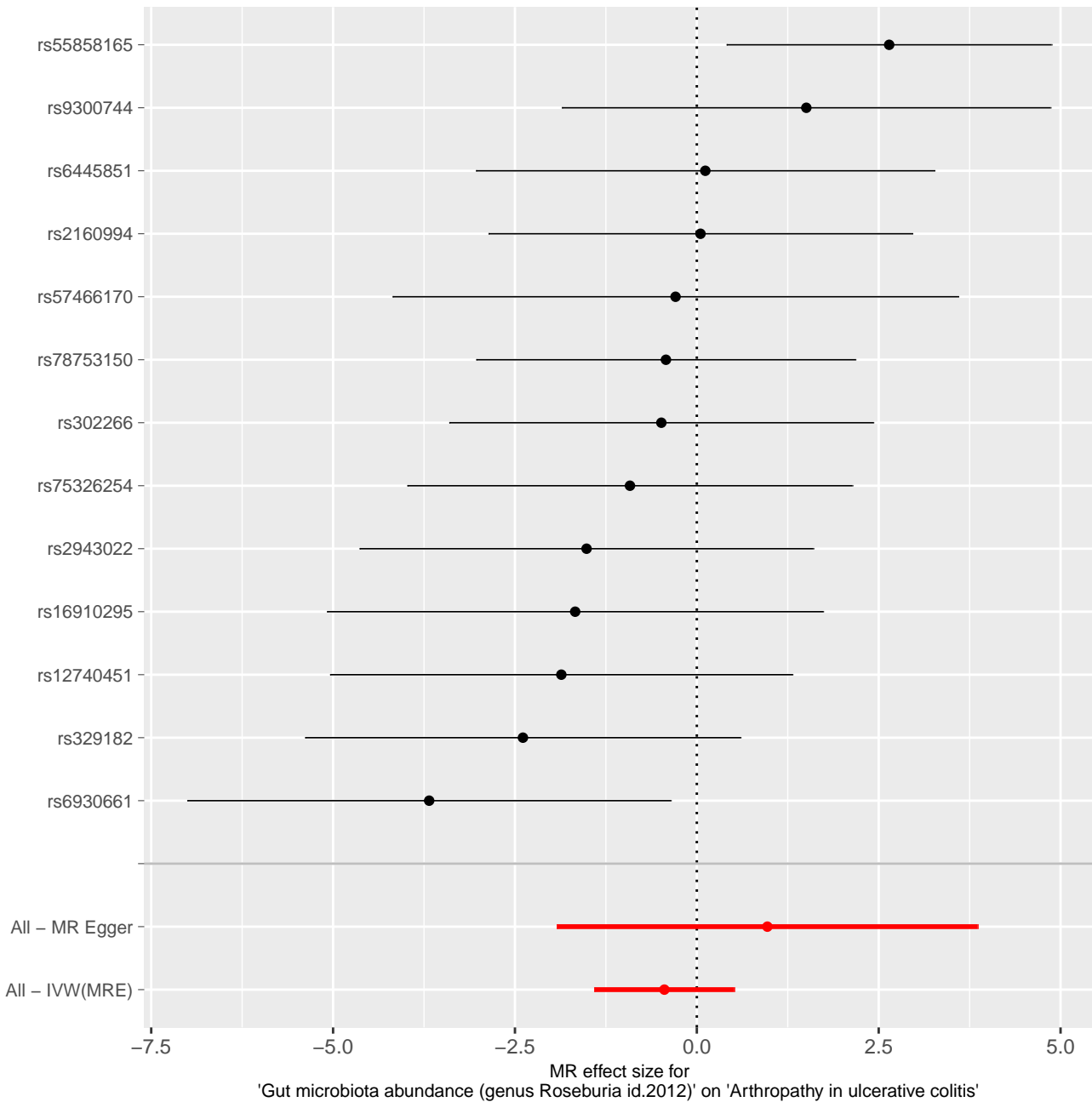
Batch 869 : Gut microbiota abundance (genus Prevotella9 id.11183) on Arthropathy in ulcerative colitis



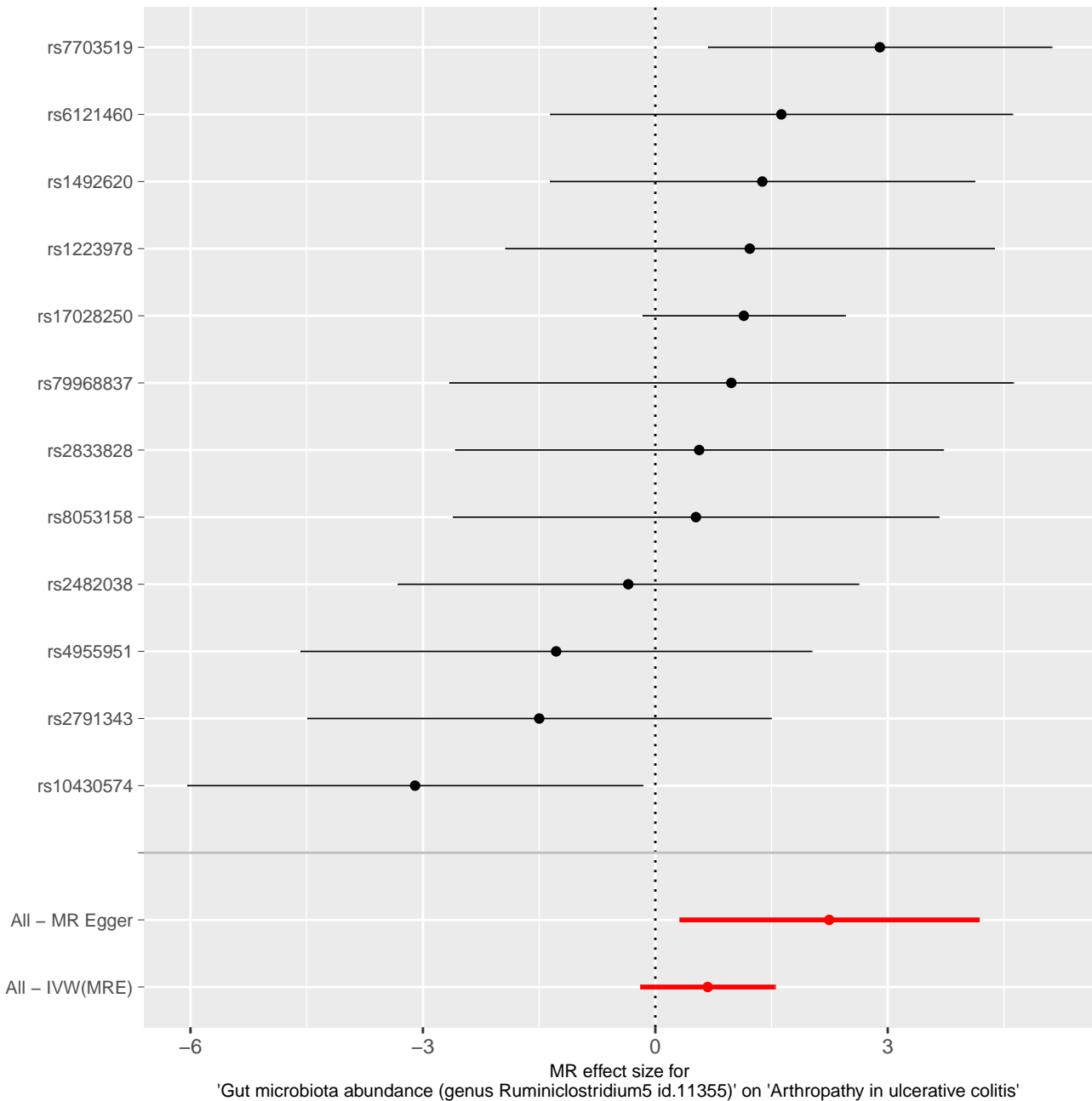


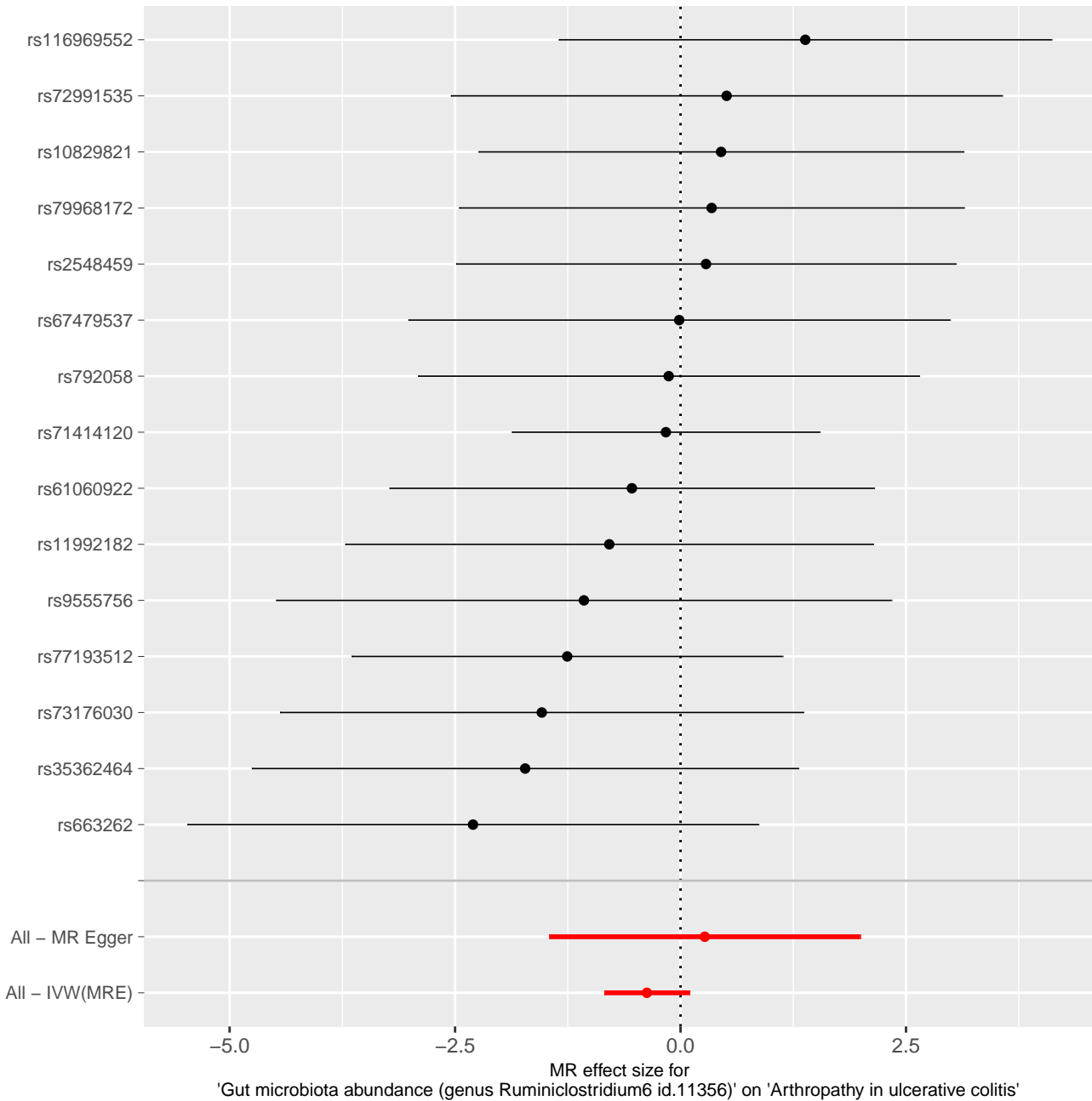
Batch 871 : Gut microbiota abundance (genus Romboutsia id.11347) on Arthropathy in ulcerative colitis

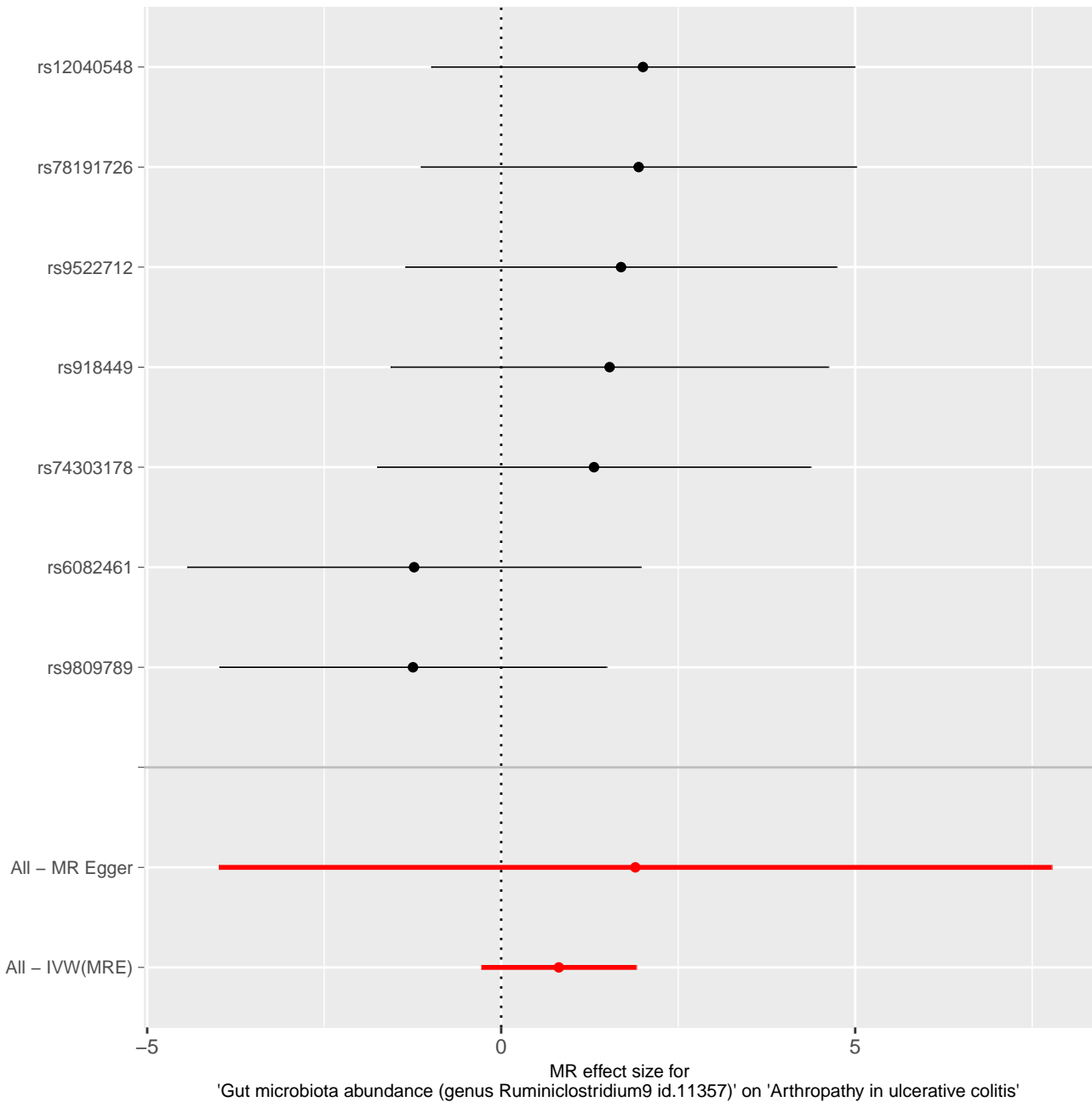


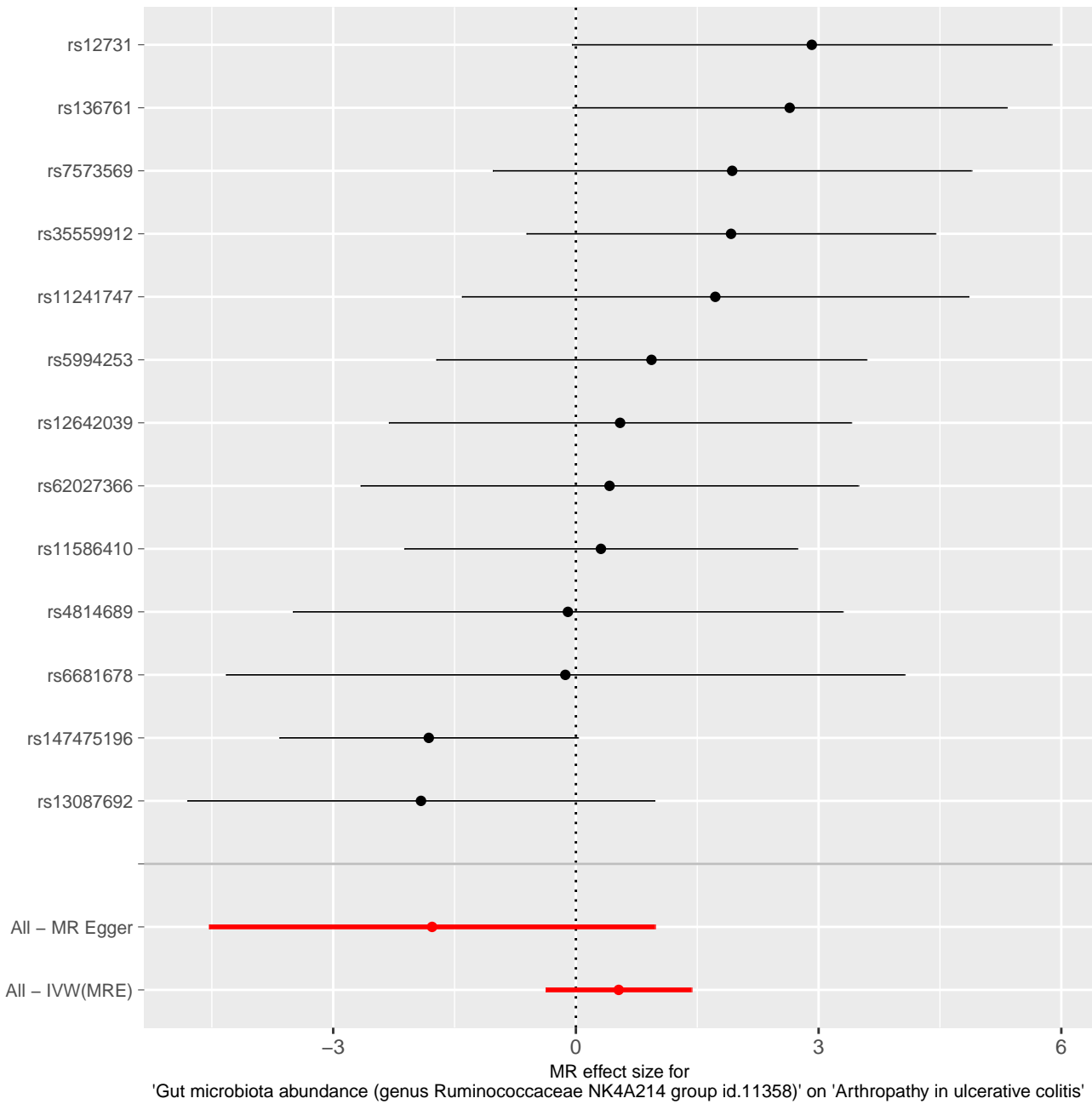


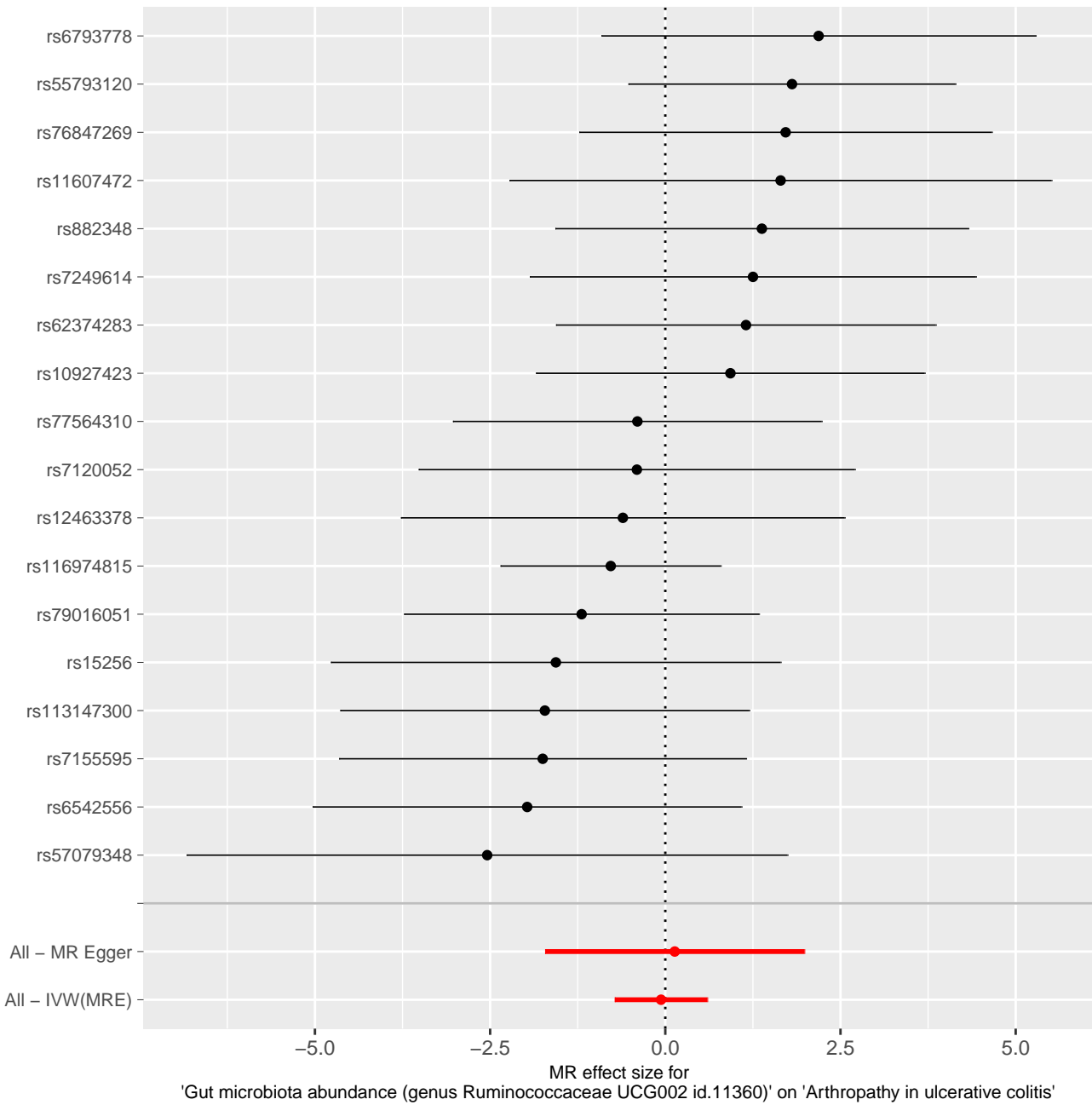
Batch 873 : Gut microbiota abundance (genus Ruminiclostridium5 id.11355) on Arthropathy in ulcerative colitis

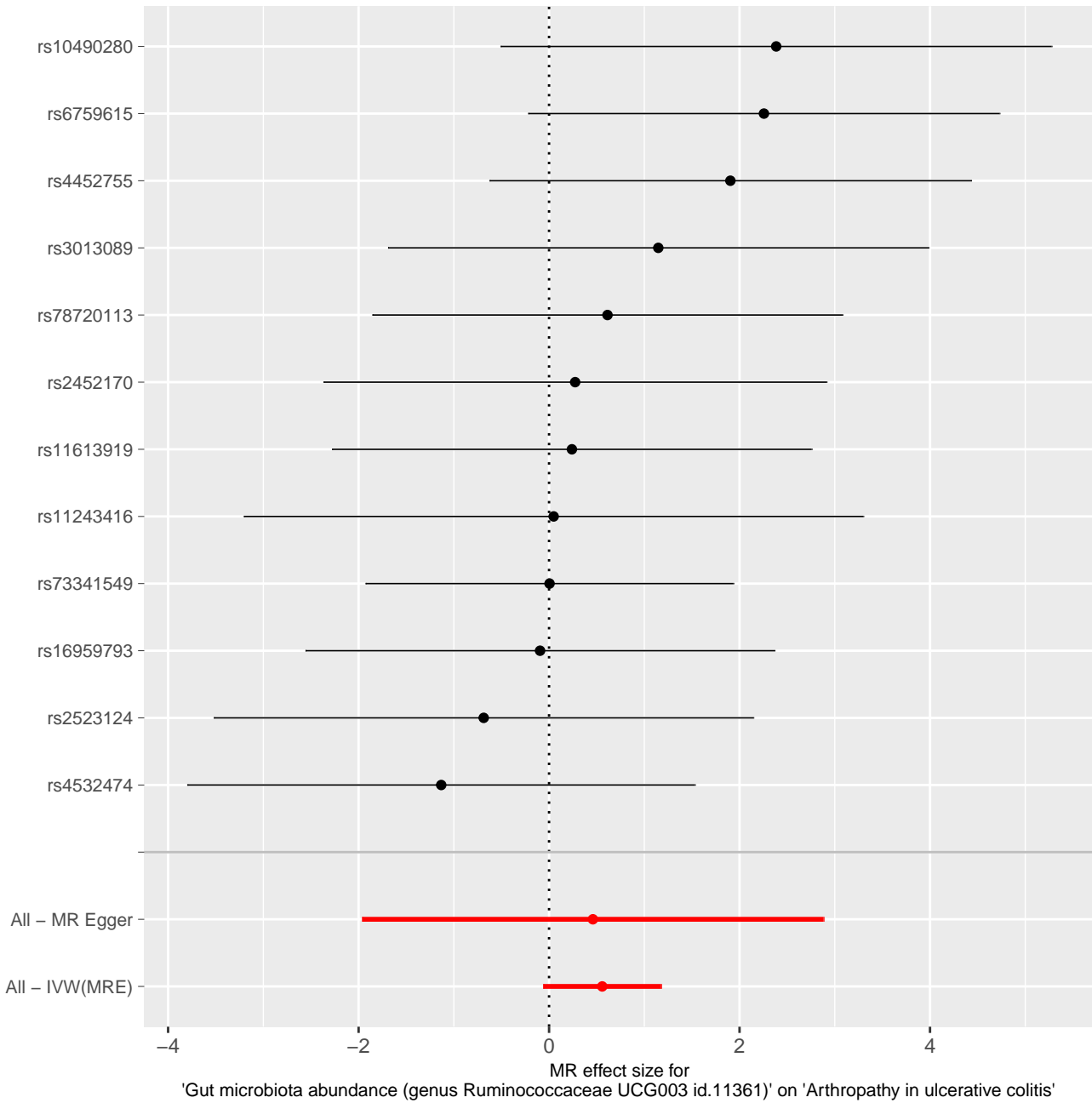


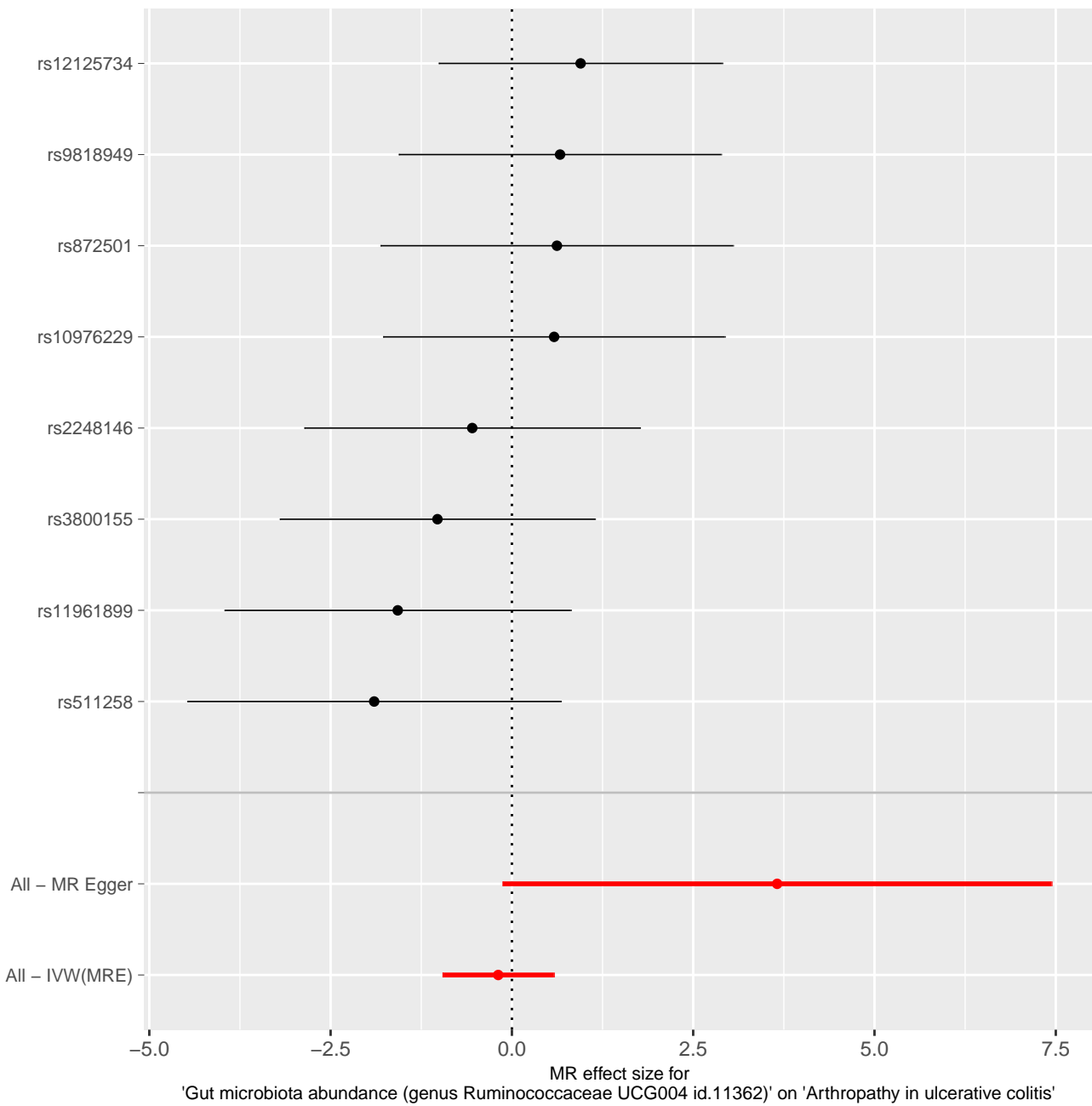


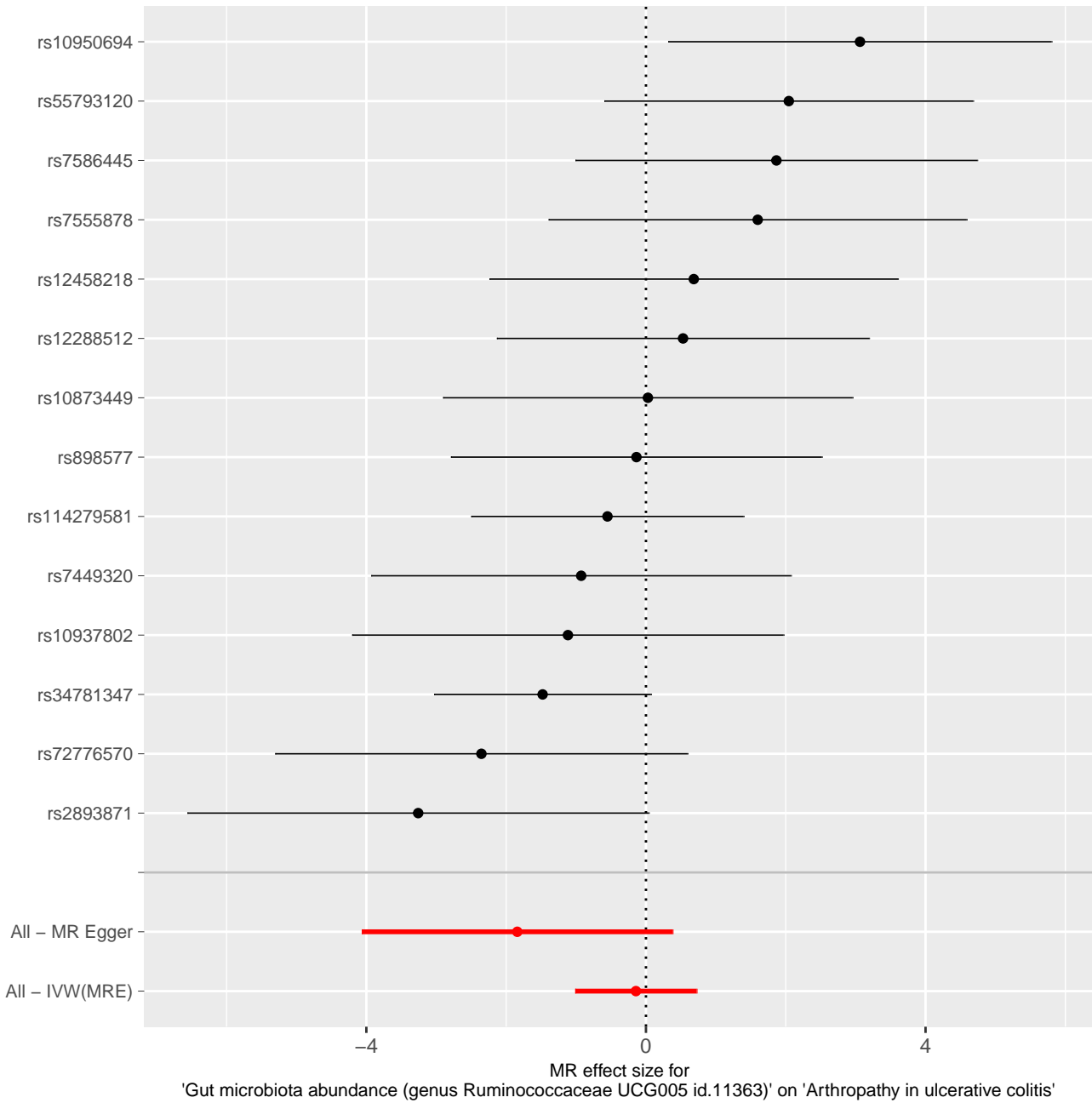


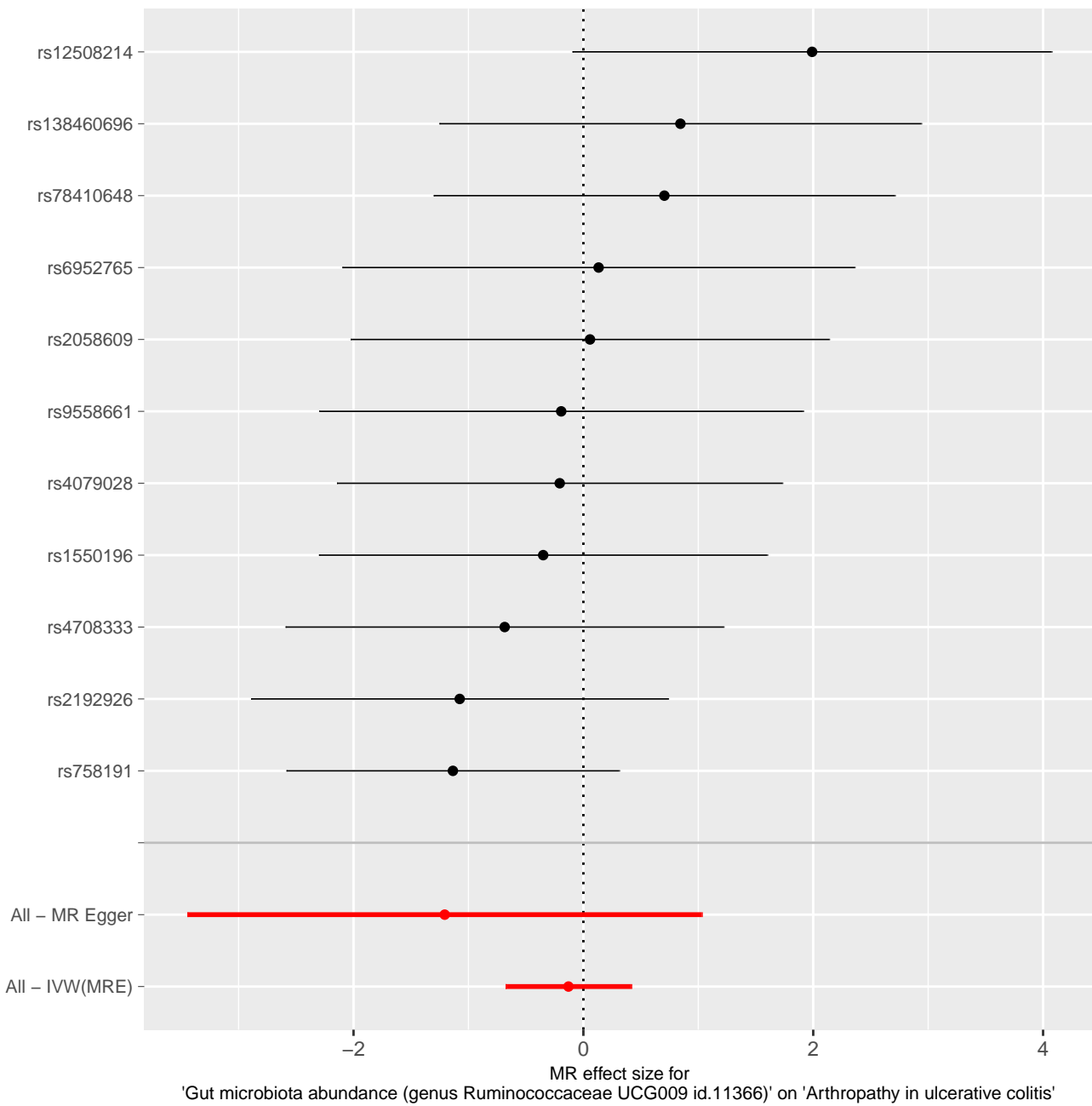


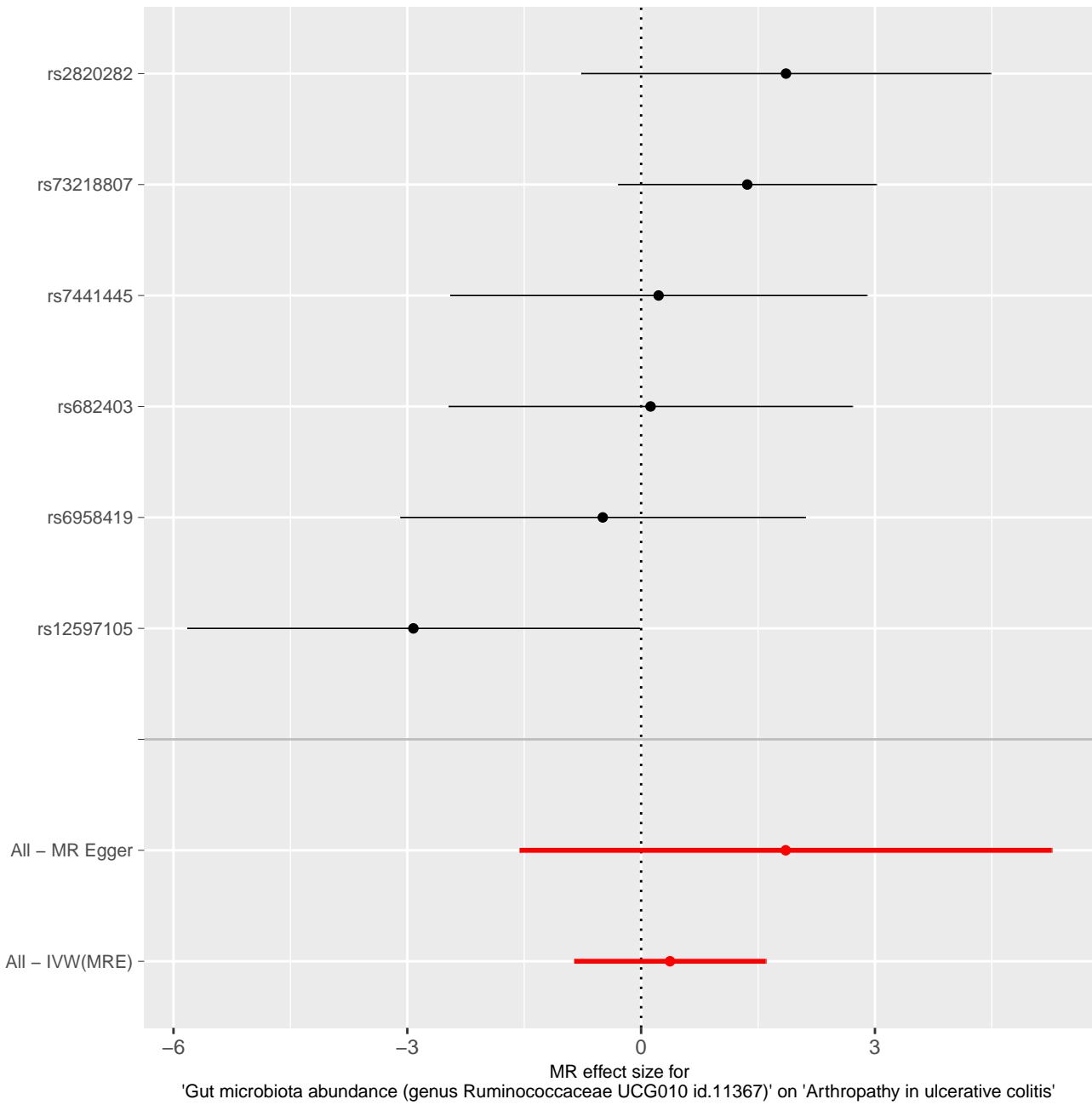


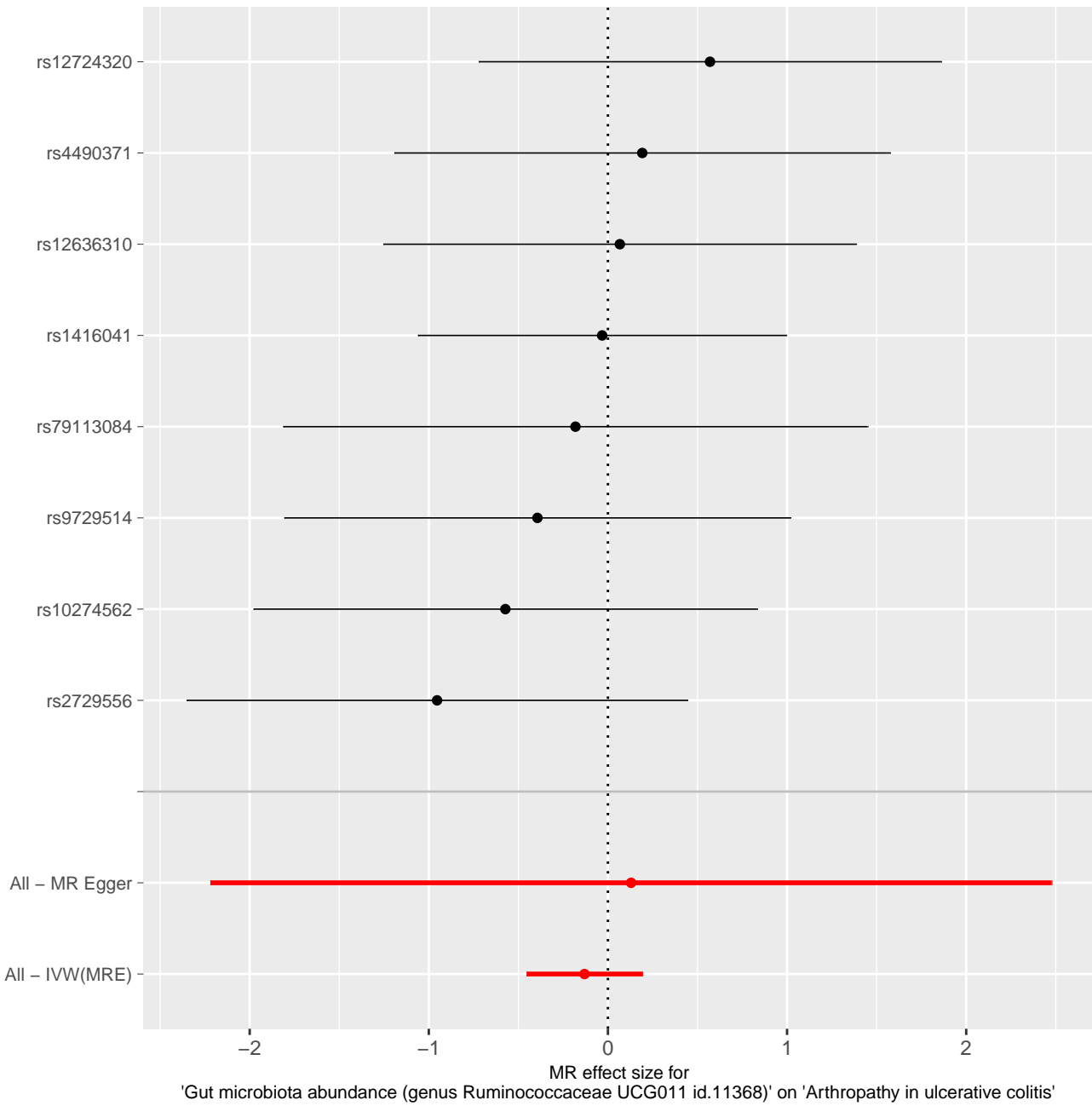


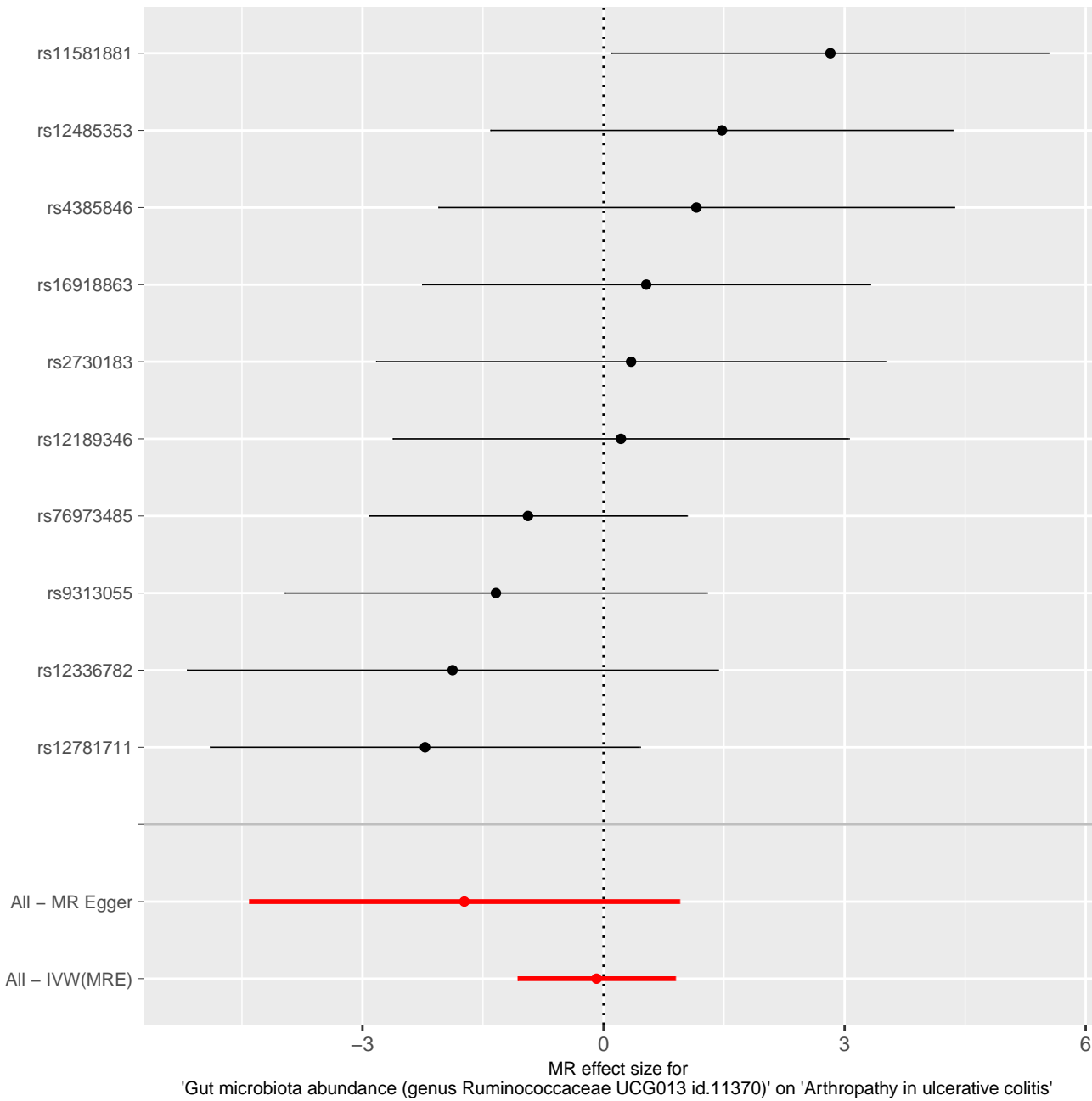




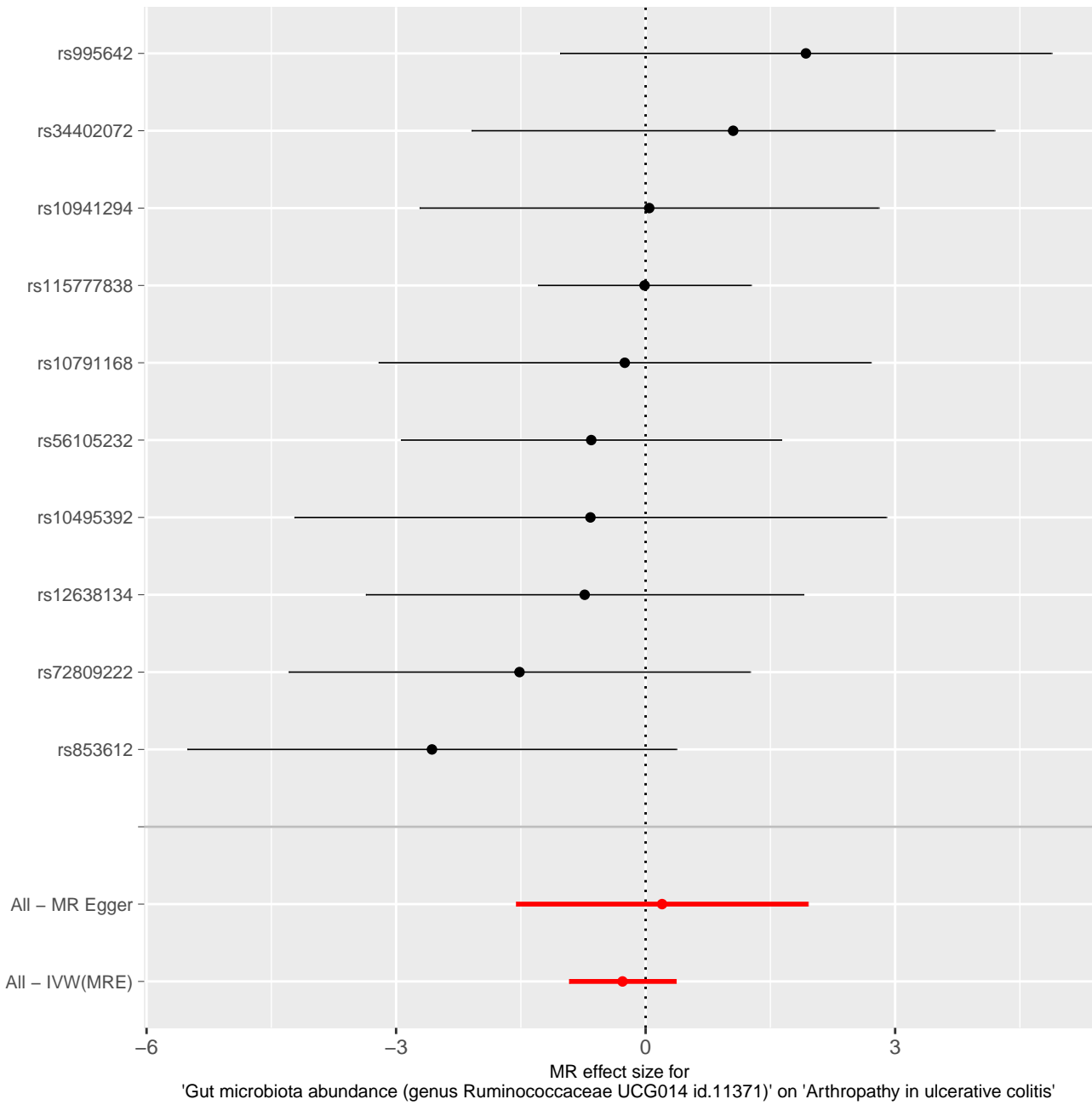


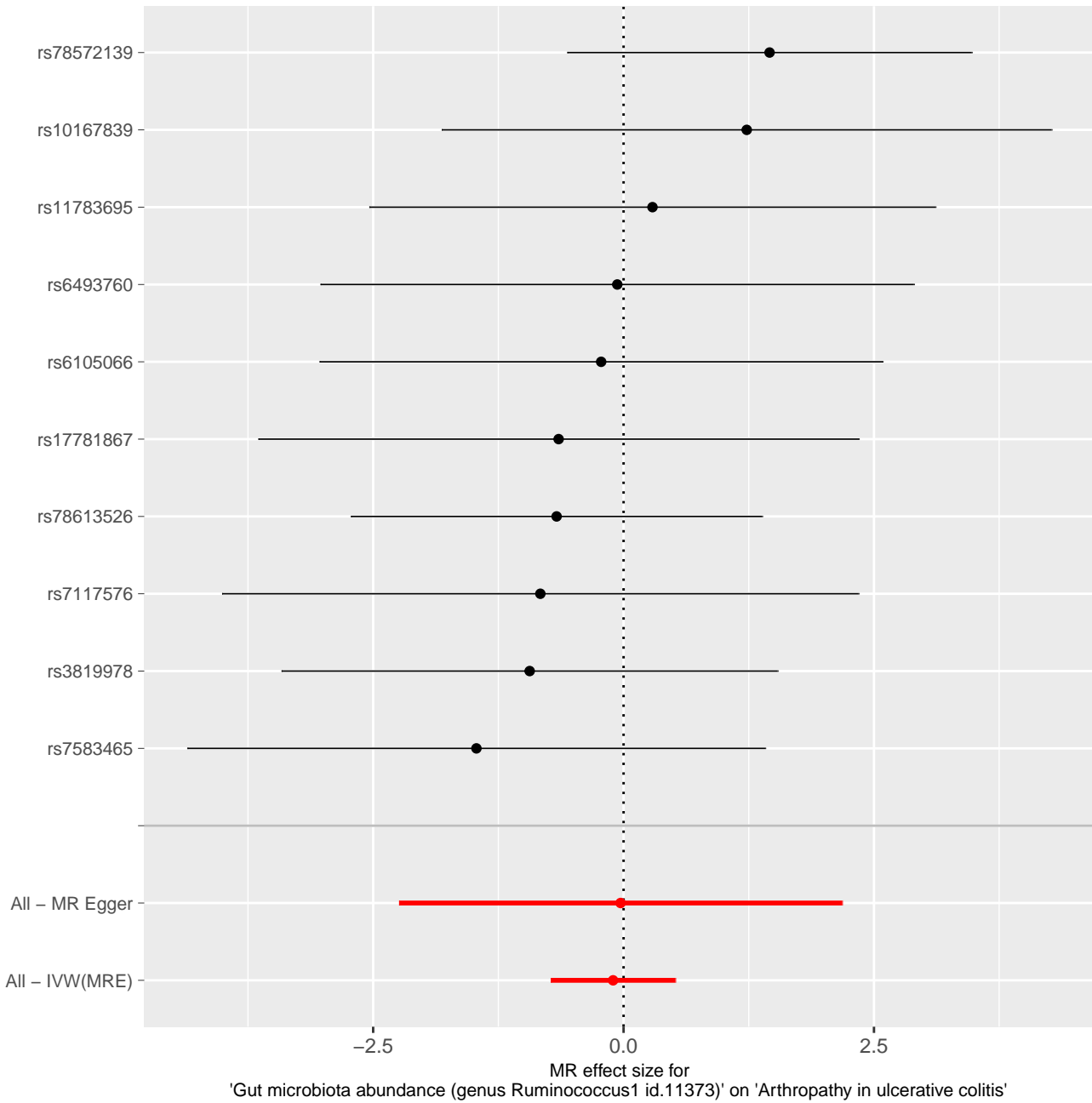




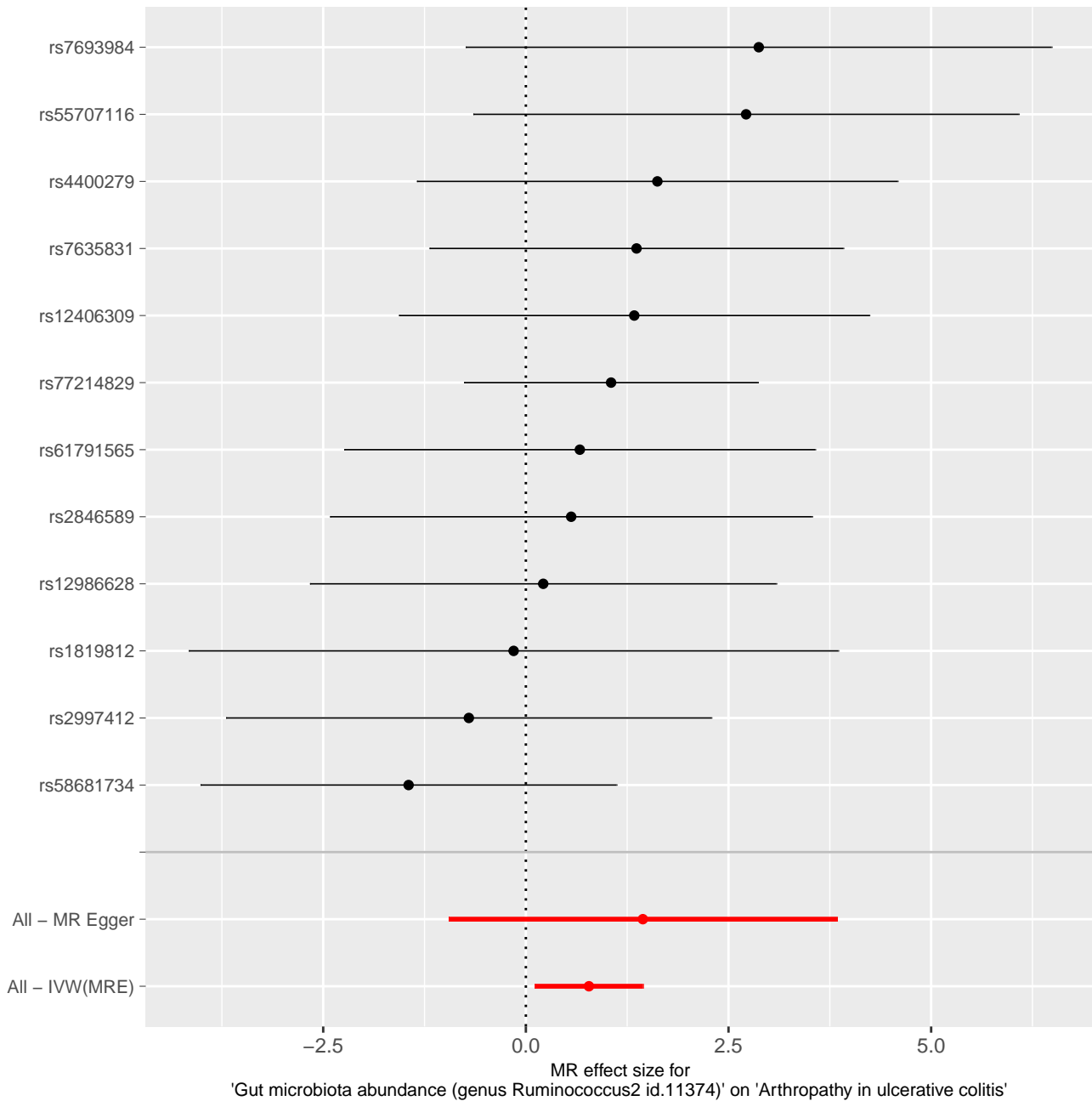


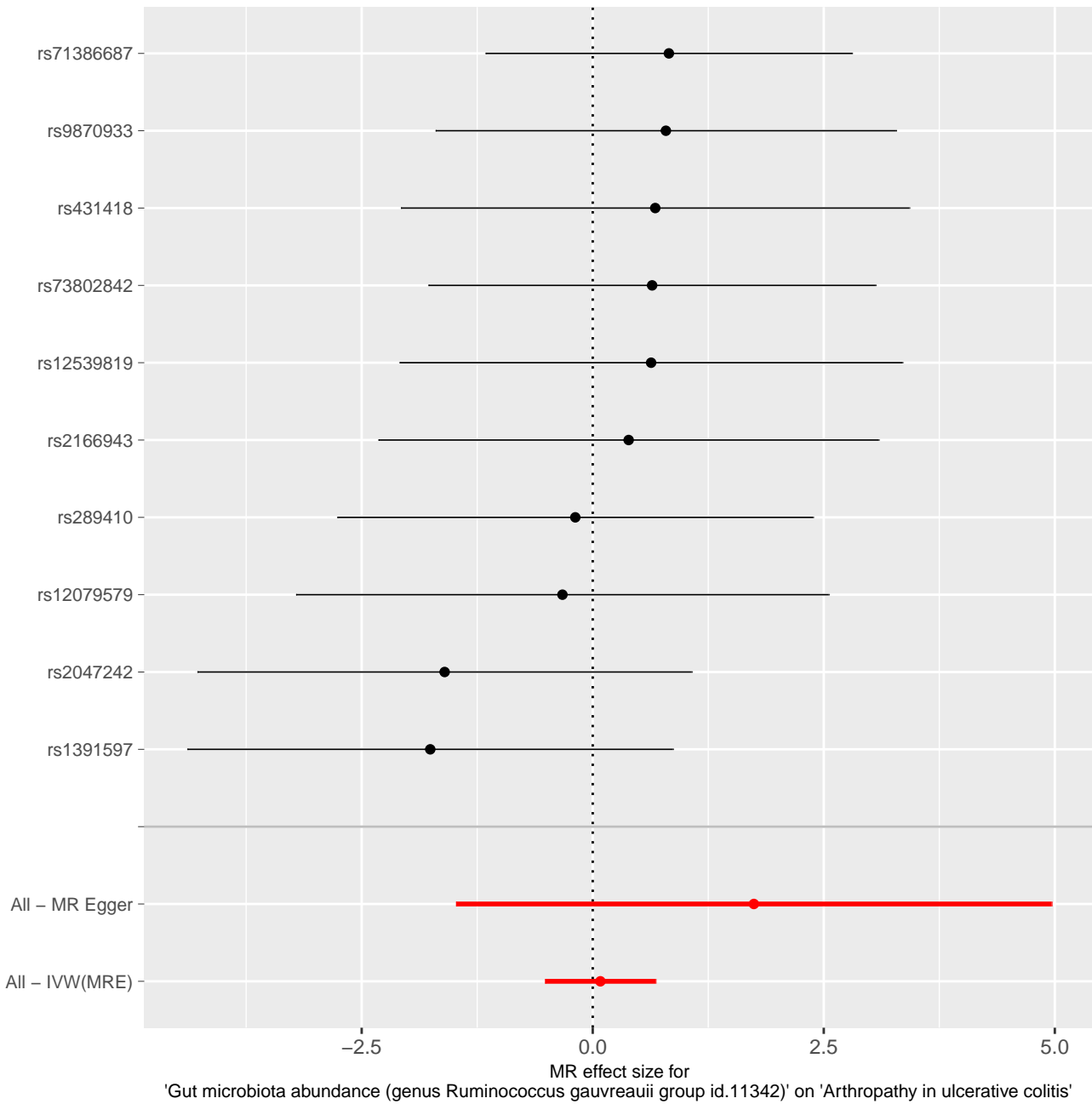
MR effect size for 'Gut microbiota abundance (genus Ruminococcaceae UCG013 id.11370)' on 'Arthropathy in ulcerative colitis'



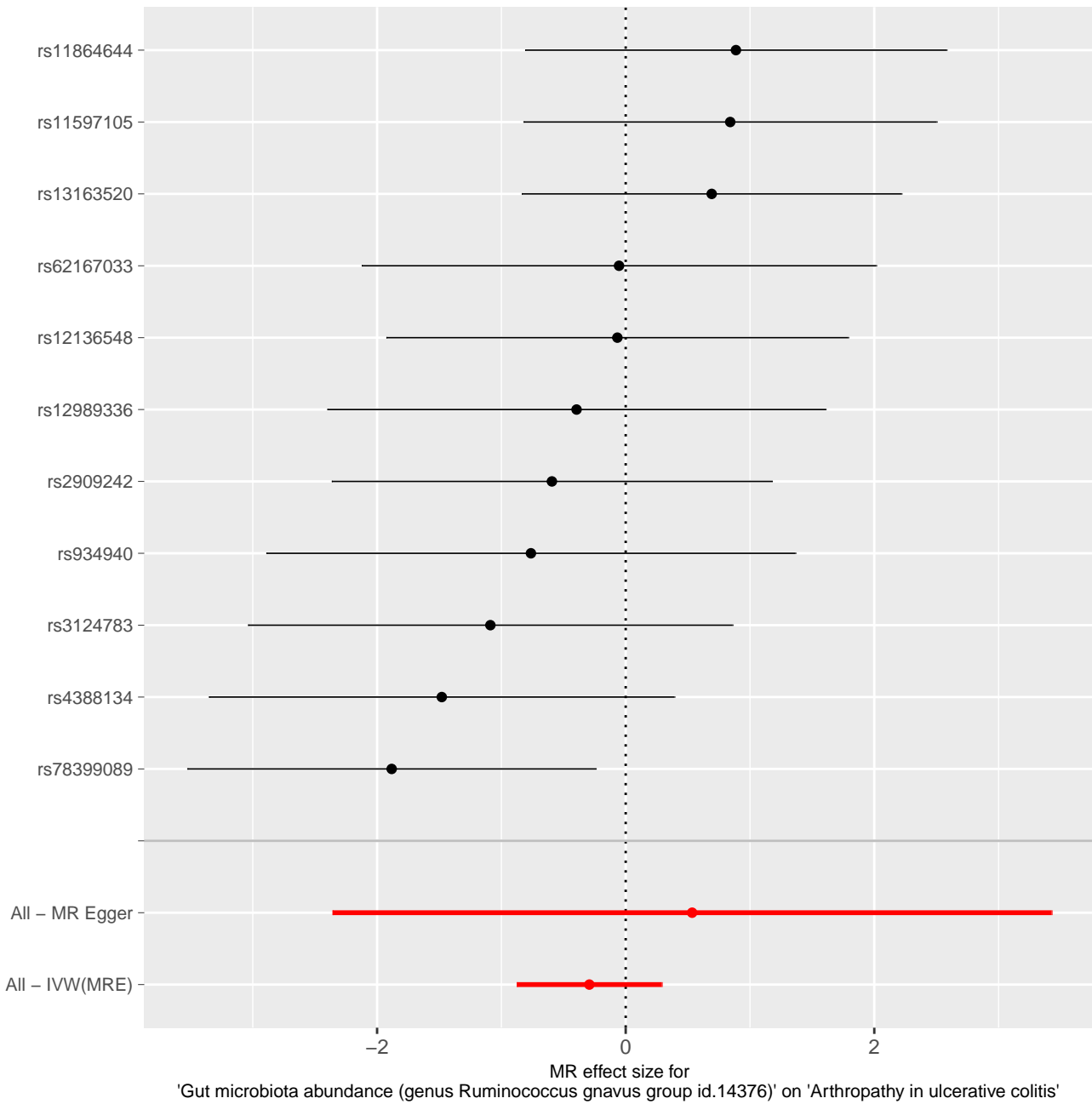


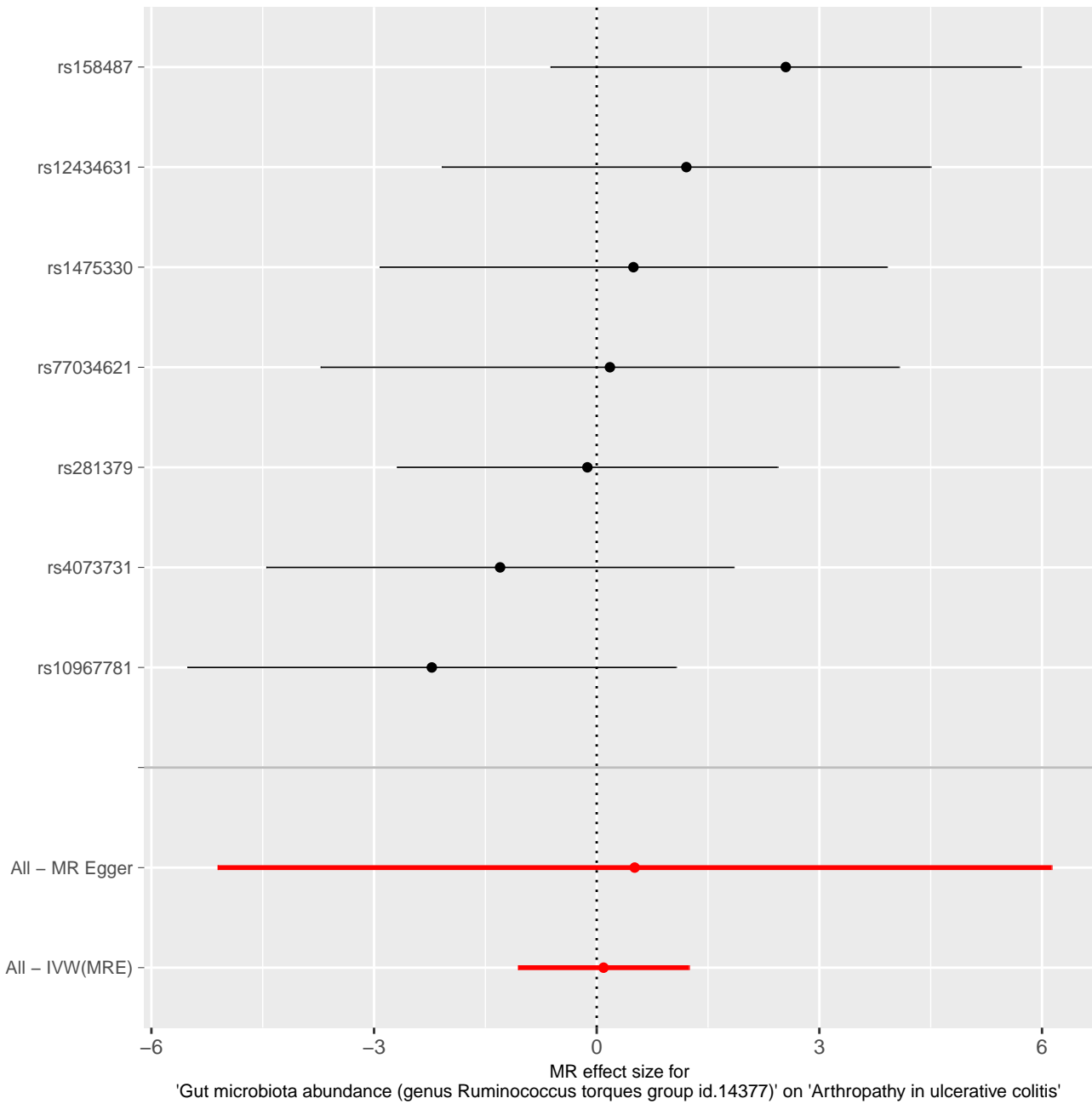
Batch 887 : Gut microbiota abundance (genus Ruminococcus2 id.11374) on Arthropathy in ulcerative colitis

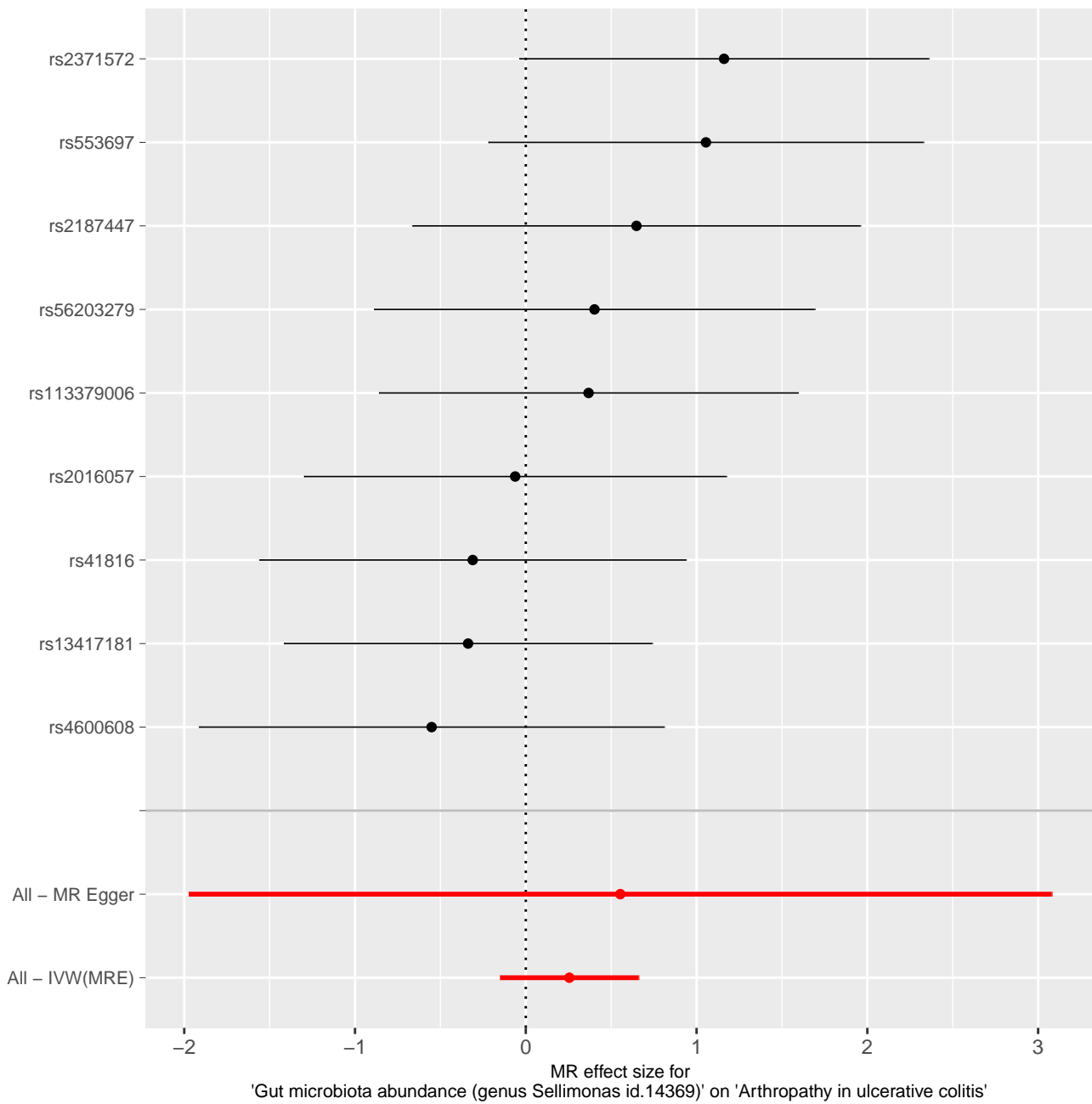


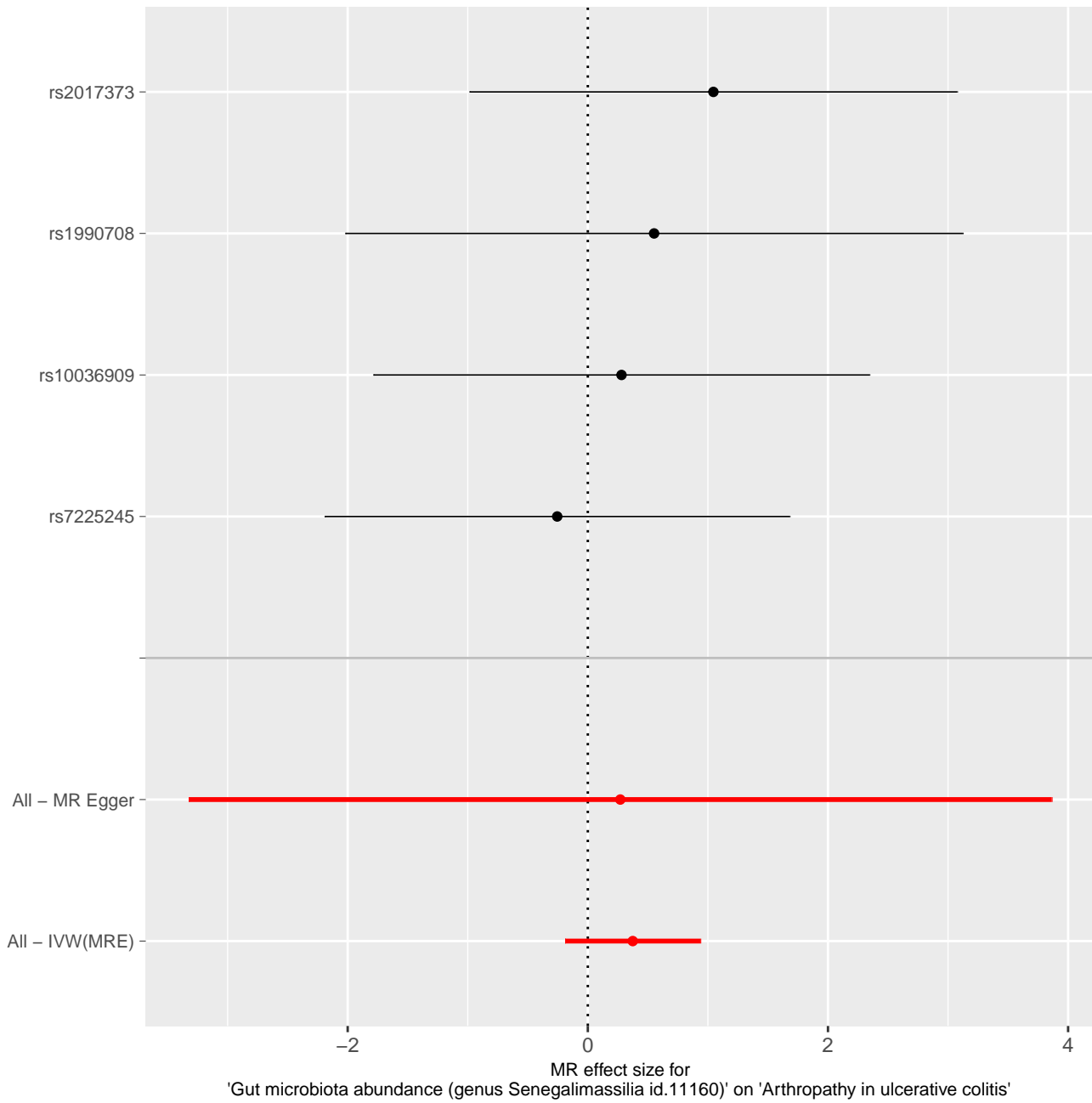


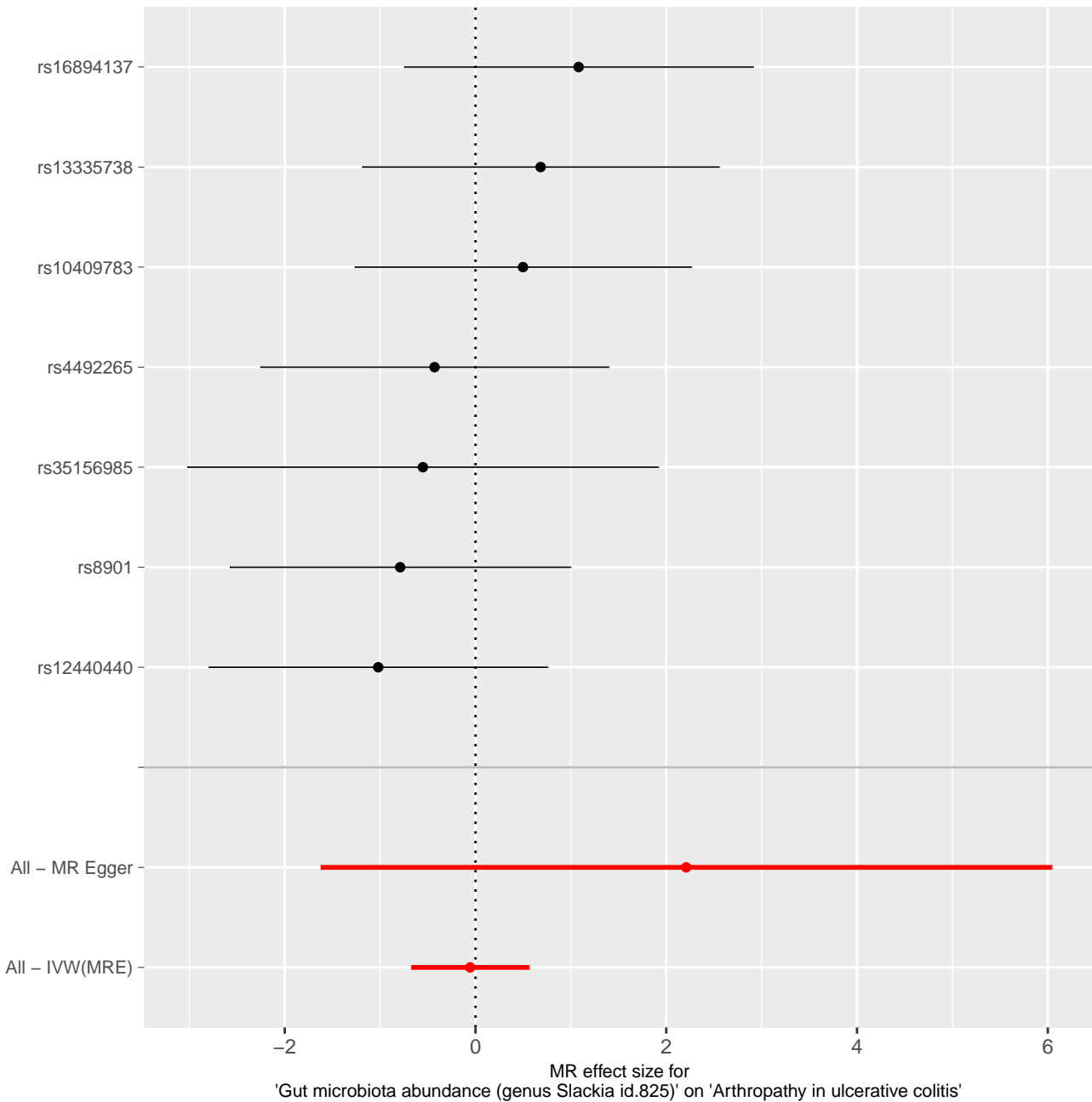
'Gut microbiota abundance (genus Ruminococcus gauvreauii group id.11342)' on 'Arthropathy in ulcerative colitis'



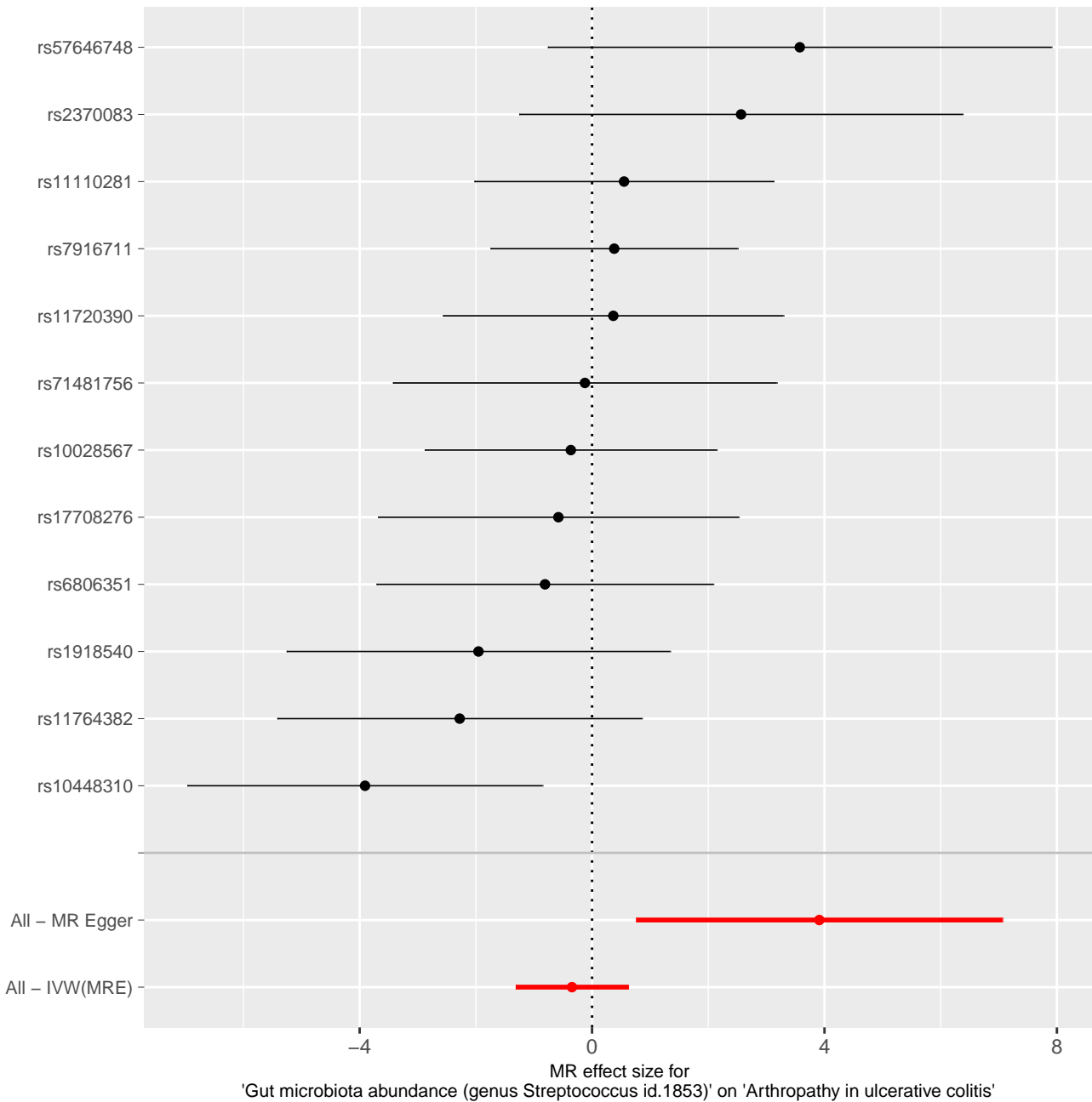


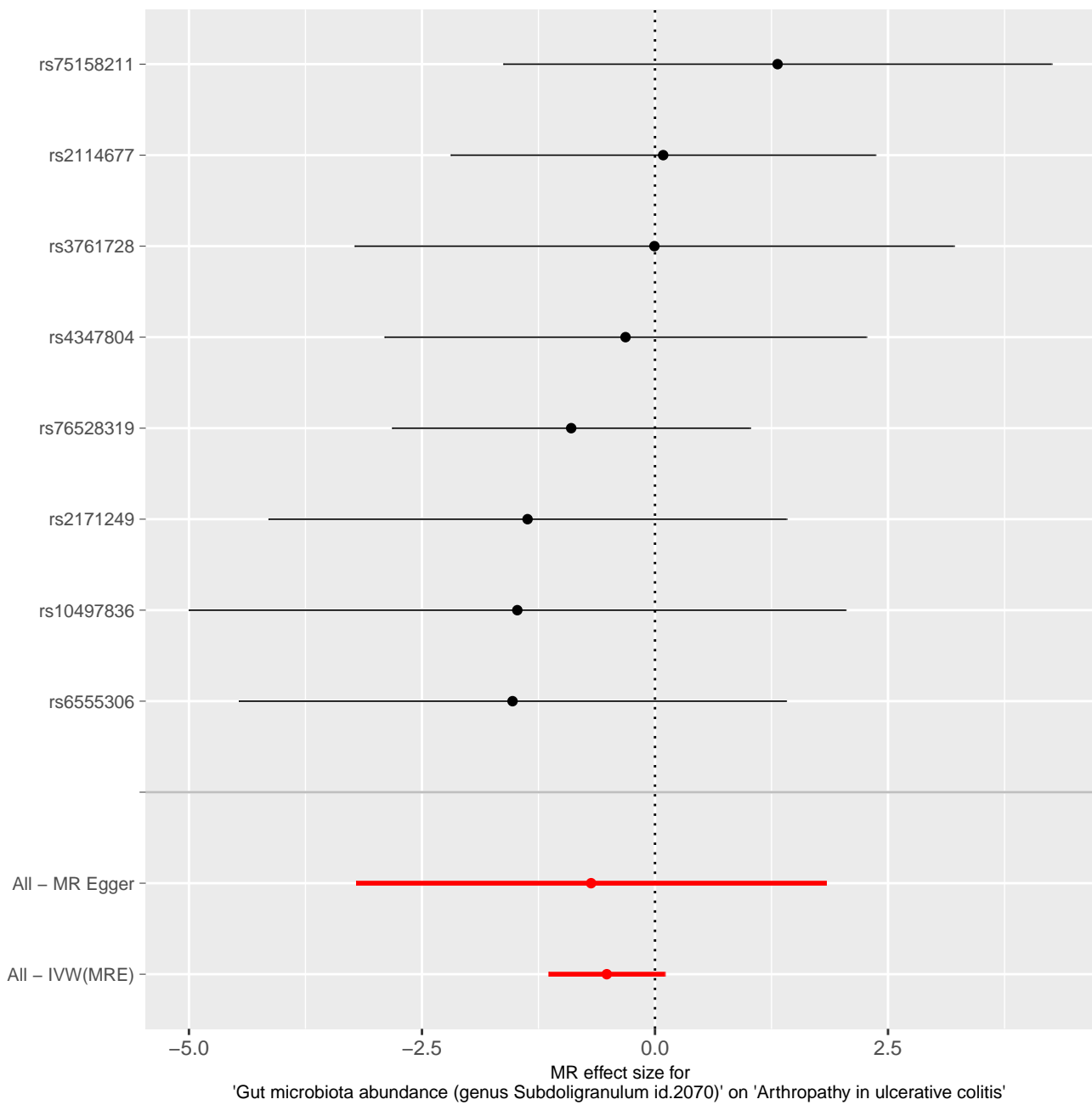




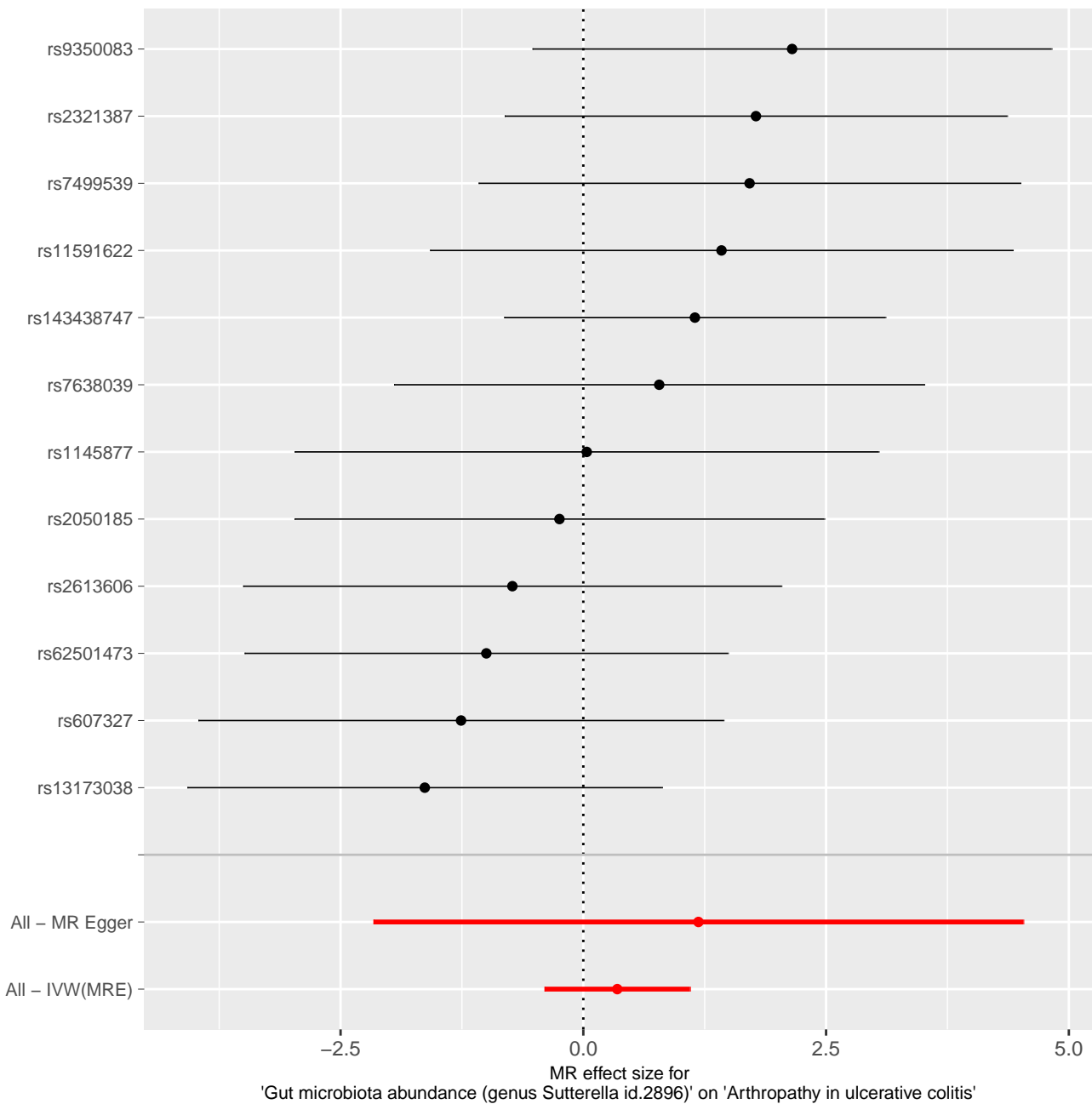


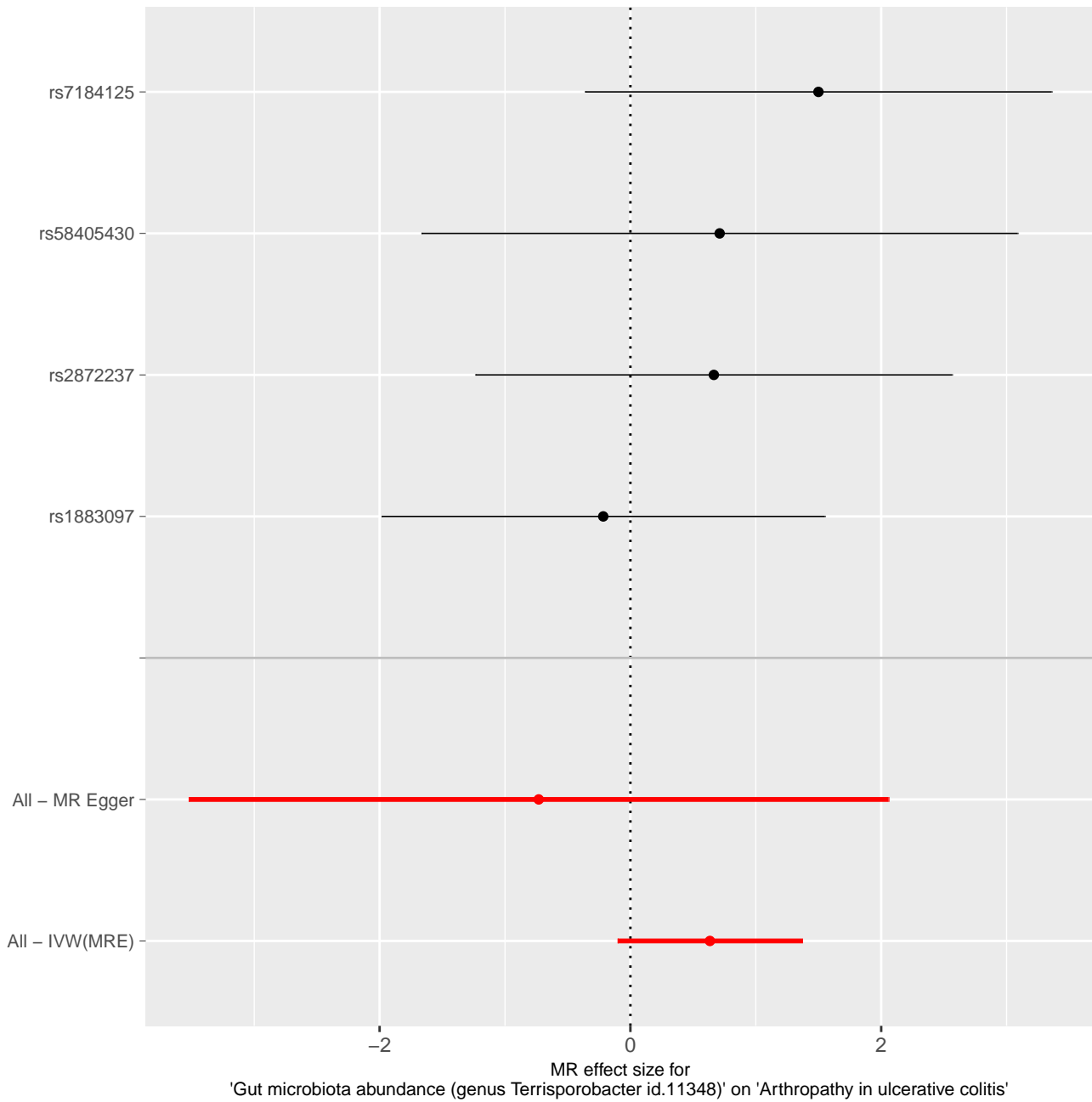
Batch 894 : Gut microbiota abundance (genus Streptococcus id.1853) on Arthropathy in ulcerative colitis

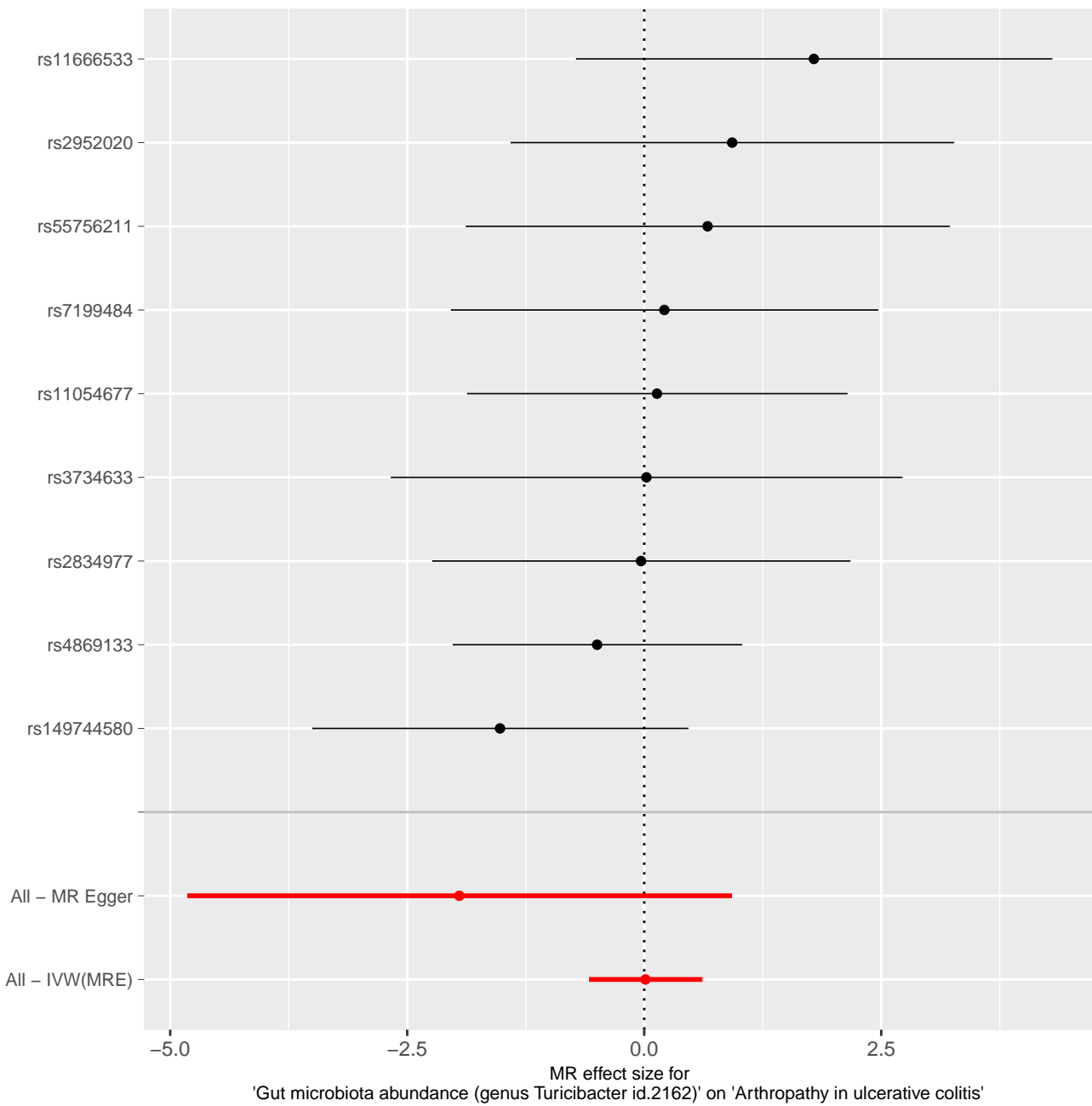




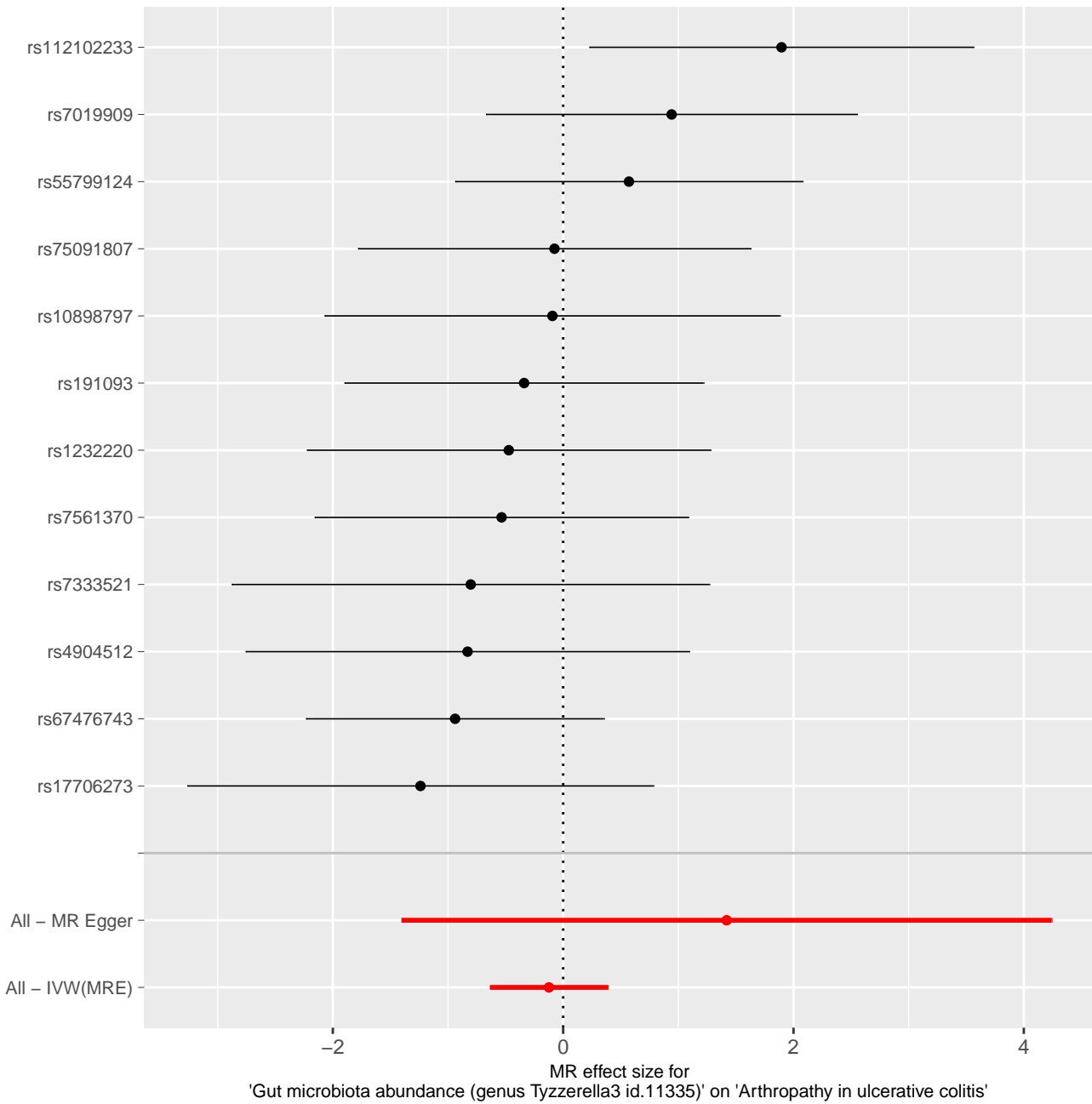
Batch 896 : Gut microbiota abundance (genus Sutterella id.2896) on Arthropathy in ulcerative colitis

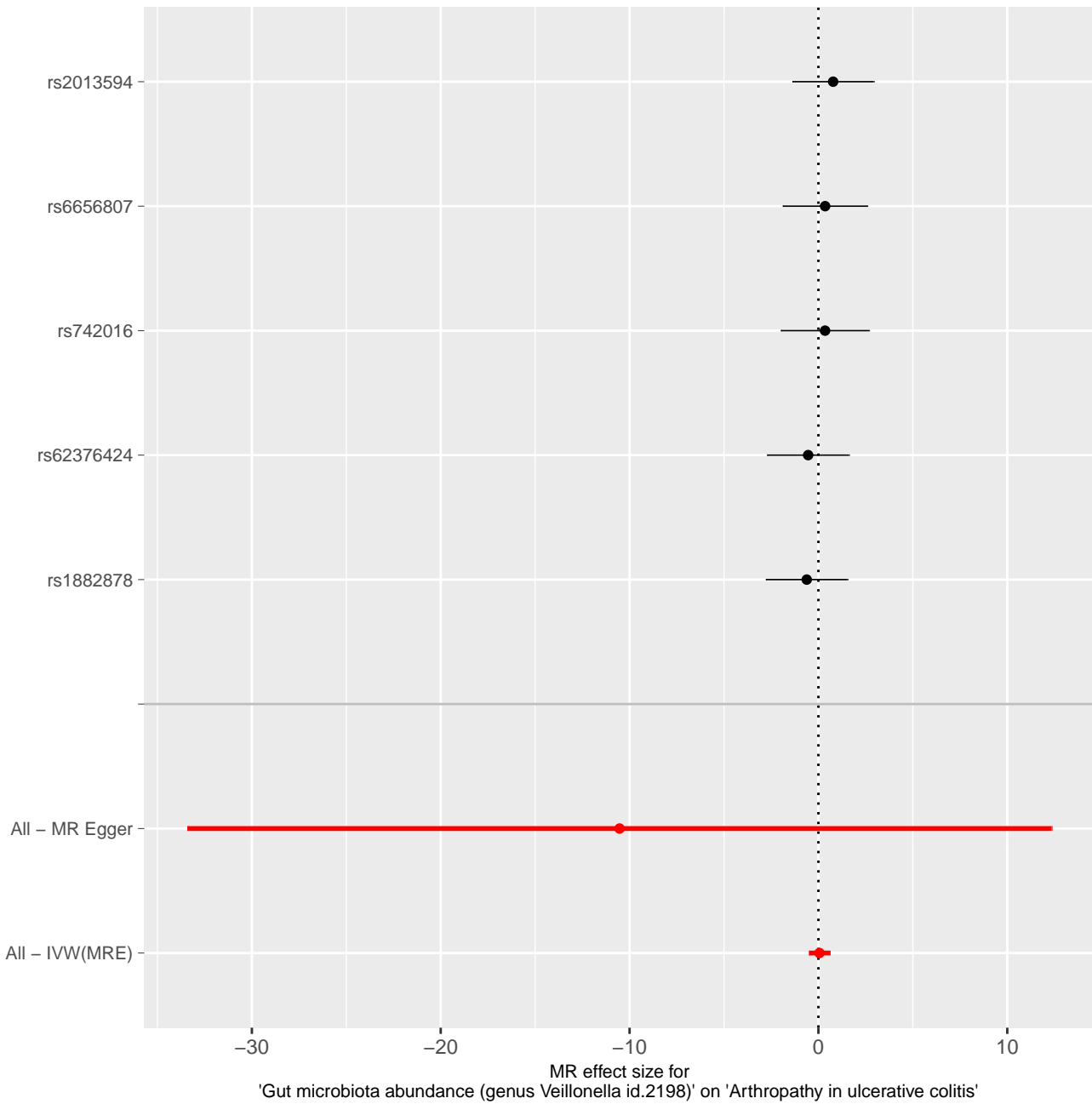


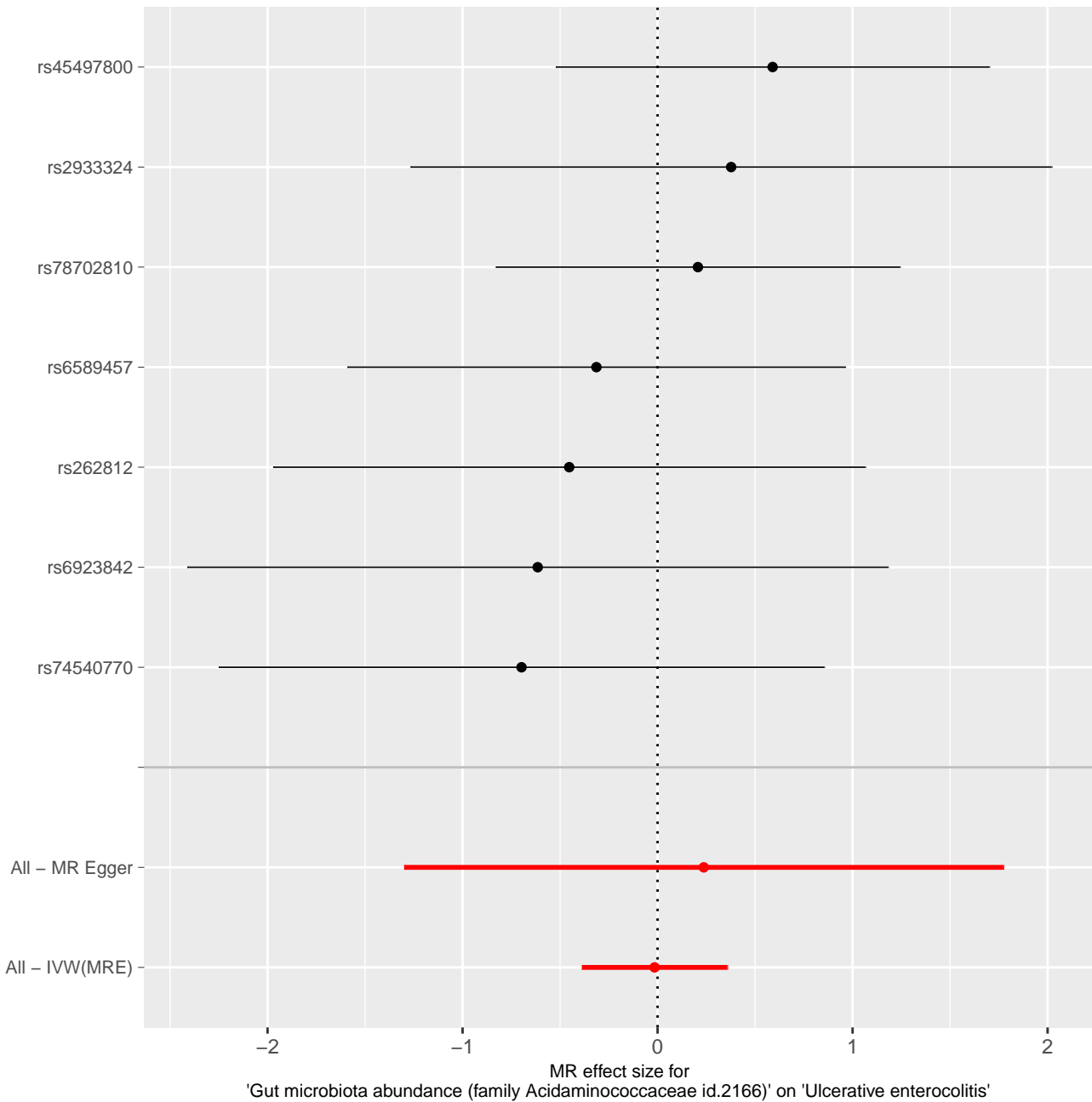


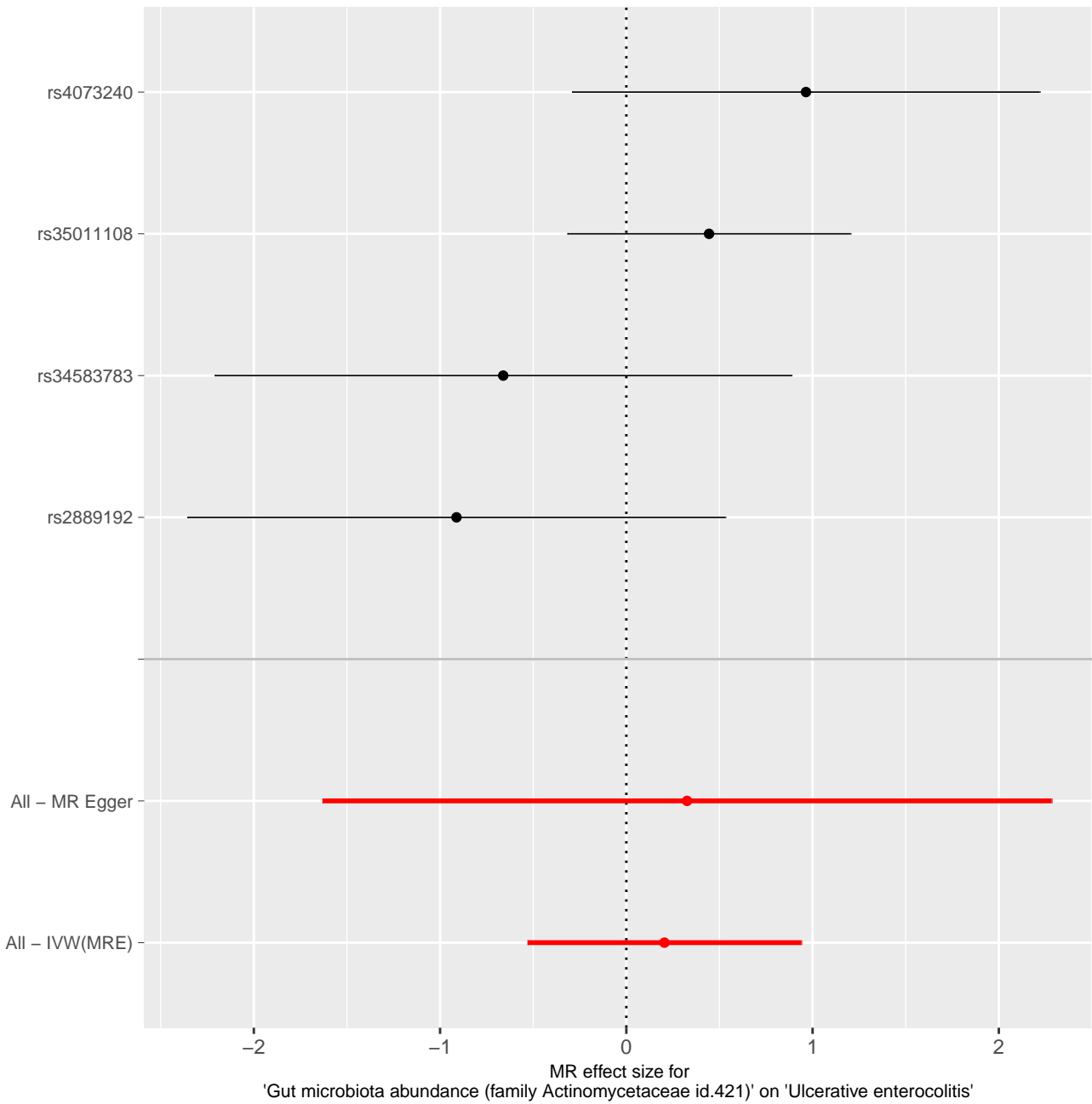


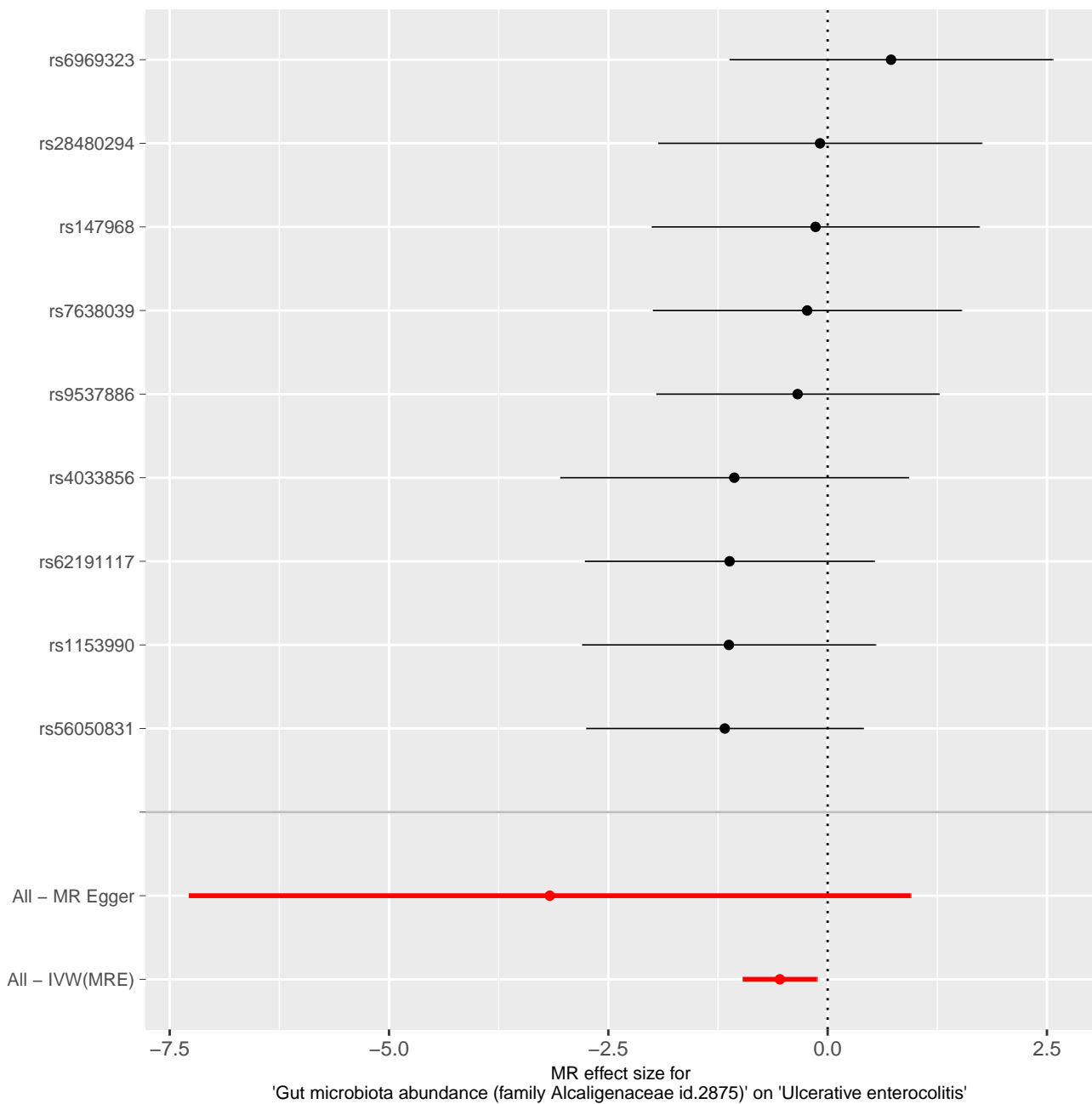
Batch 899 : Gut microbiota abundance (genus Tyzzerella3 id.11335) on Arthropathy in ulcerative colitis

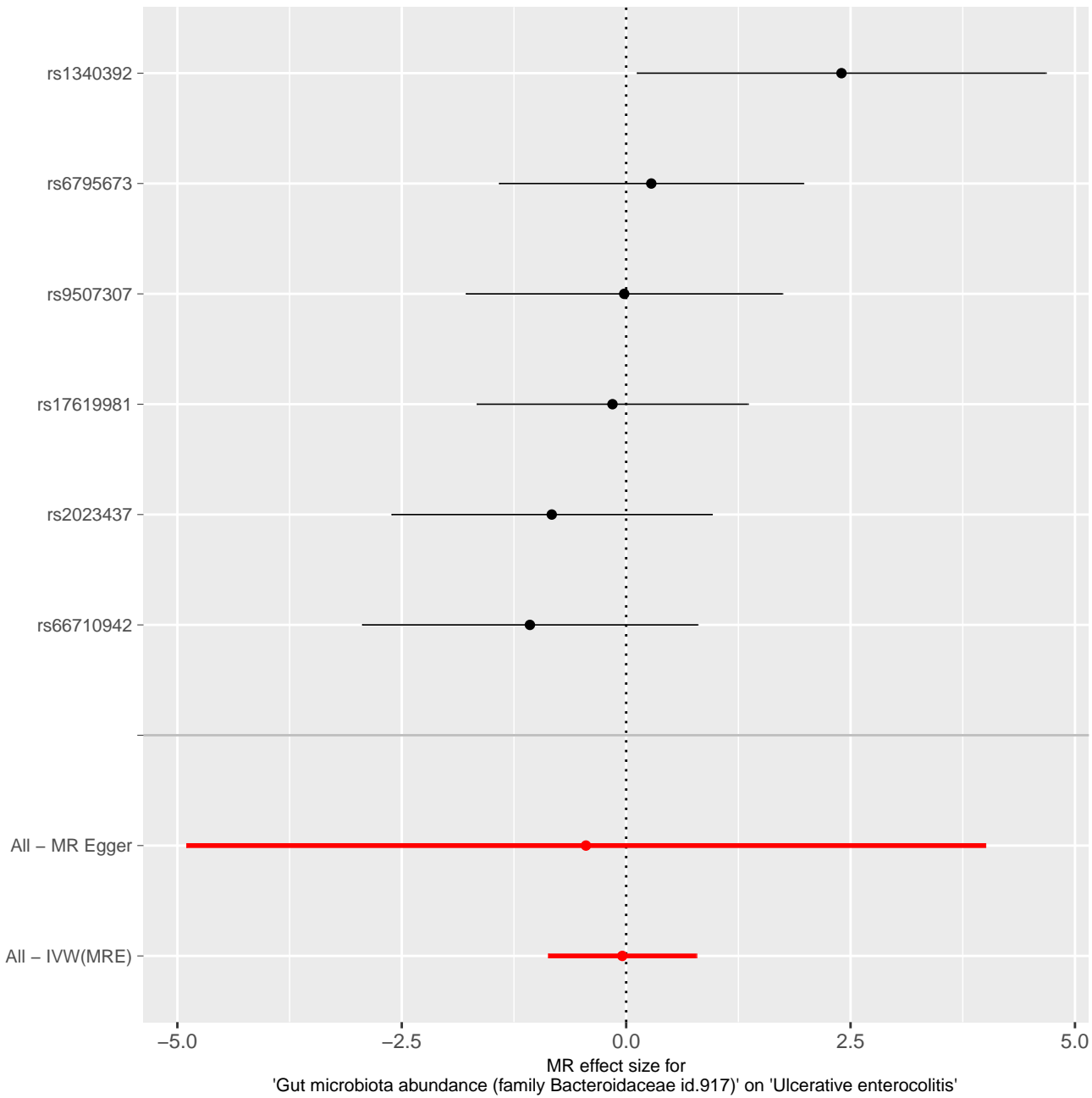


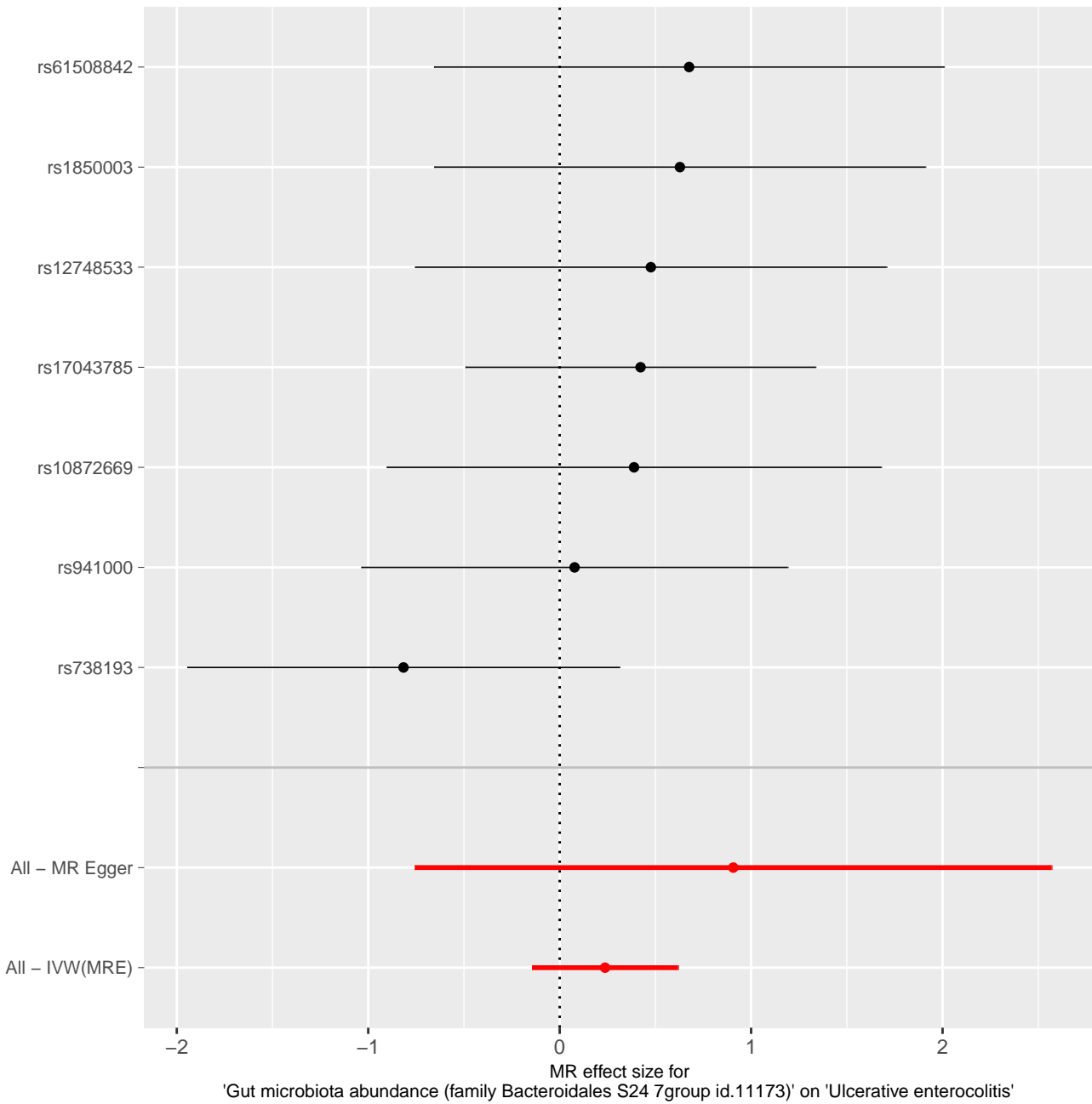


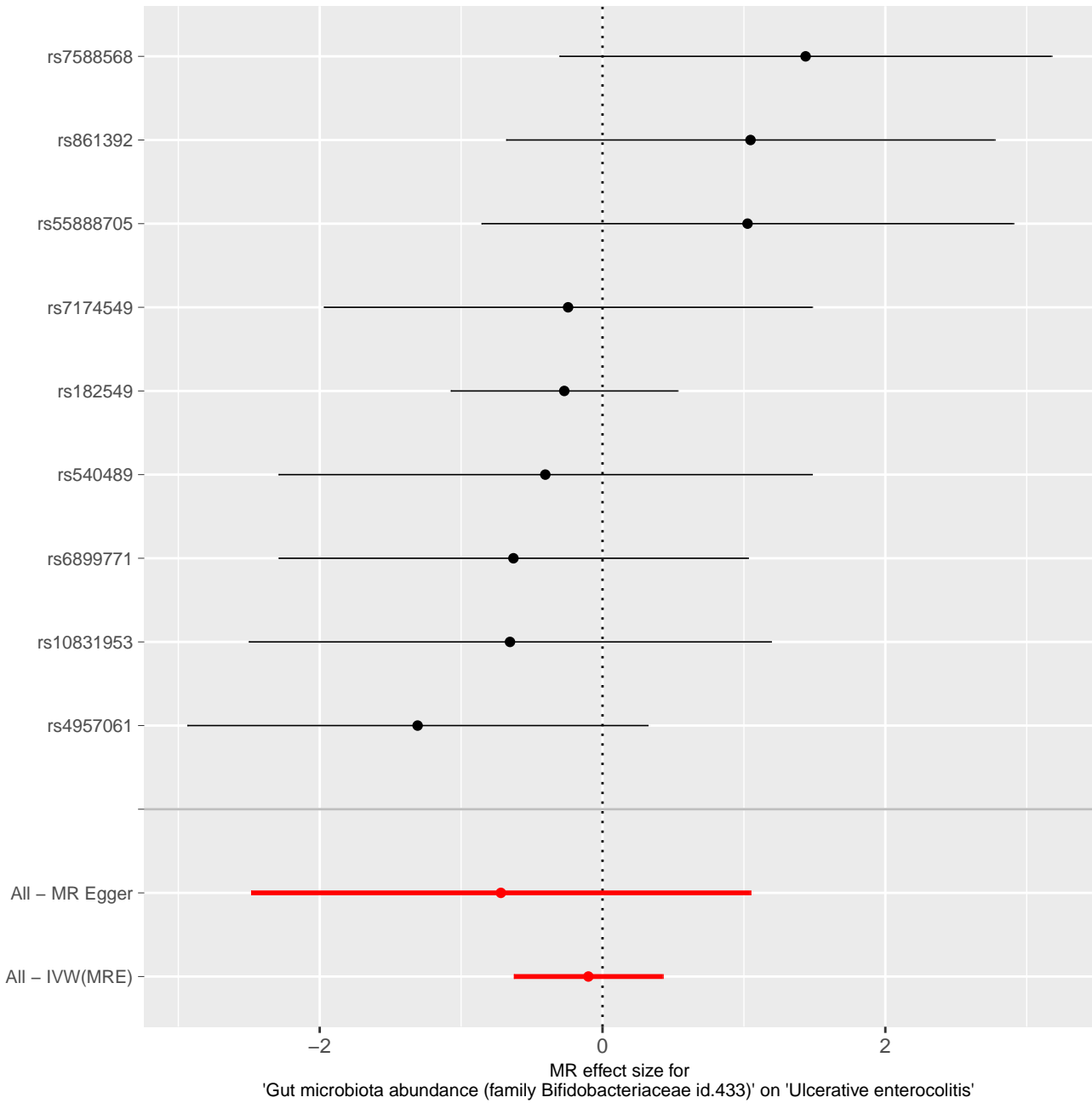


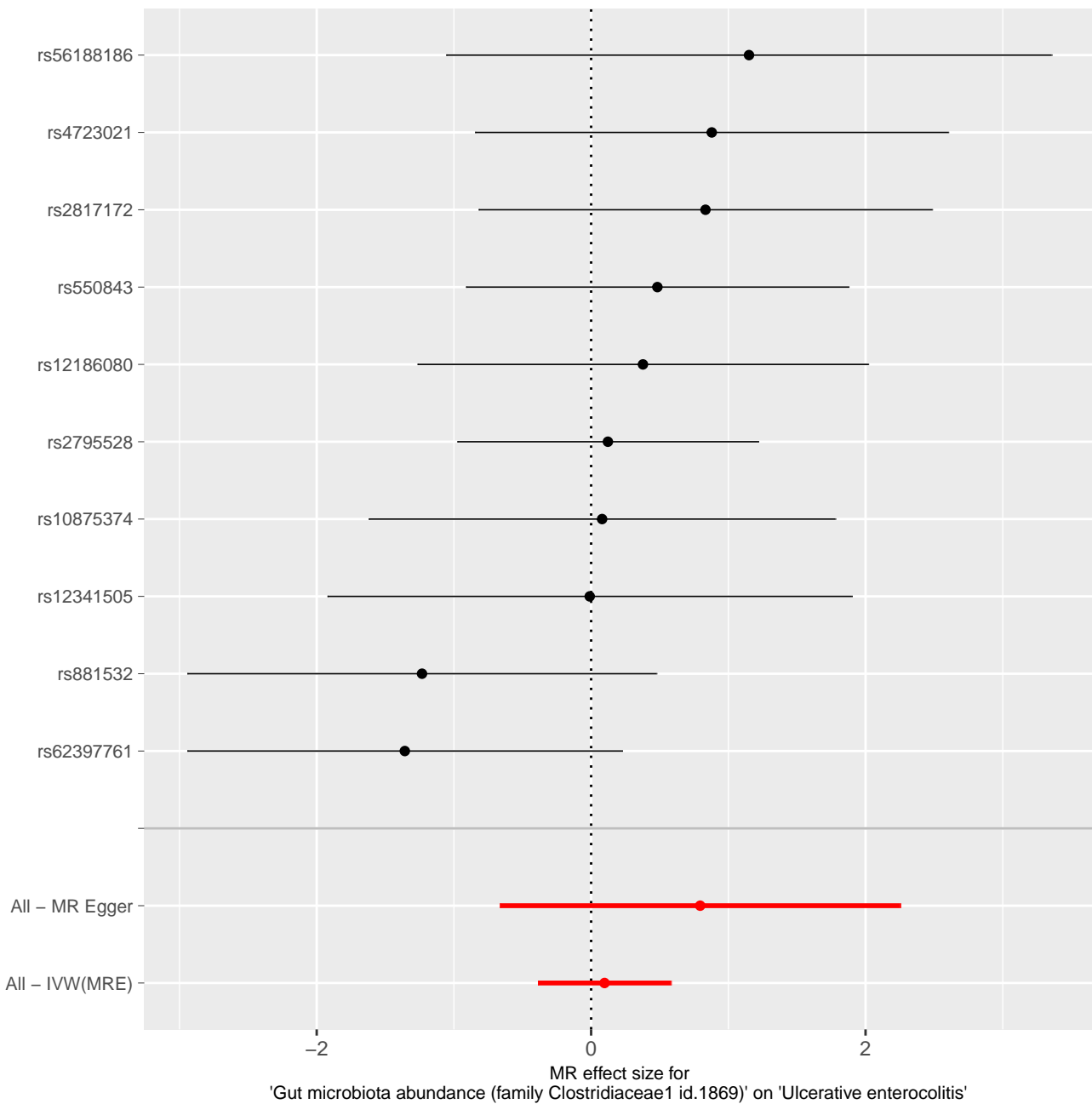


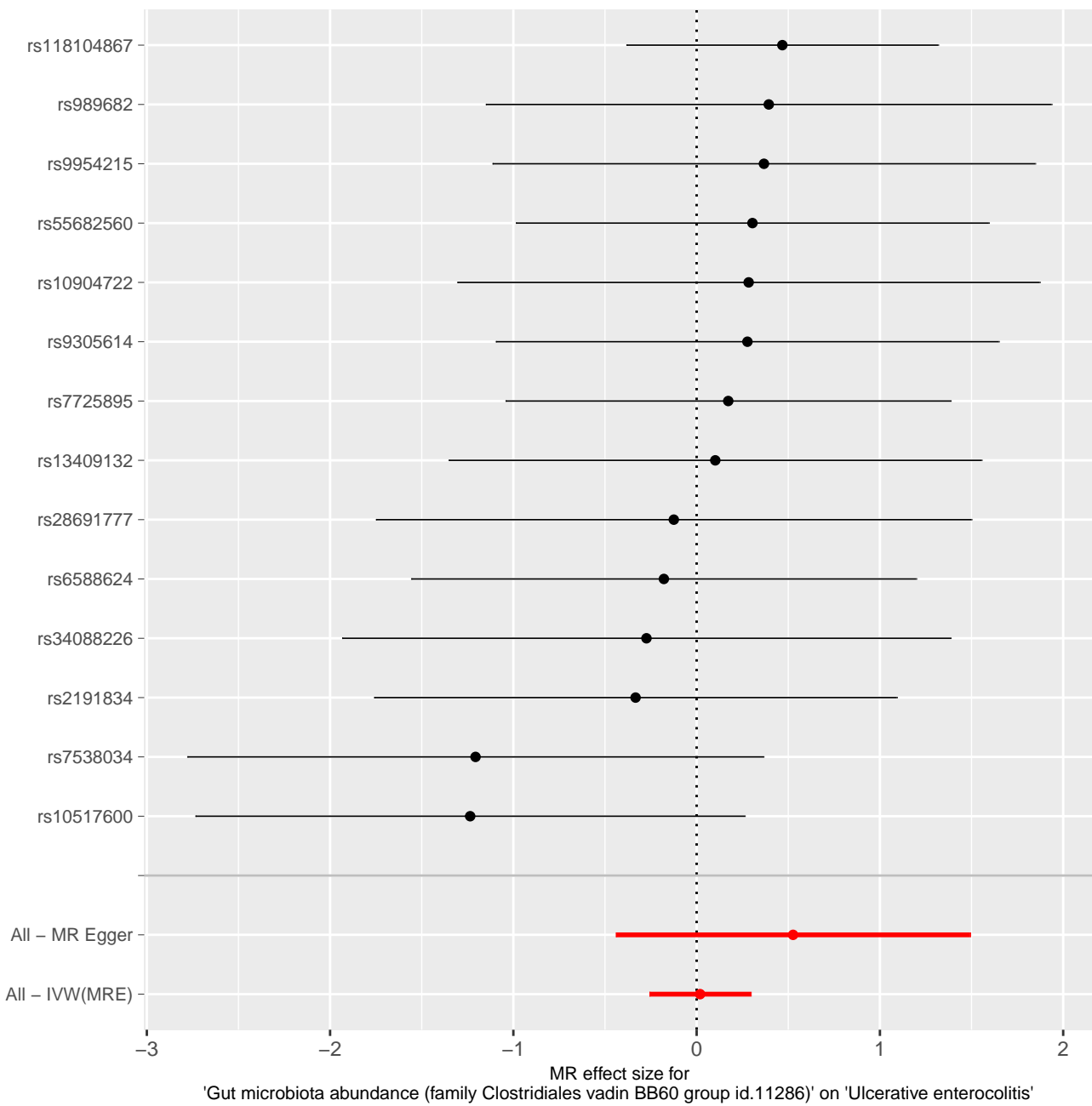


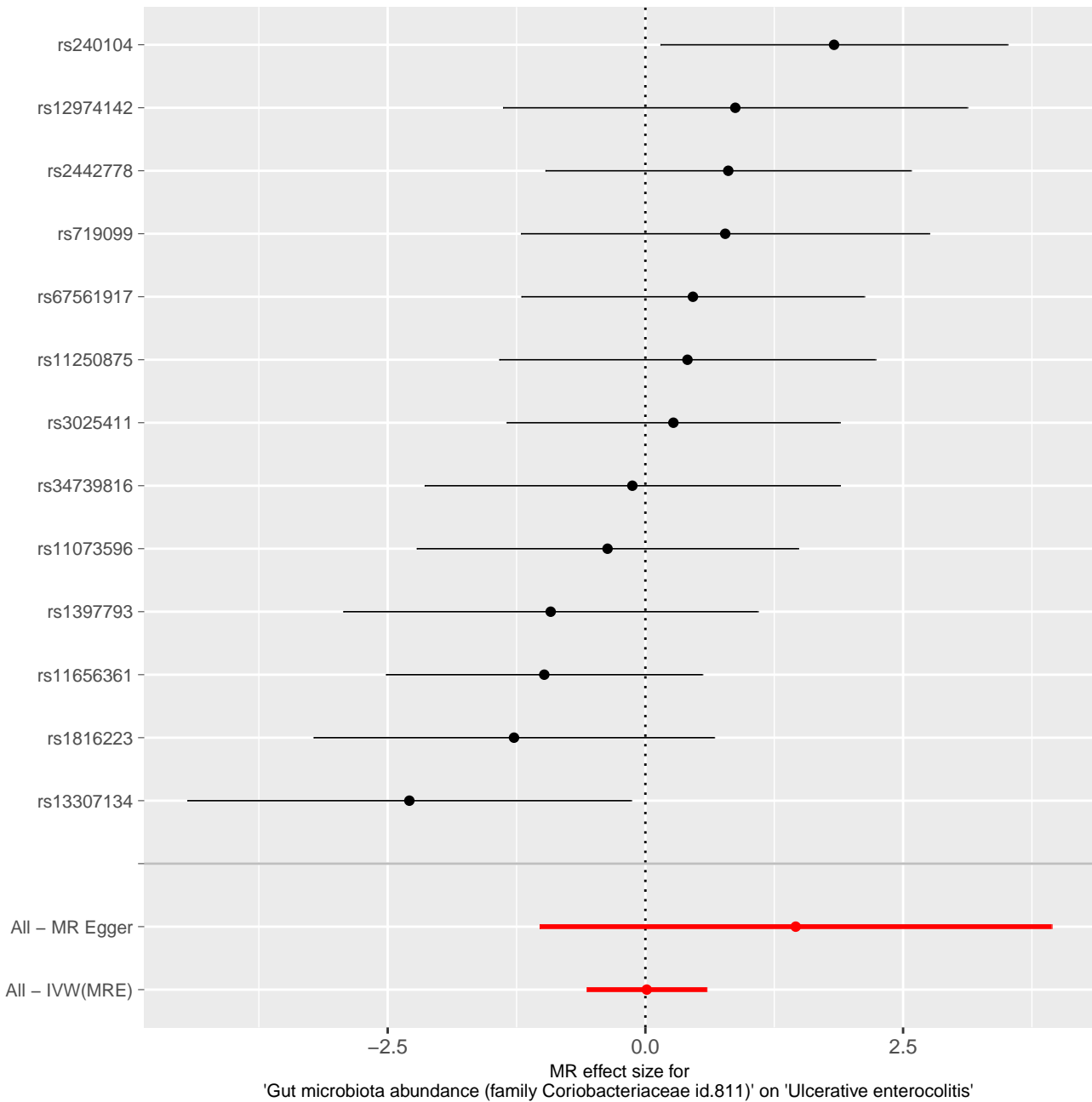


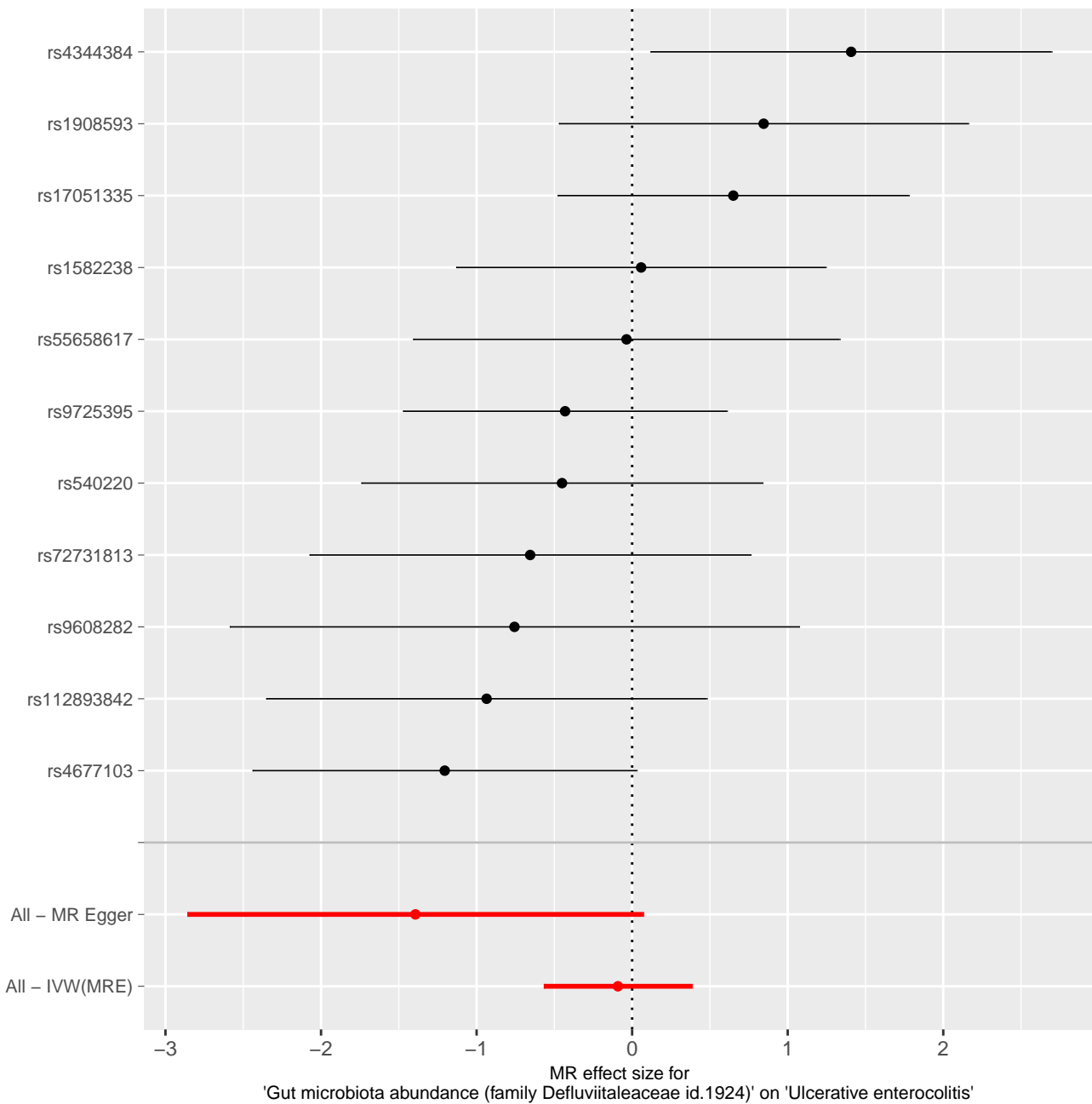


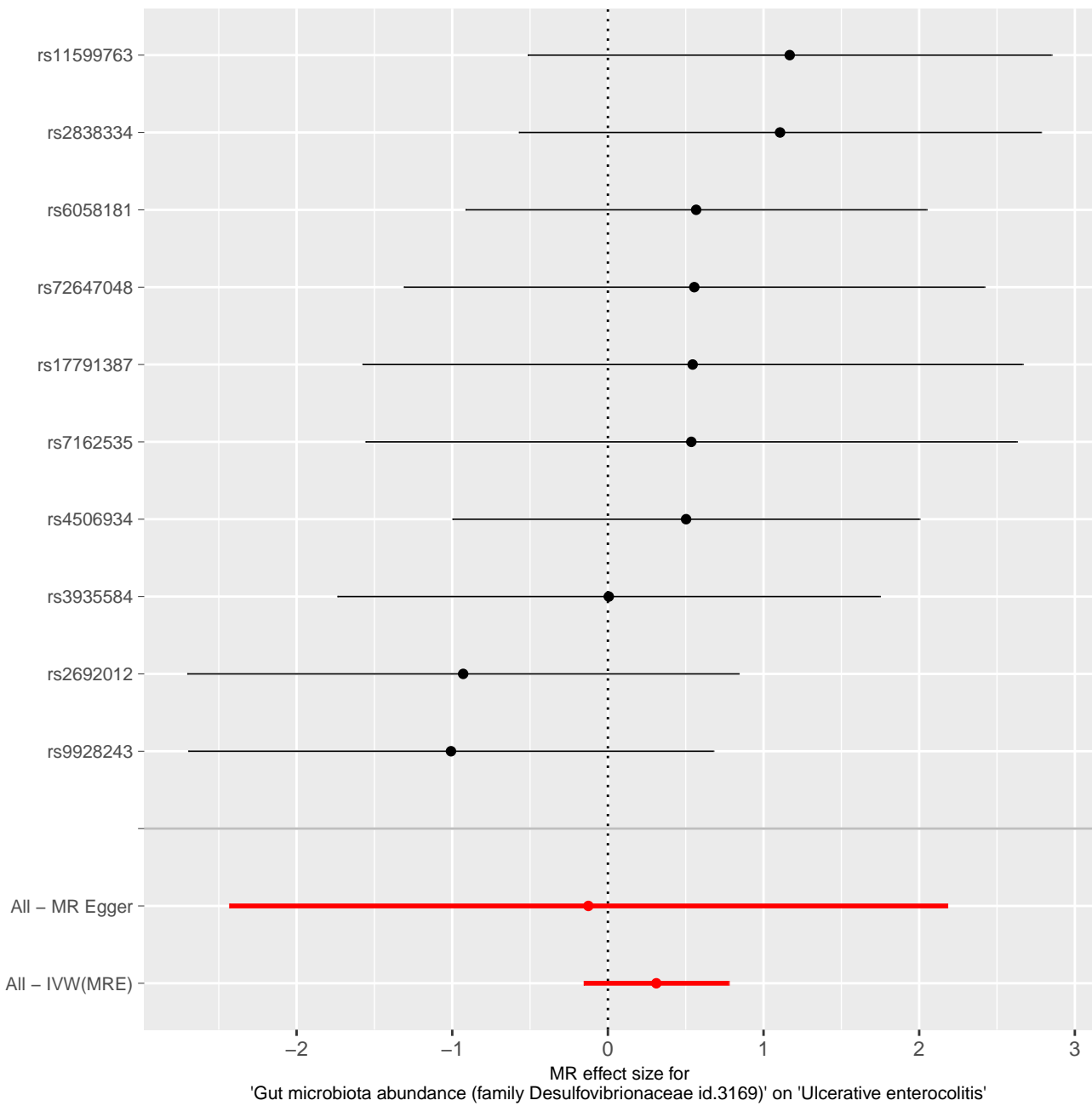


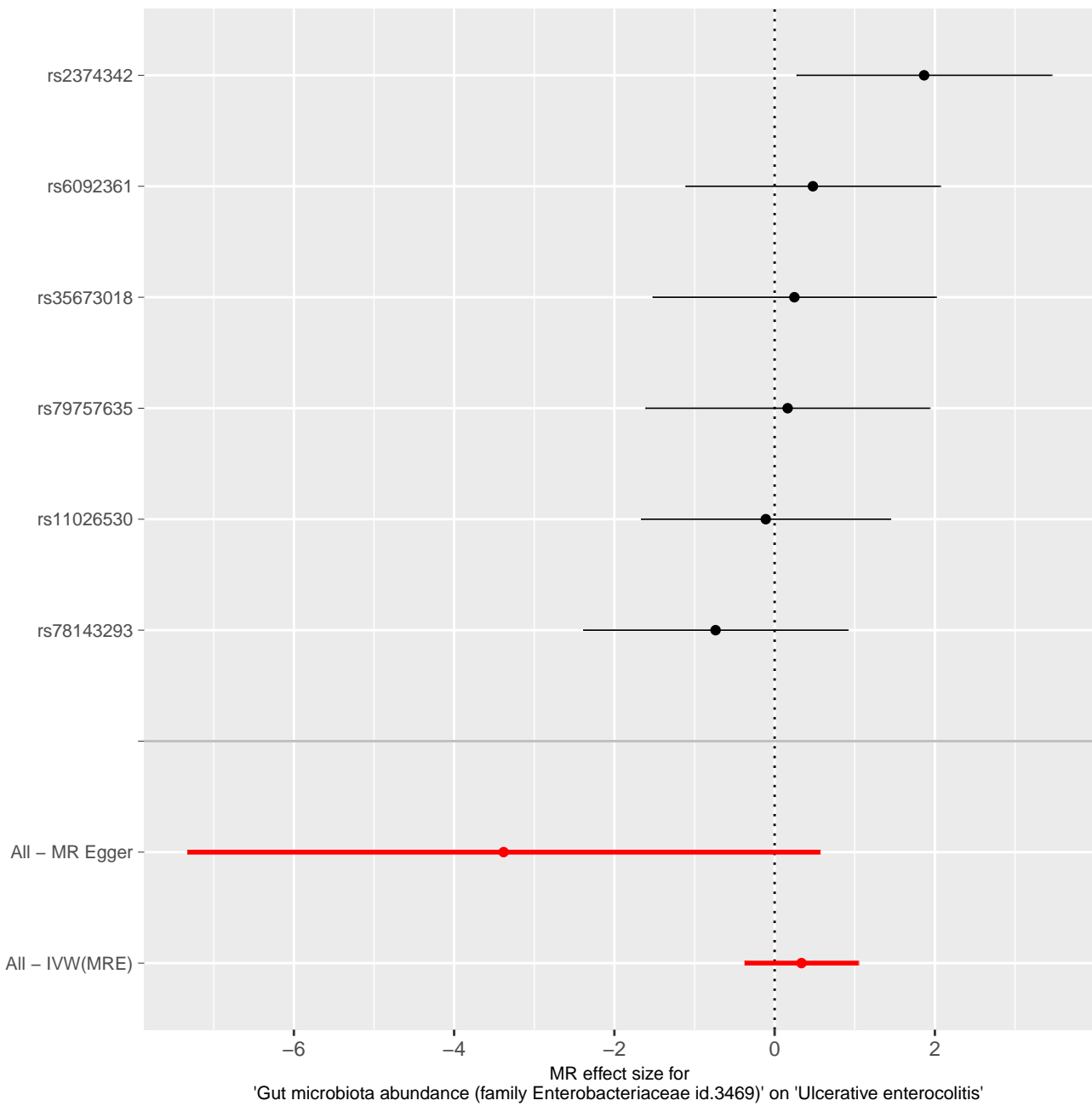




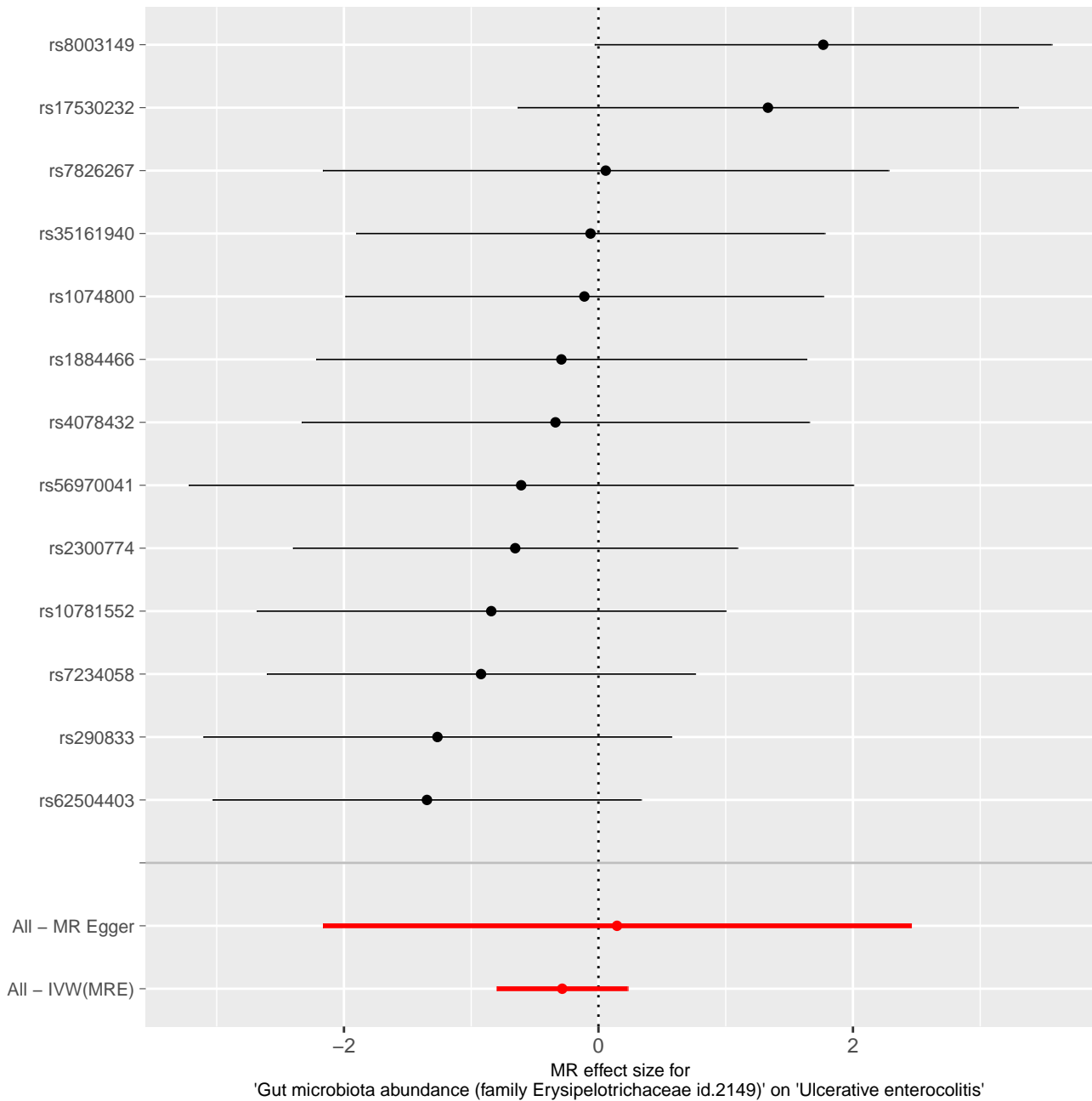


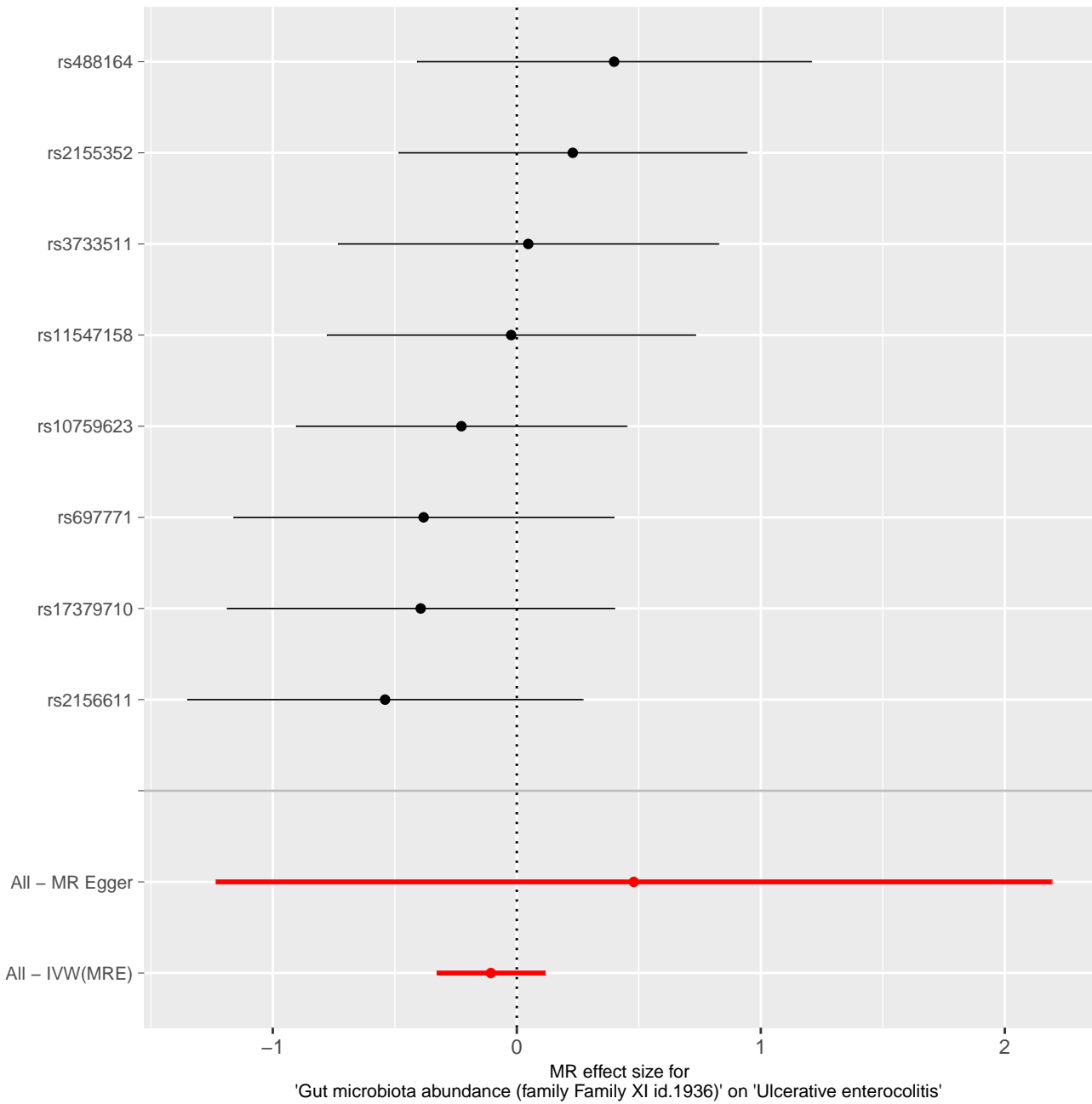


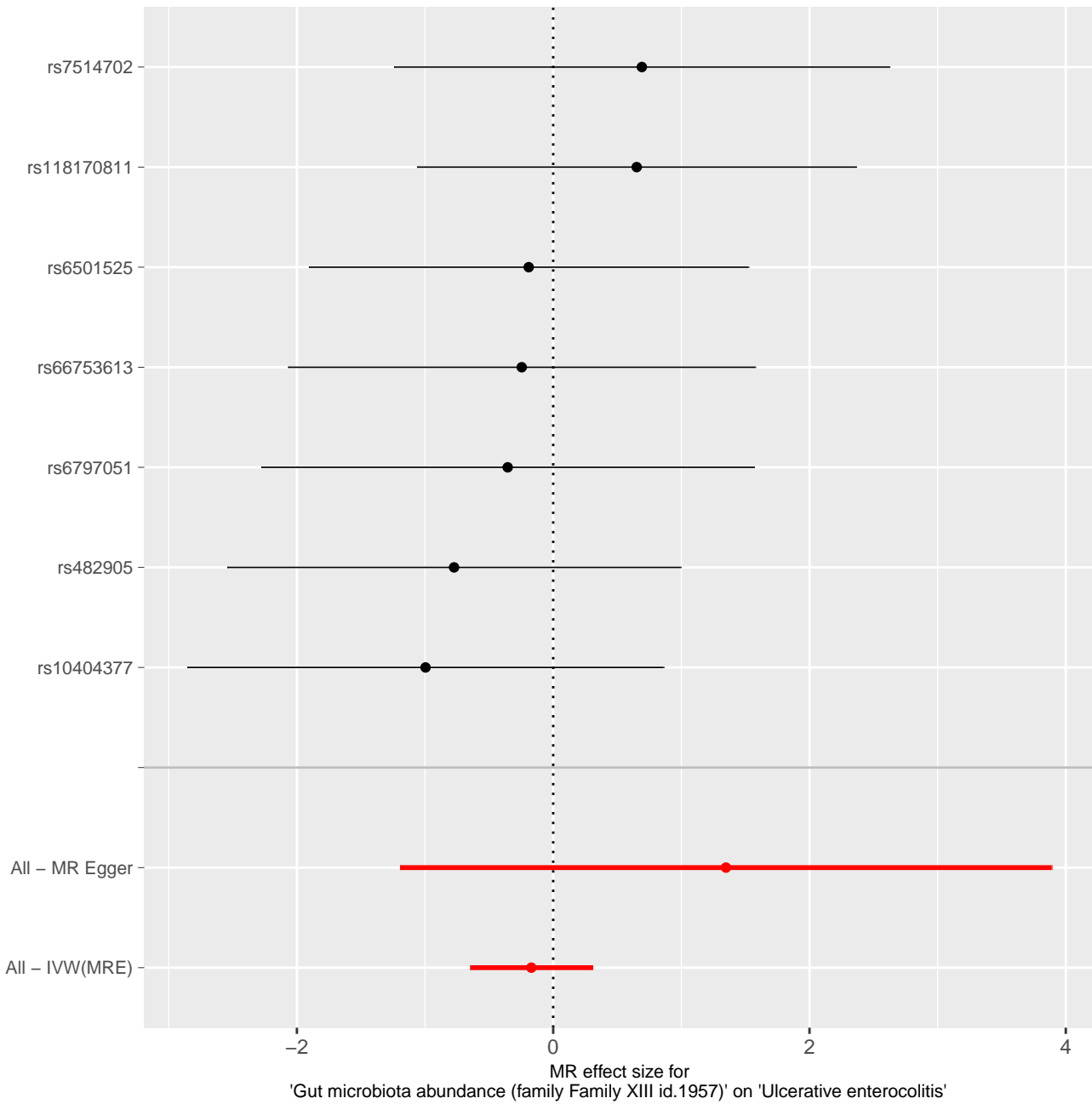


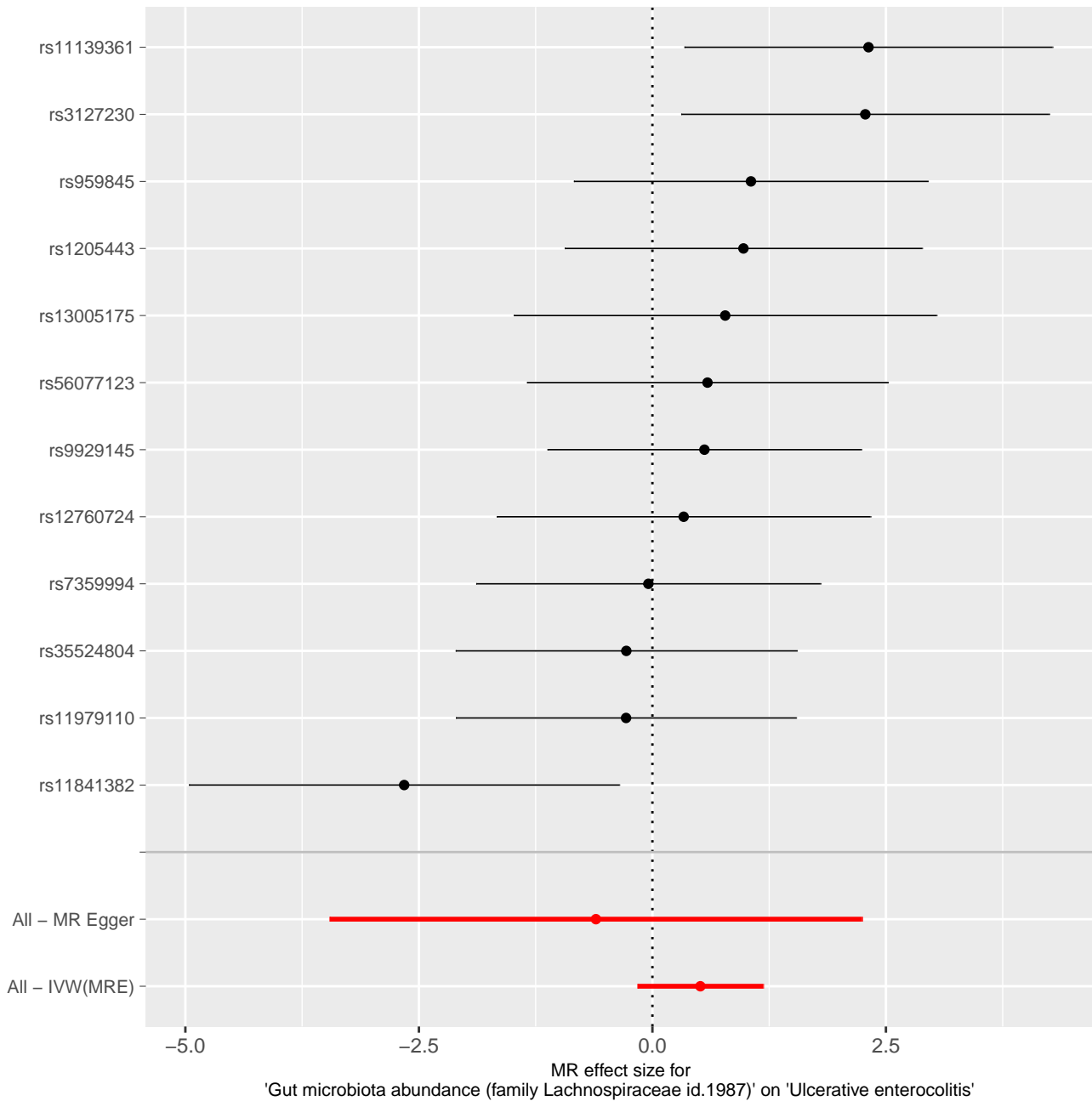


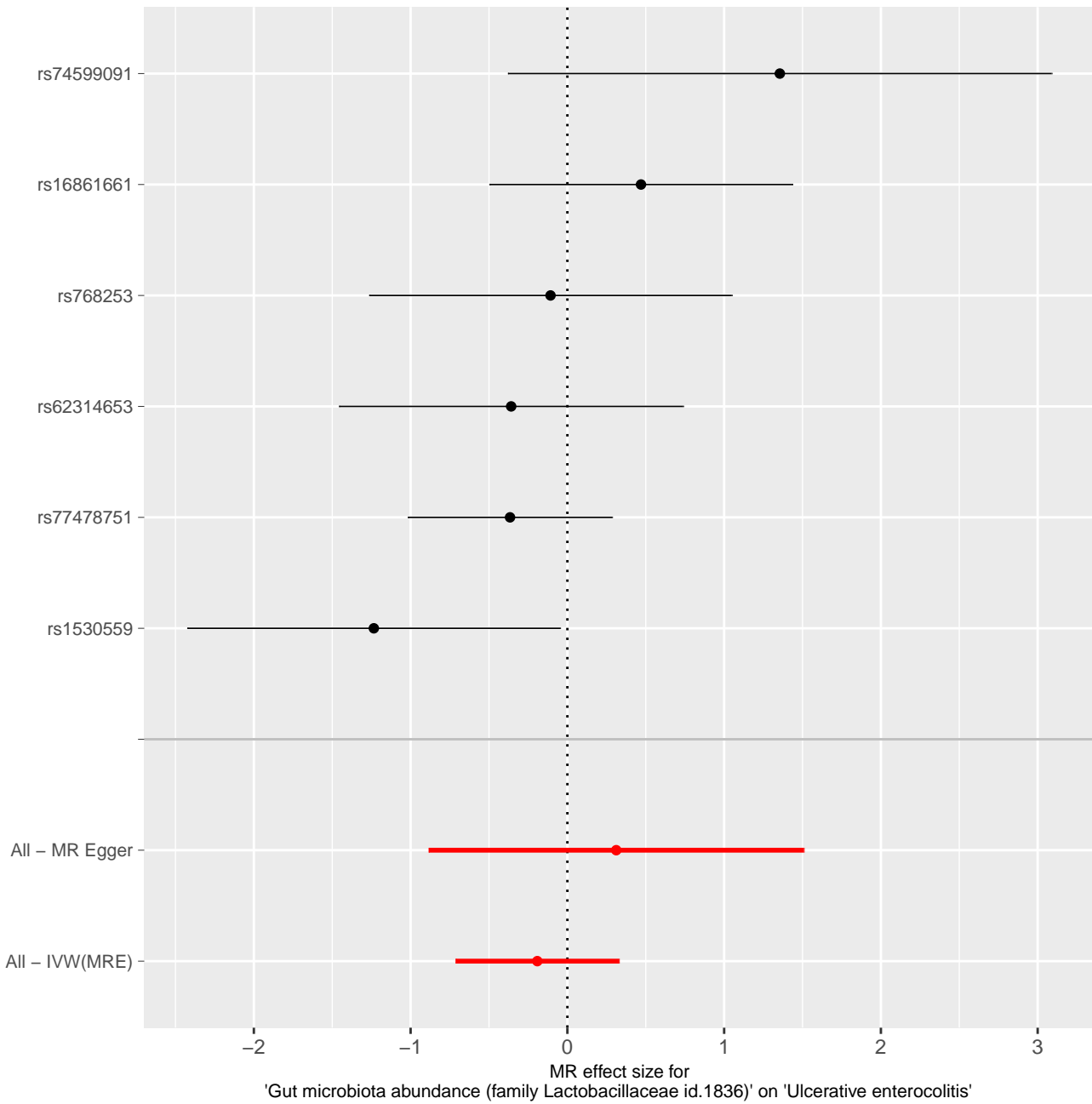
Batch 914 : Gut microbiota abundance (family Erysipelotrichaceae id.2149) on Ulcerative enterocolitis

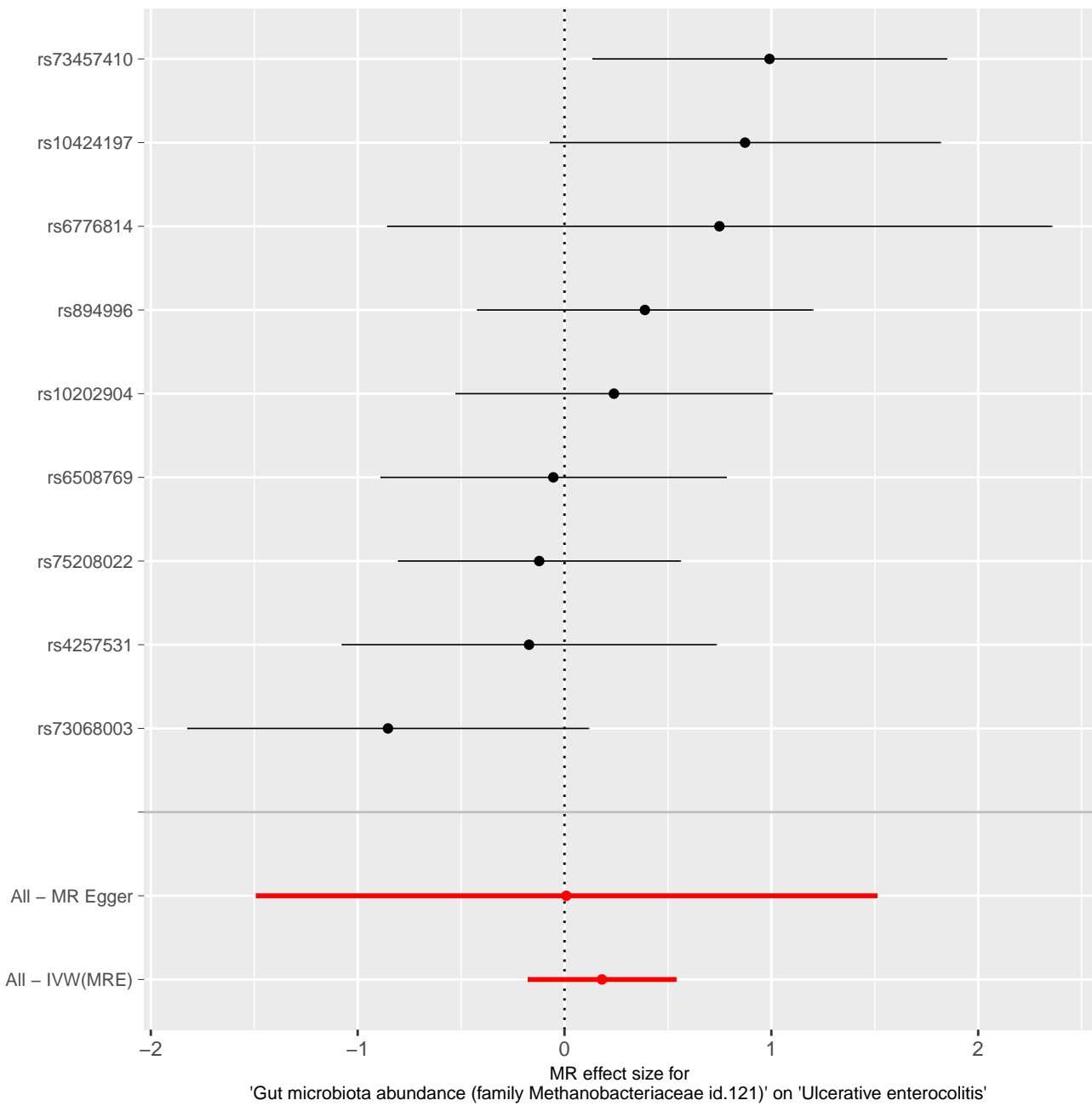


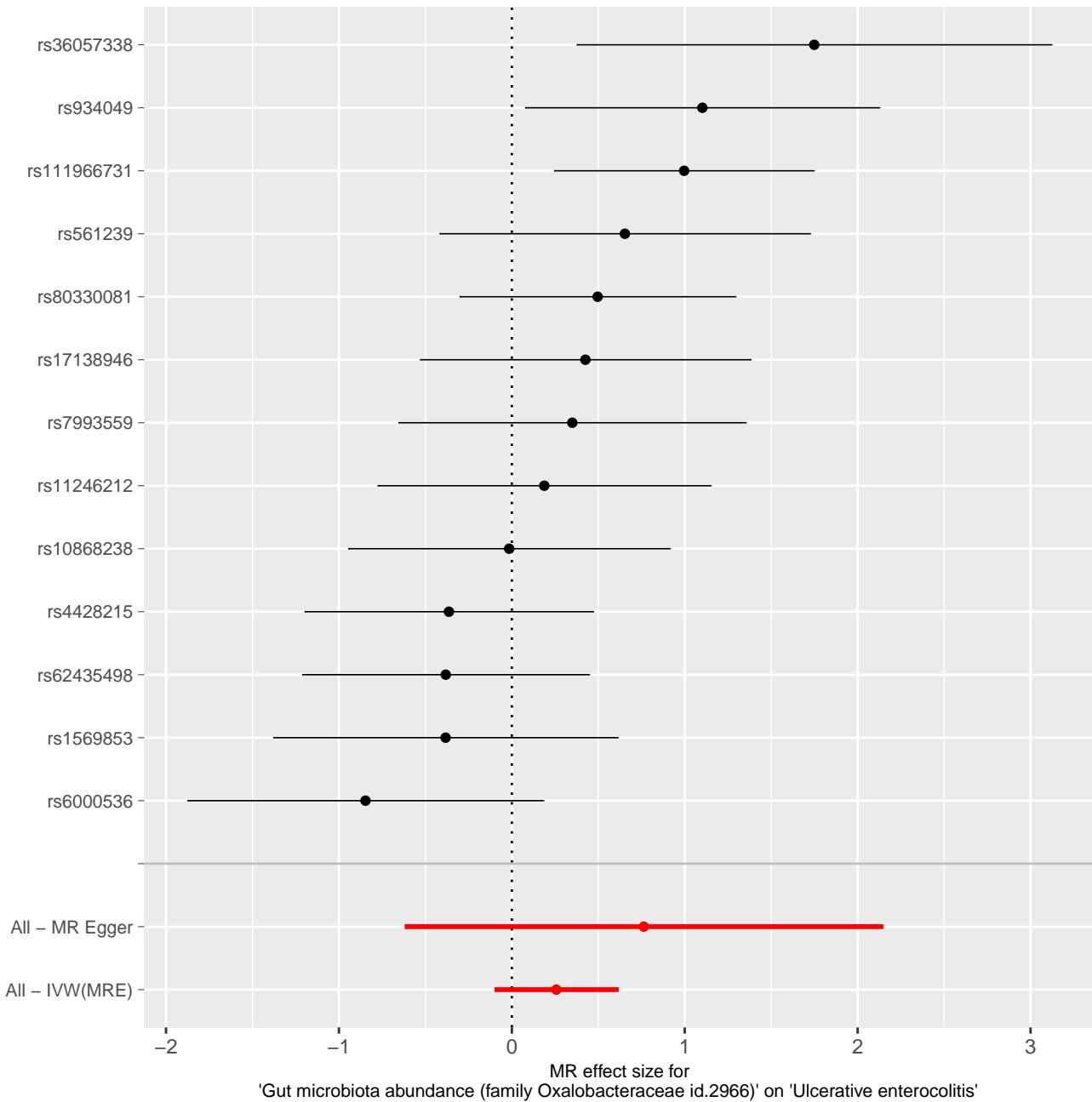




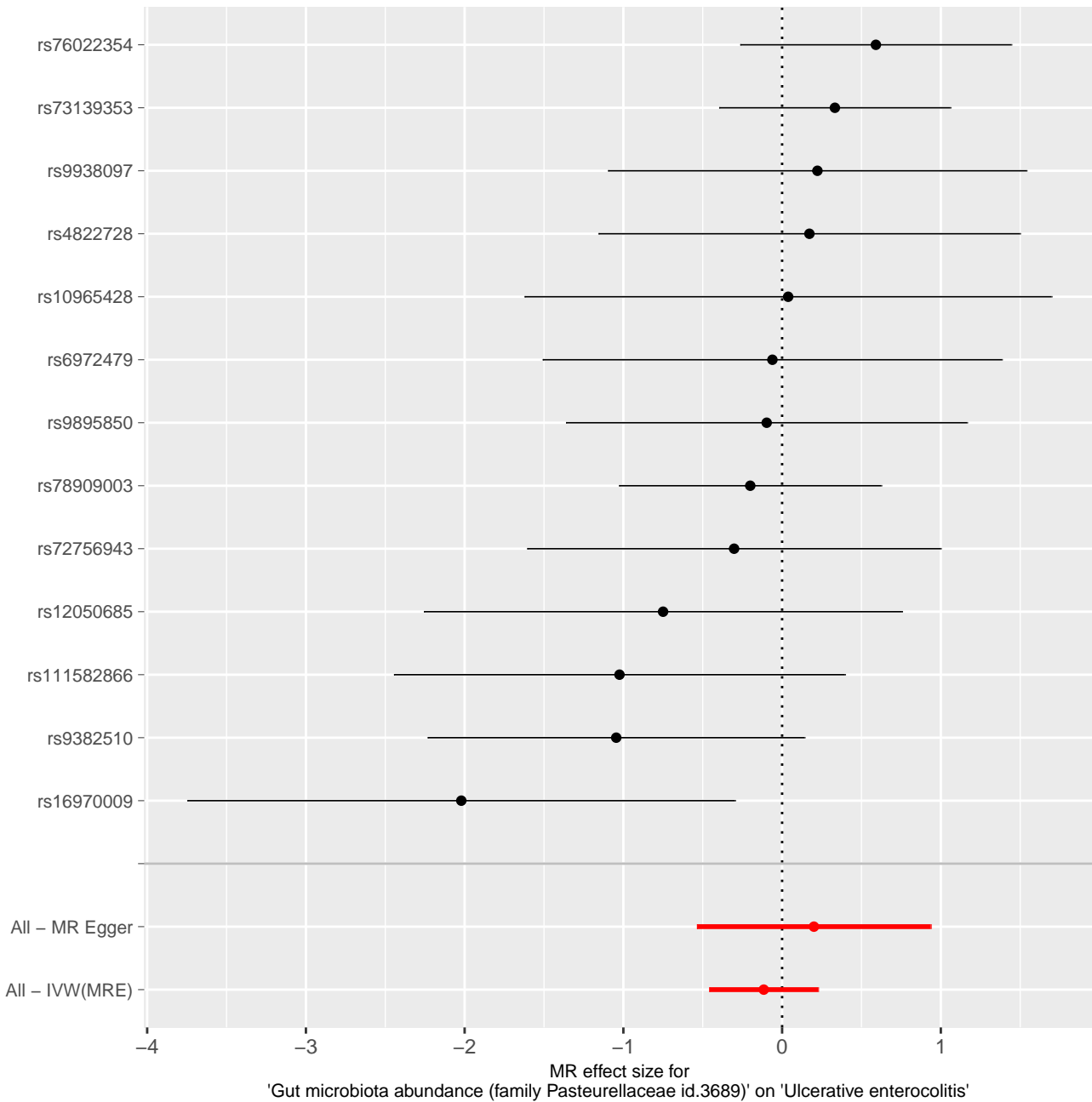


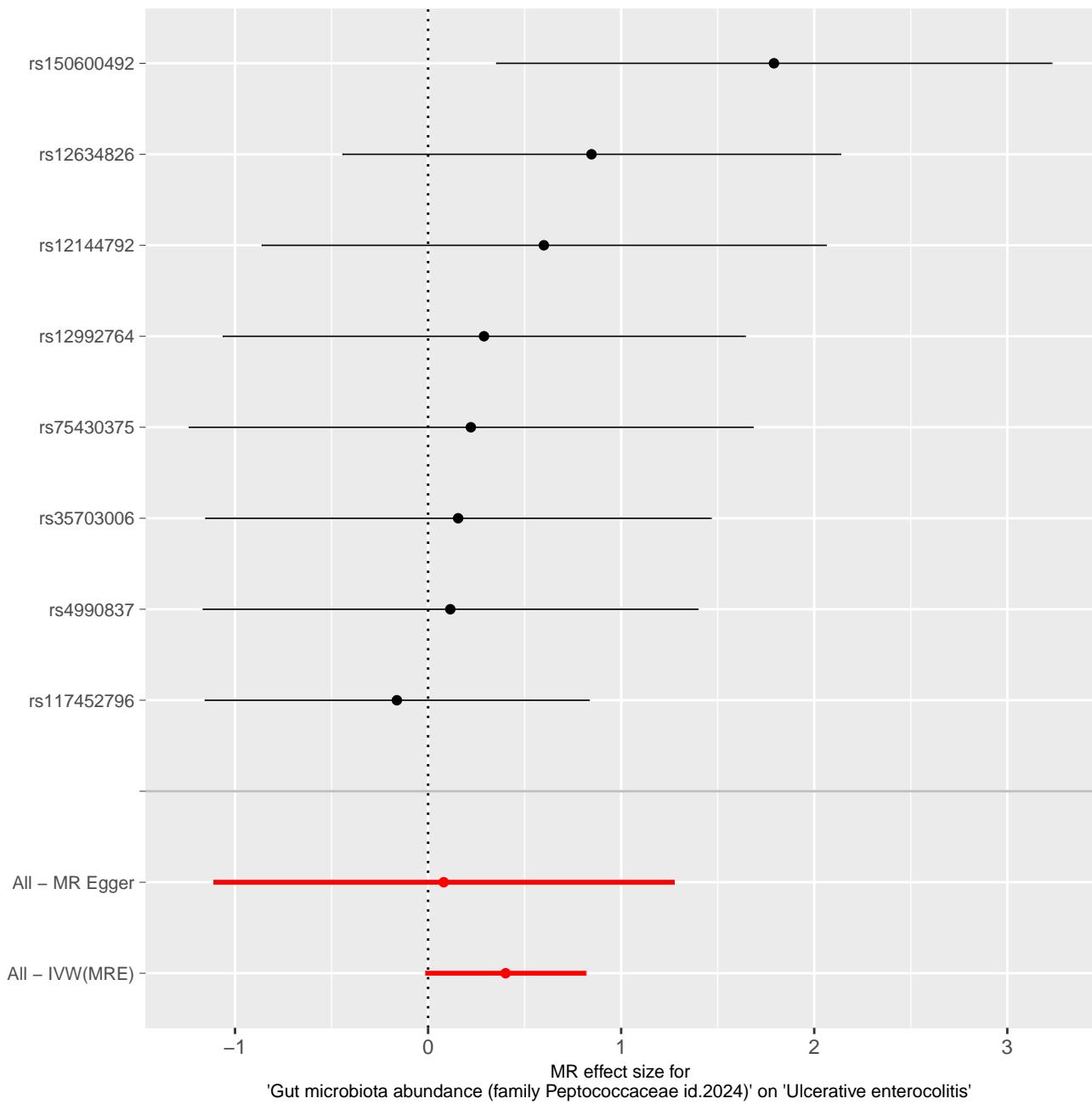


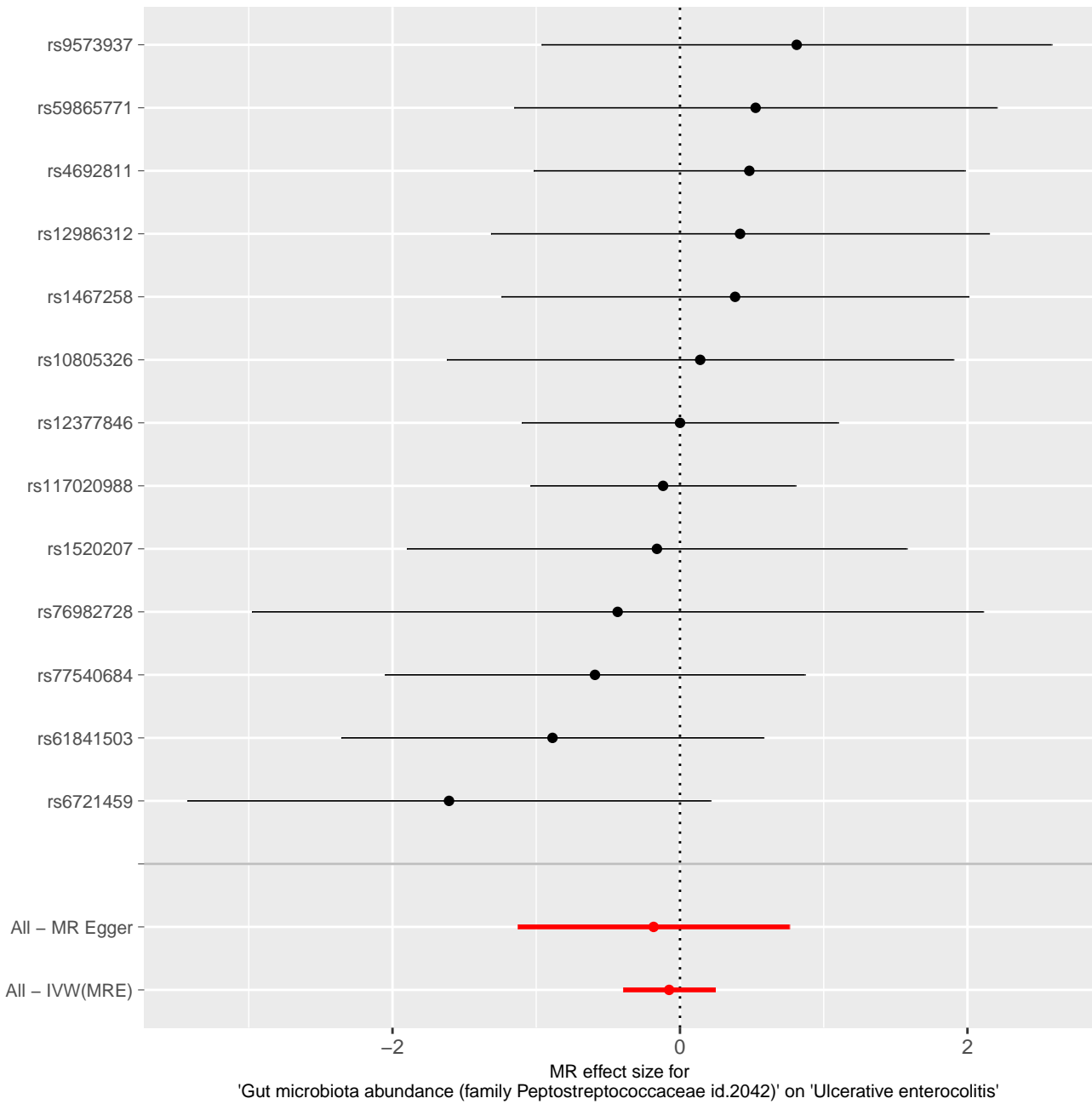


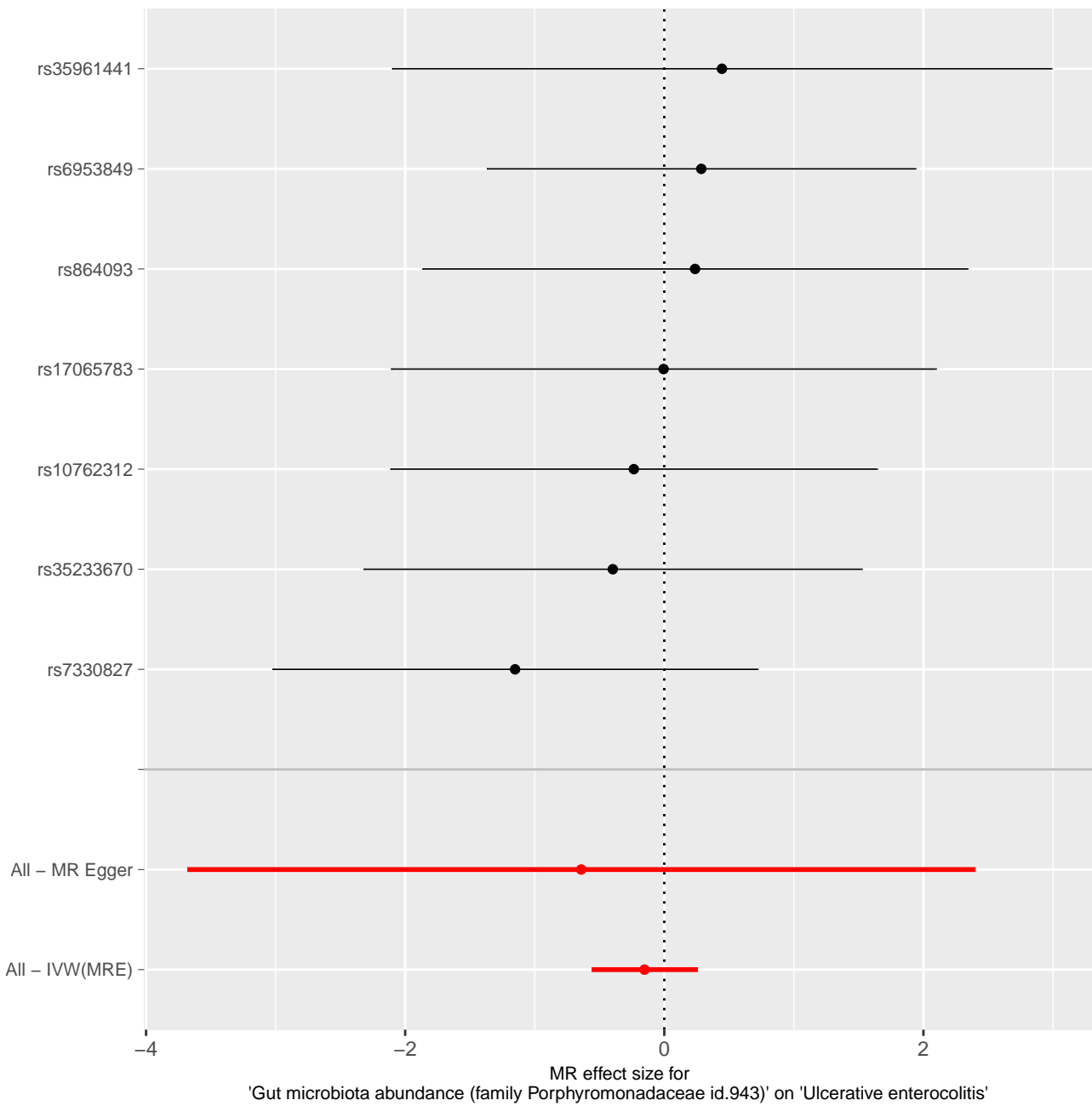


Batch 921 : Gut microbiota abundance (family Pasteurellaceae id.3689) on Ulcerative enterocolitis

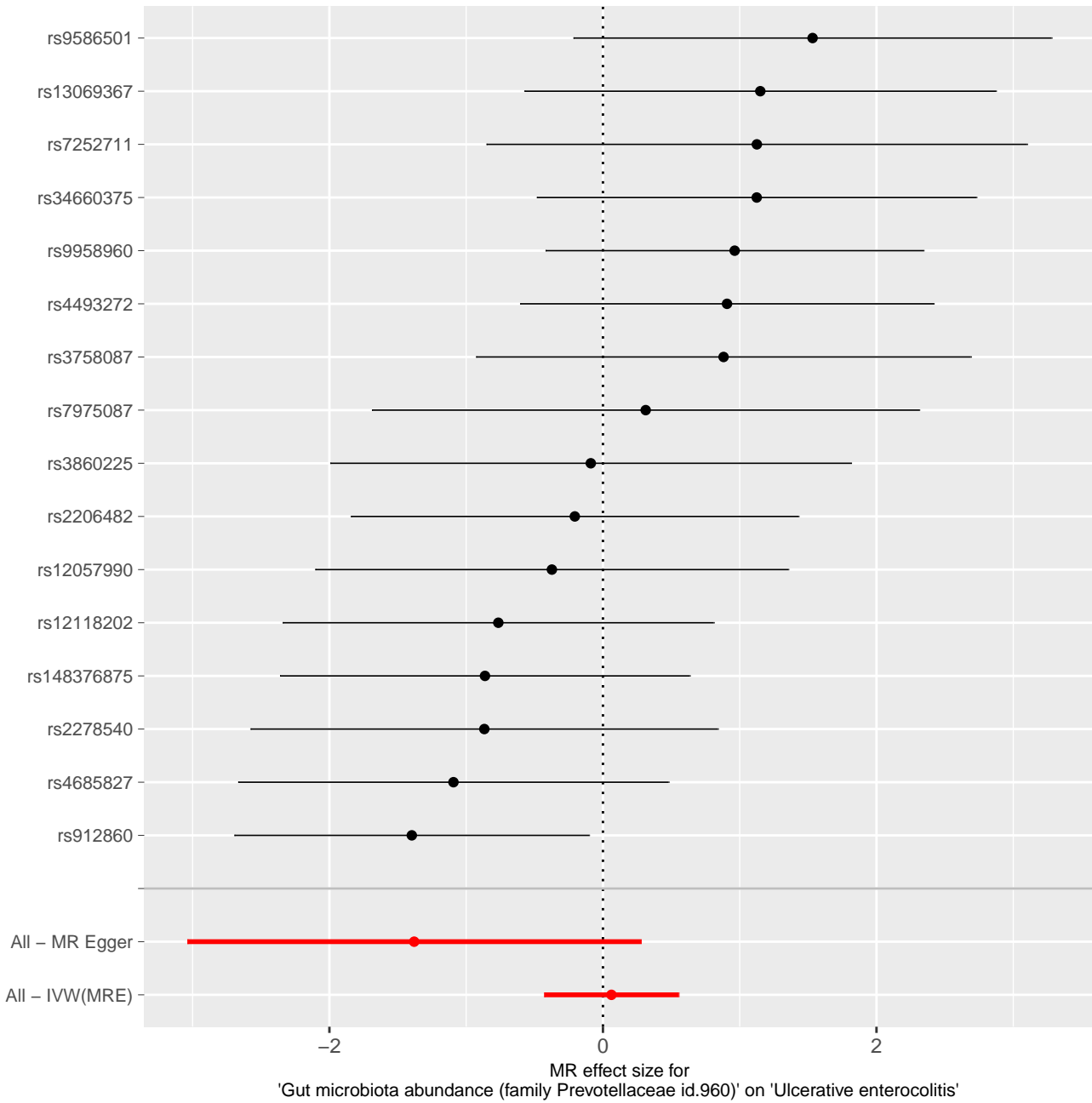


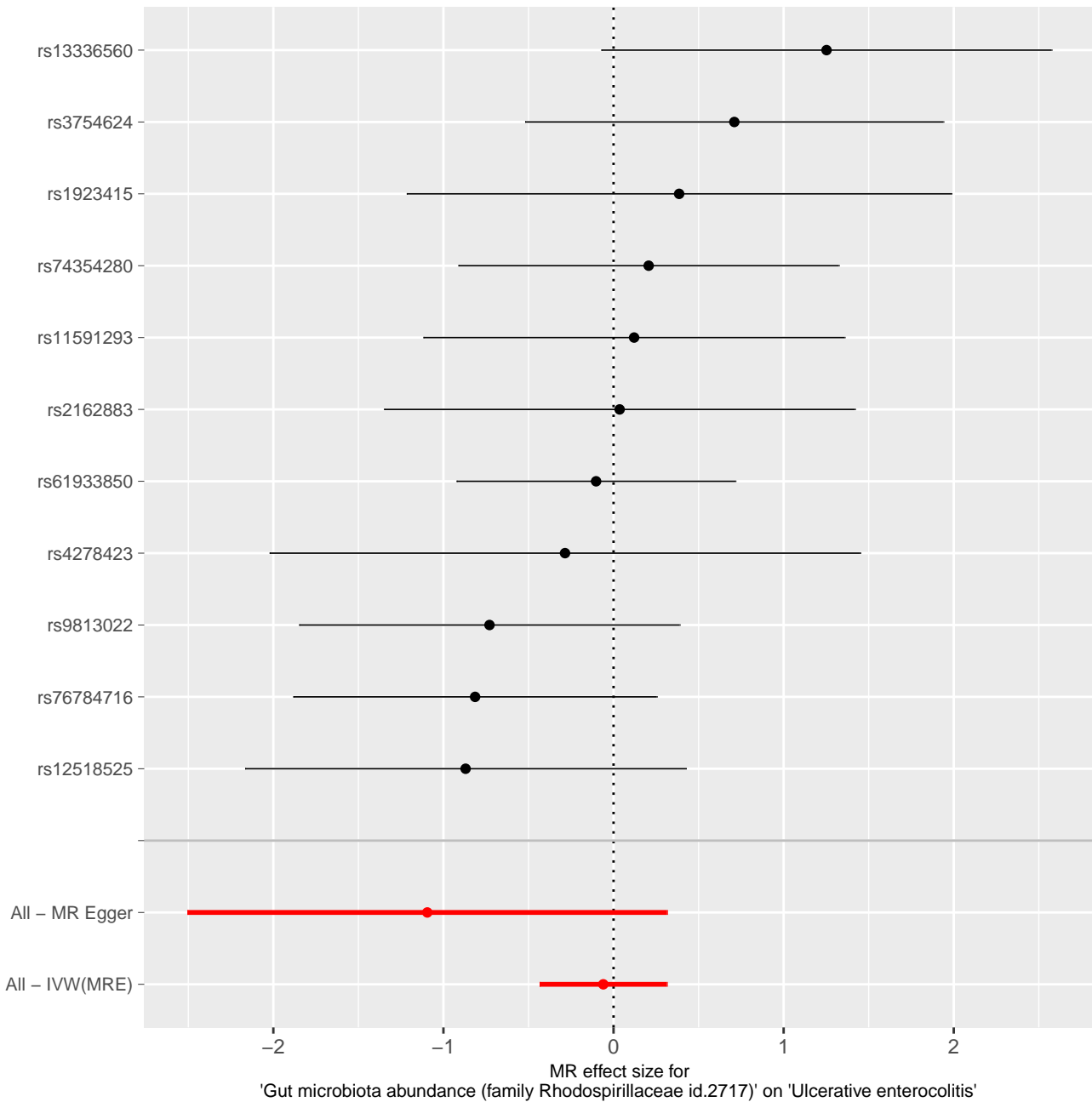


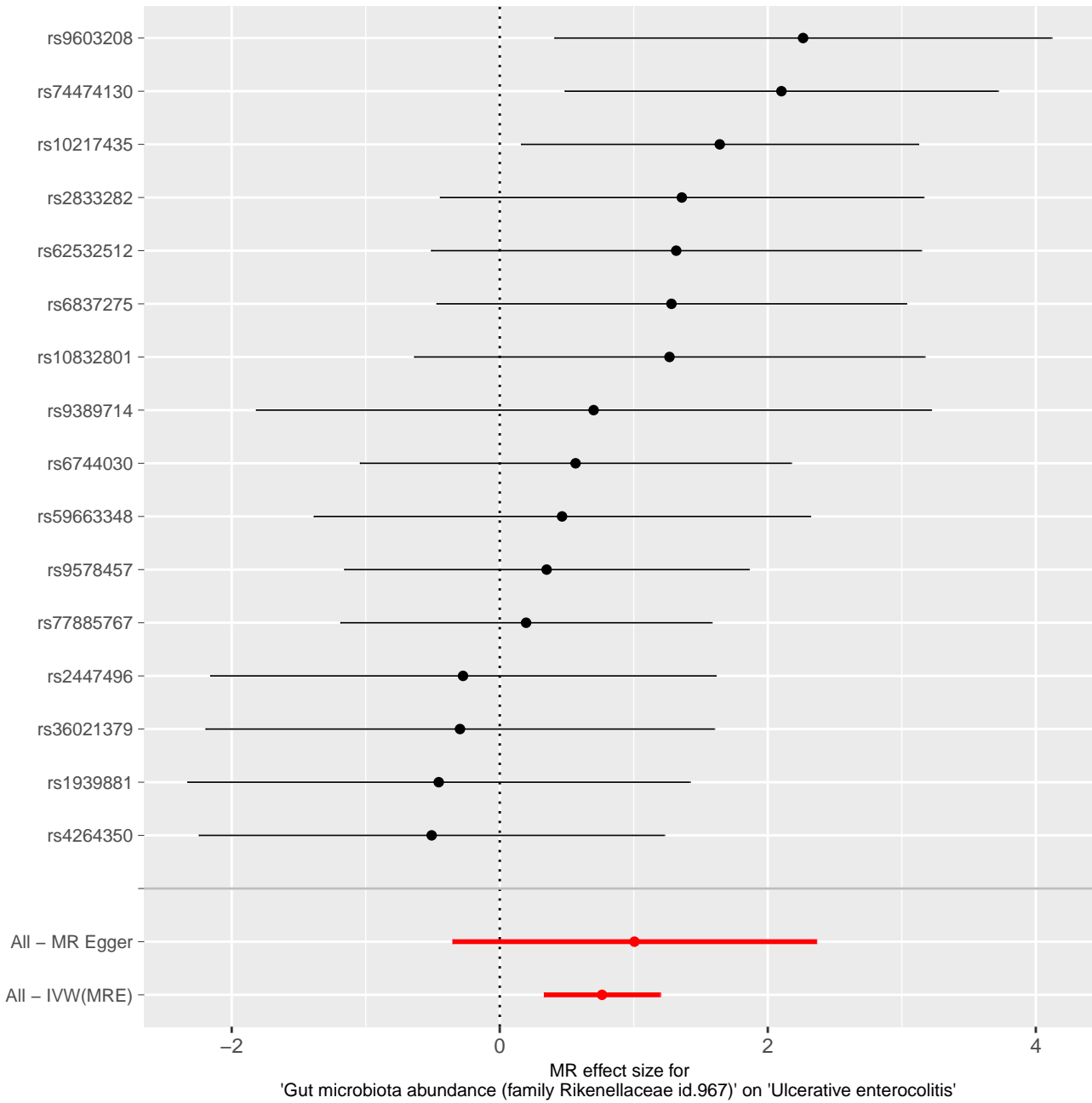


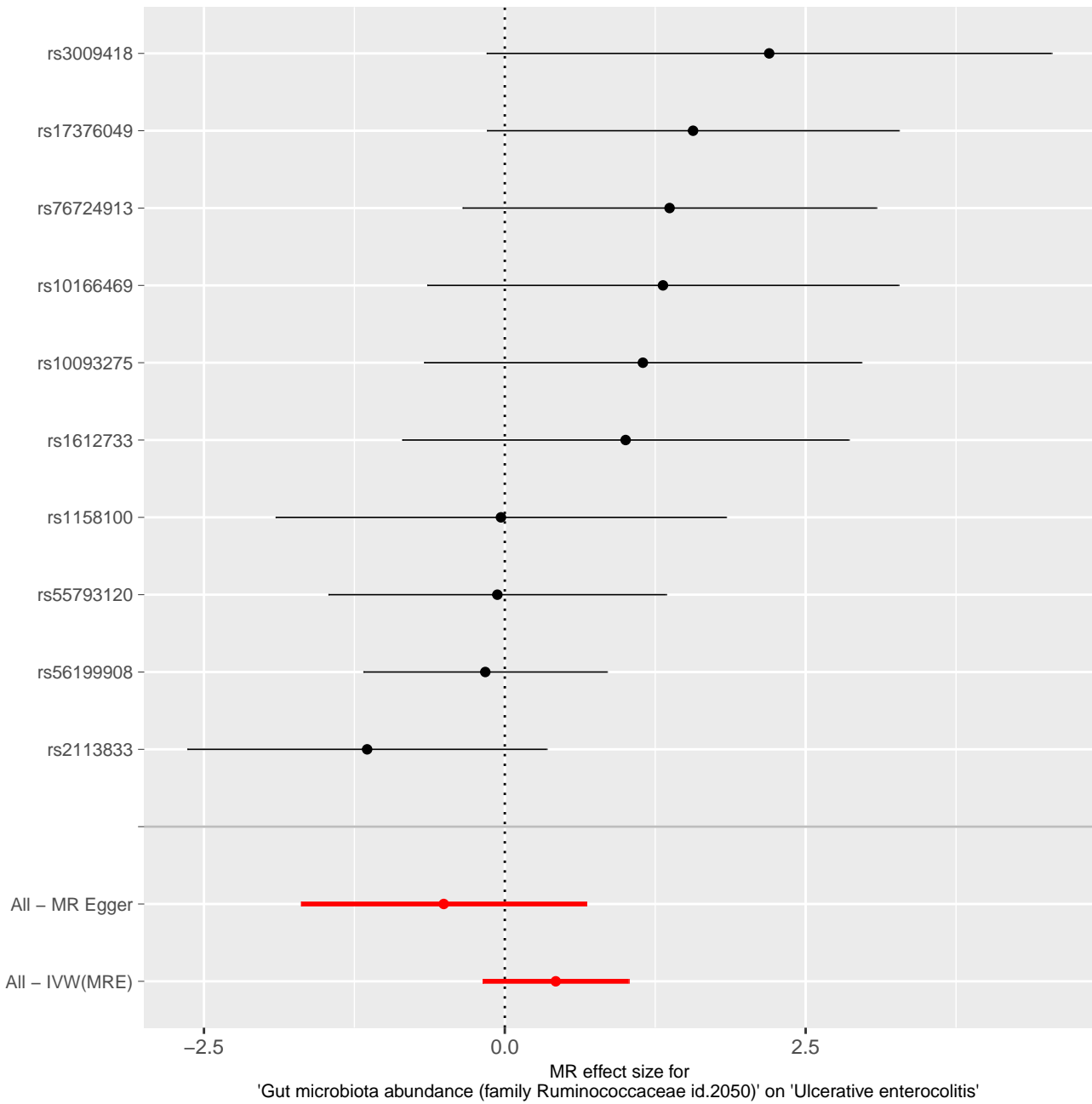


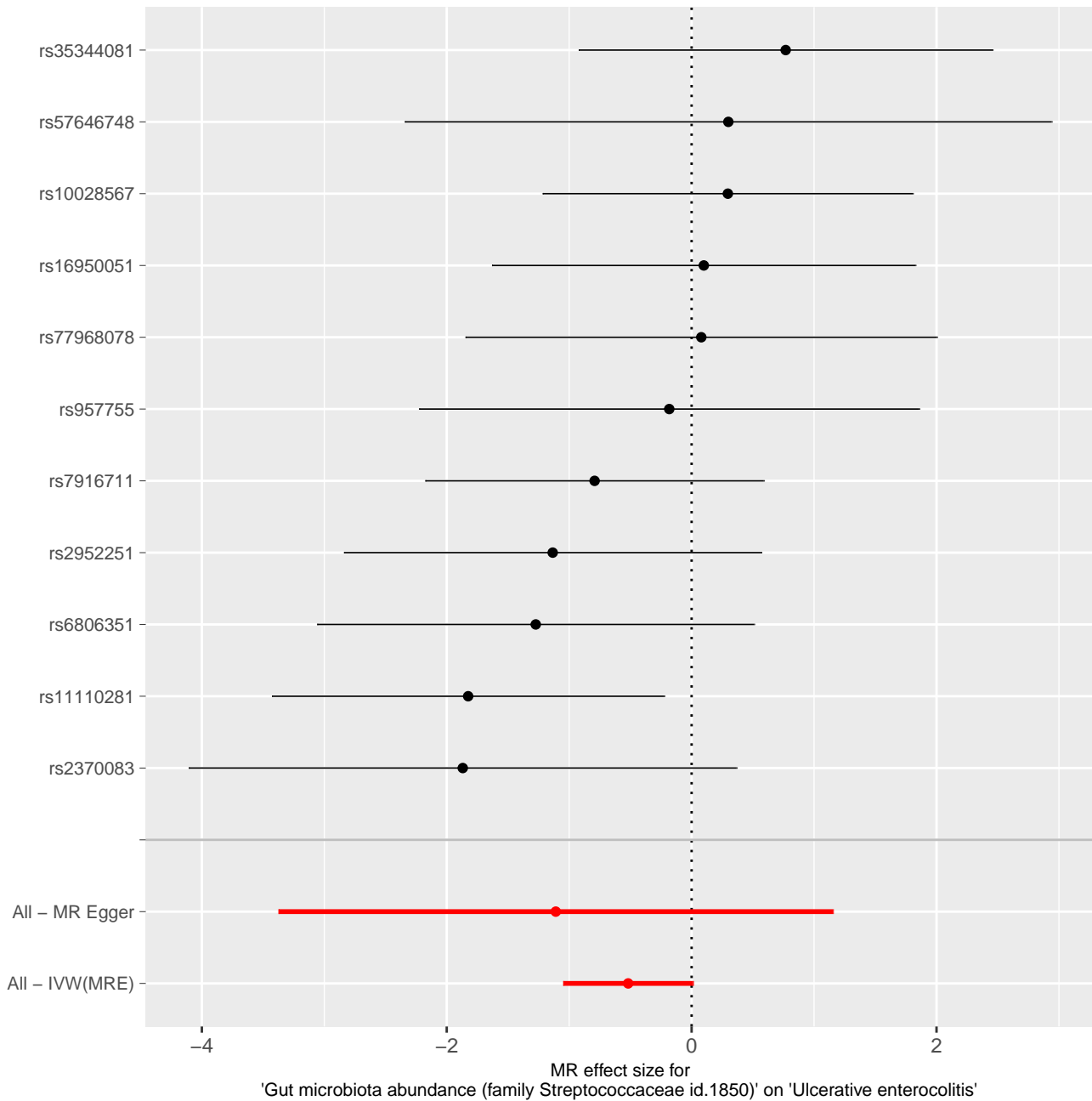
Batch 925 : Gut microbiota abundance (family Prevotellaceae id.960) on Ulcerative enterocolitis

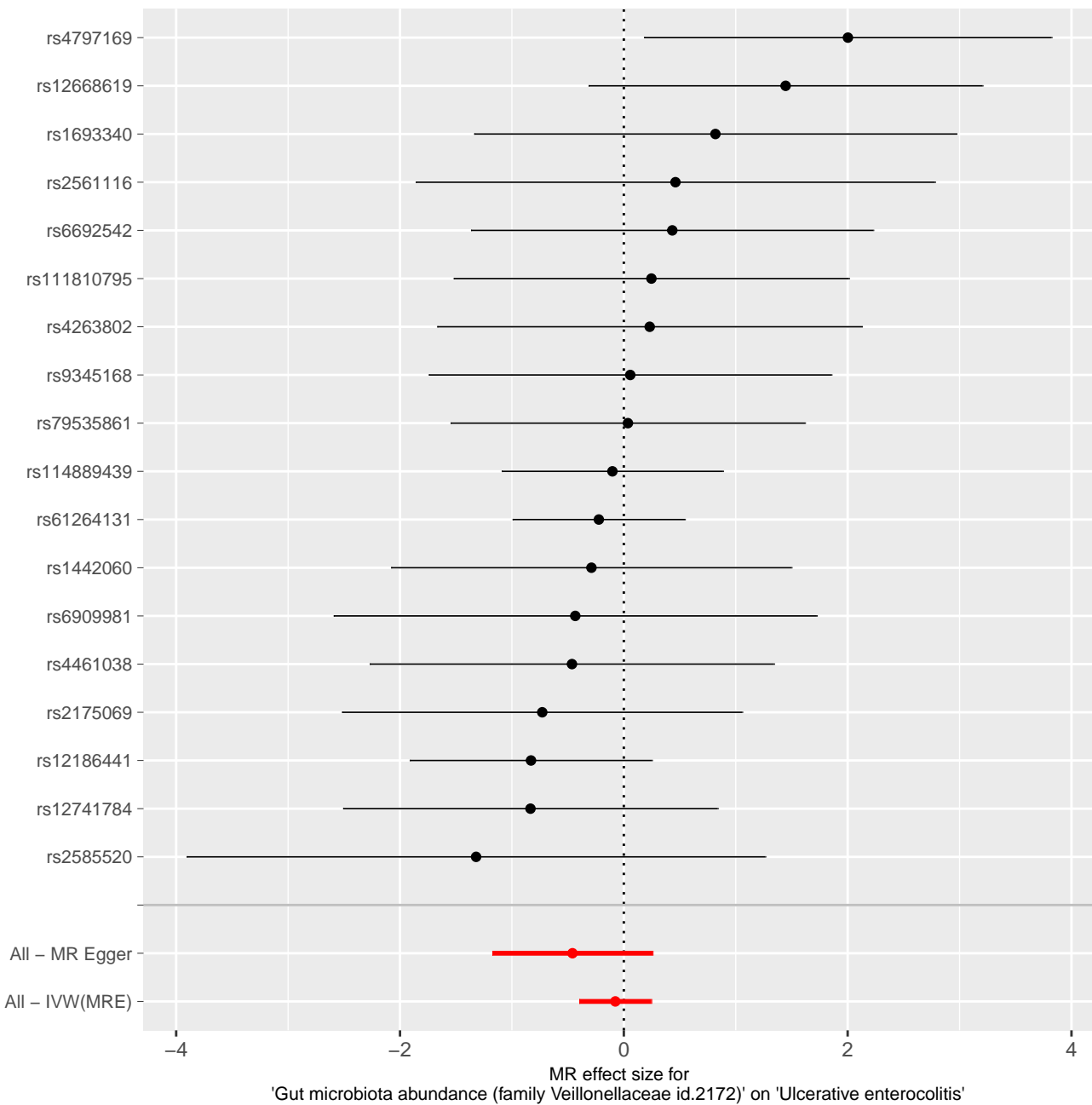


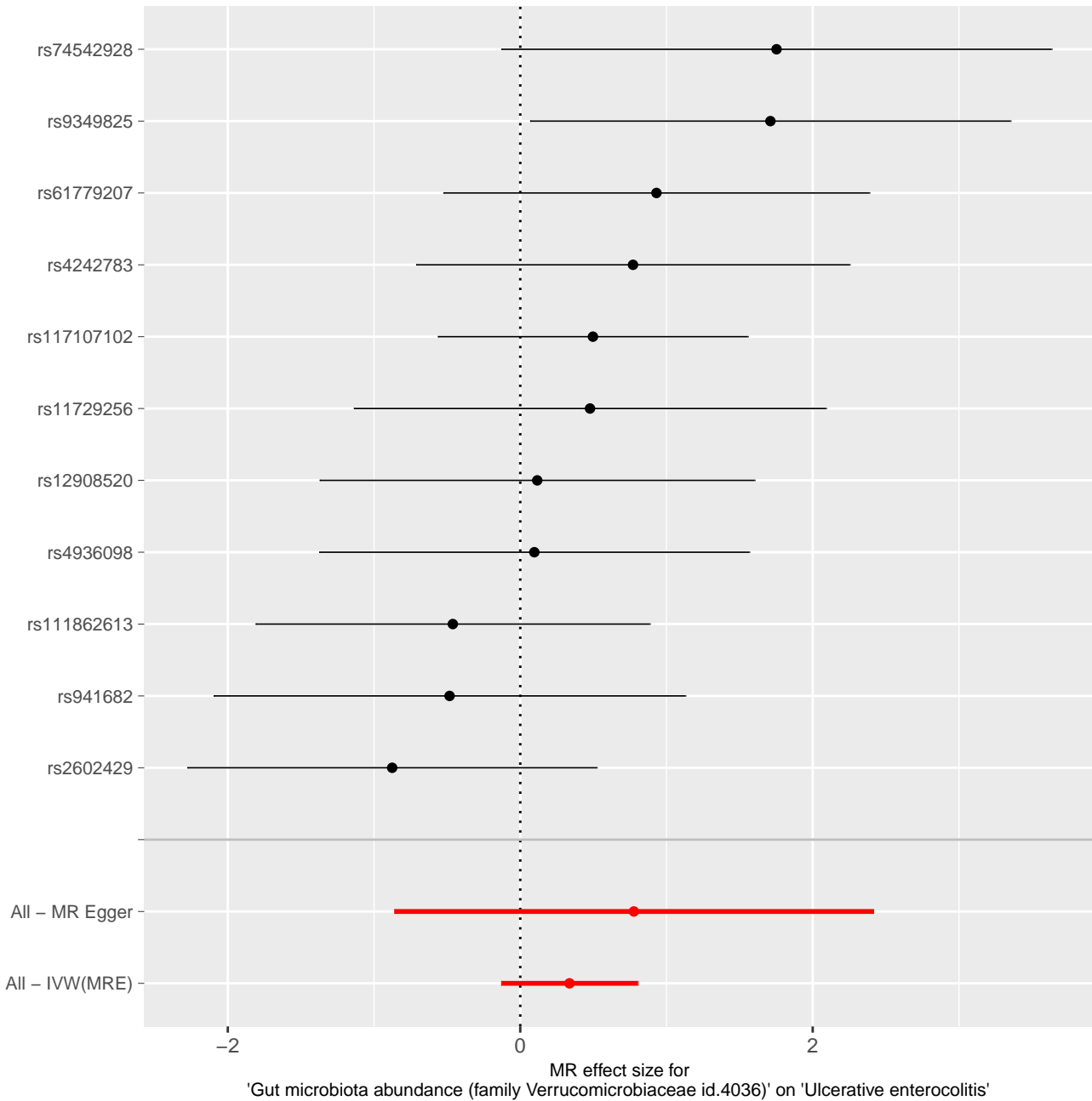




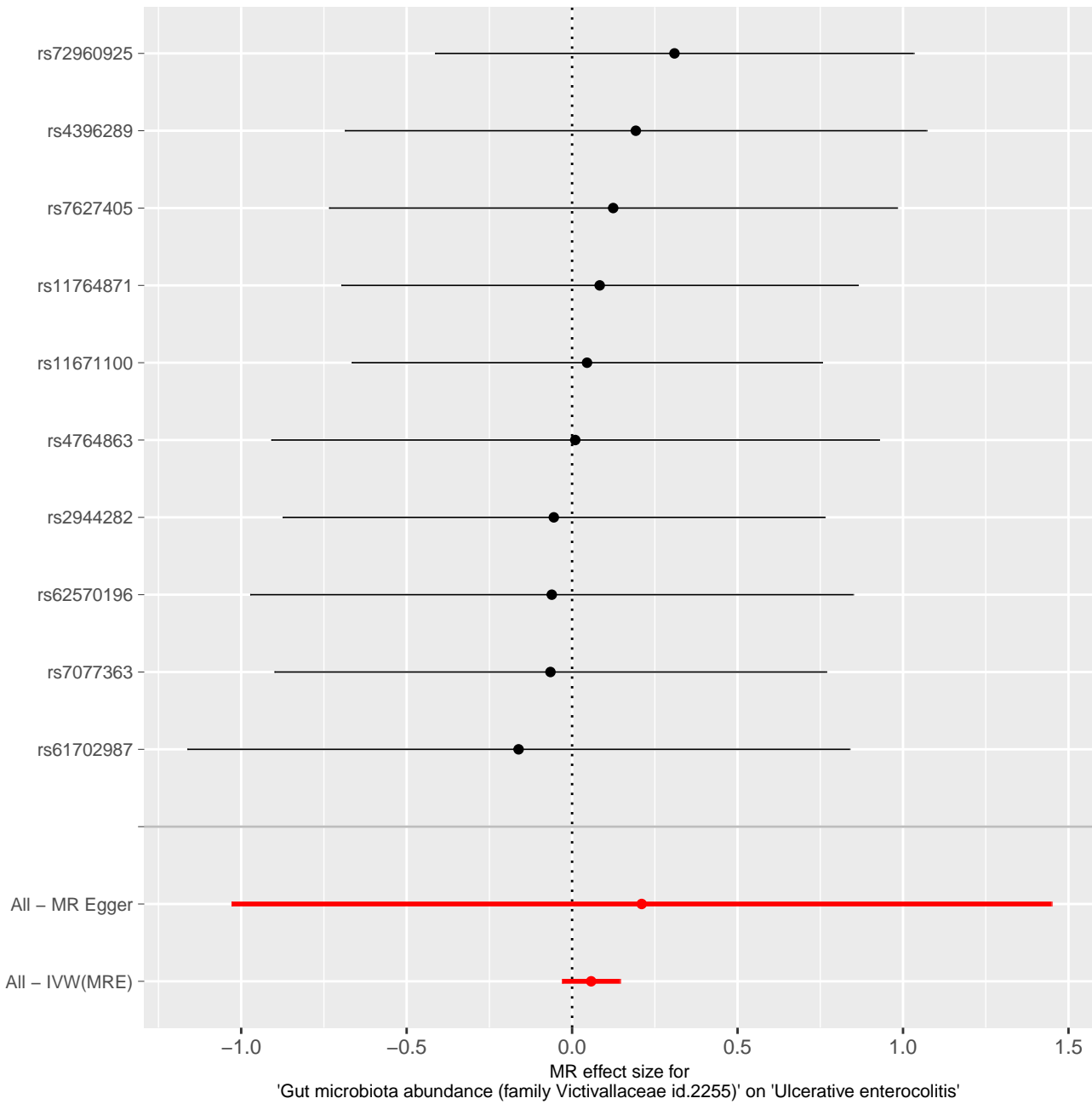


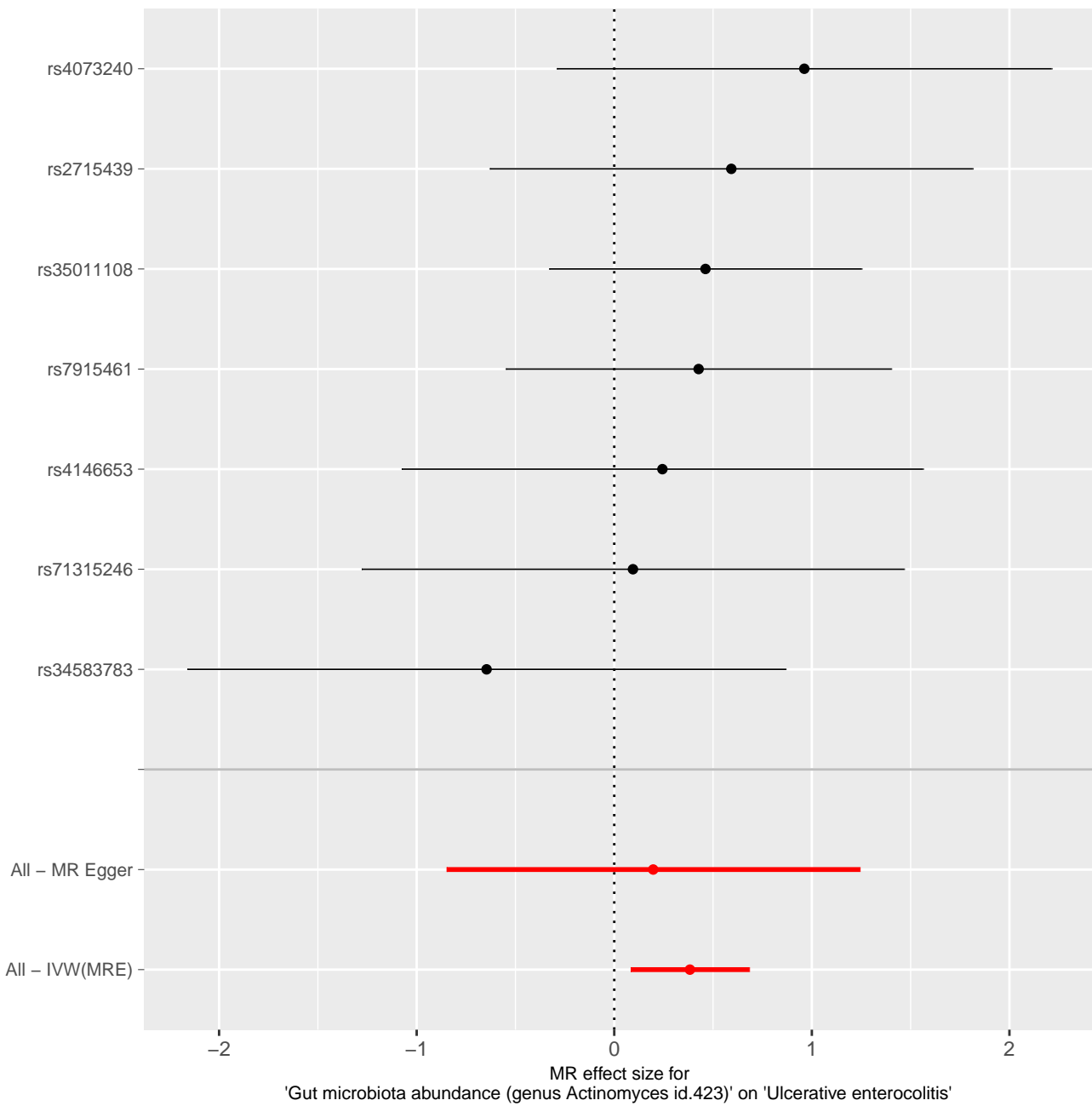


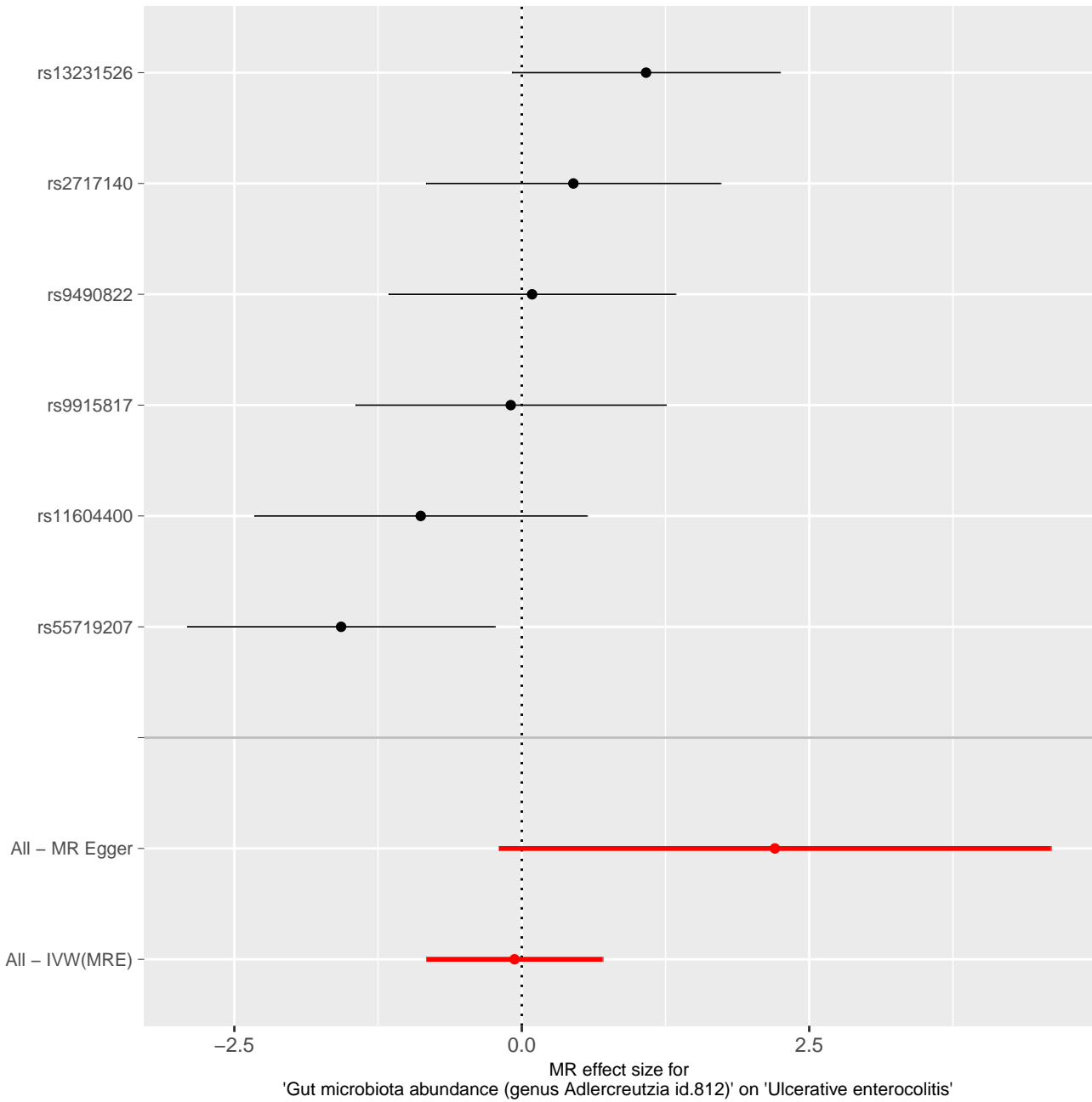


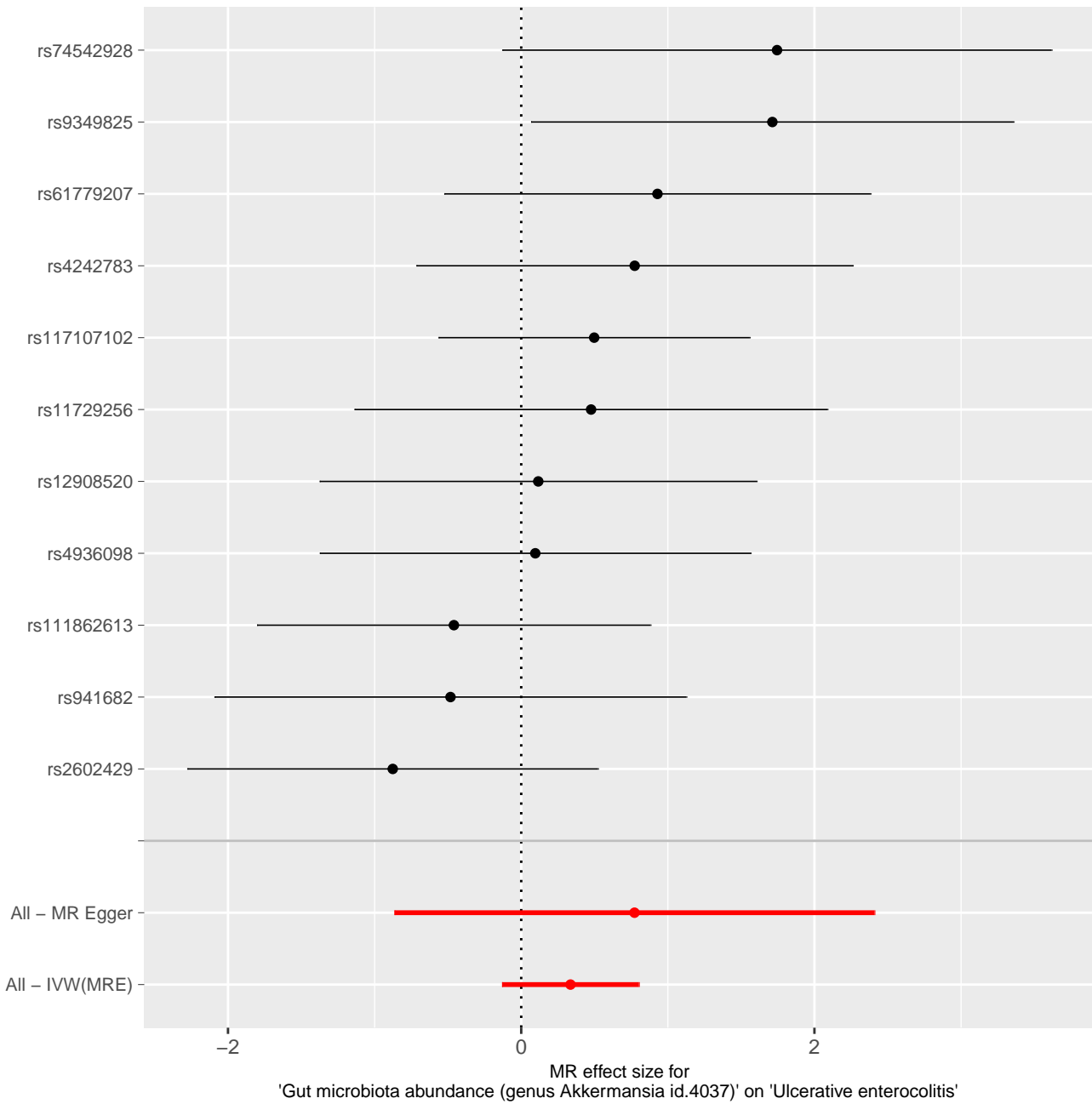


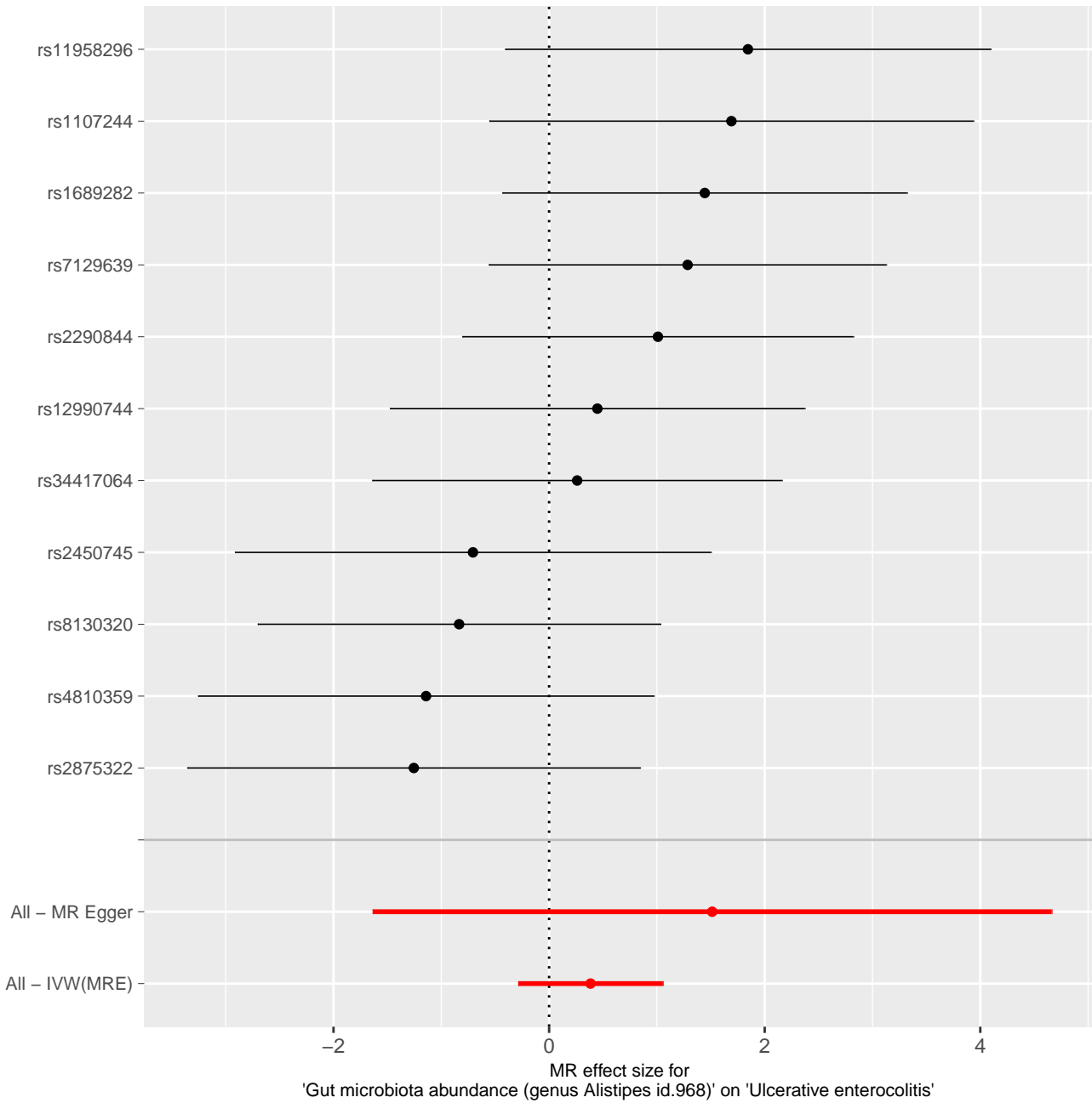
Batch 932 : Gut microbiota abundance (family Victivallaceae id.2255) on Ulcerative enterocolitis

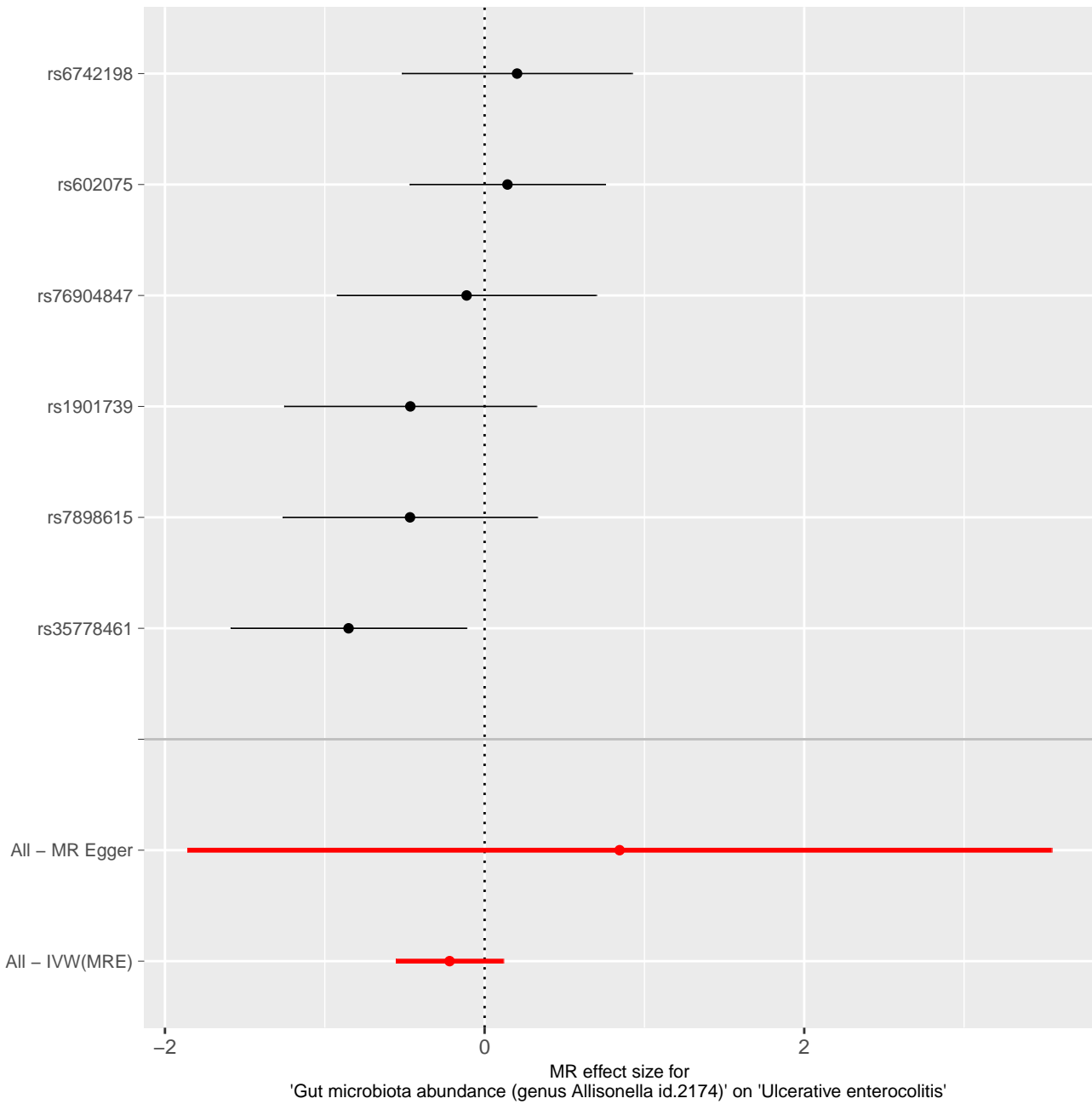


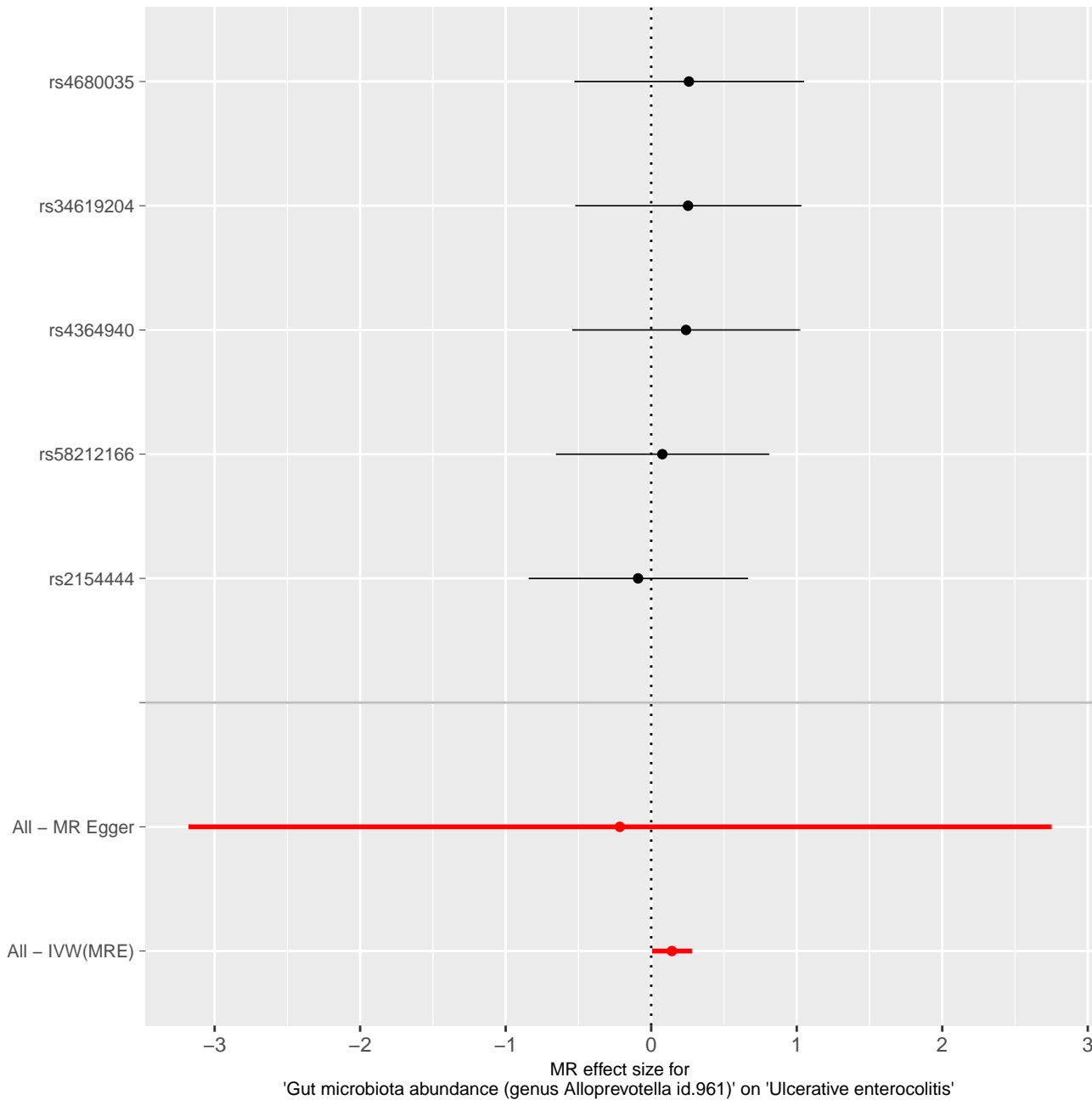


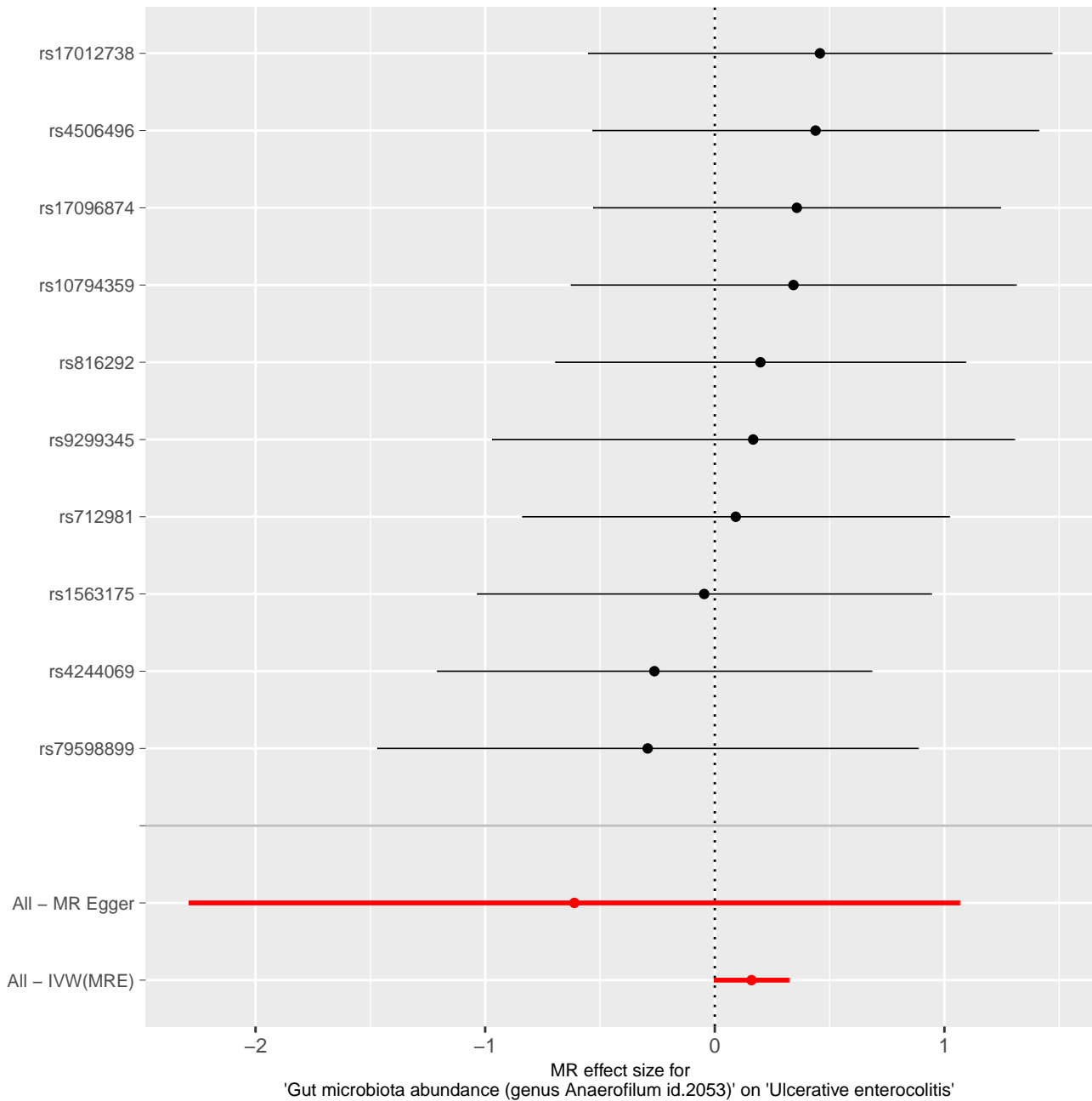


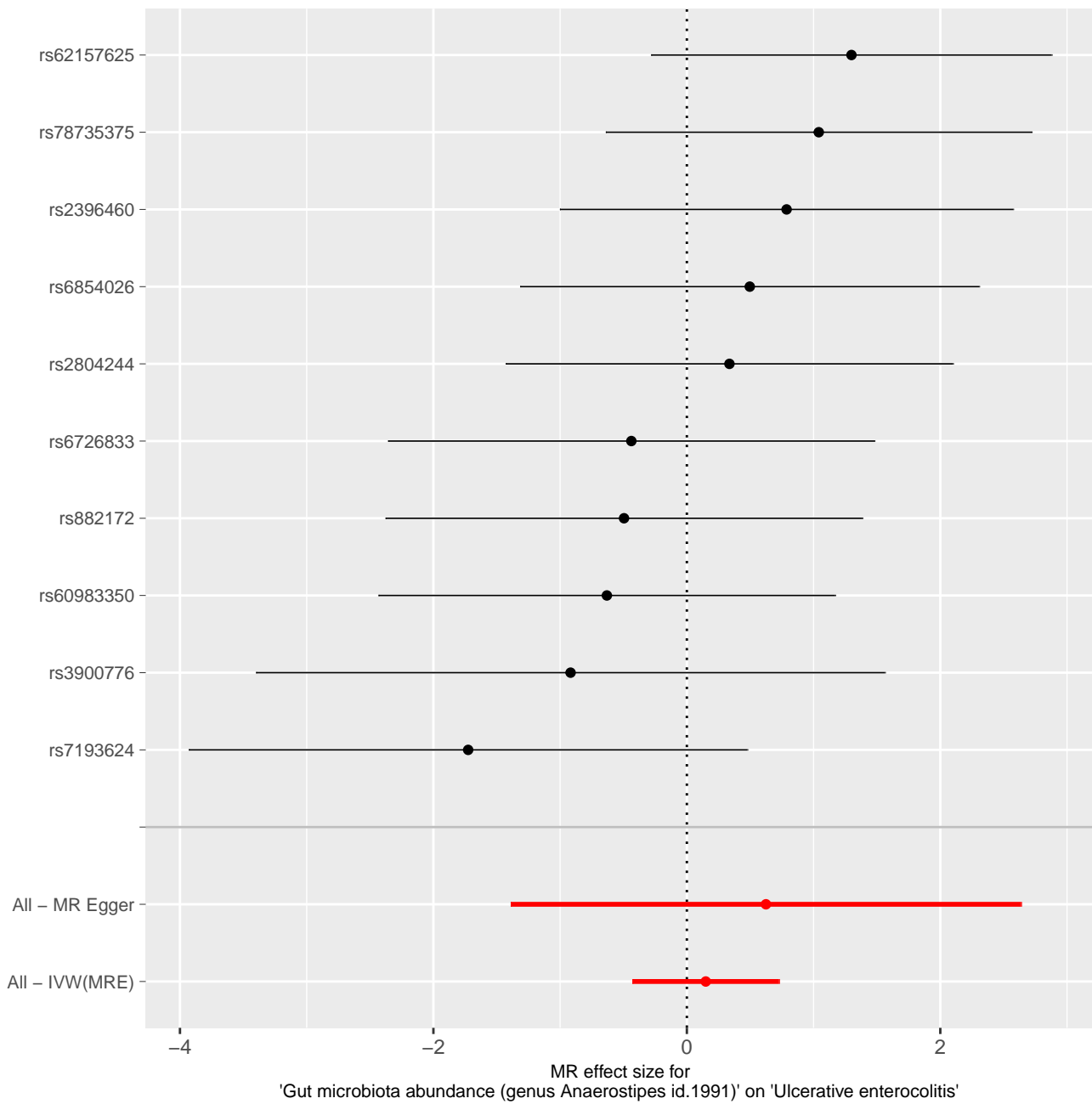




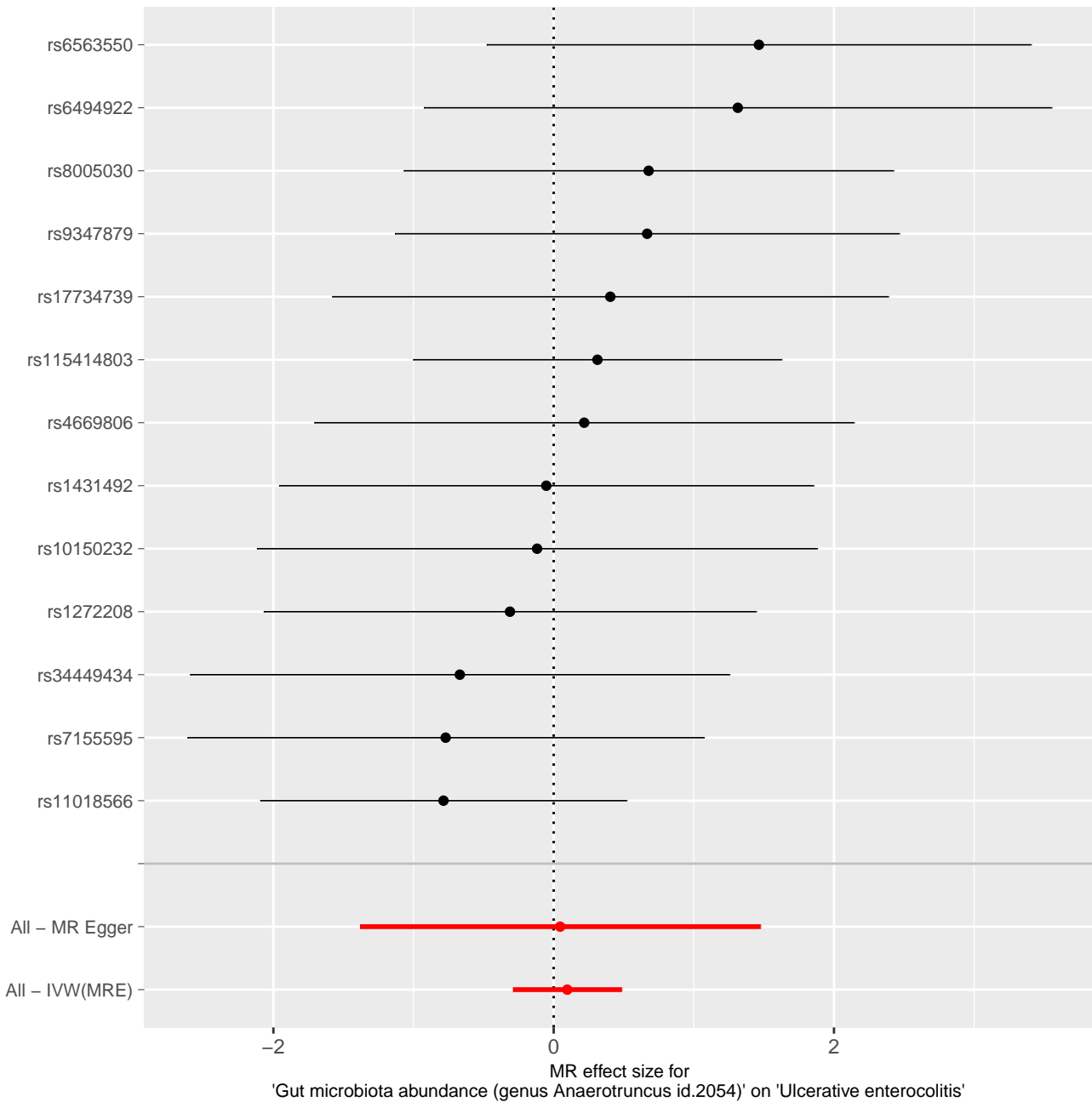


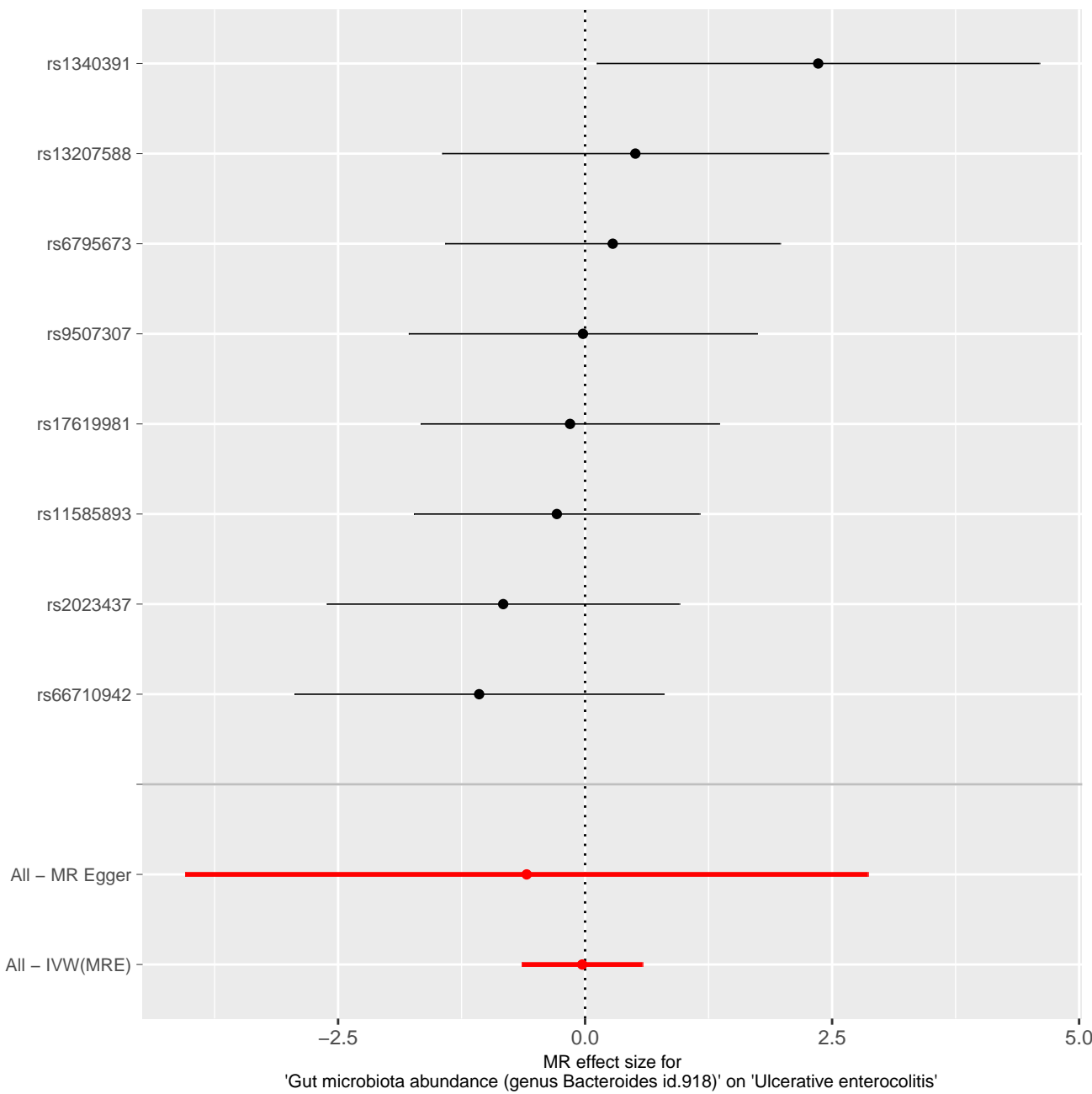


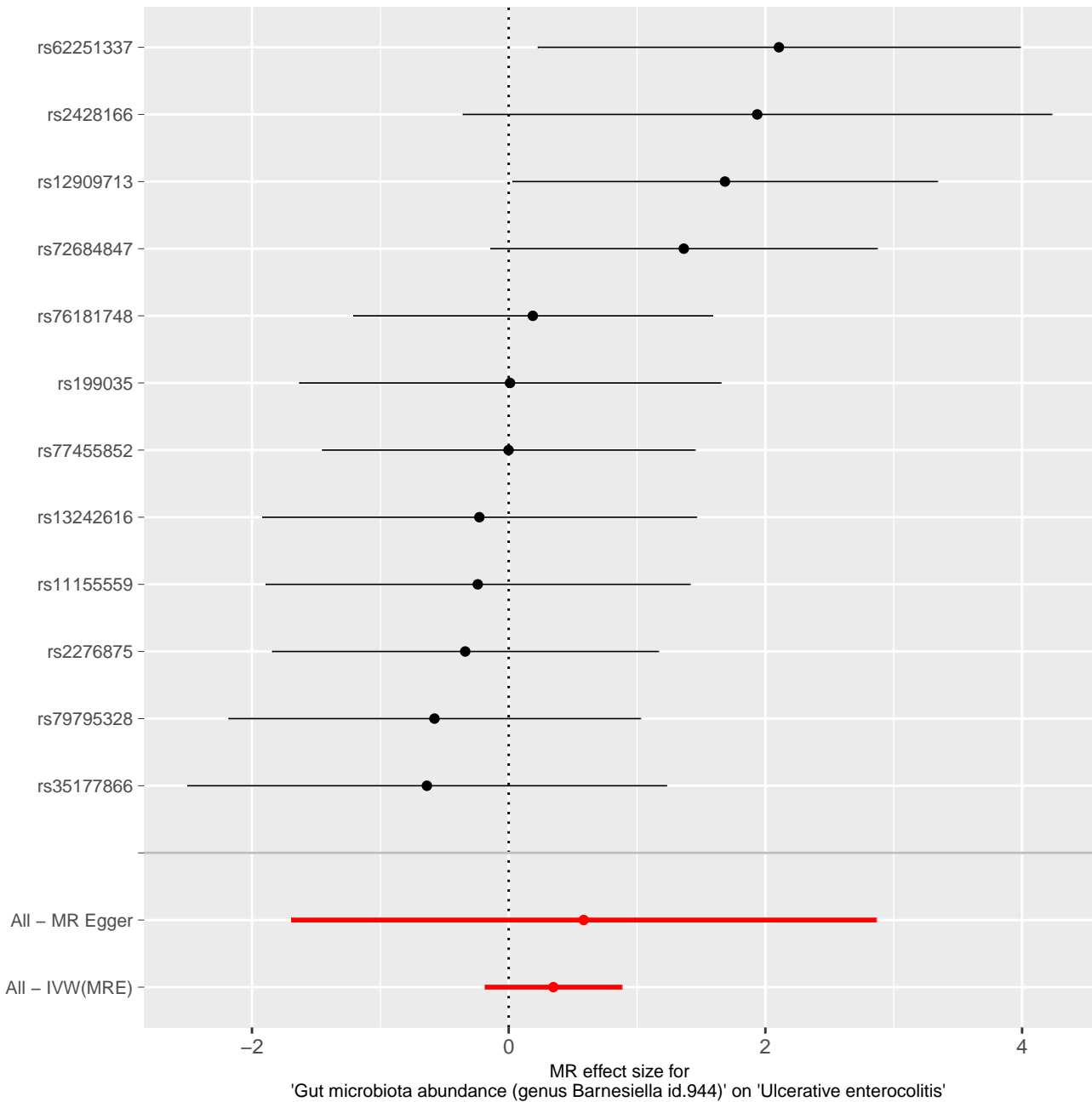


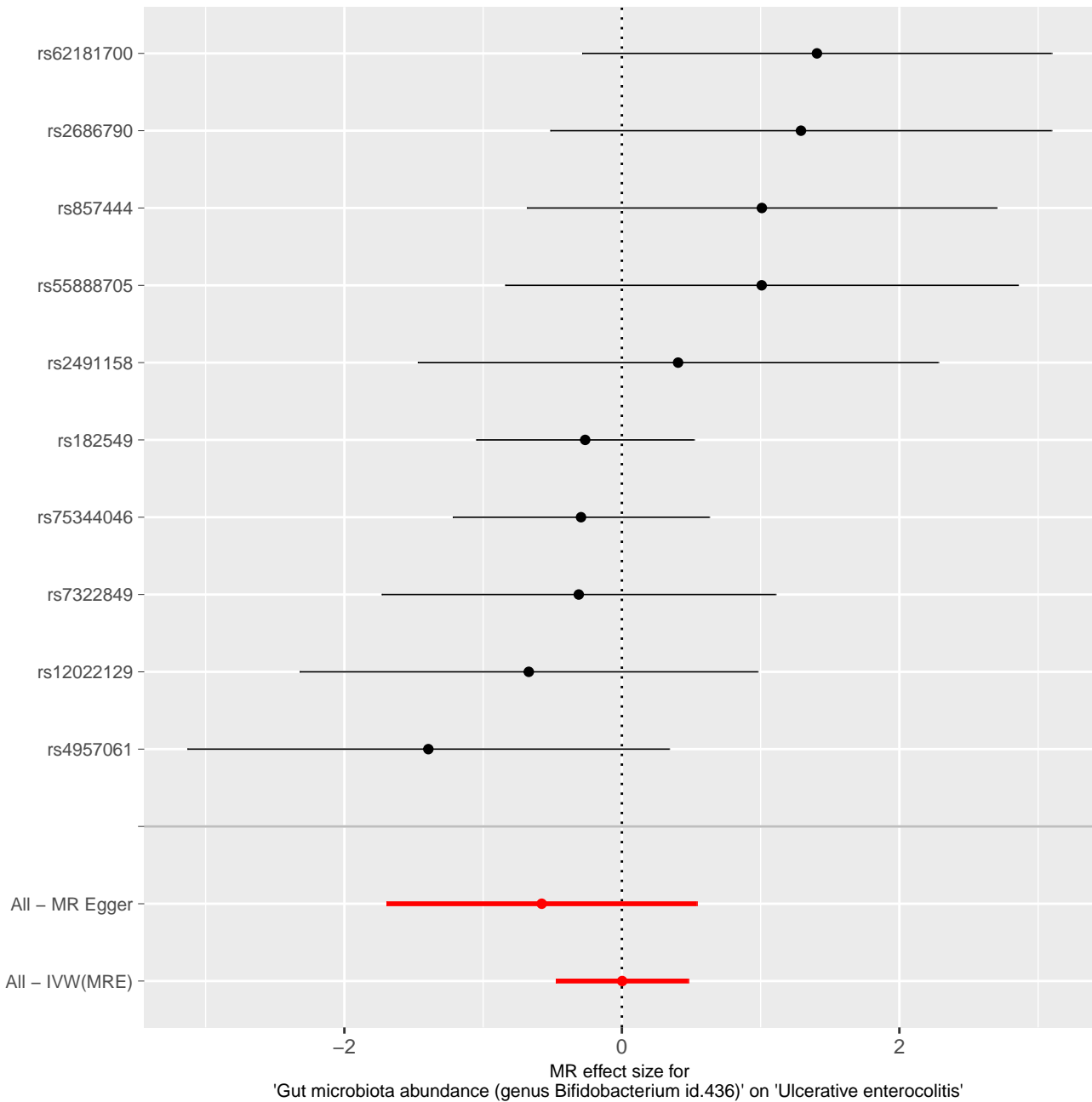


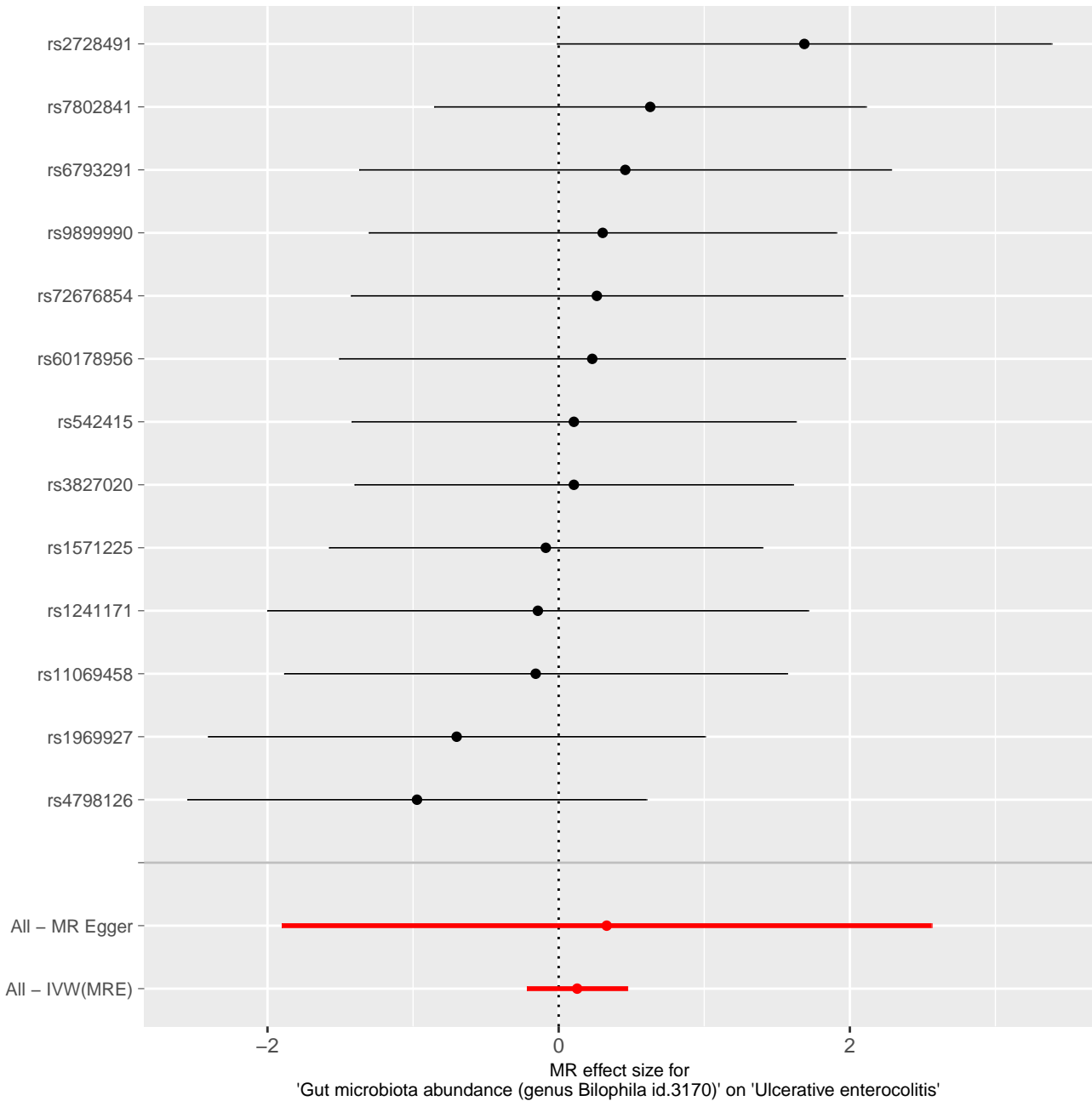
Batch 941 : Gut microbiota abundance (genus Anaerotruncus id.2054) on Ulcerative enterocolitis

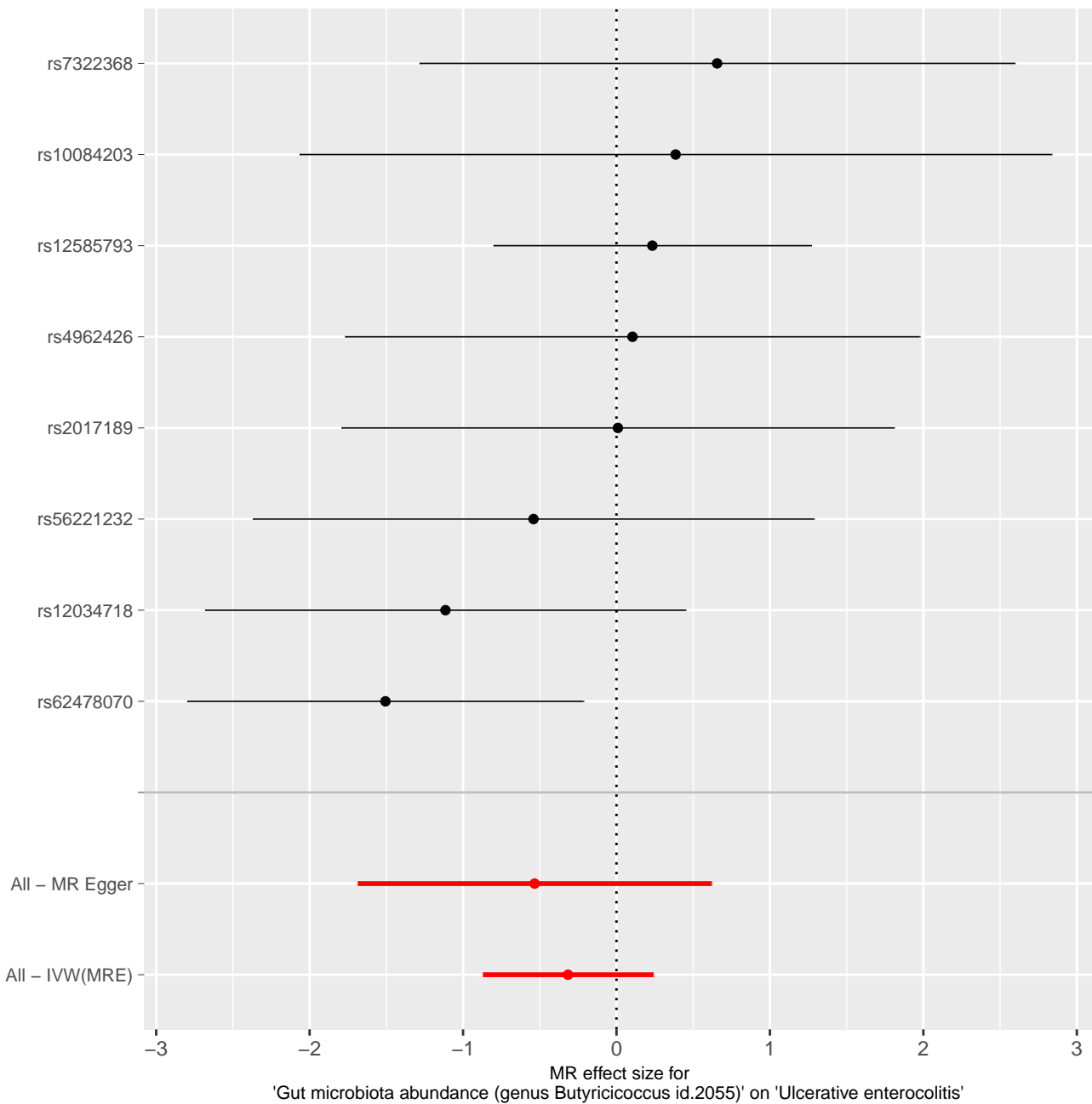




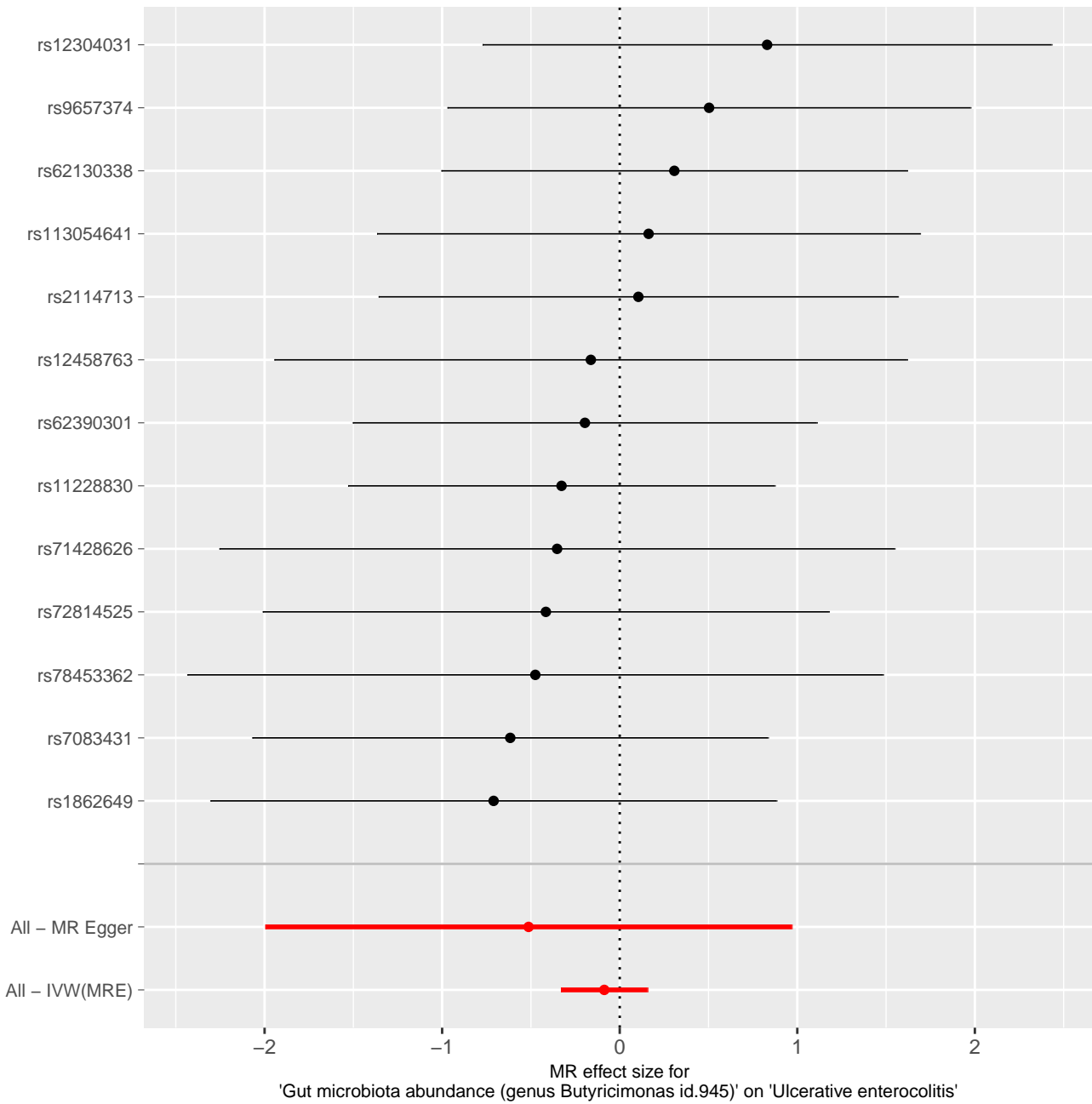




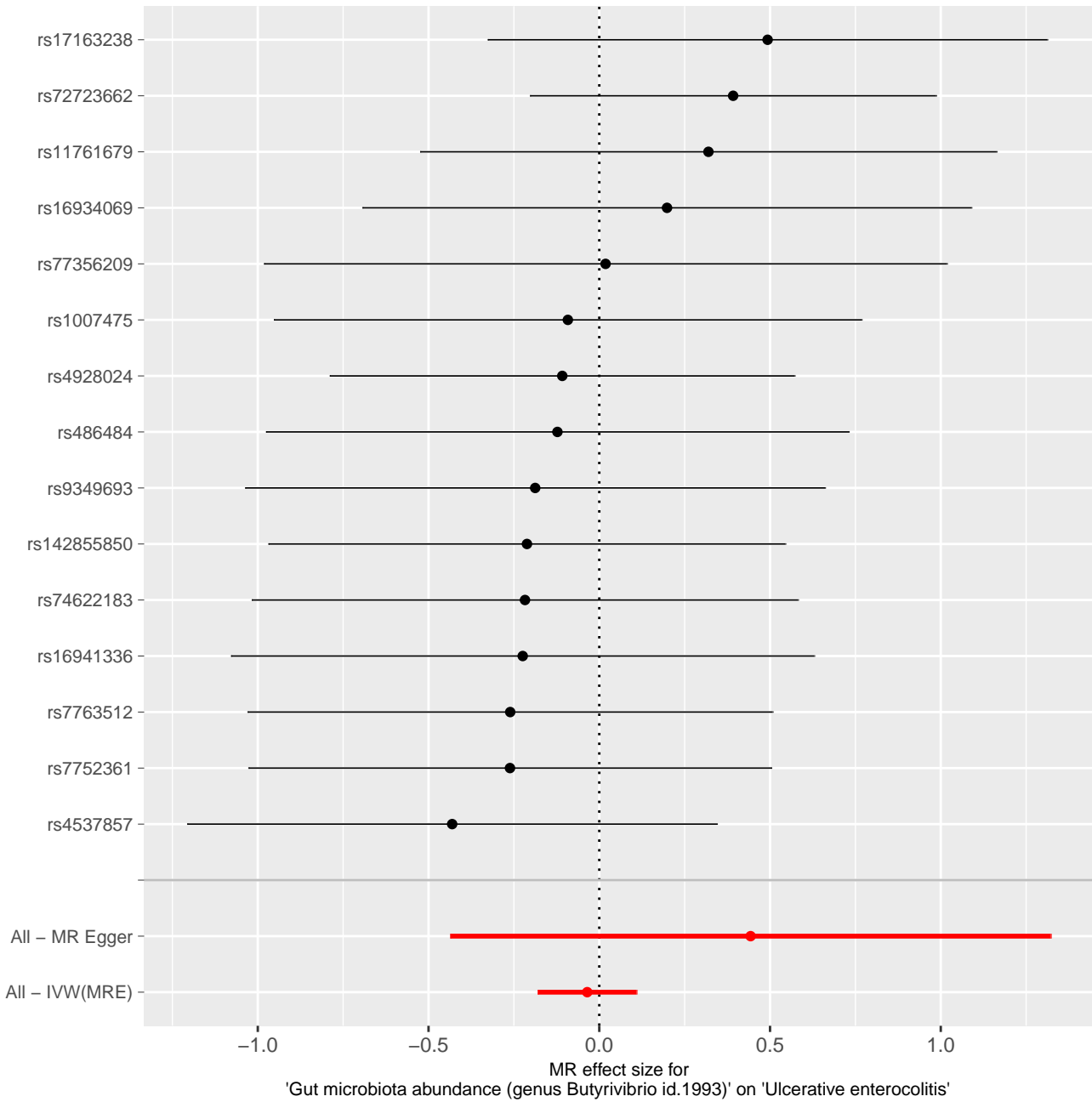


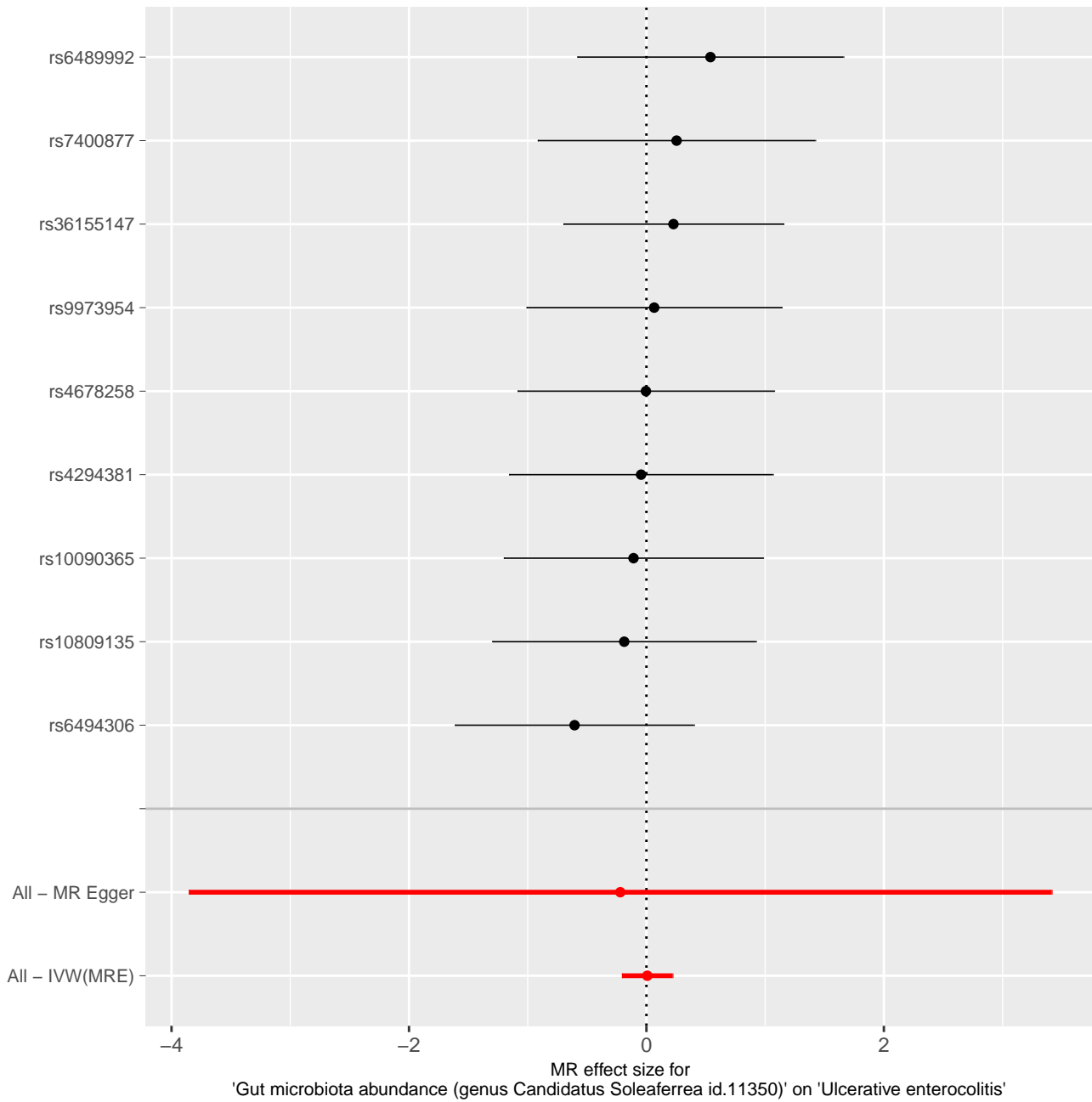


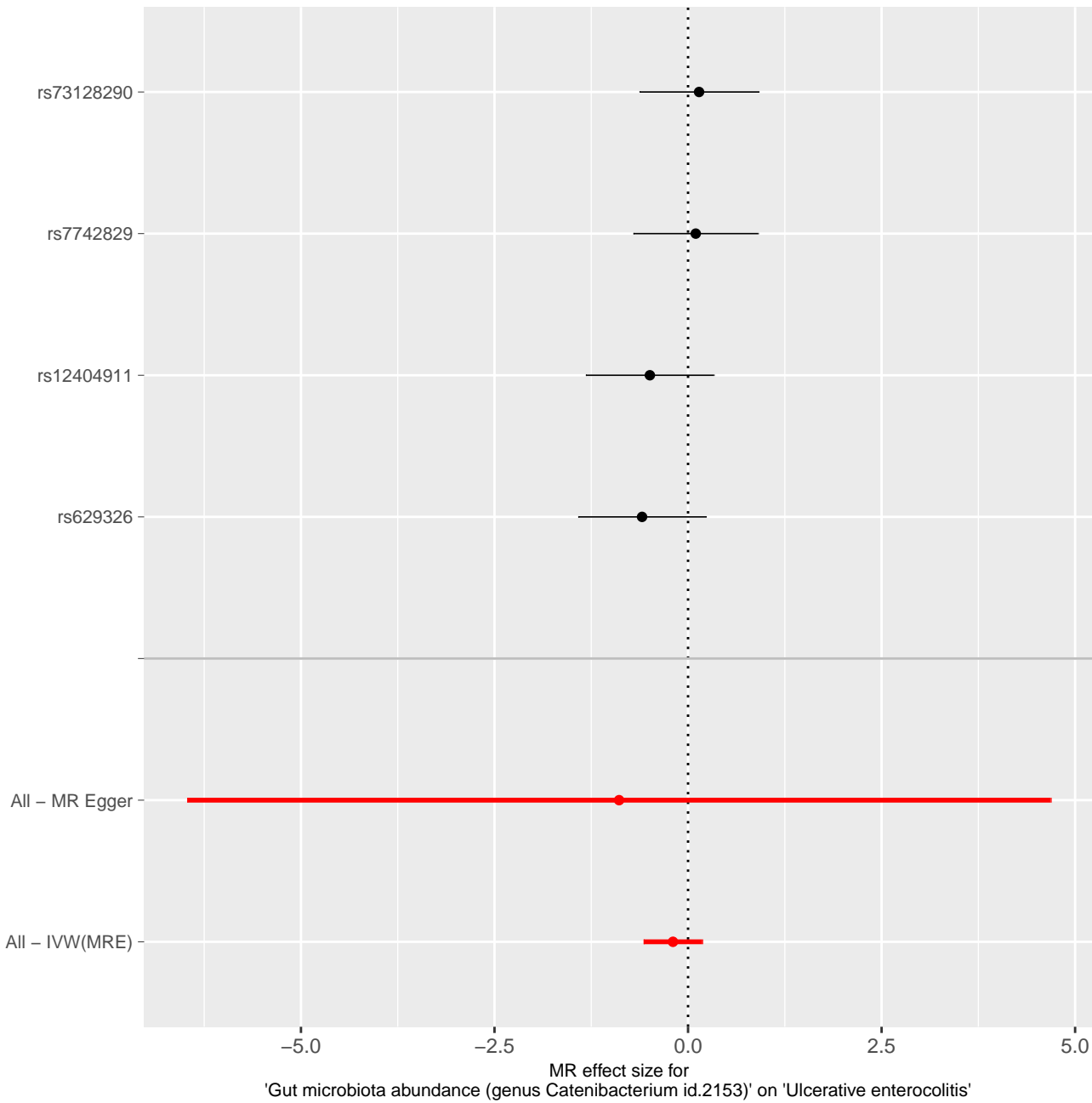
Batch 948 : Gut microbiota abundance (genus Butyricimonas id.945) on Ulcerative enterocolitis

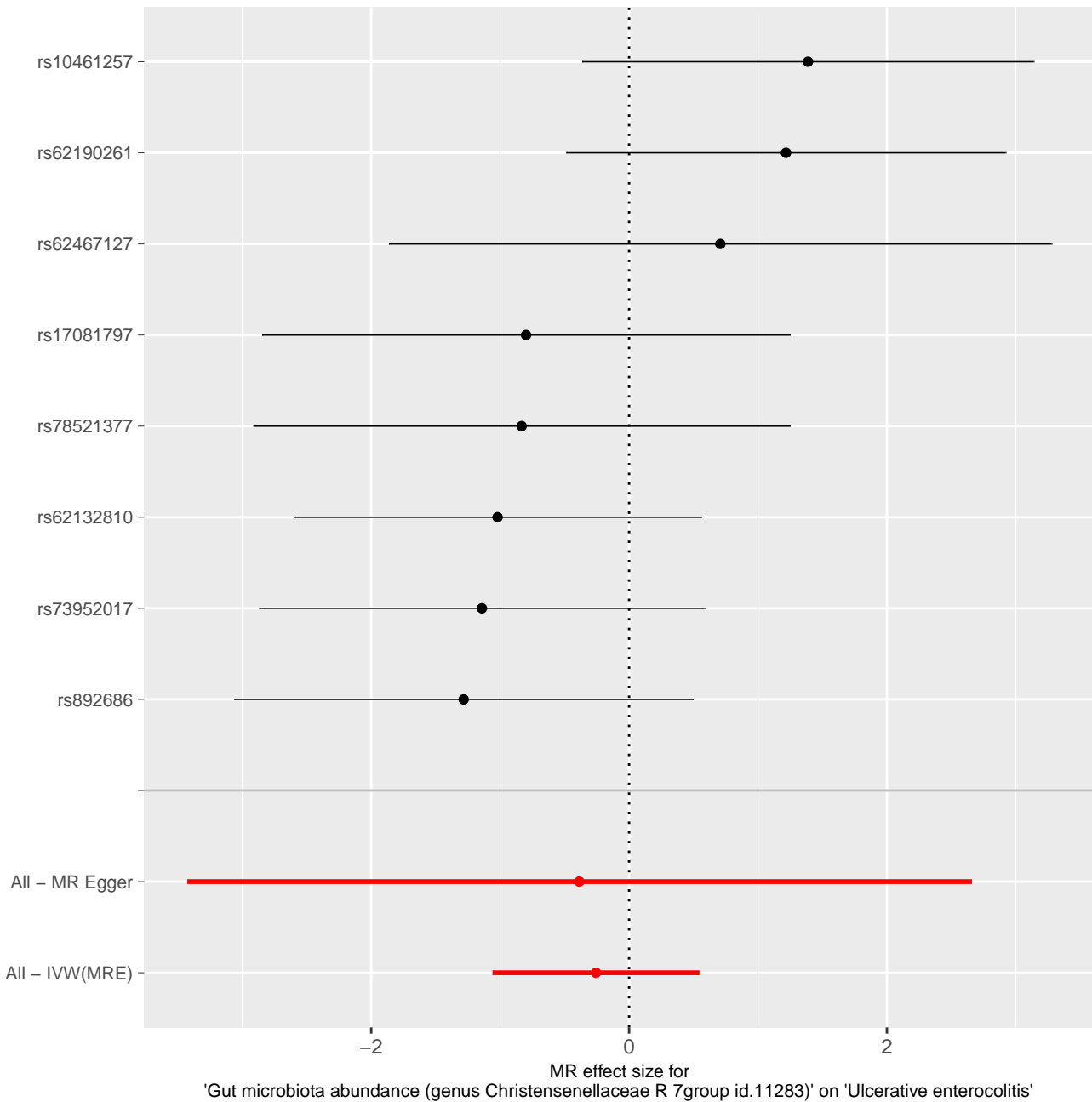


Batch 949 : Gut microbiota abundance (genus Butyrivibrio id.1993) on Ulcerative enterocolitis

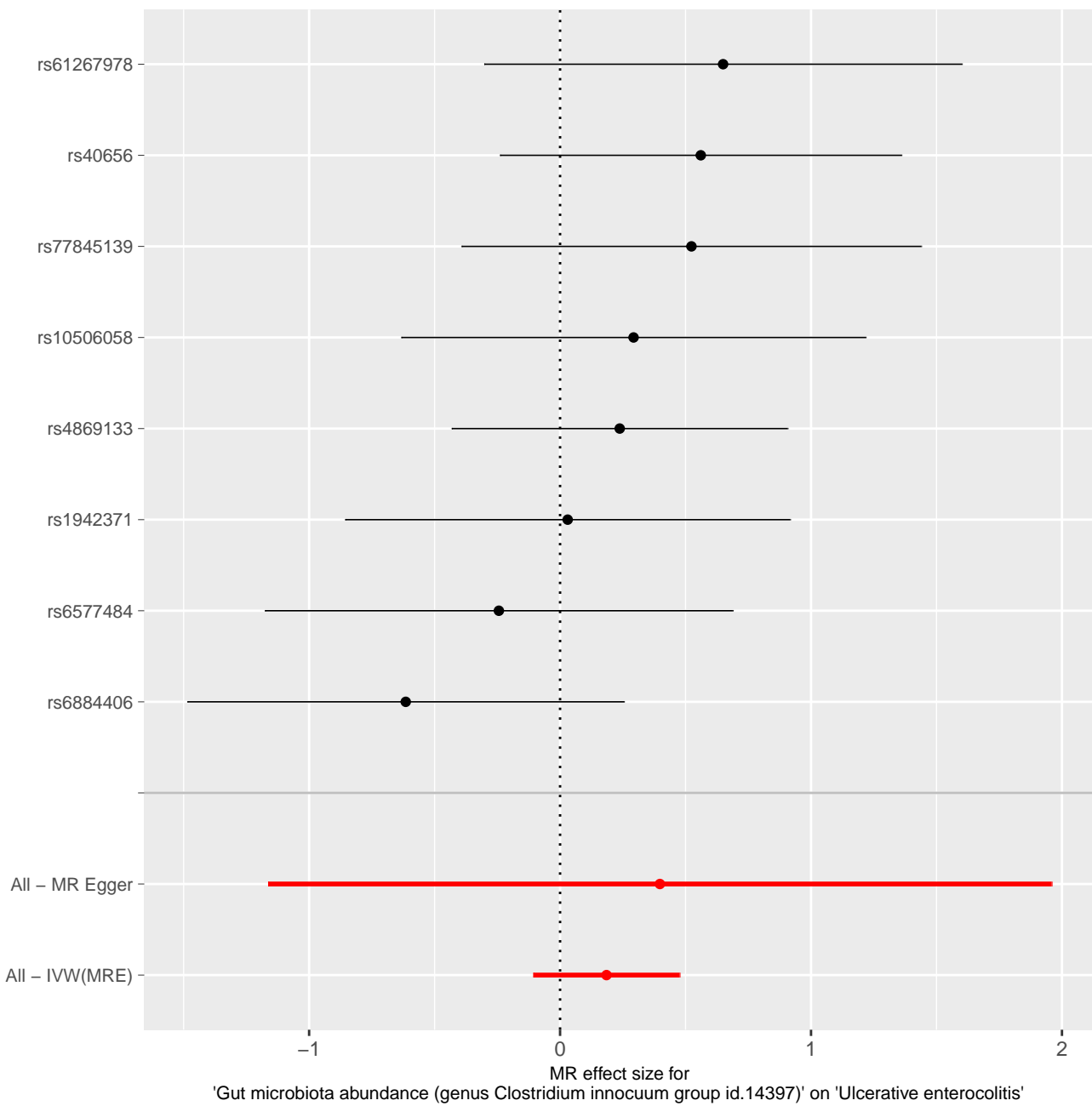


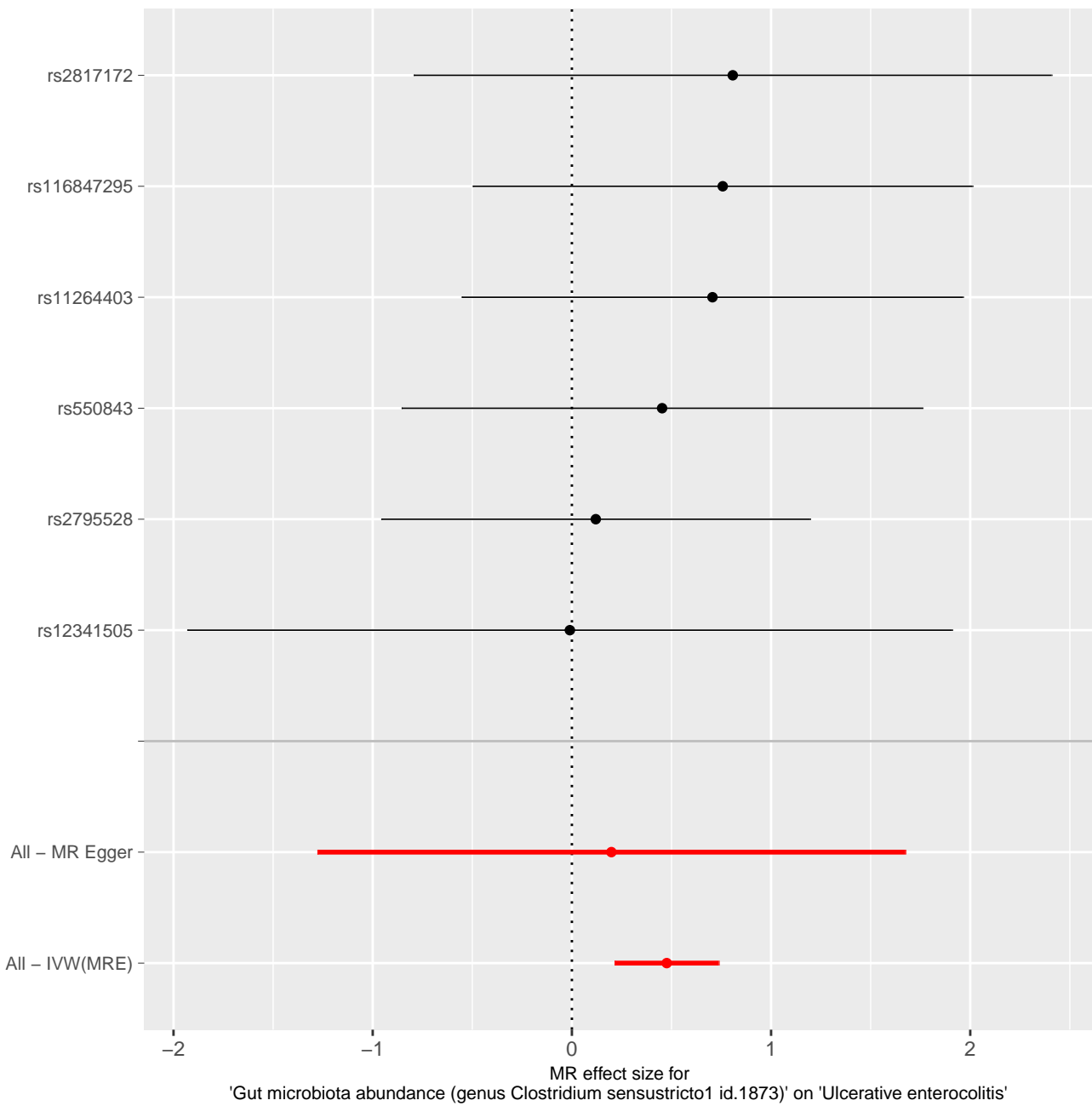


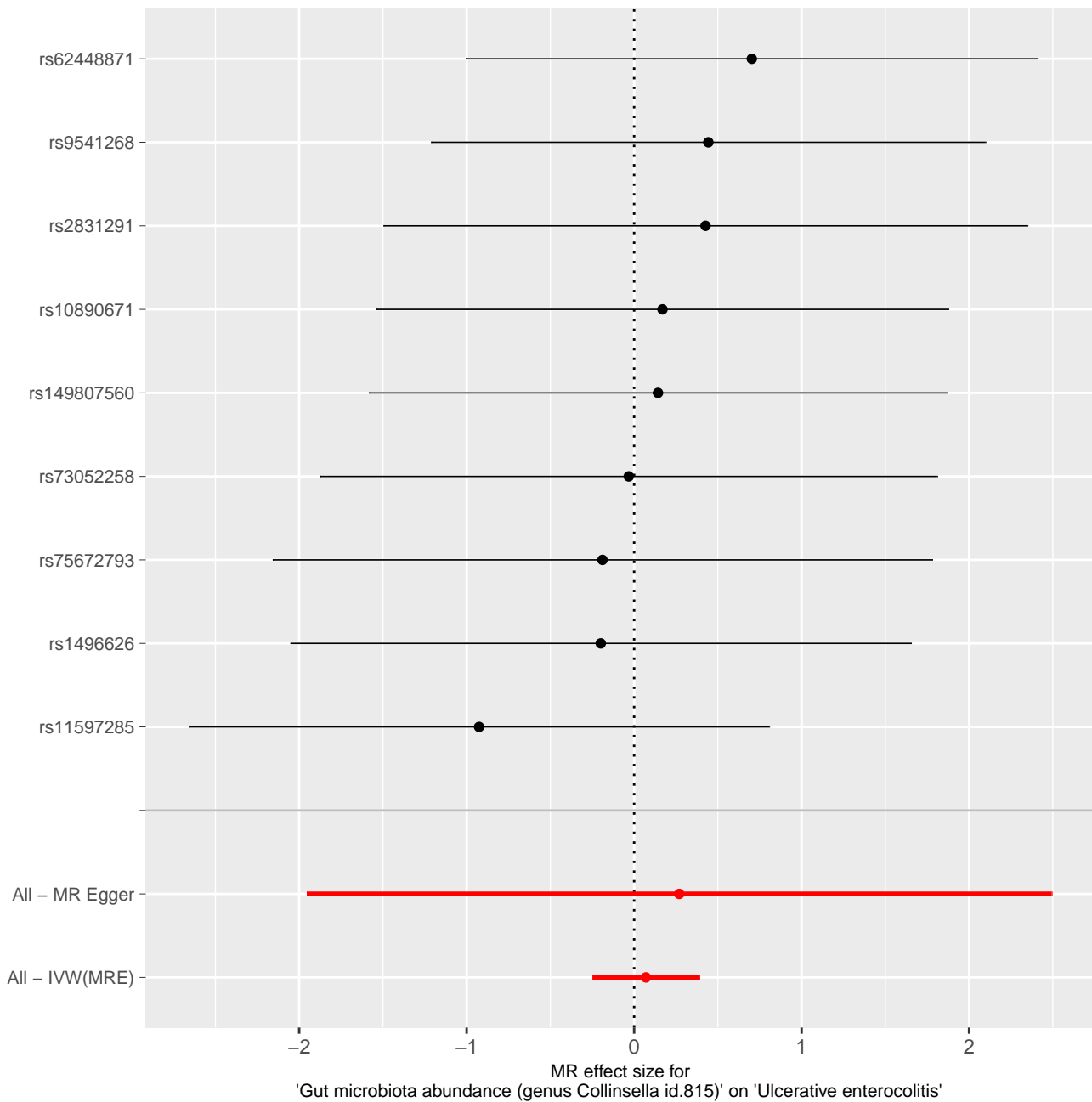




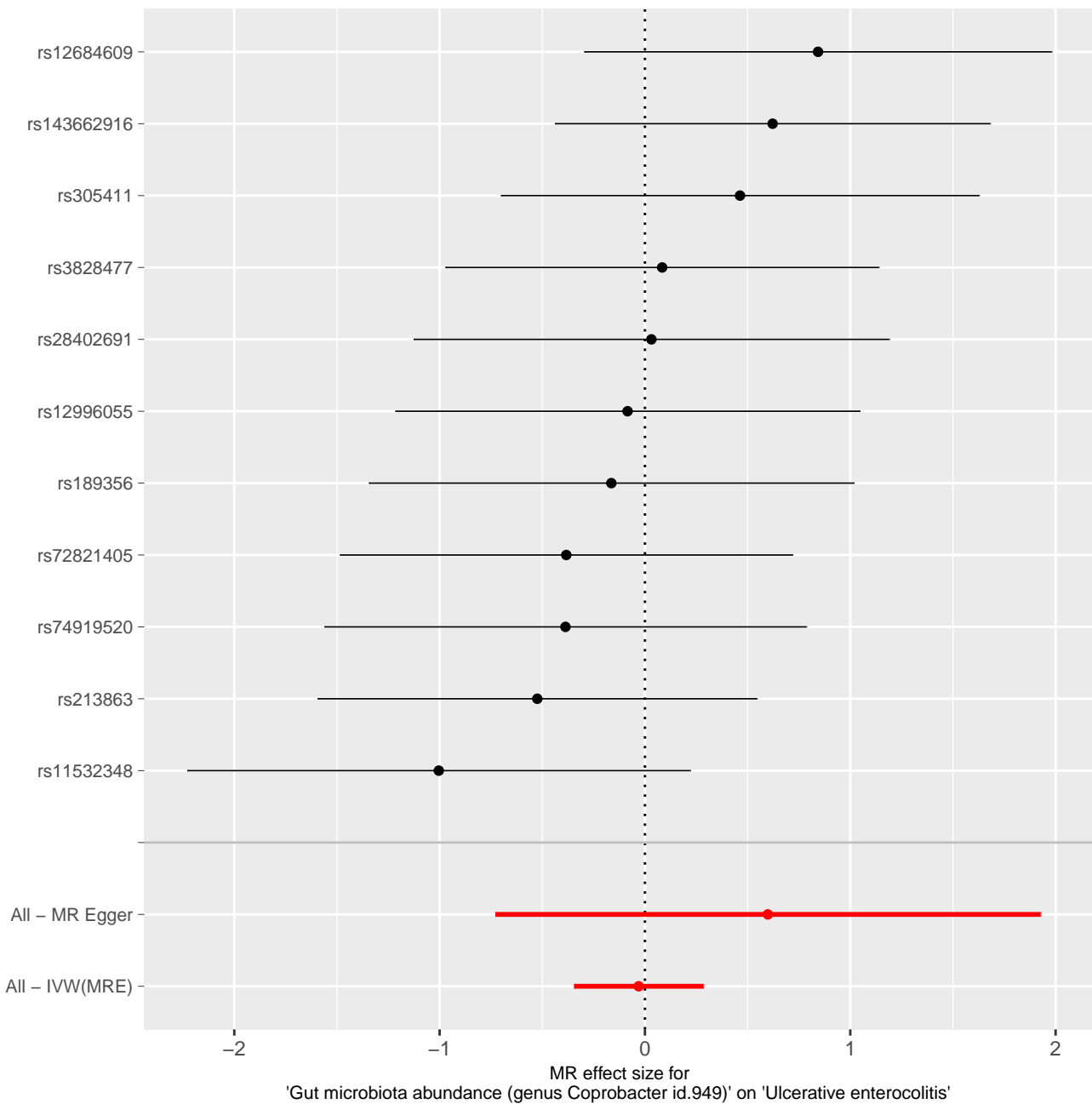
MR effect size for 'Gut microbiota abundance (genus Christensenellaceae R 7group id.11283)' on 'Ulcerative enterocolitis'

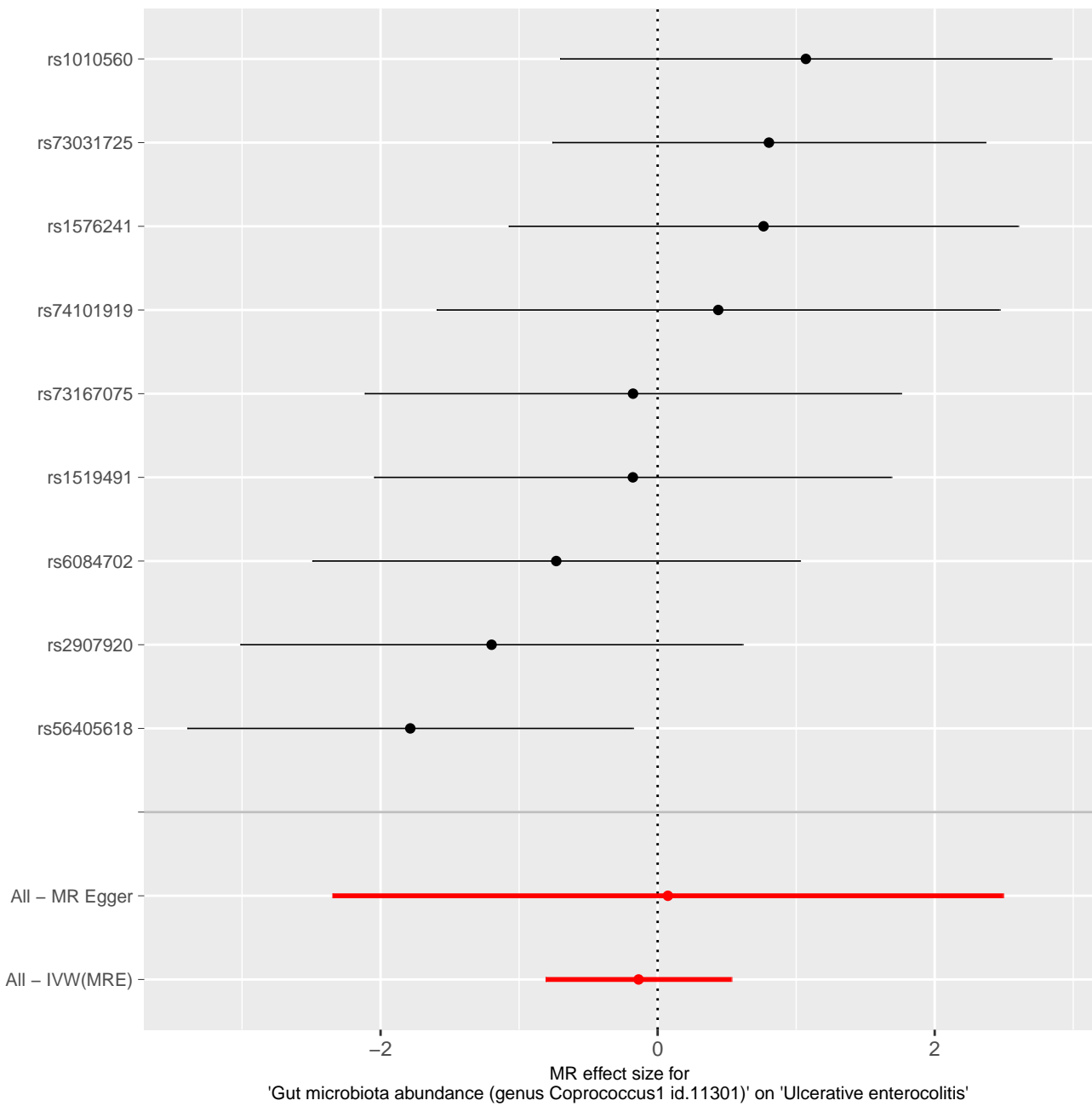


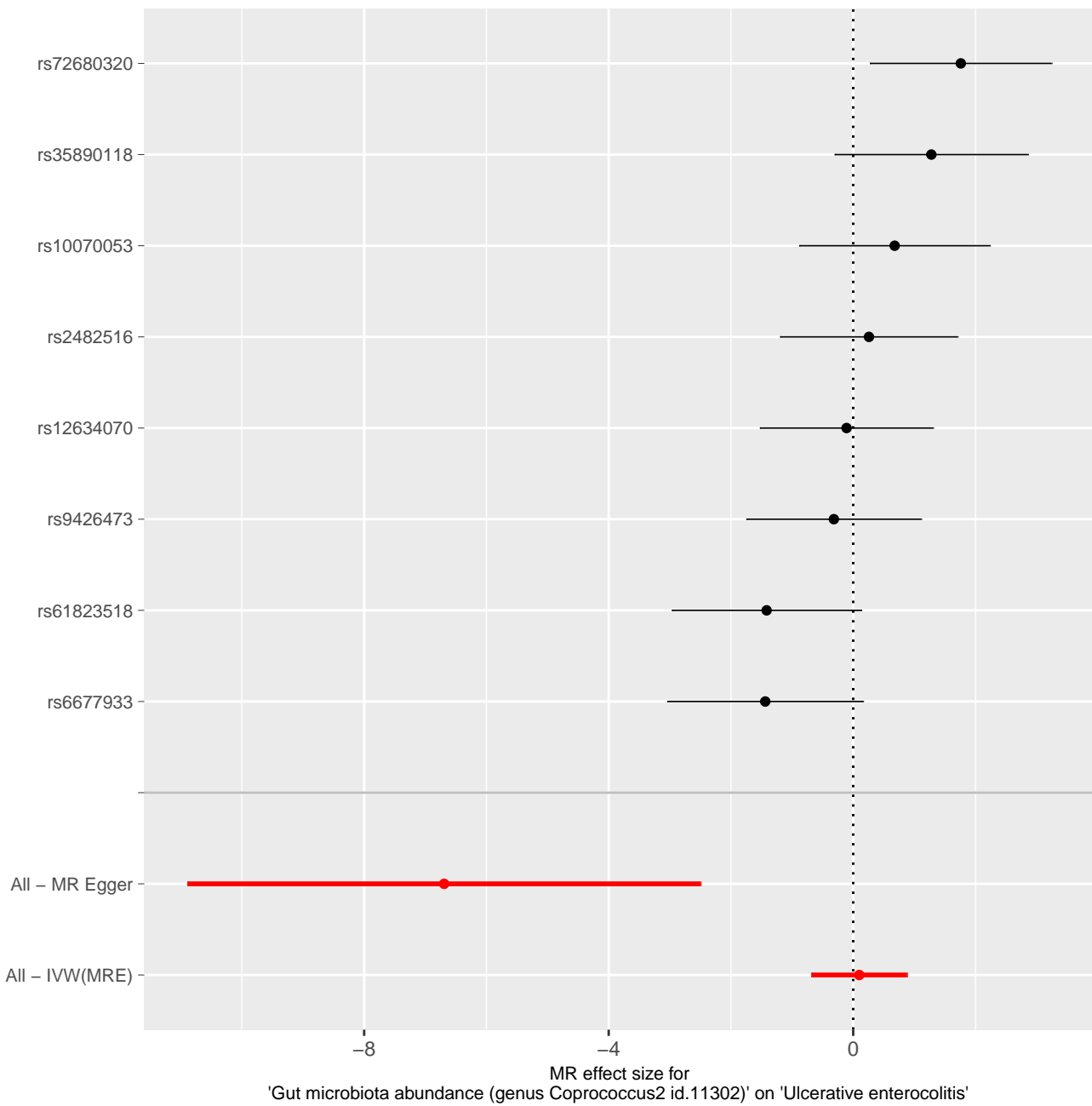


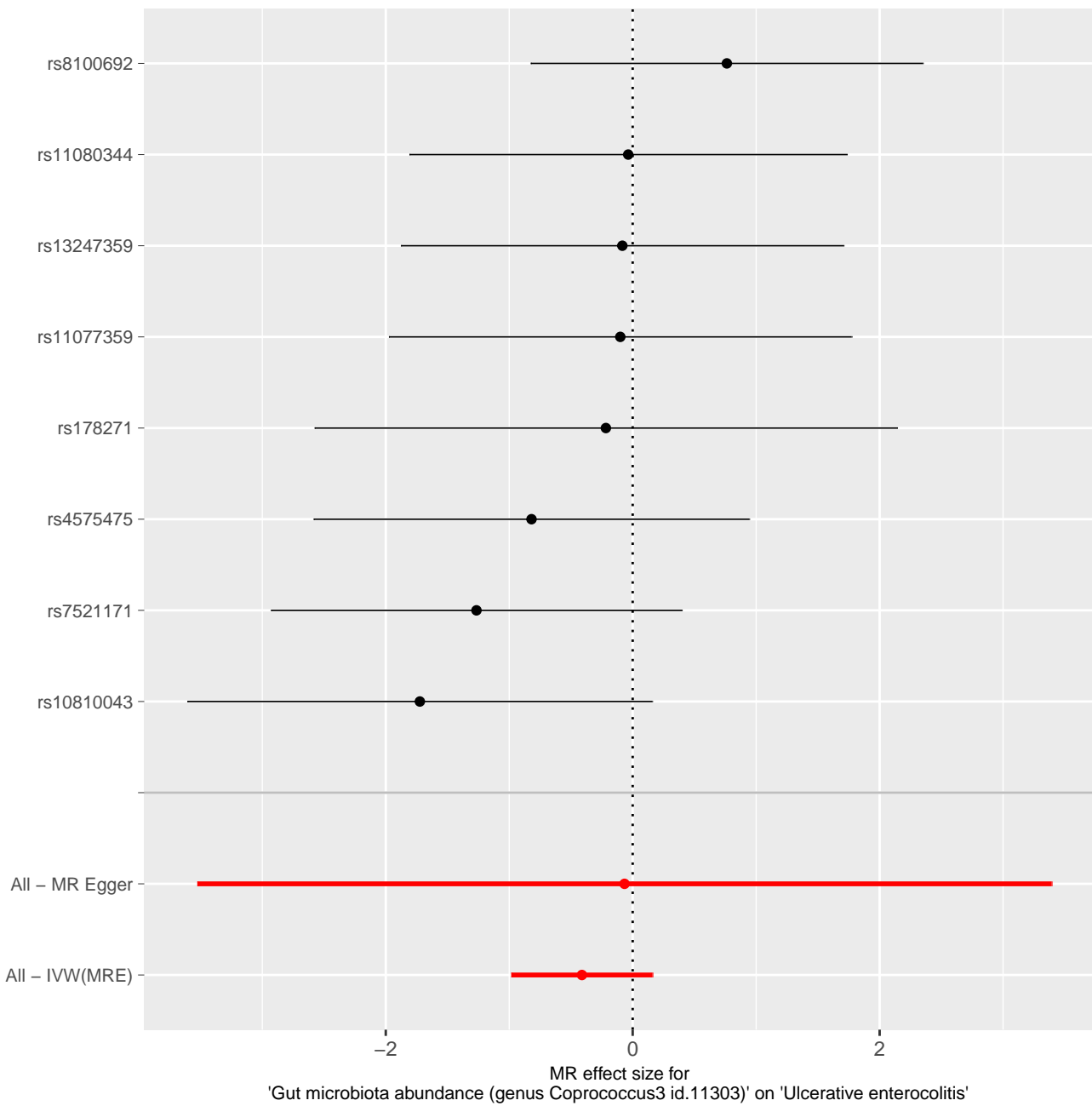


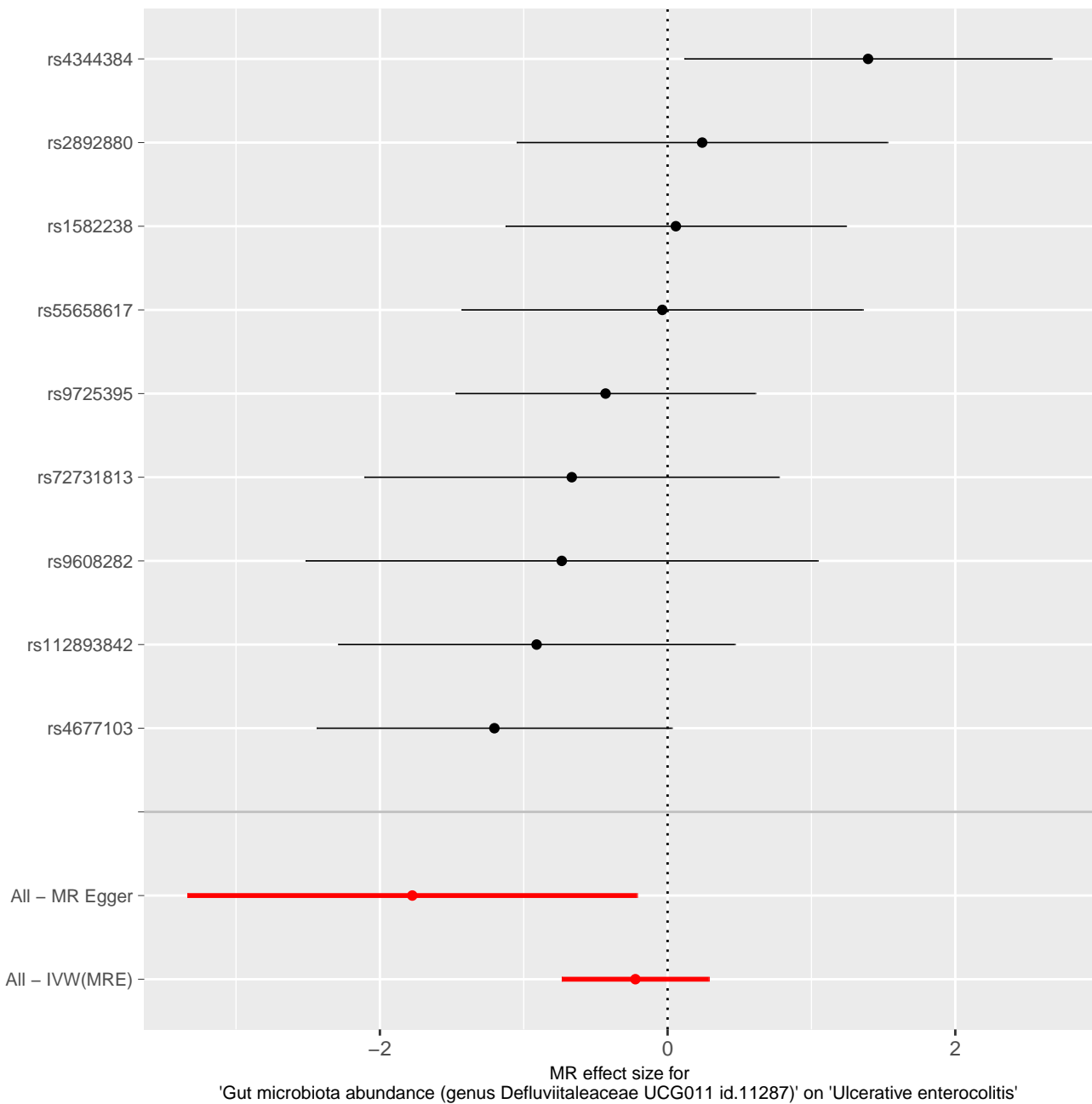
Batch 956 : Gut microbiota abundance (genus Coprobacter id.949) on Ulcerative enterocolitis

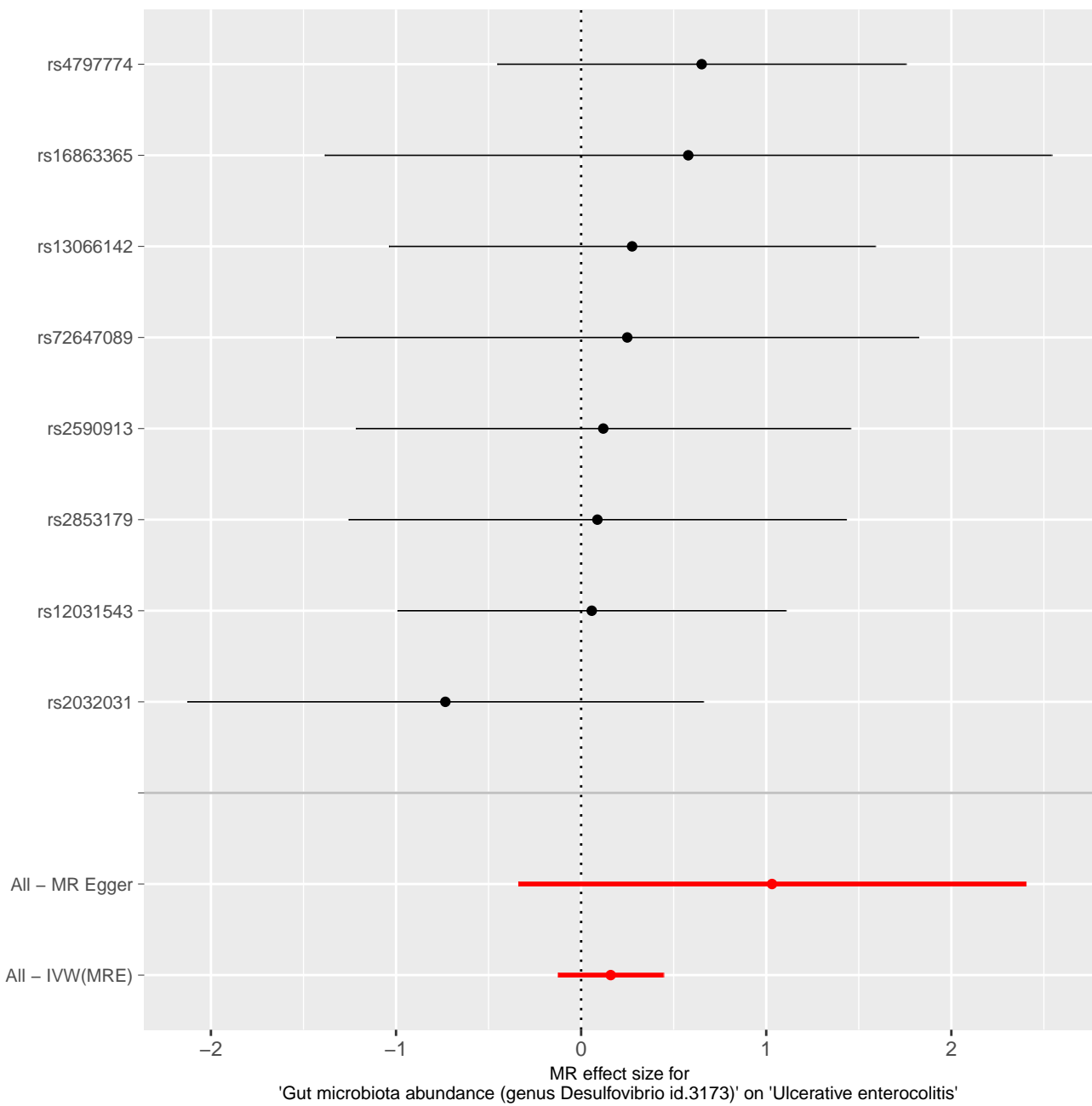


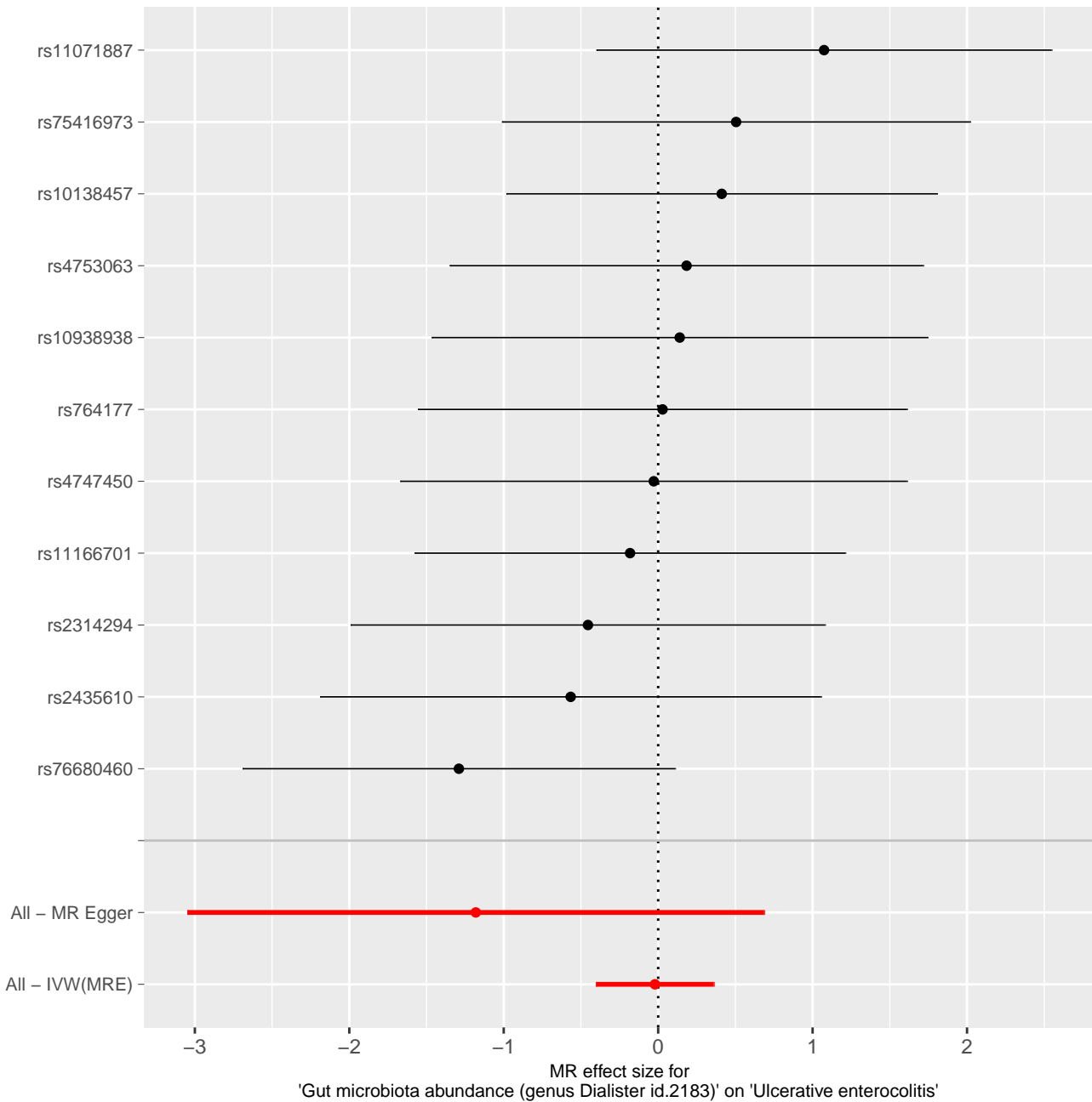




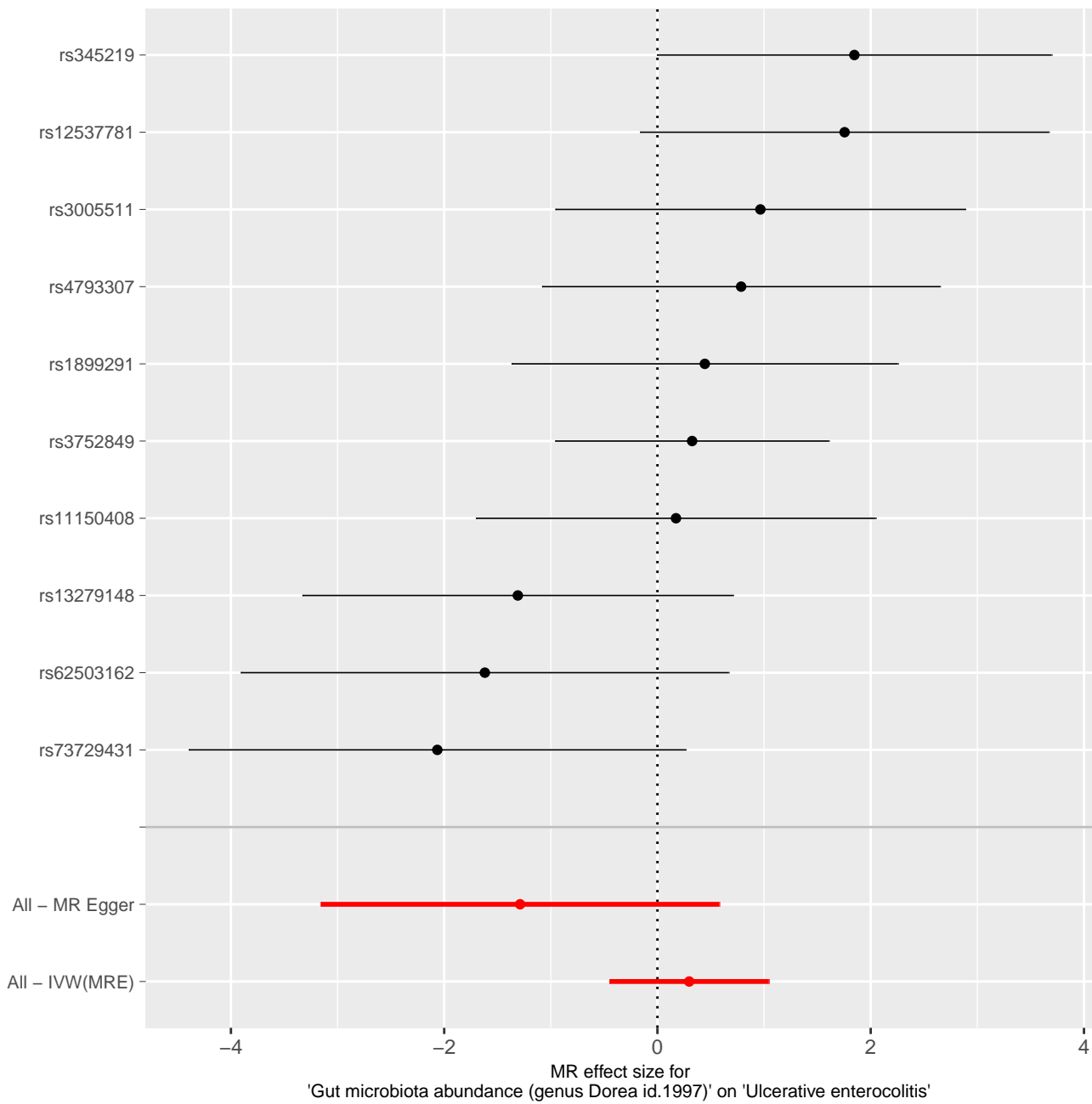




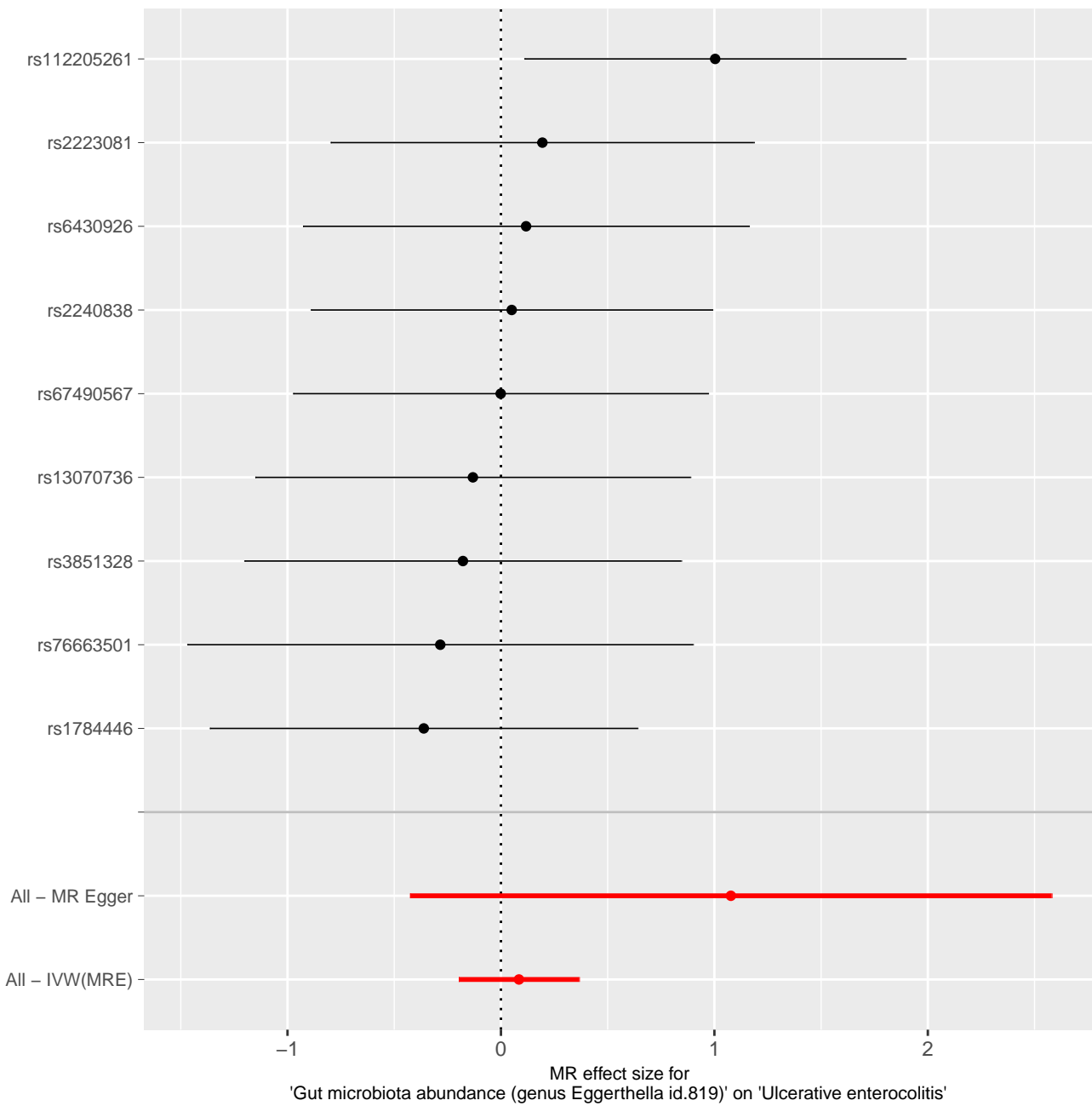


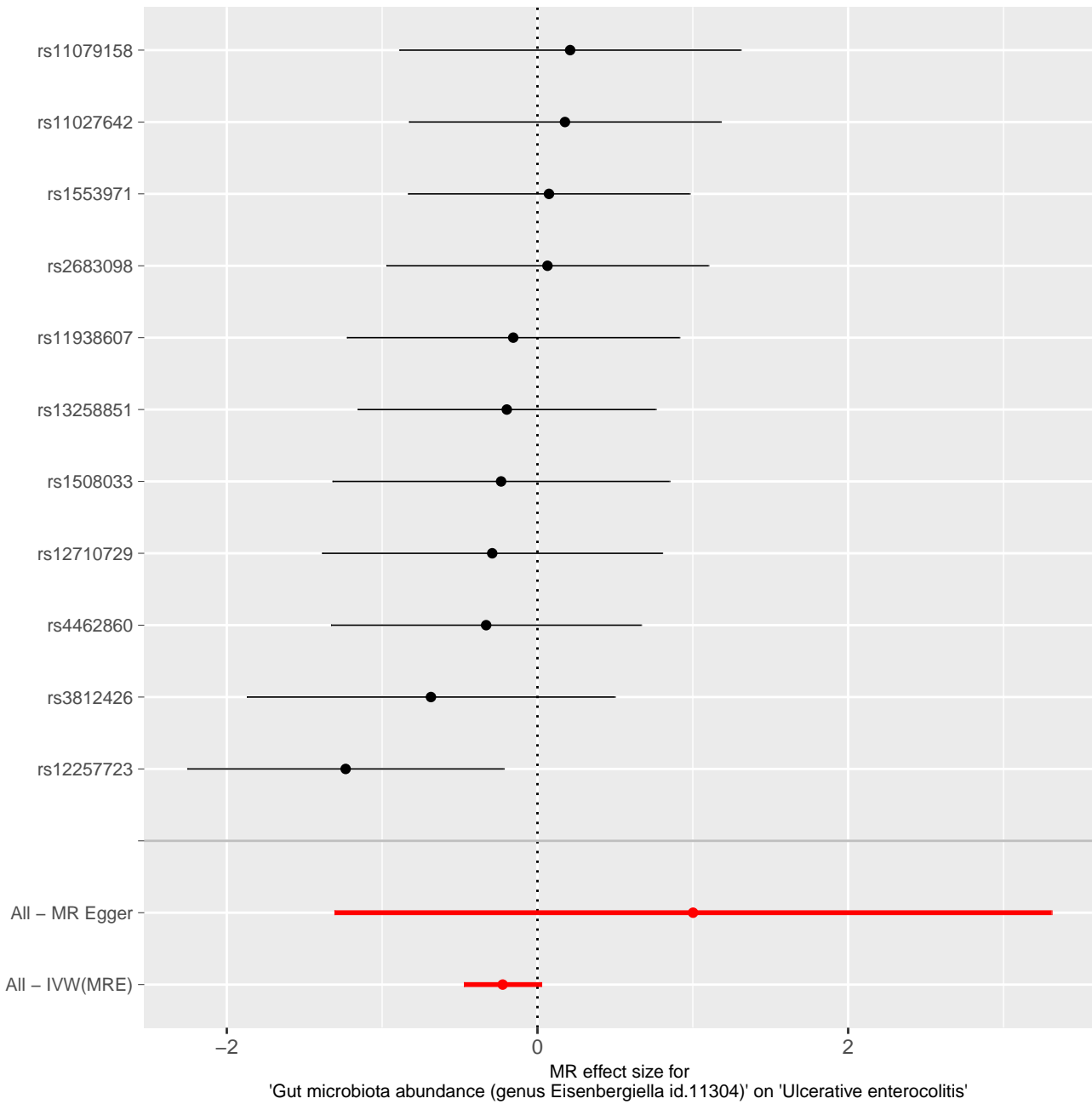


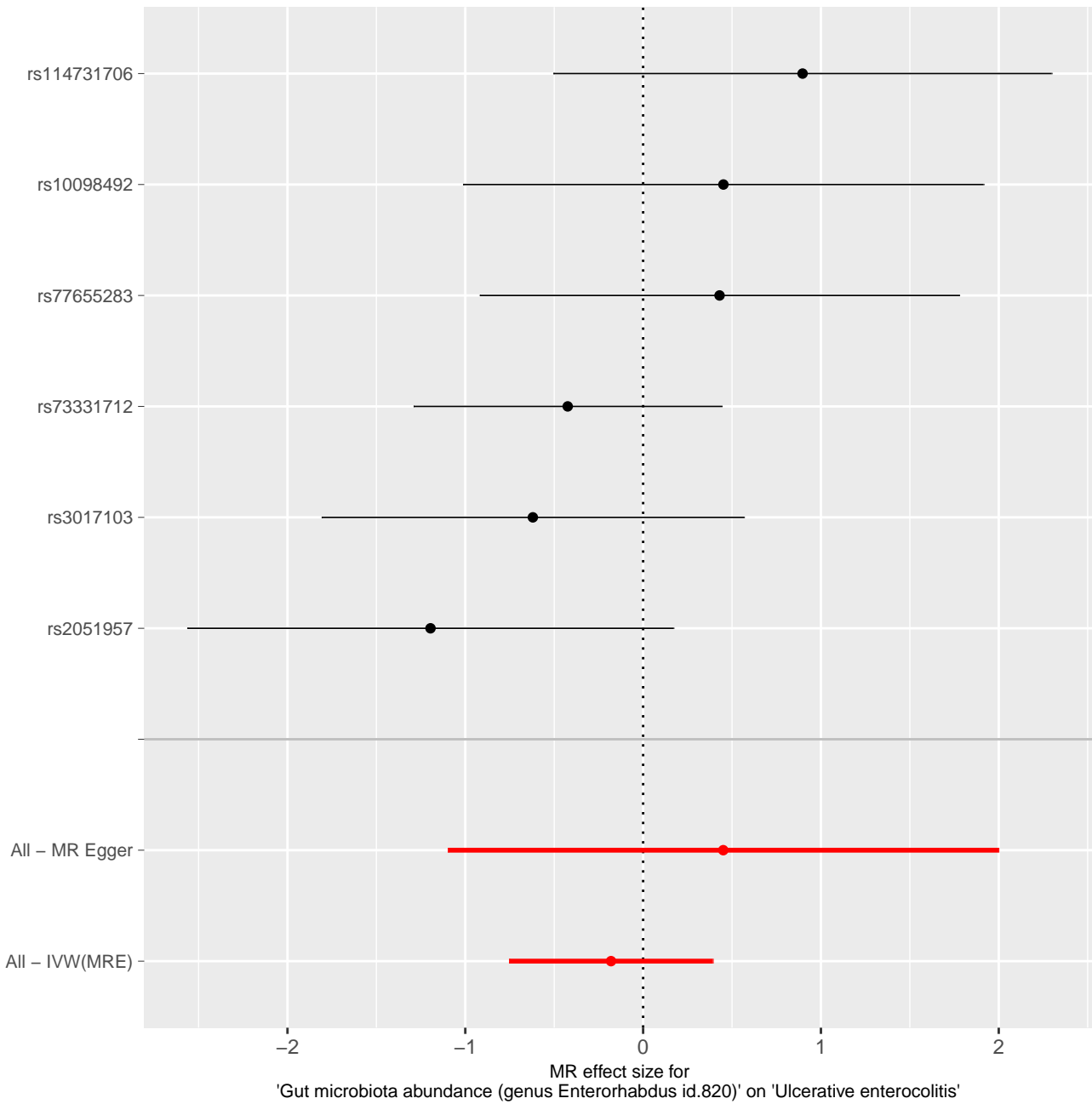
Batch 963 : Gut microbiota abundance (genus Dorea id.1997) on Ulcerative enterocolitis



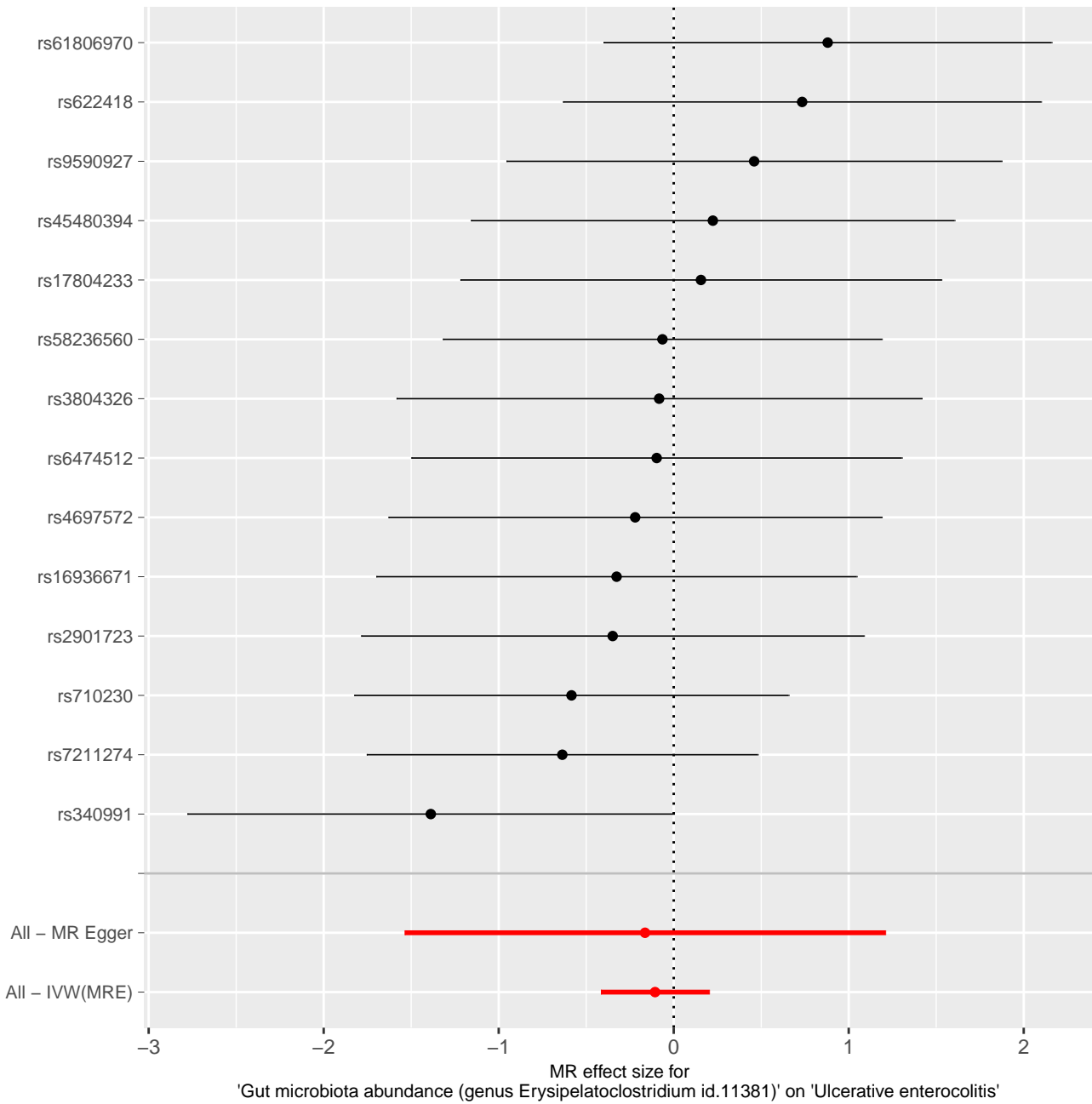
Batch 964 : Gut microbiota abundance (genus Eggerthella id.819) on Ulcerative enterocolitis

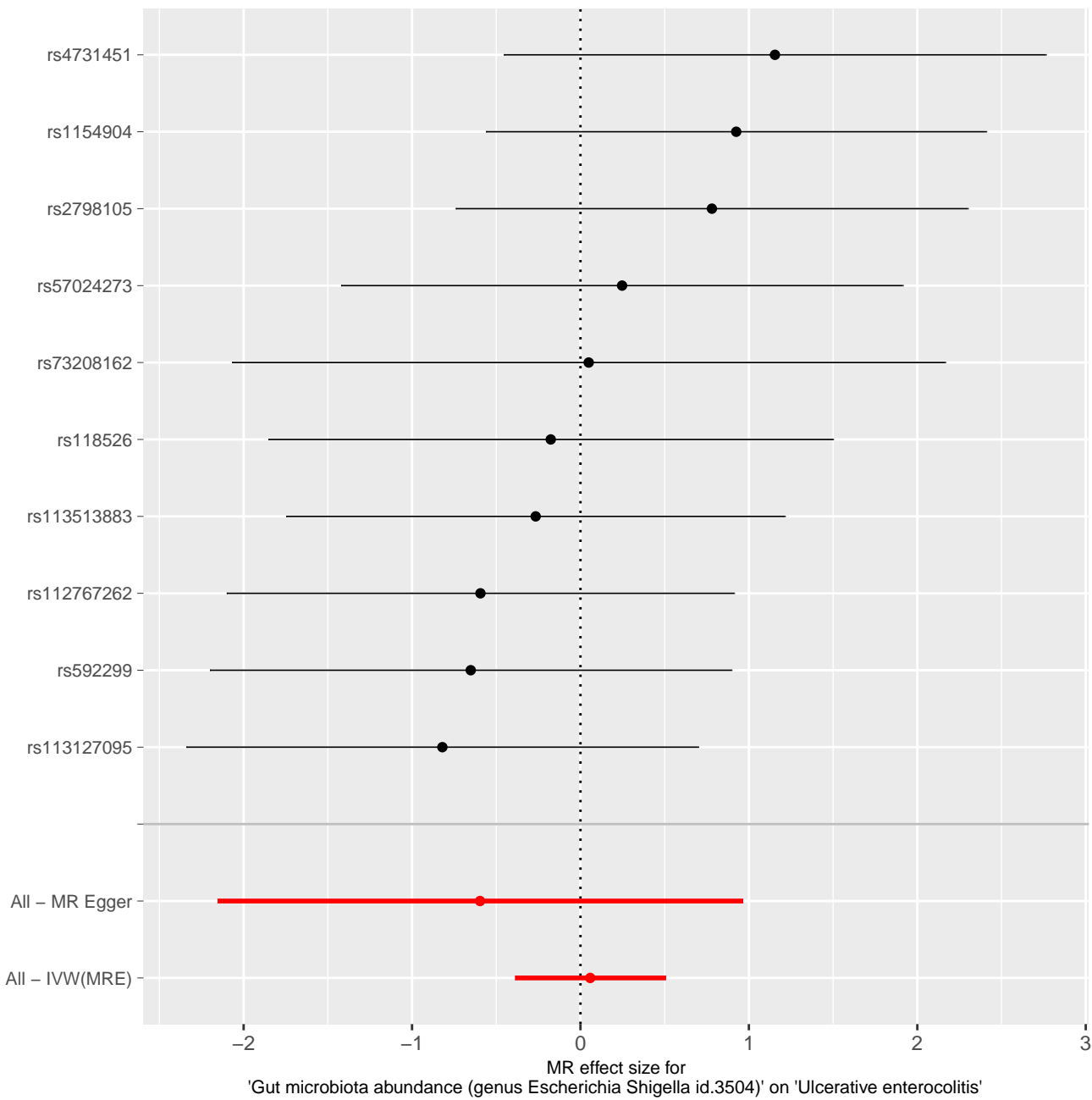




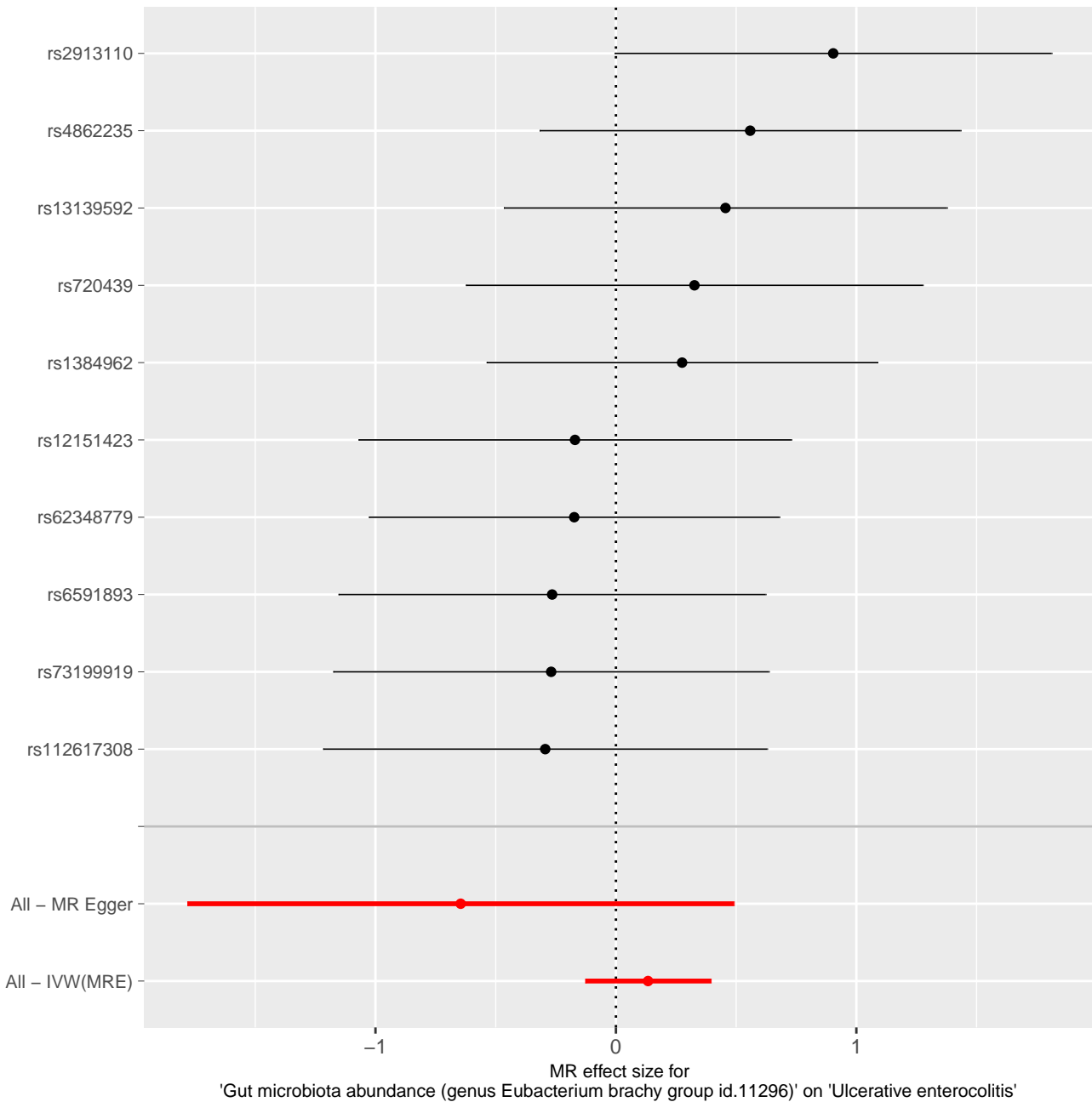


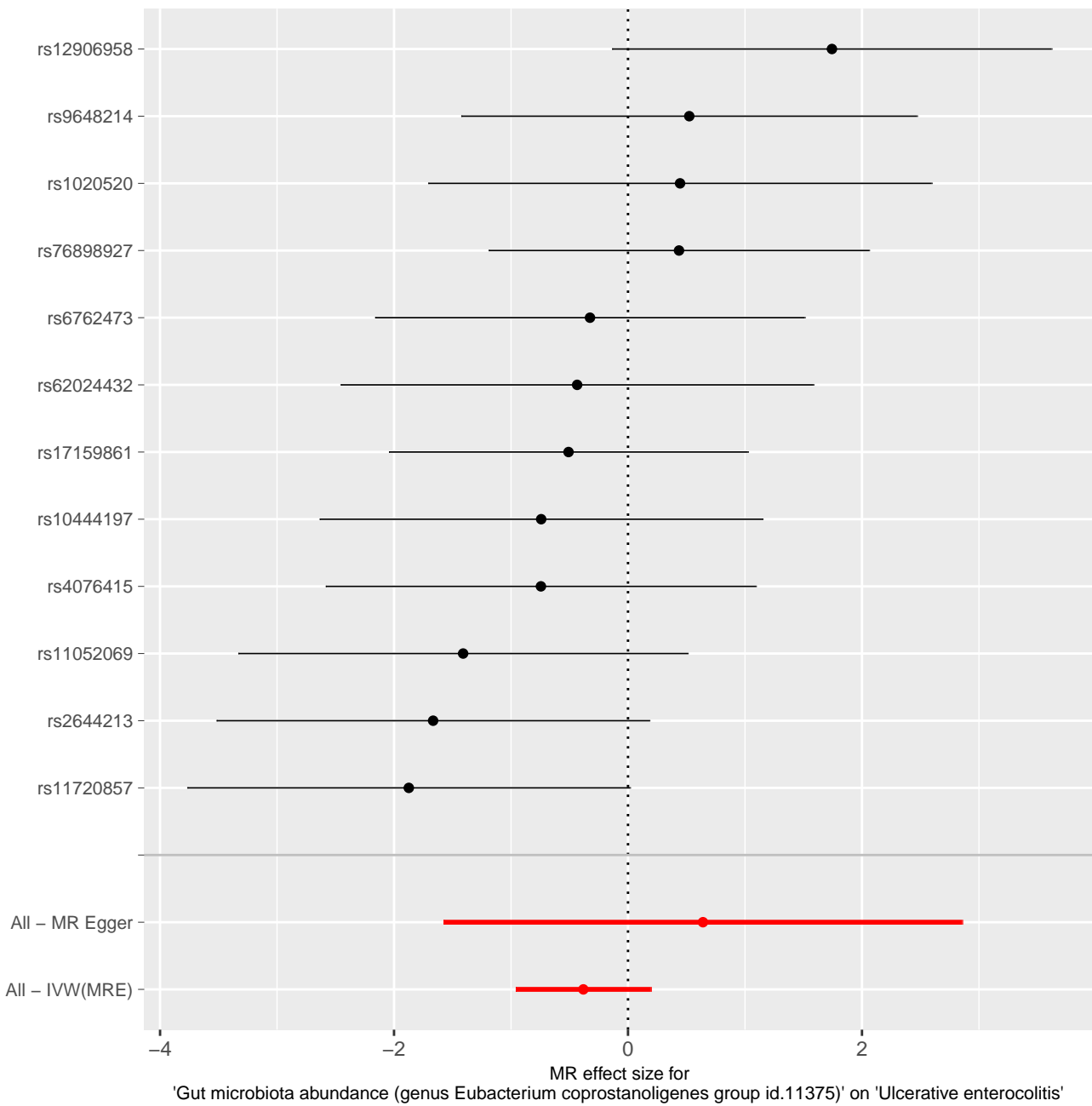
Batch 967 : Gut microbiota abundance (genus Erysipelatoclostridium id.11381) on Ulcerative enterocolitis

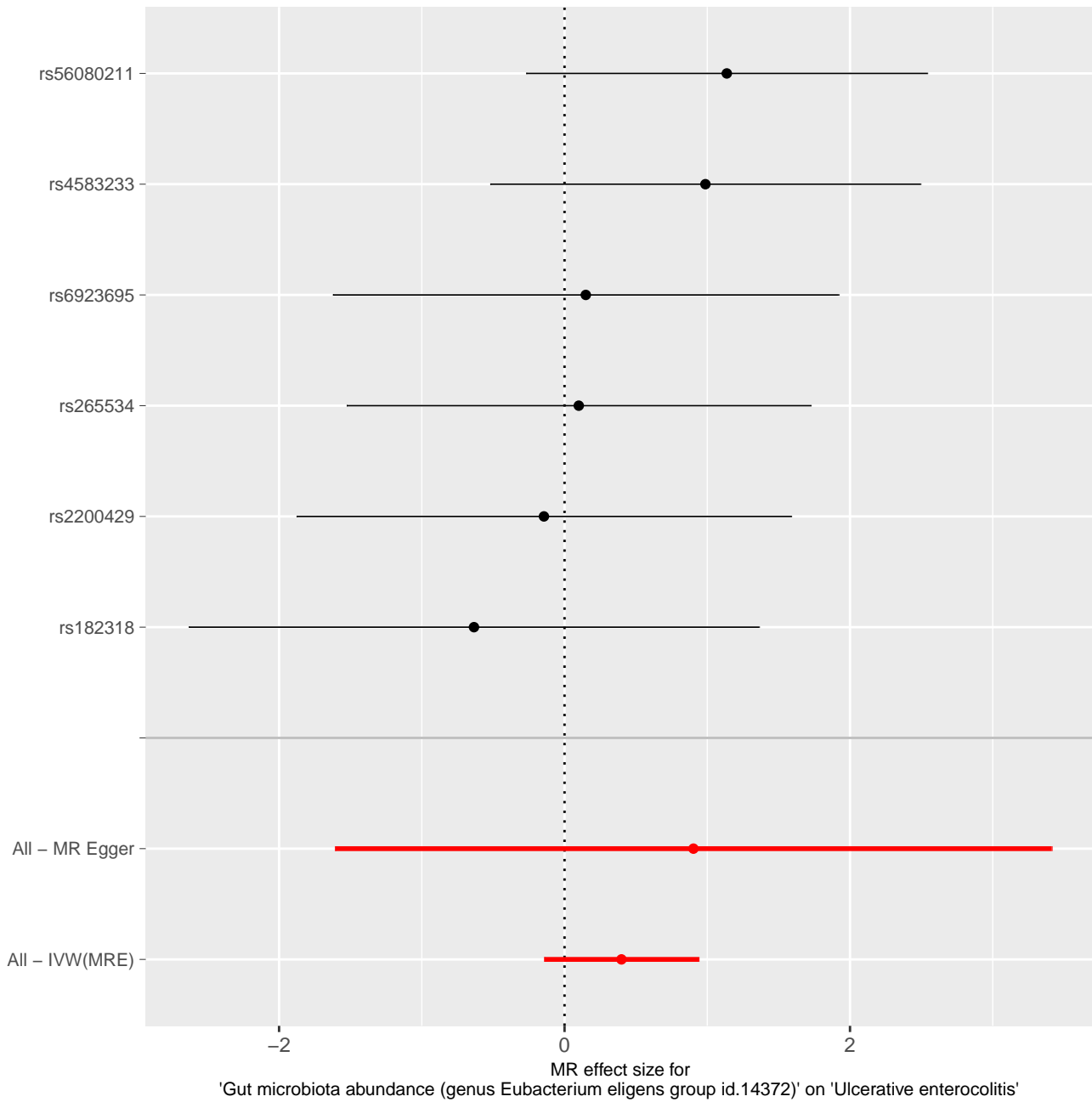


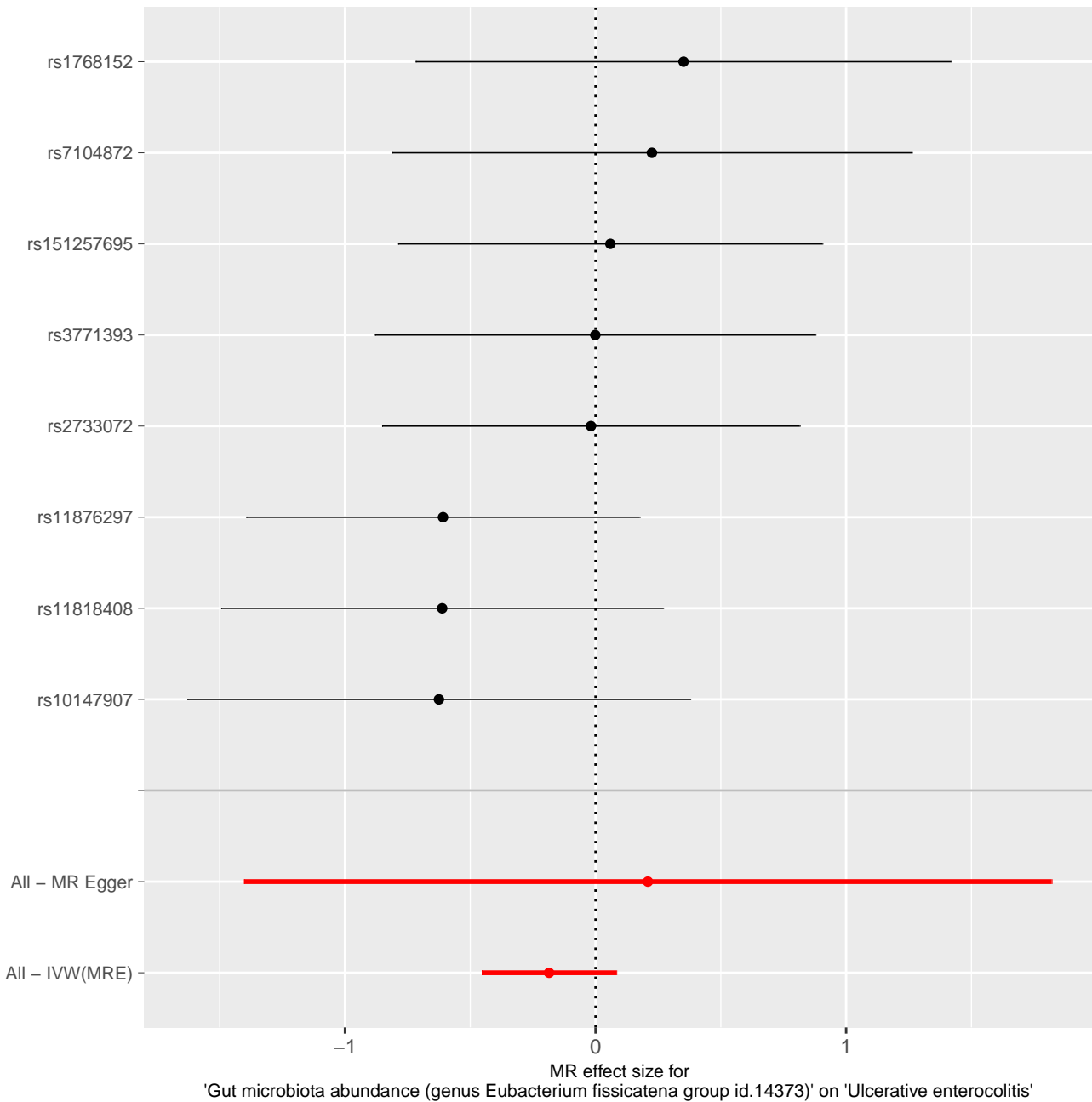


Batch 970 : Gut microbiota abundance (genus Eubacterium brachy group id.11296) on Ulcerative enterocolitis

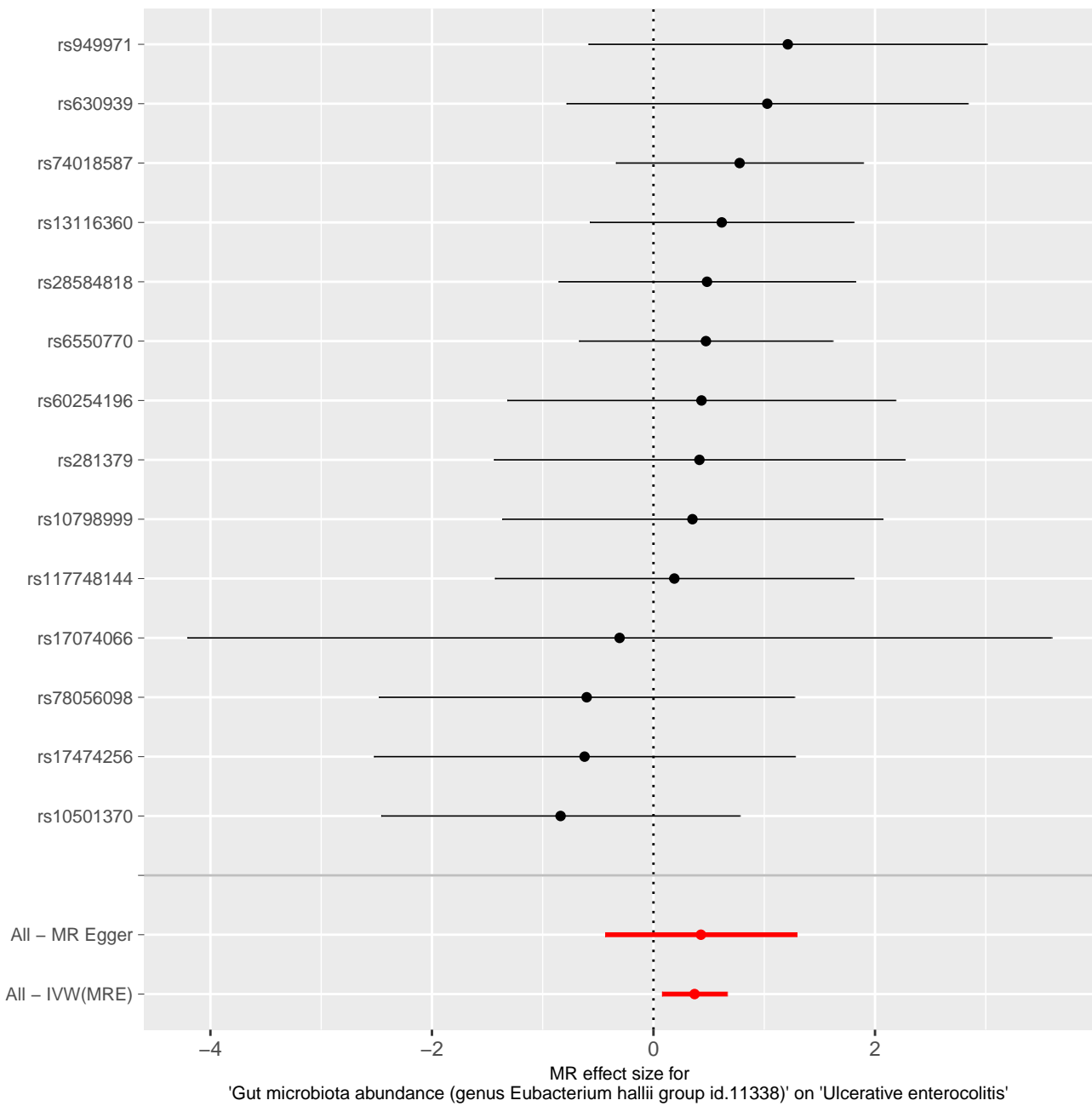


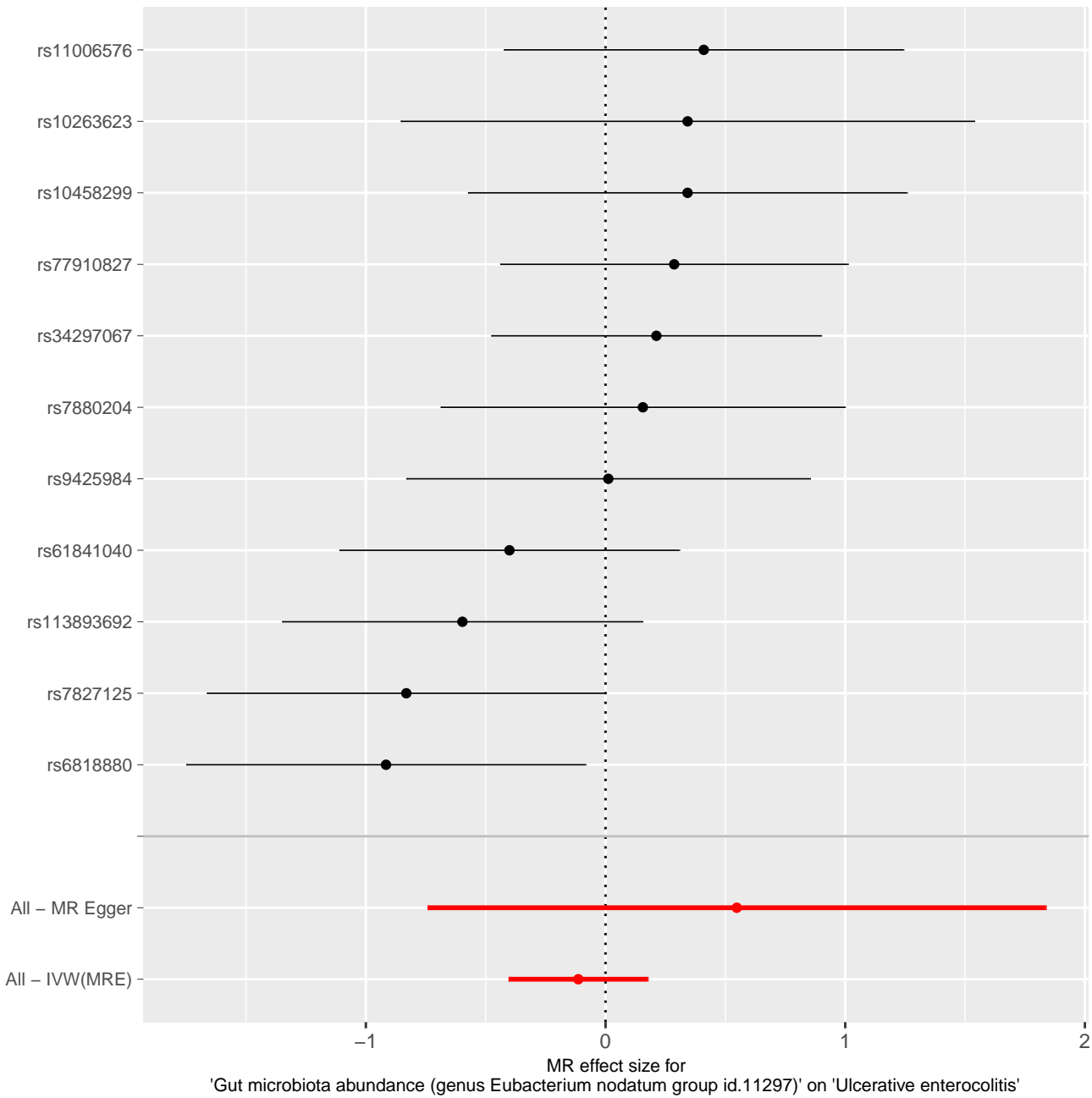


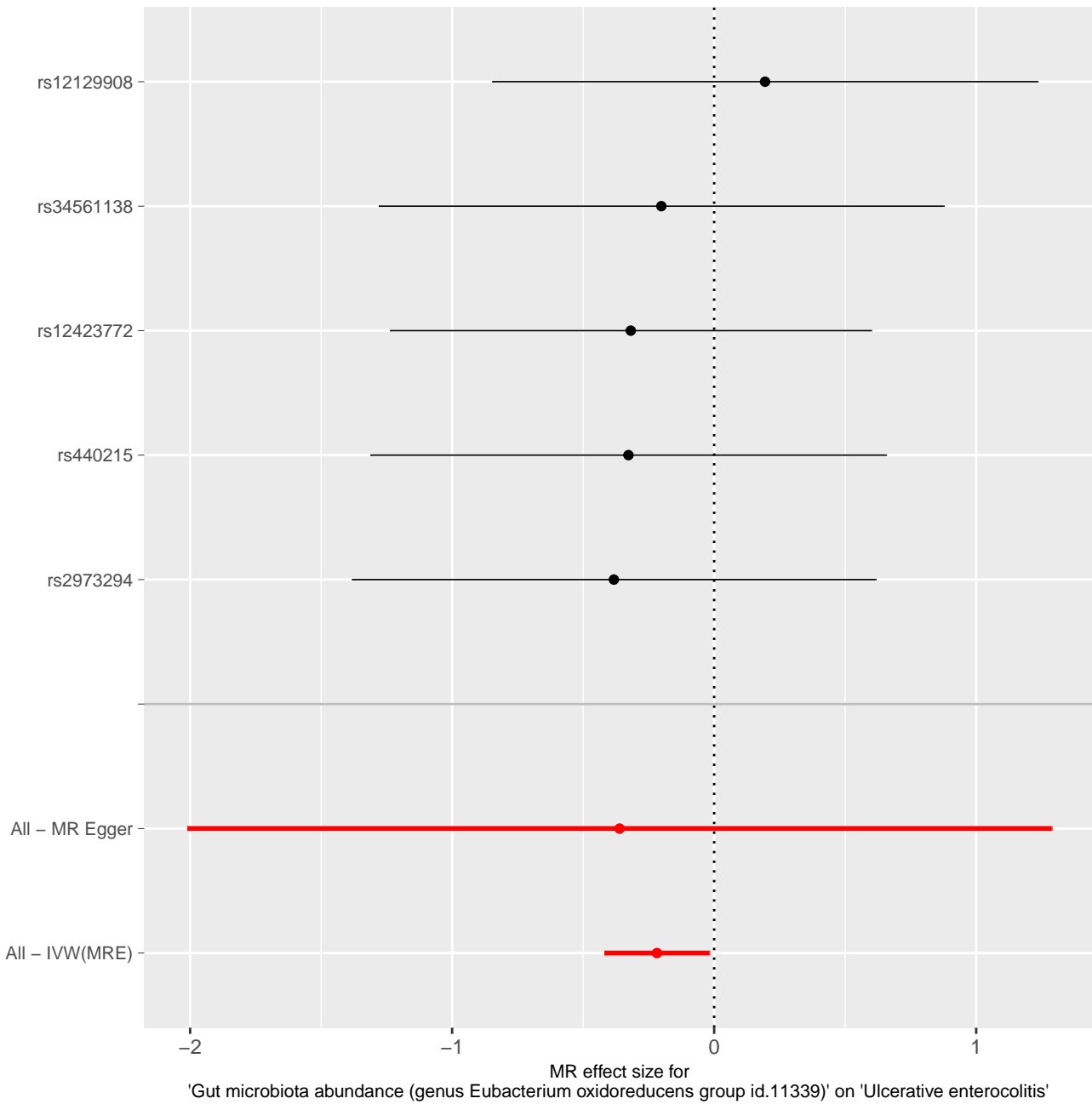


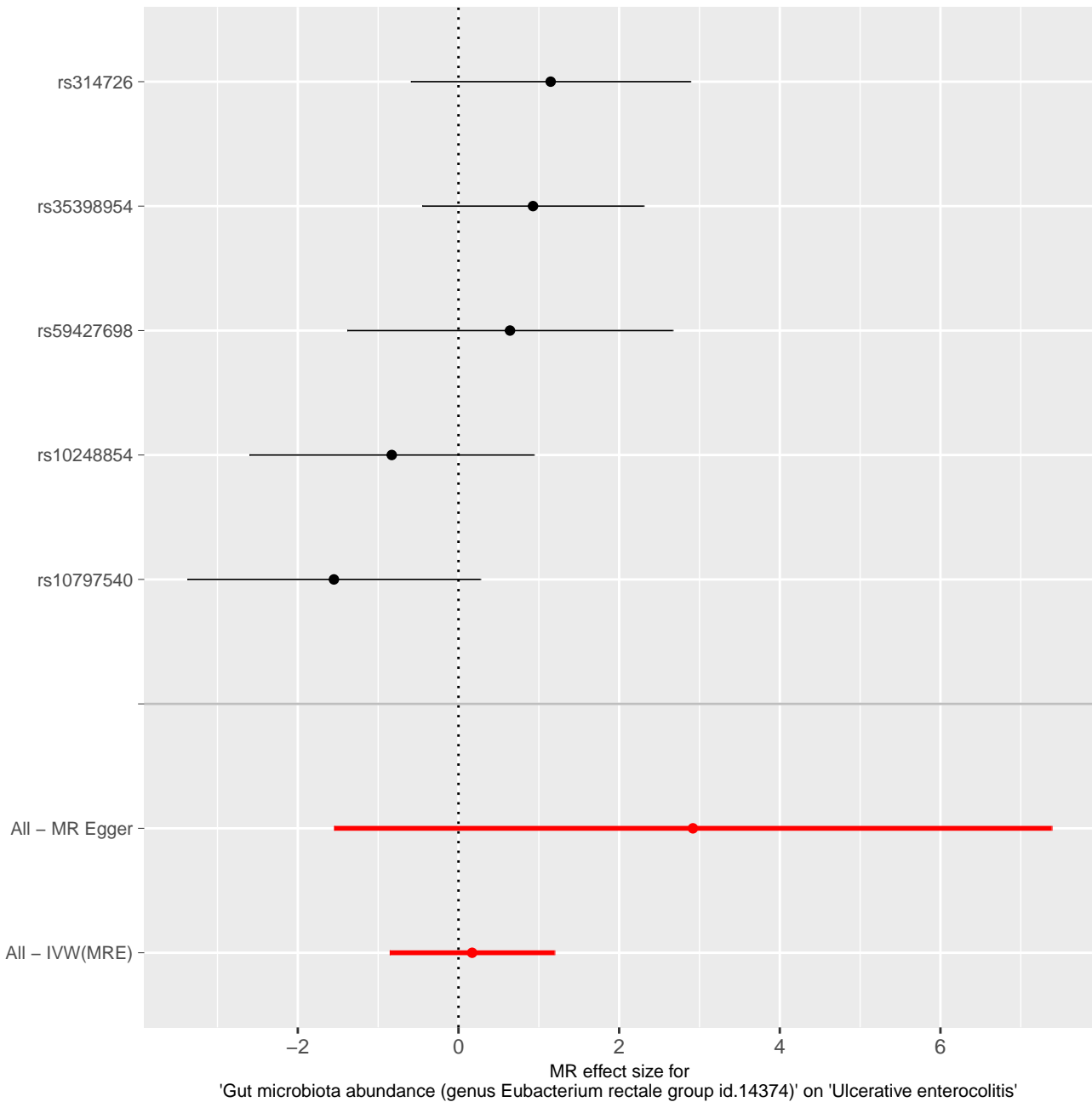


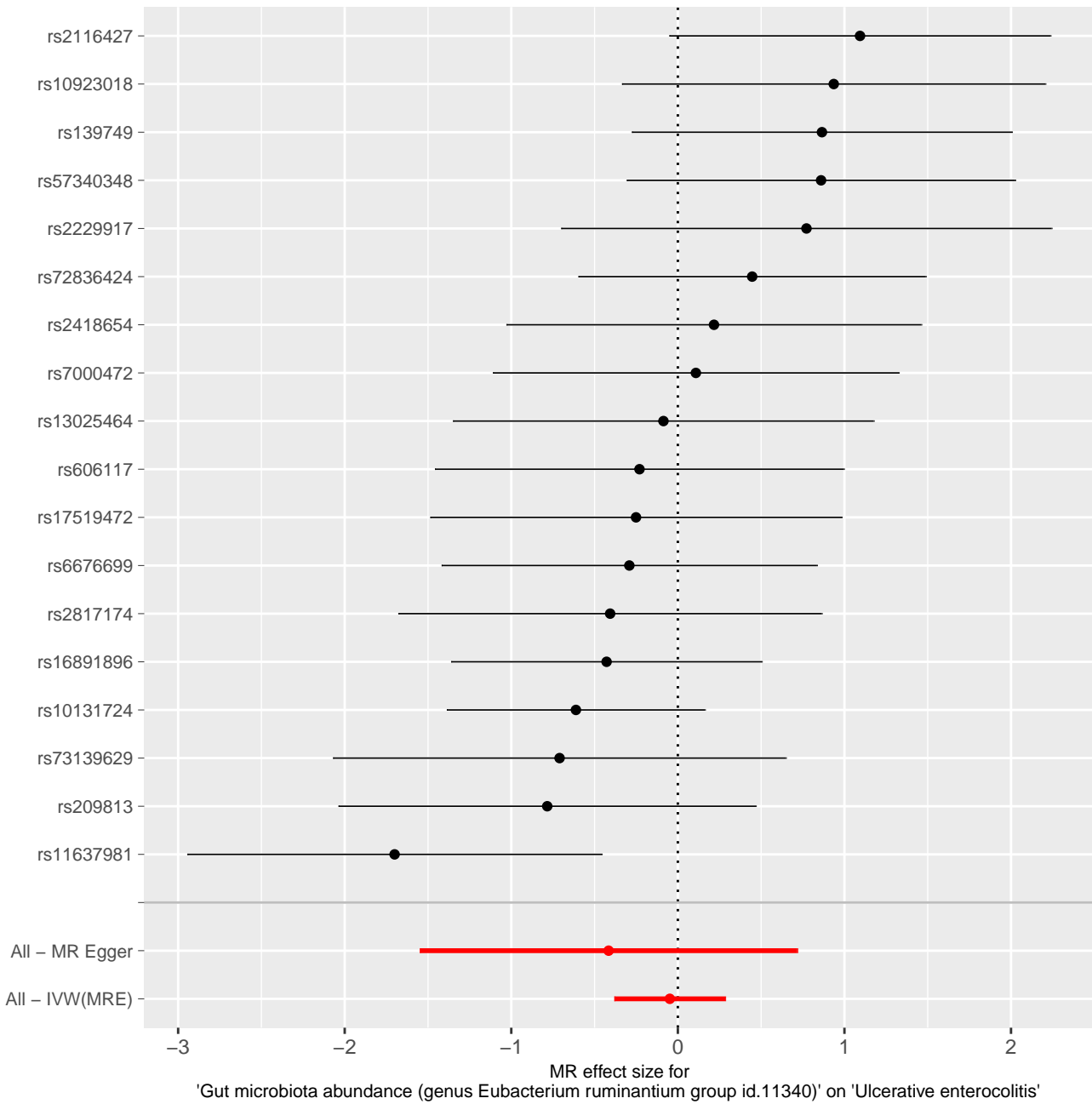
Batch 974 : Gut microbiota abundance (genus Eubacterium hallii group id.11338) on Ulcerative enterocolitis

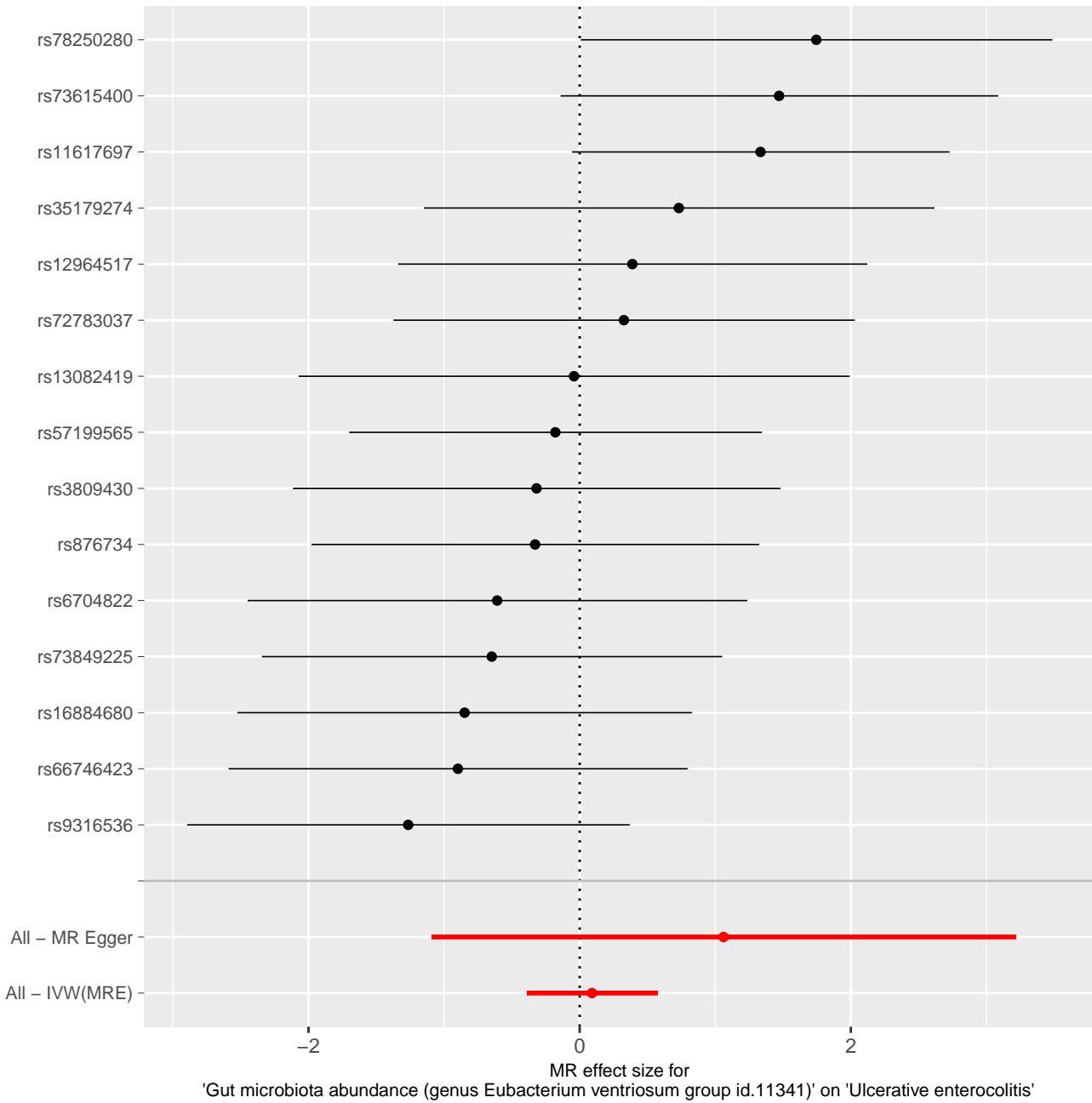


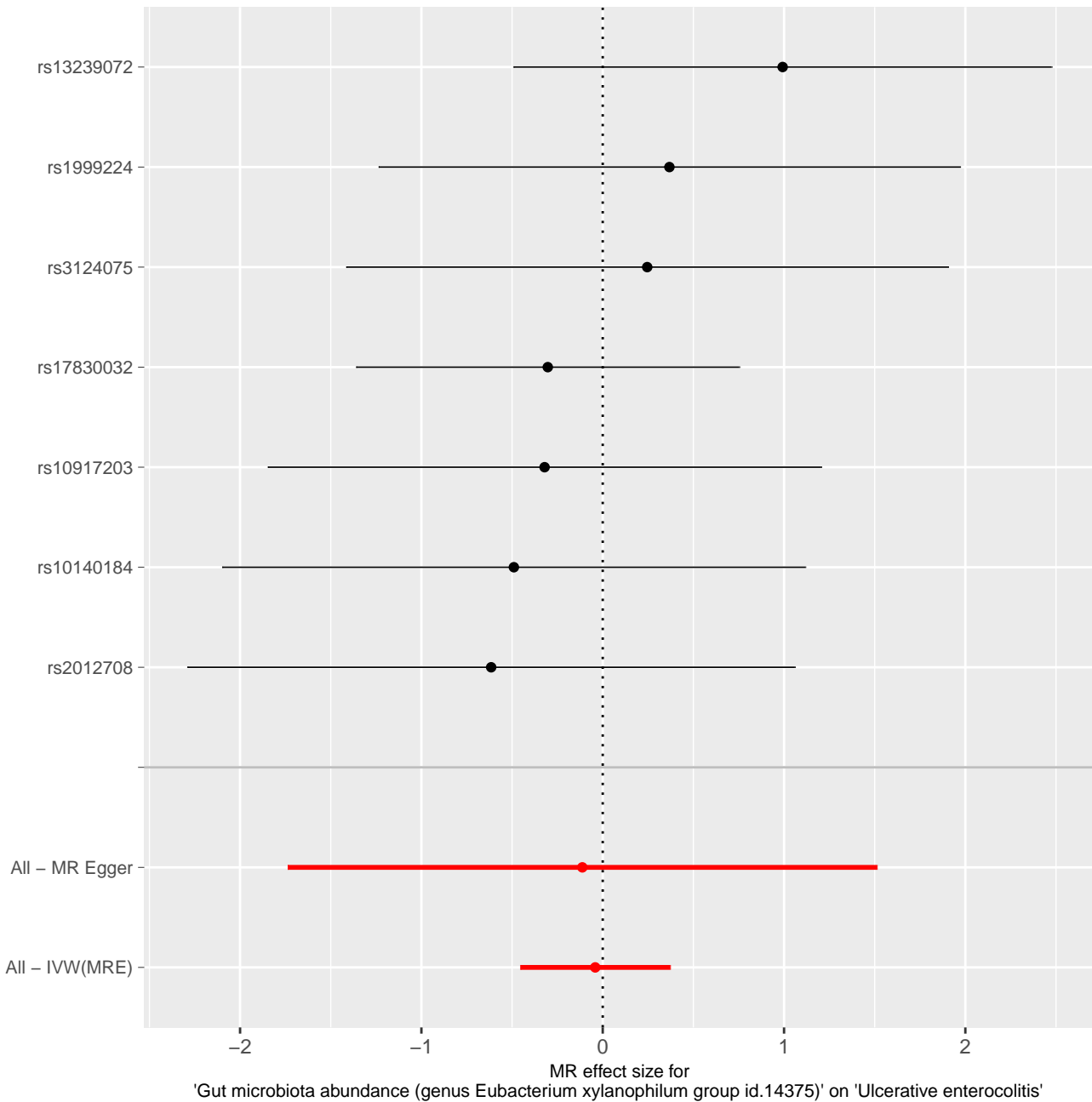


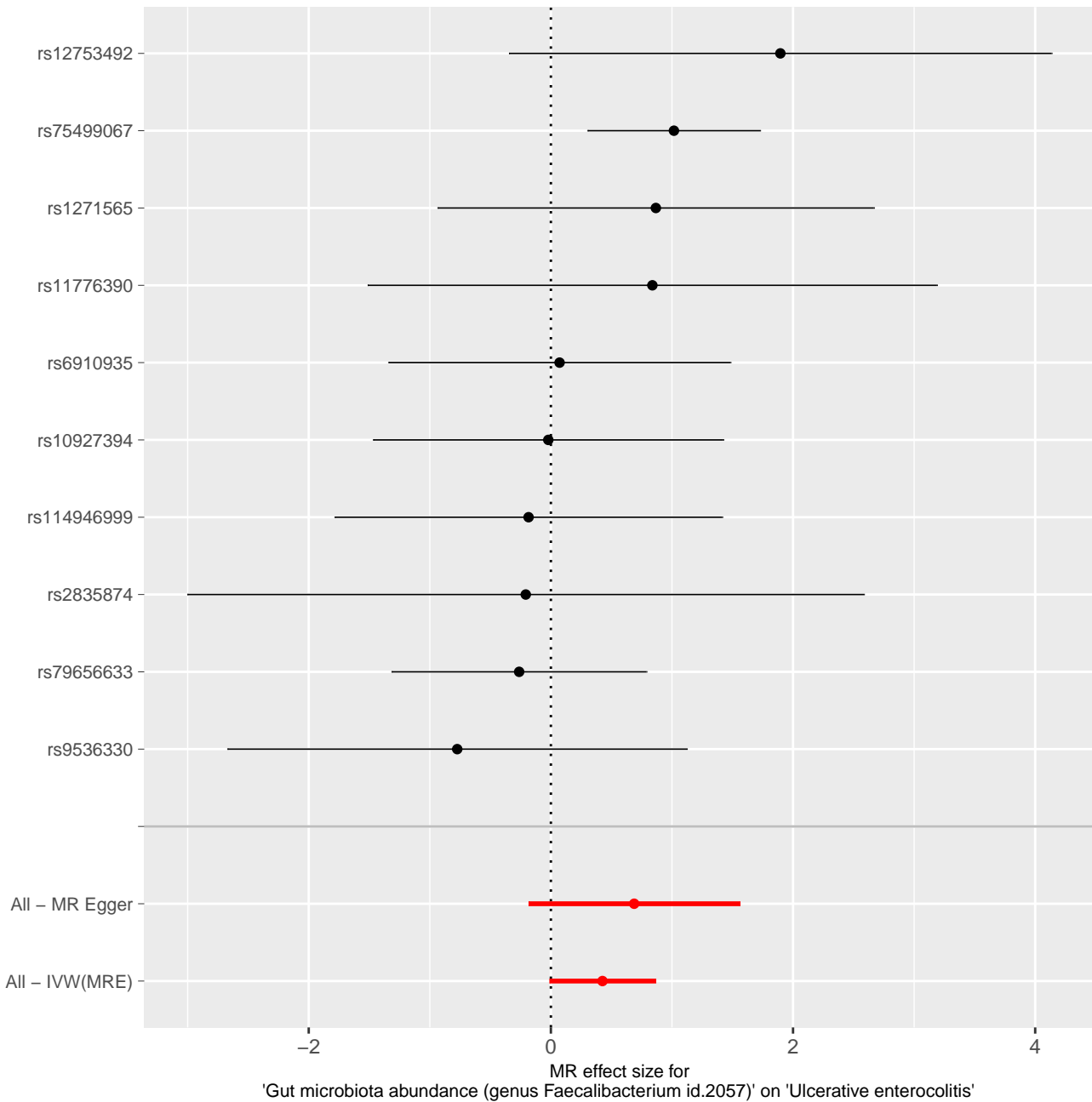


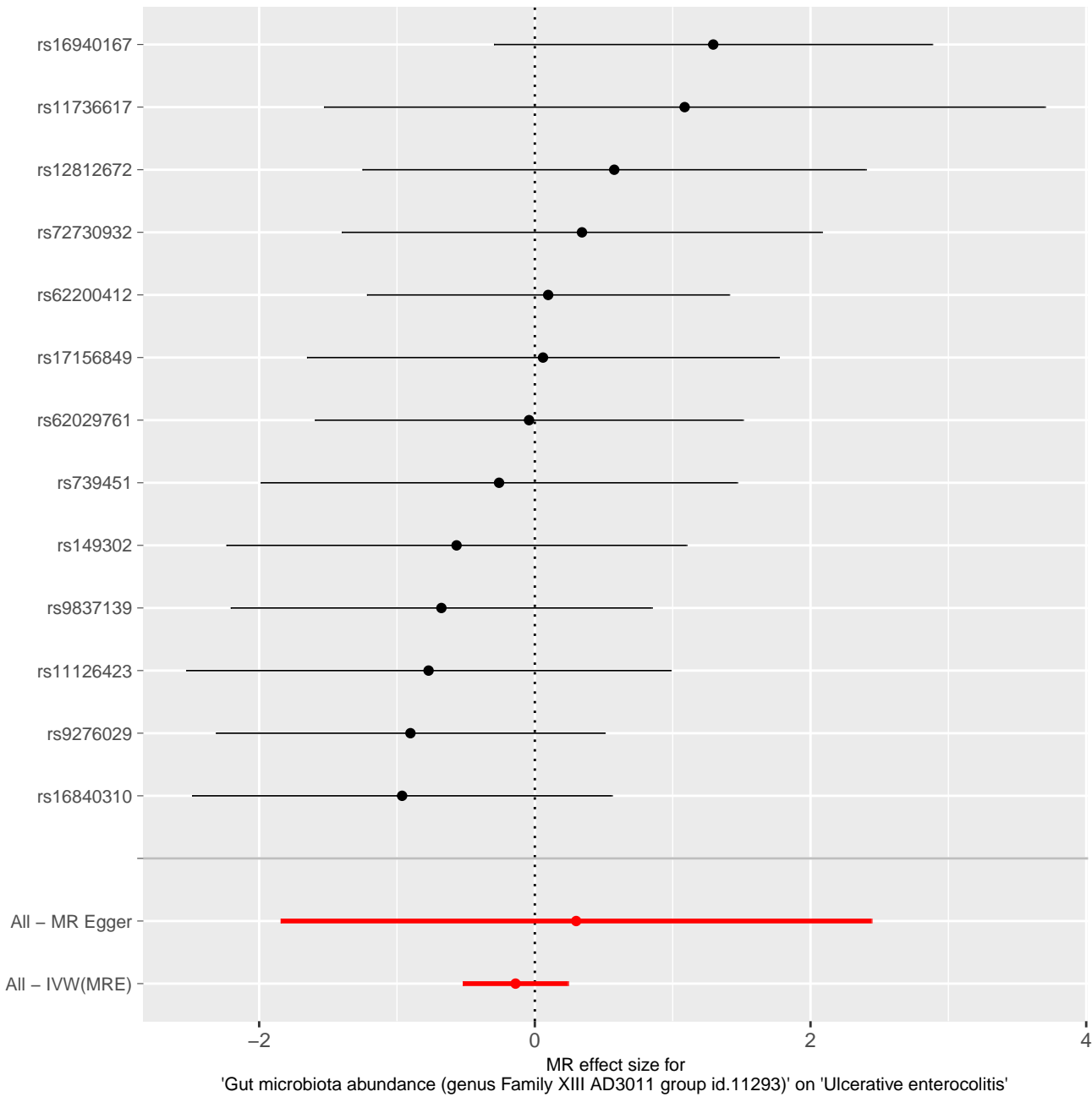


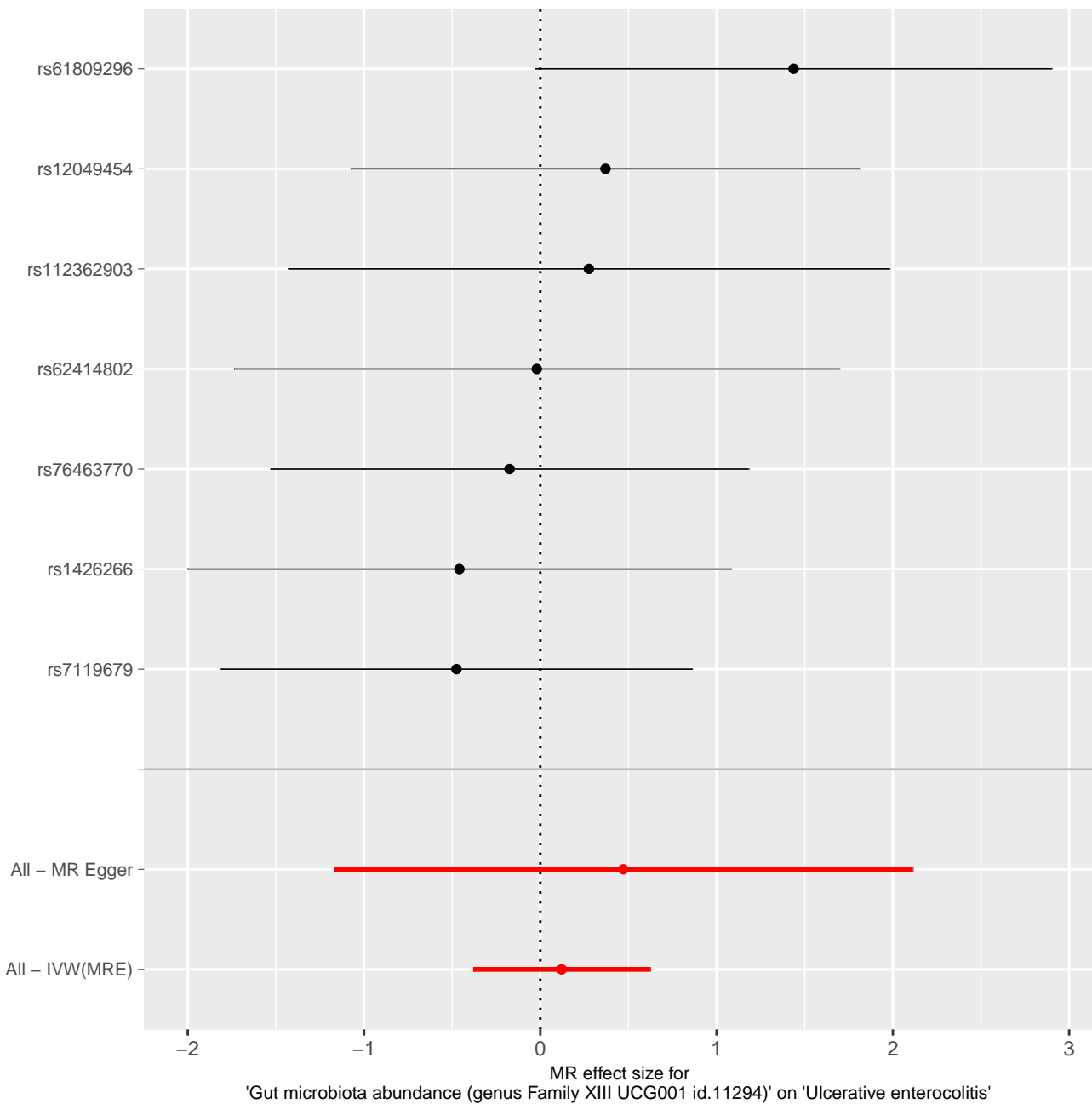


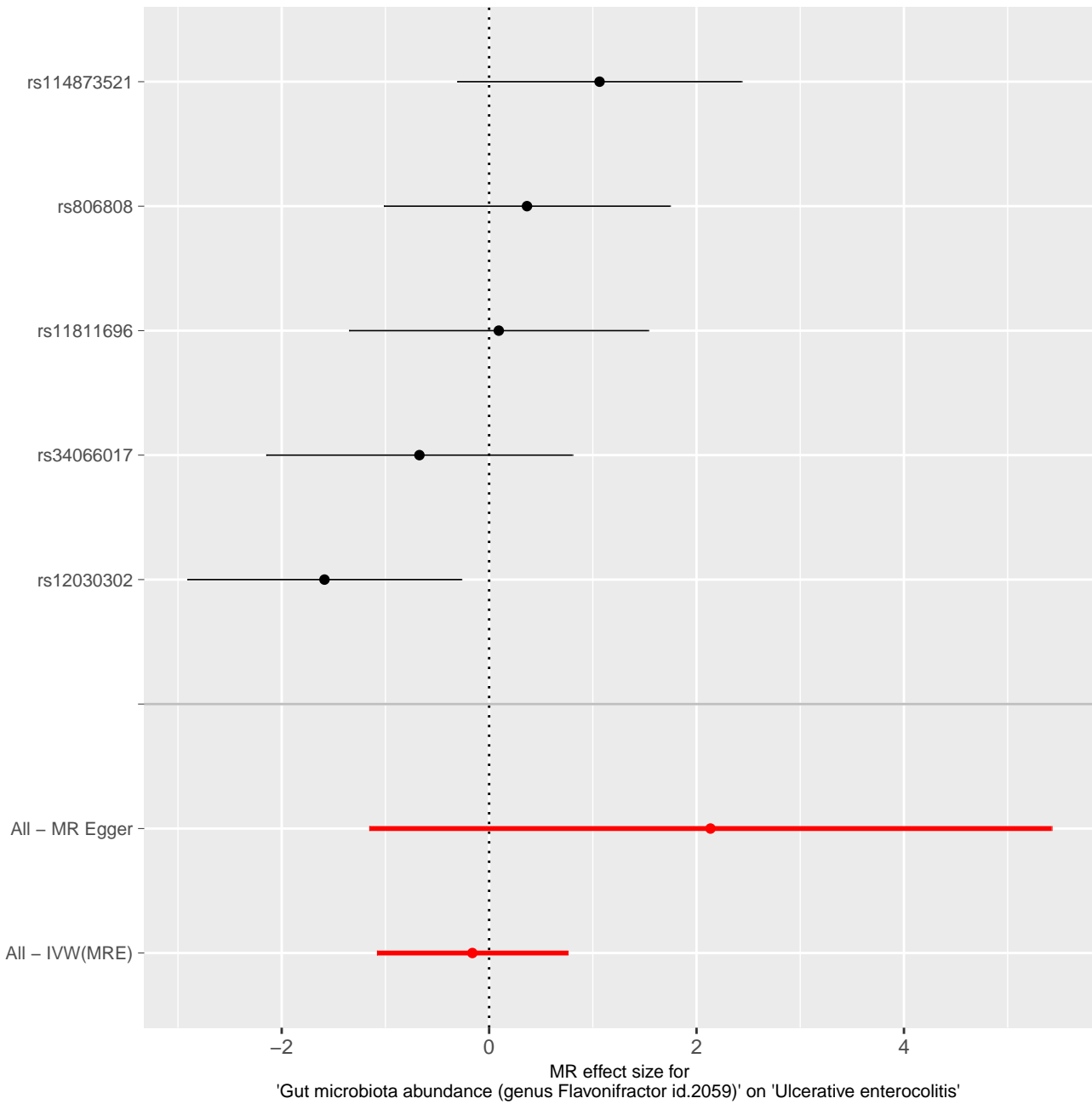


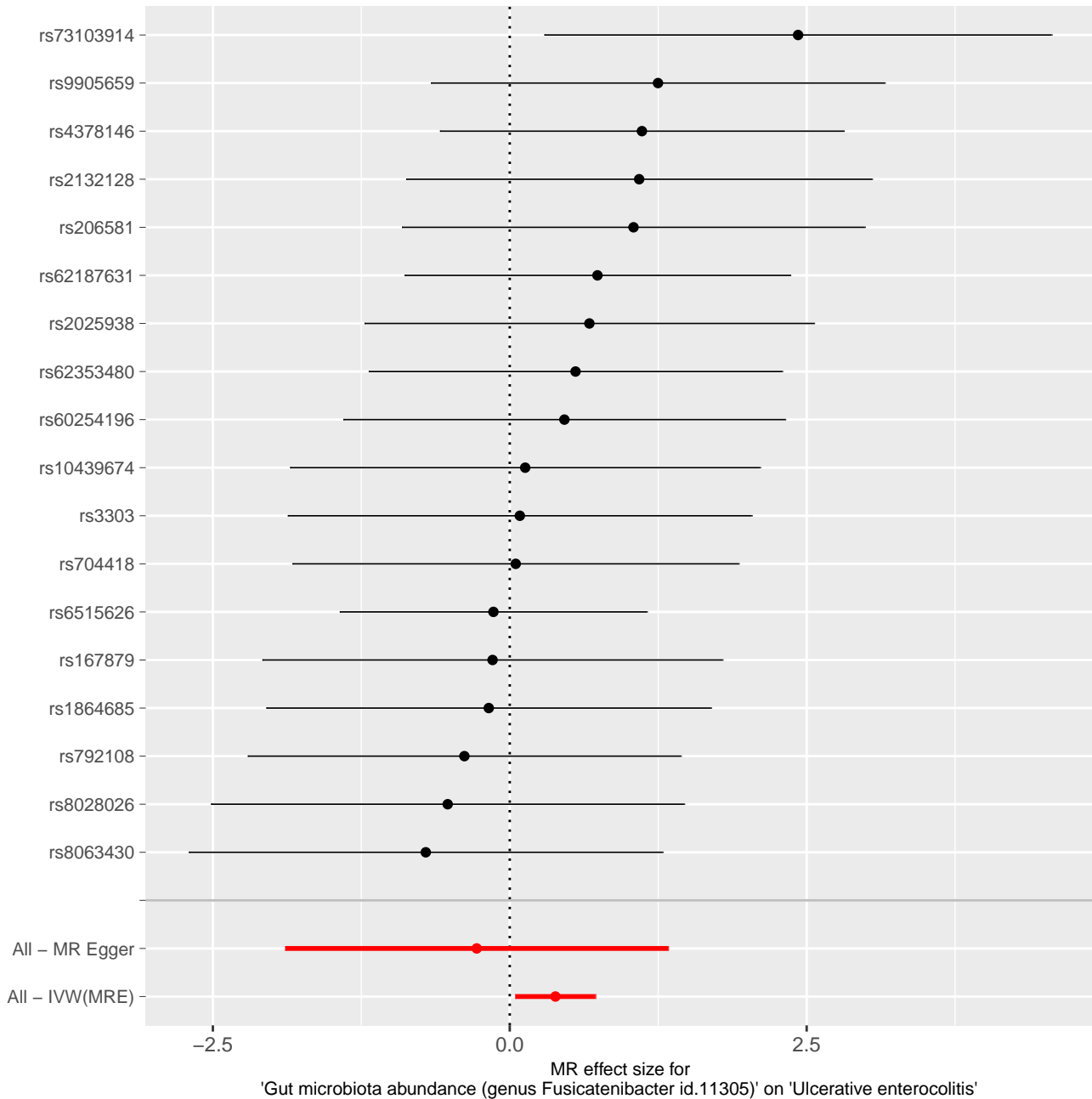


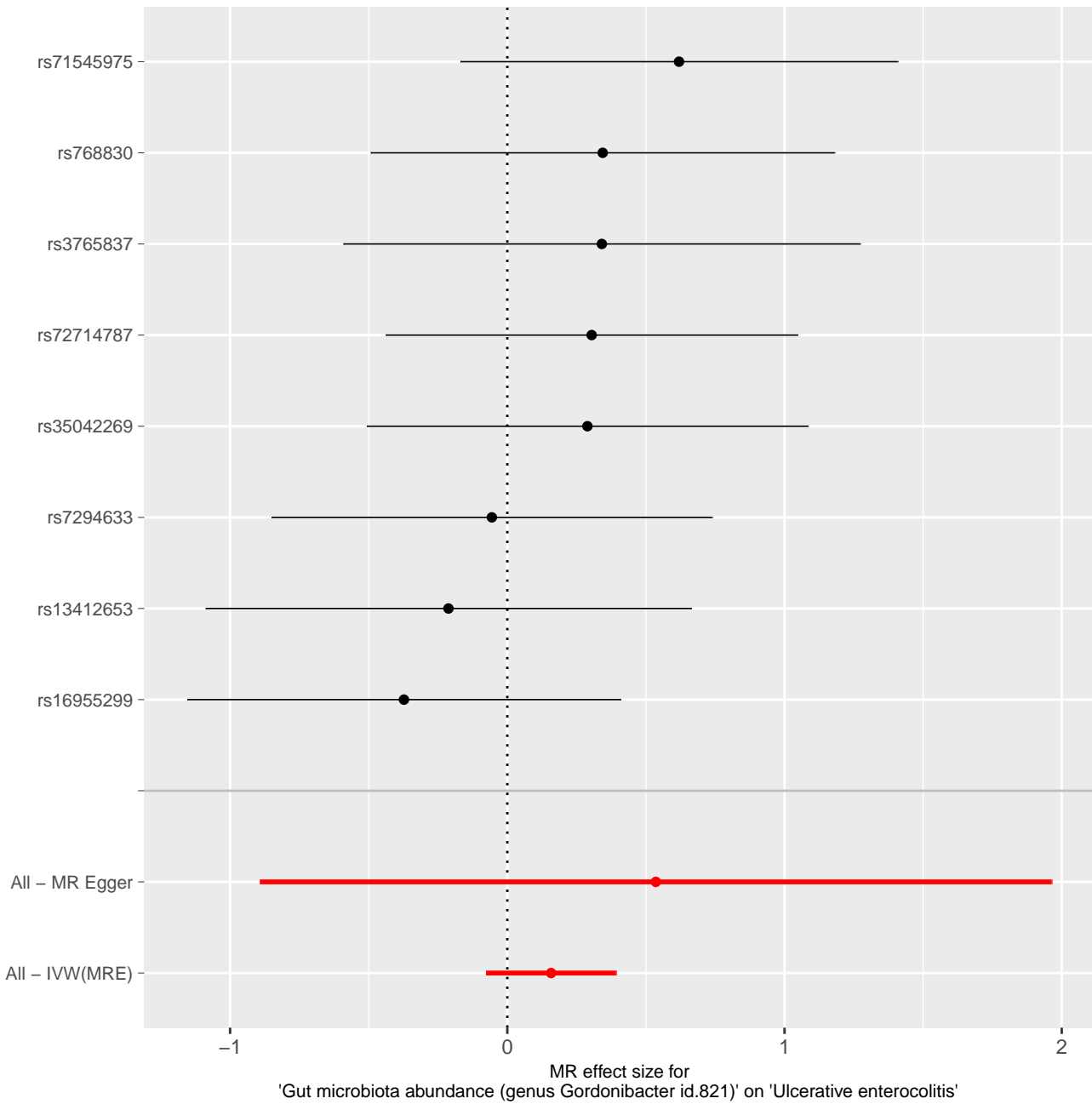




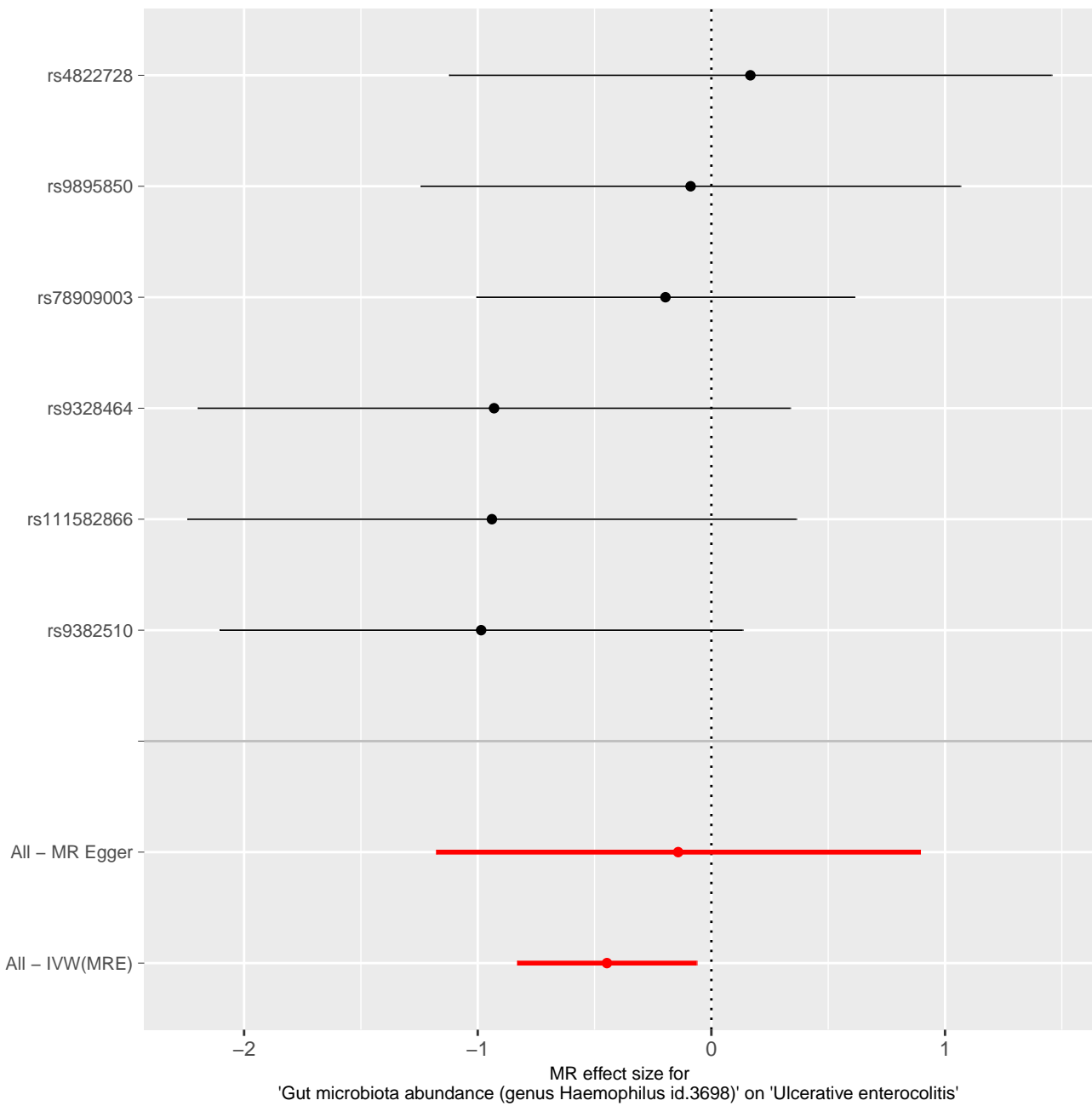


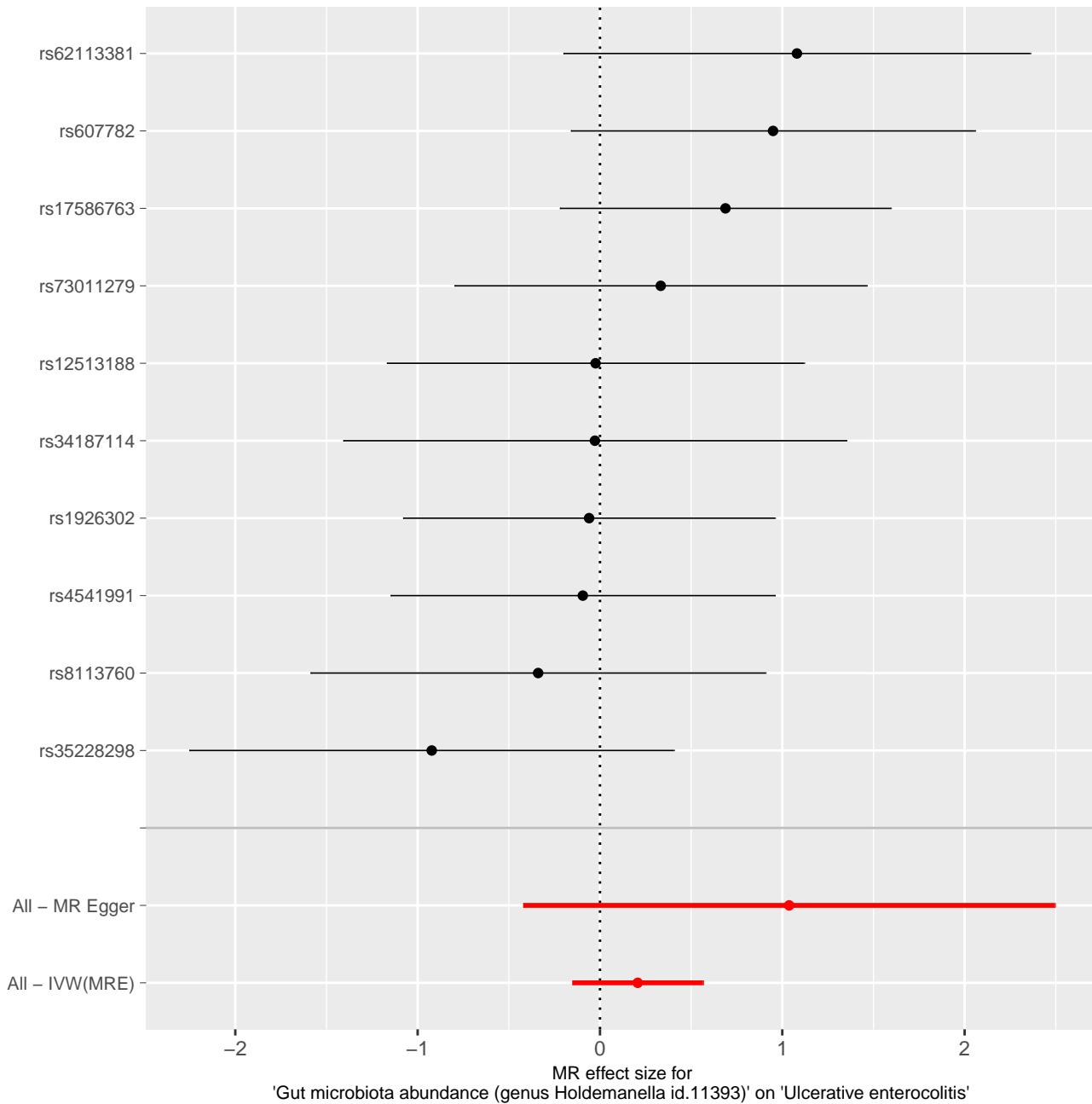




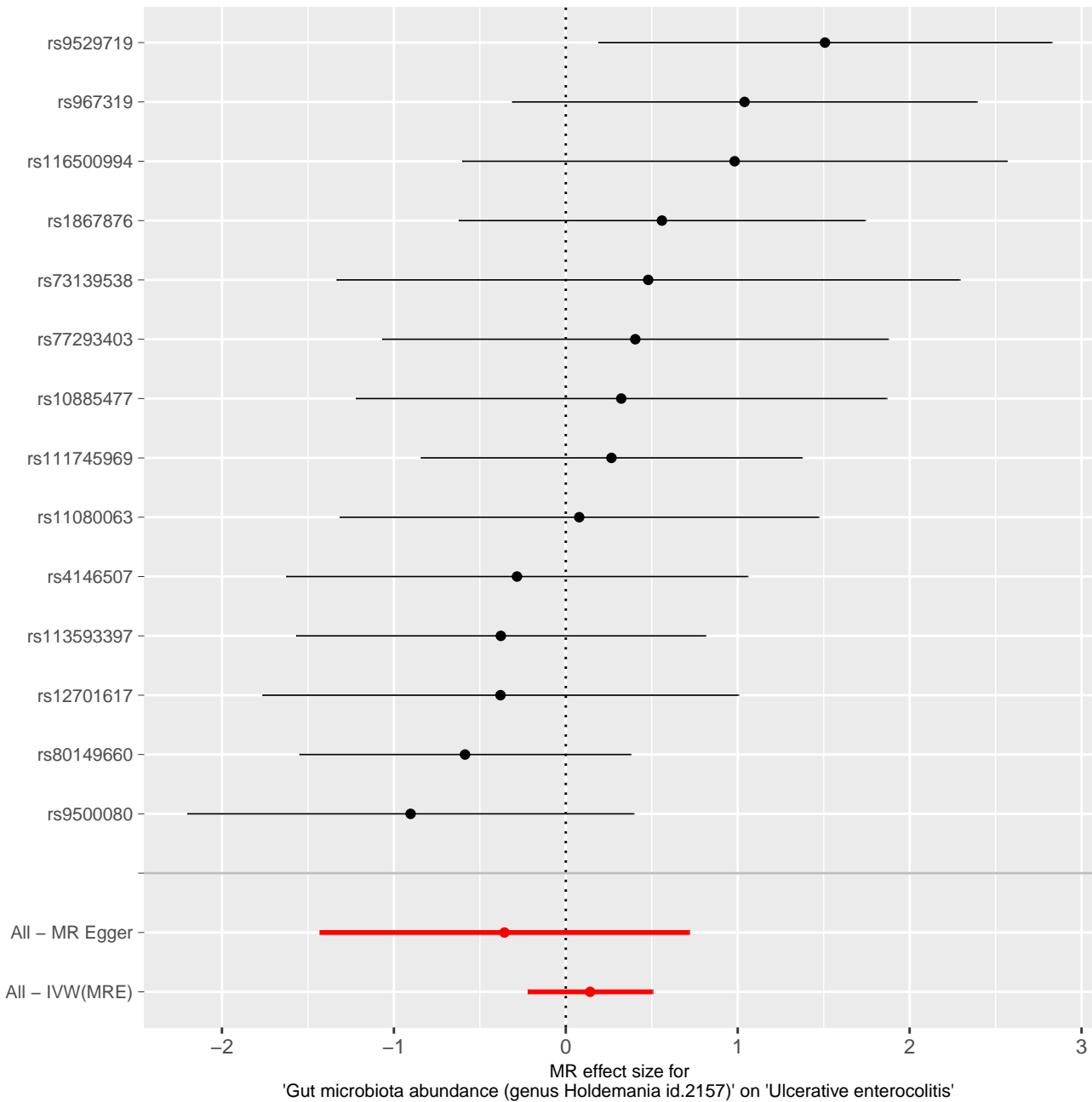


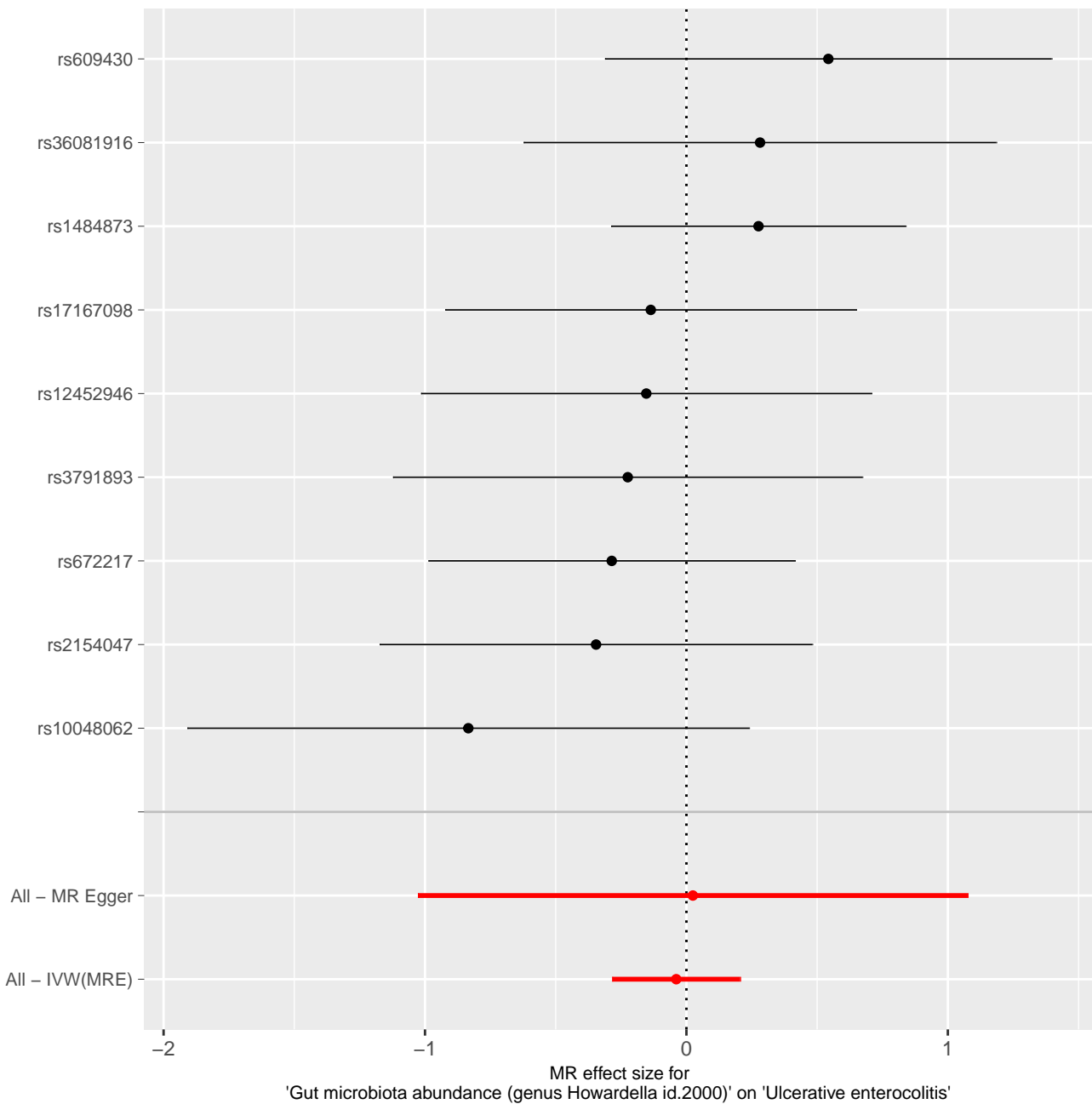
Batch 987 : Gut microbiota abundance (genus Haemophilus id.3698) on Ulcerative enterocolitis

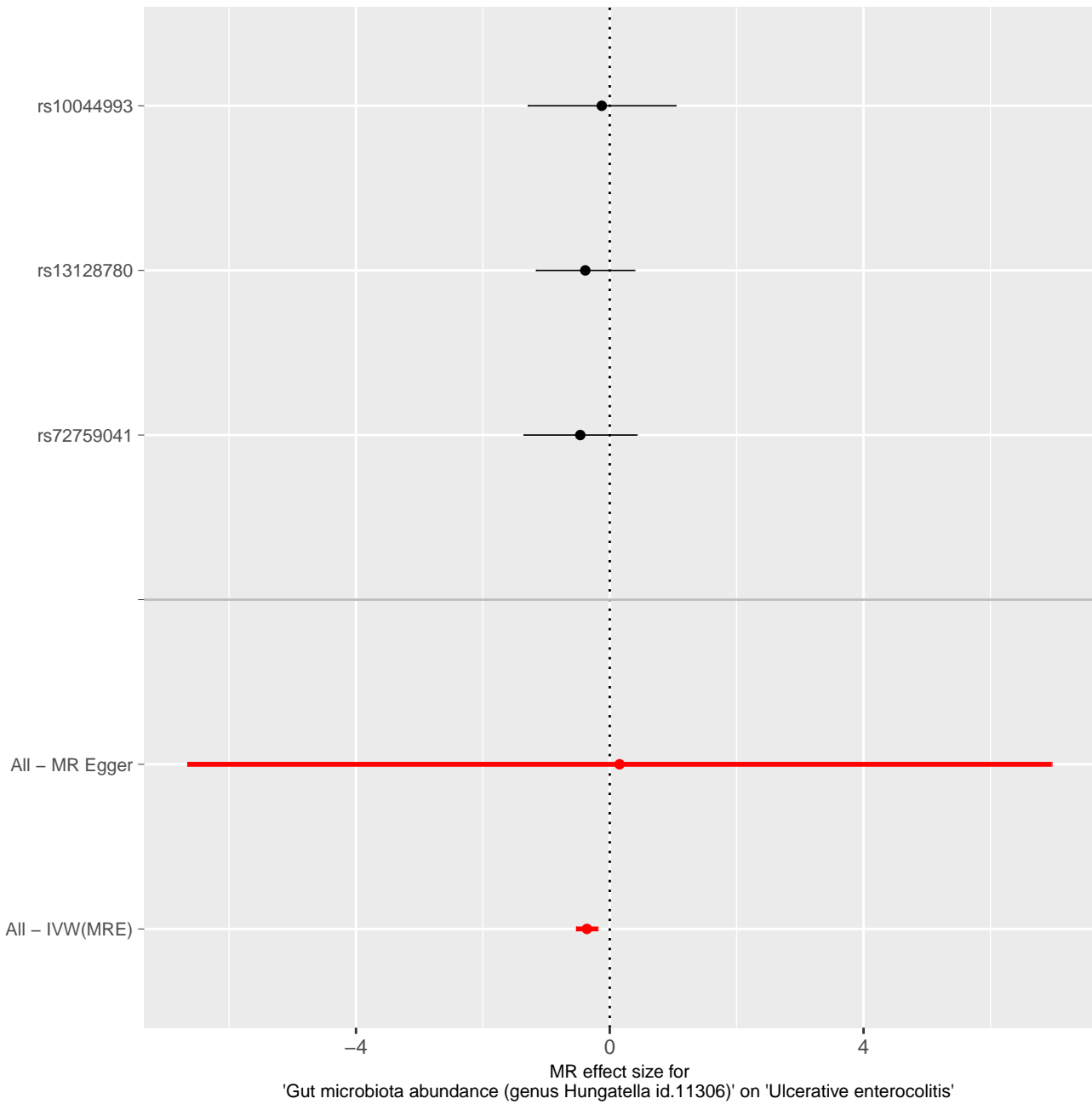




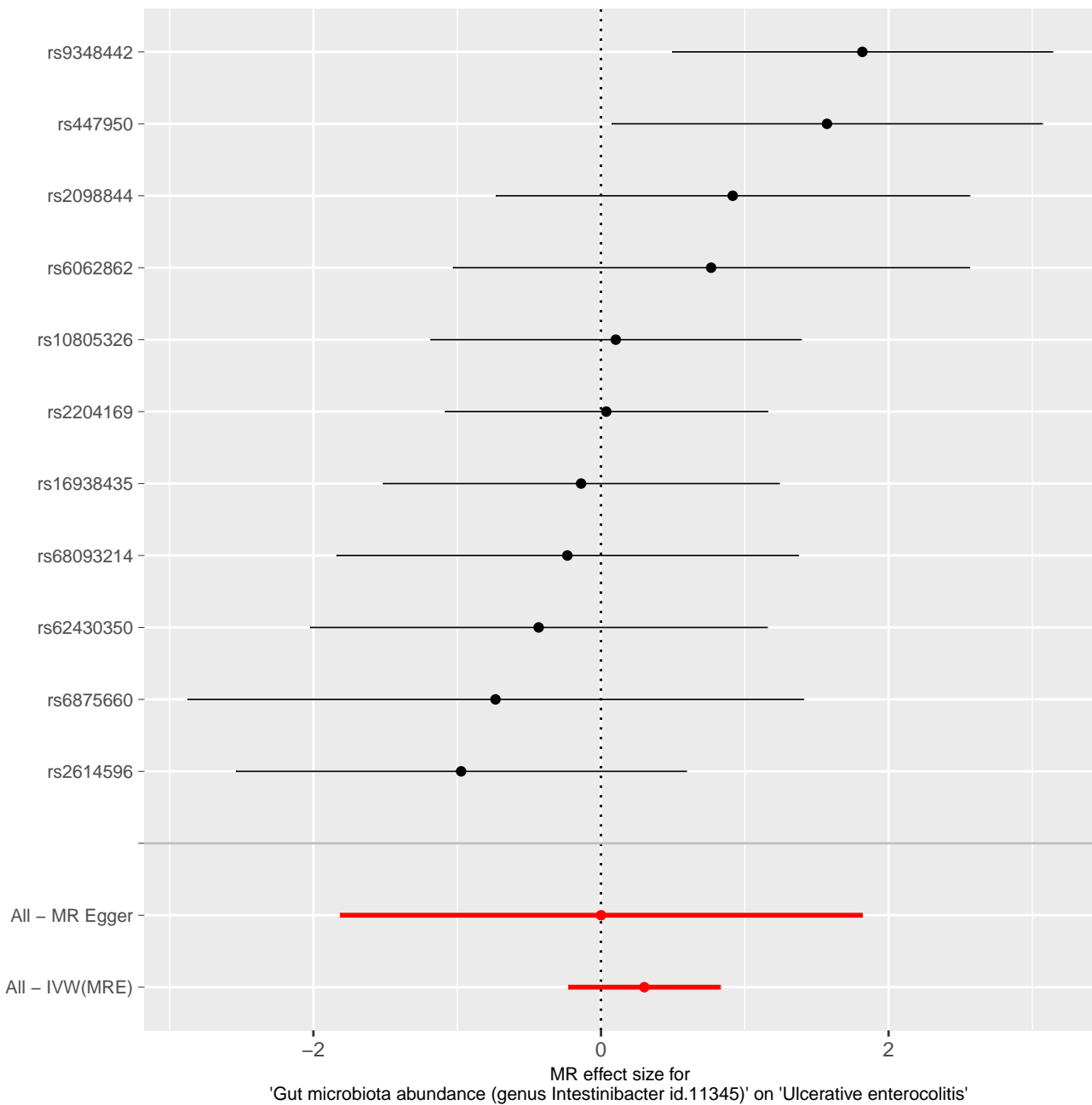
Batch 989 : Gut microbiota abundance (genus Holdemania id.2157) on Ulcerative enterocolitis

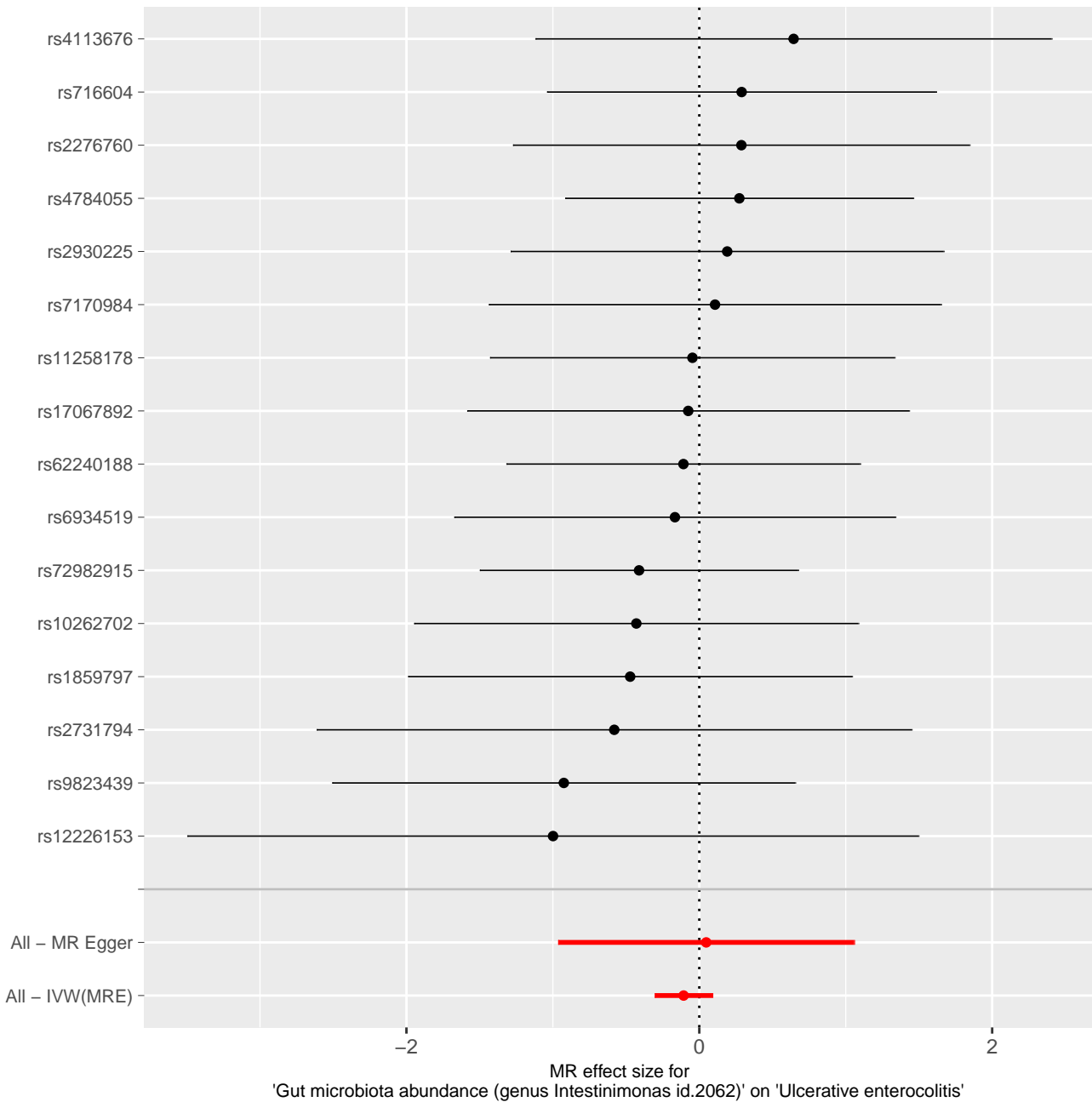


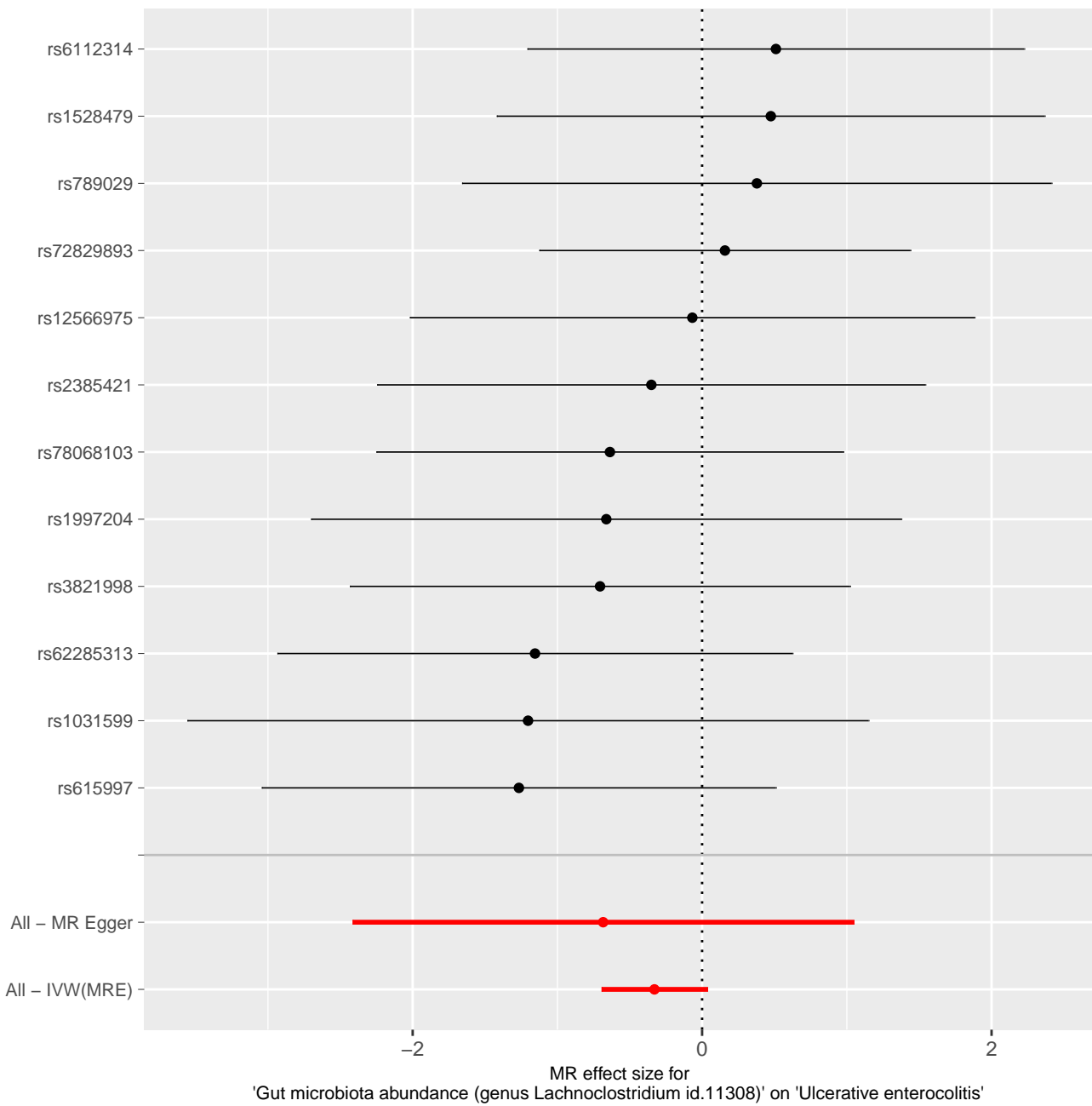


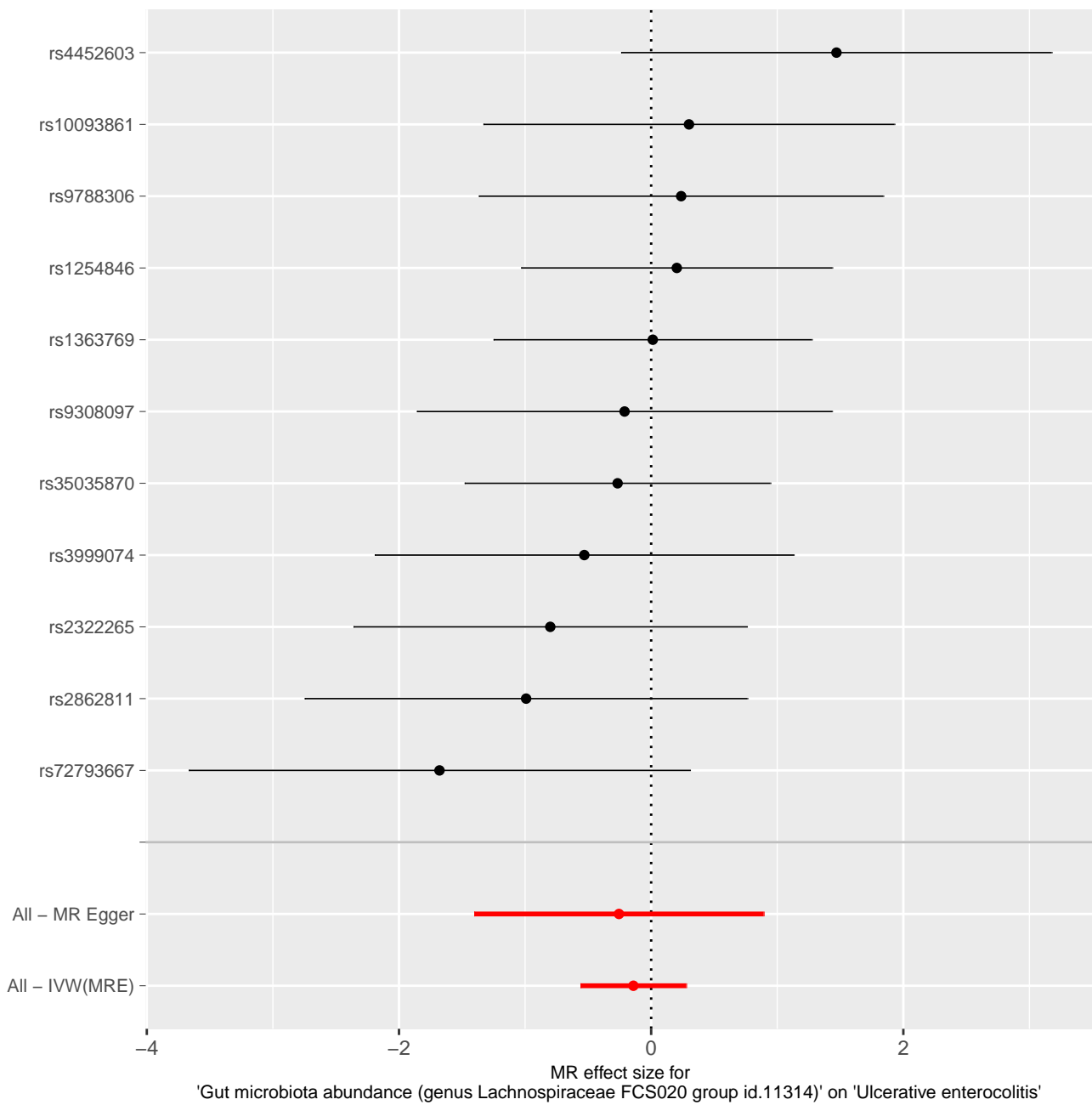


Batch 992 : Gut microbiota abundance (genus Intestinibacter id.11345) on Ulcerative enterocolitis

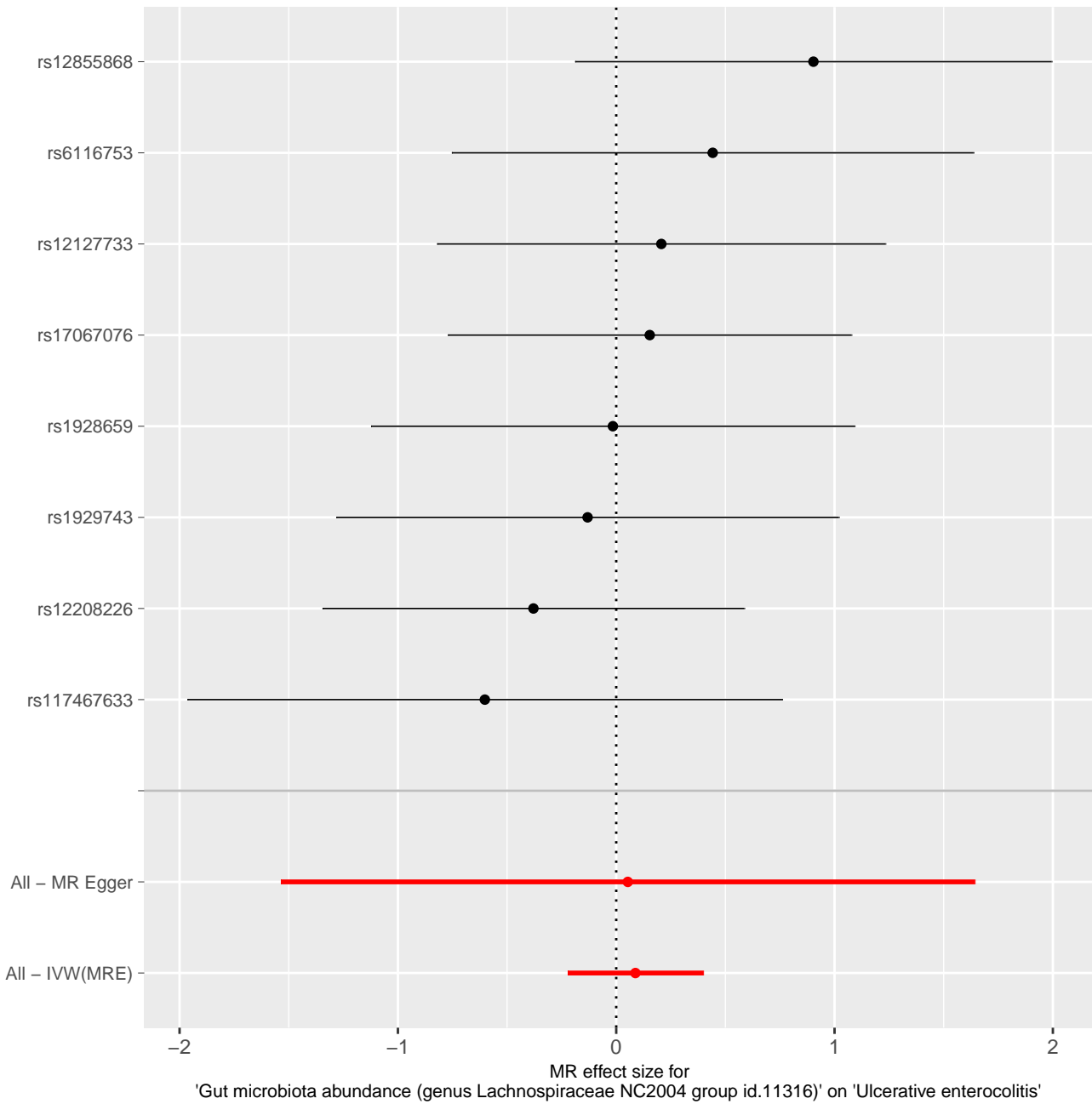


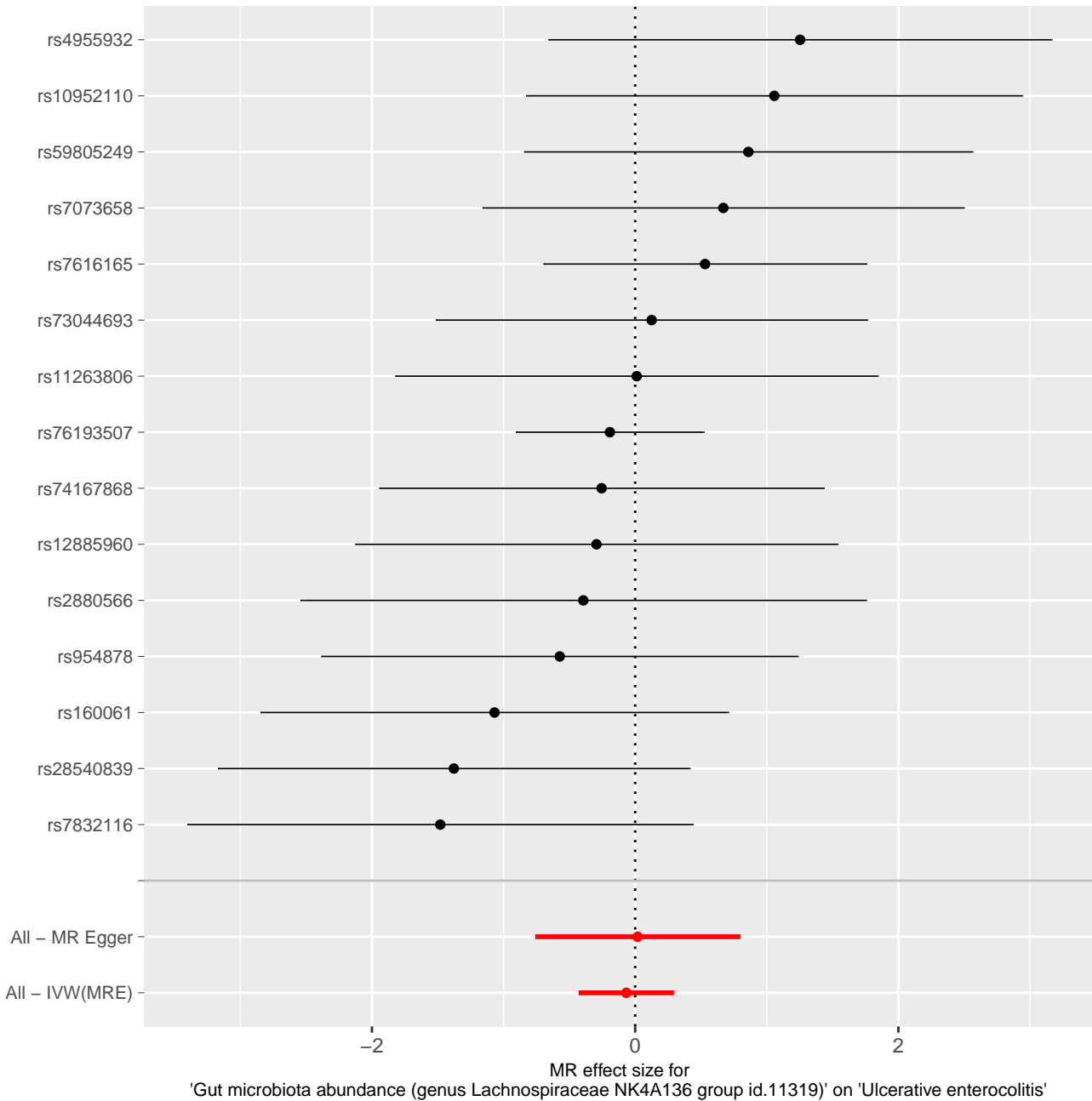


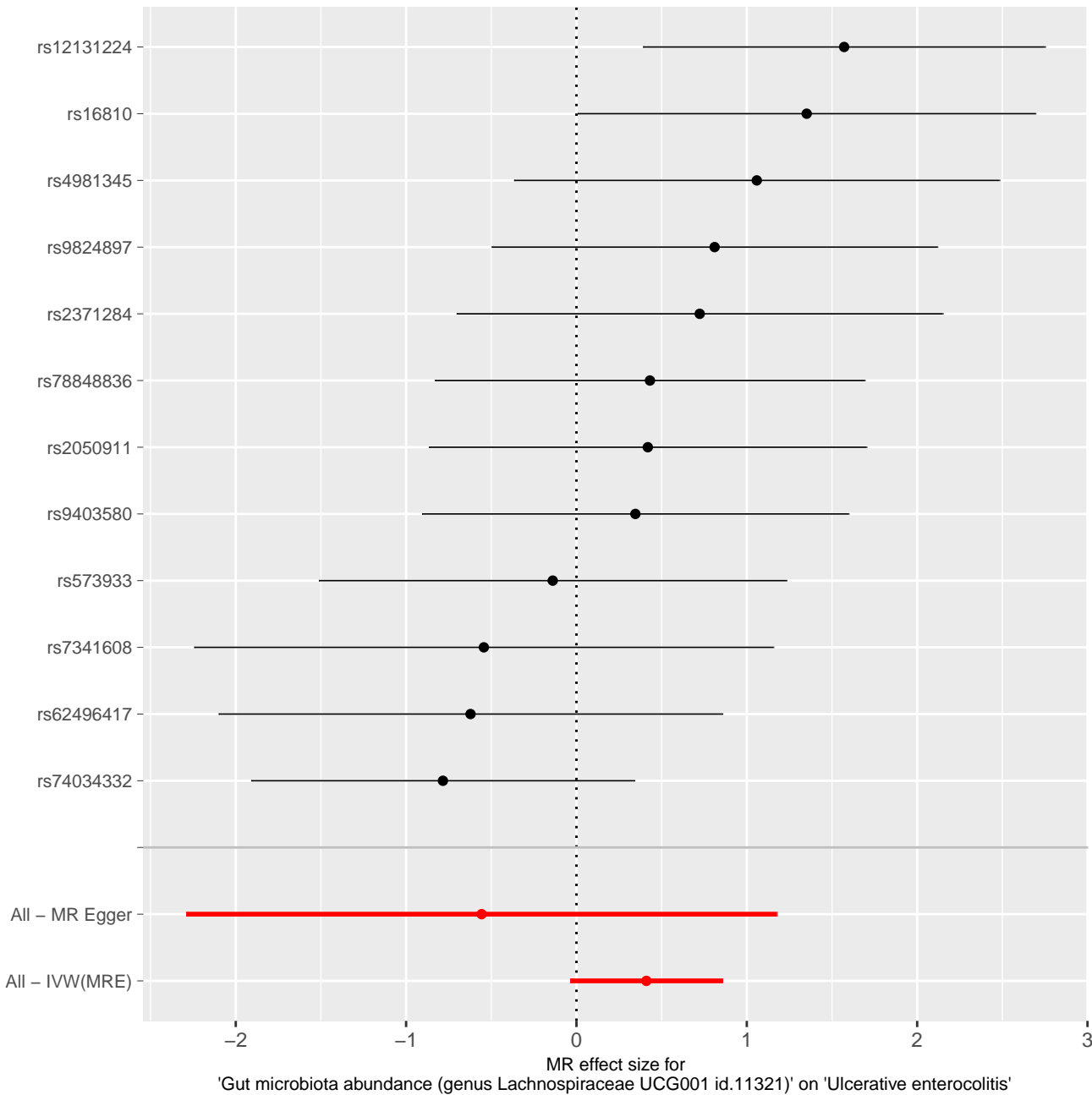


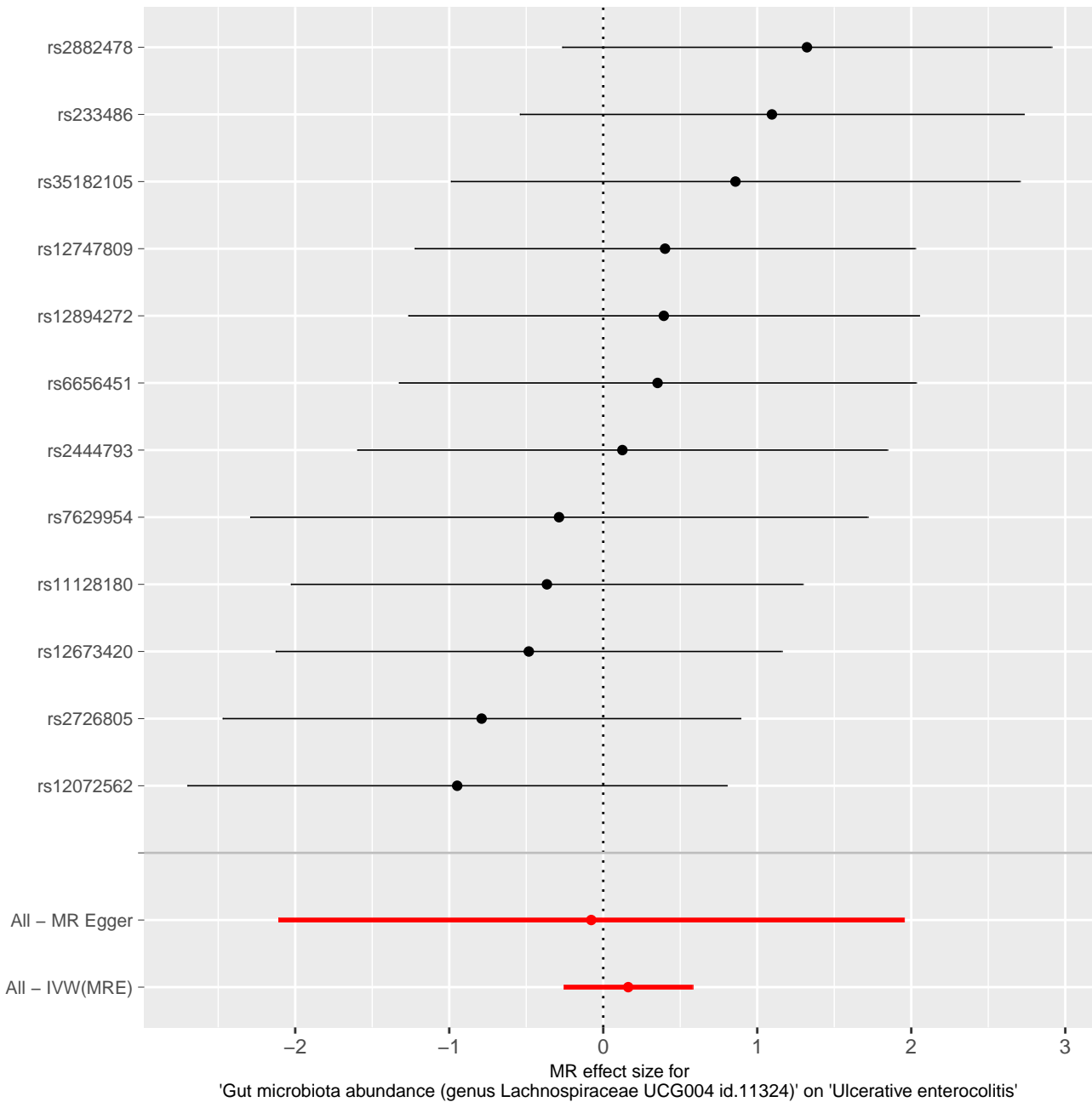


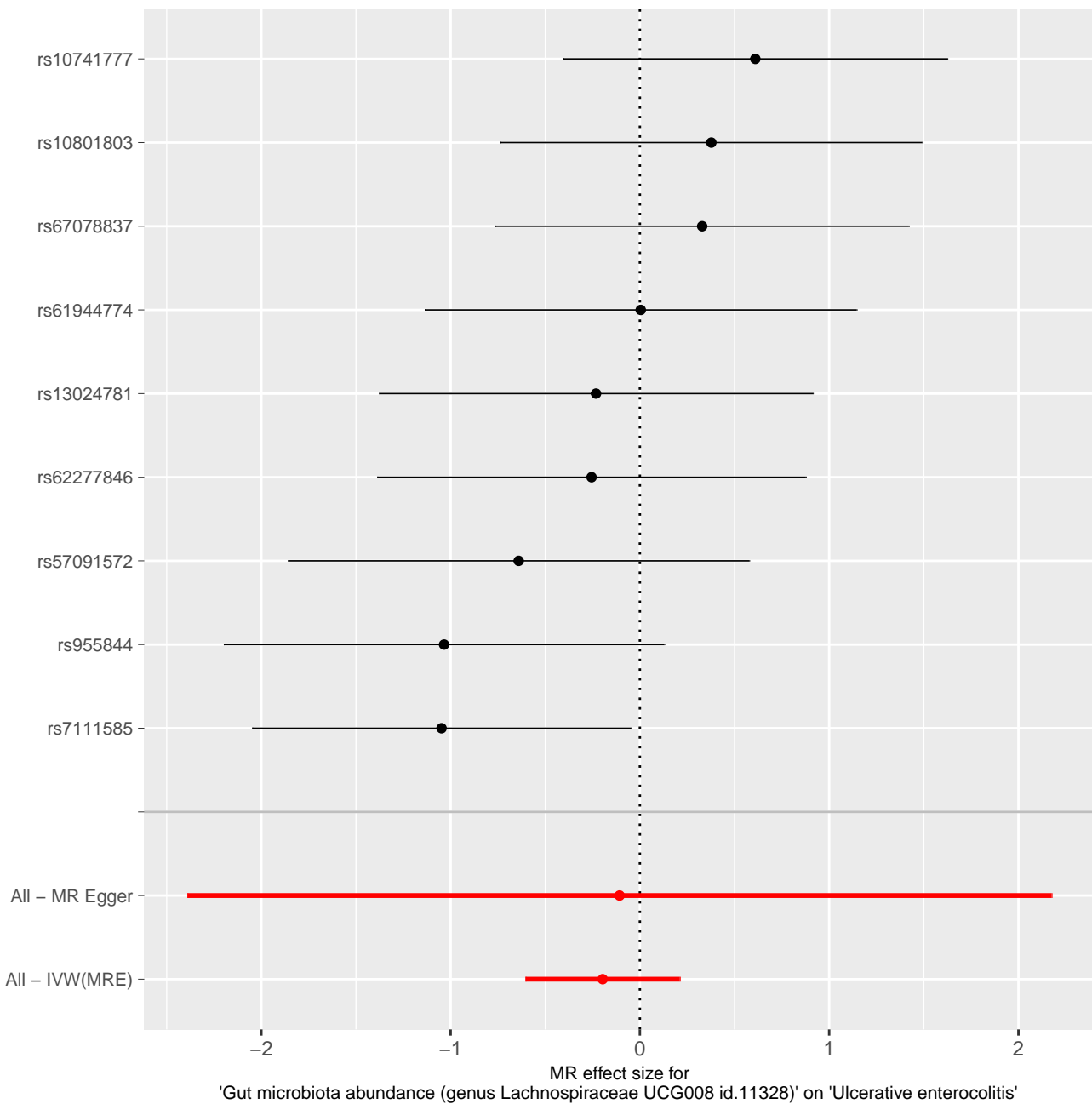
MR effect size for
'Gut microbiota abundance (genus Lachnospiraceae FCS020 group id.11314)' on 'Ulcerative enterocolitis'

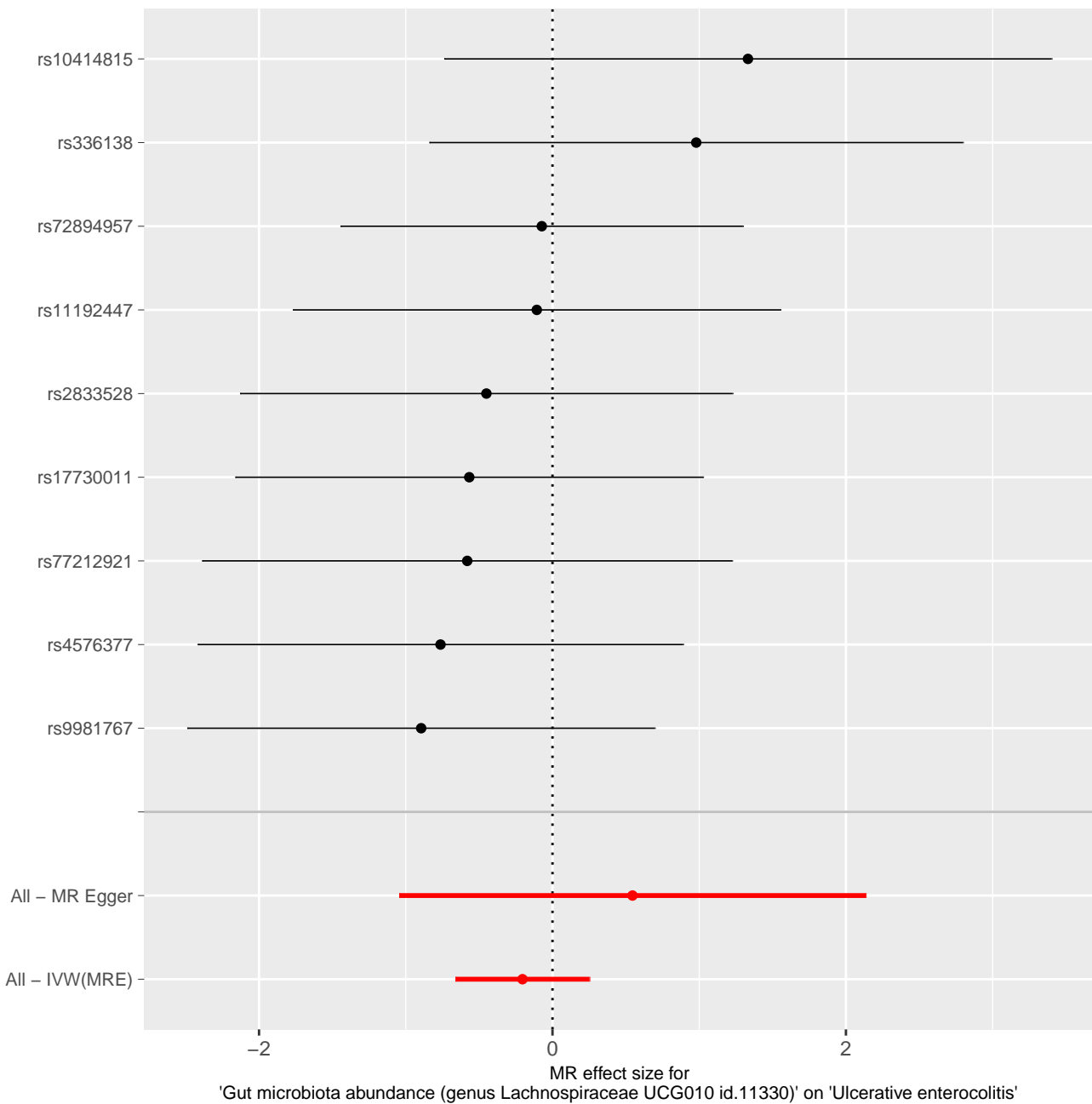


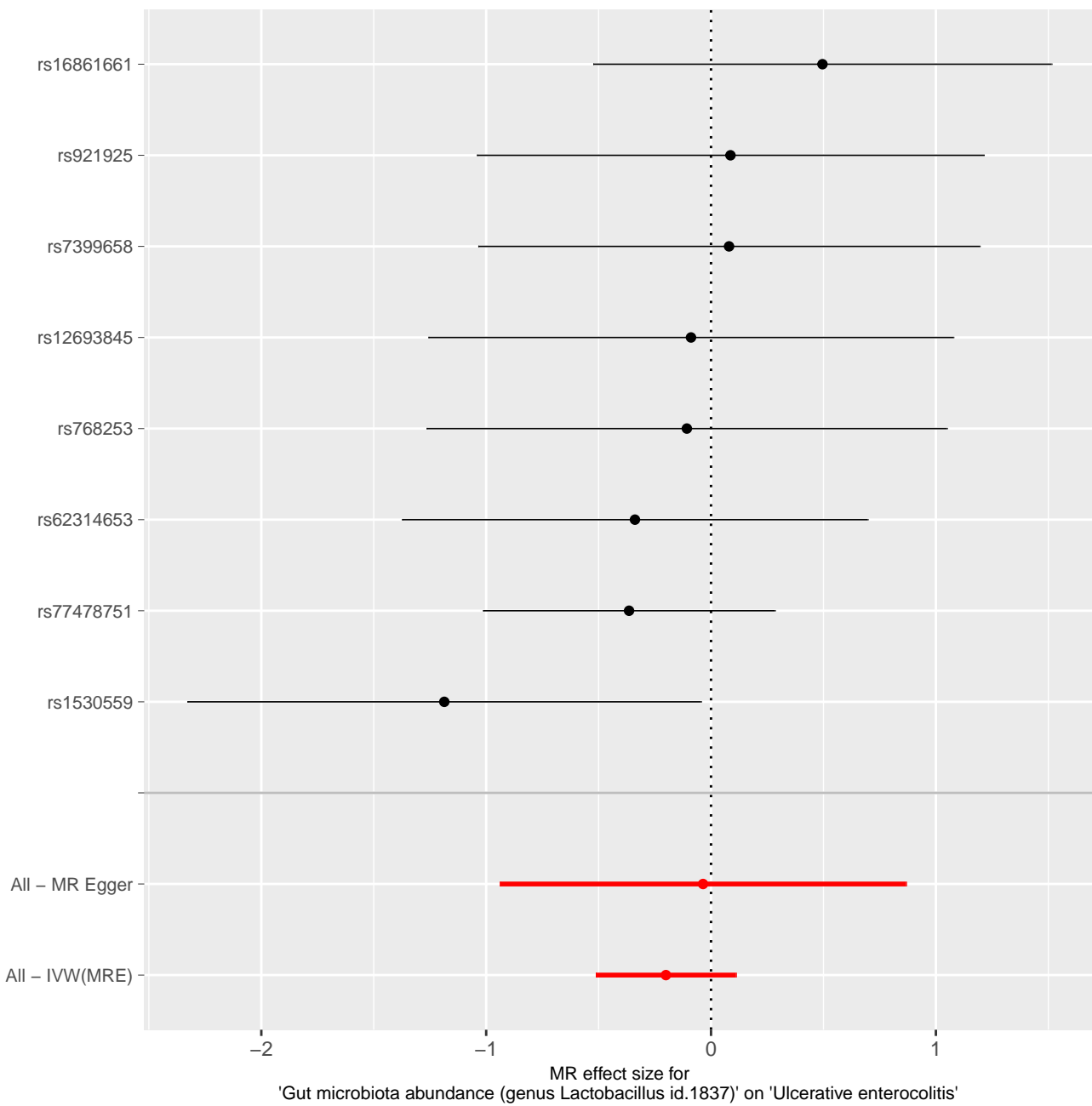


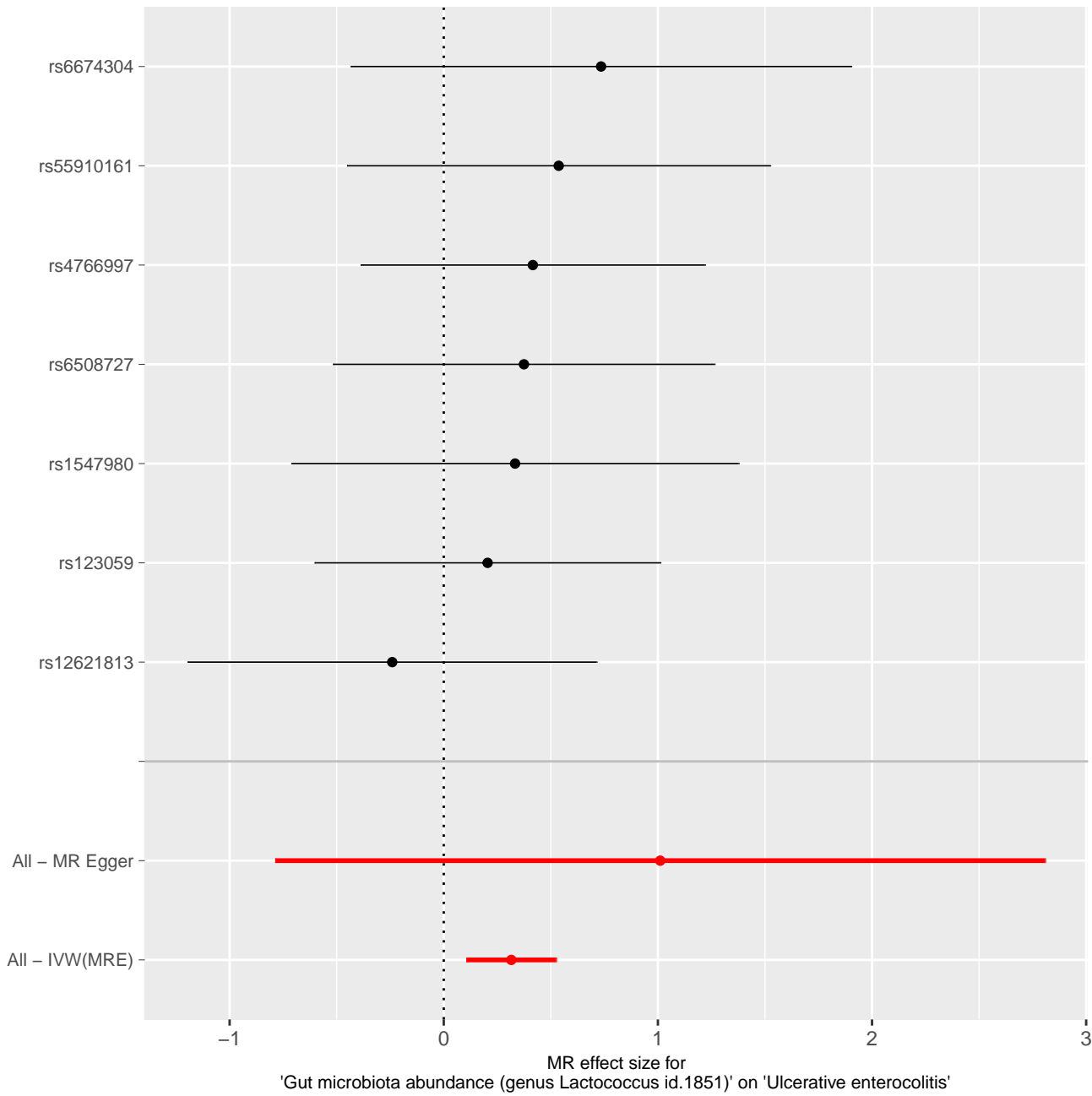


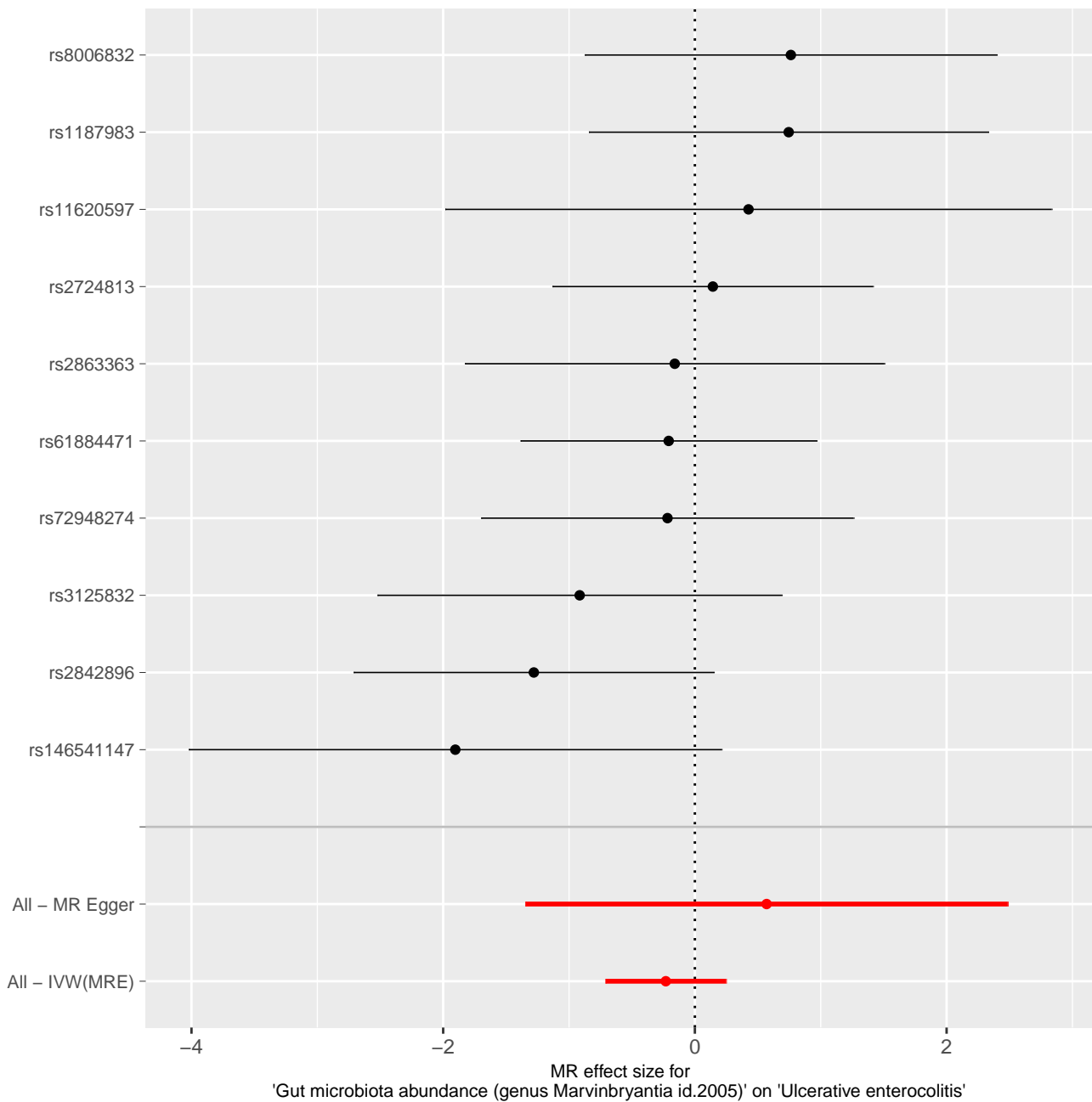


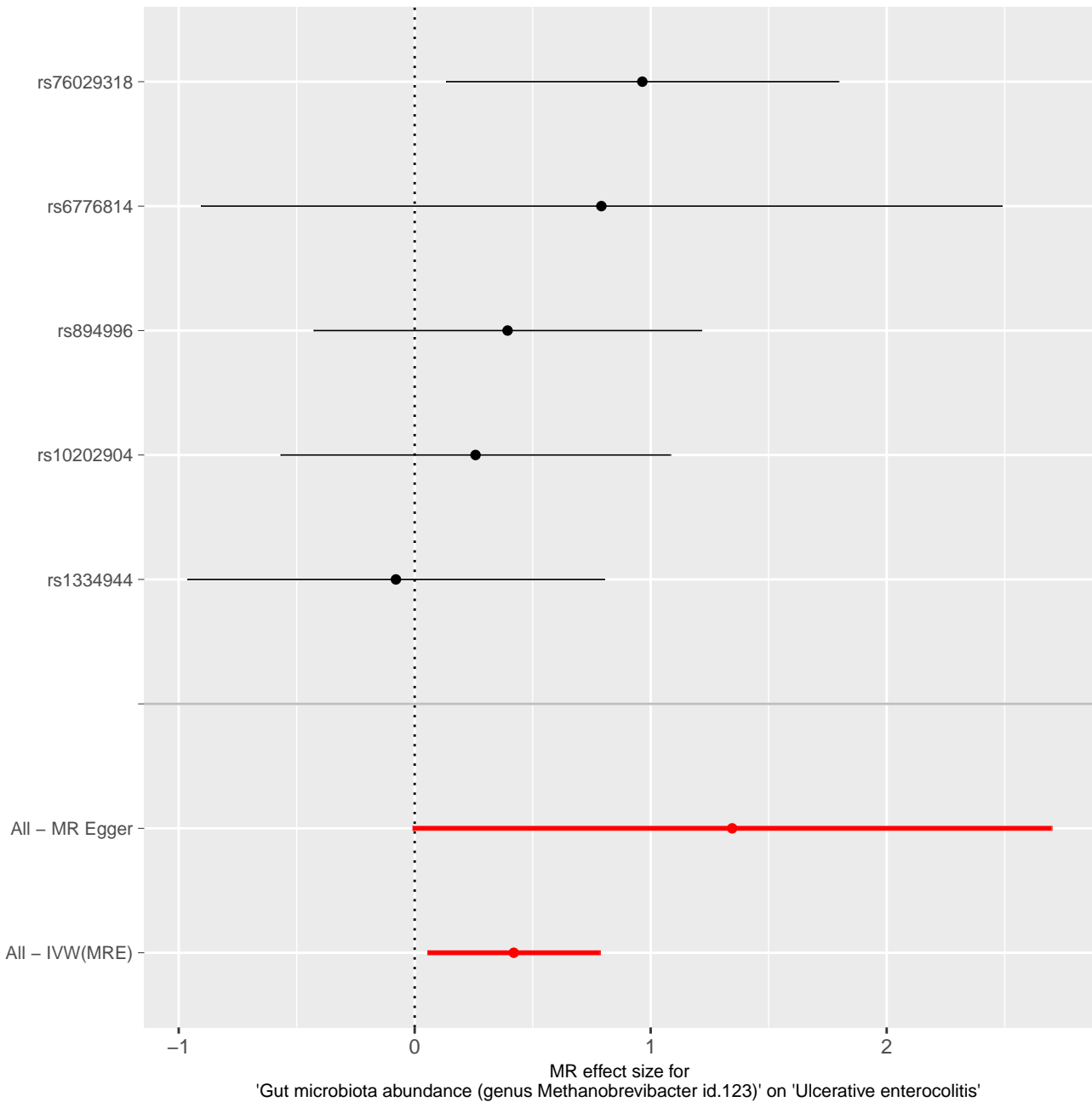


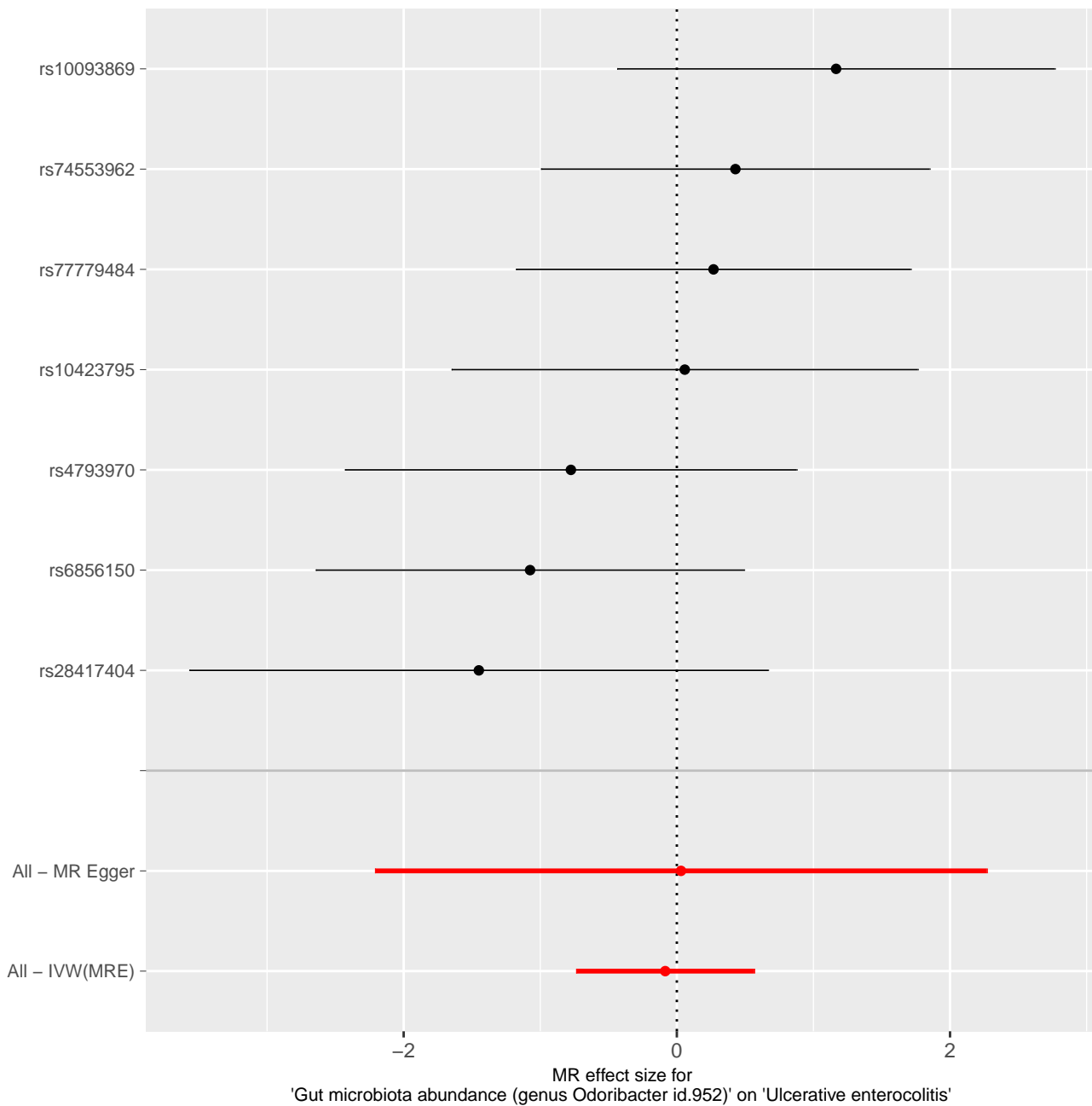


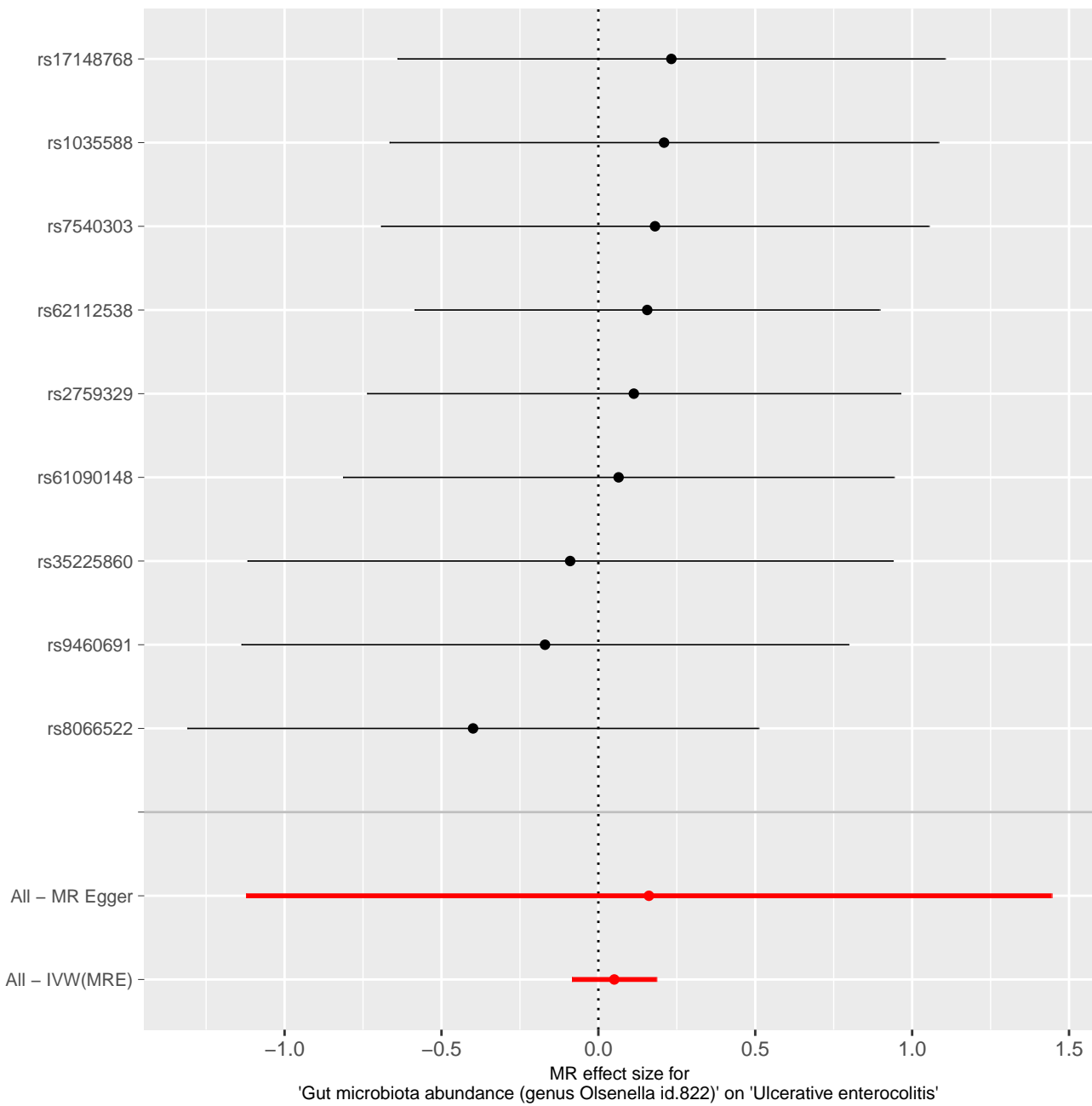


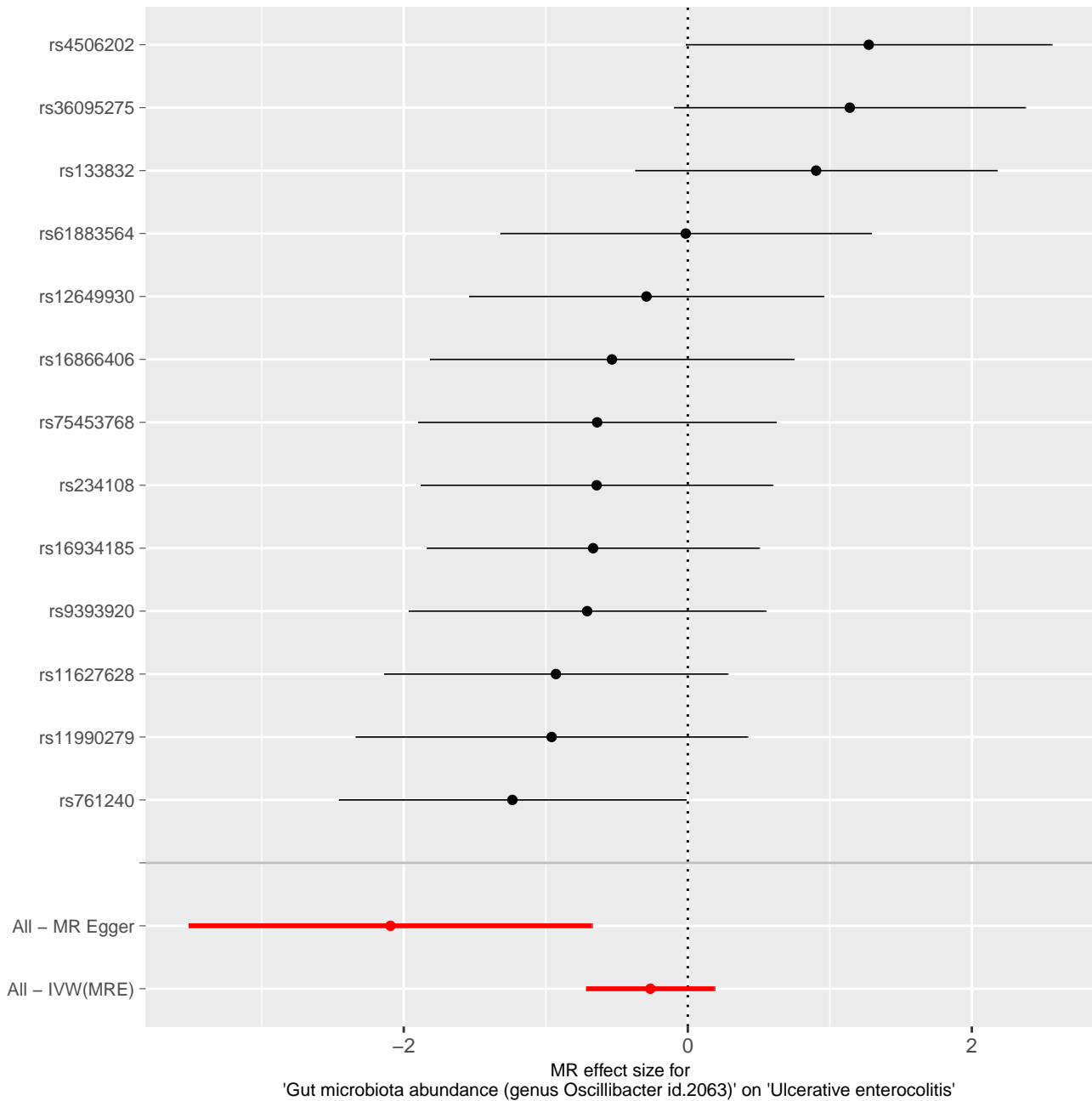


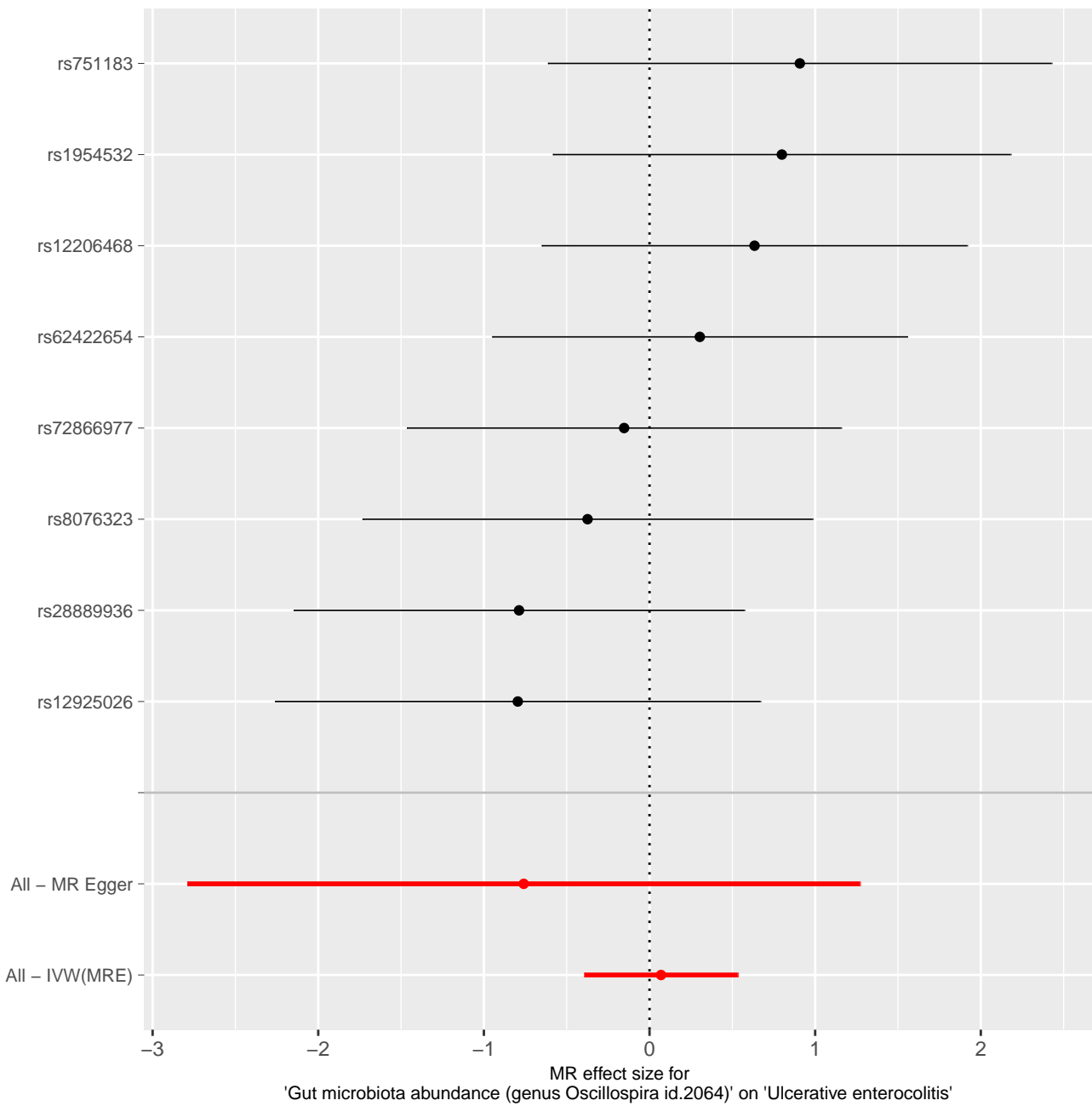




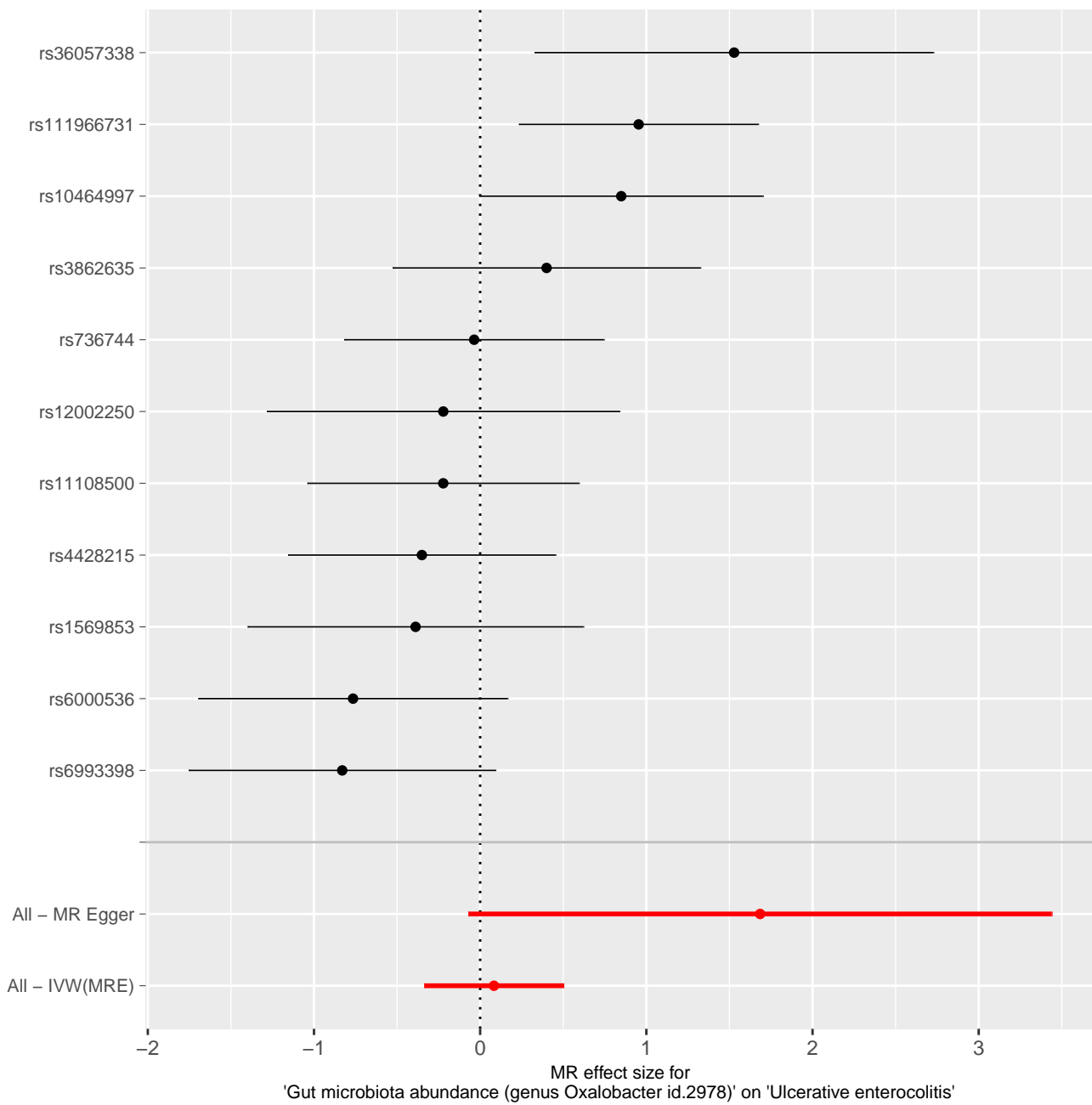


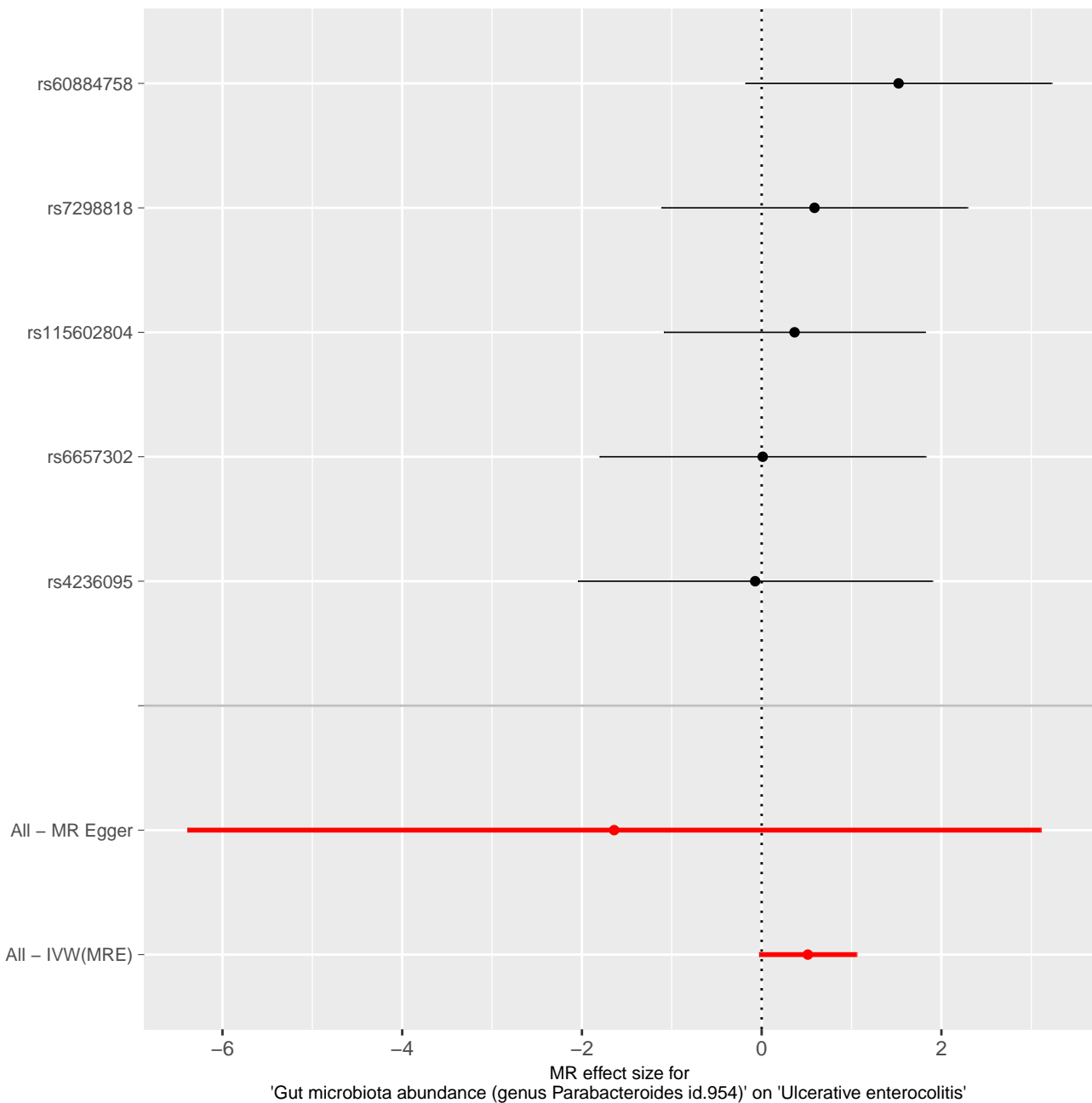


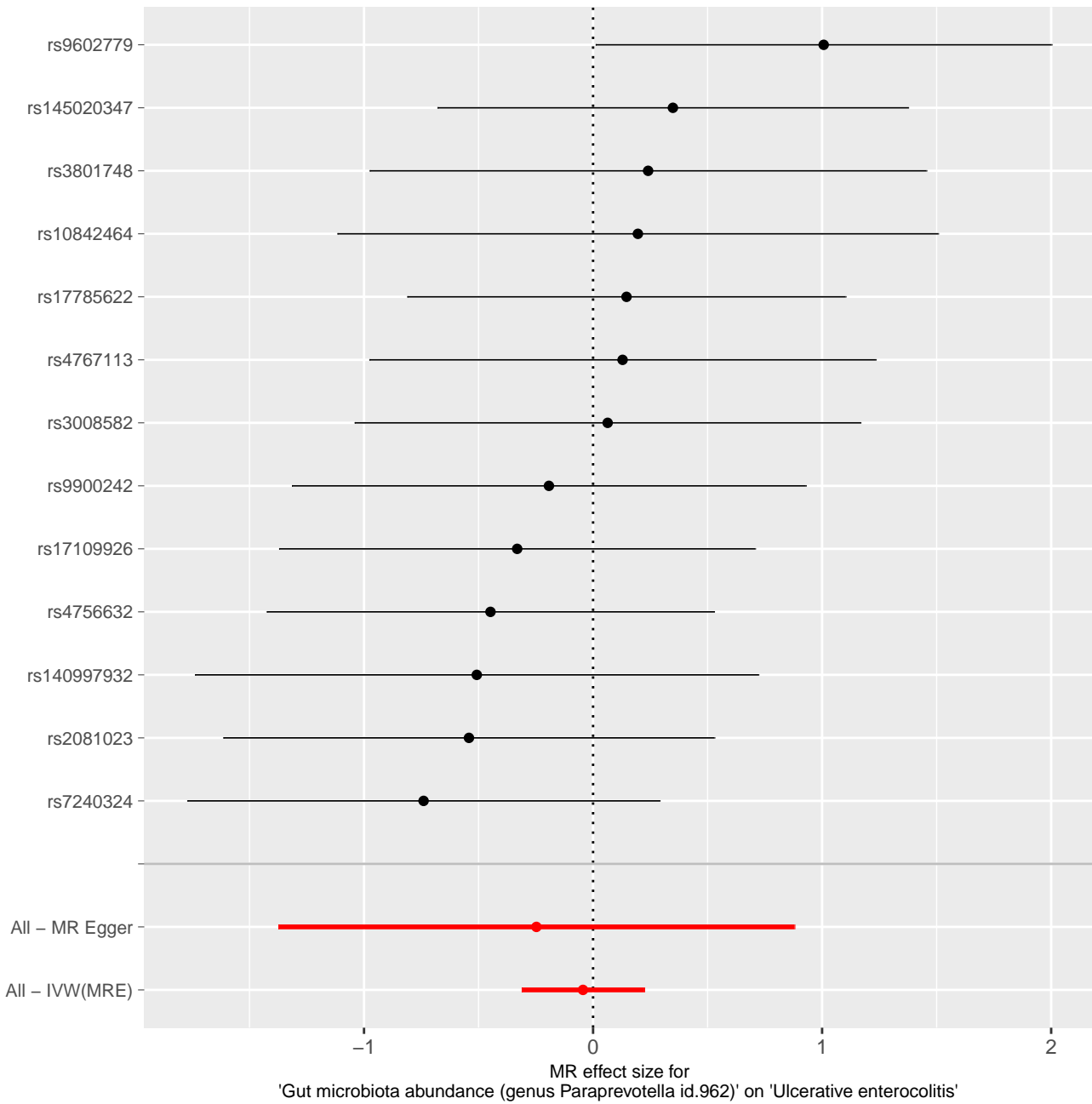


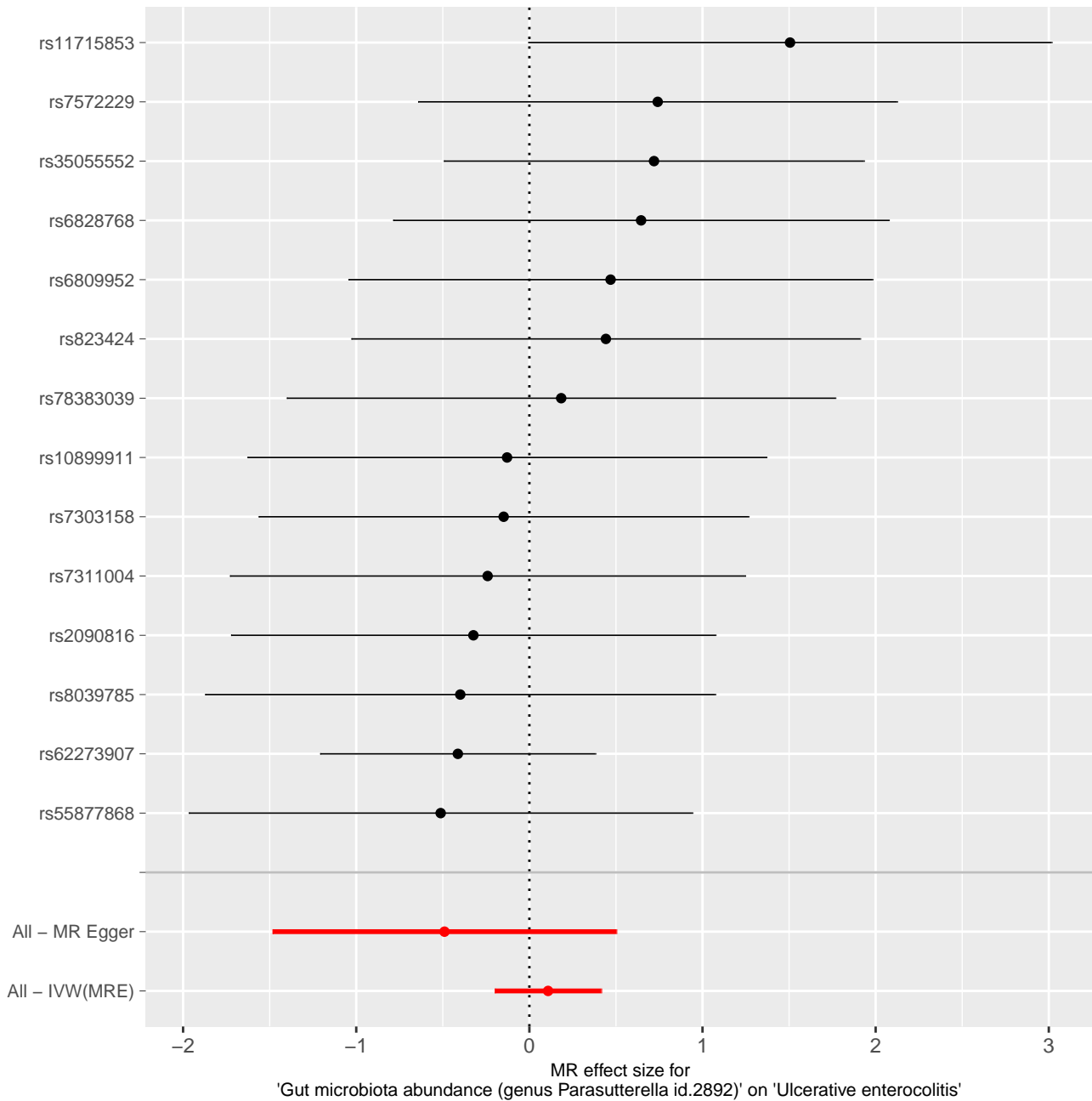


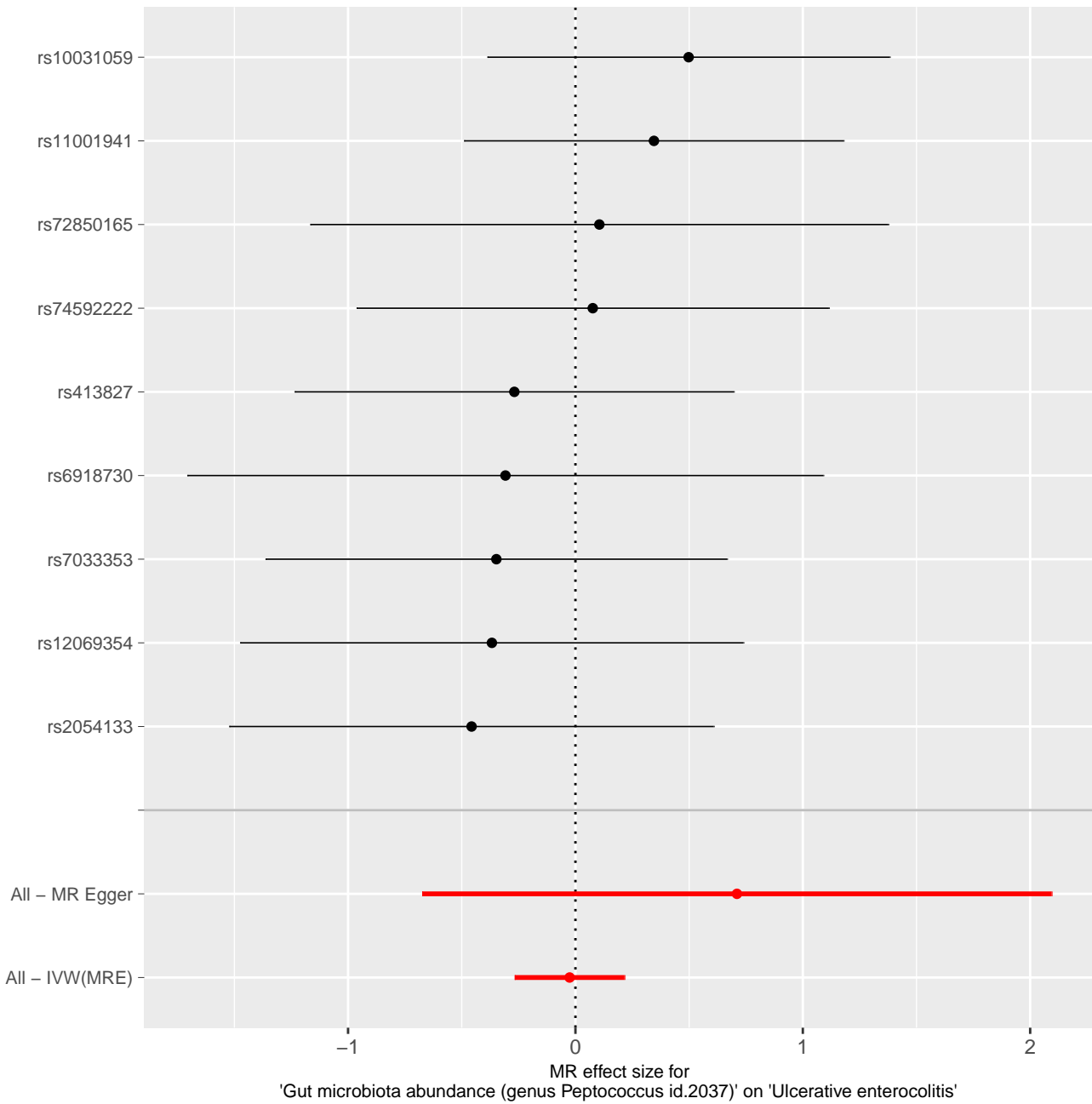
Batch 1012 : Gut microbiota abundance (genus Oxalobacter id.2978) on Ulcerative enterocolitis

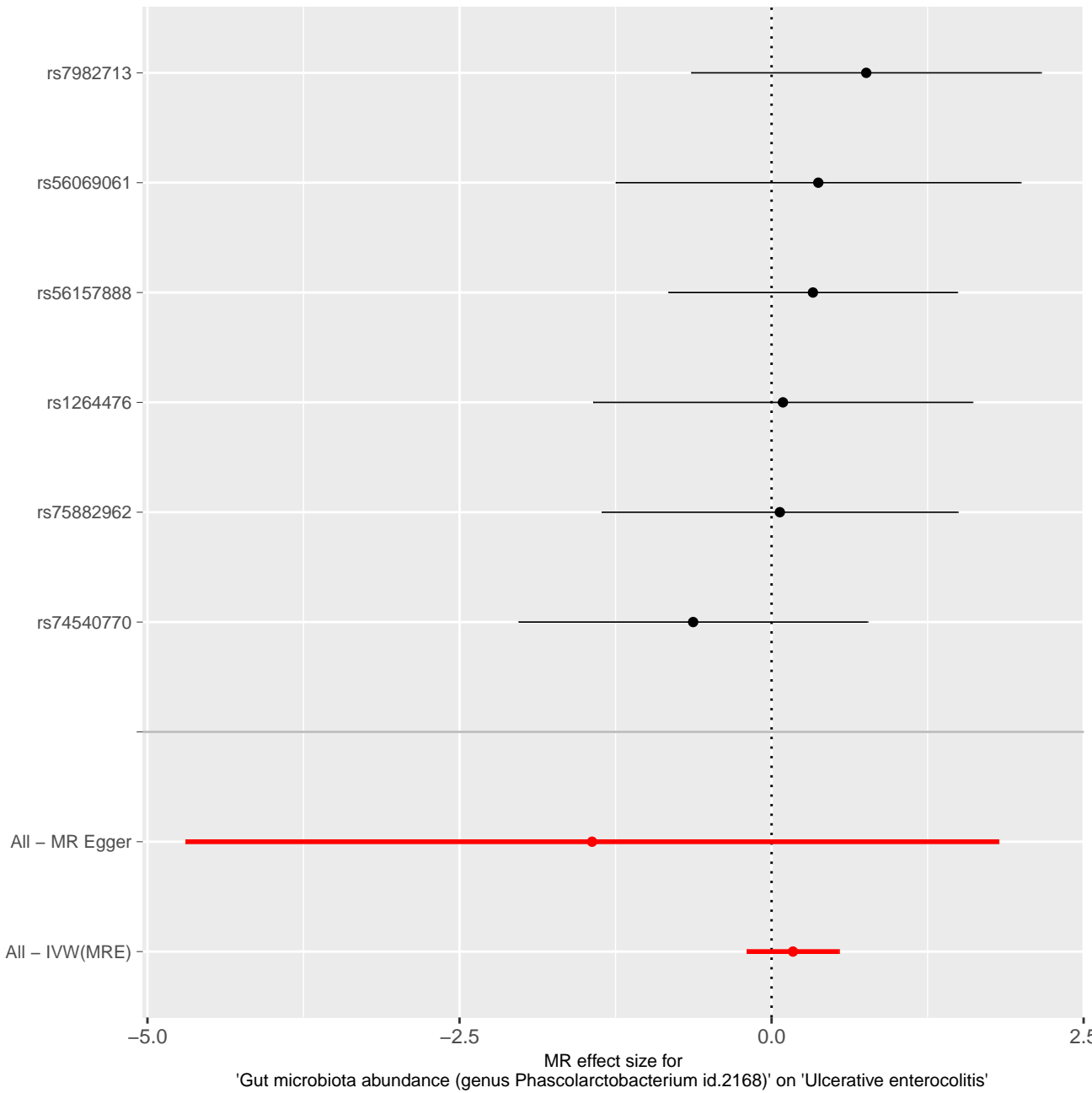




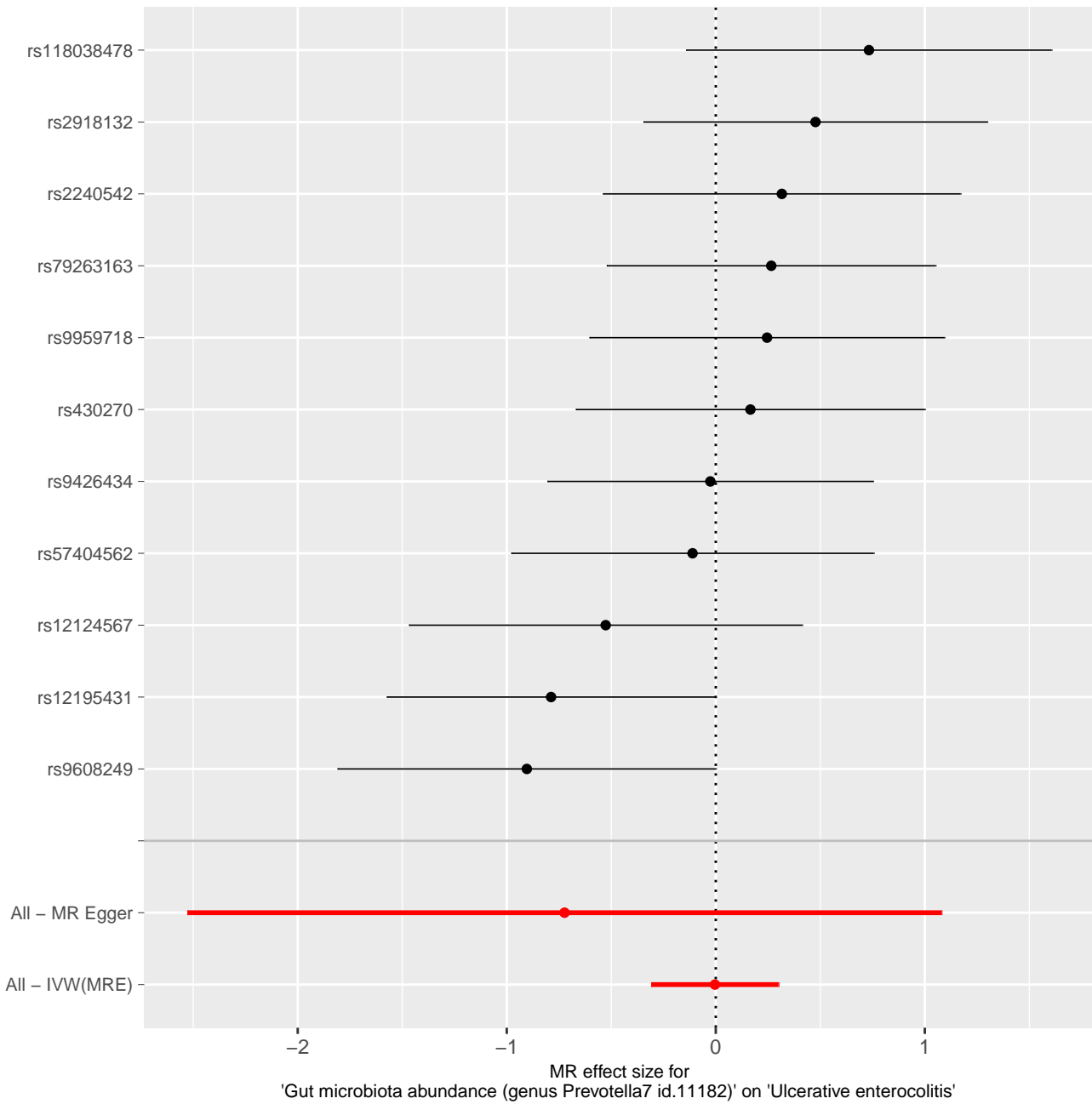


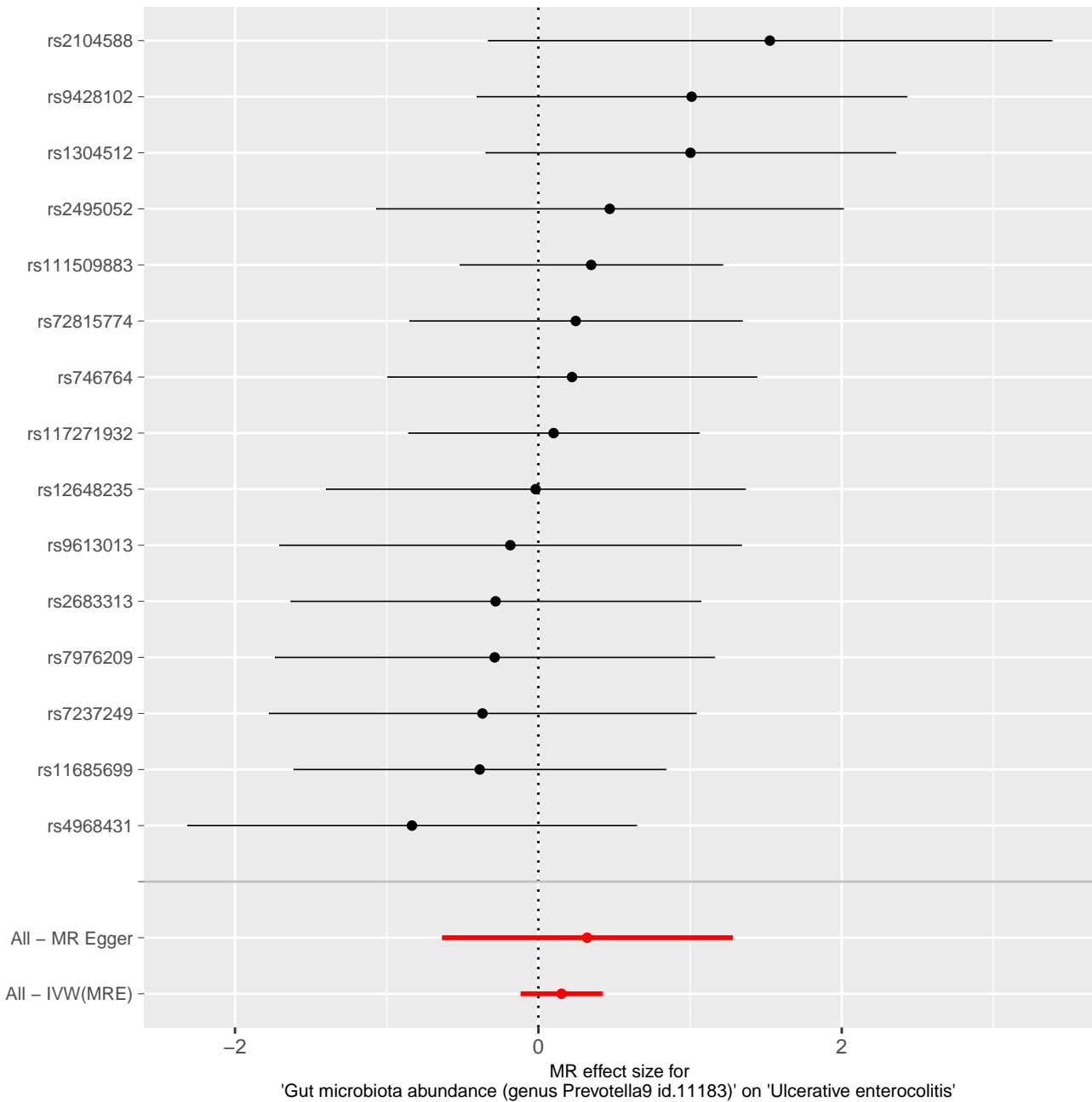


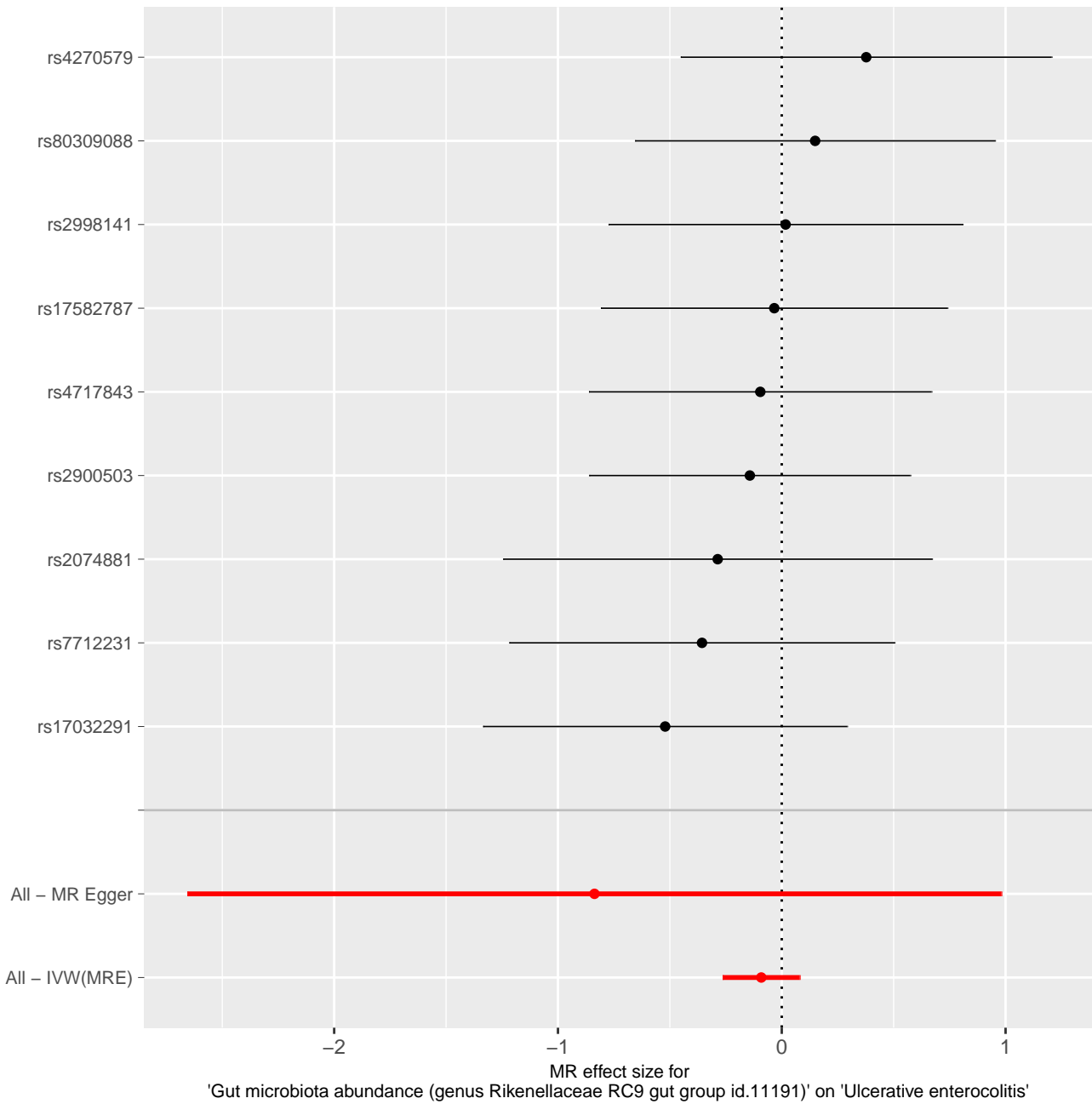


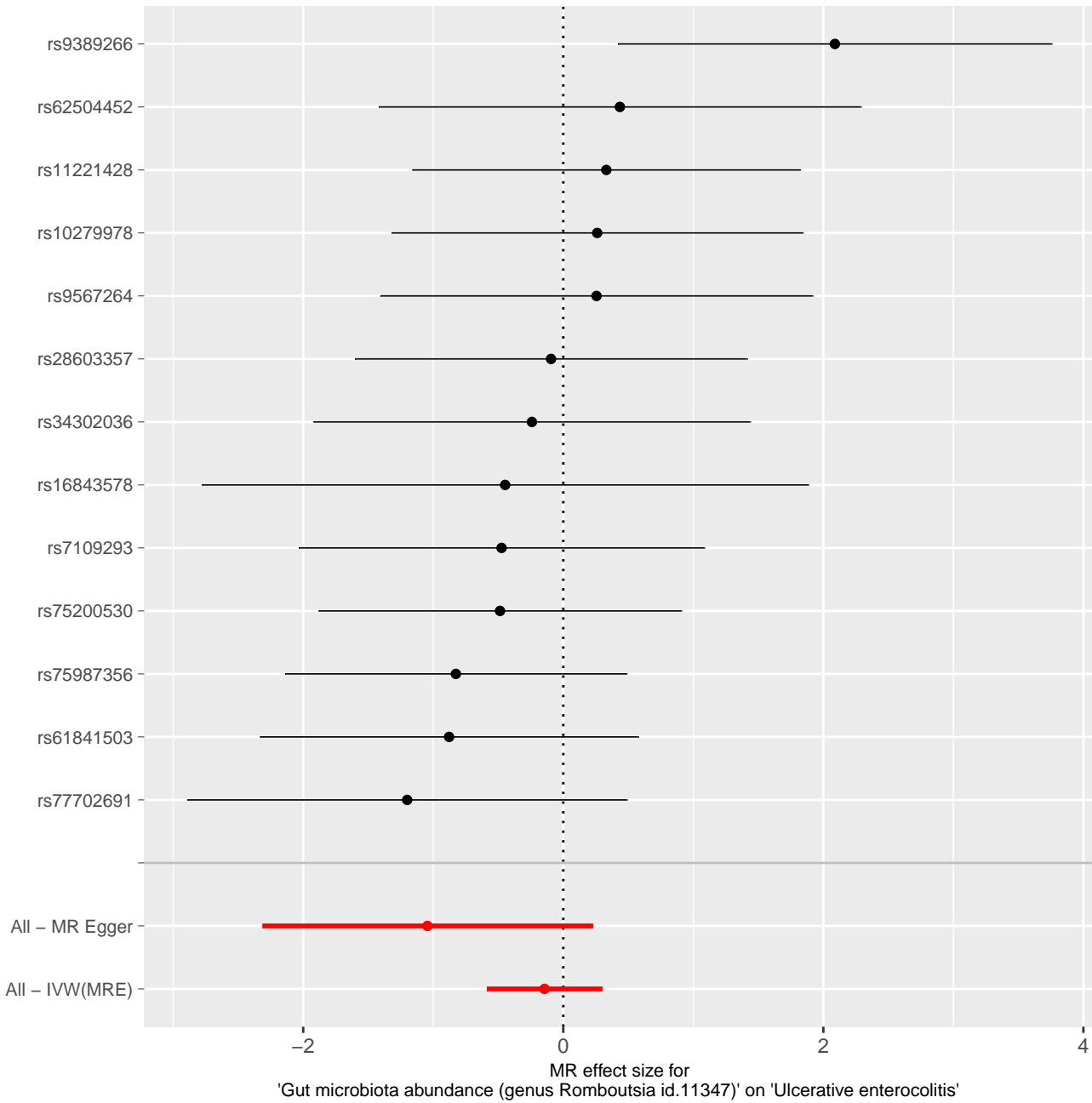


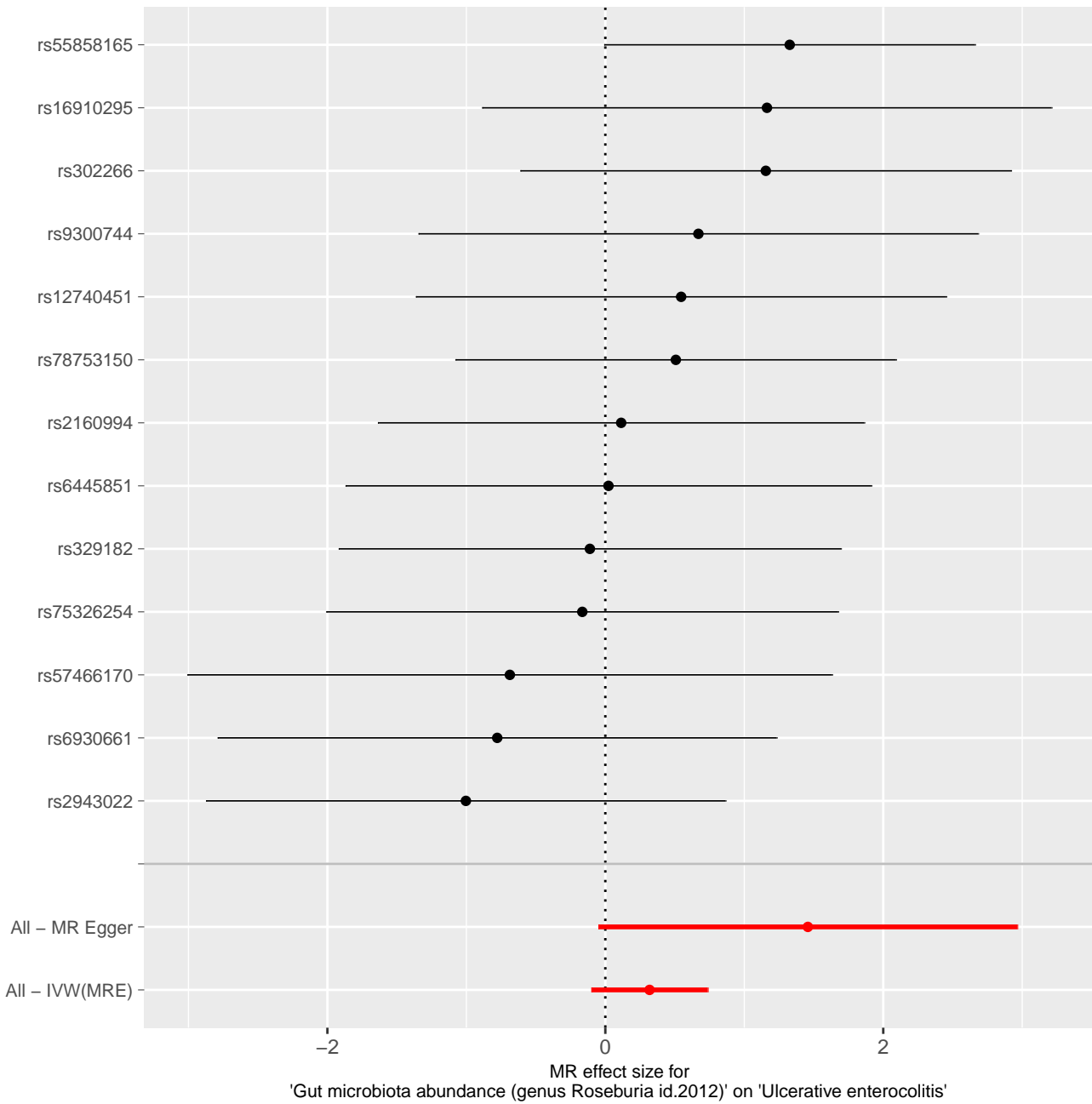
Batch 1018 : Gut microbiota abundance (genus Prevotella7 id.11182) on Ulcerative enterocolitis

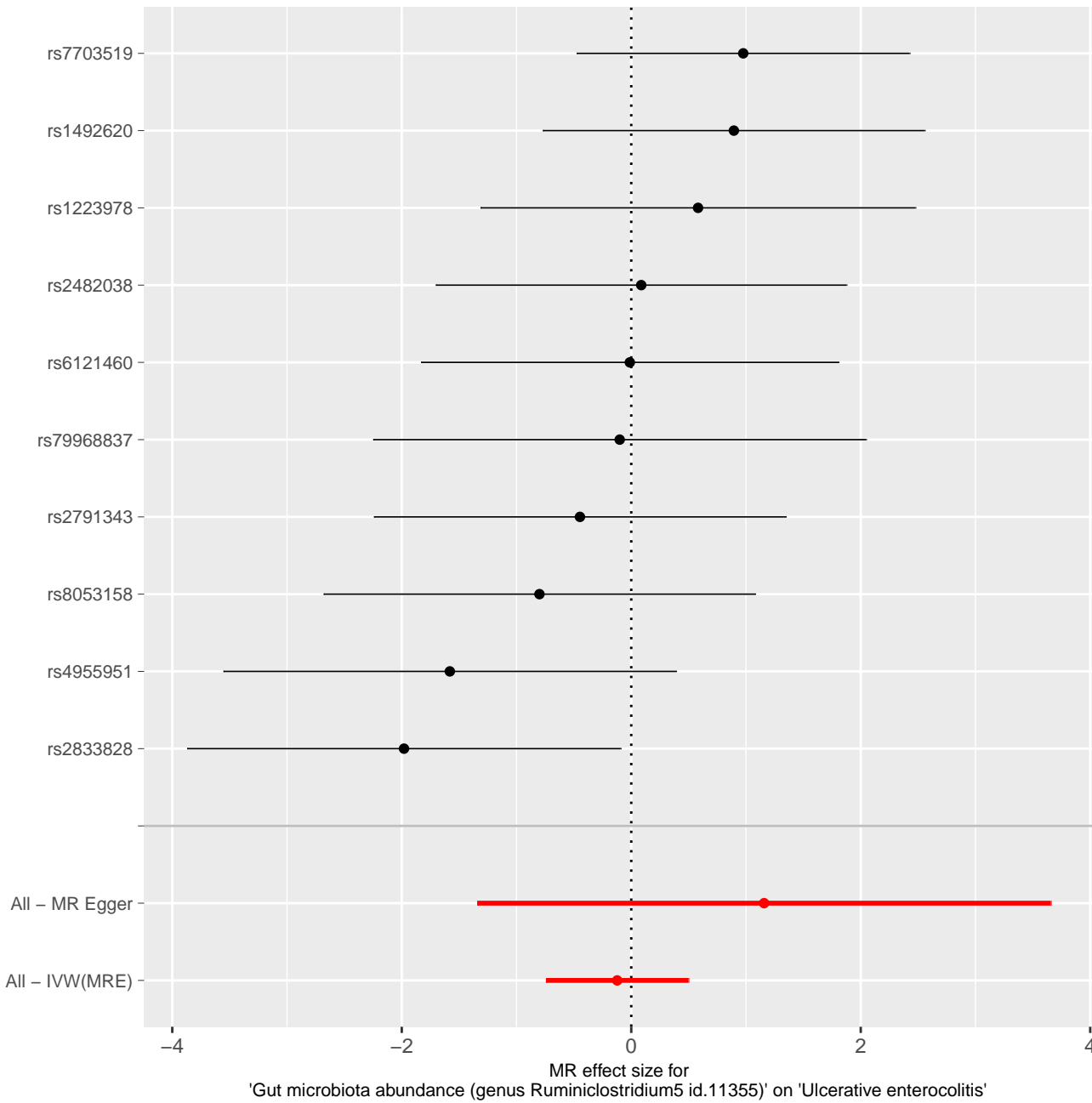


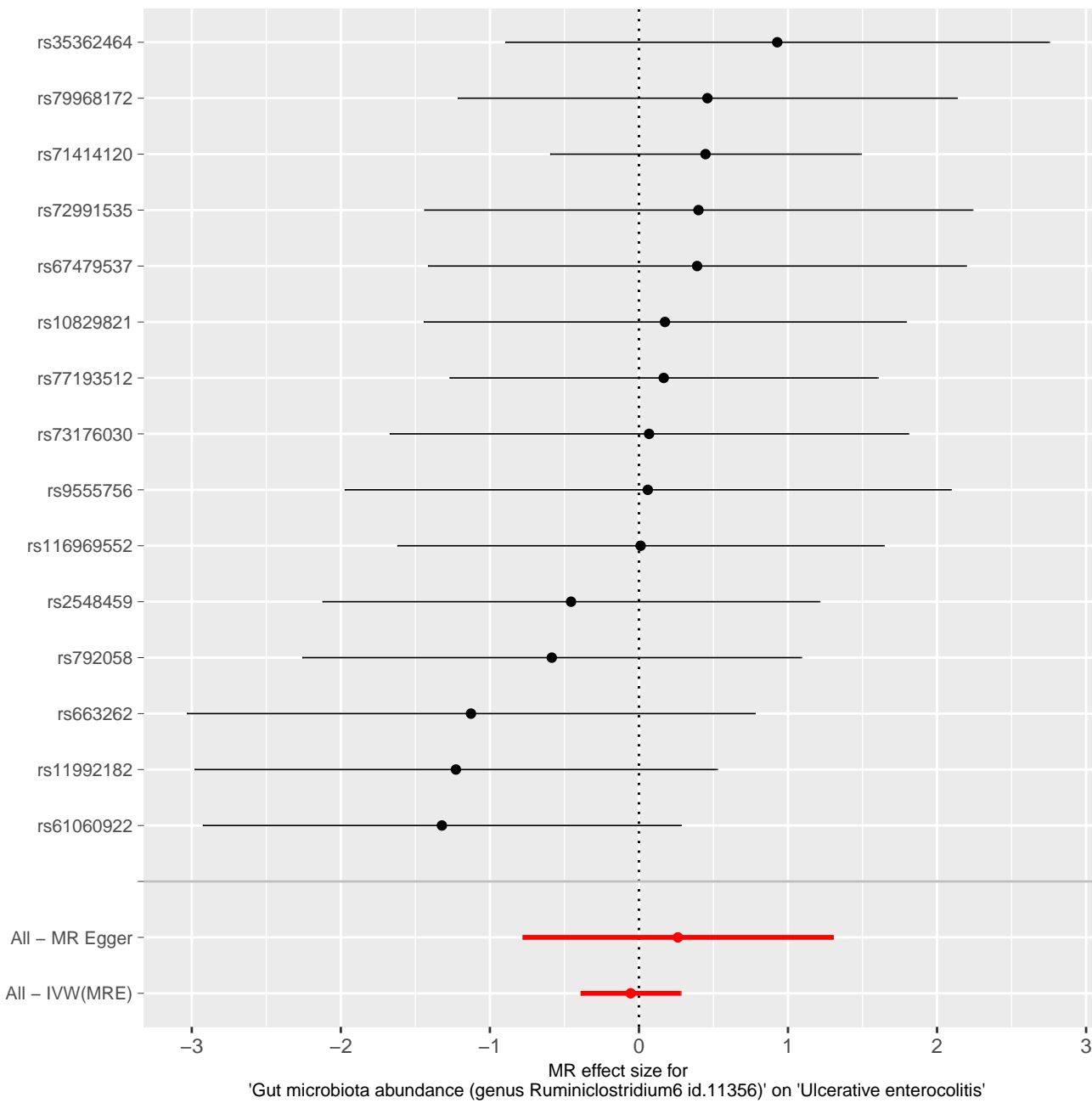


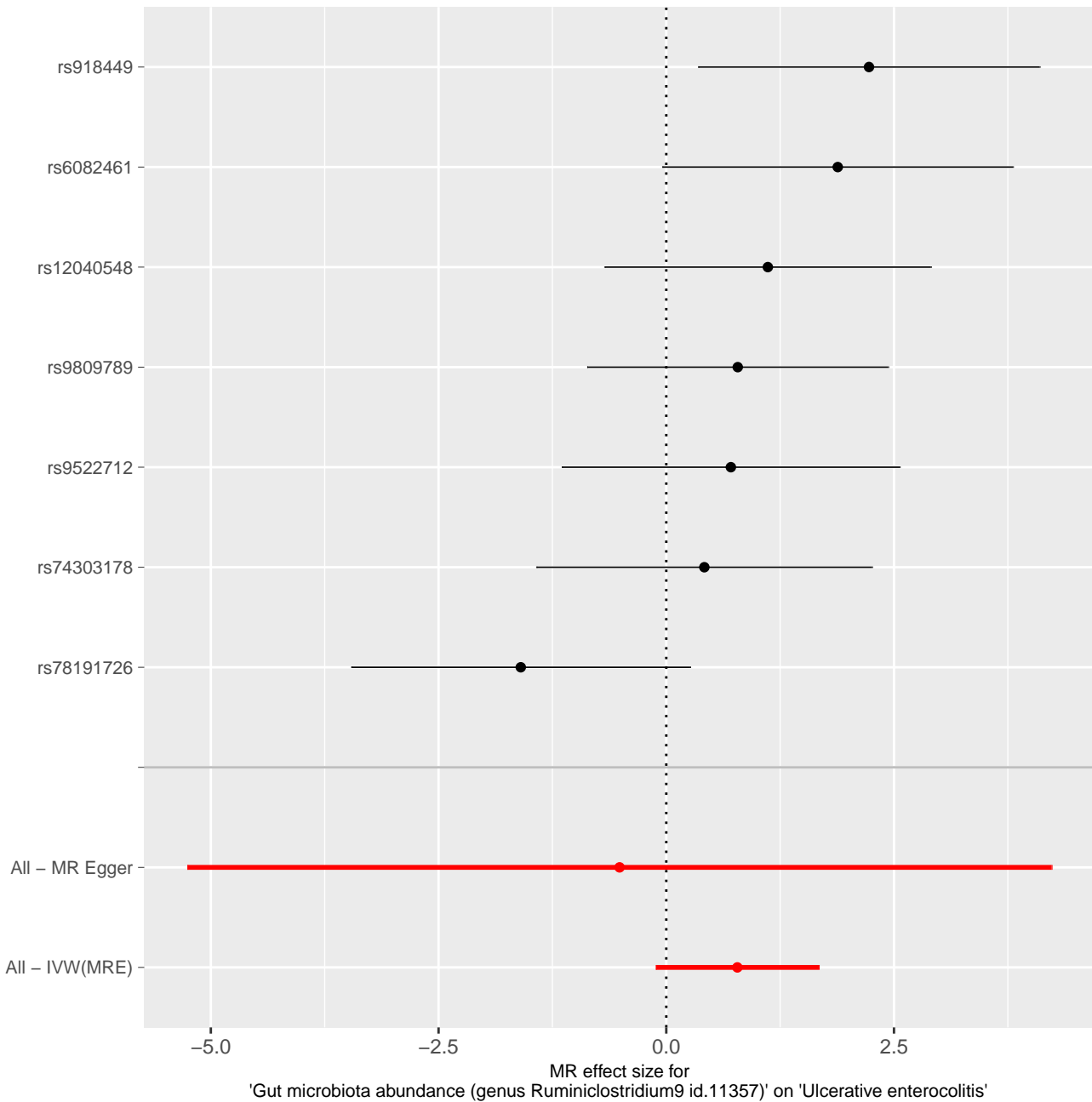


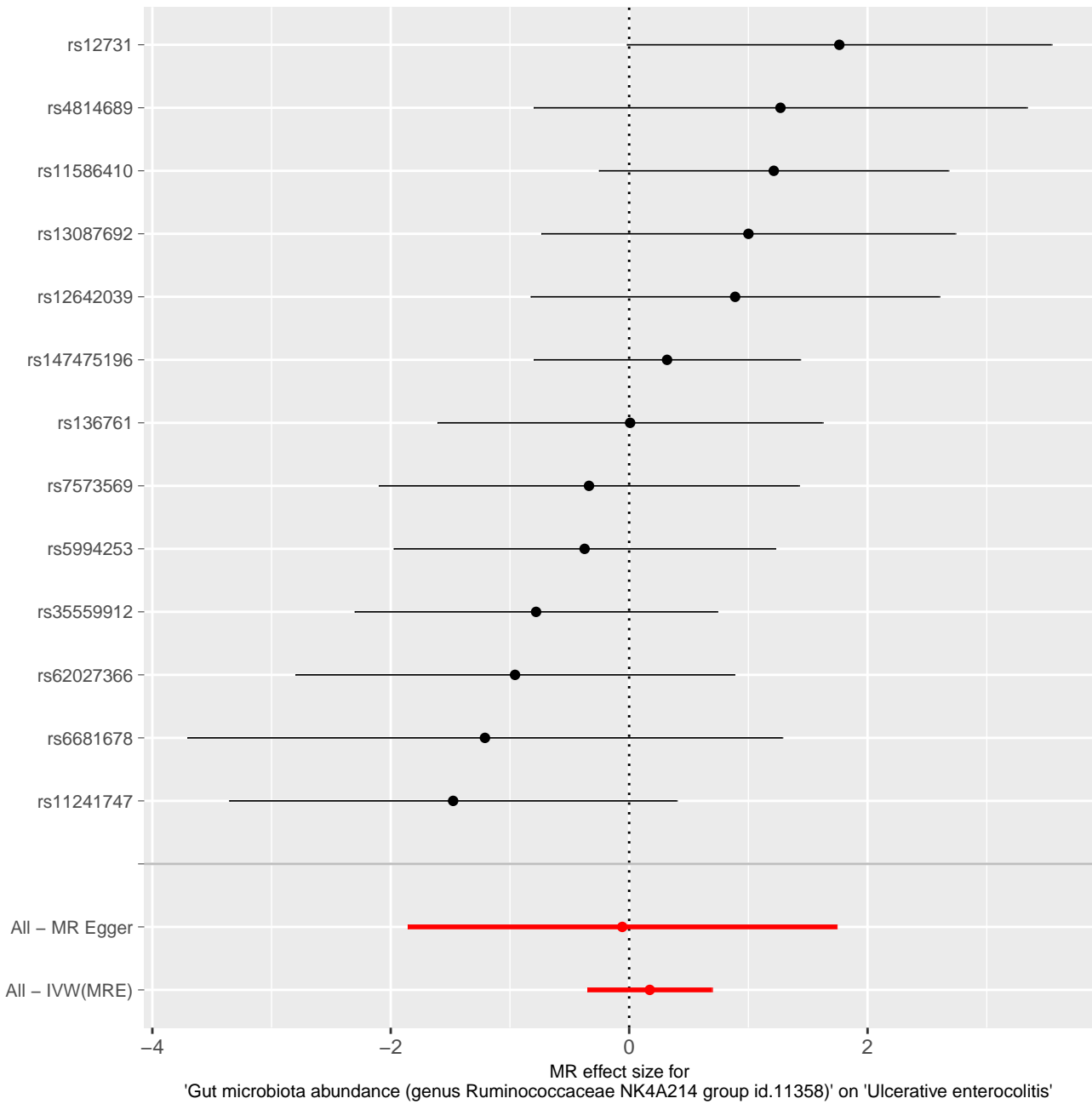


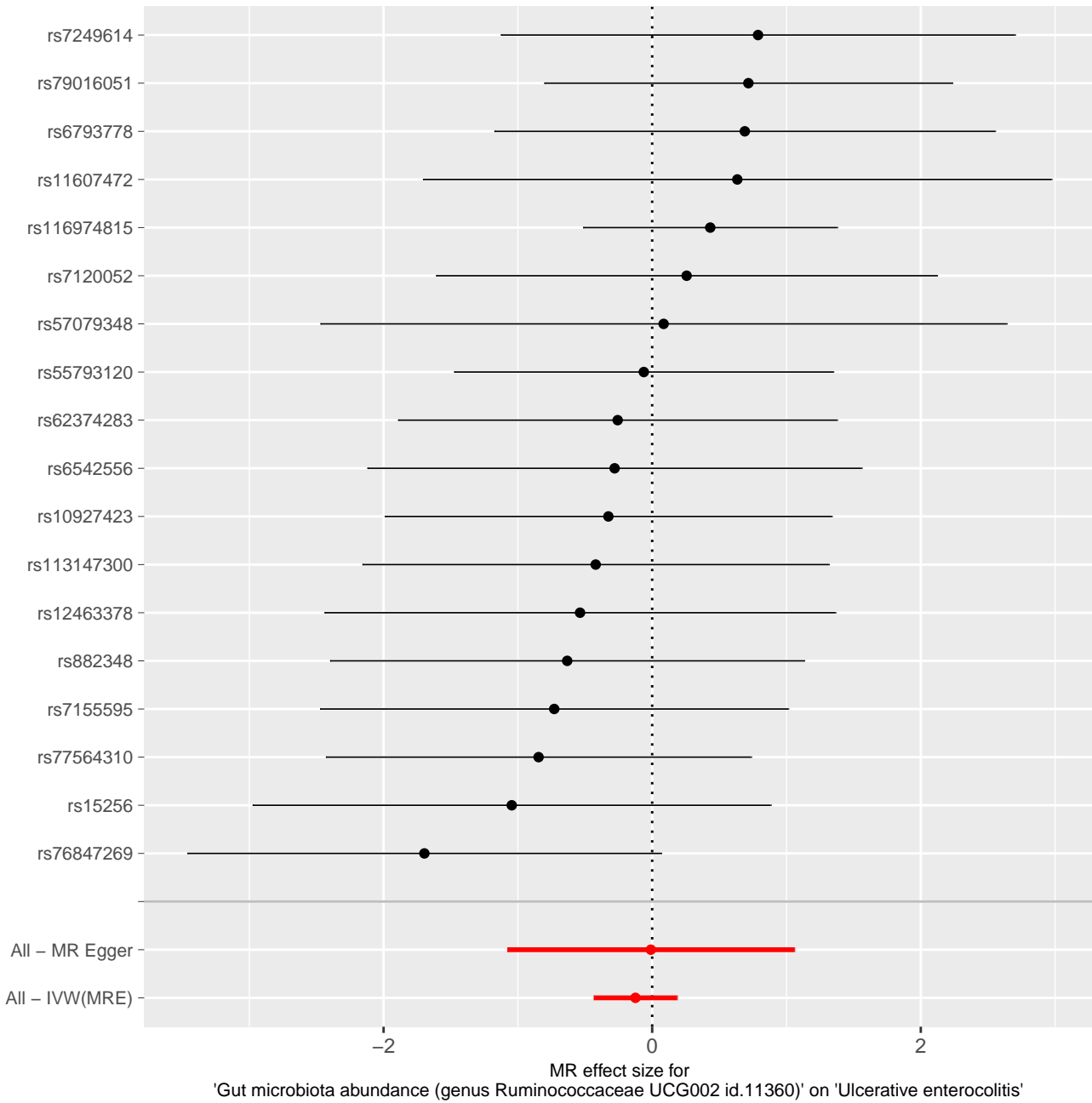


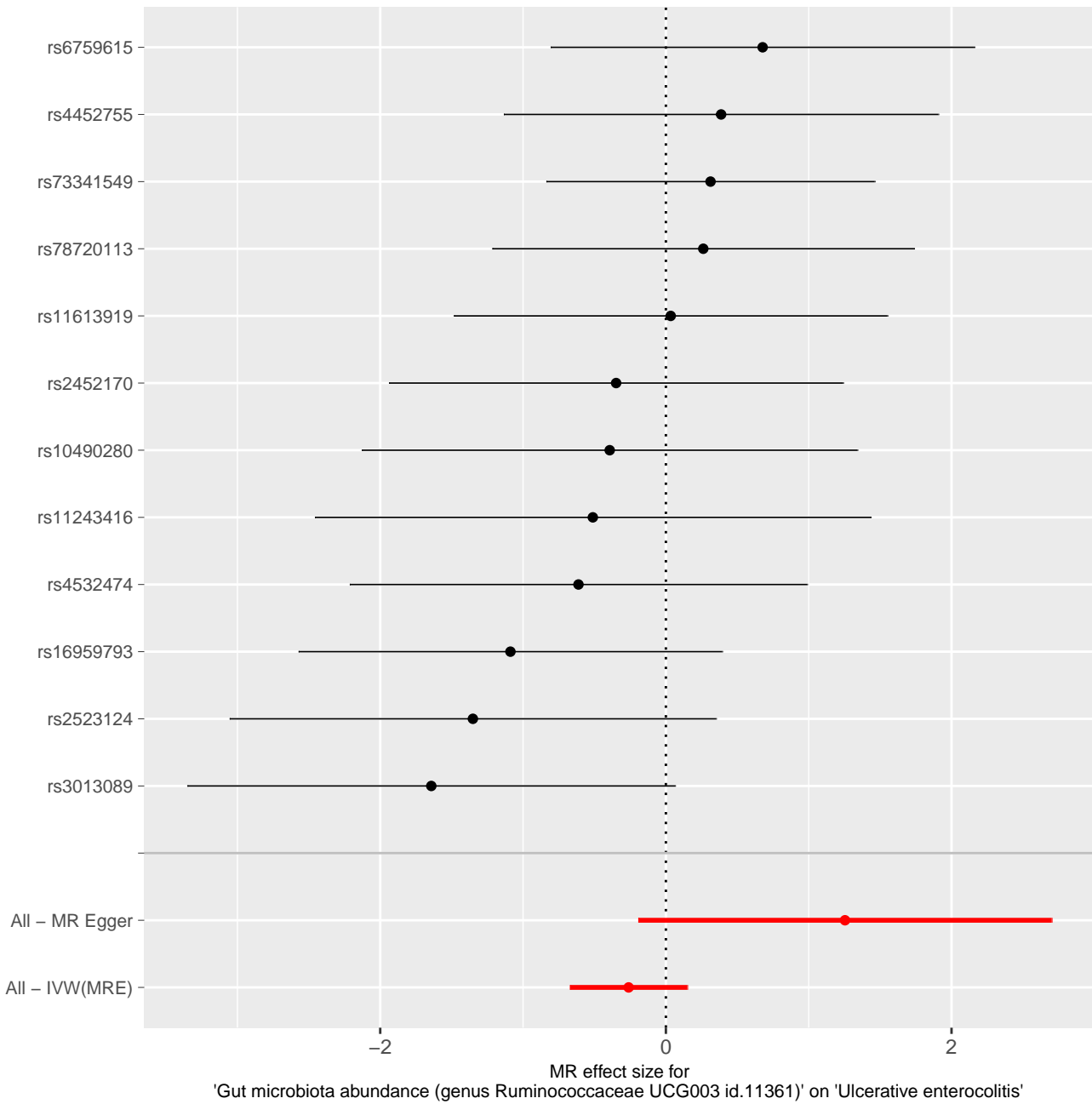


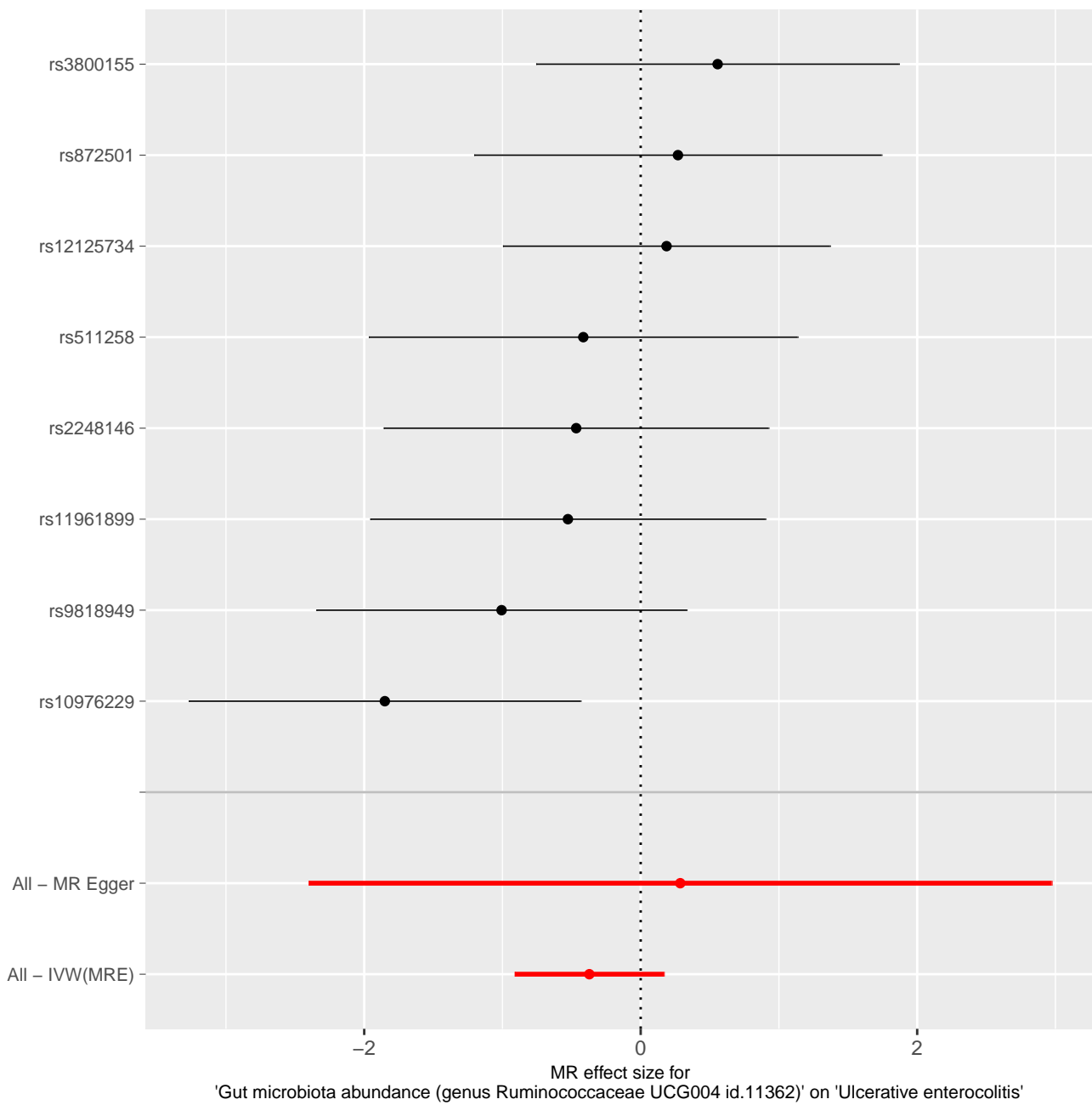




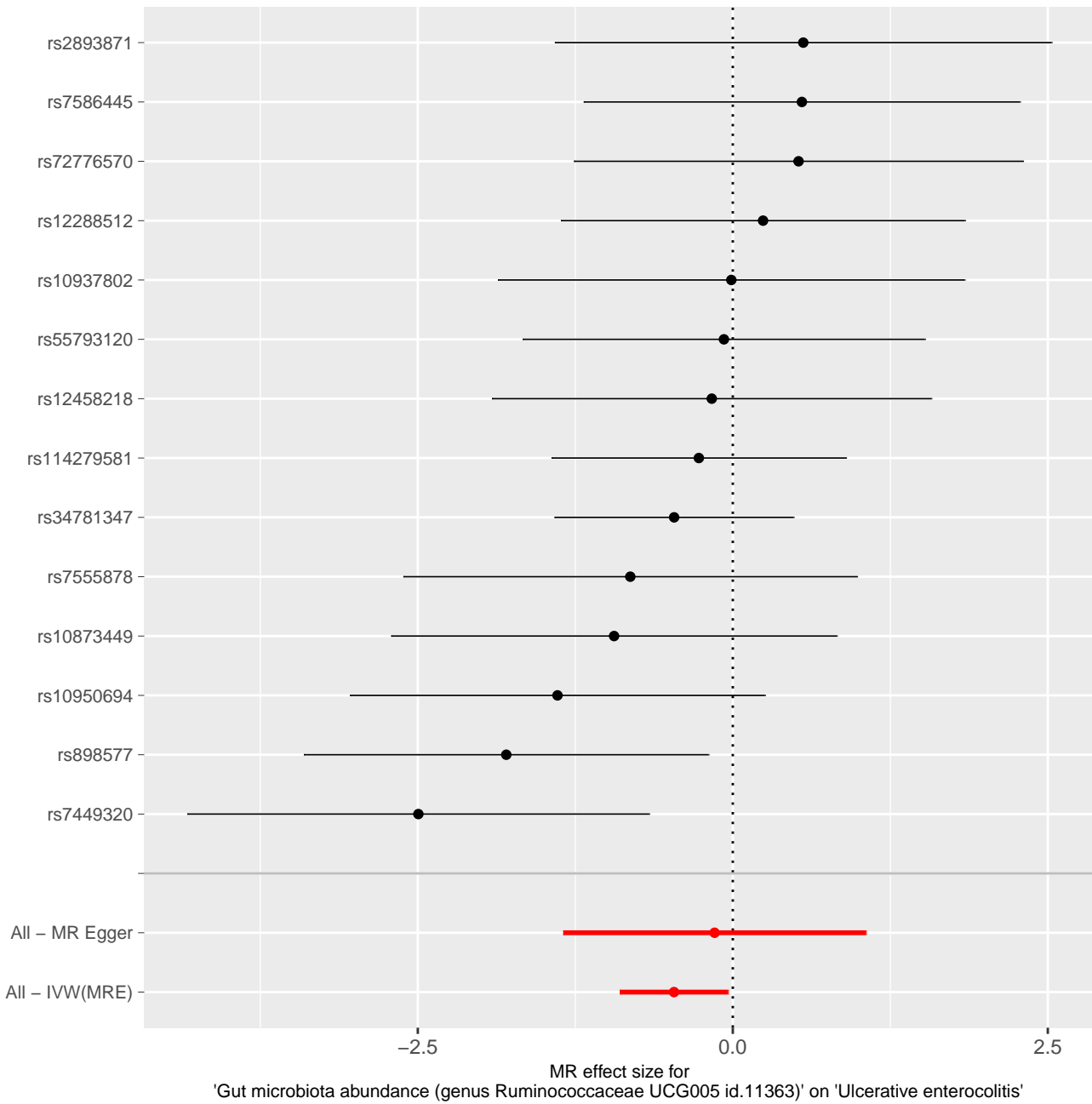


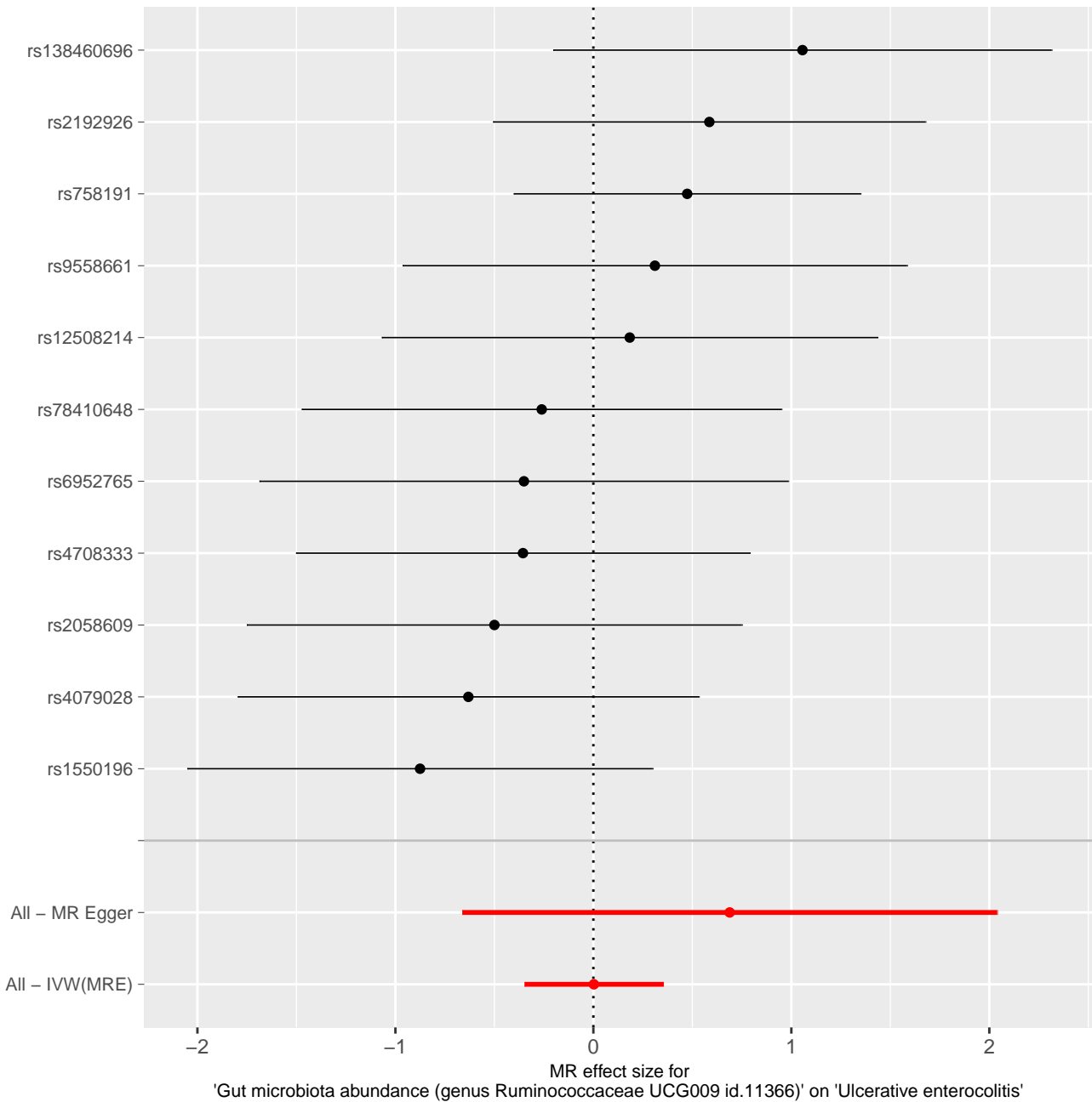


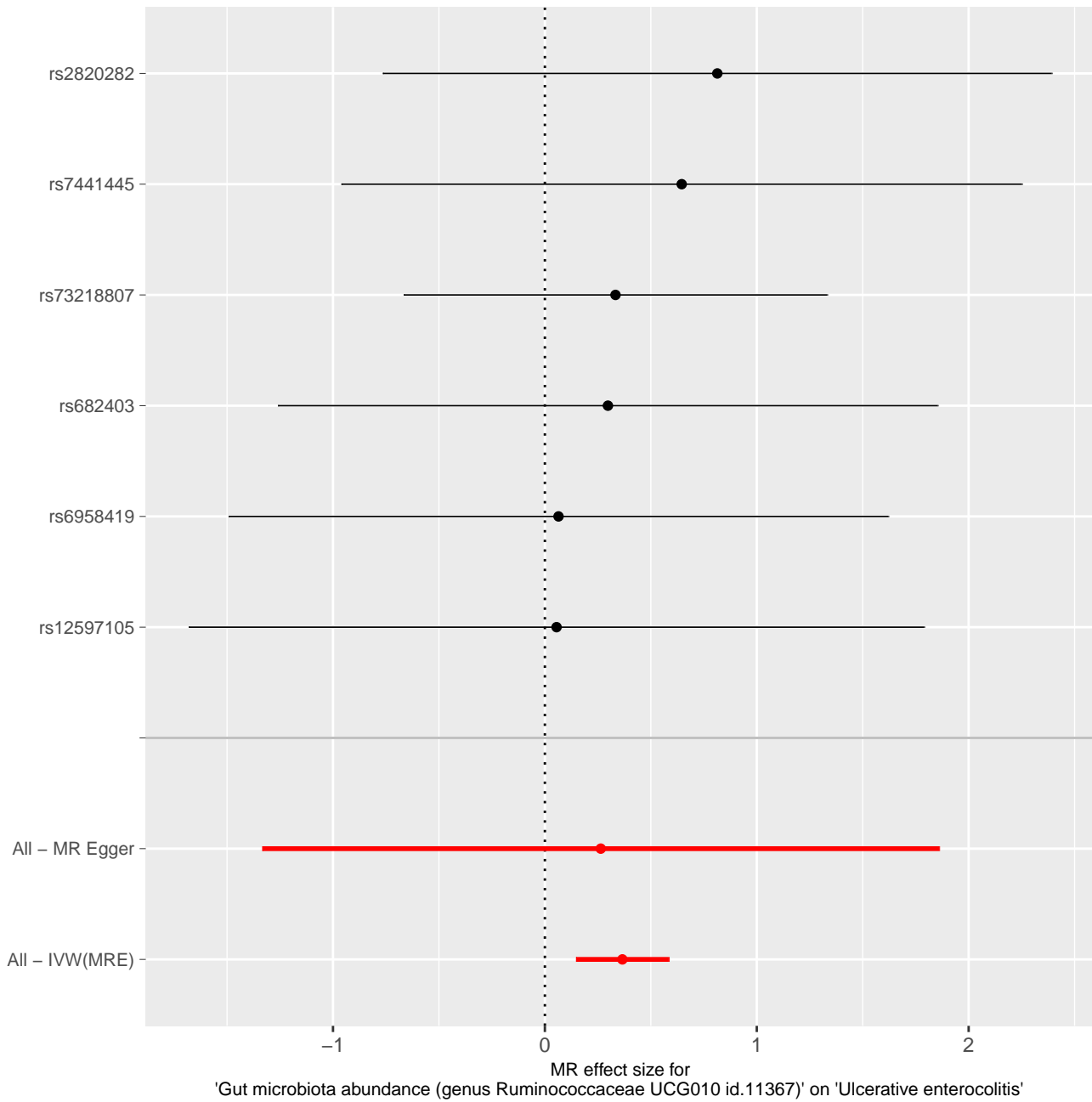


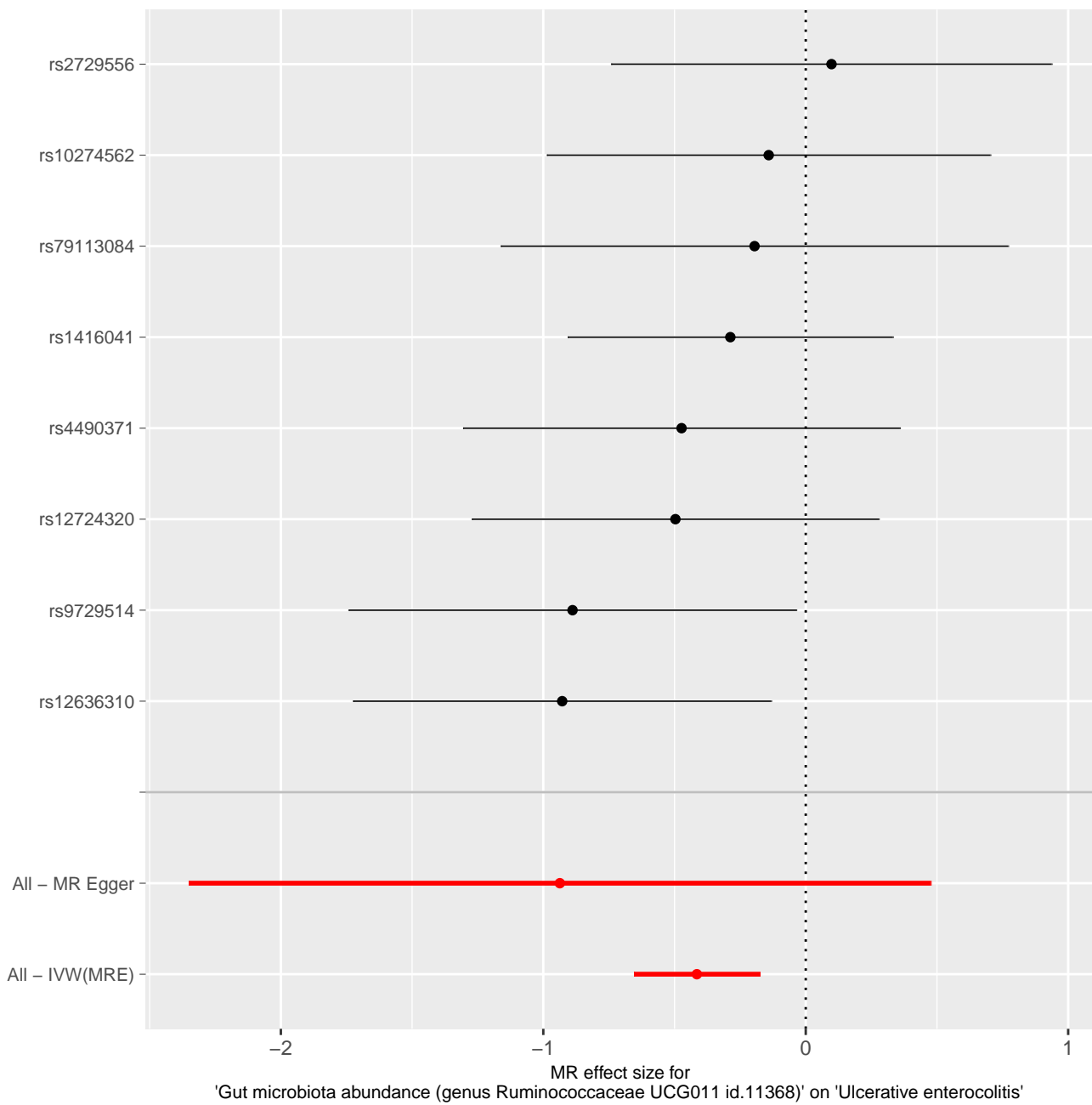


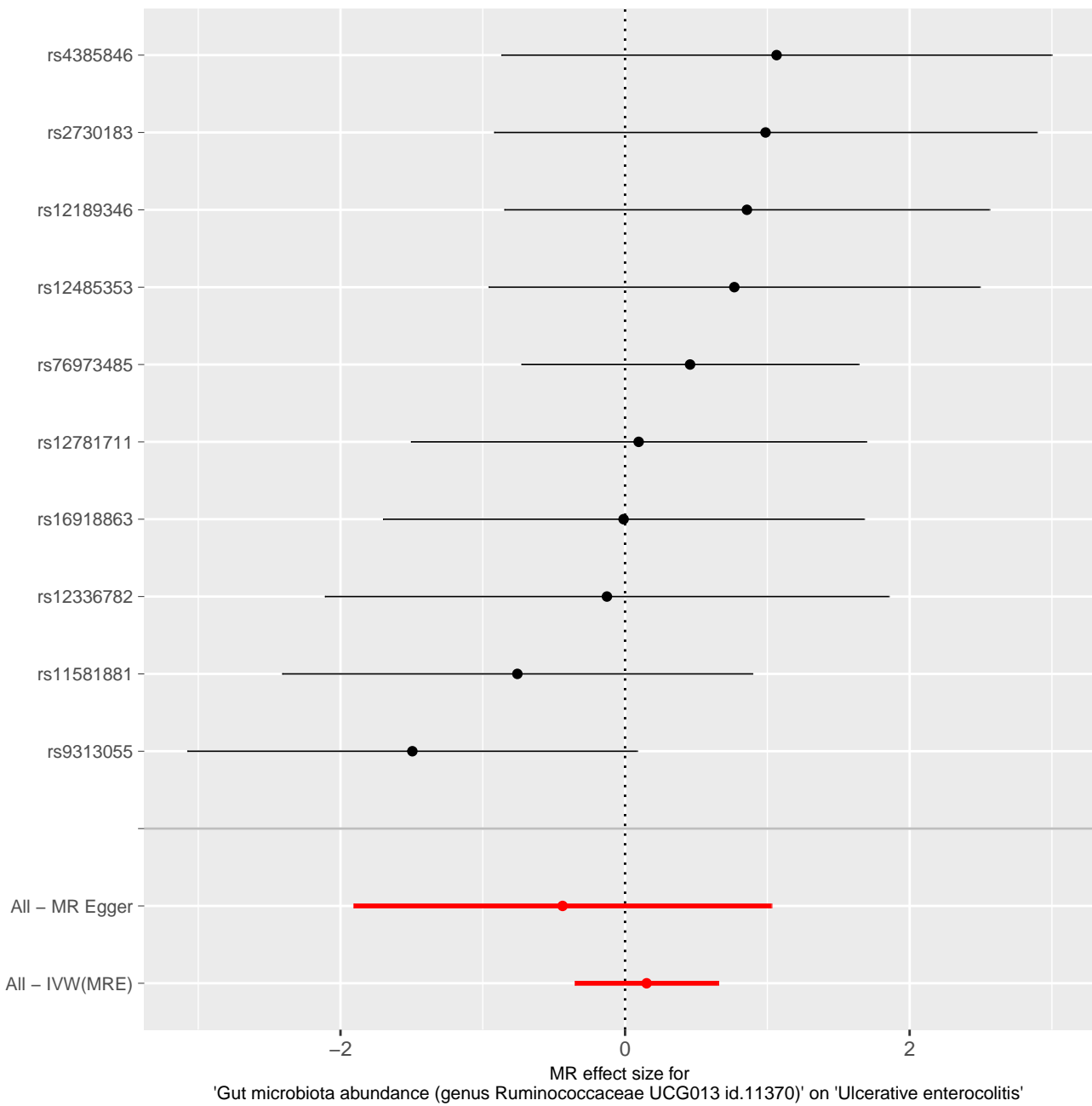
MR effect size for
'Gut microbiota abundance (genus Ruminococcaceae UCG004 id.11362)' on 'Ulcerative enterocolitis'

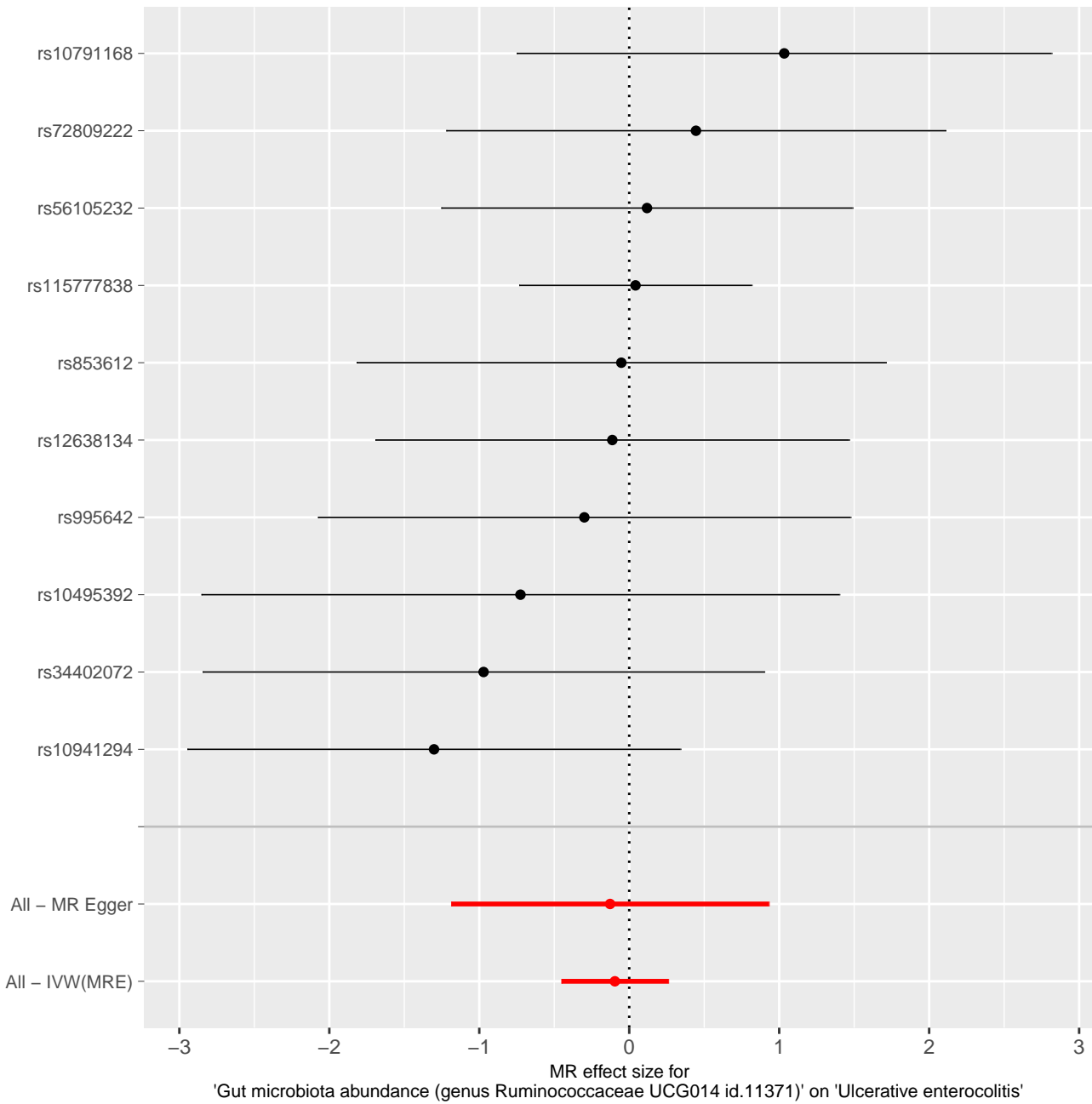


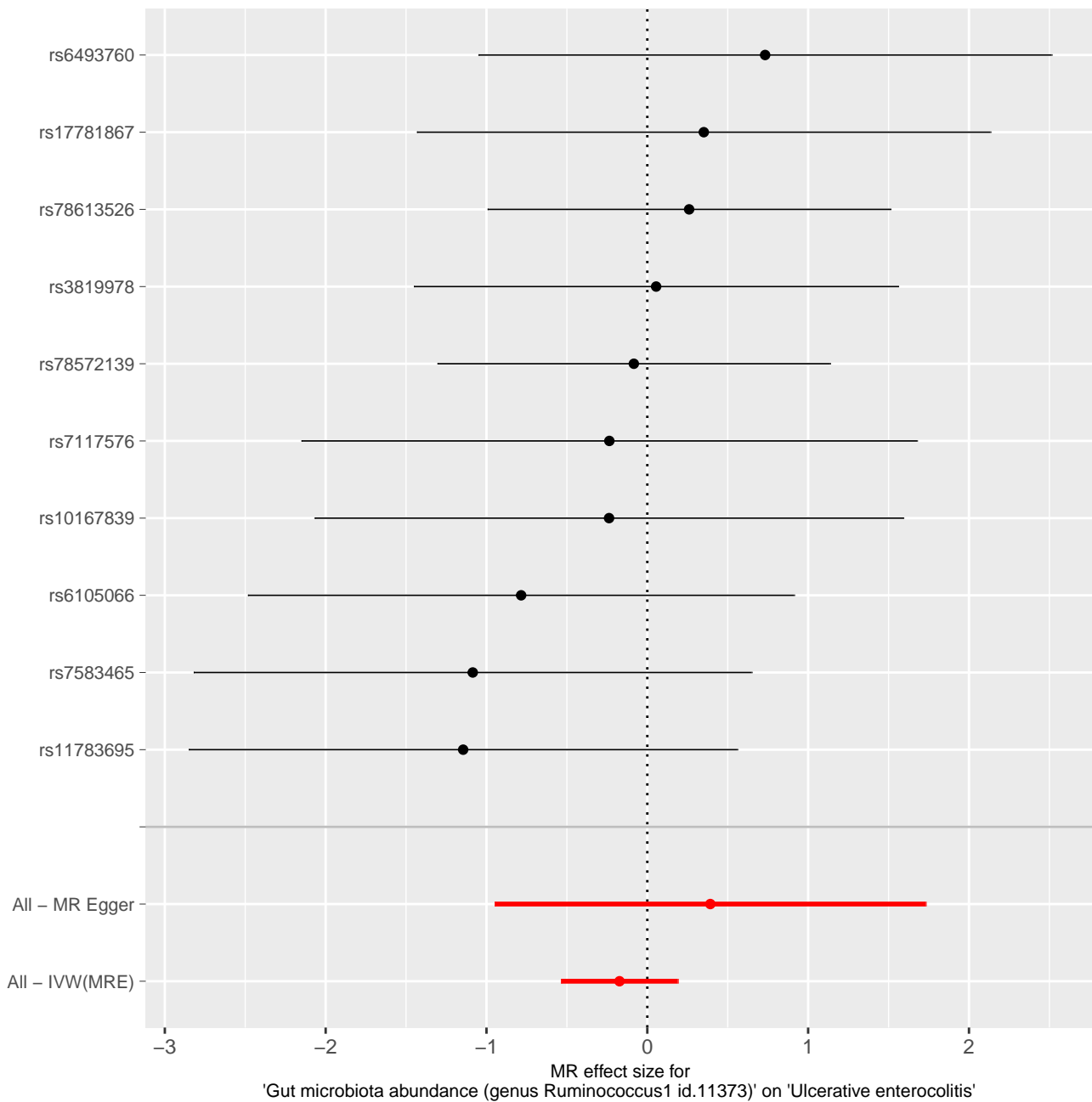


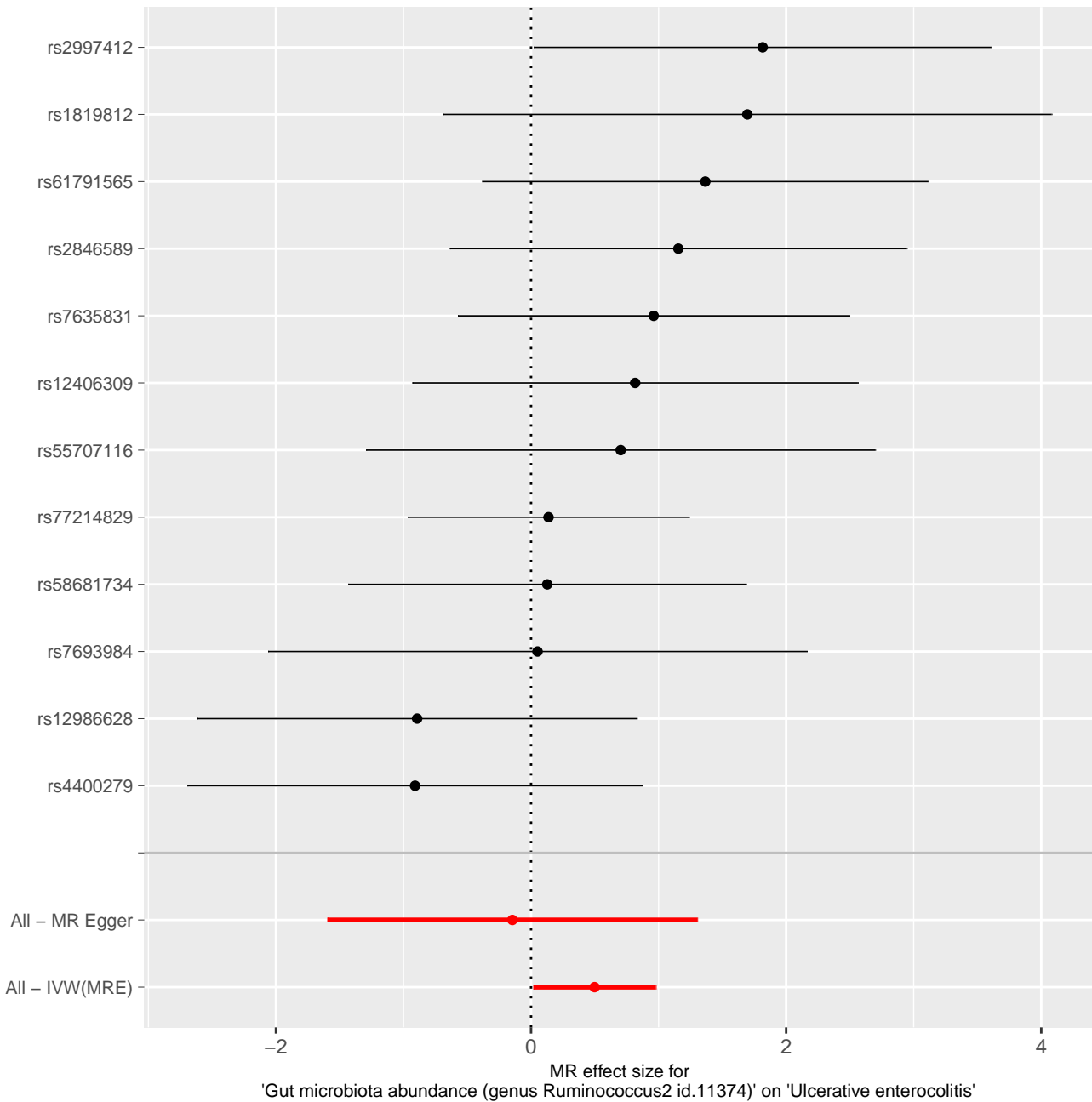


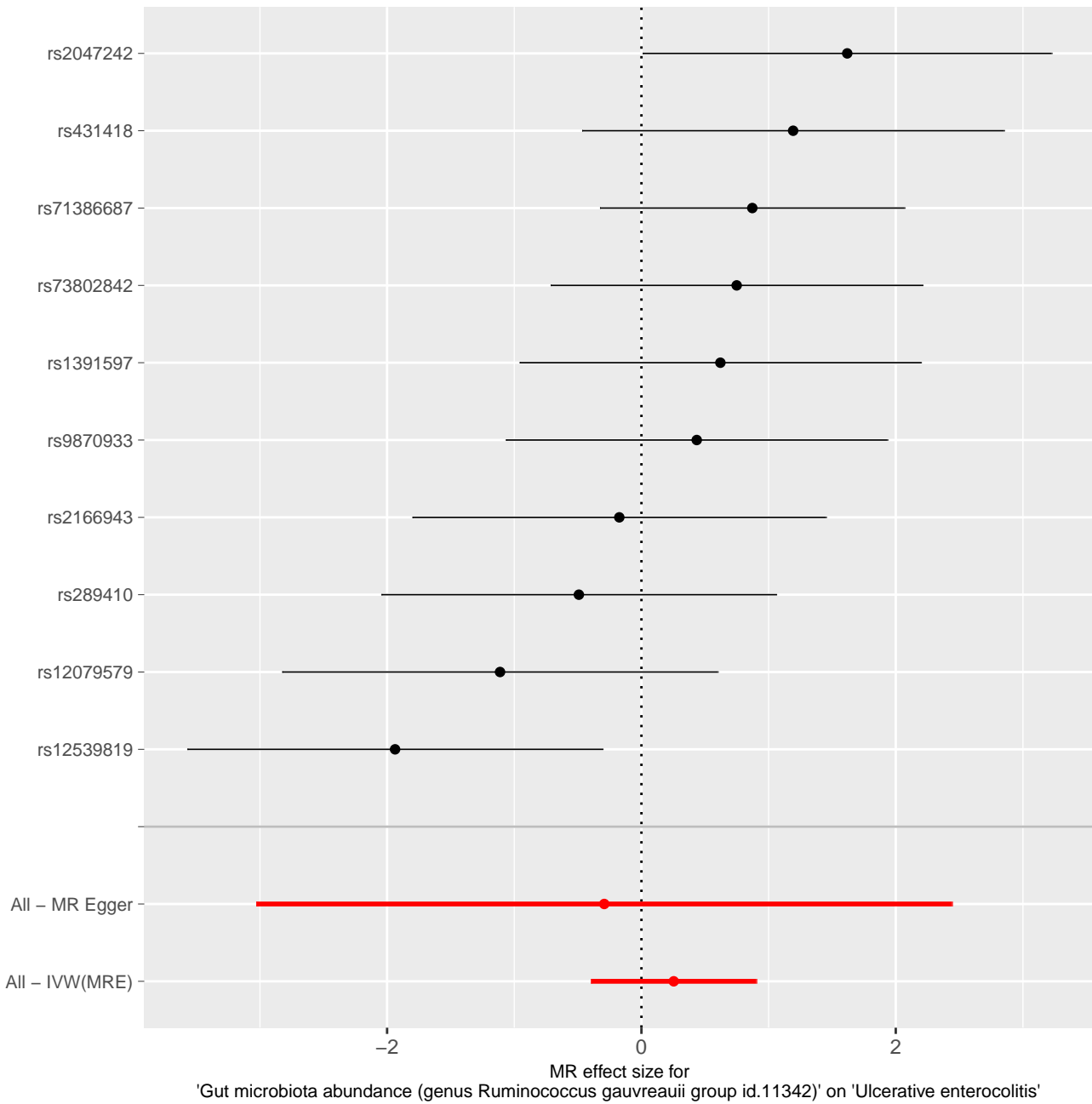


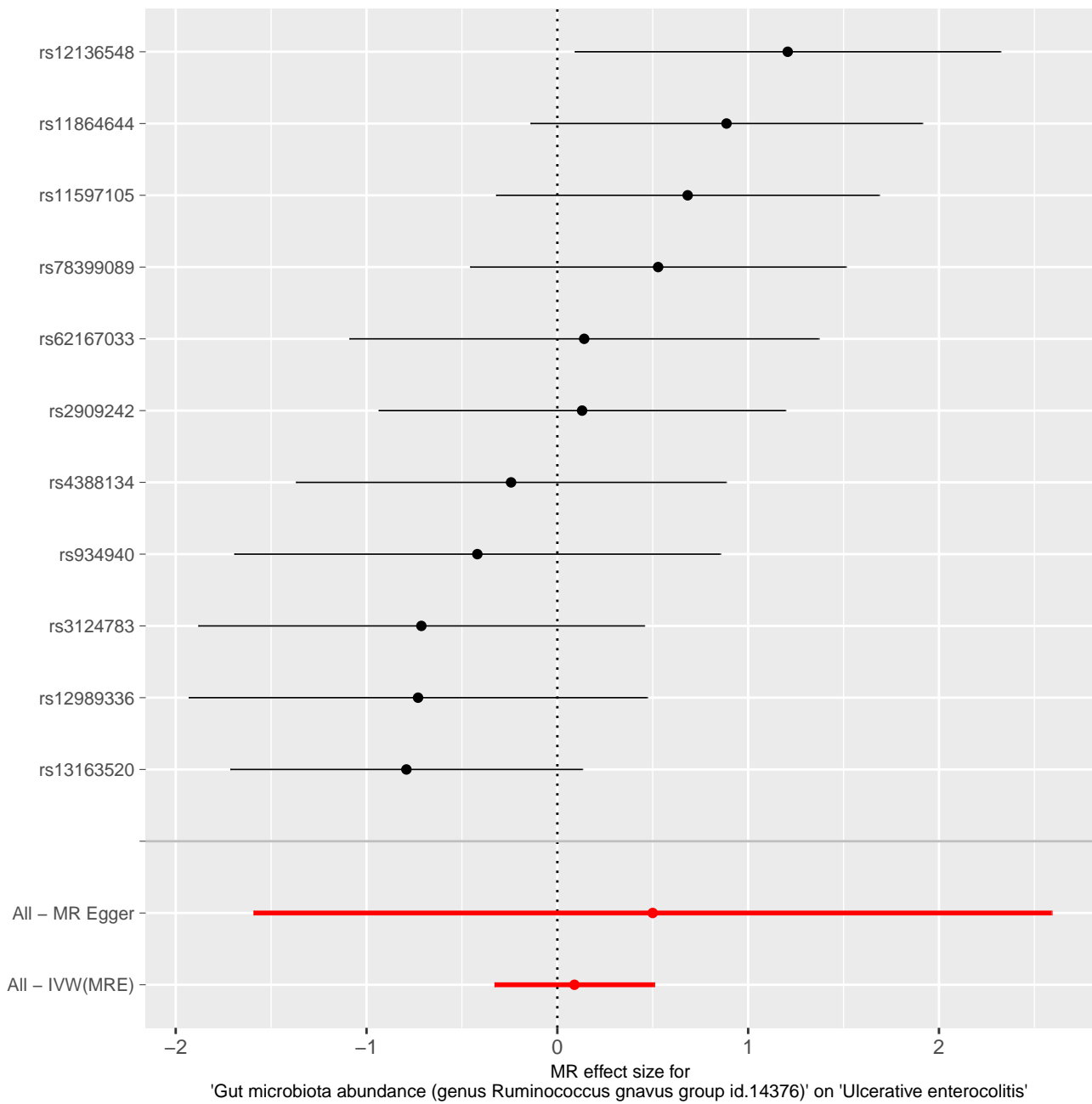


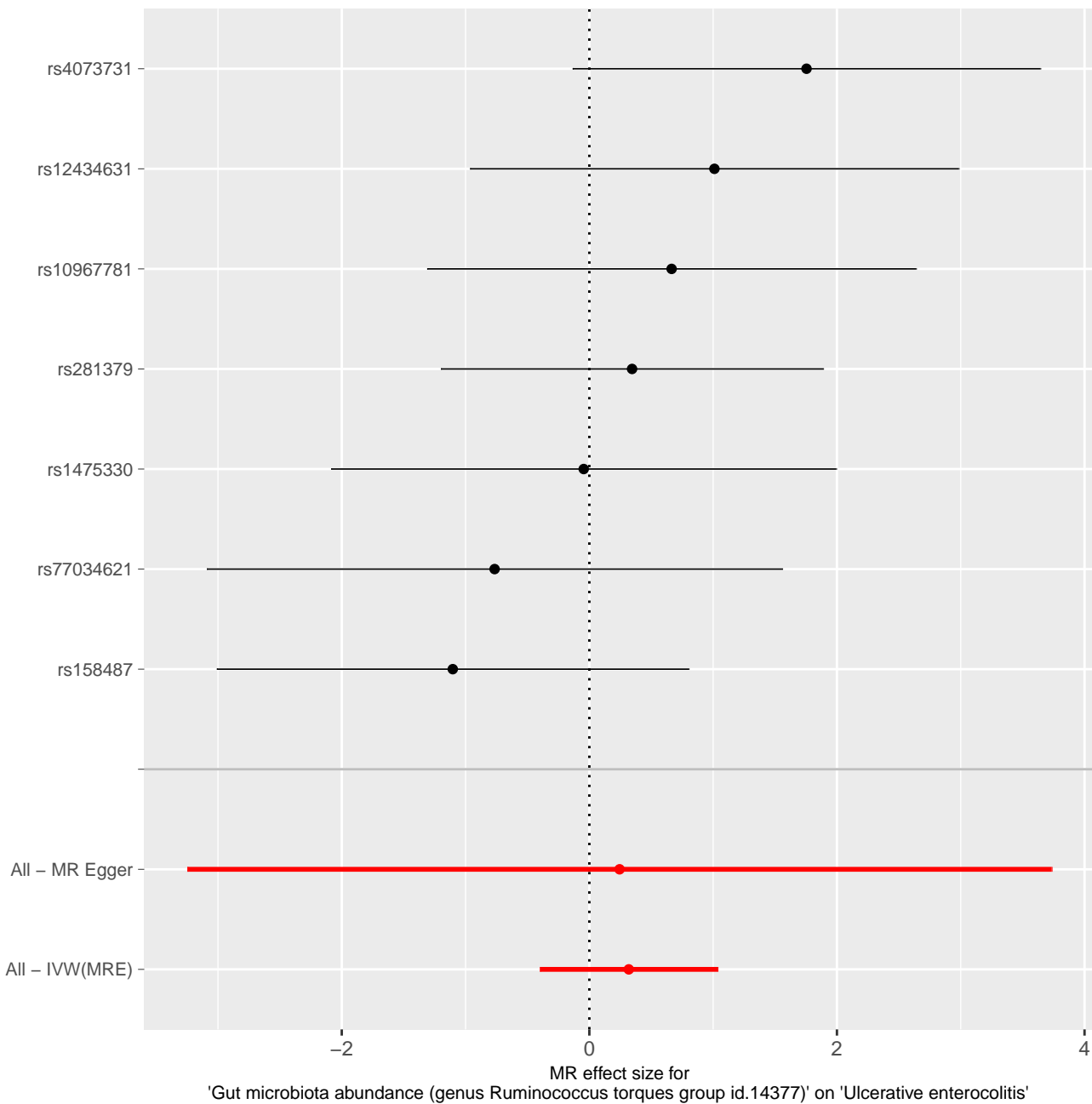


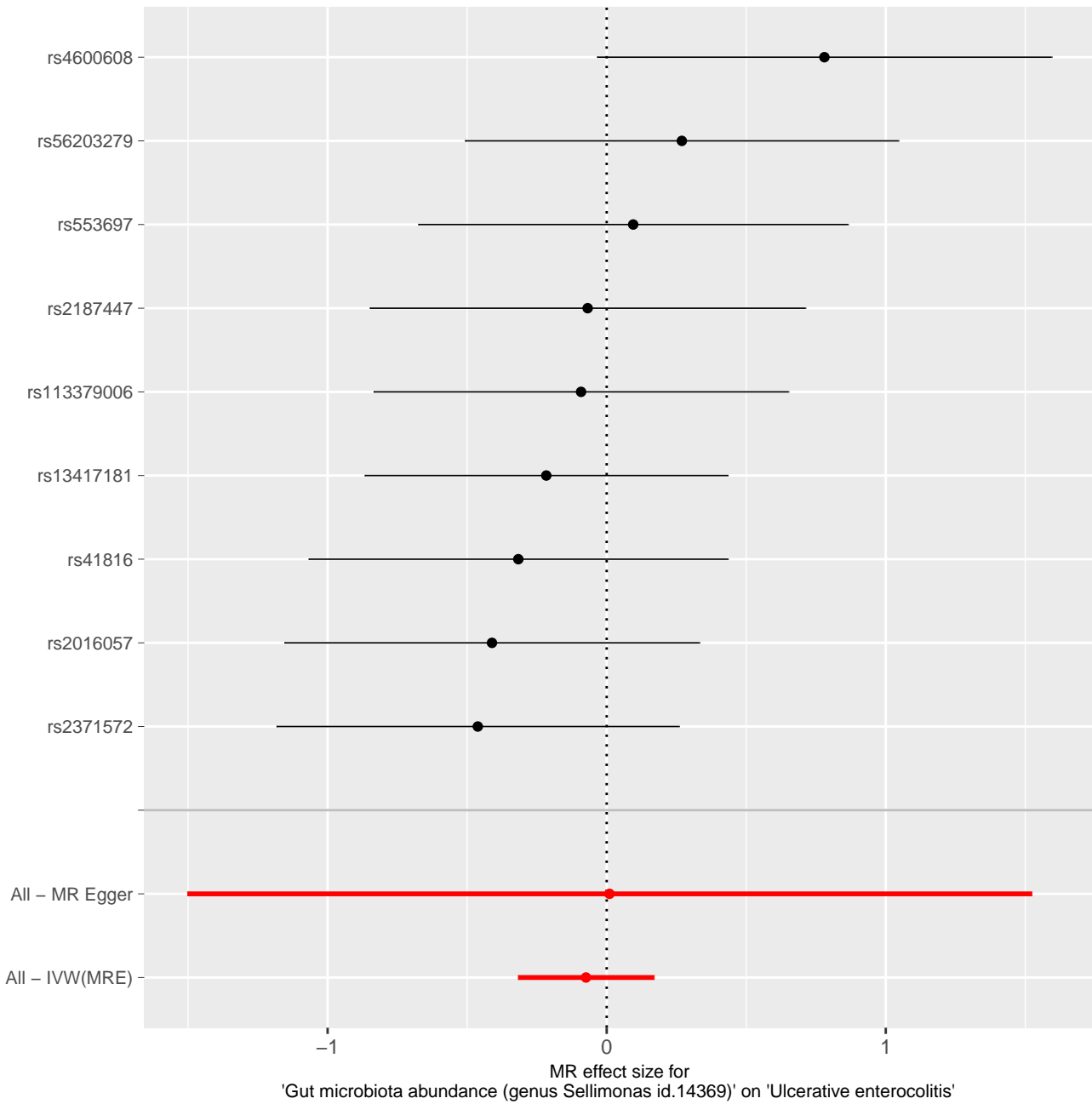


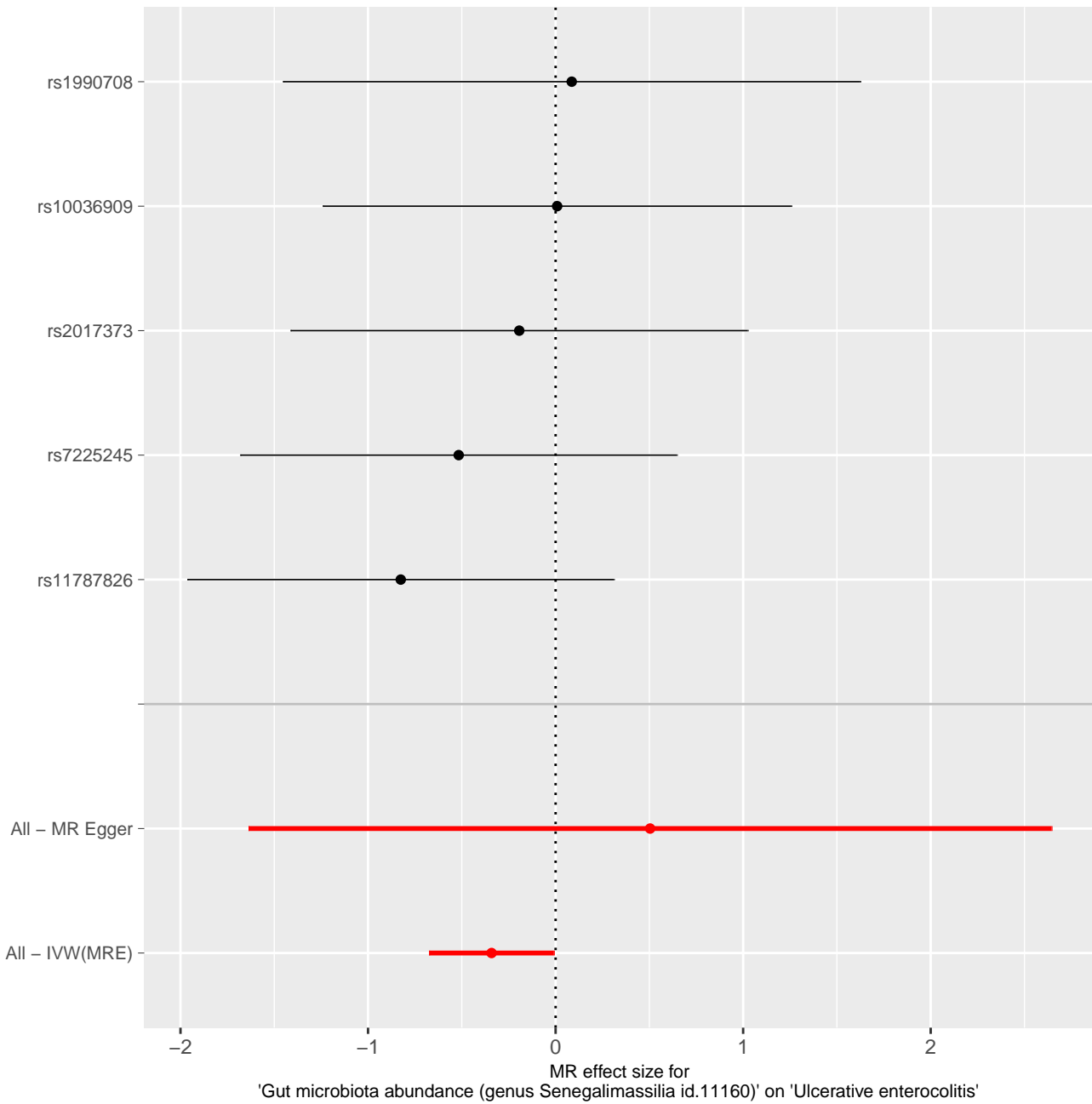


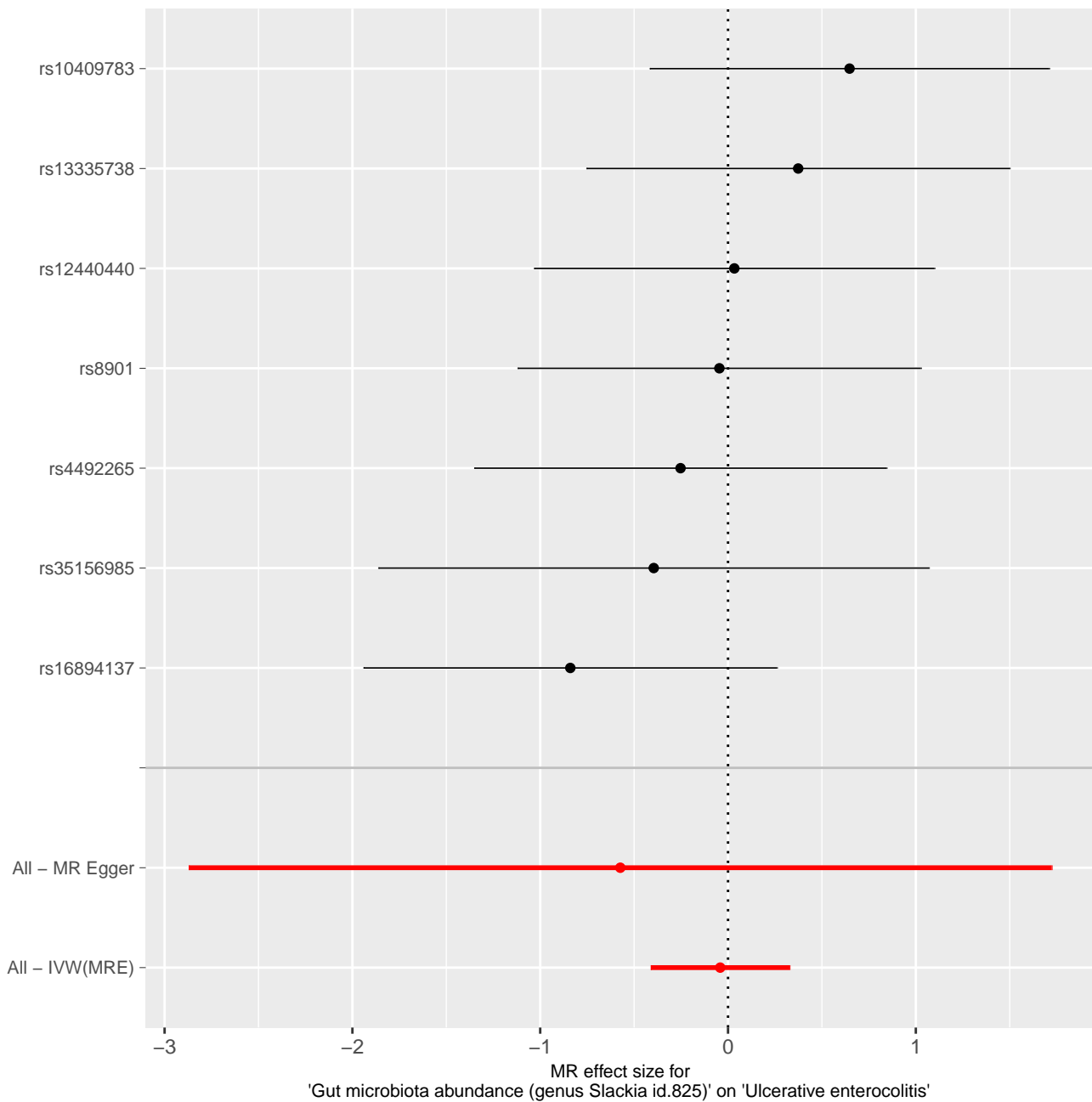


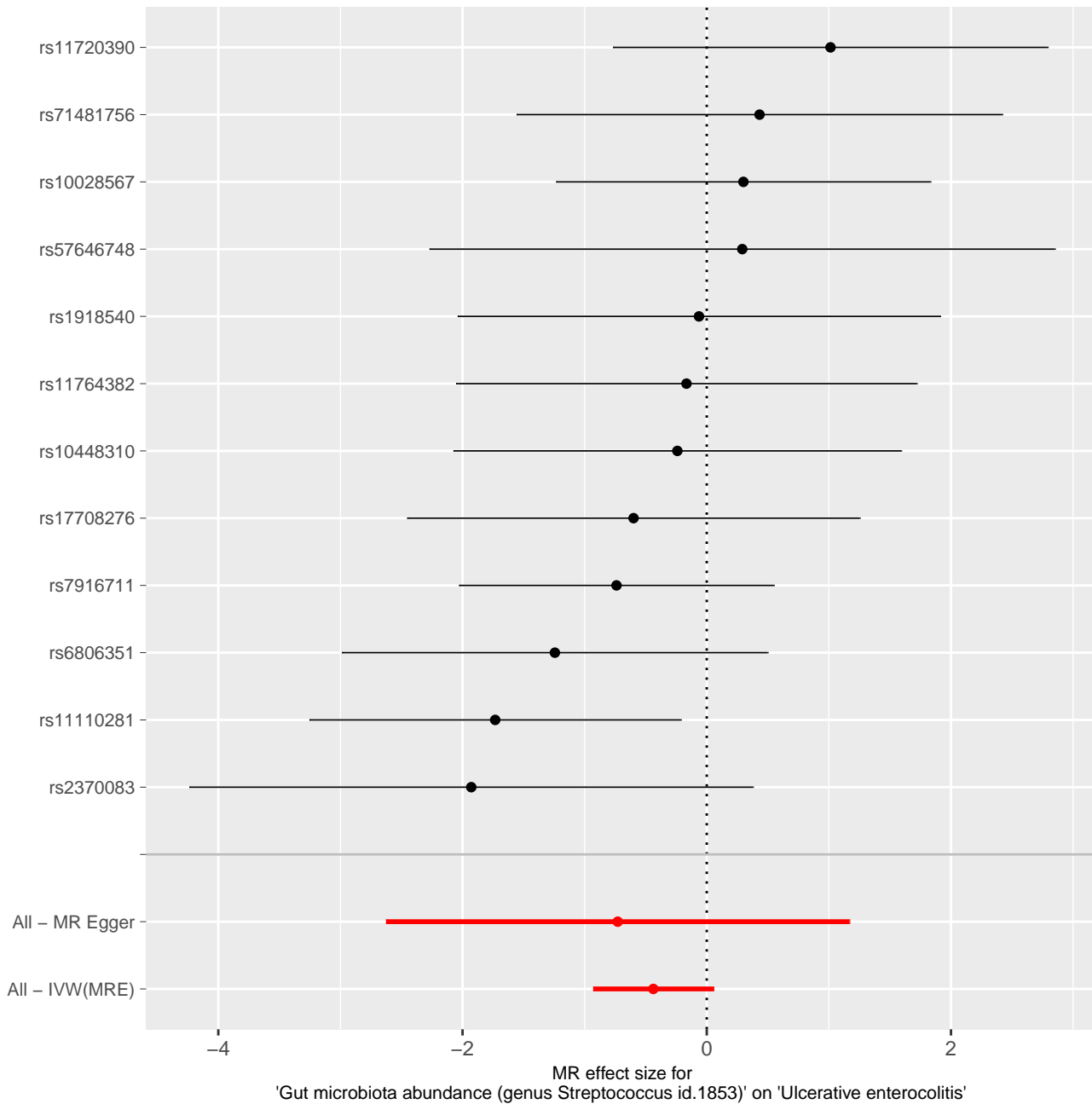


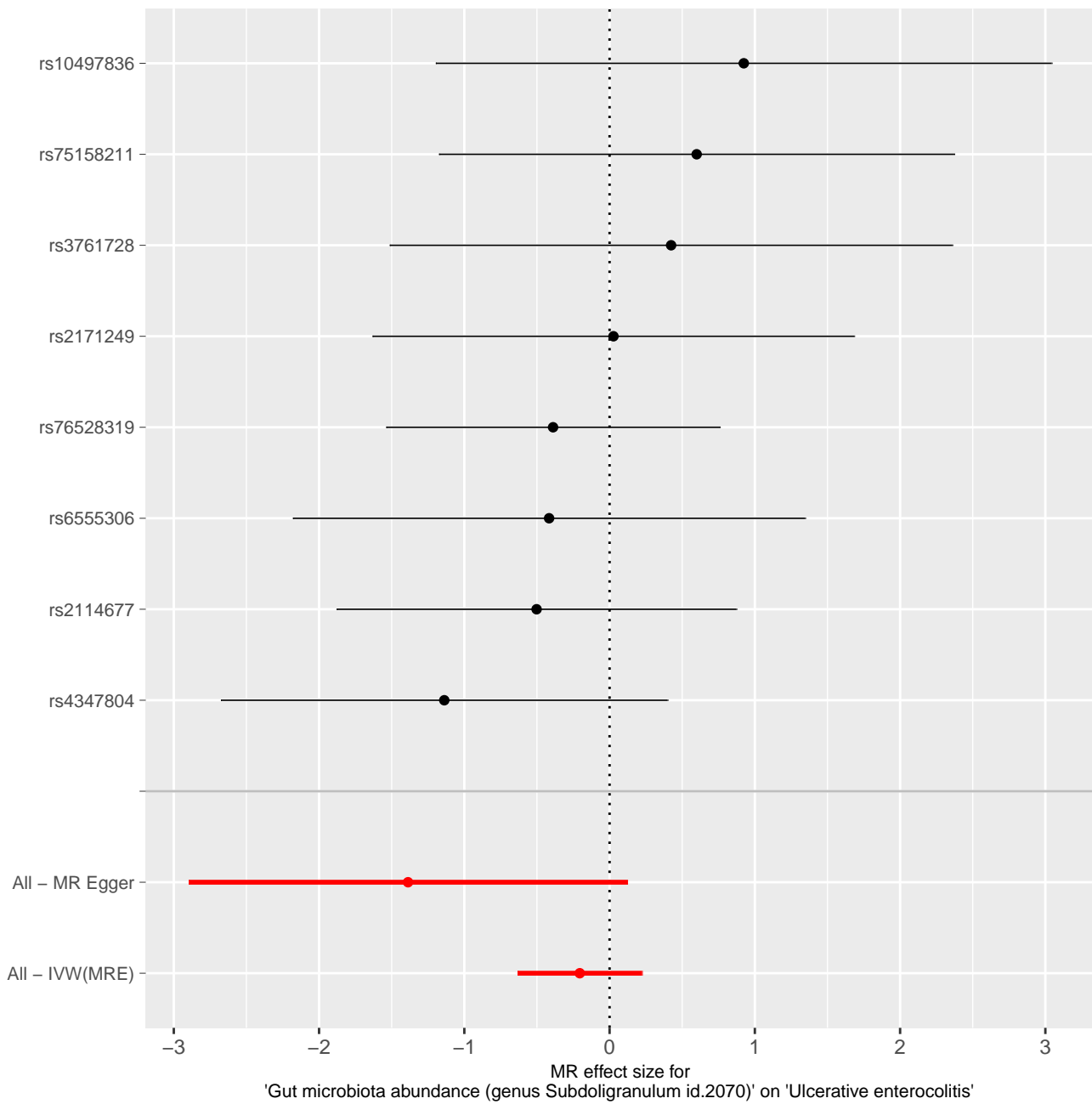


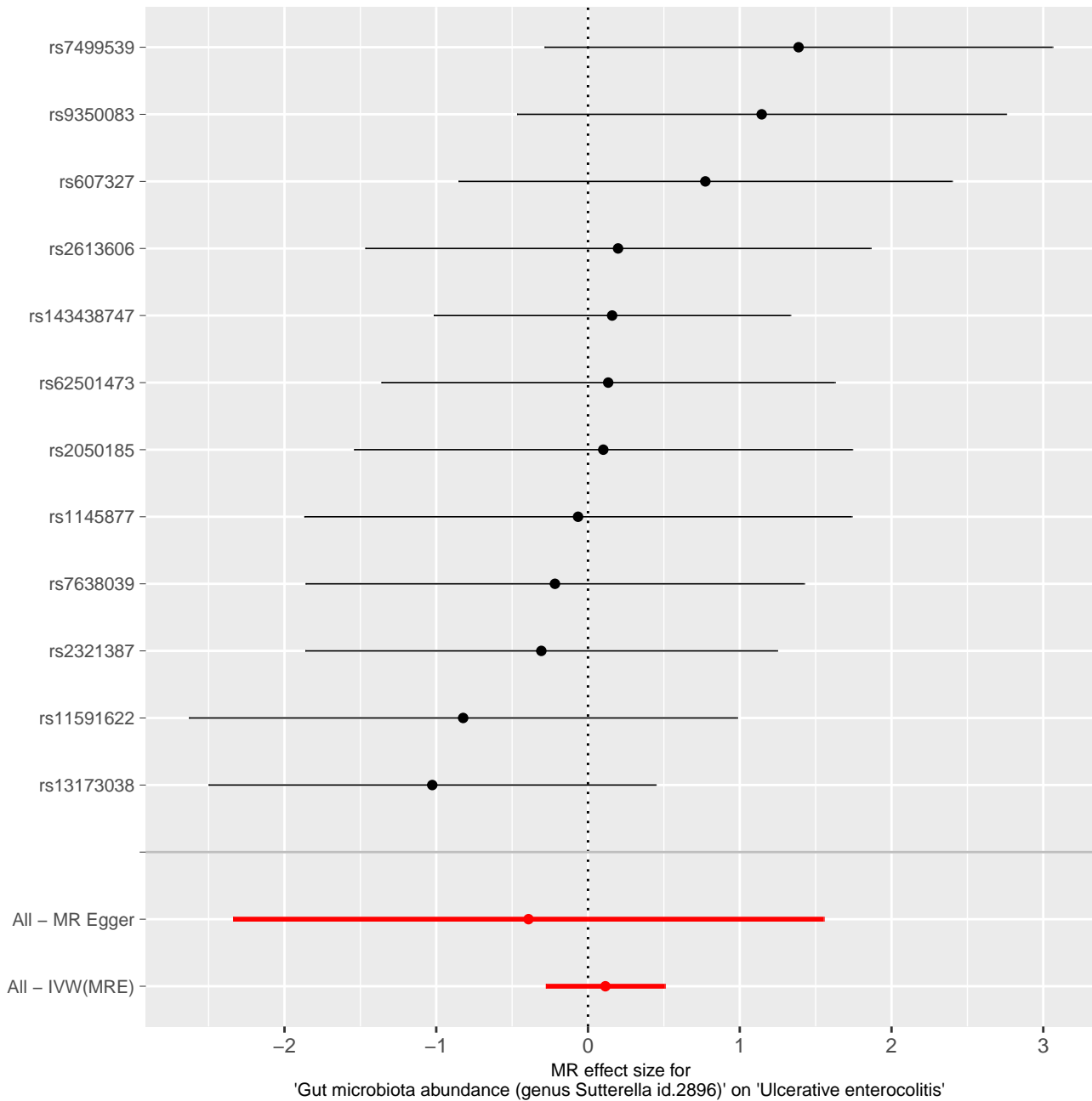


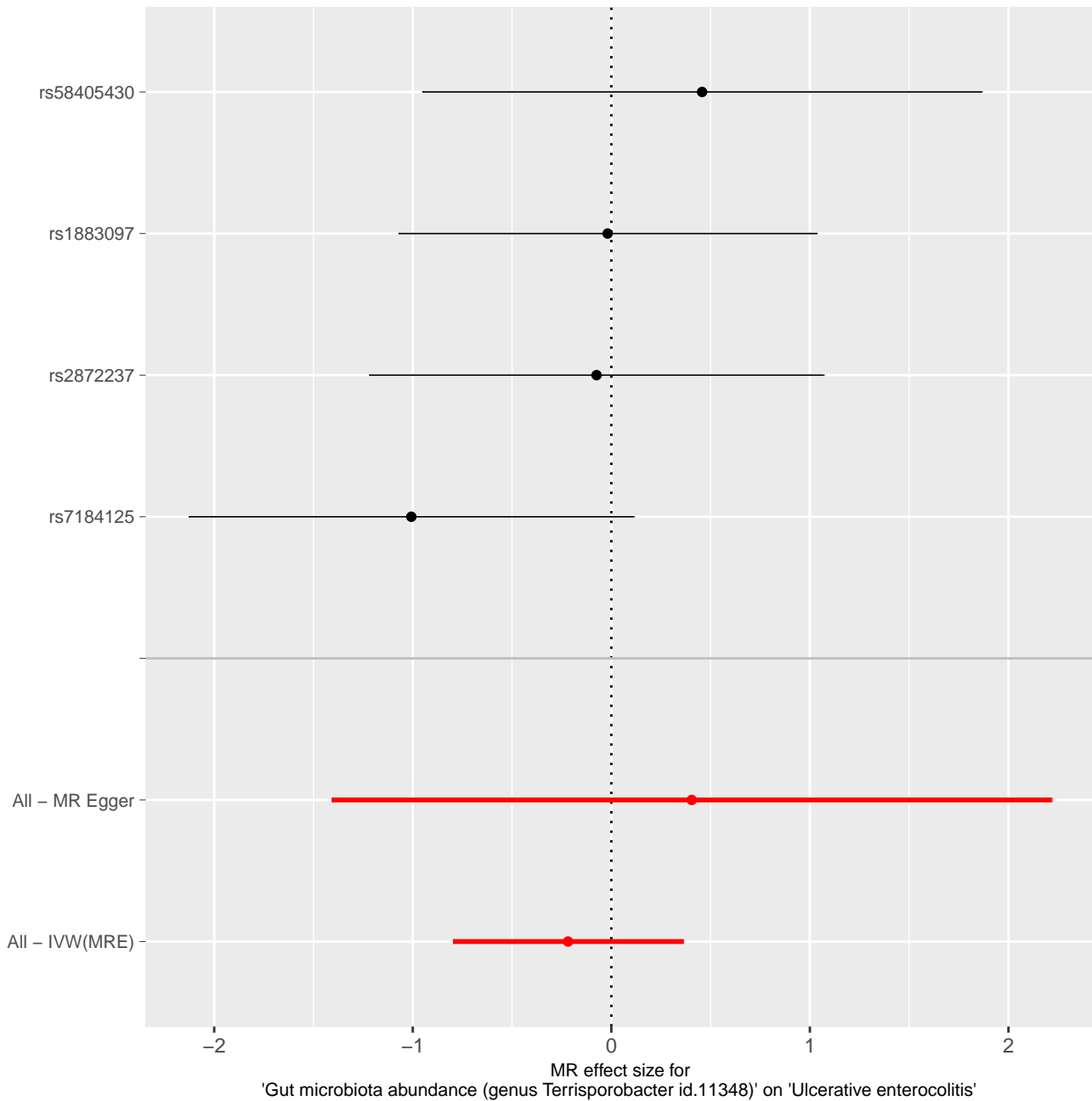


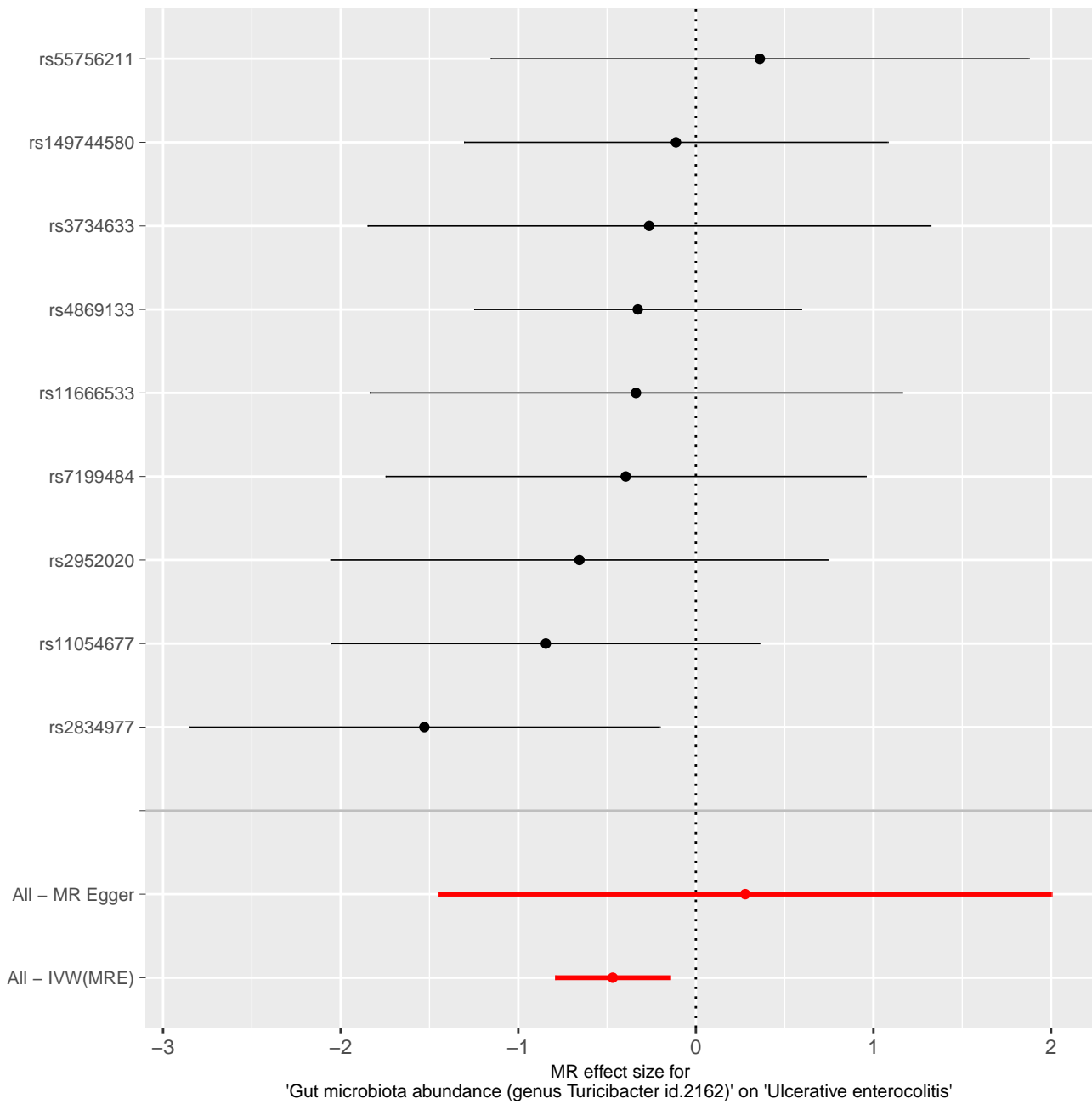


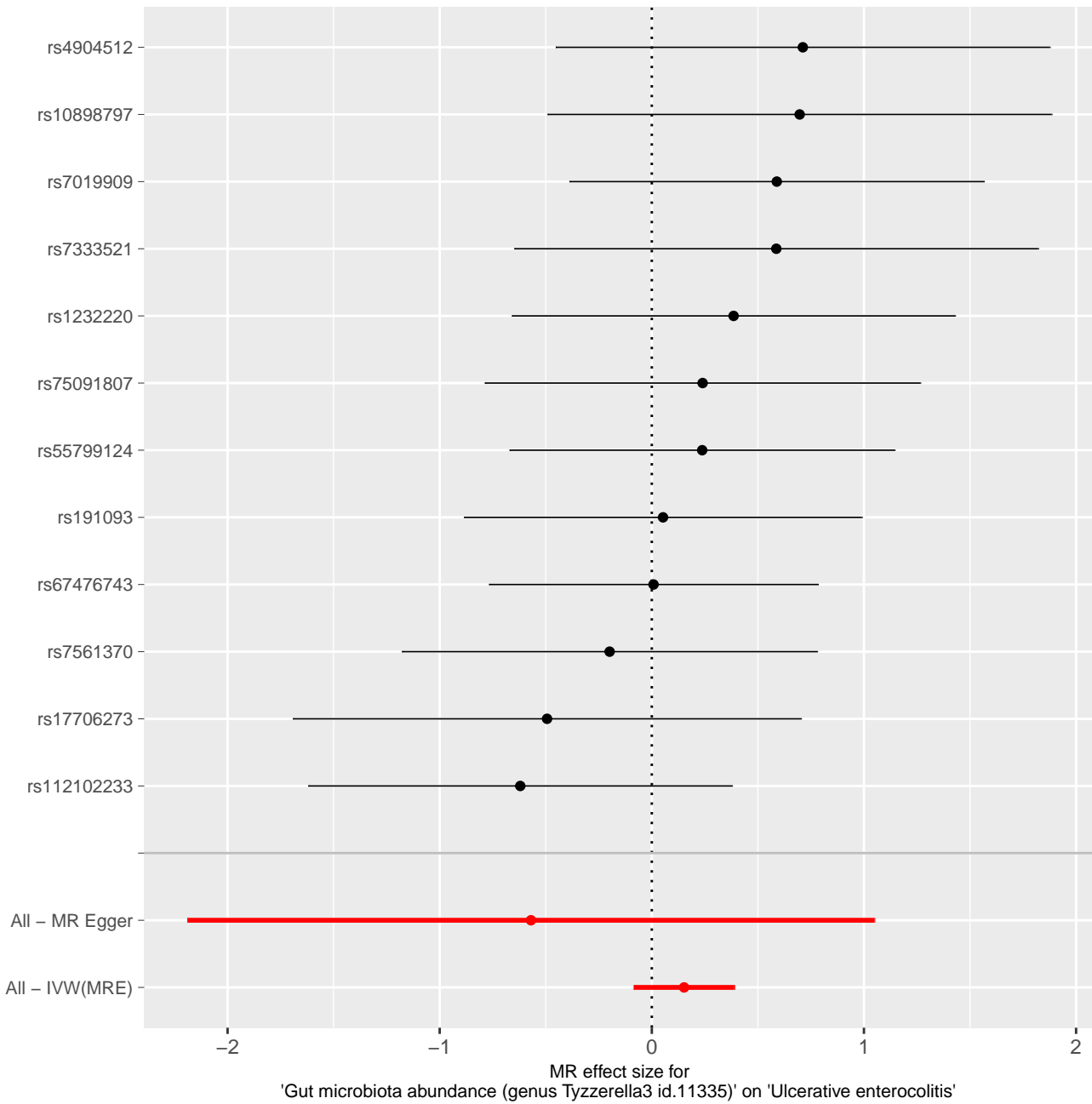


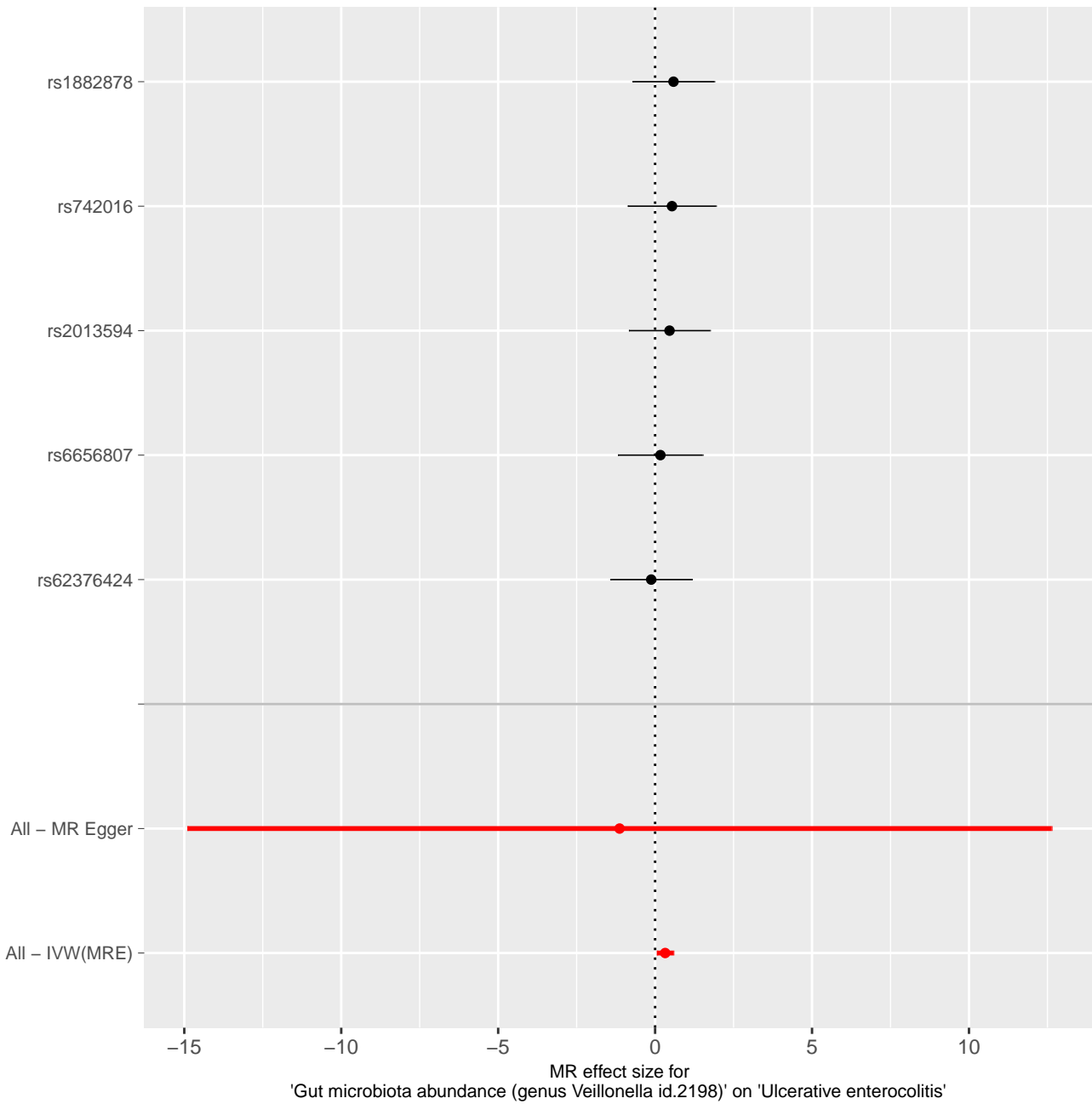


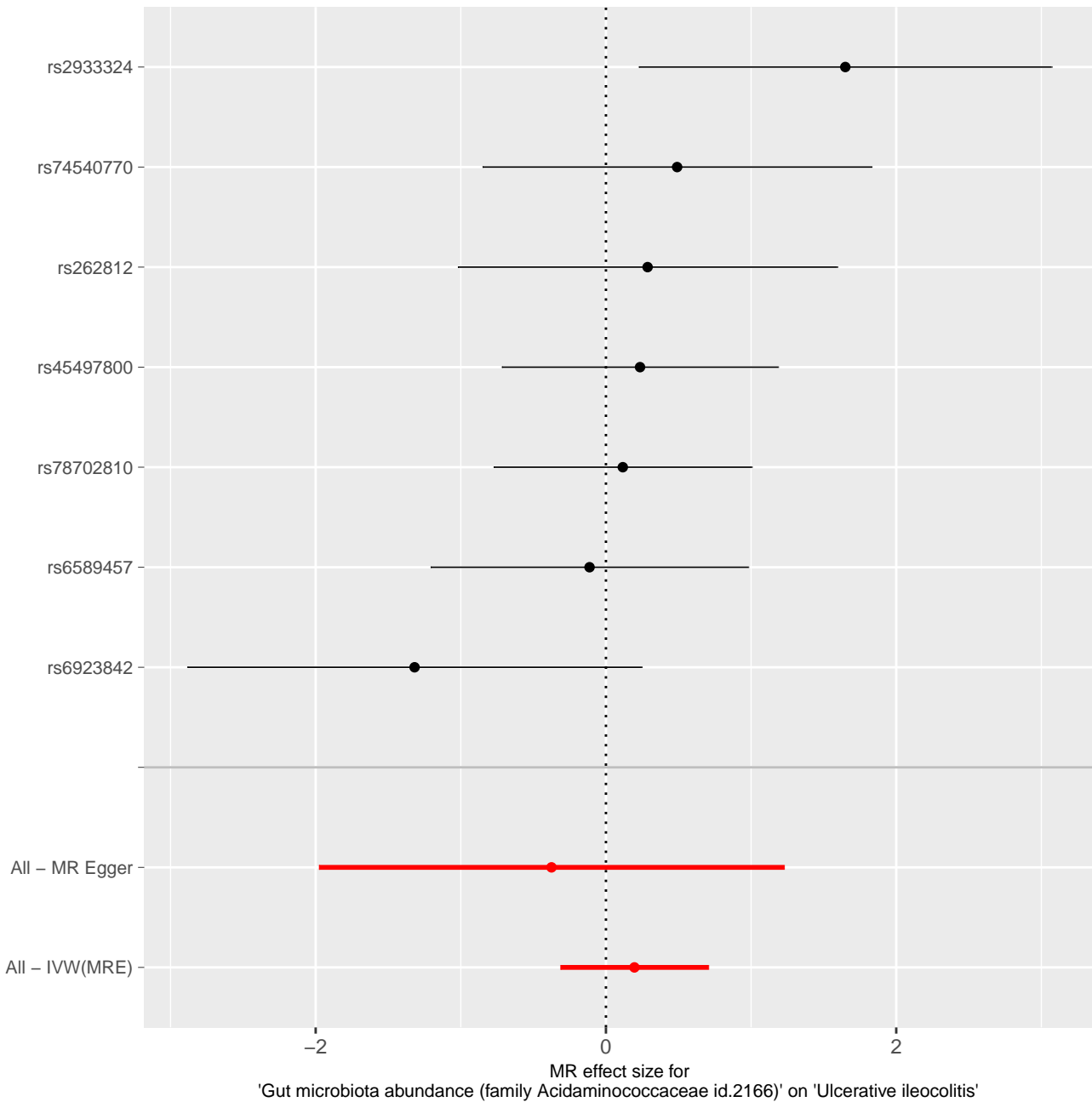


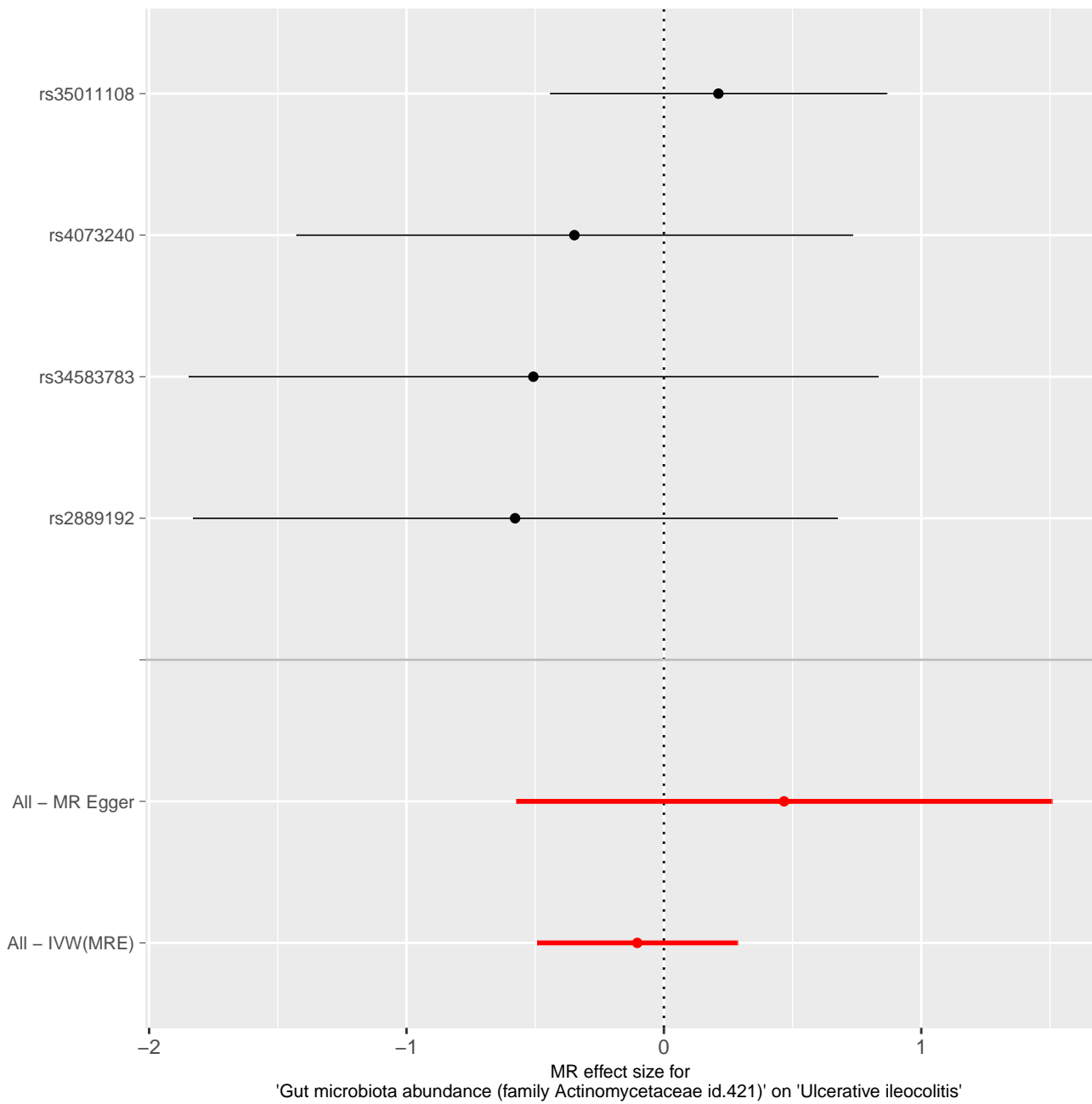


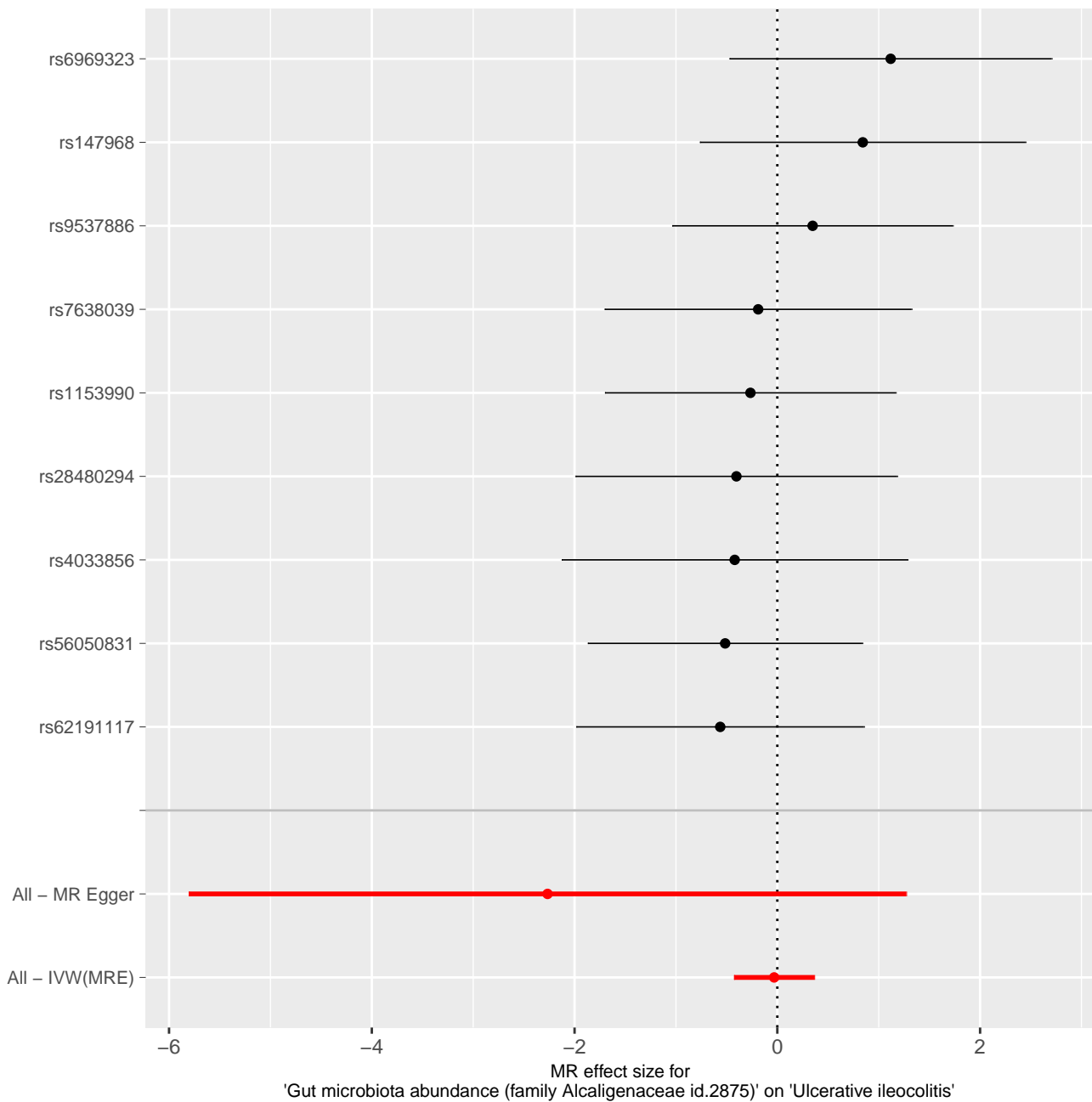


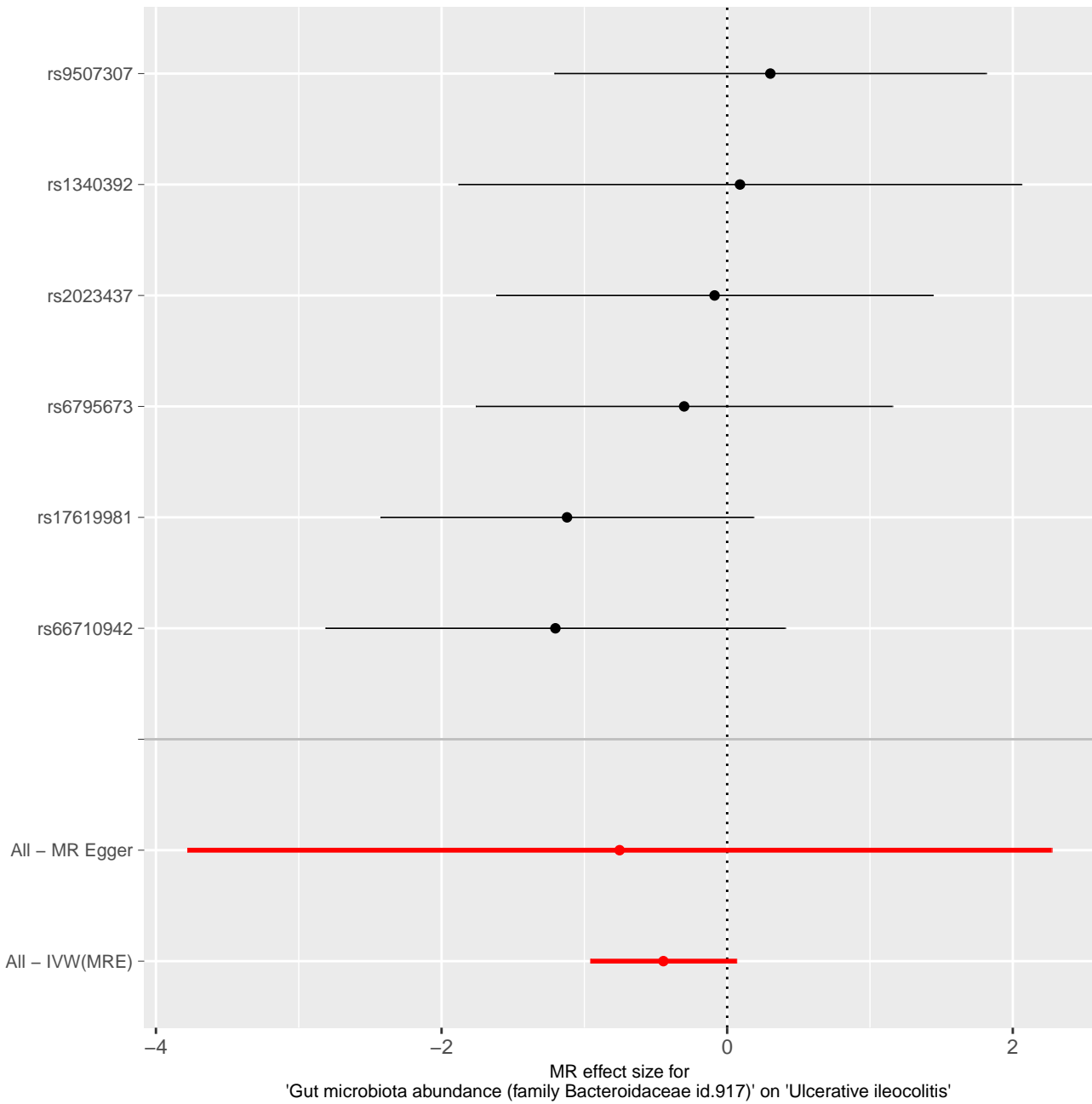


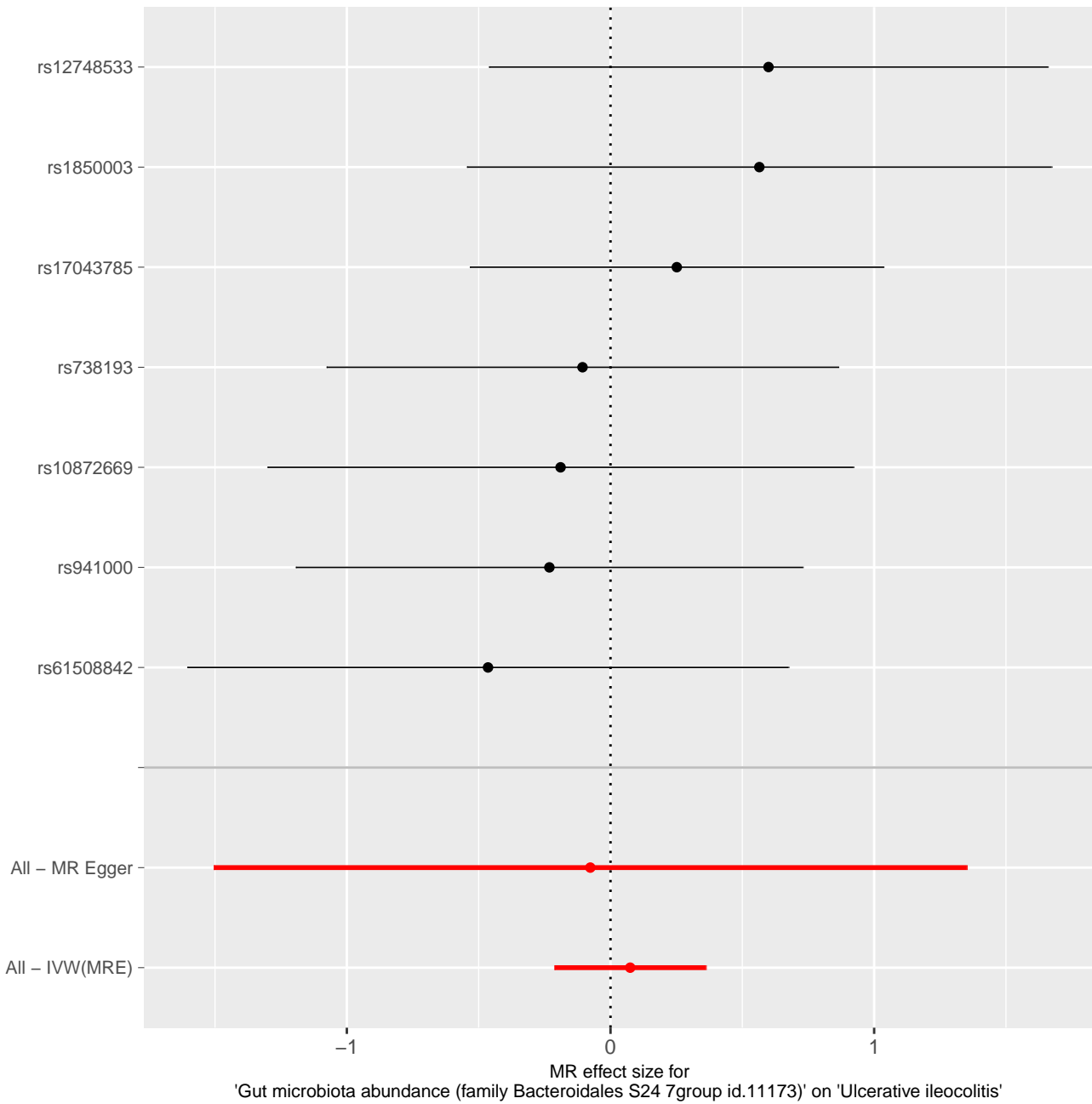




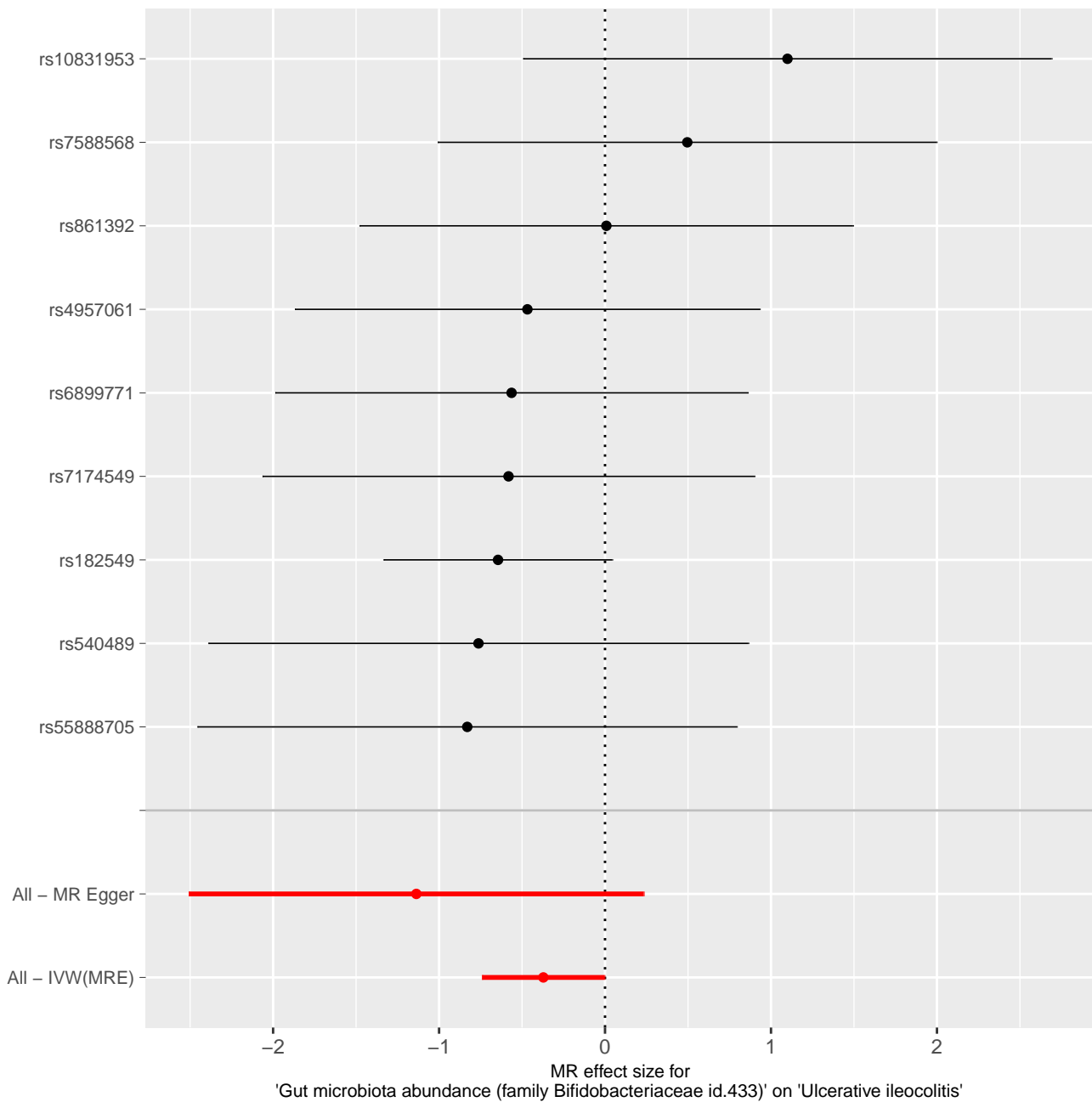


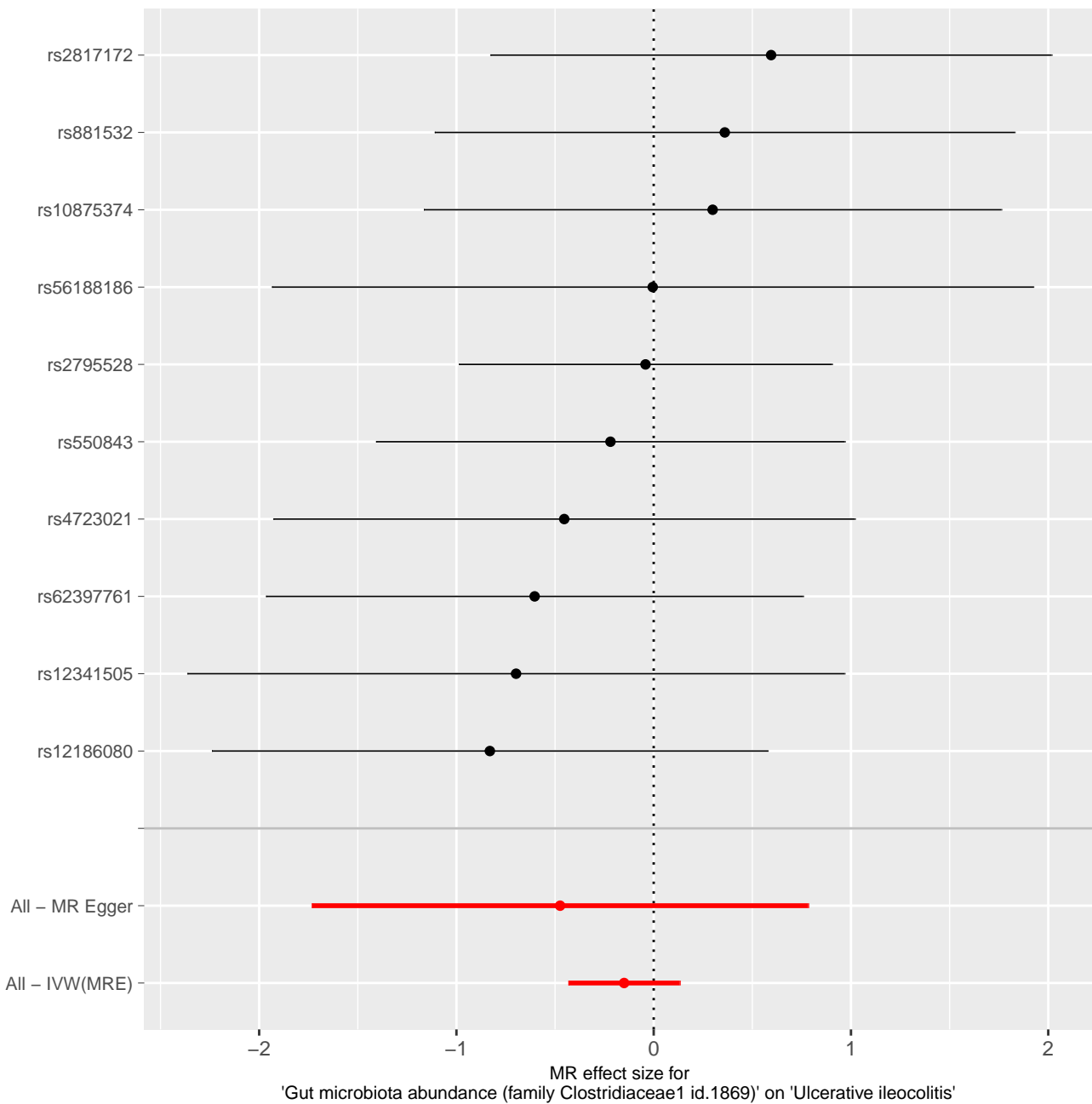


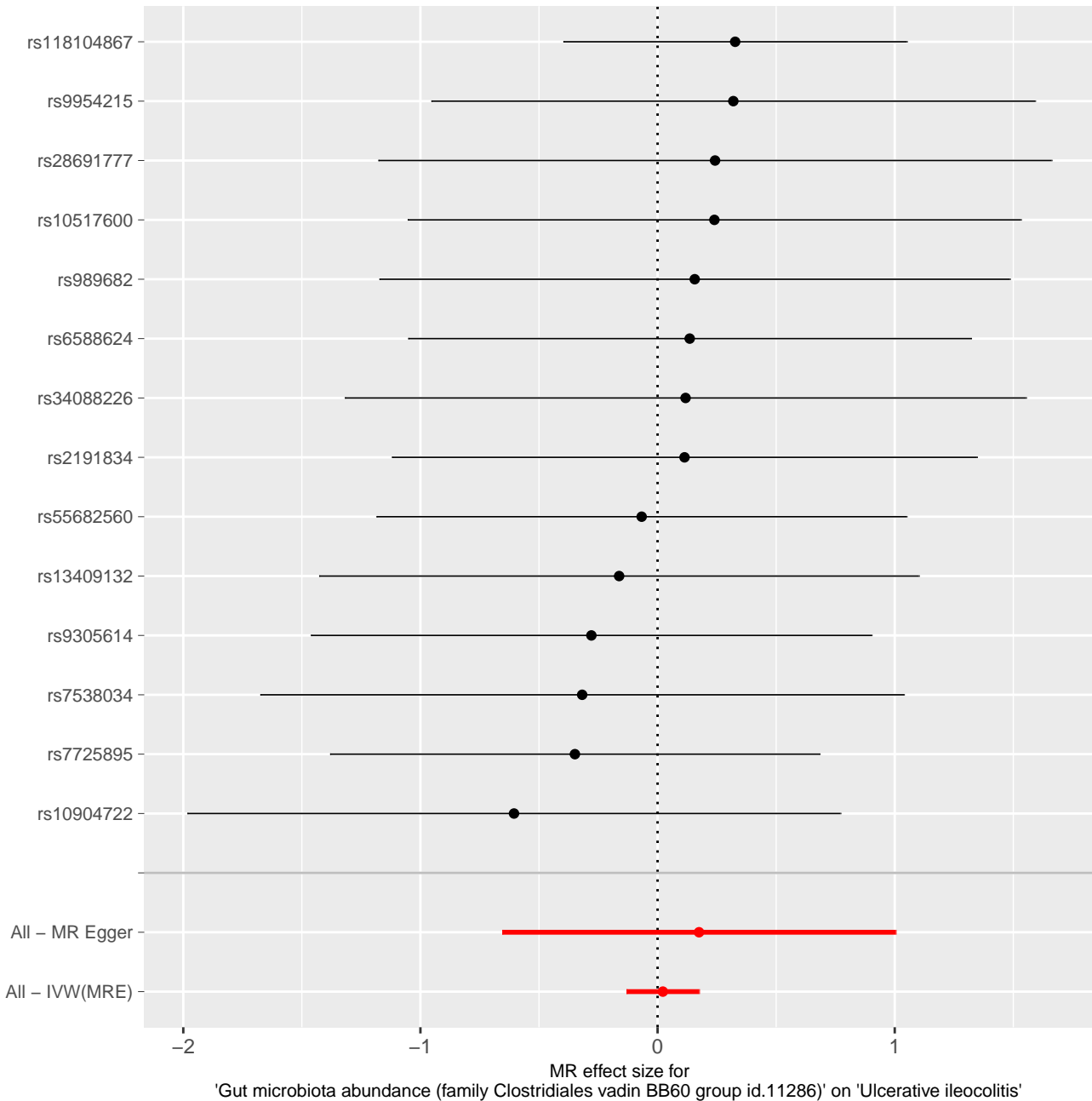


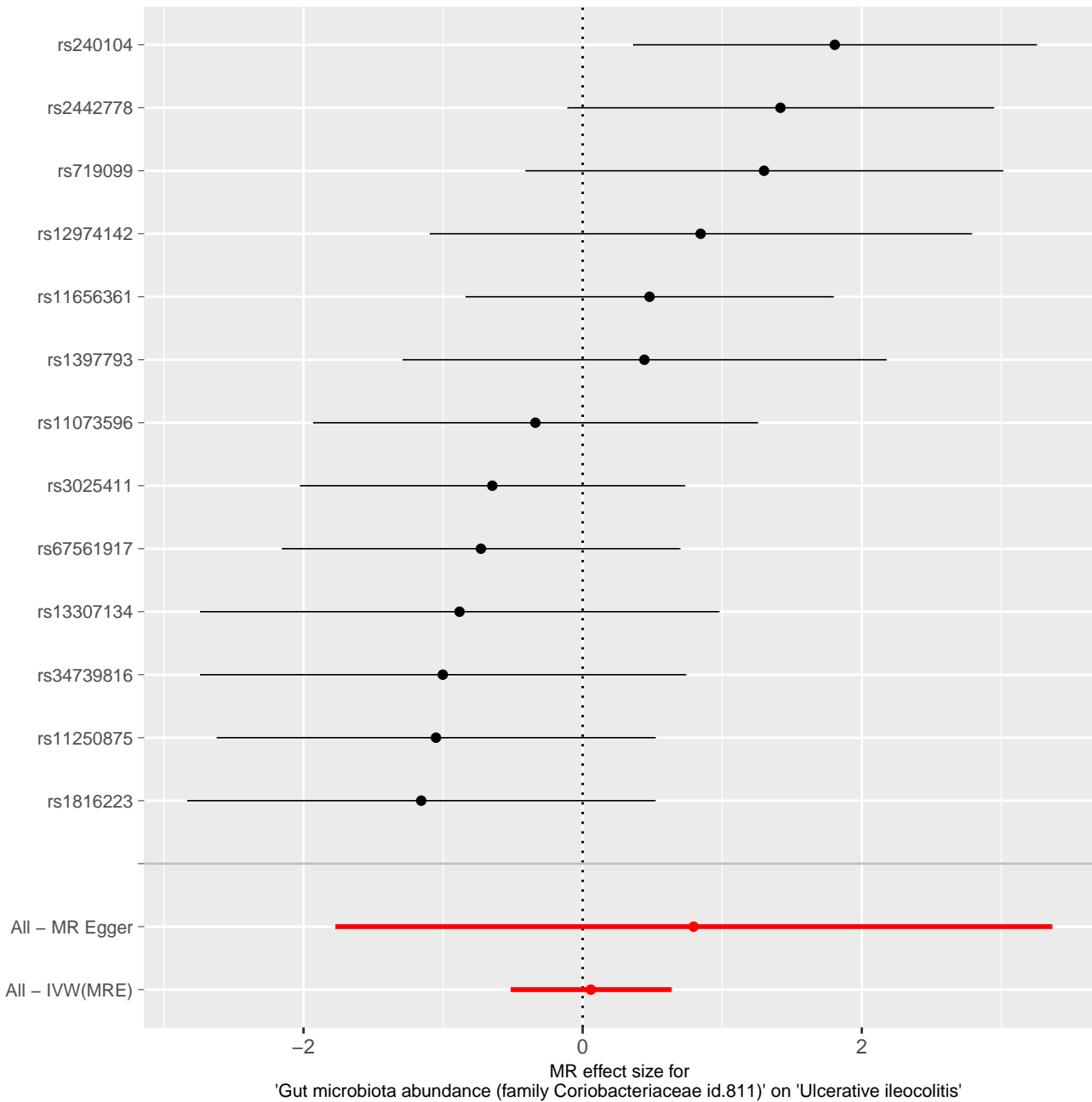


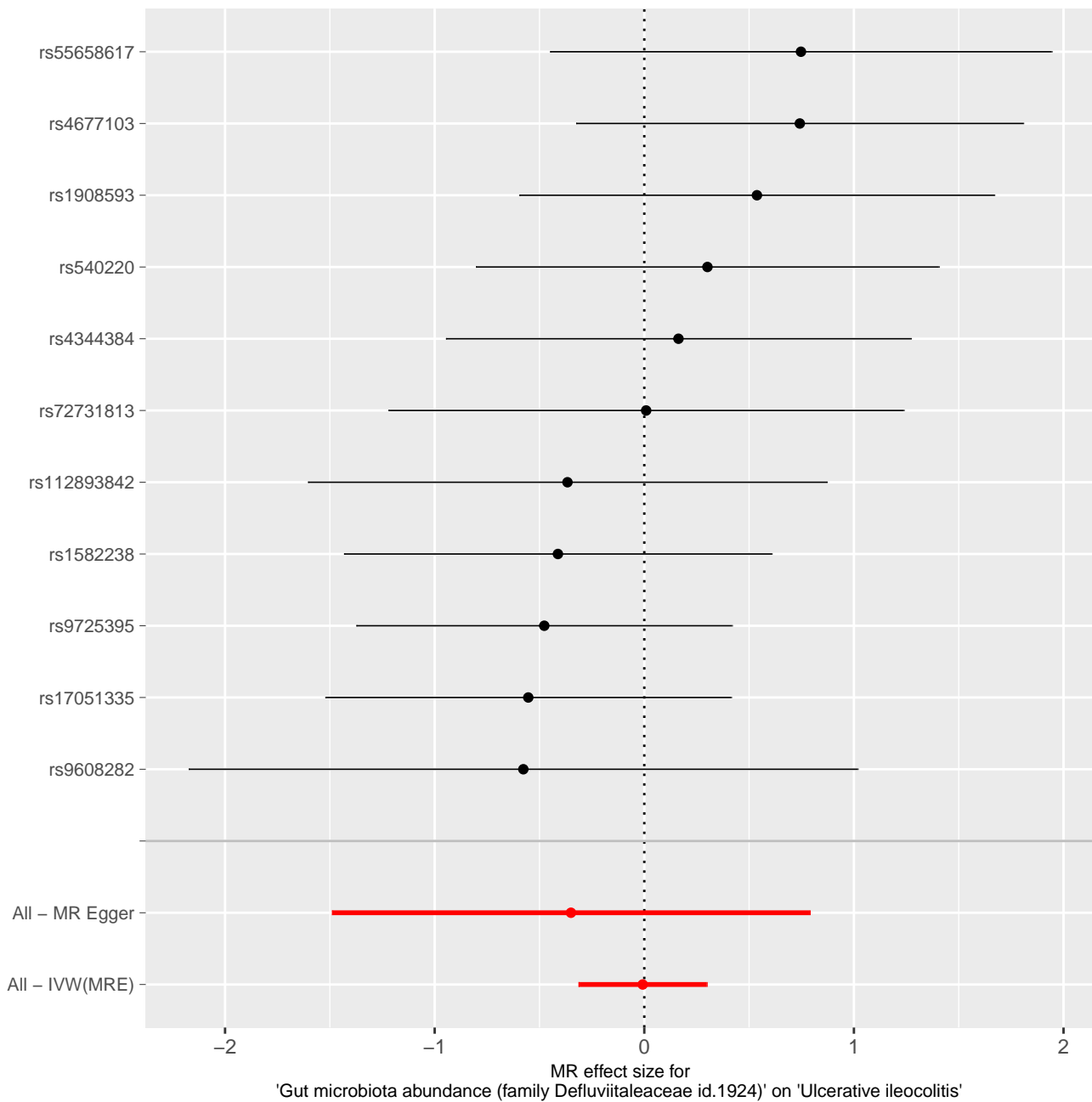
MR effect size for 'Gut microbiota abundance (family Bacteroidales S24 7group id.11173)' on 'Ulcerative ileocolitis'

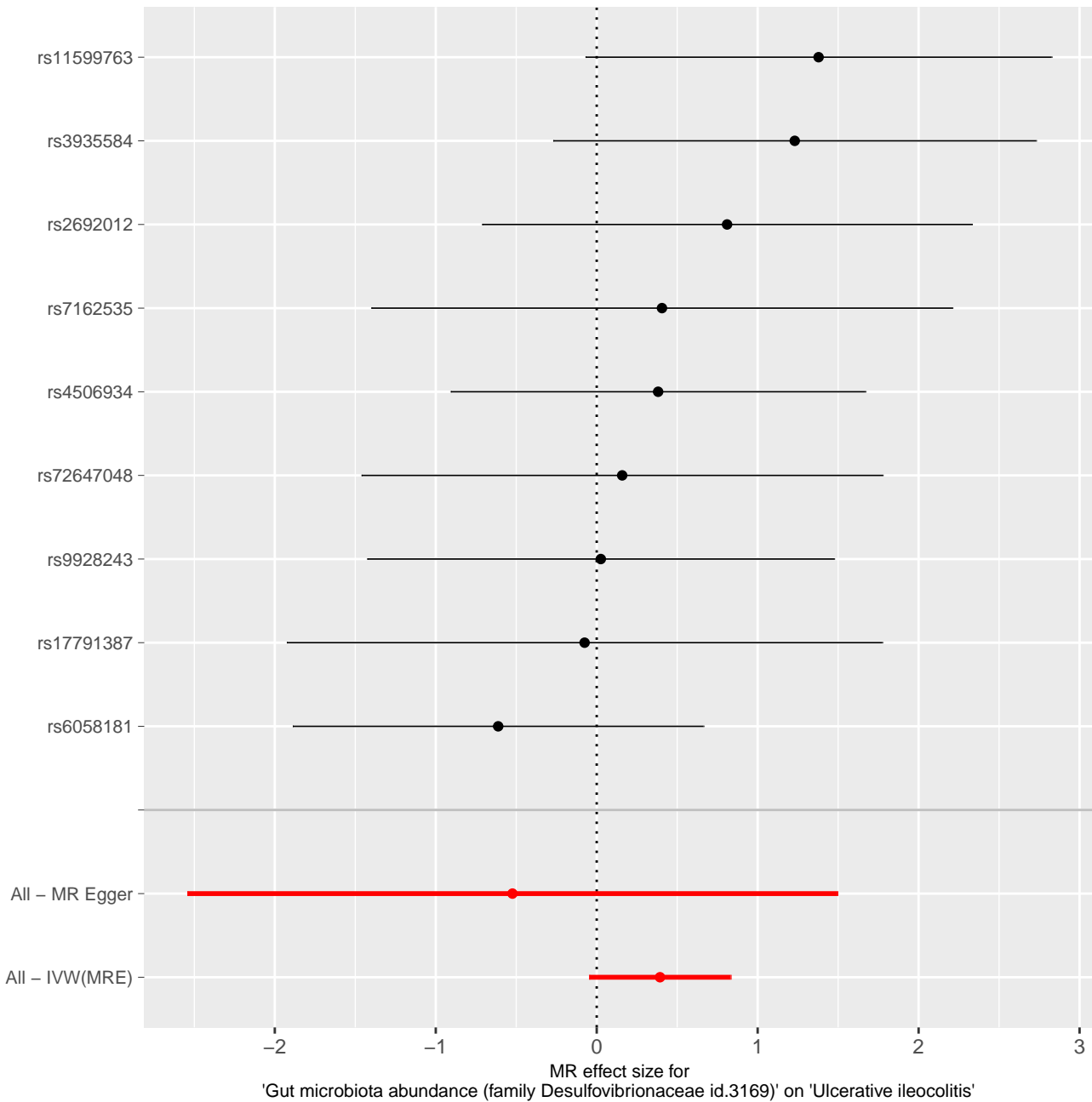


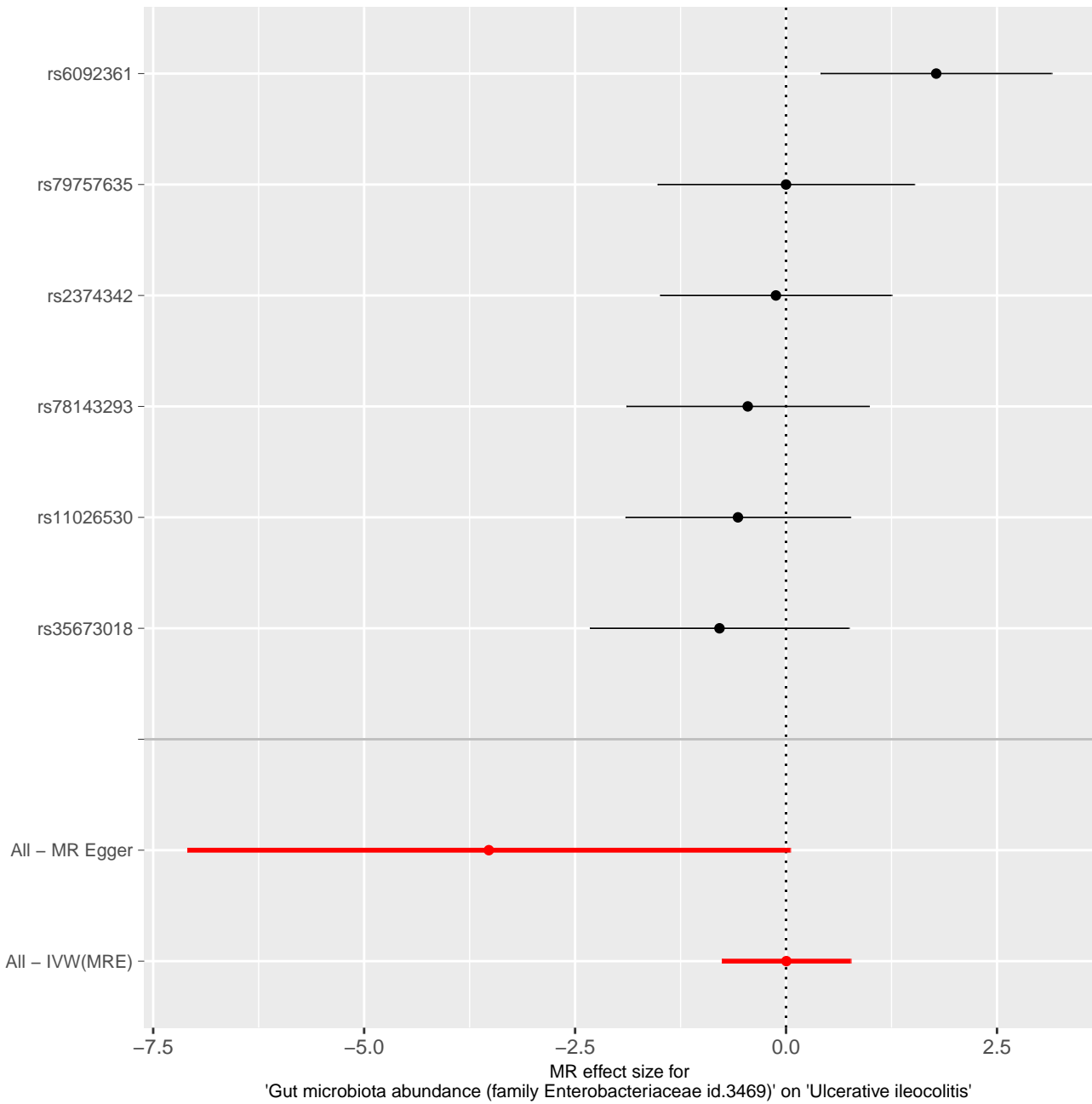


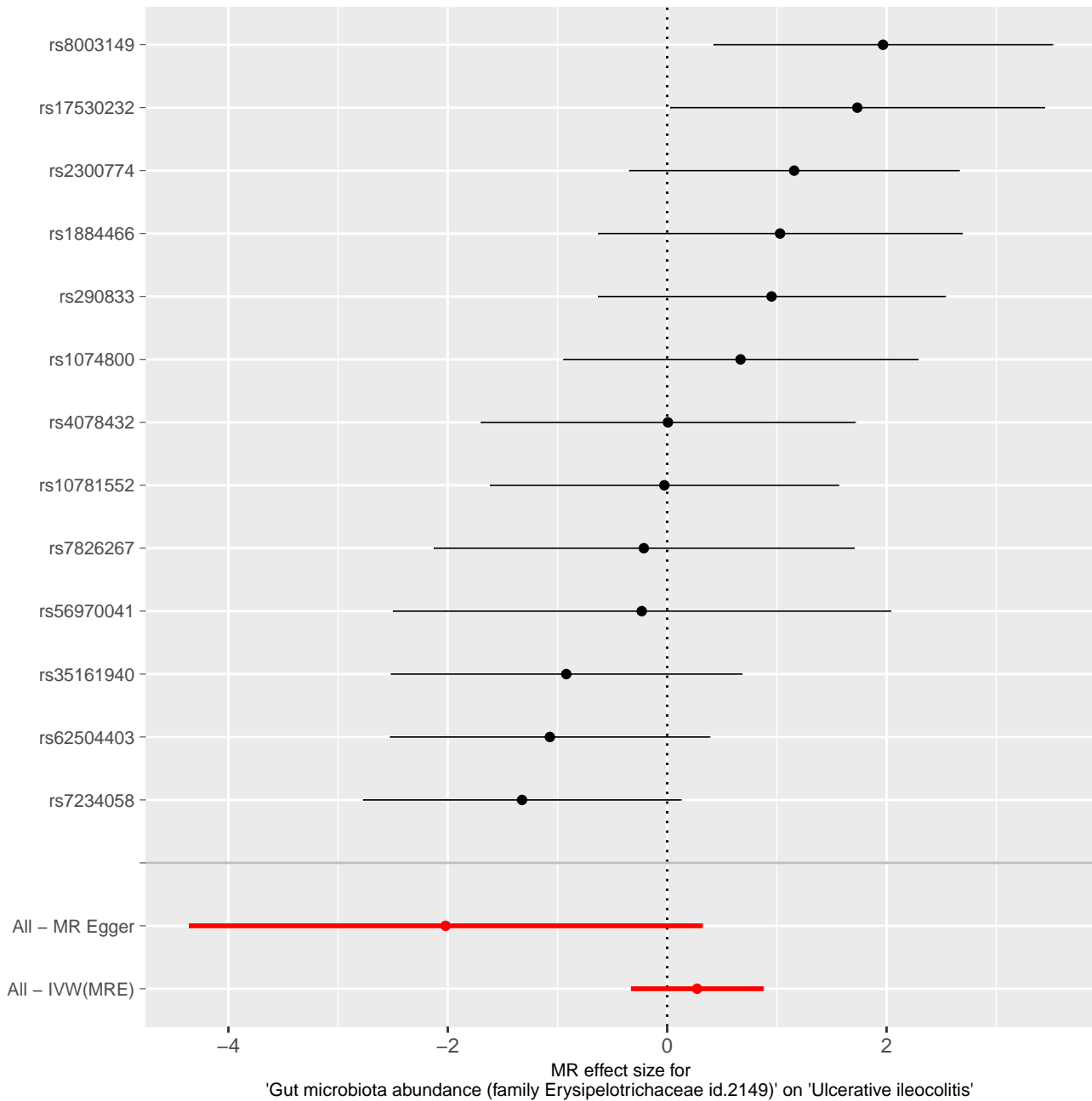


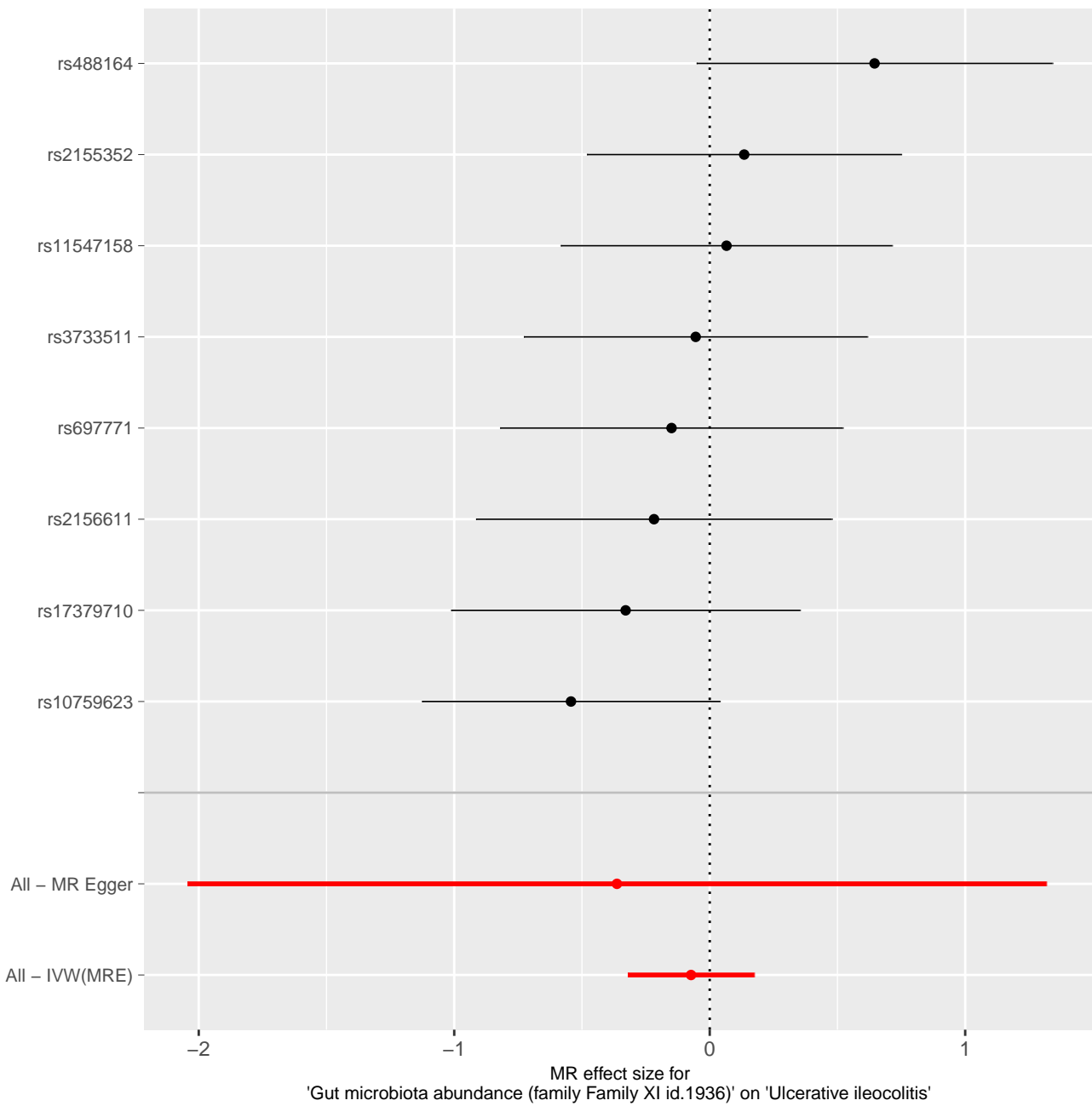


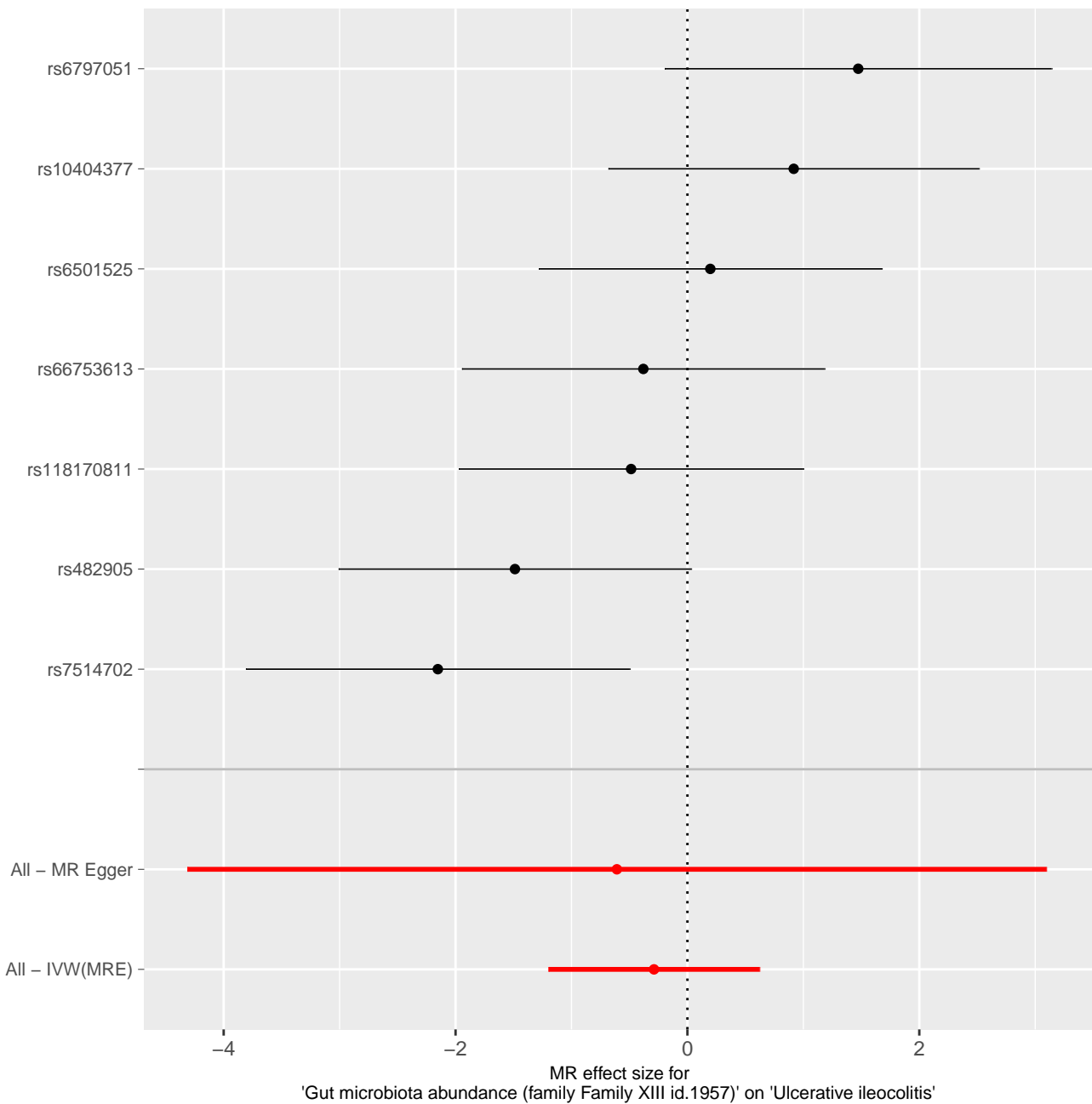


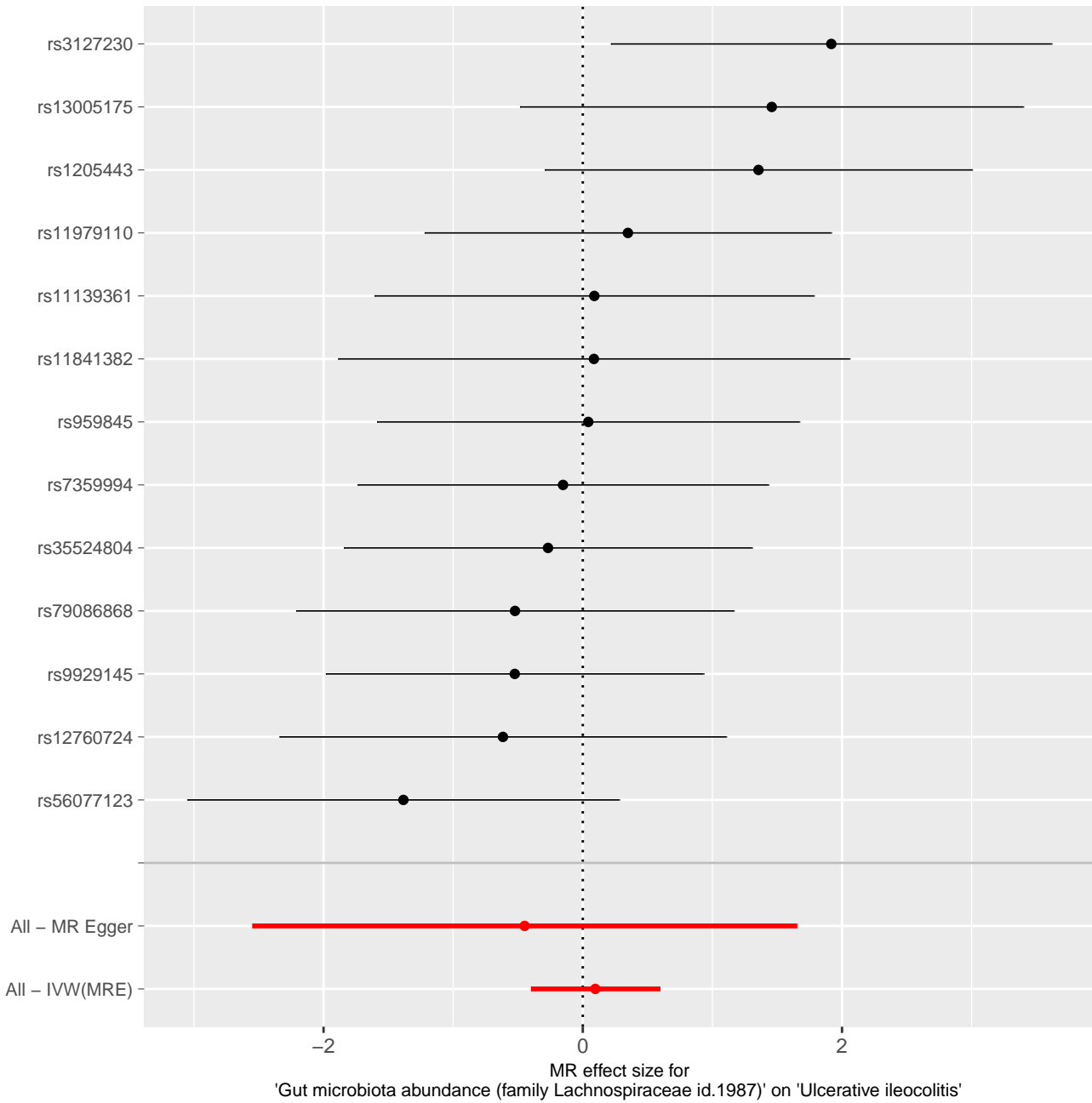


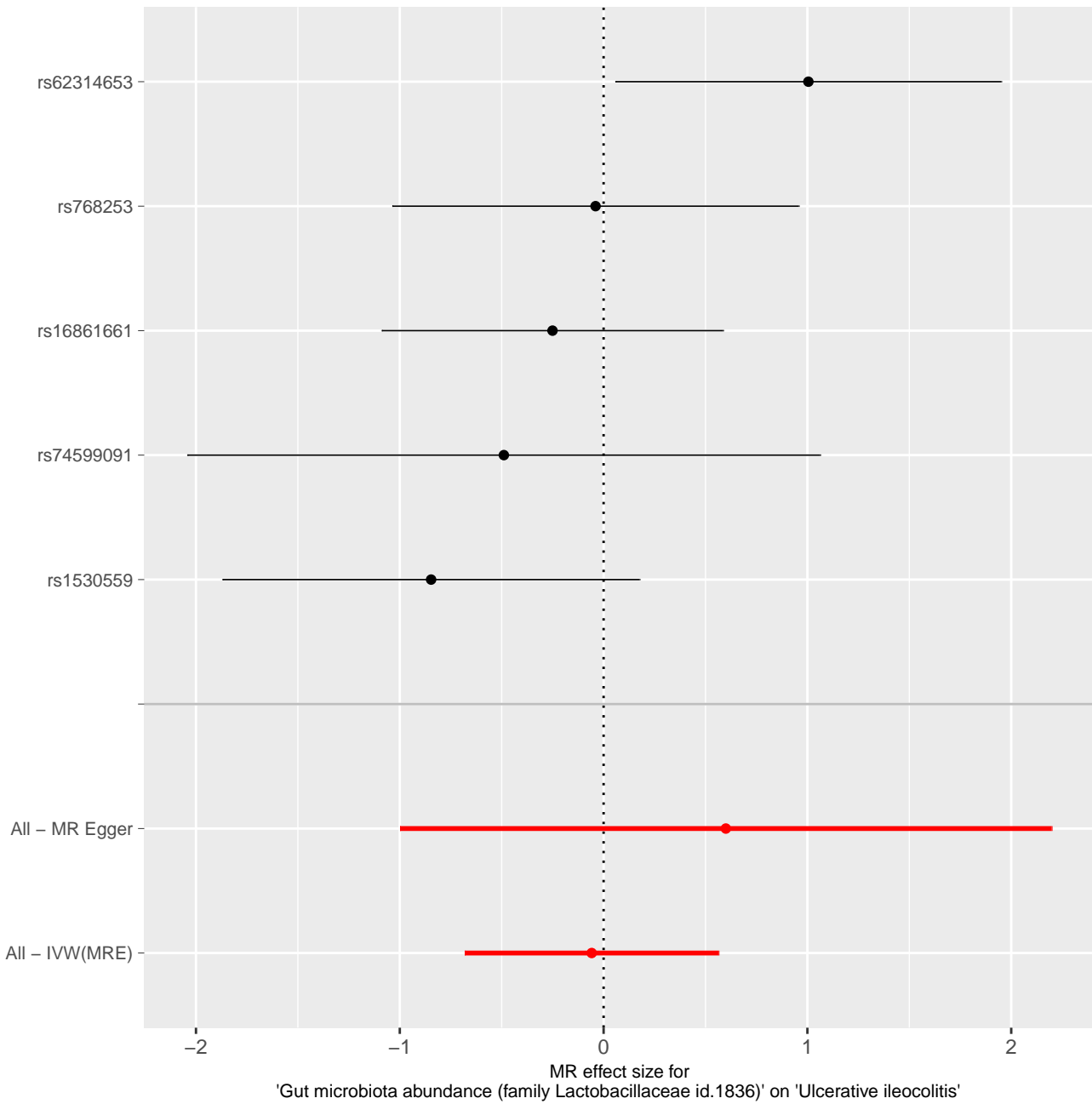


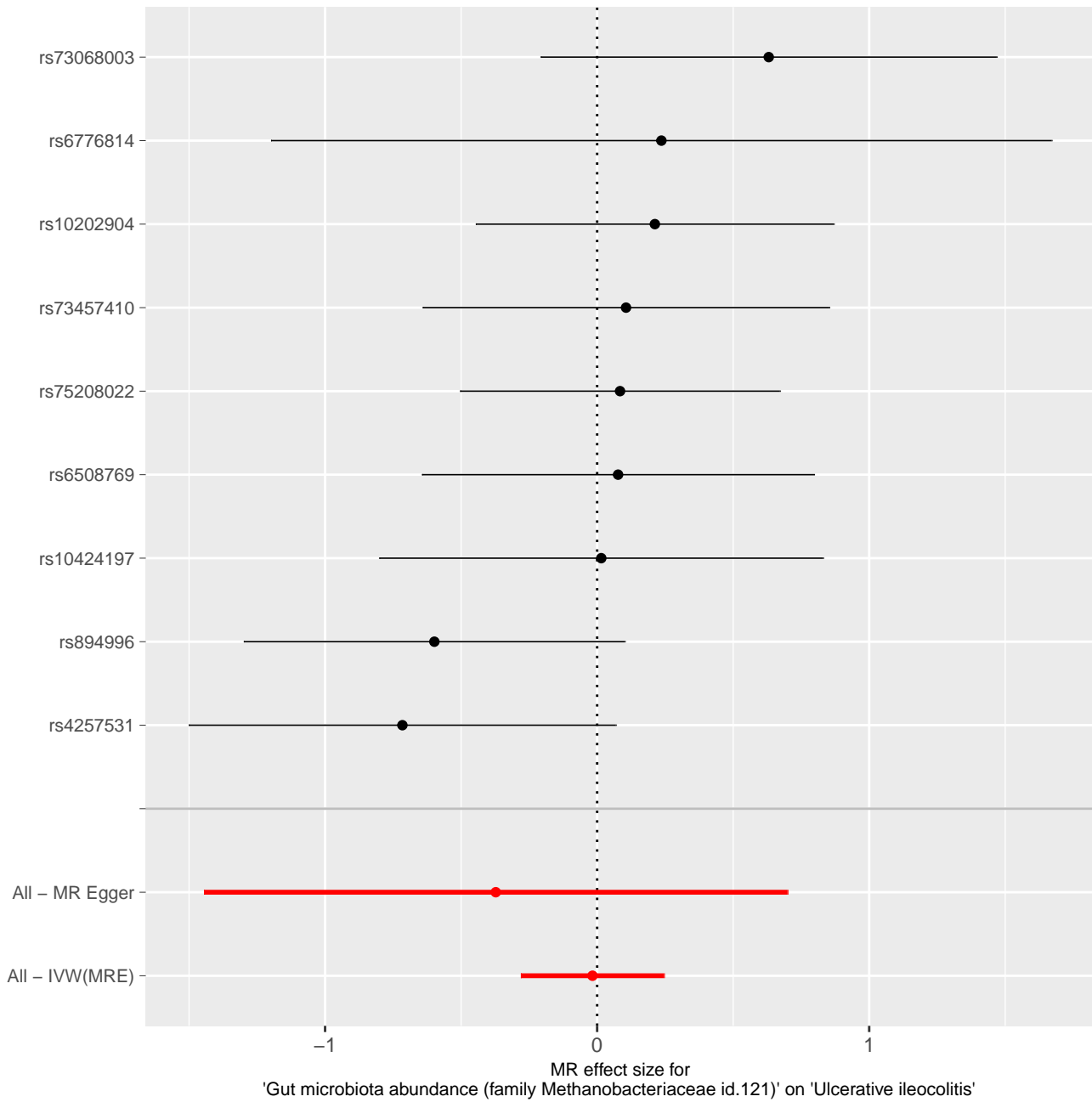


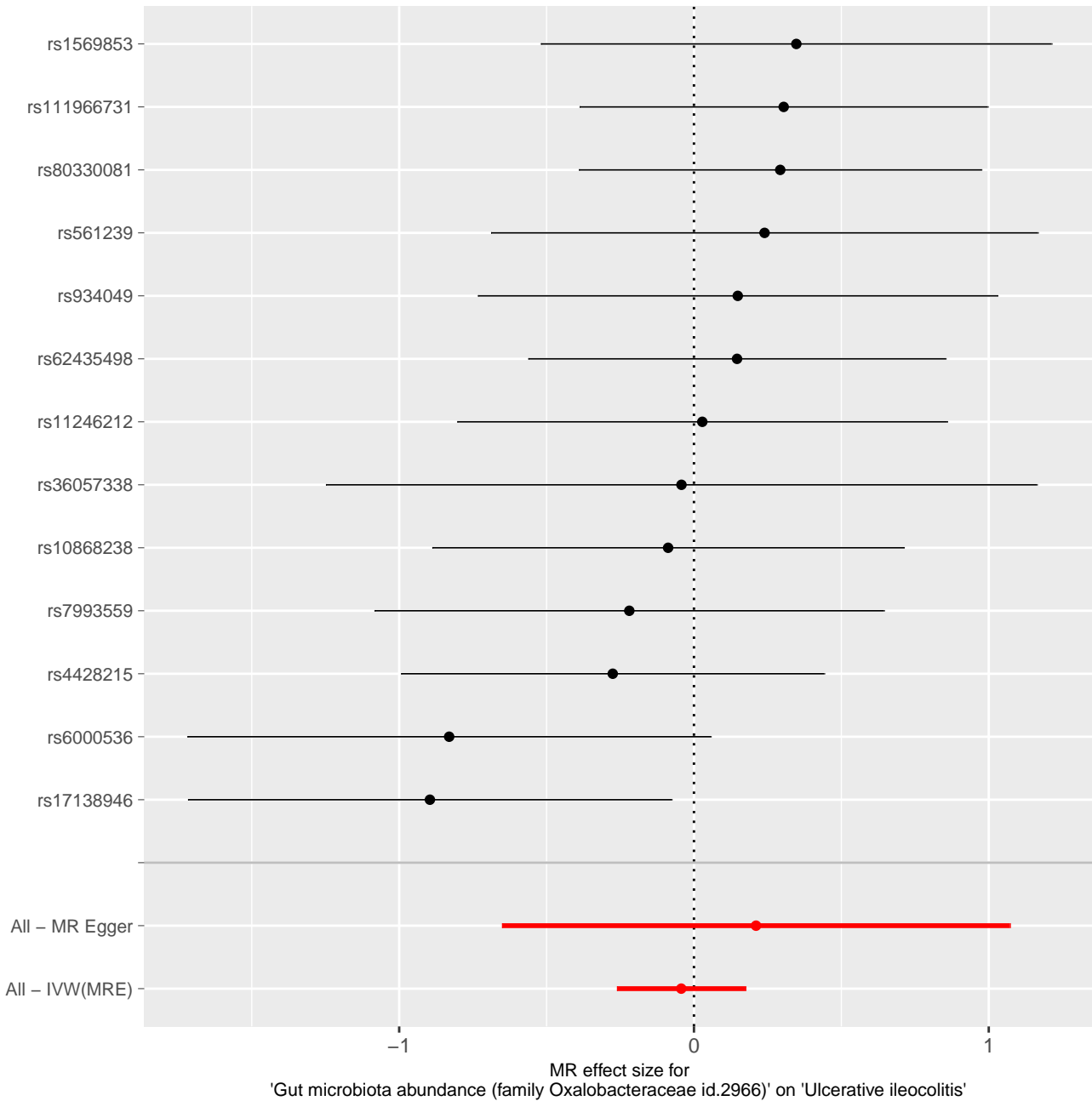


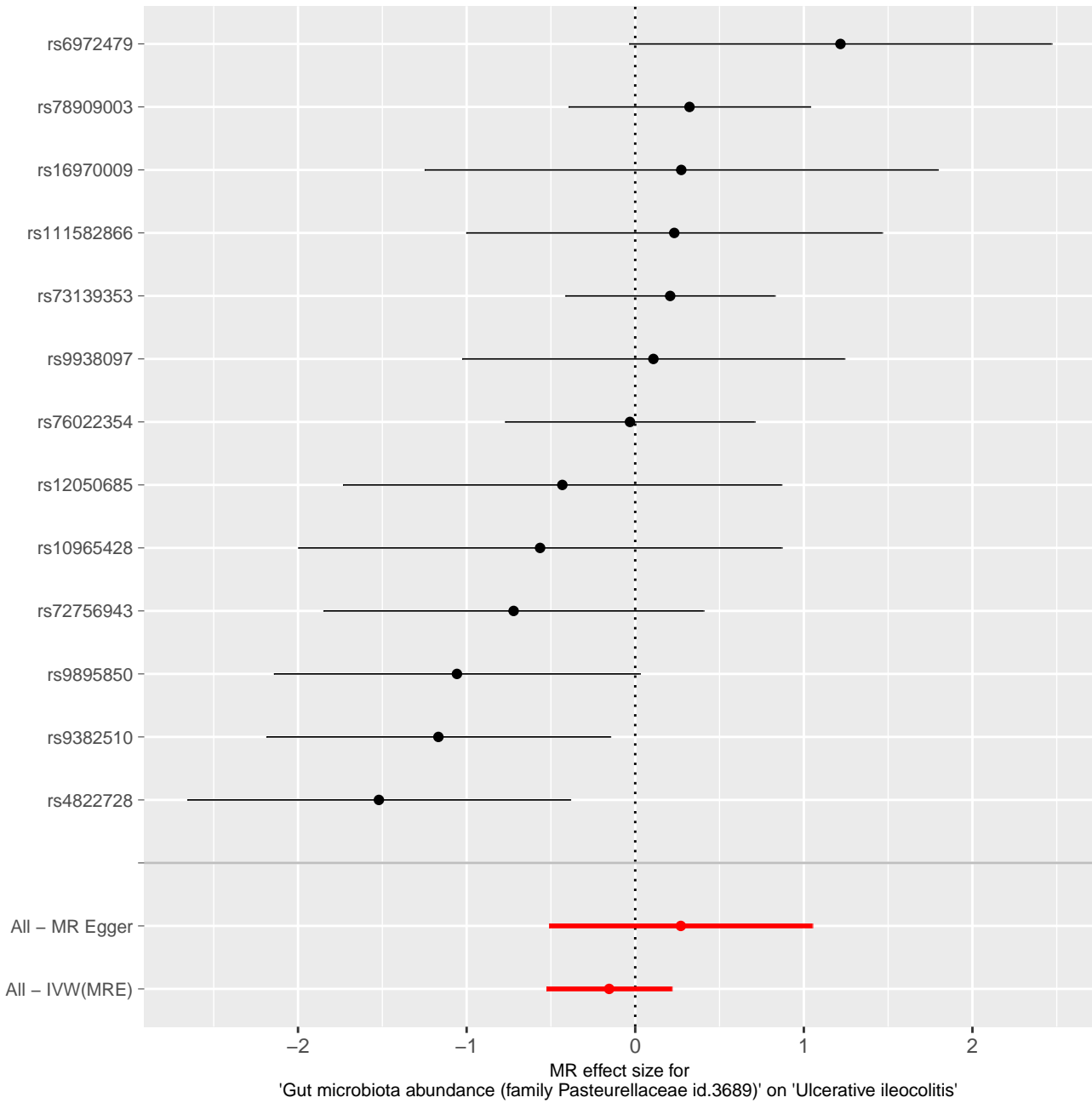


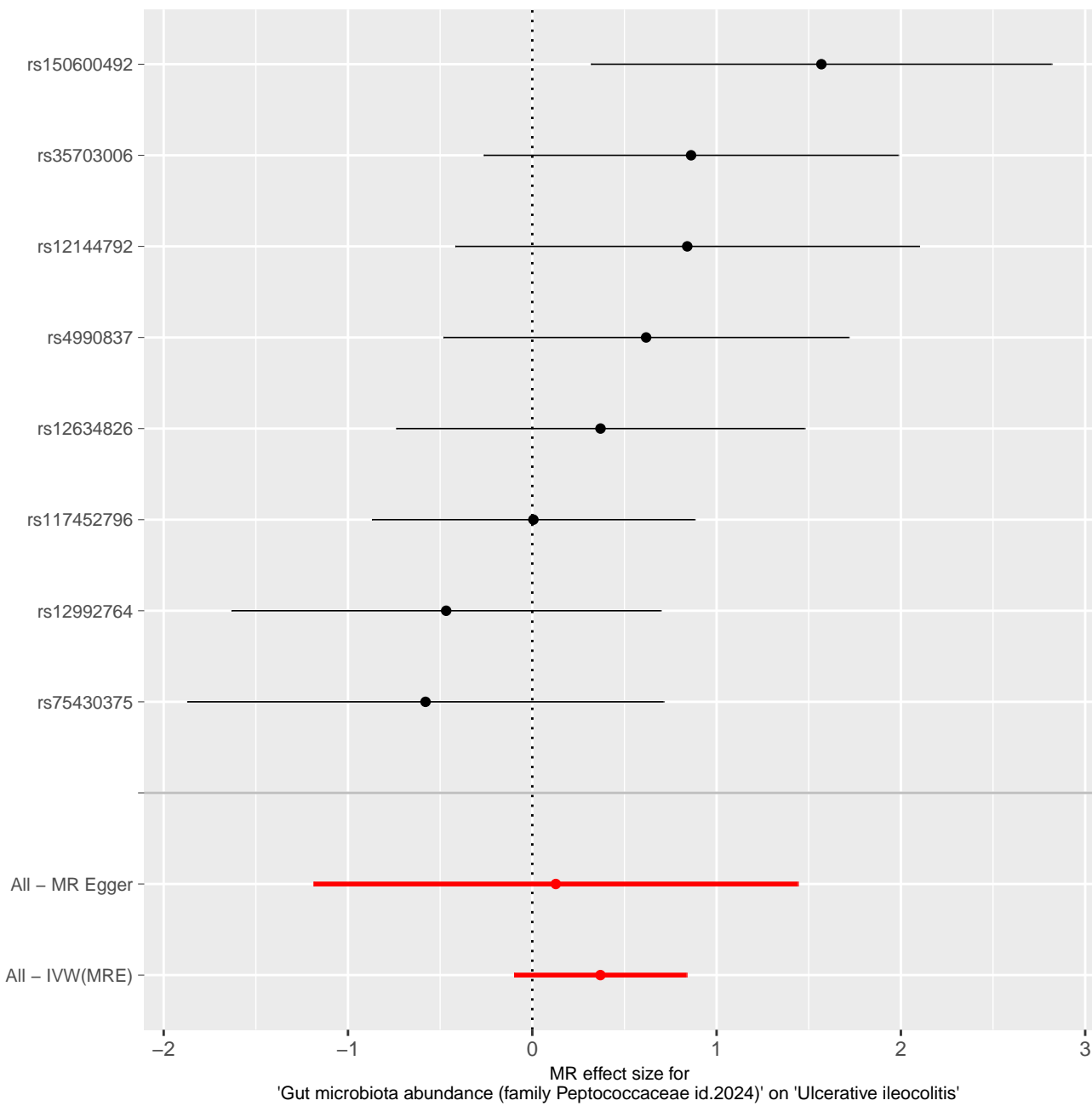


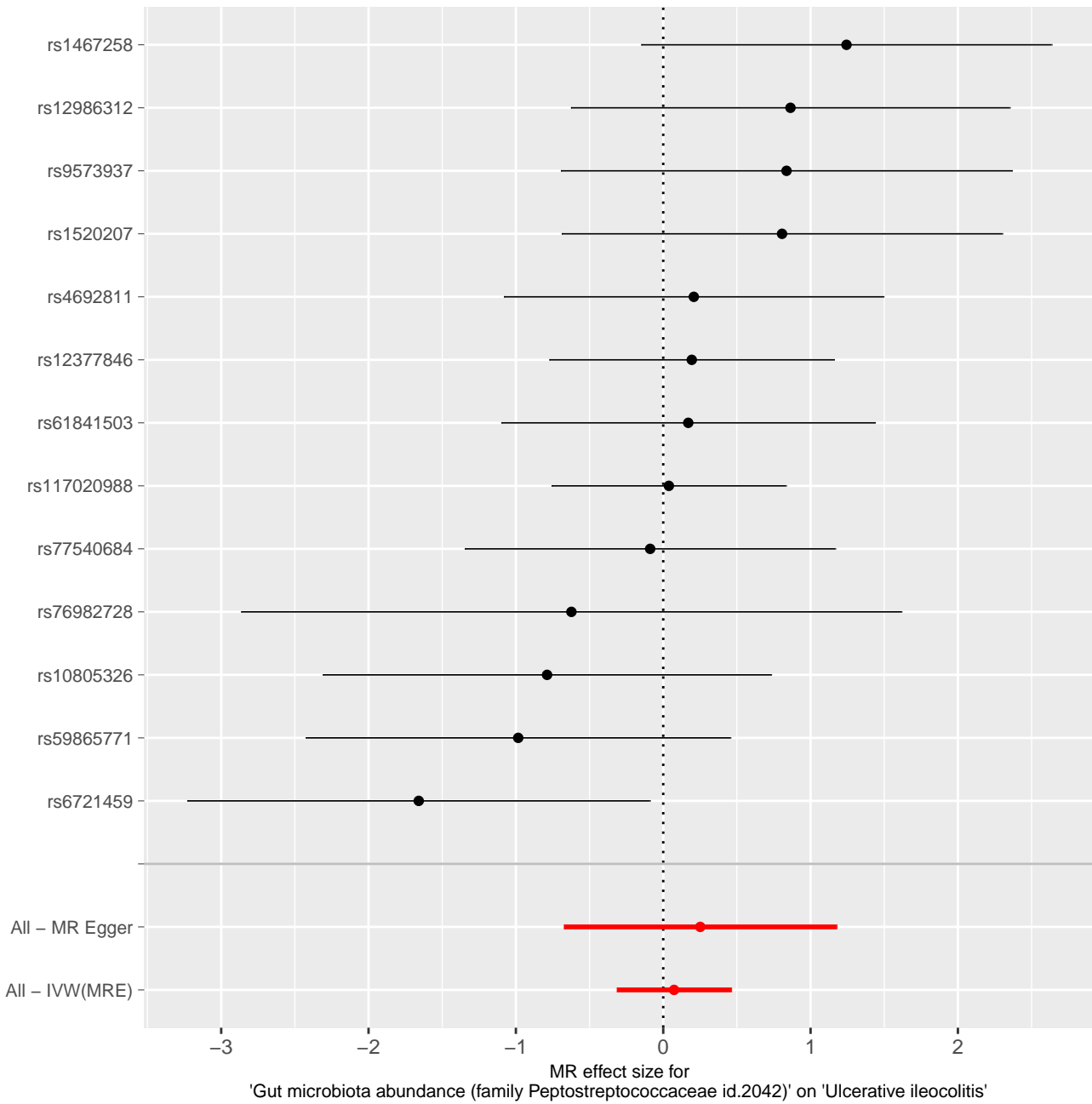


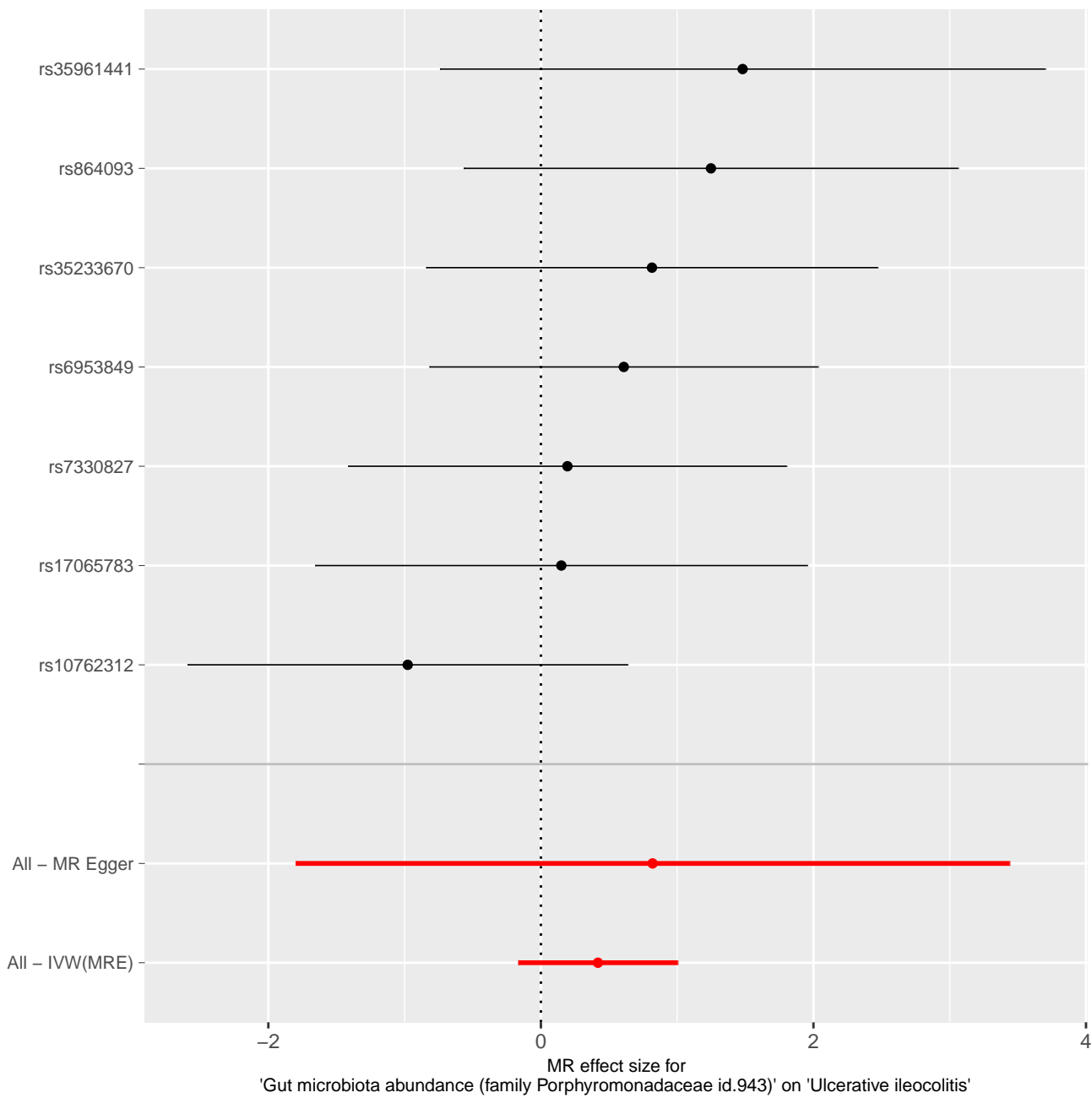


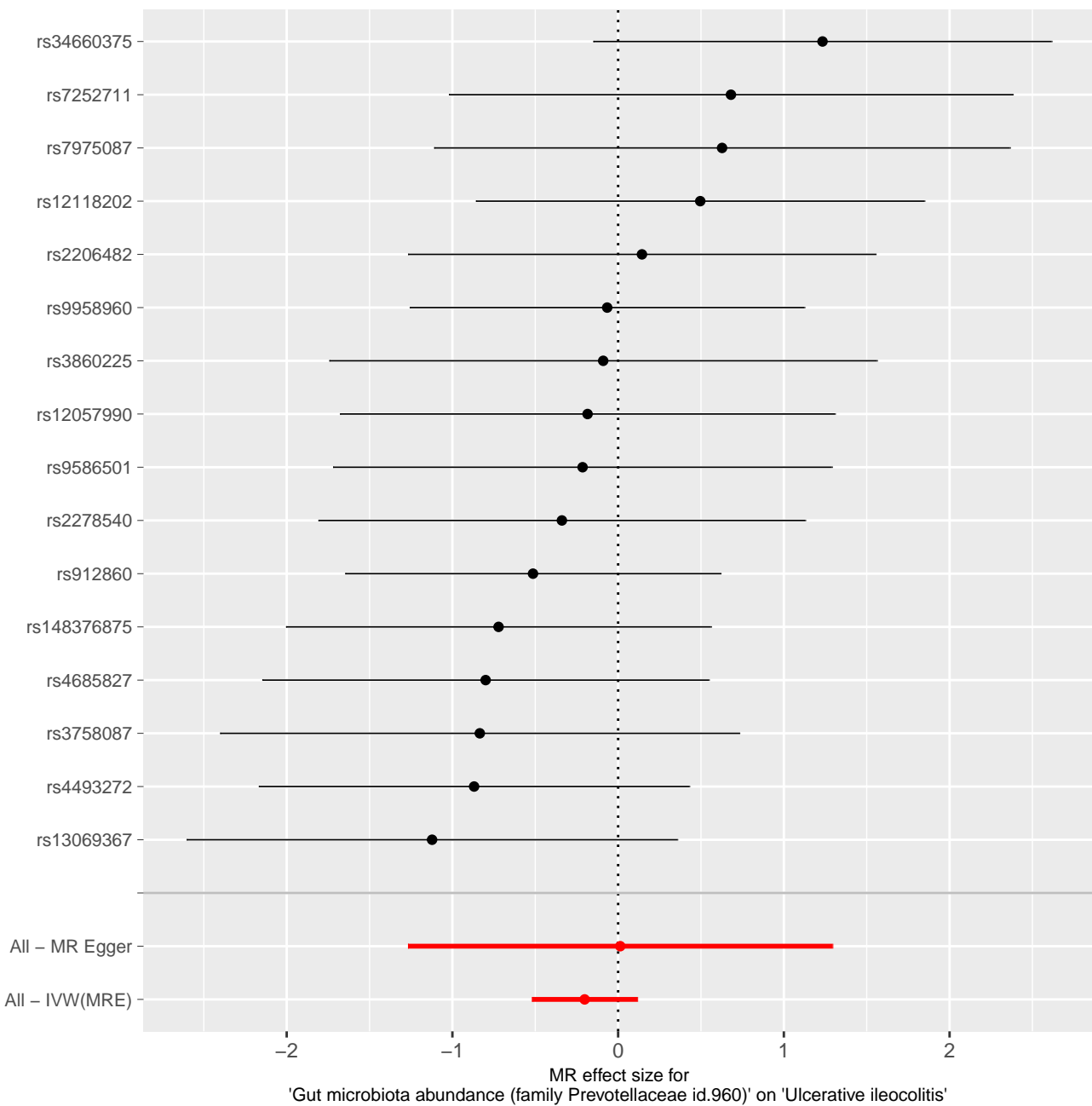


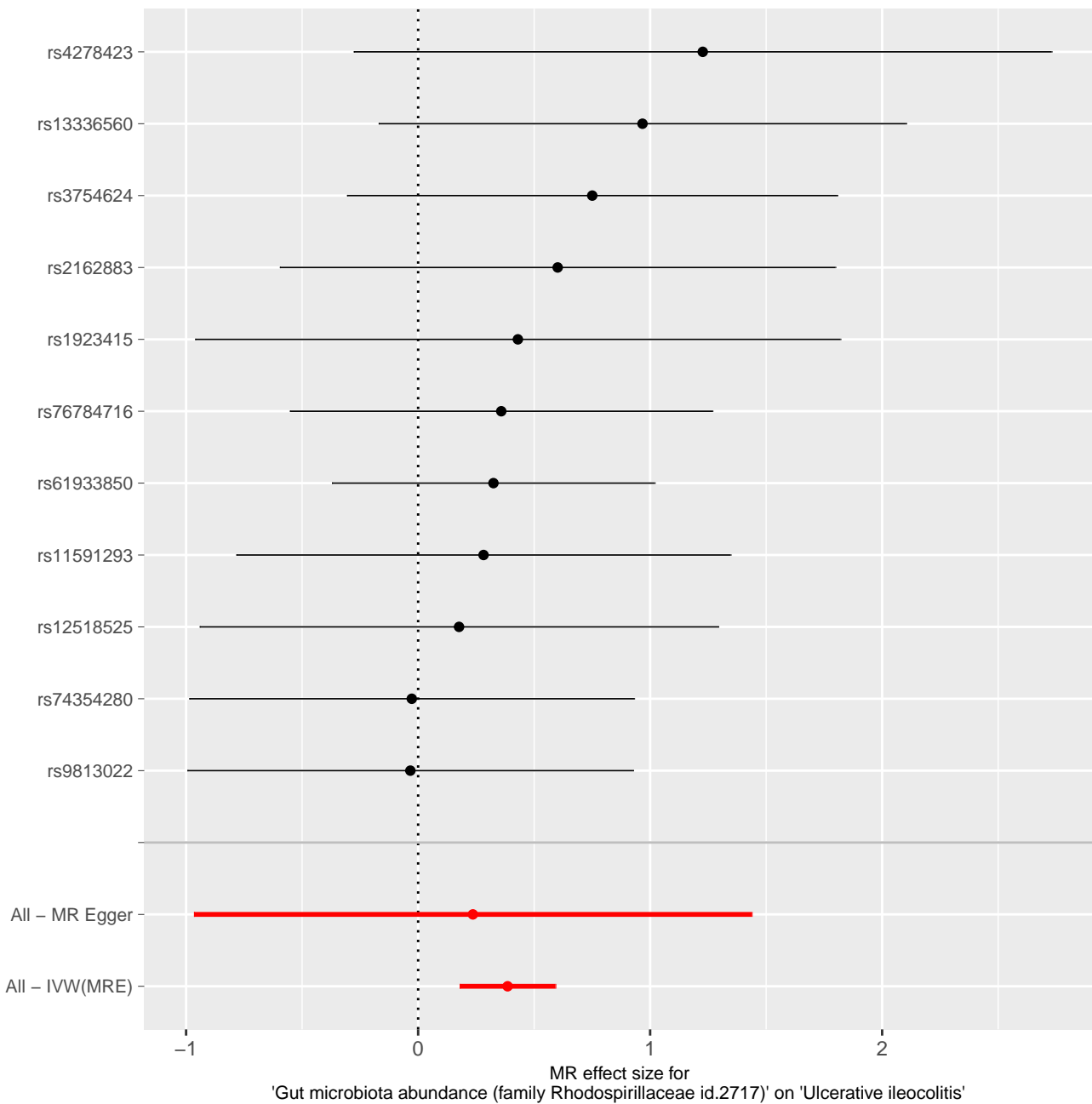




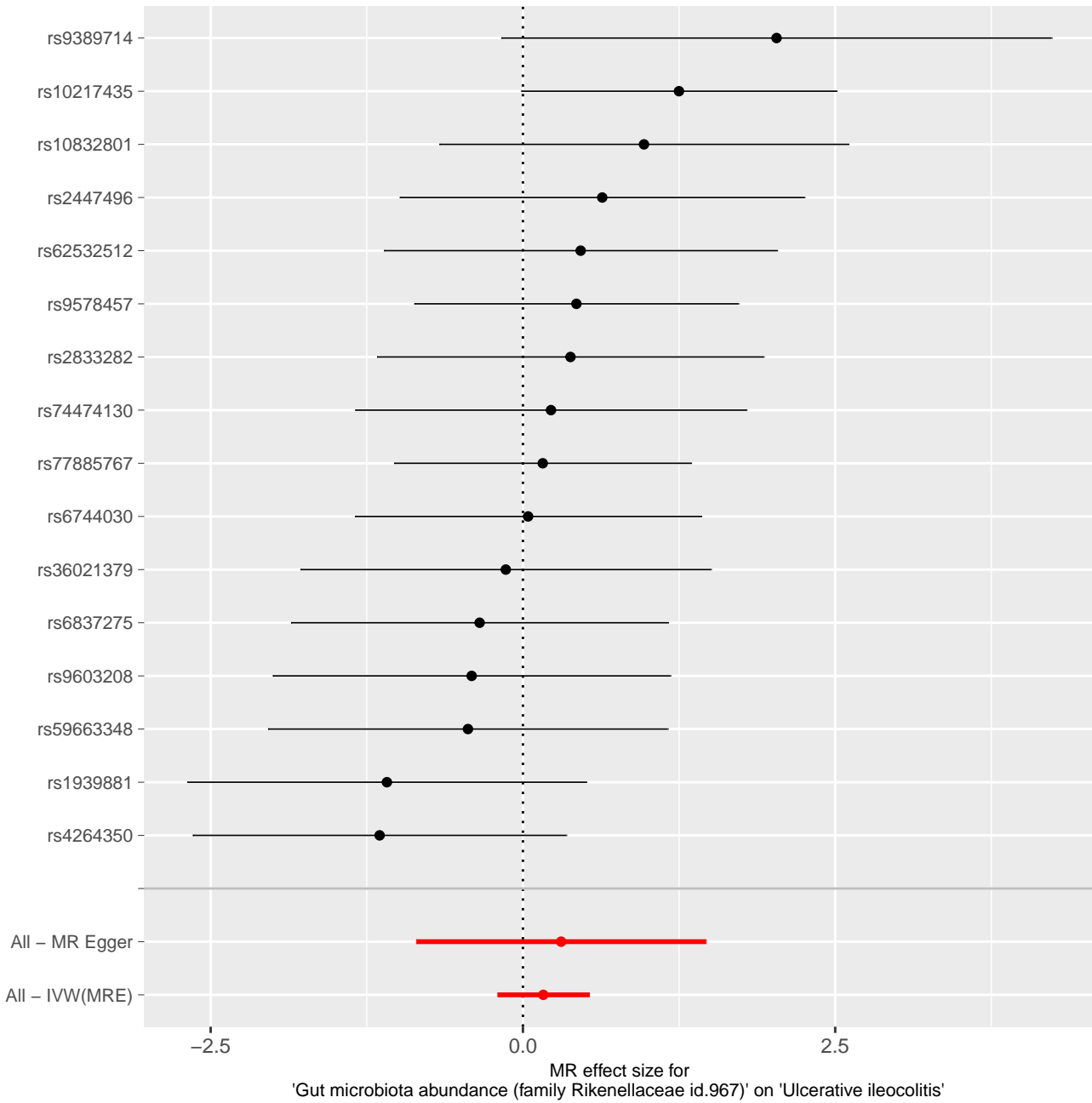


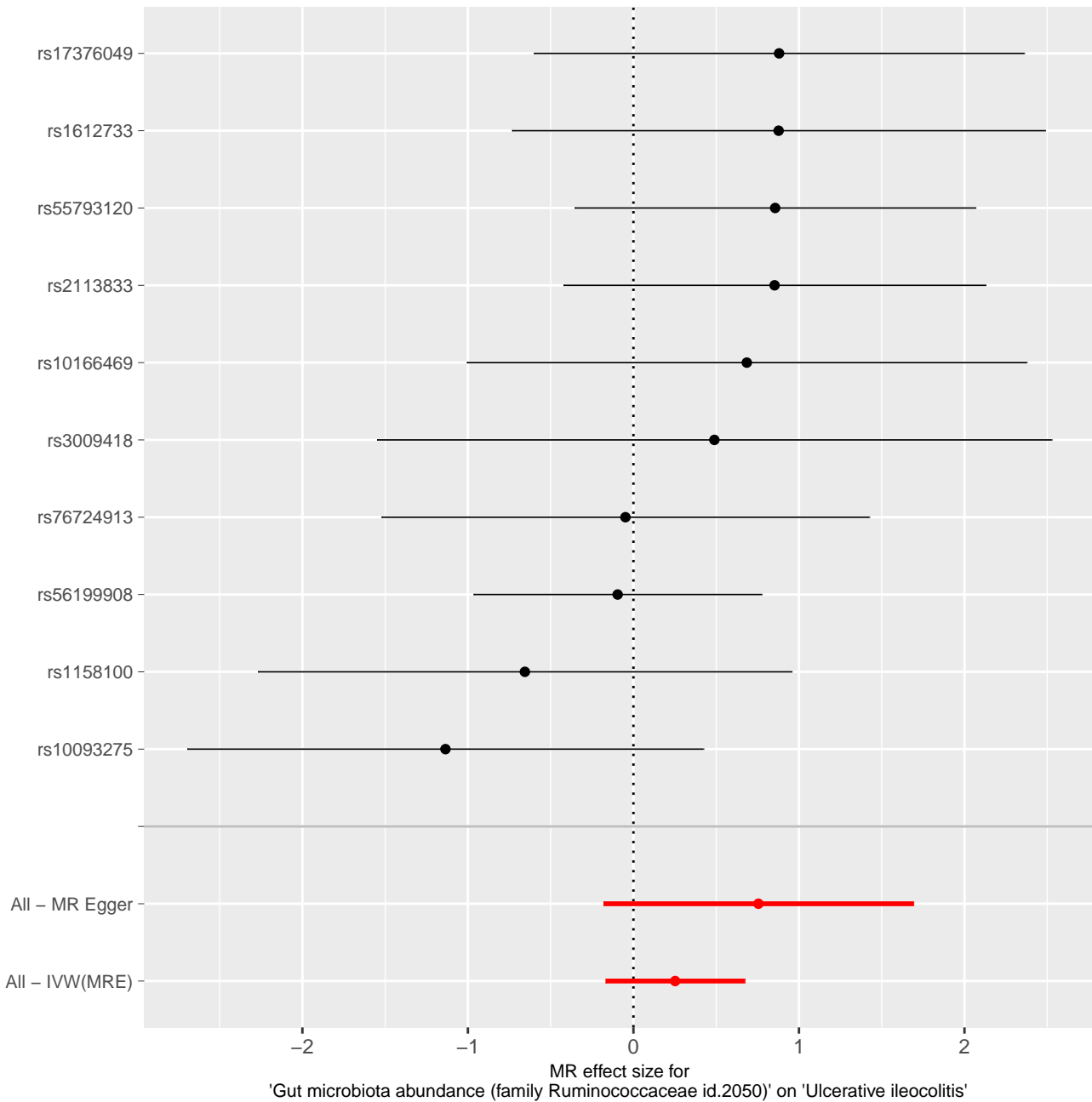


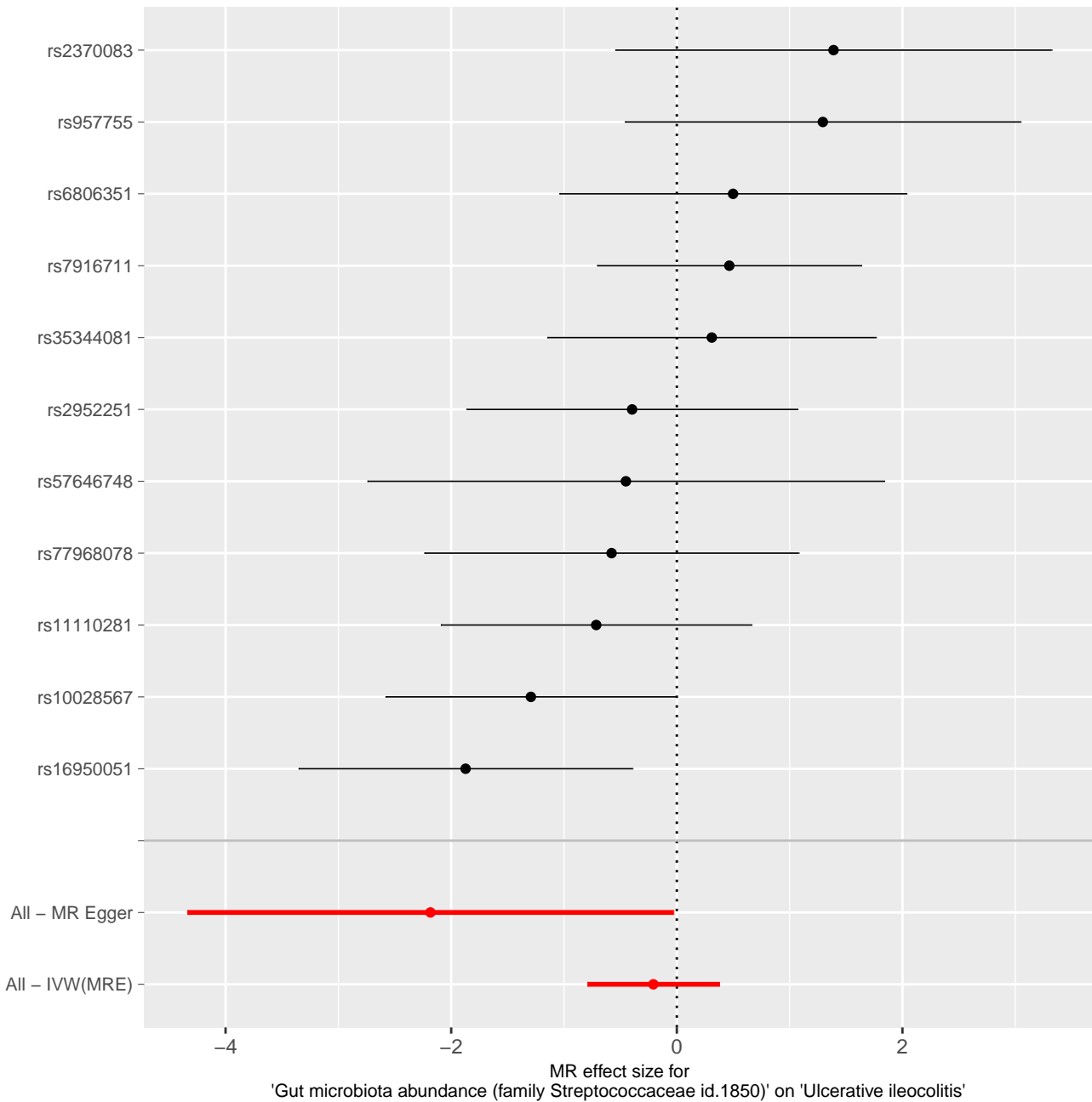


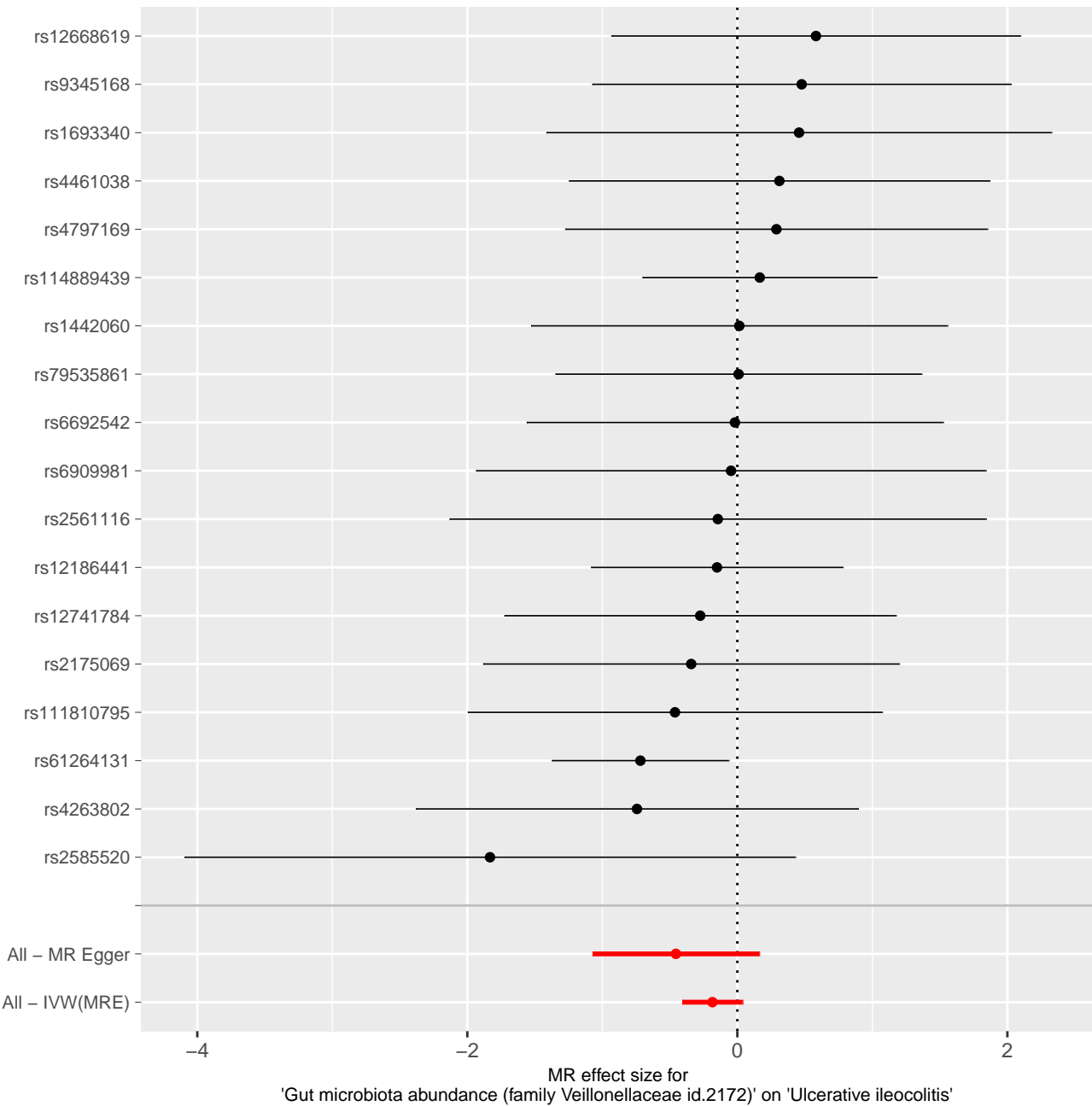


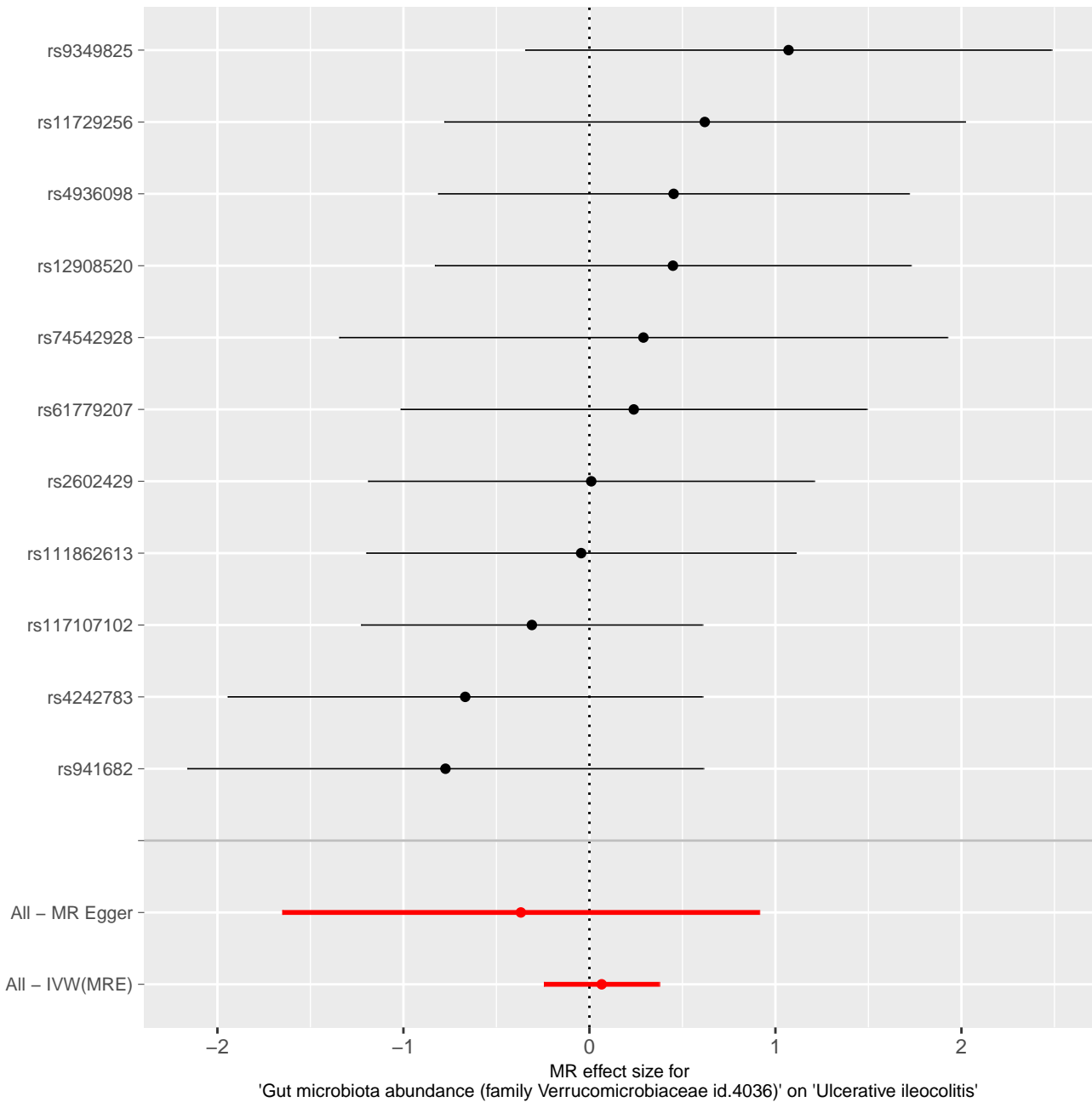
Batch 1077 : Gut microbiota abundance (family Rikenellaceae id.967) on Ulcerative ileocolitis

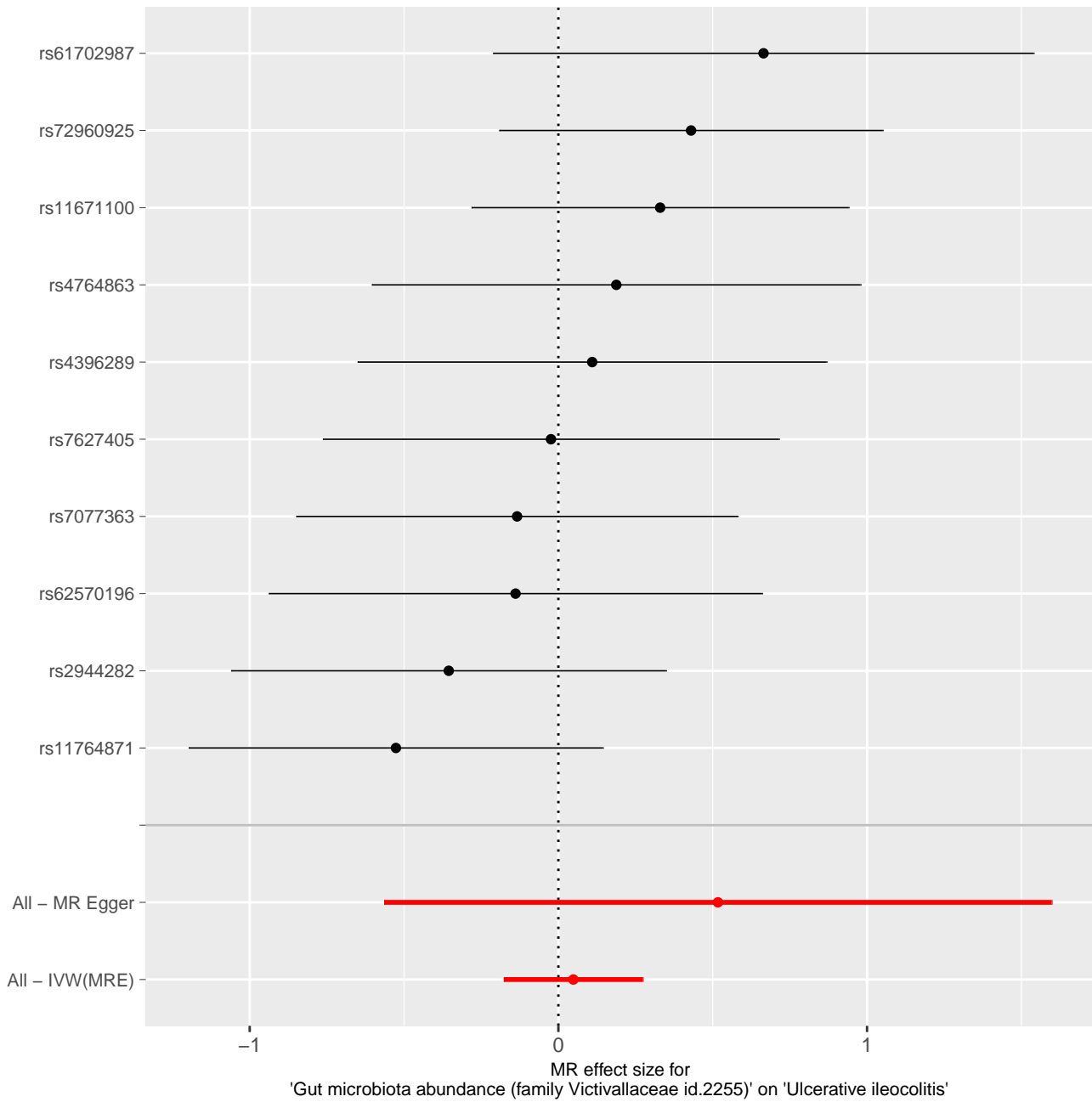


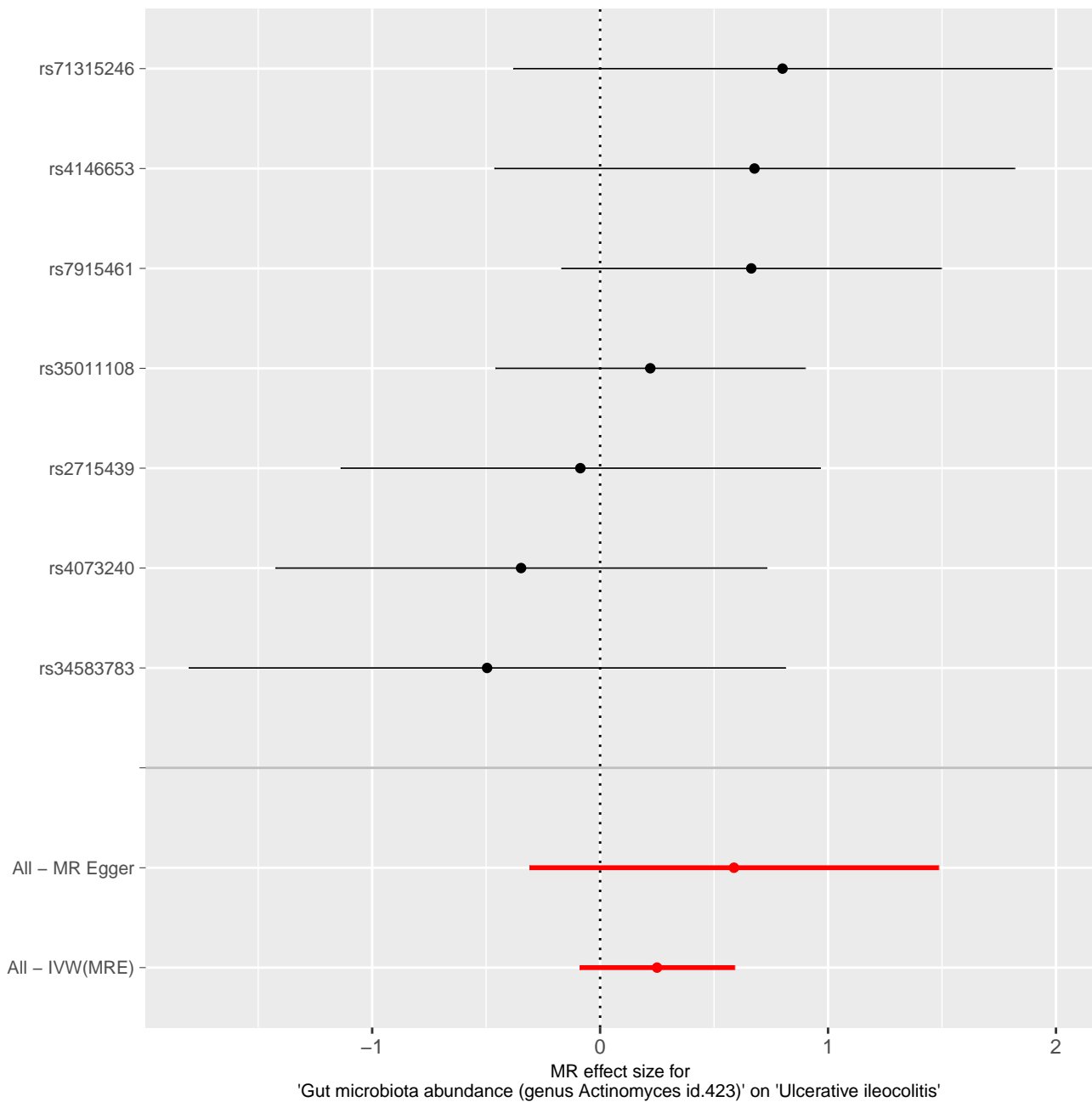


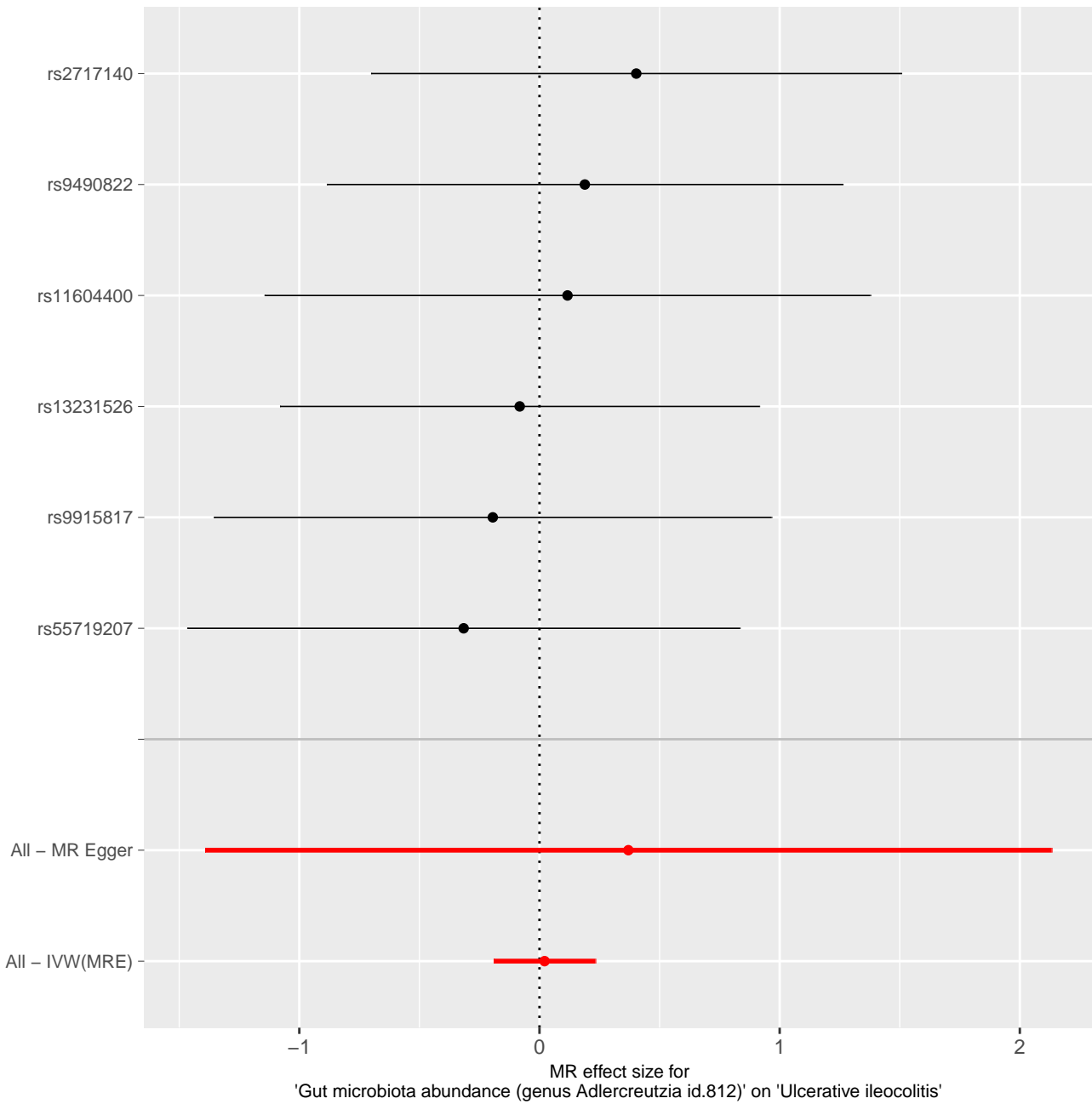




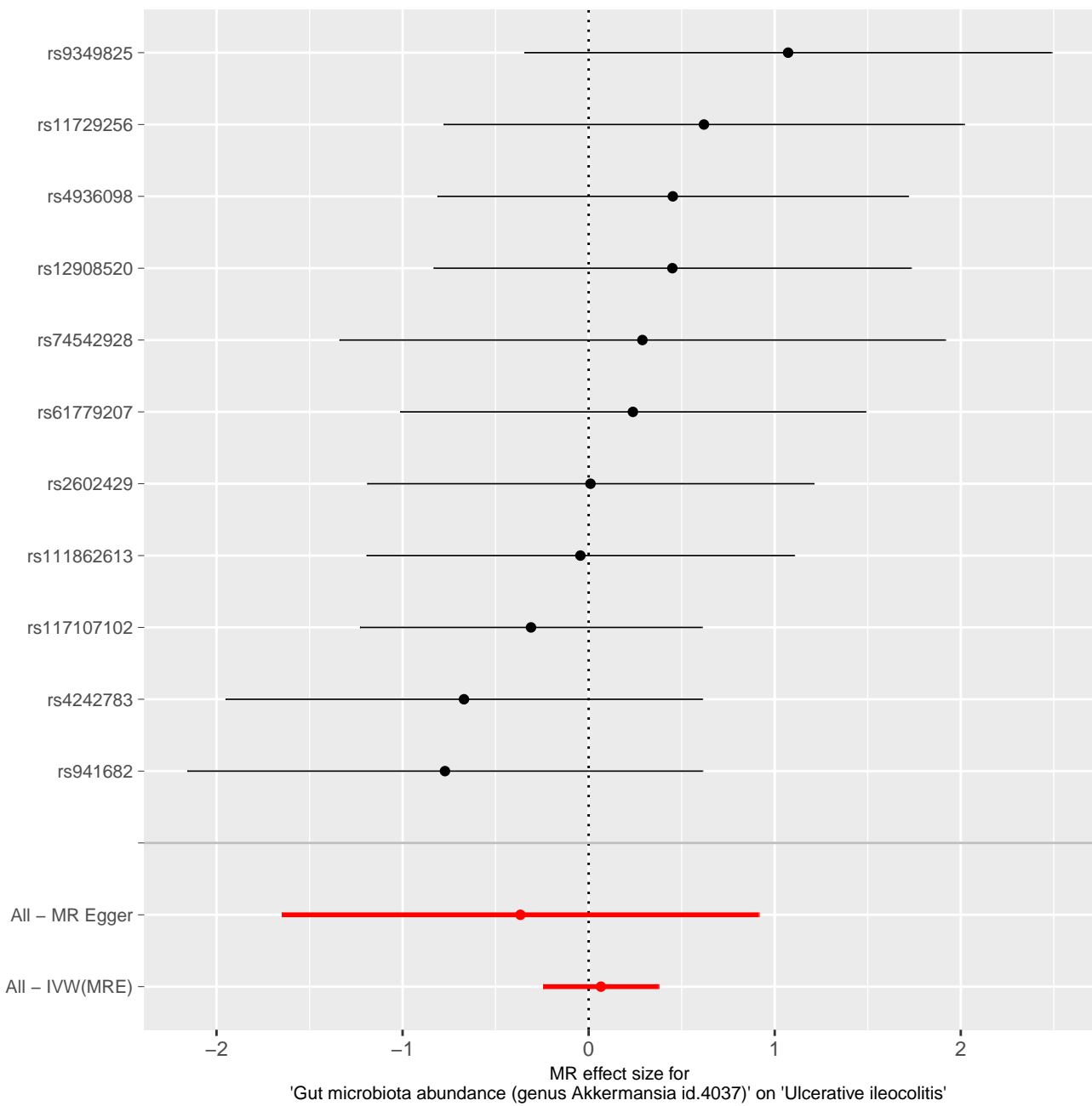


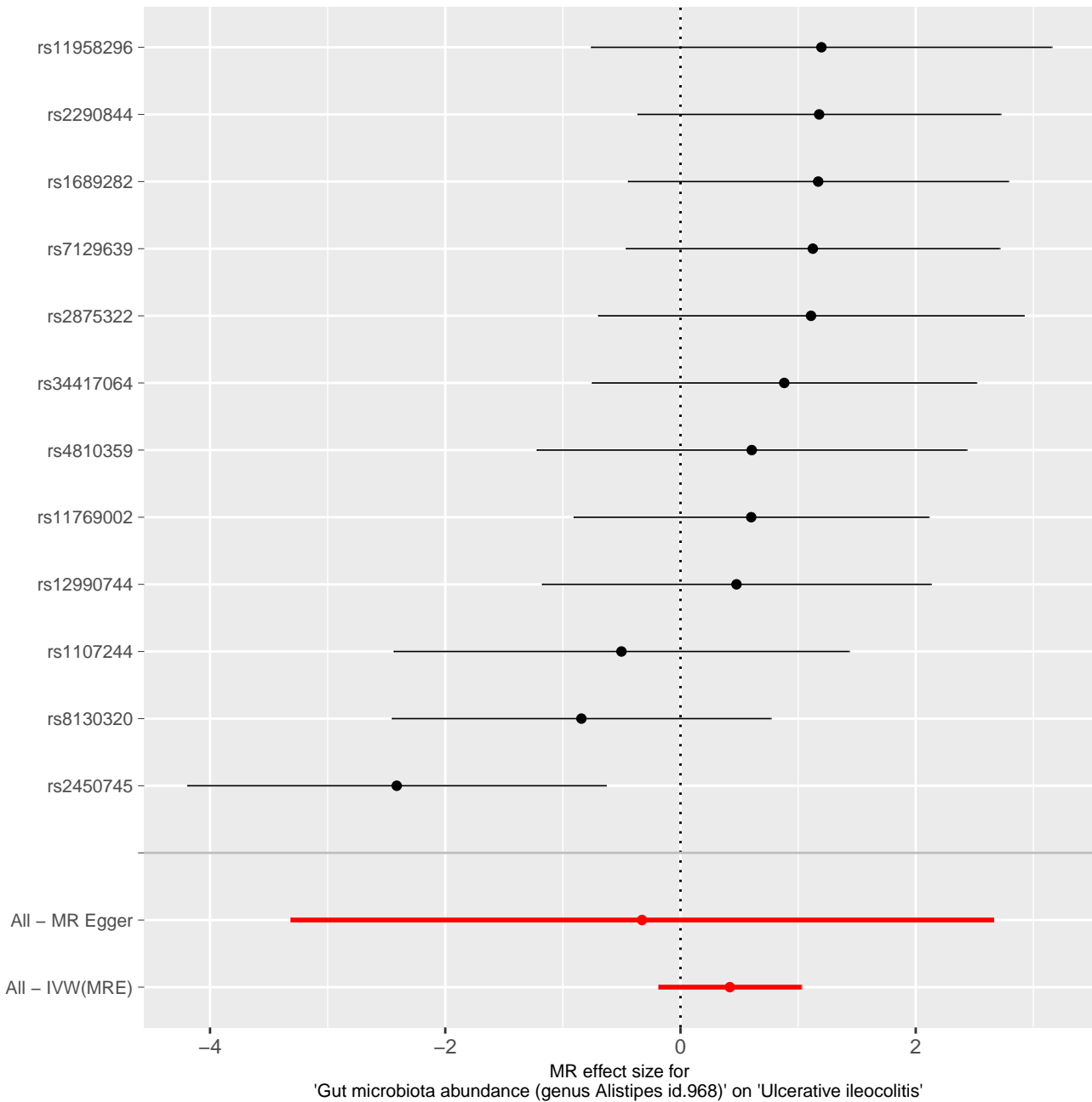


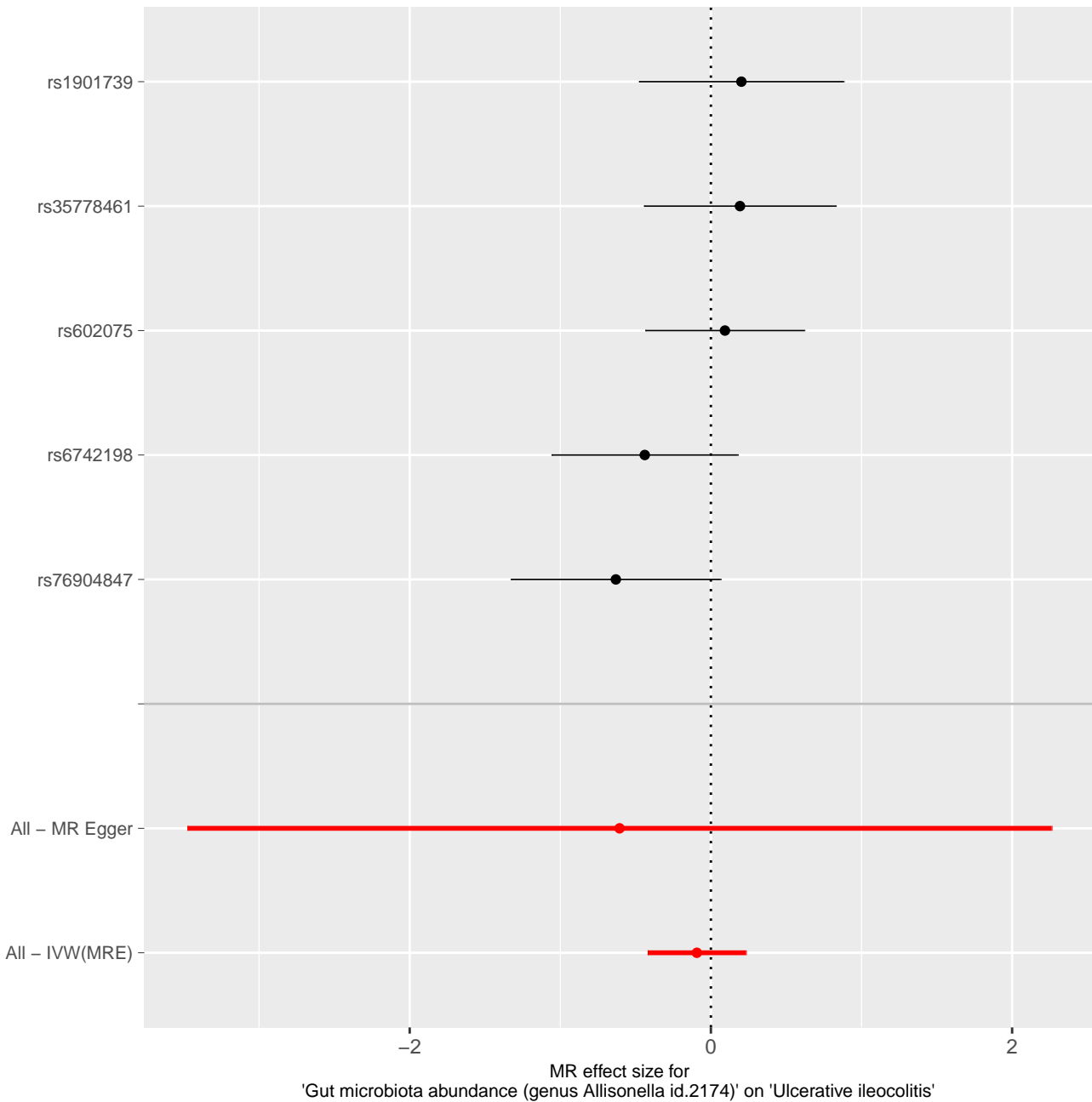


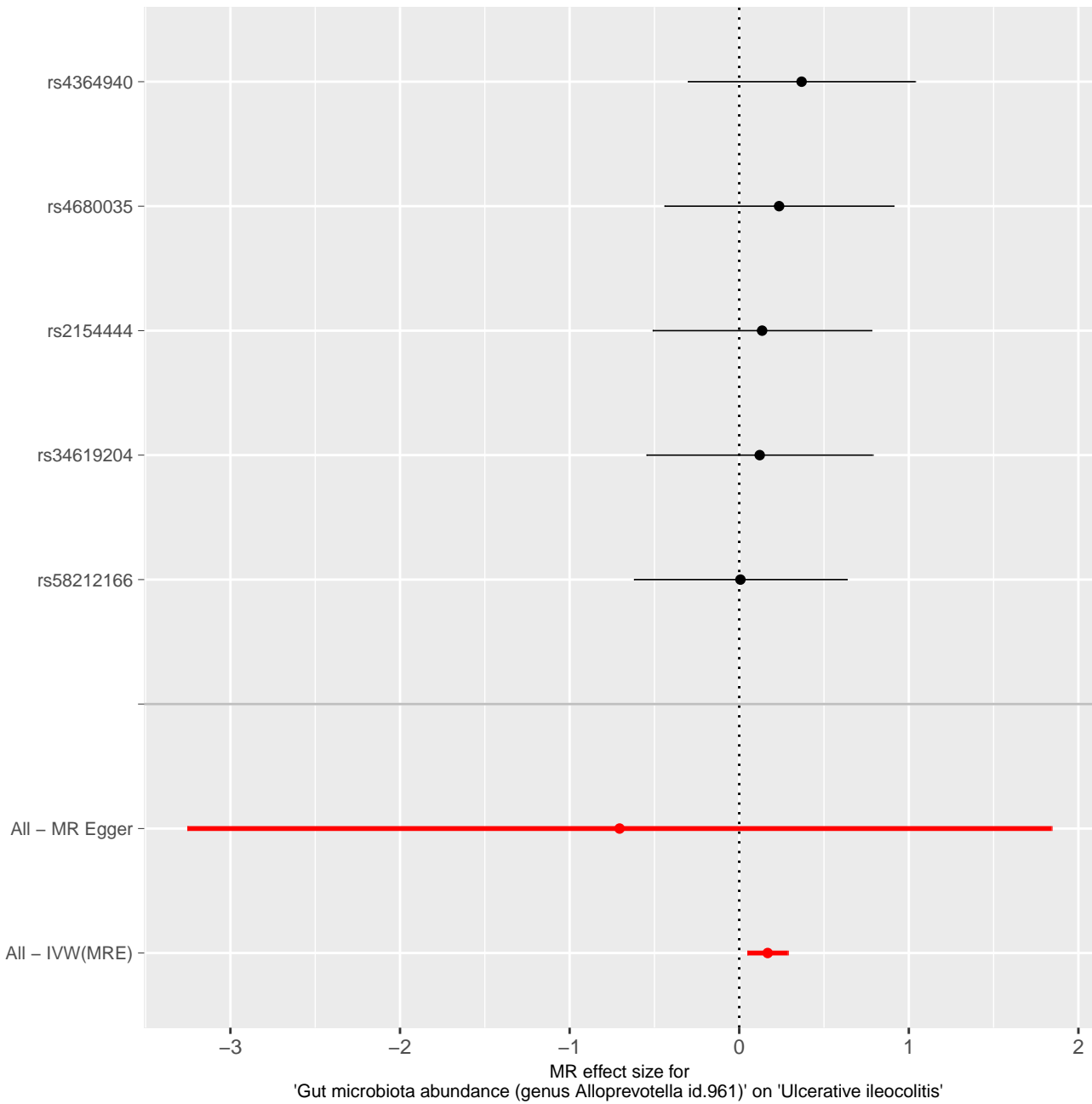


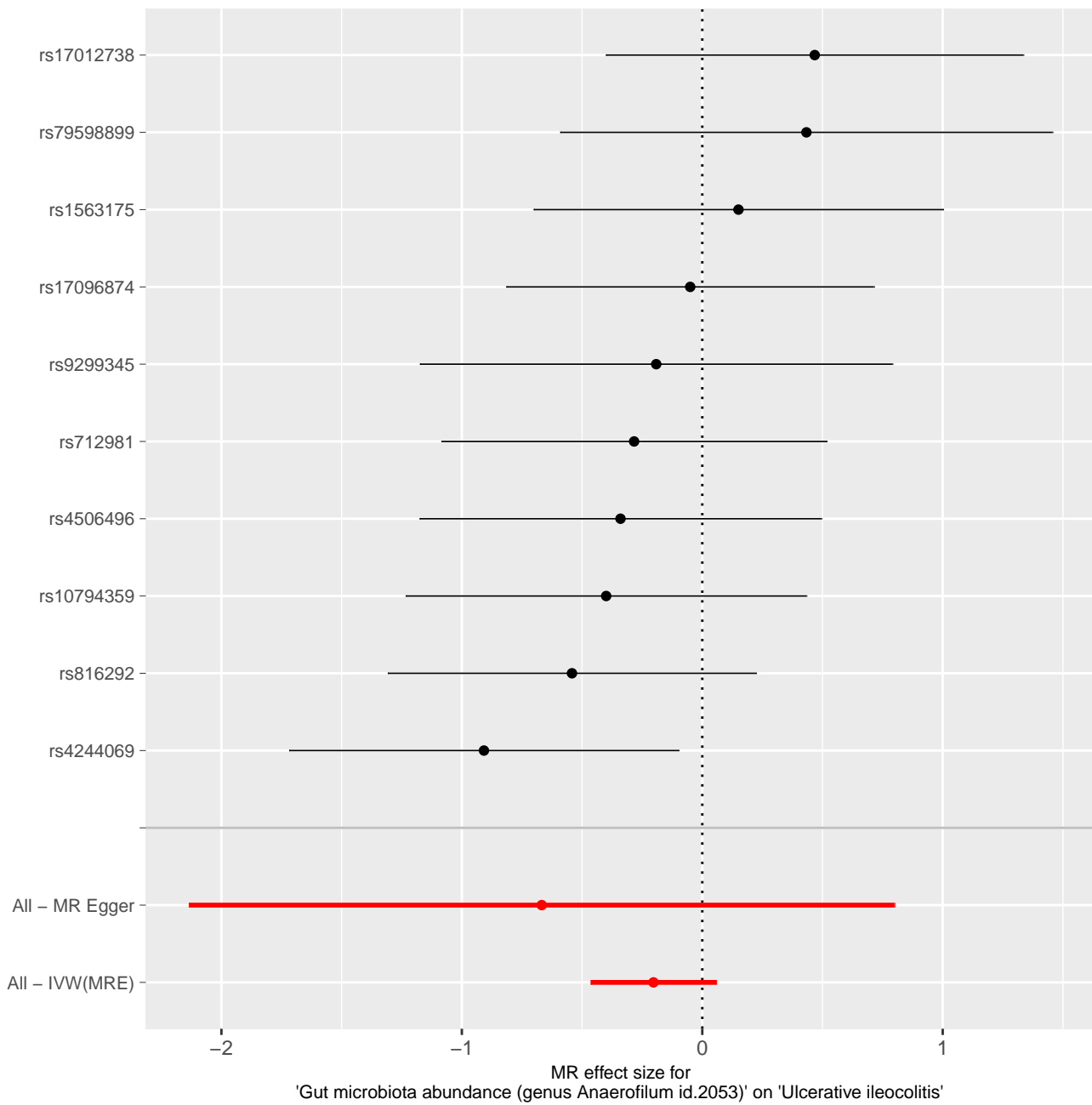
Batch 1085 : Gut microbiota abundance (genus Akkermansia id.4037) on Ulcerative ileocolitis

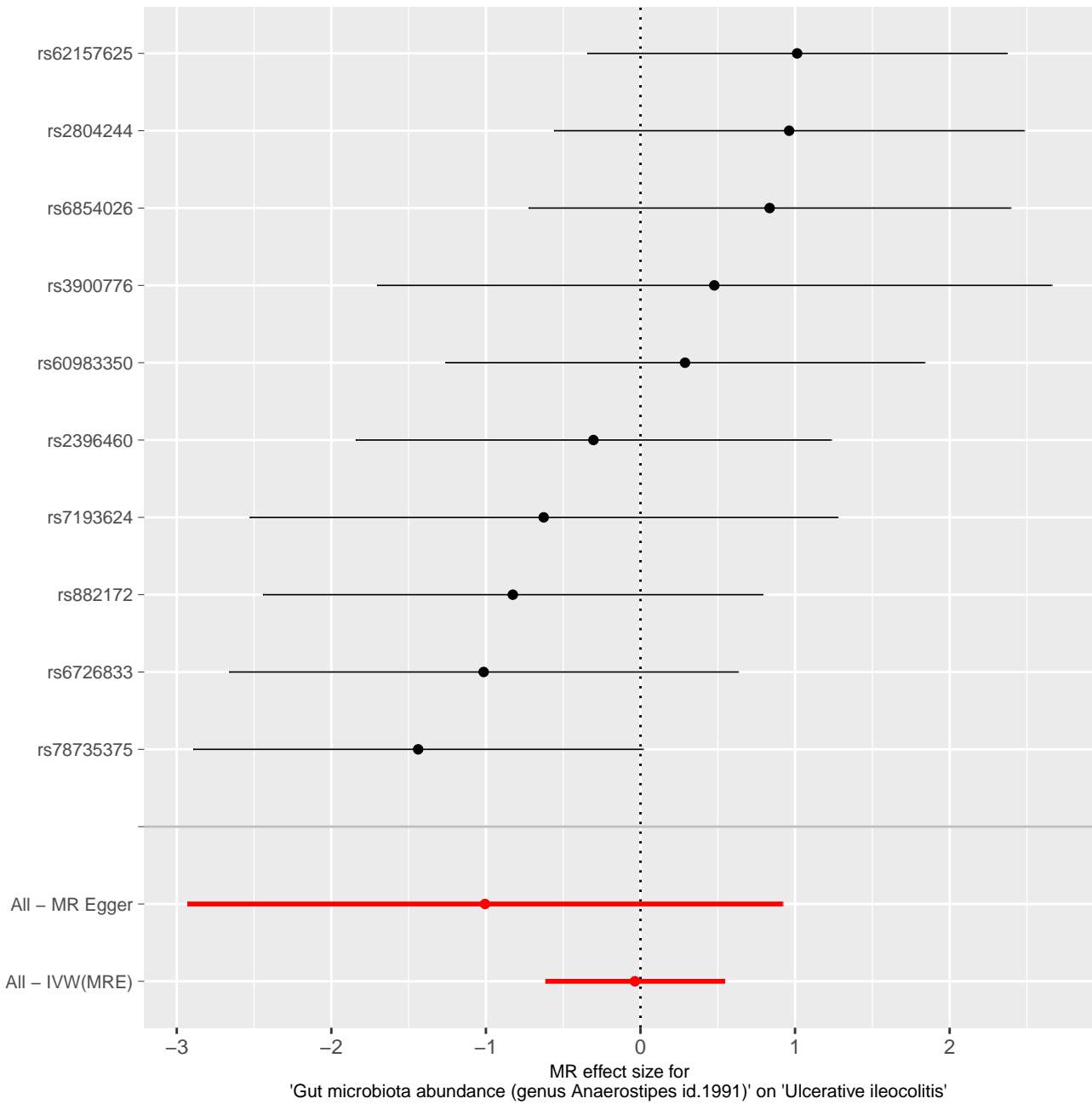


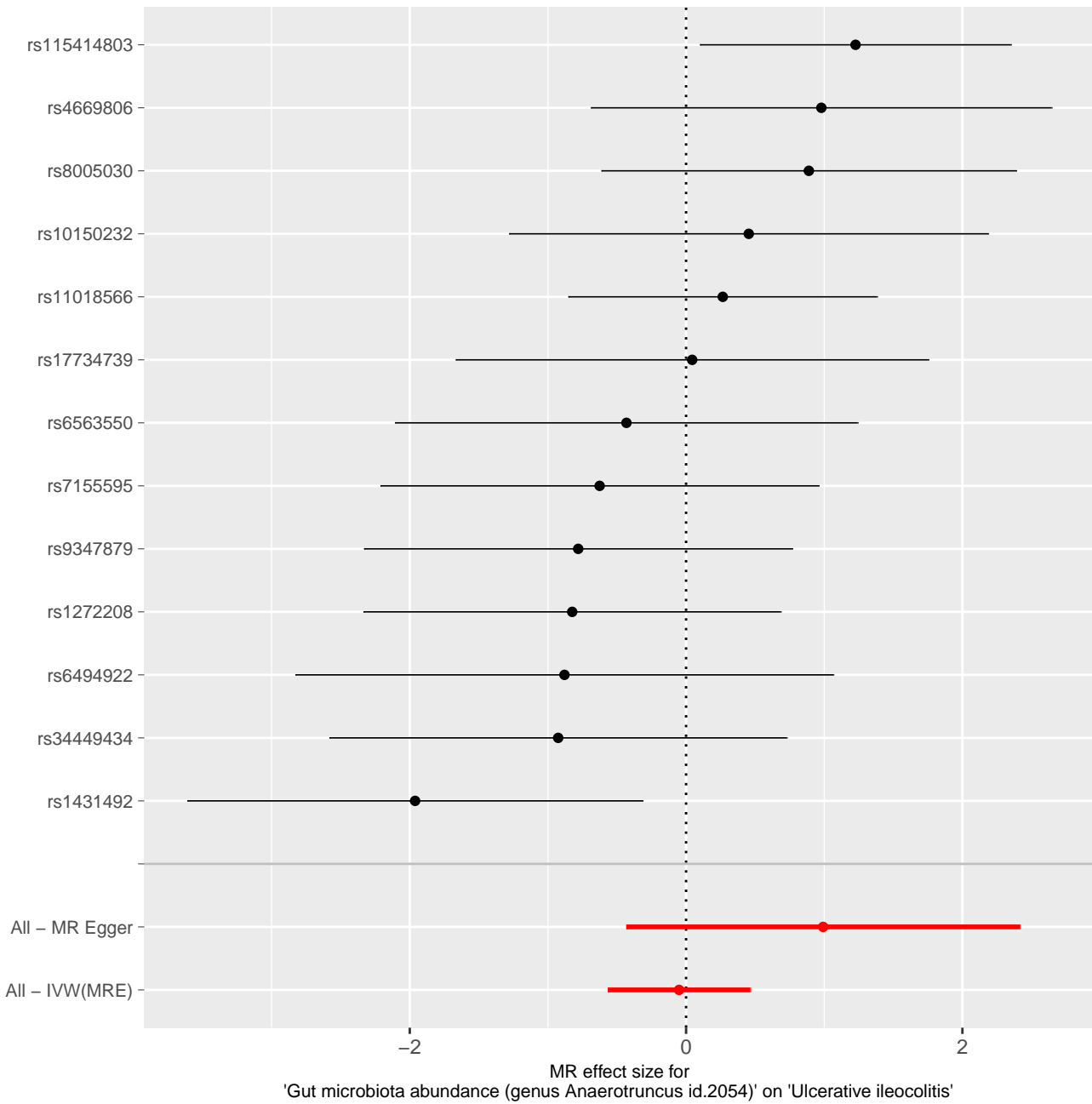


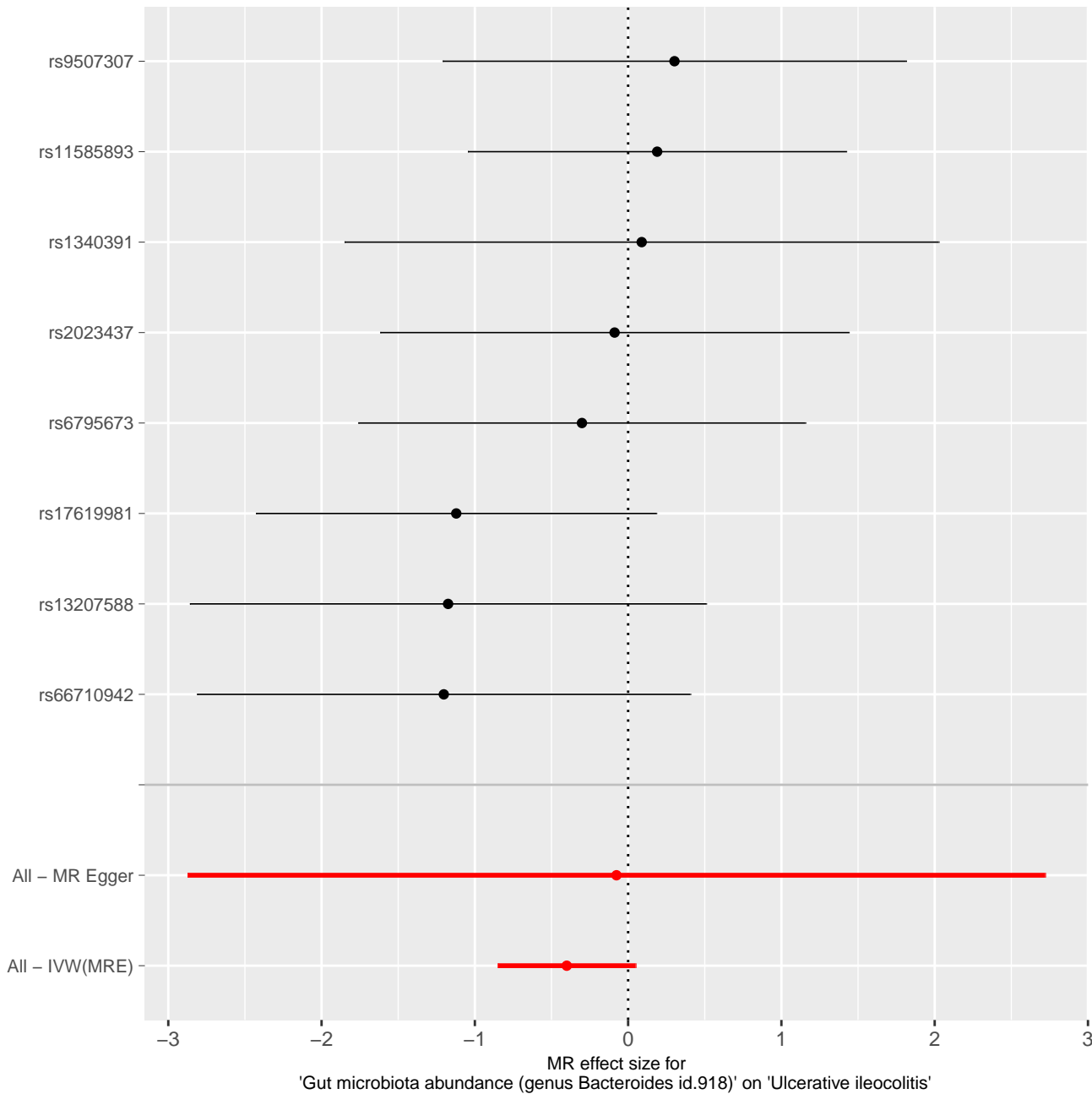




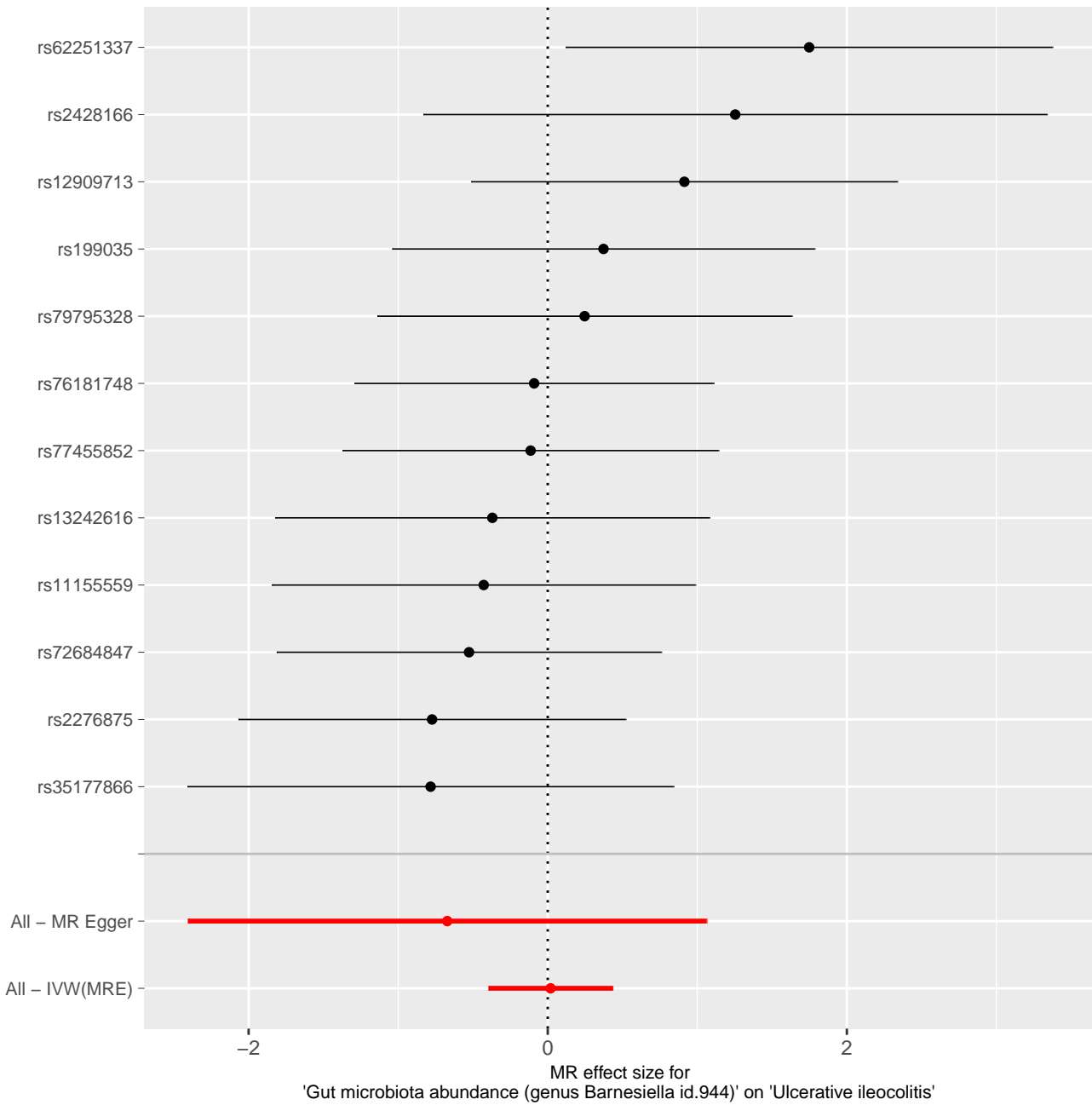


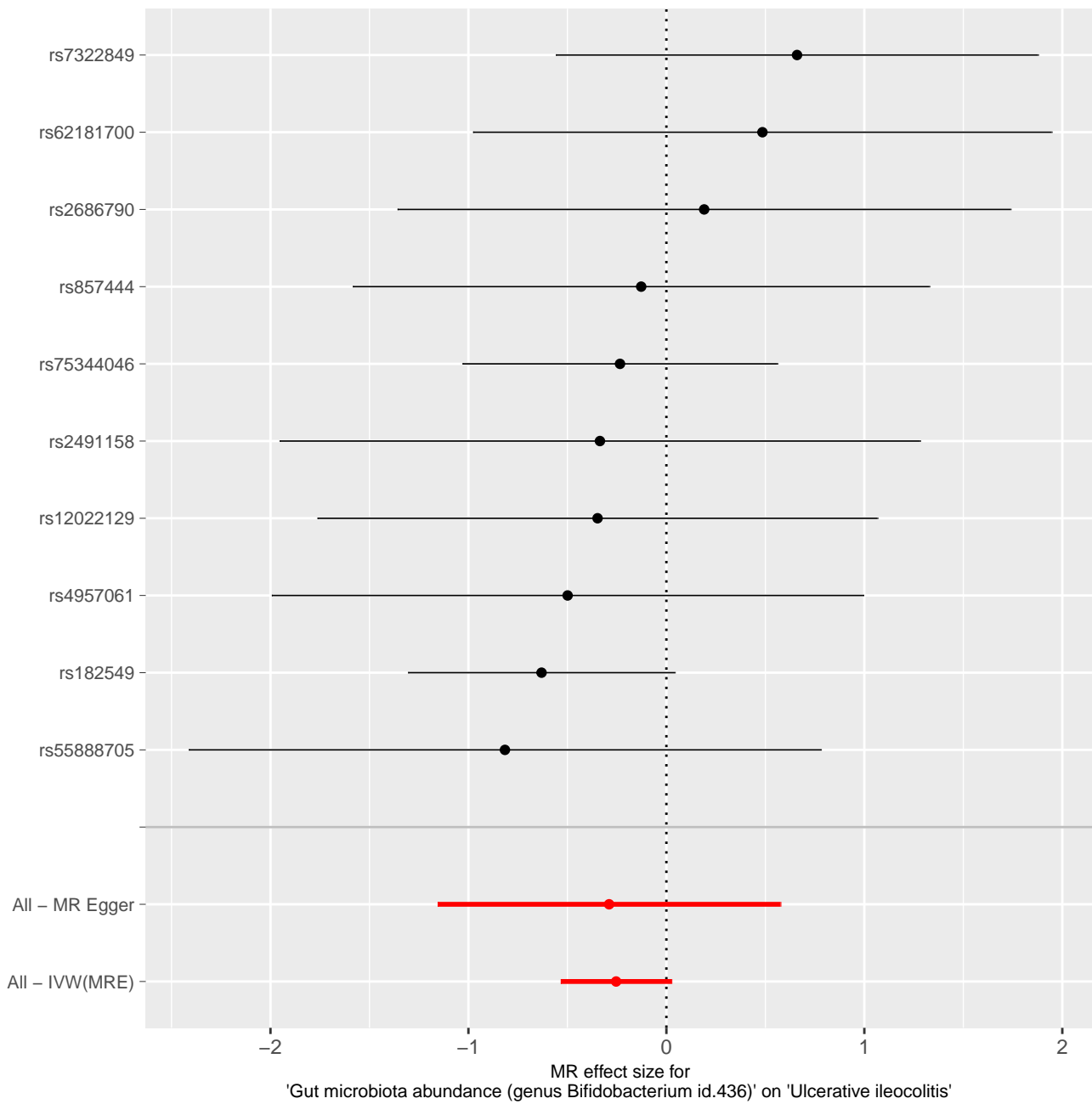


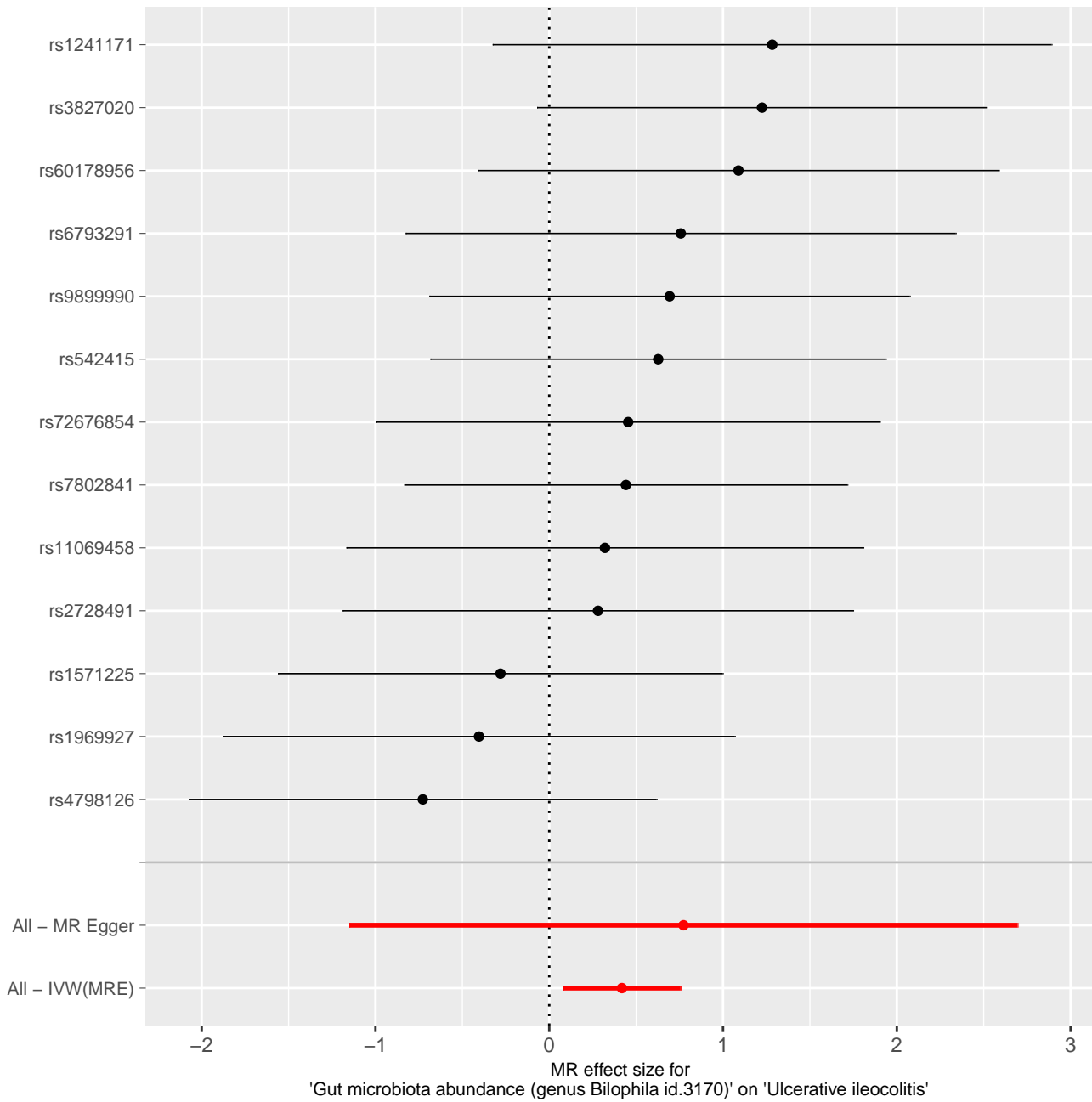


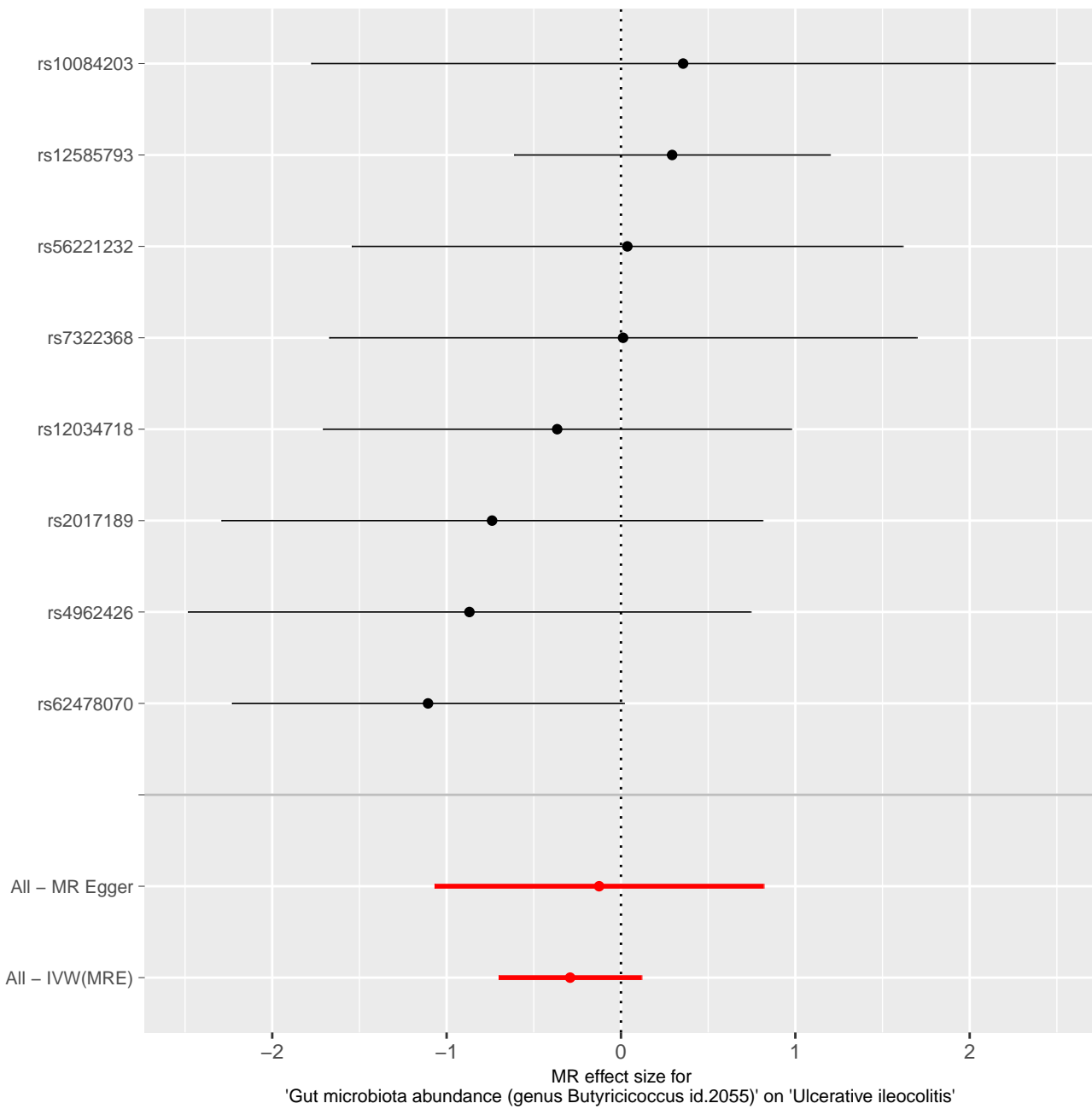


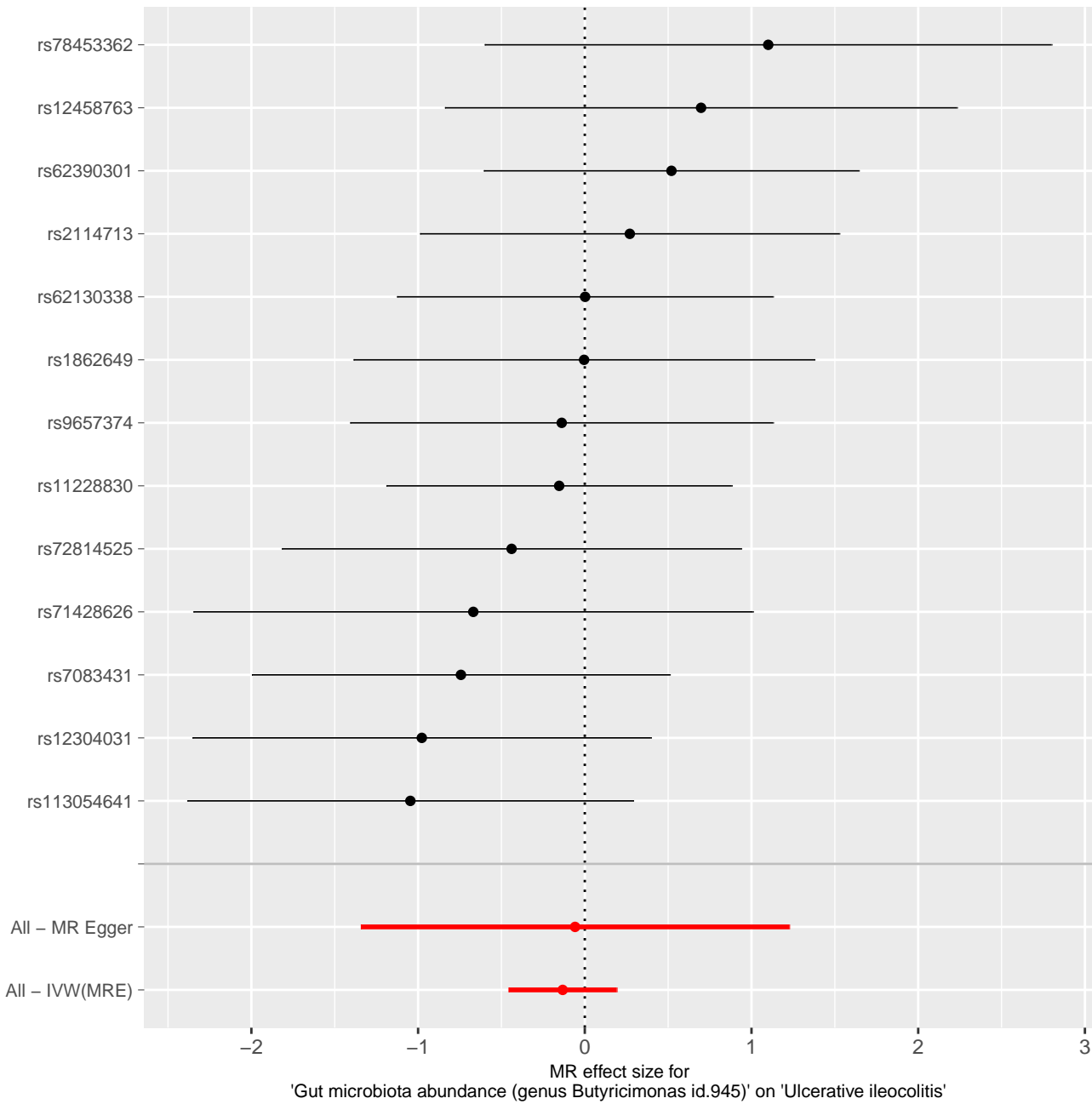
MR effect size for
'Gut microbiota abundance (genus Bacteroides id.918)' on 'Ulcerative ileocolitis'



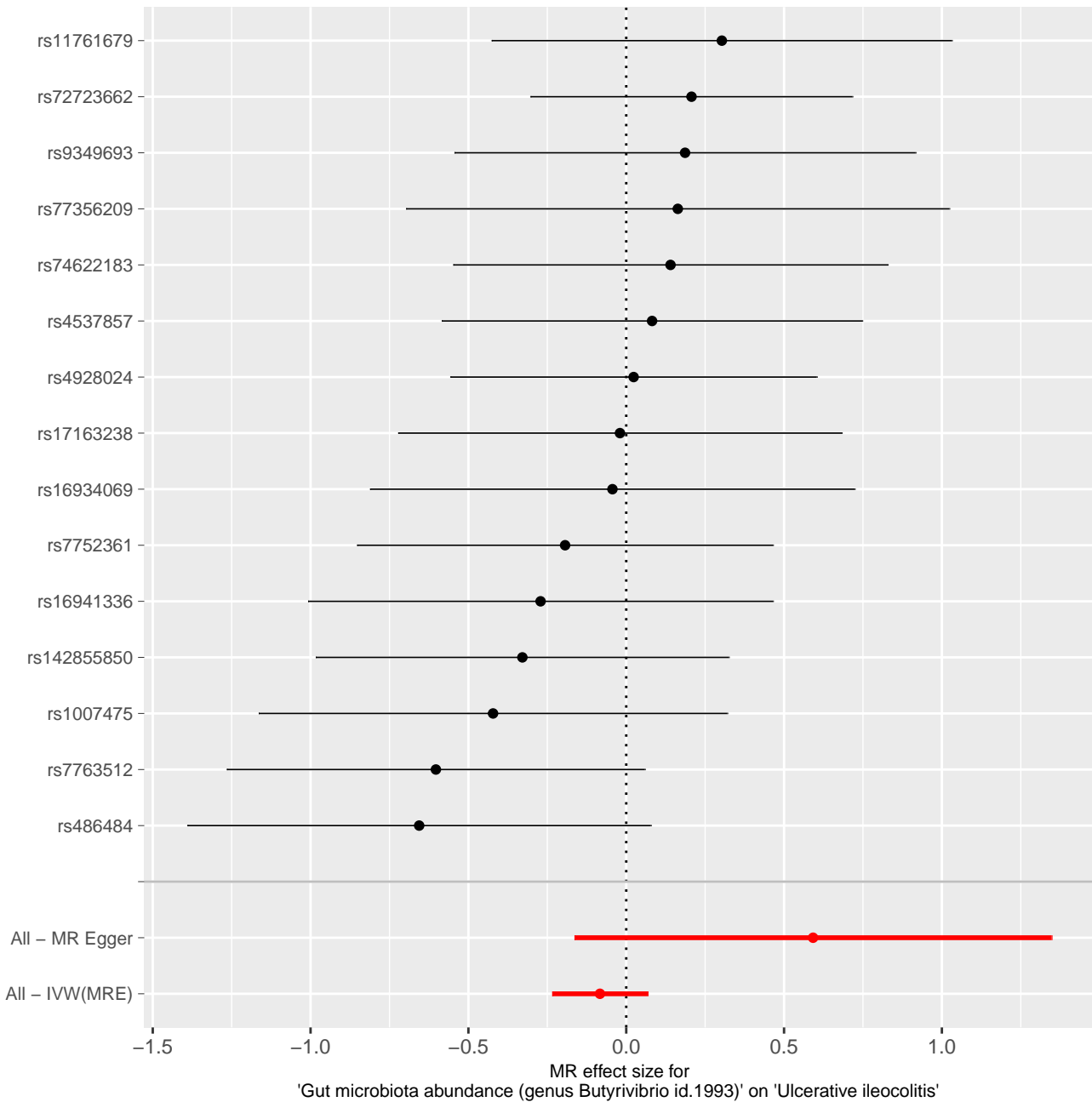


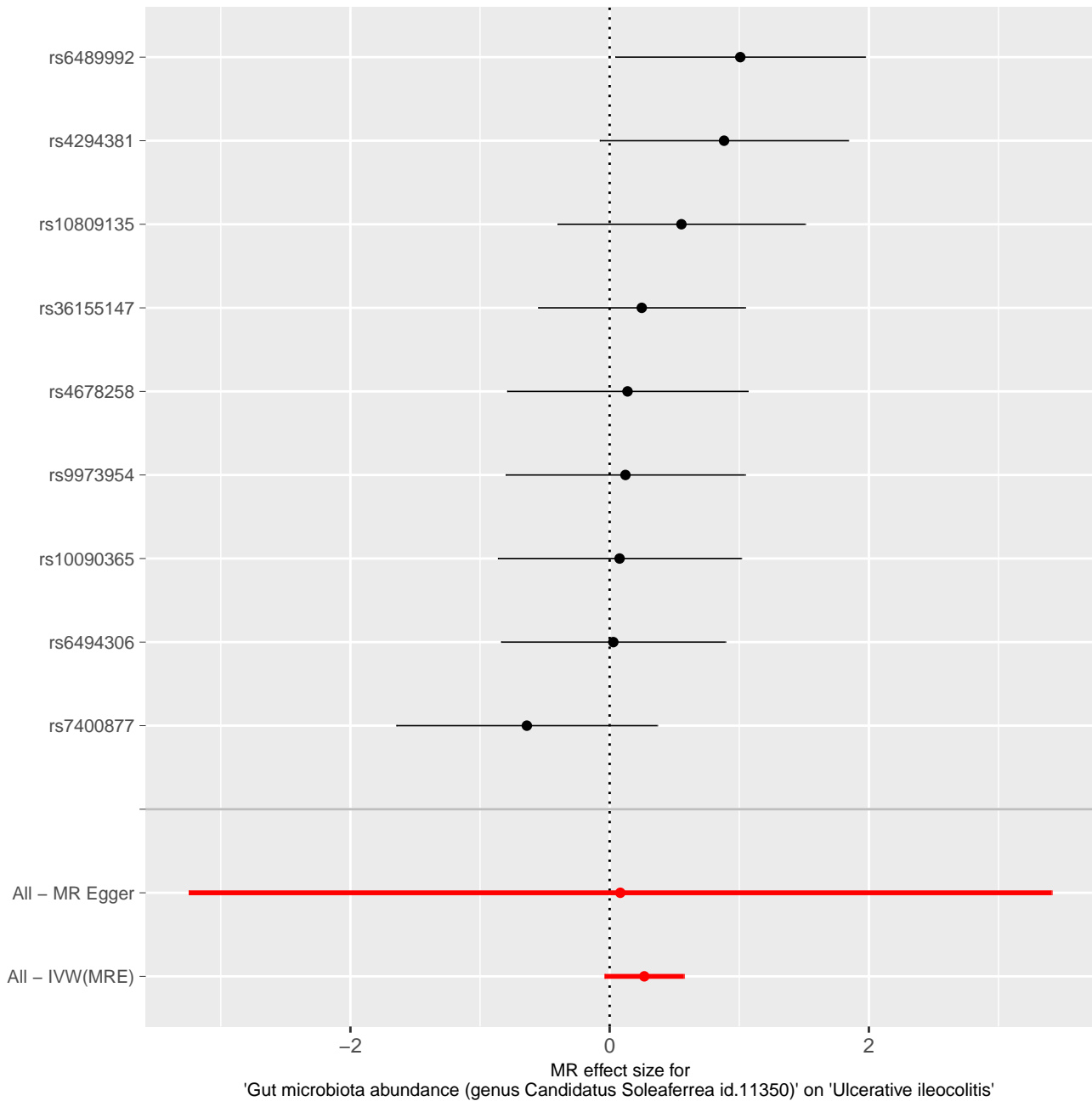


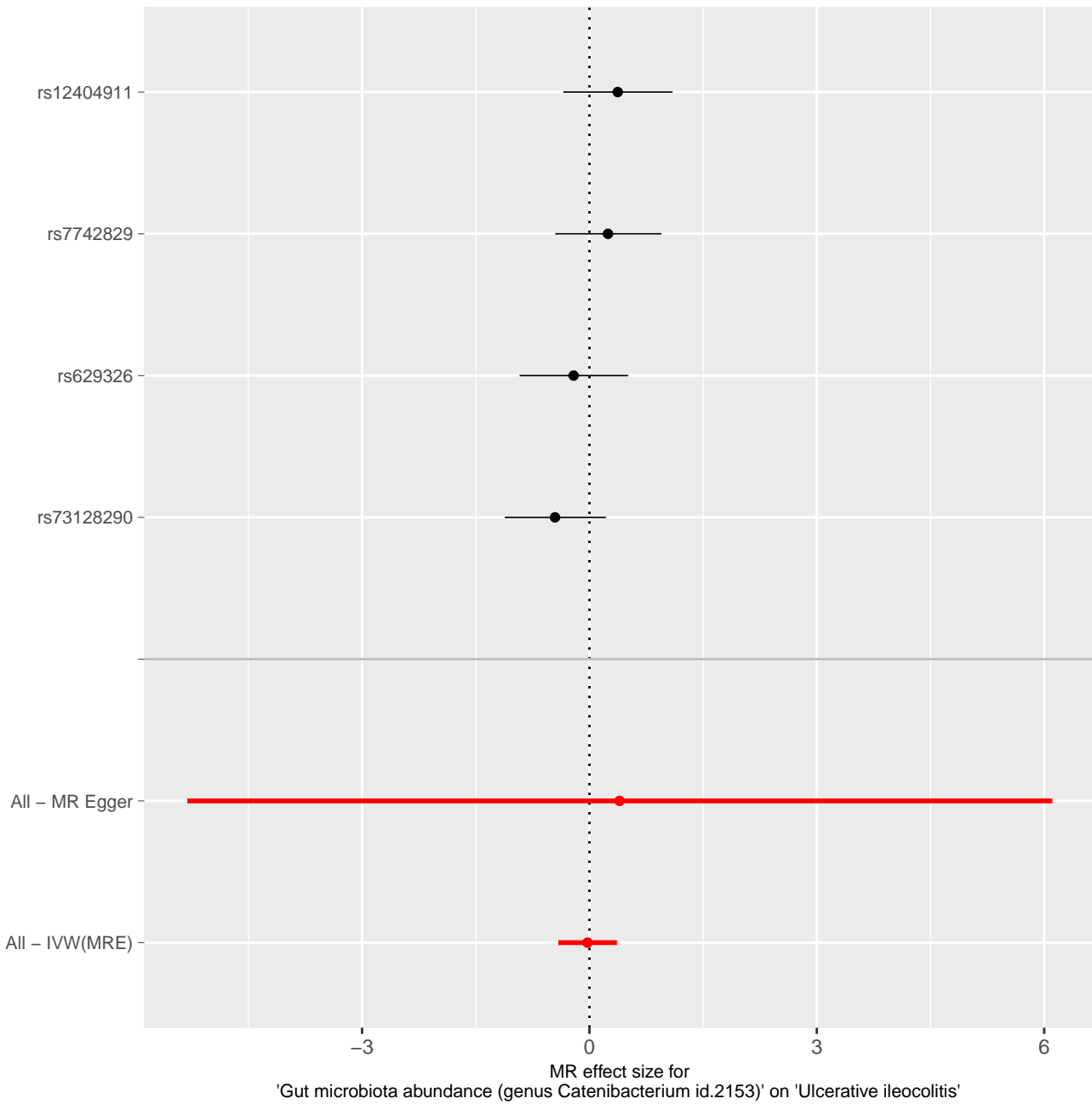


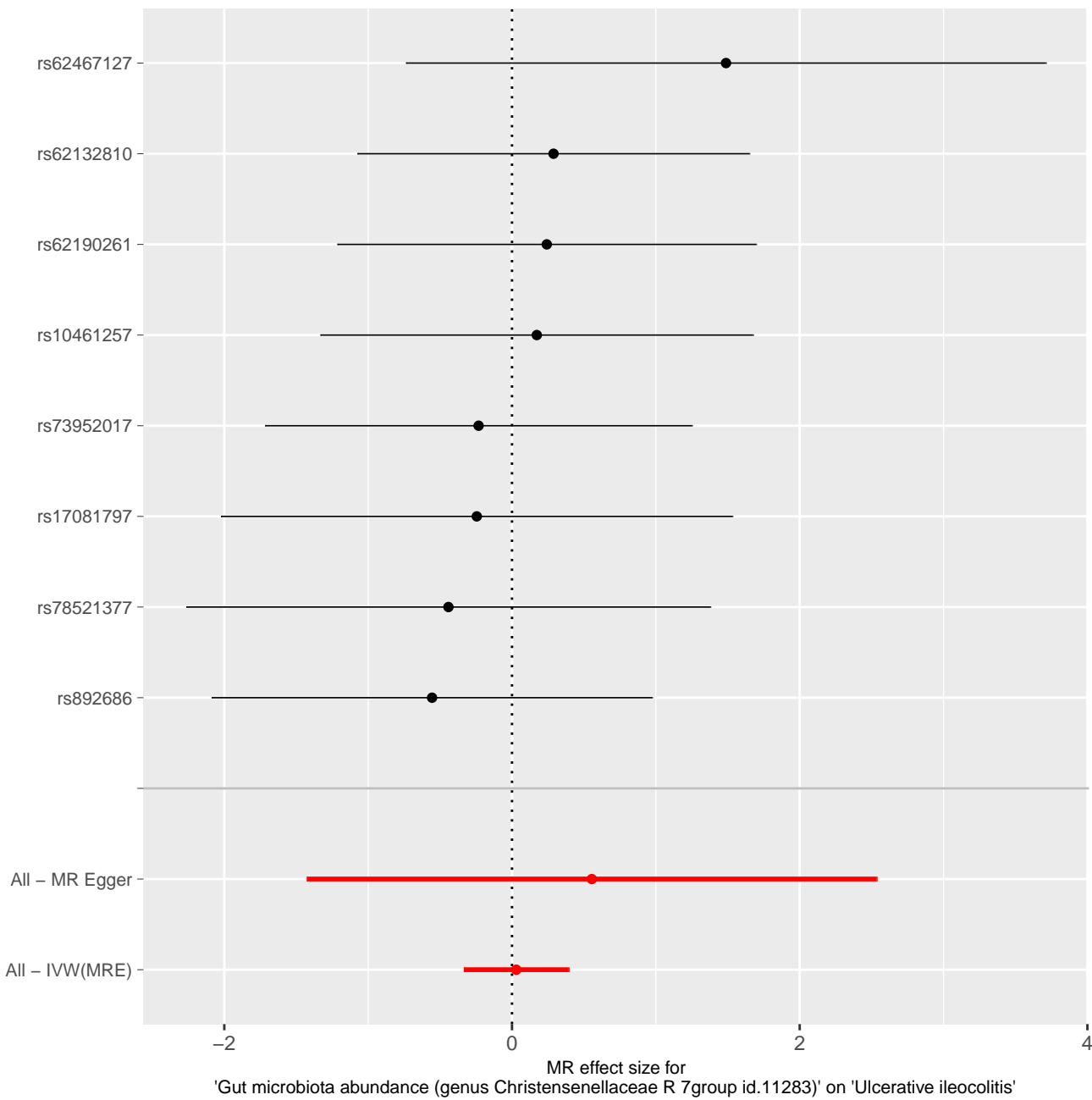


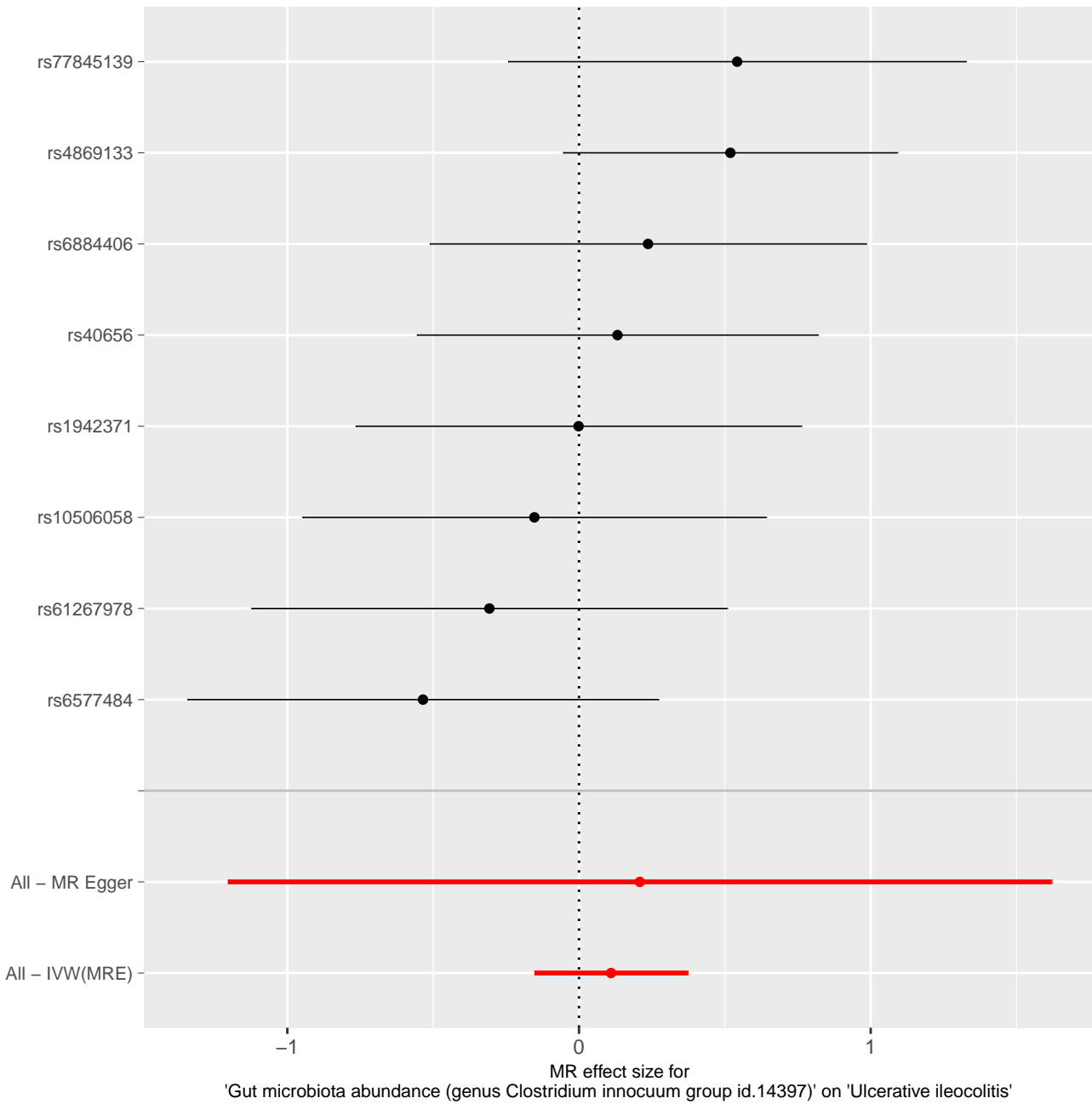
Batch 1099 : Gut microbiota abundance (genus Butyrivibrio id.1993) on Ulcerative ileocolitis

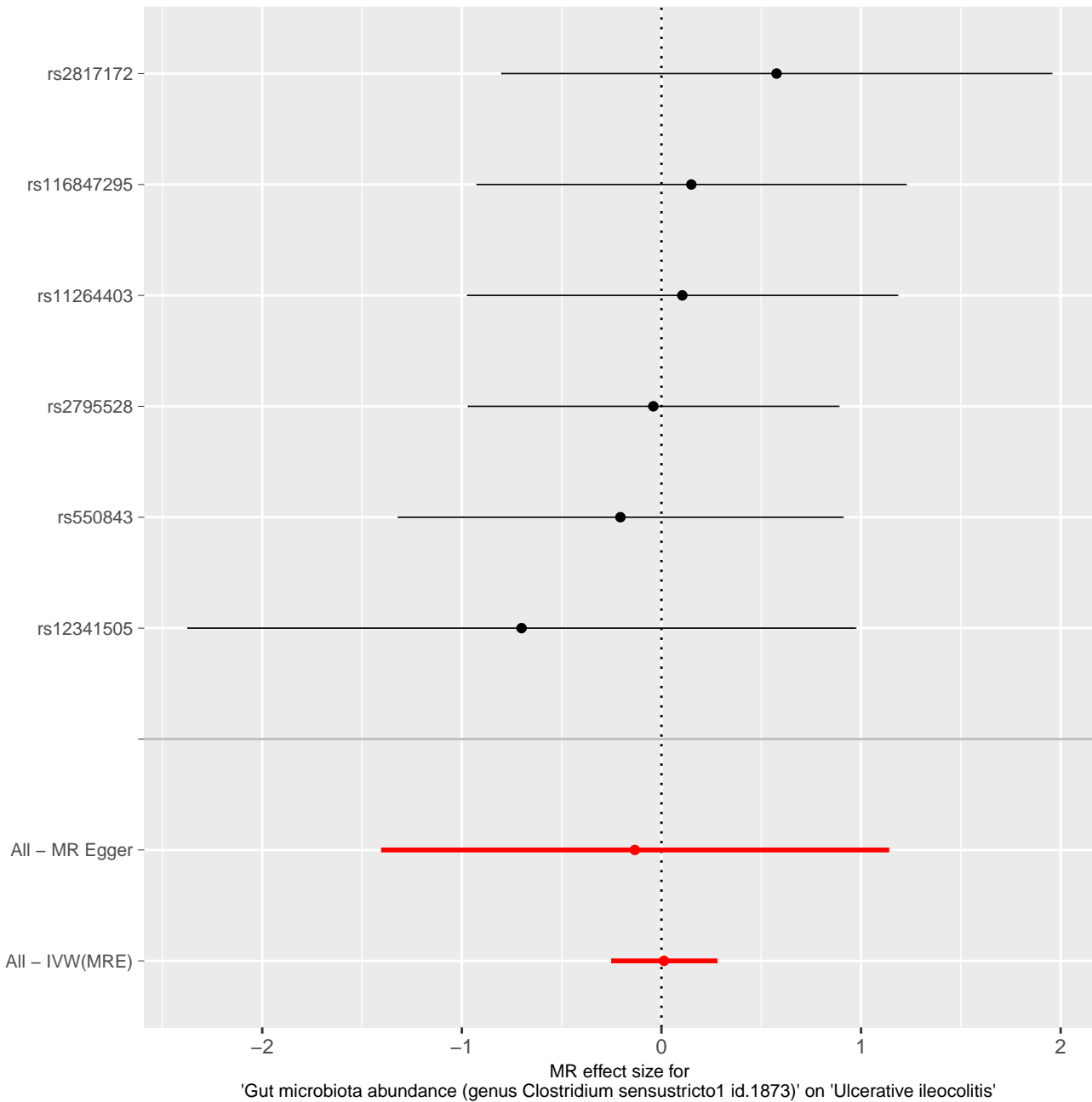


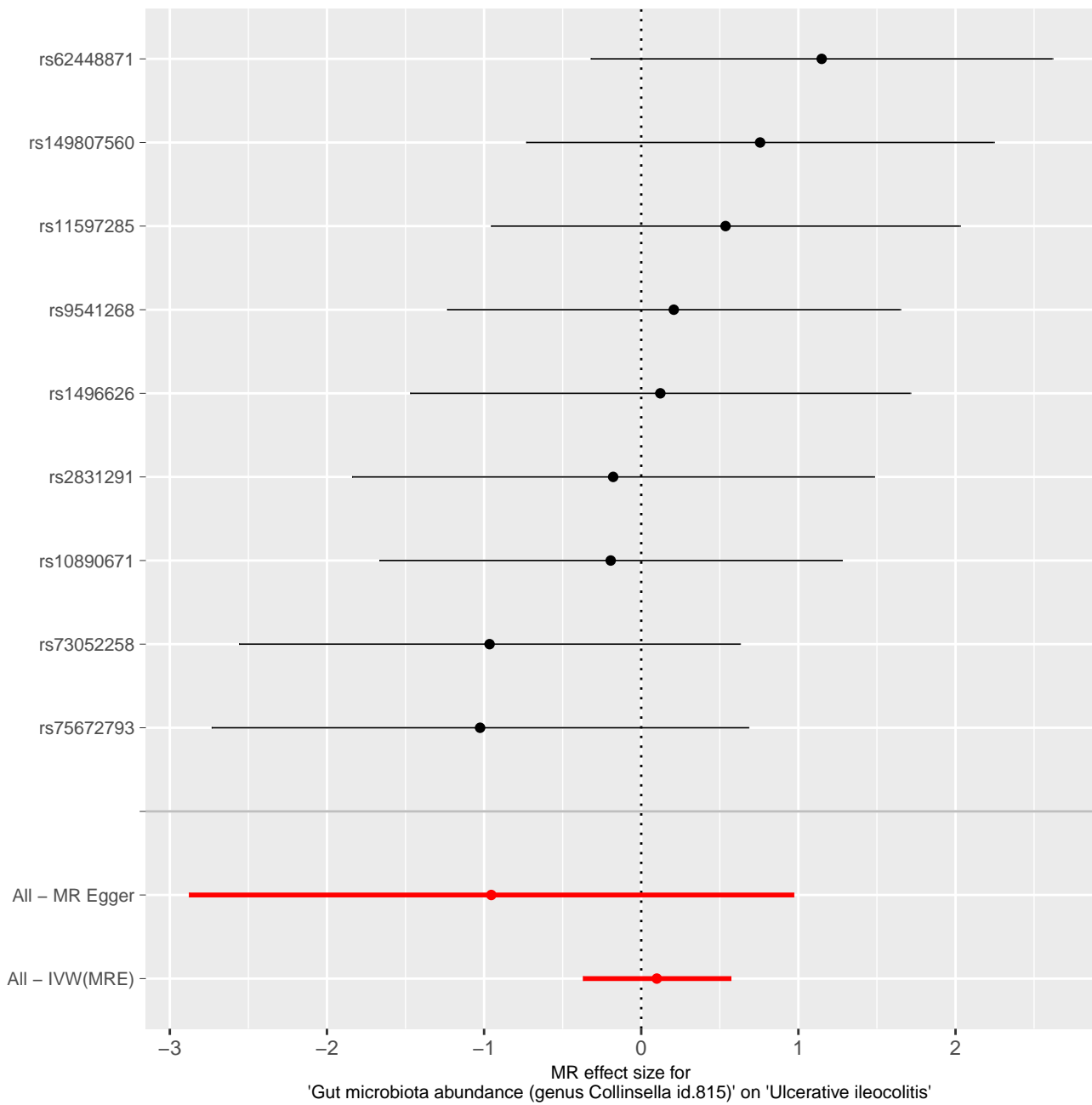


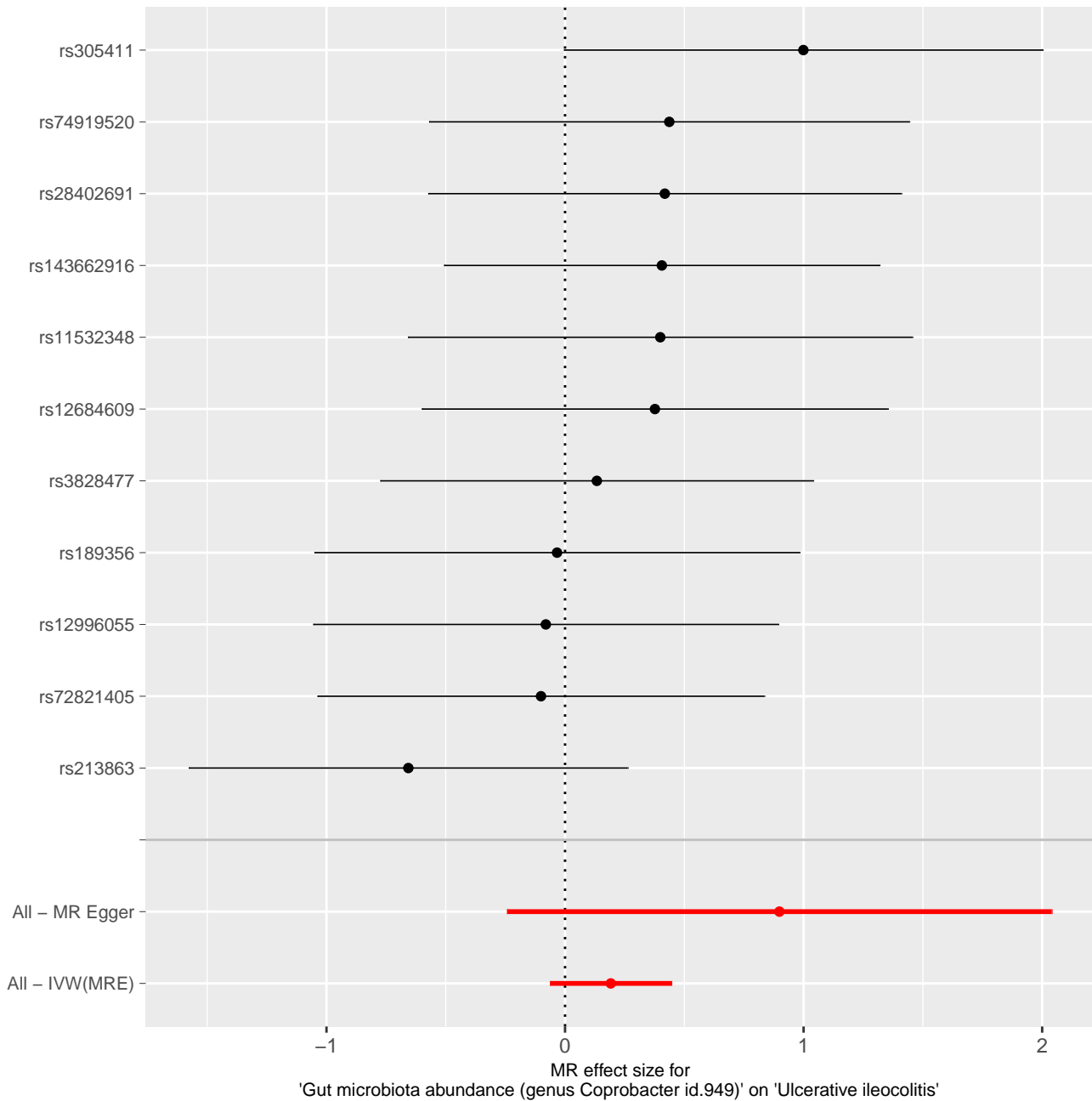


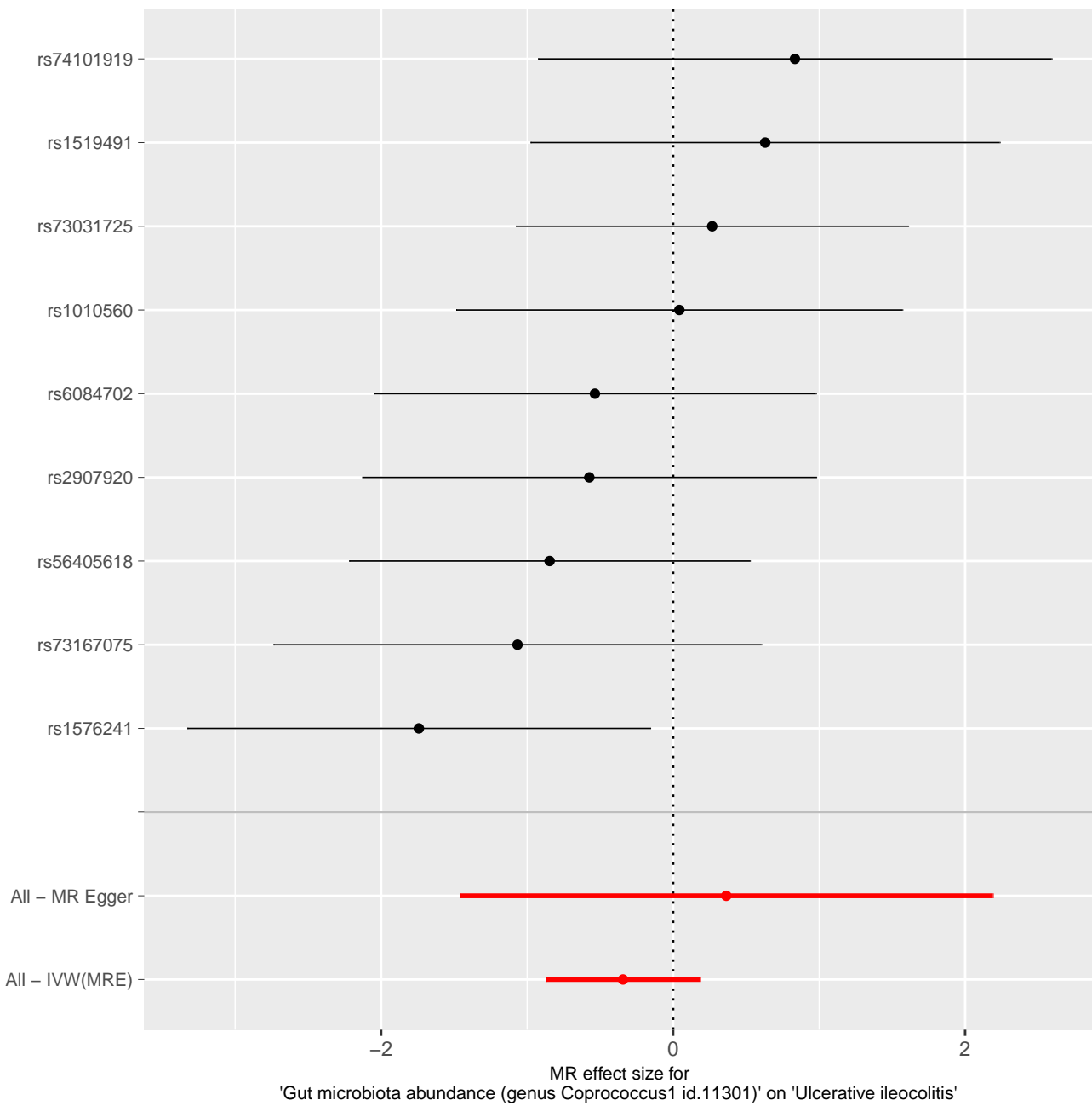


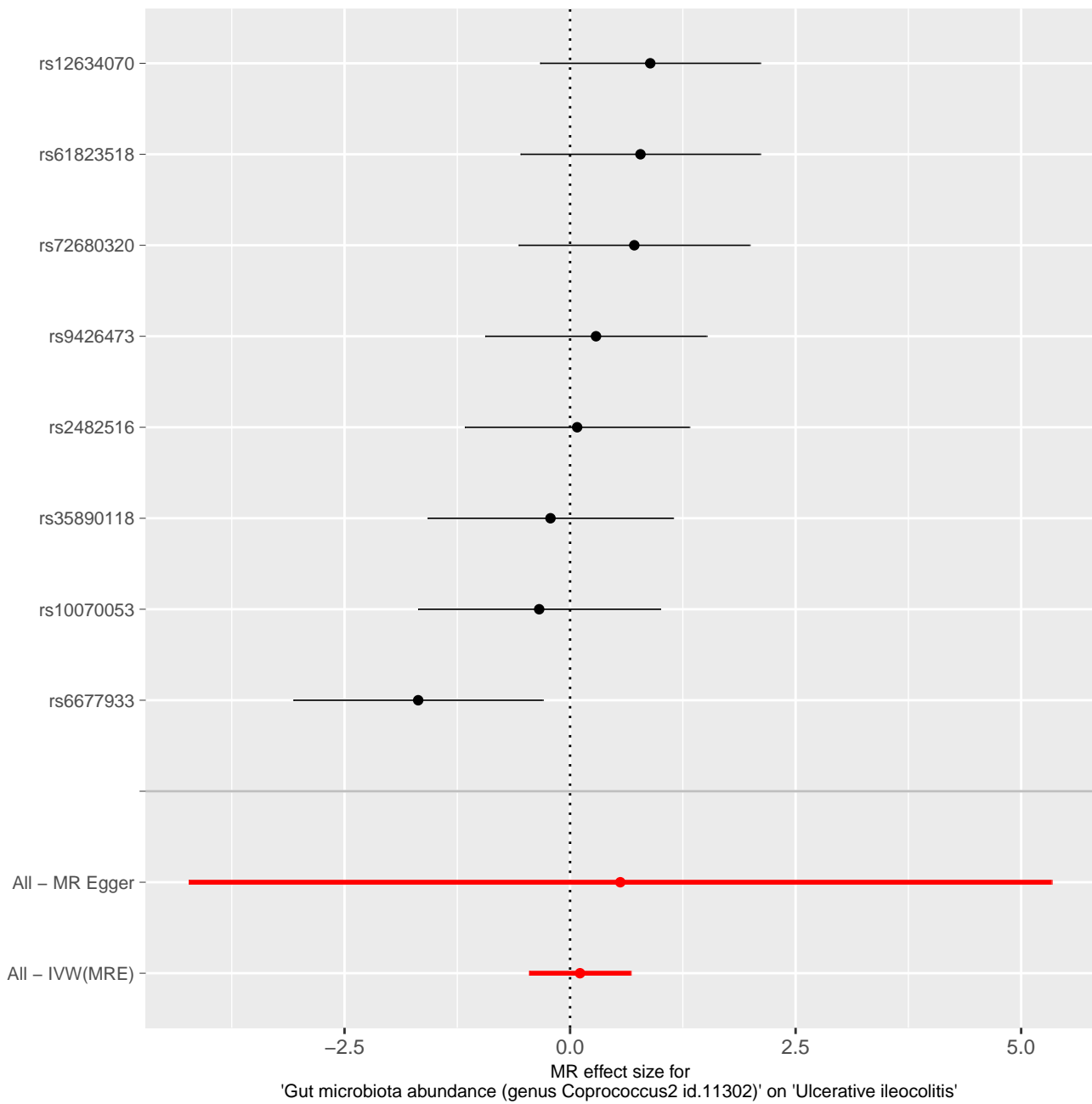


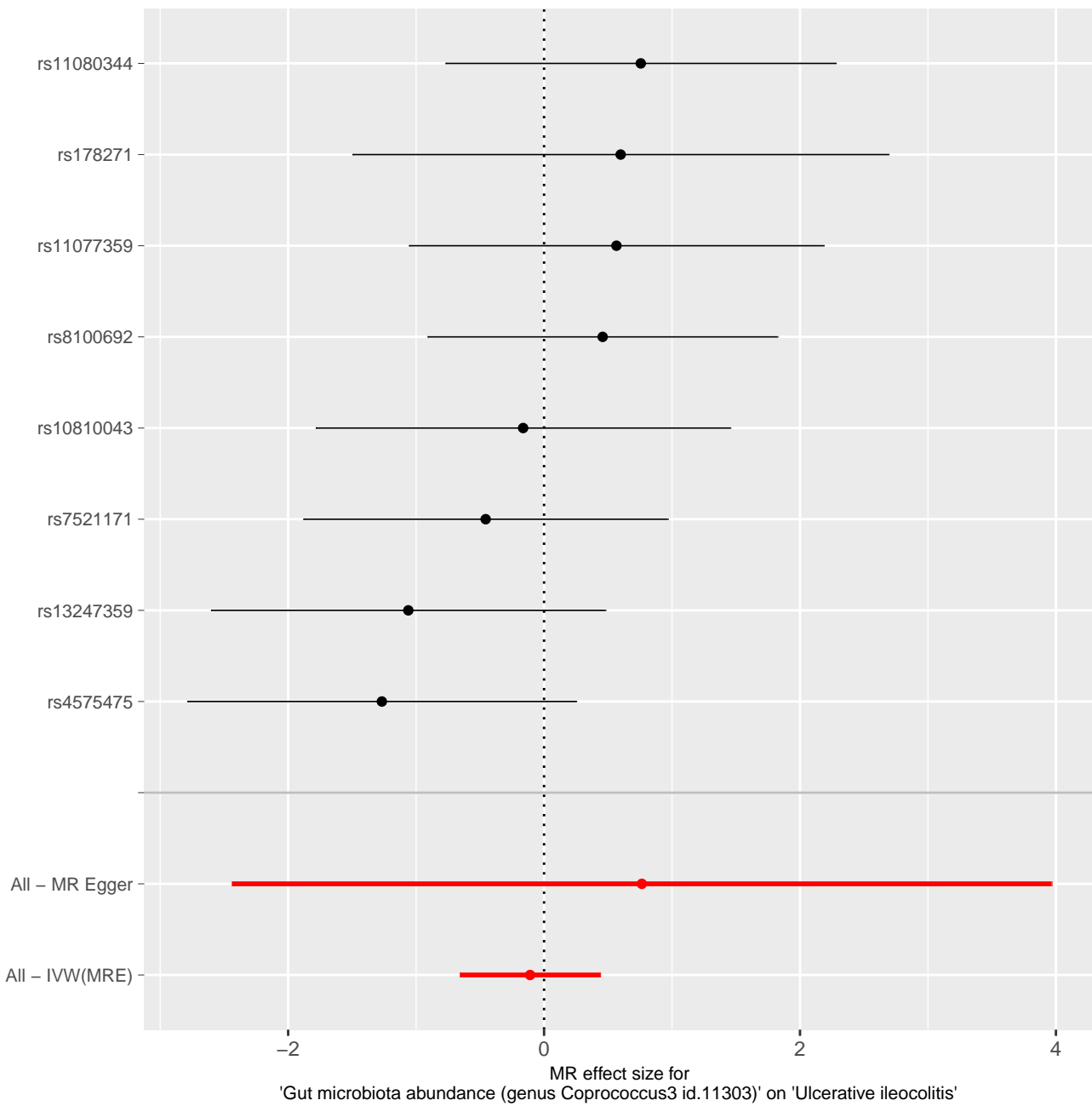


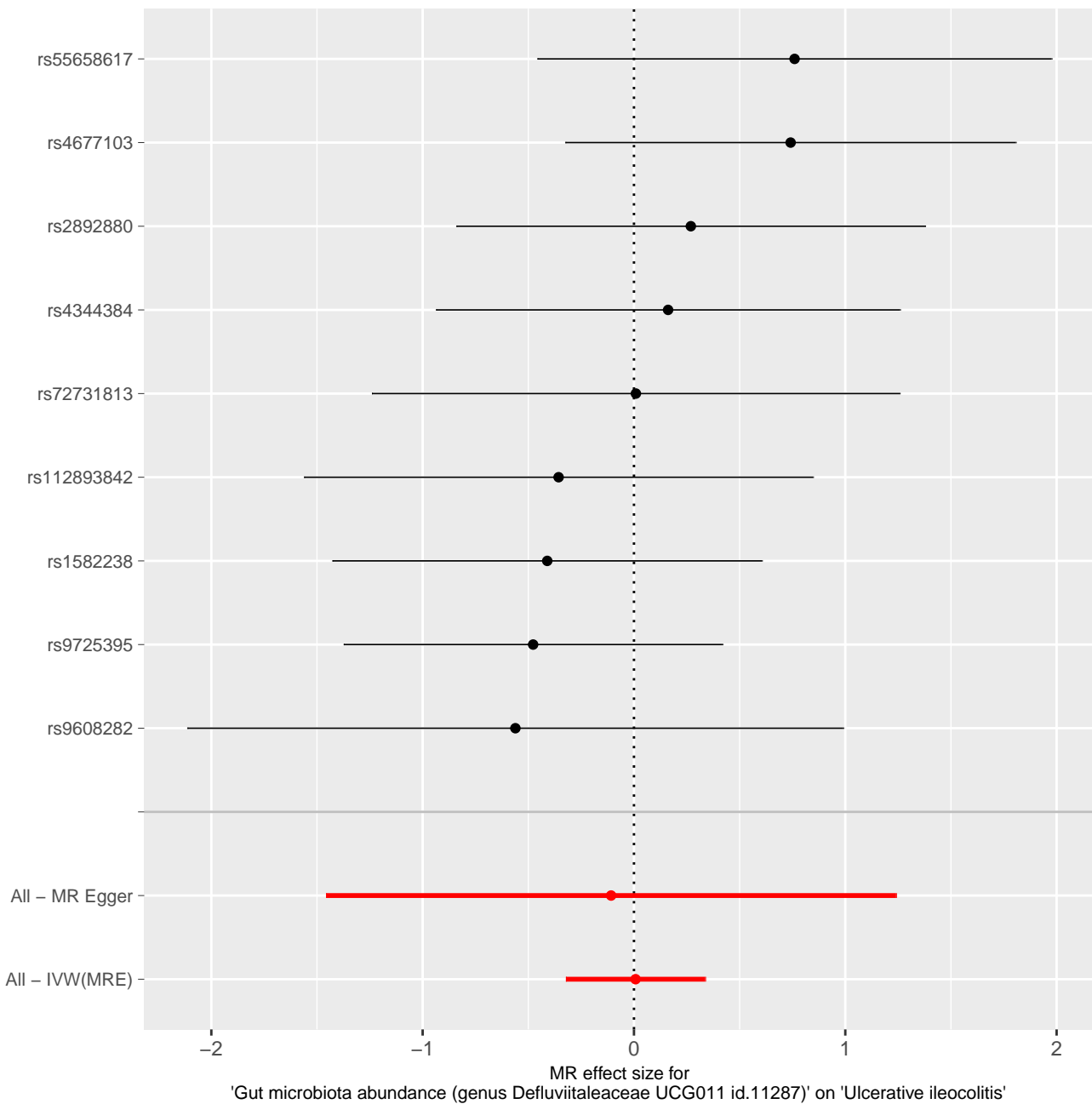


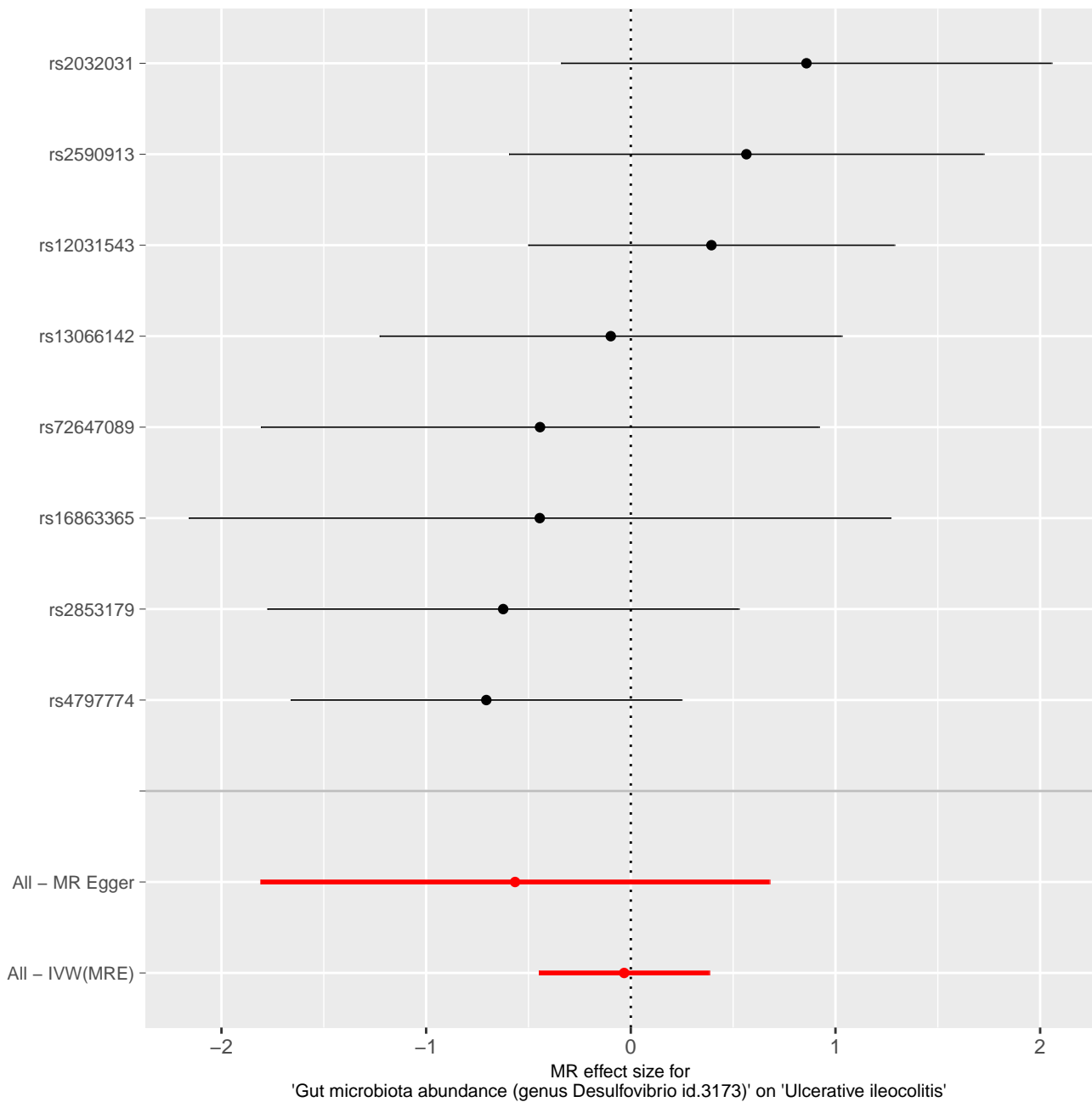


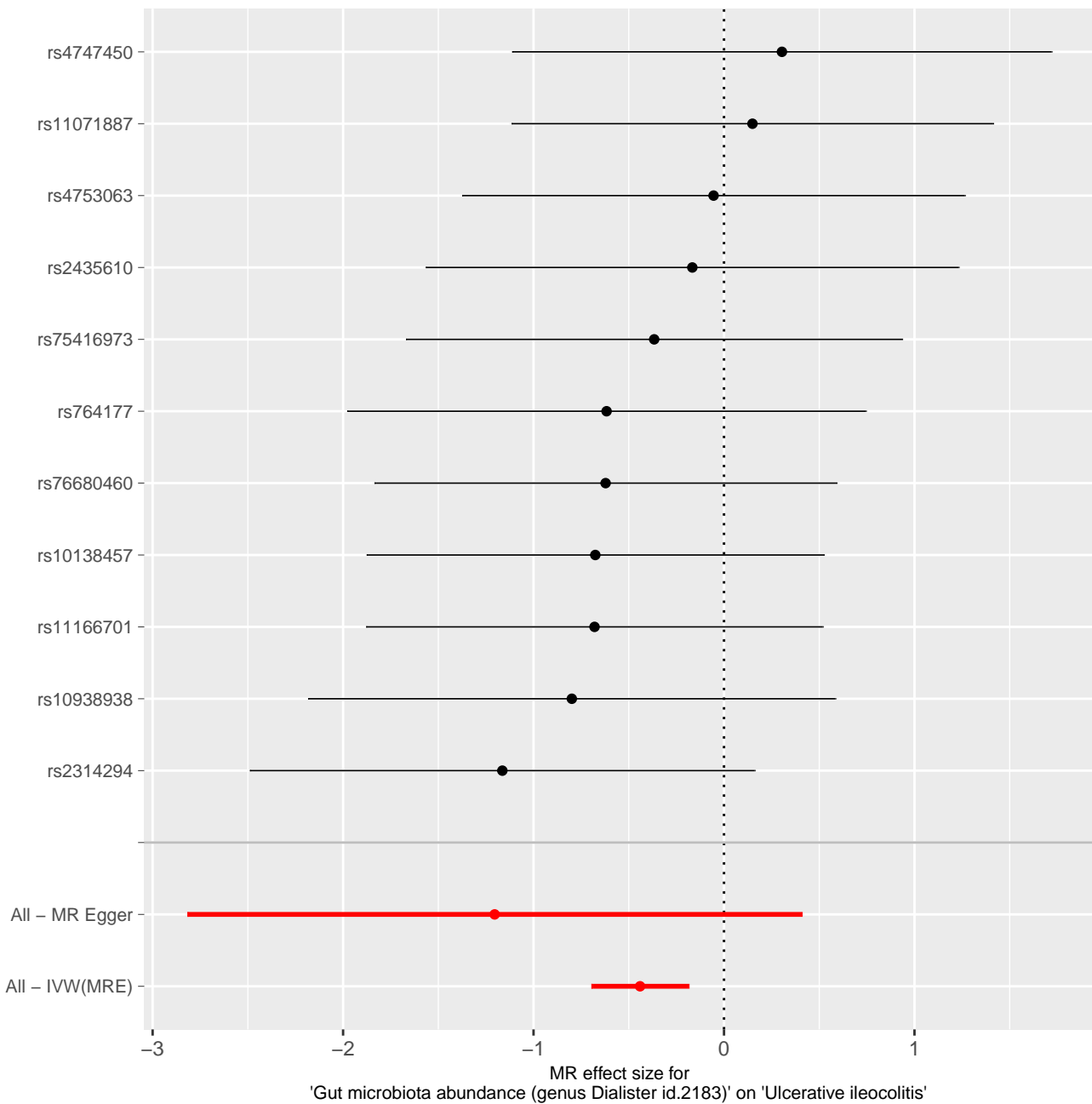


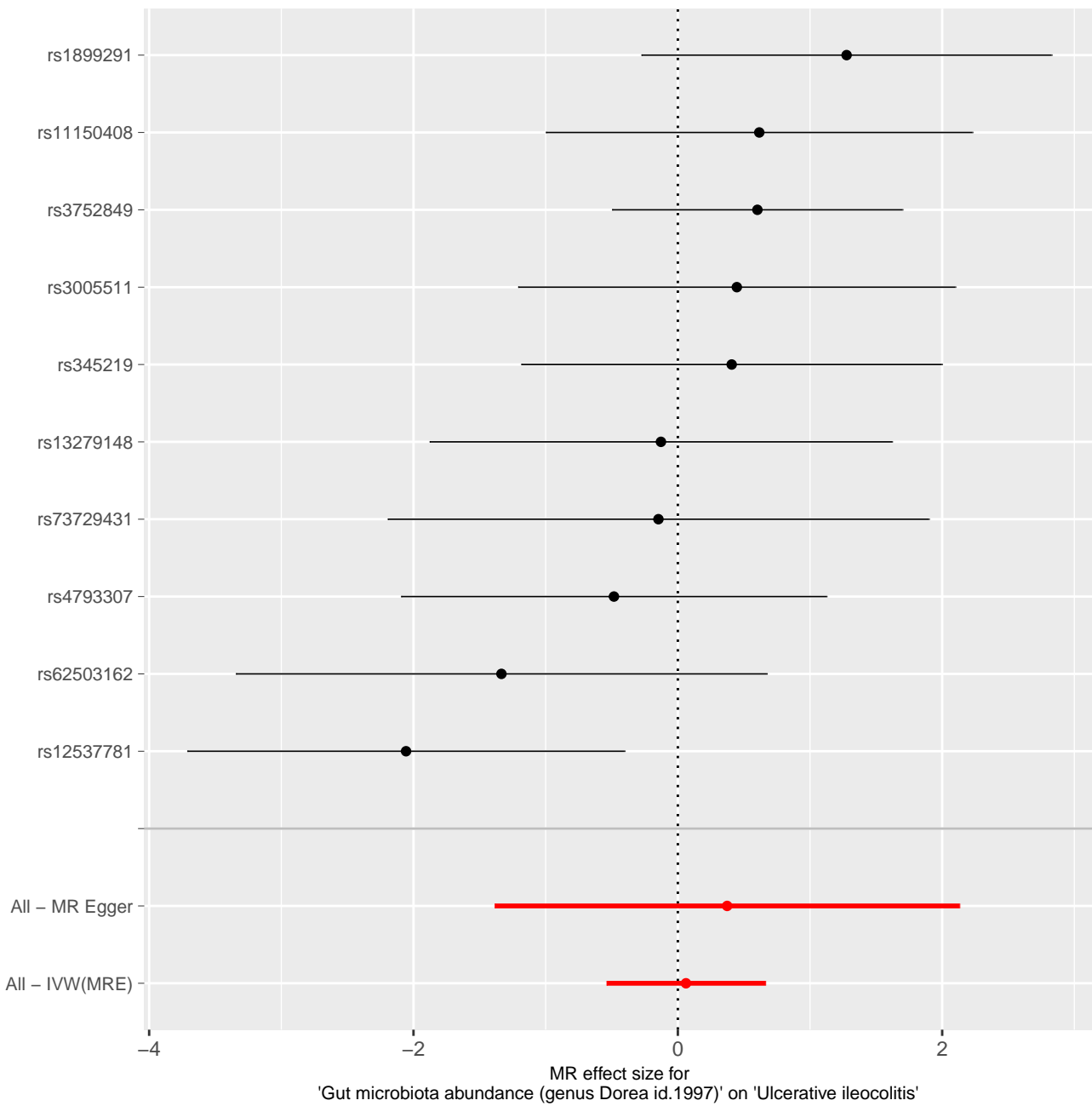


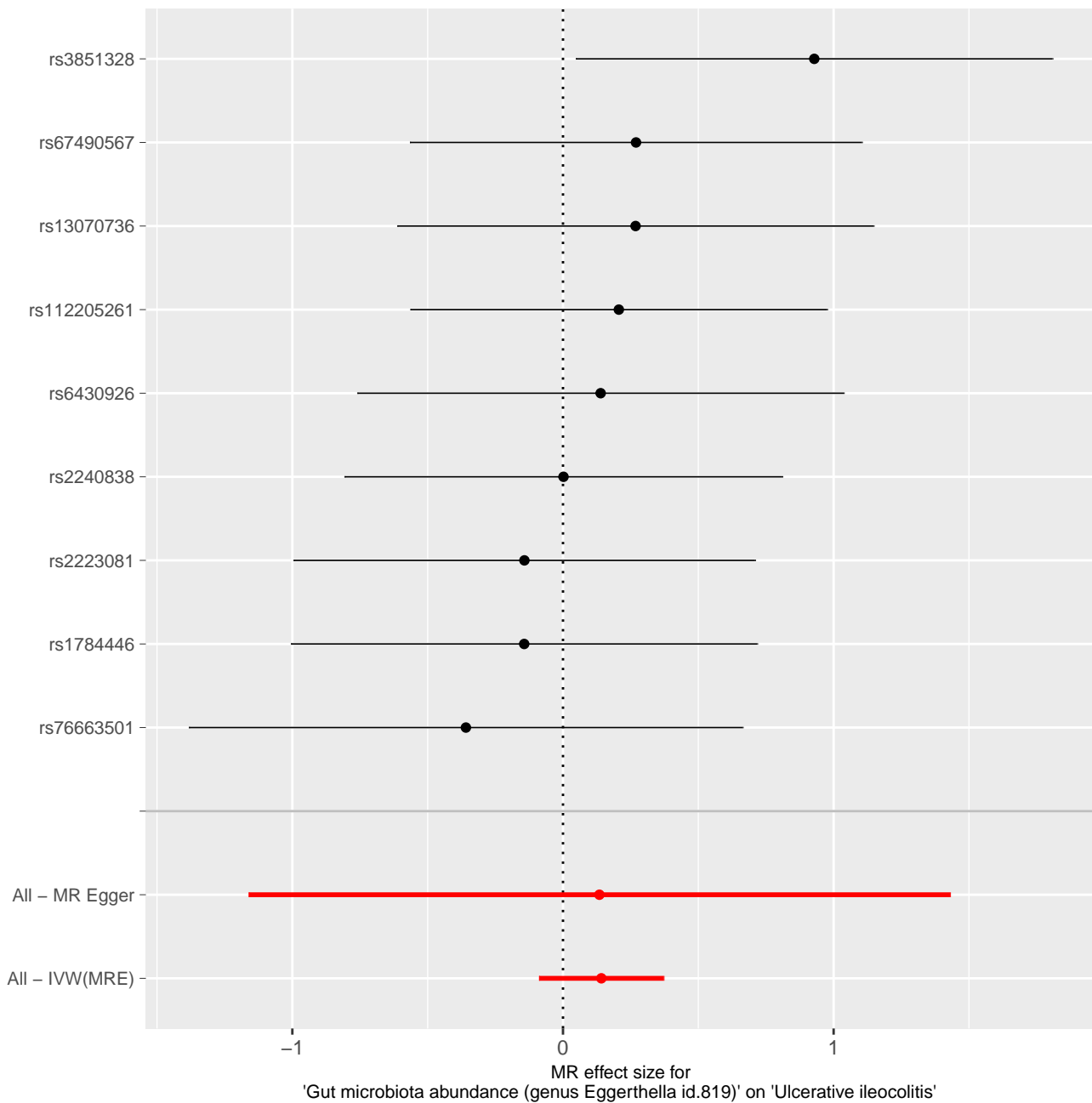


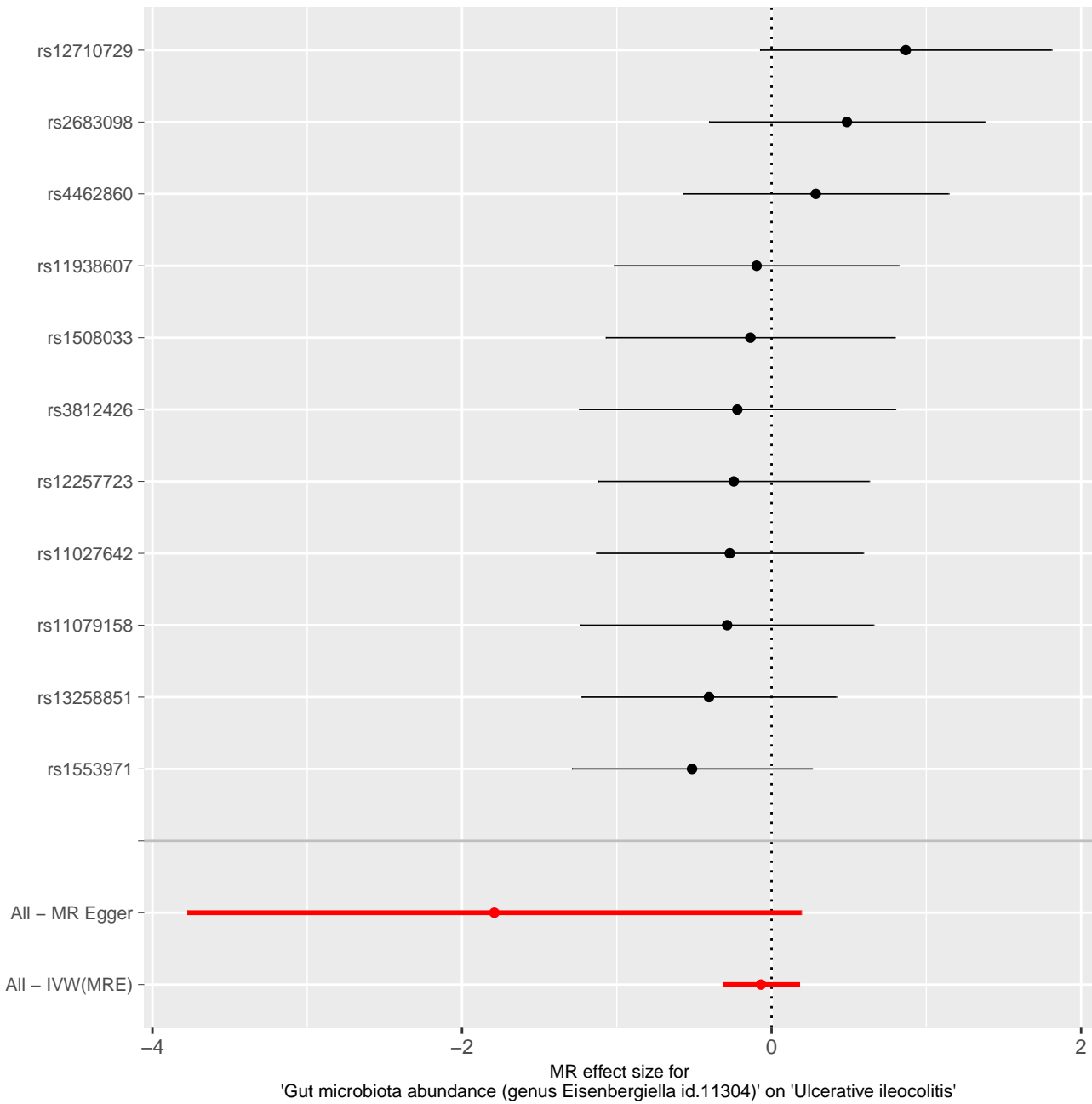


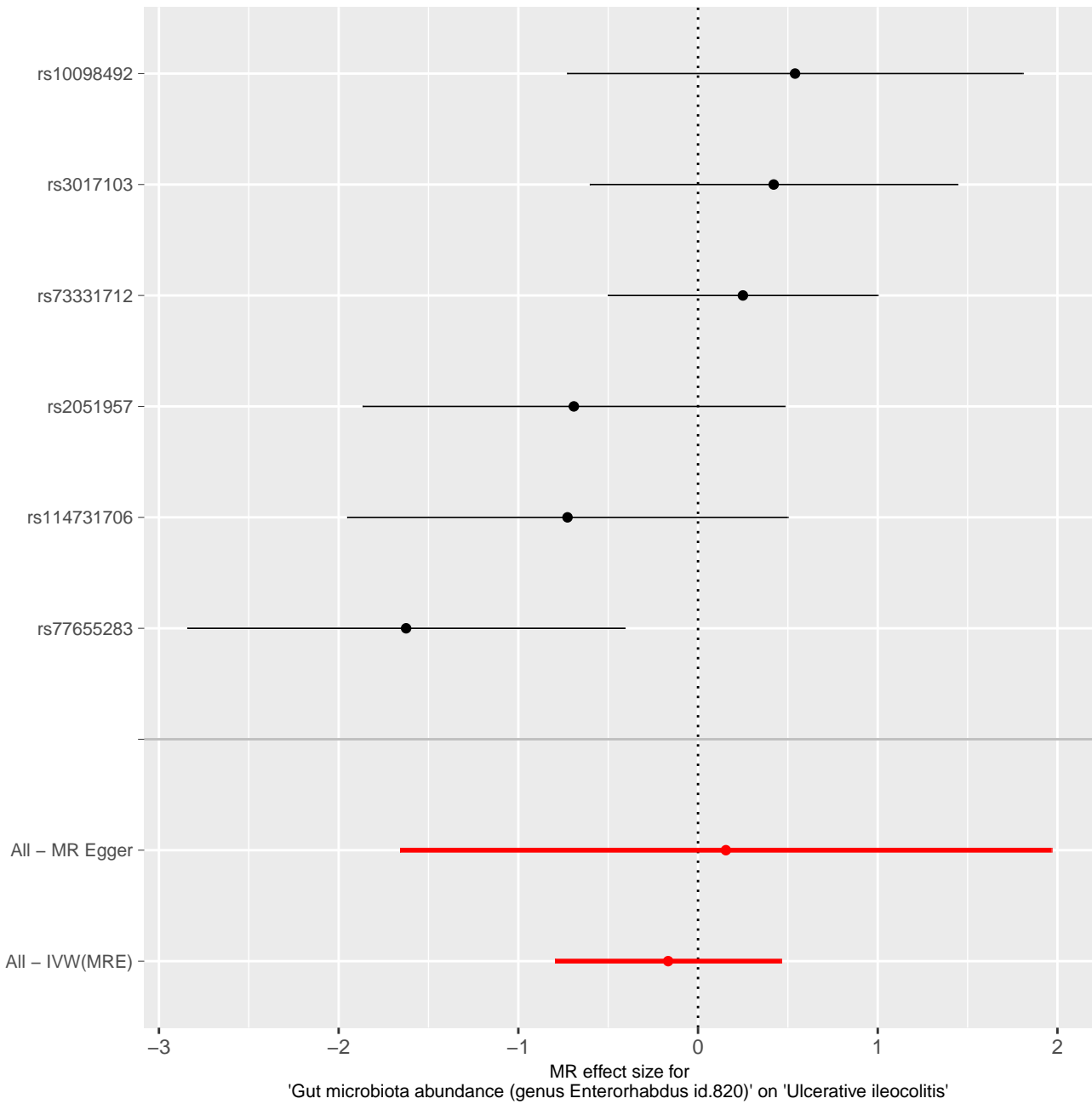


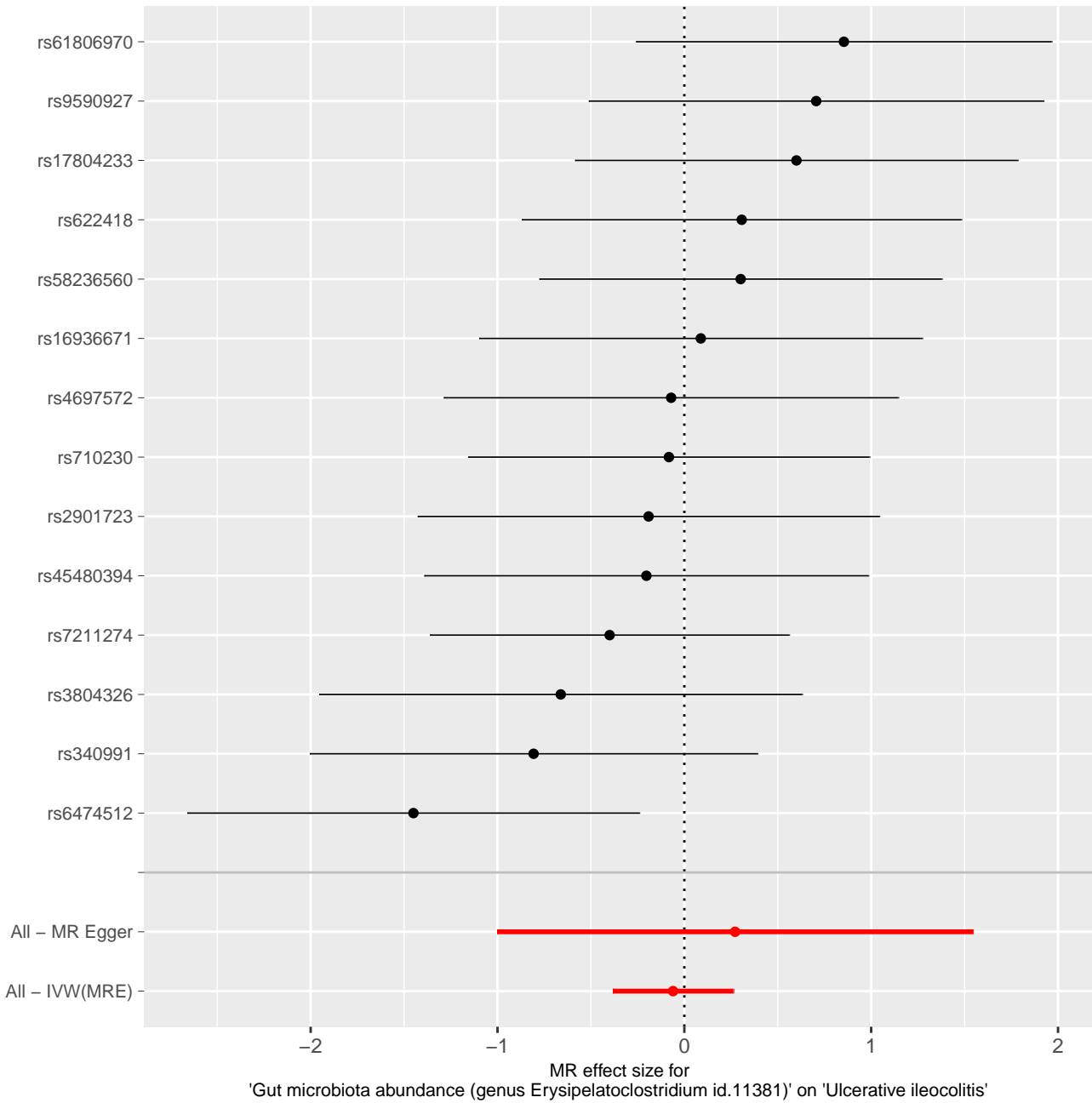


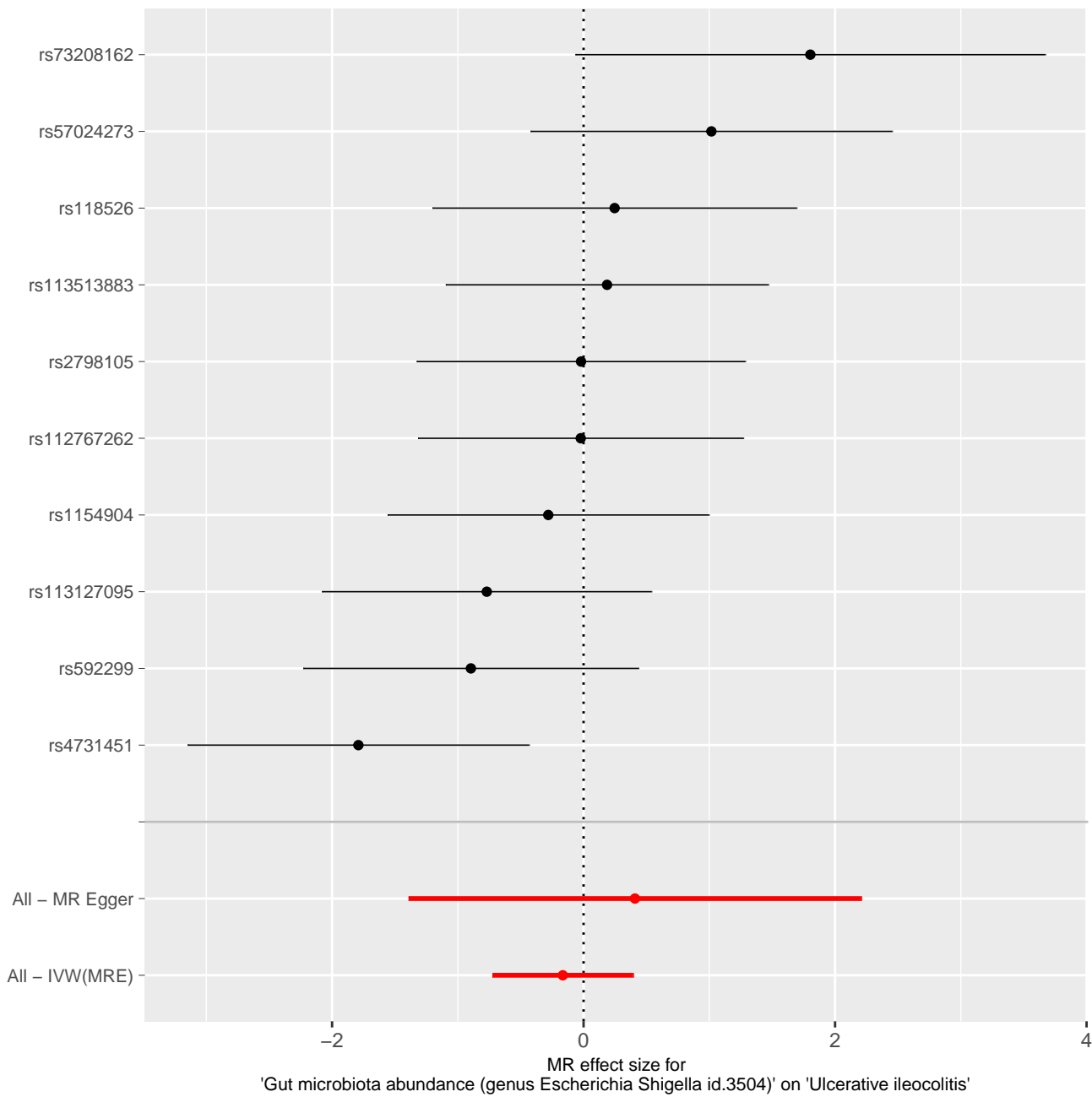


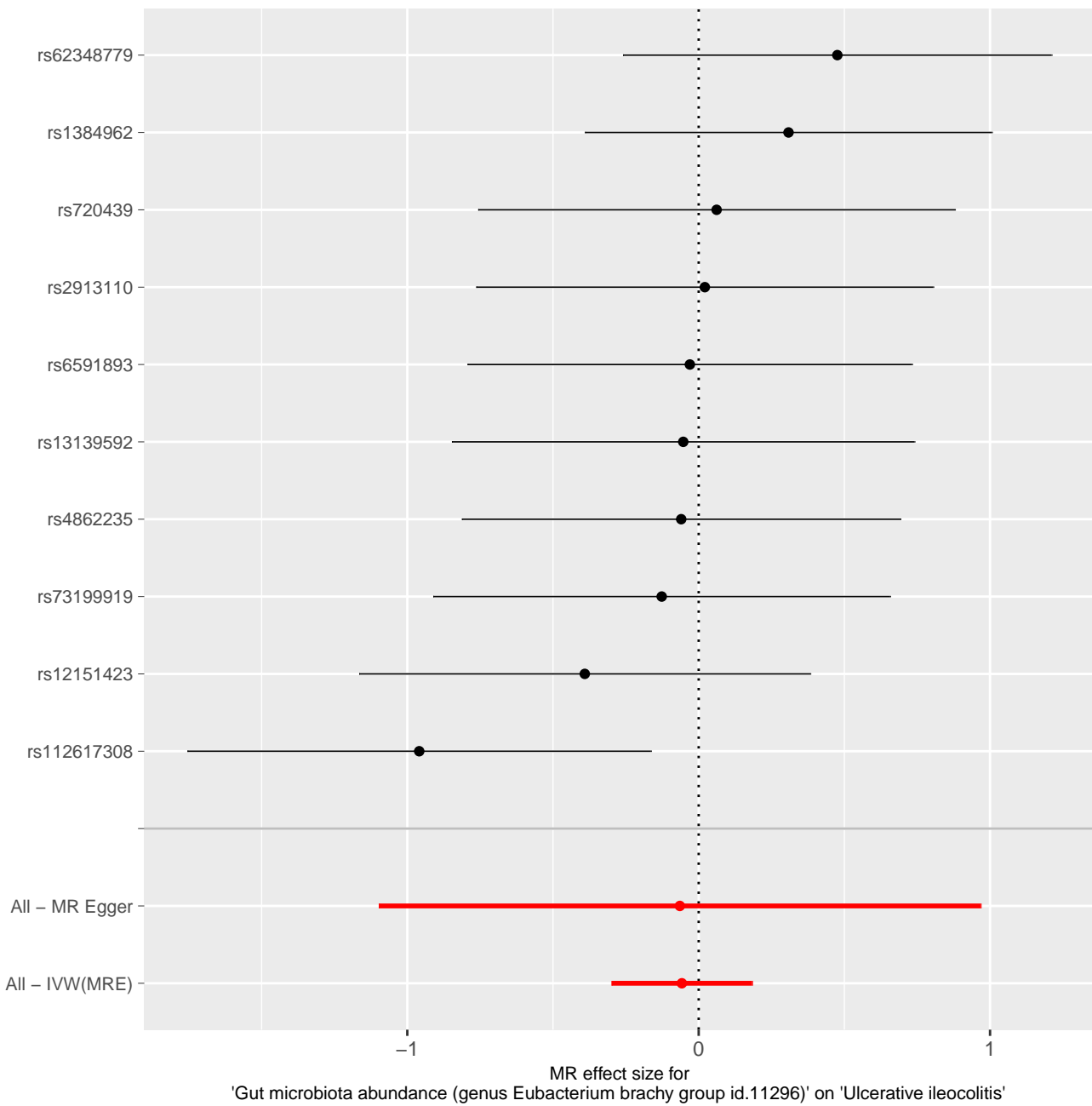


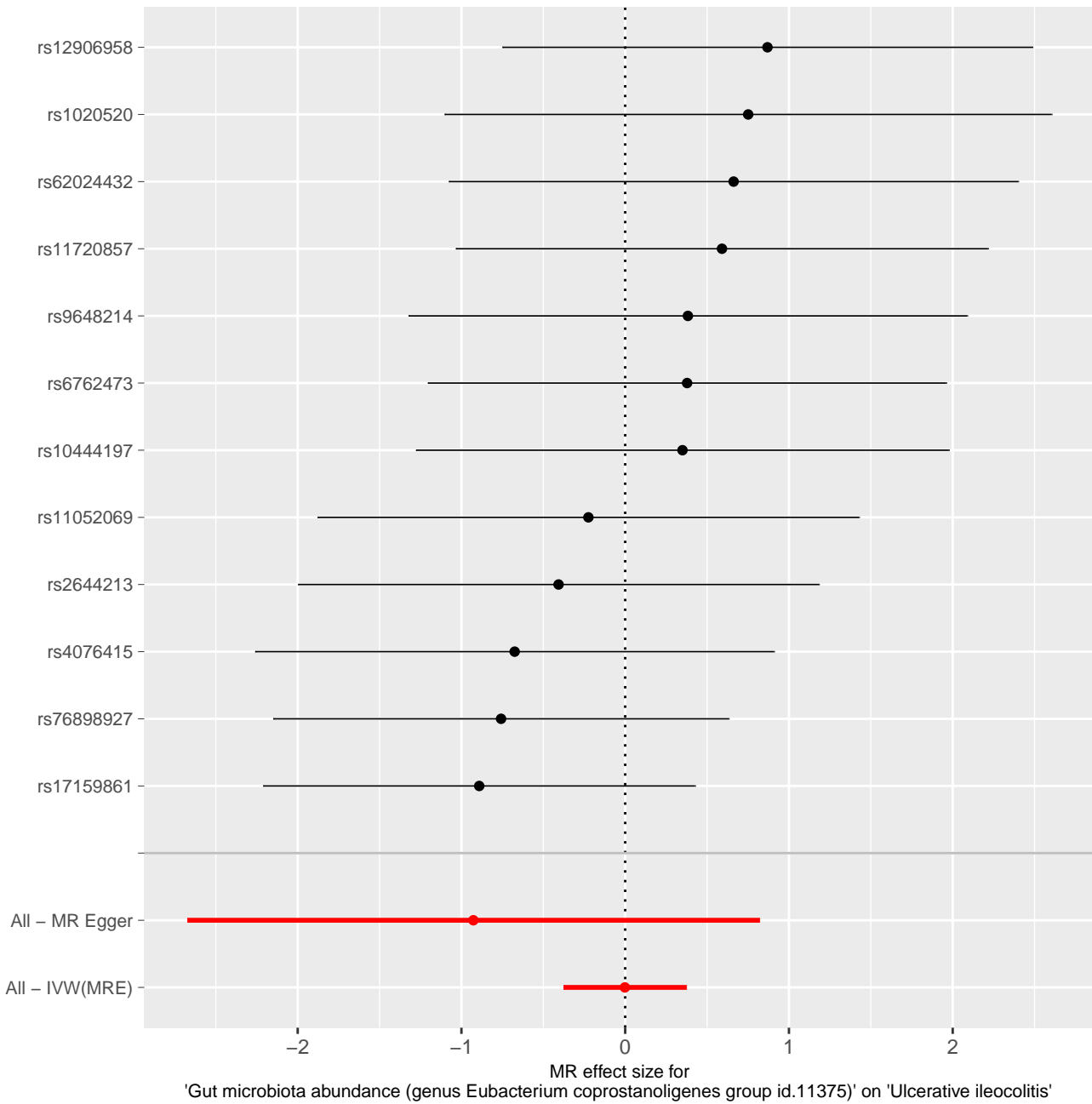


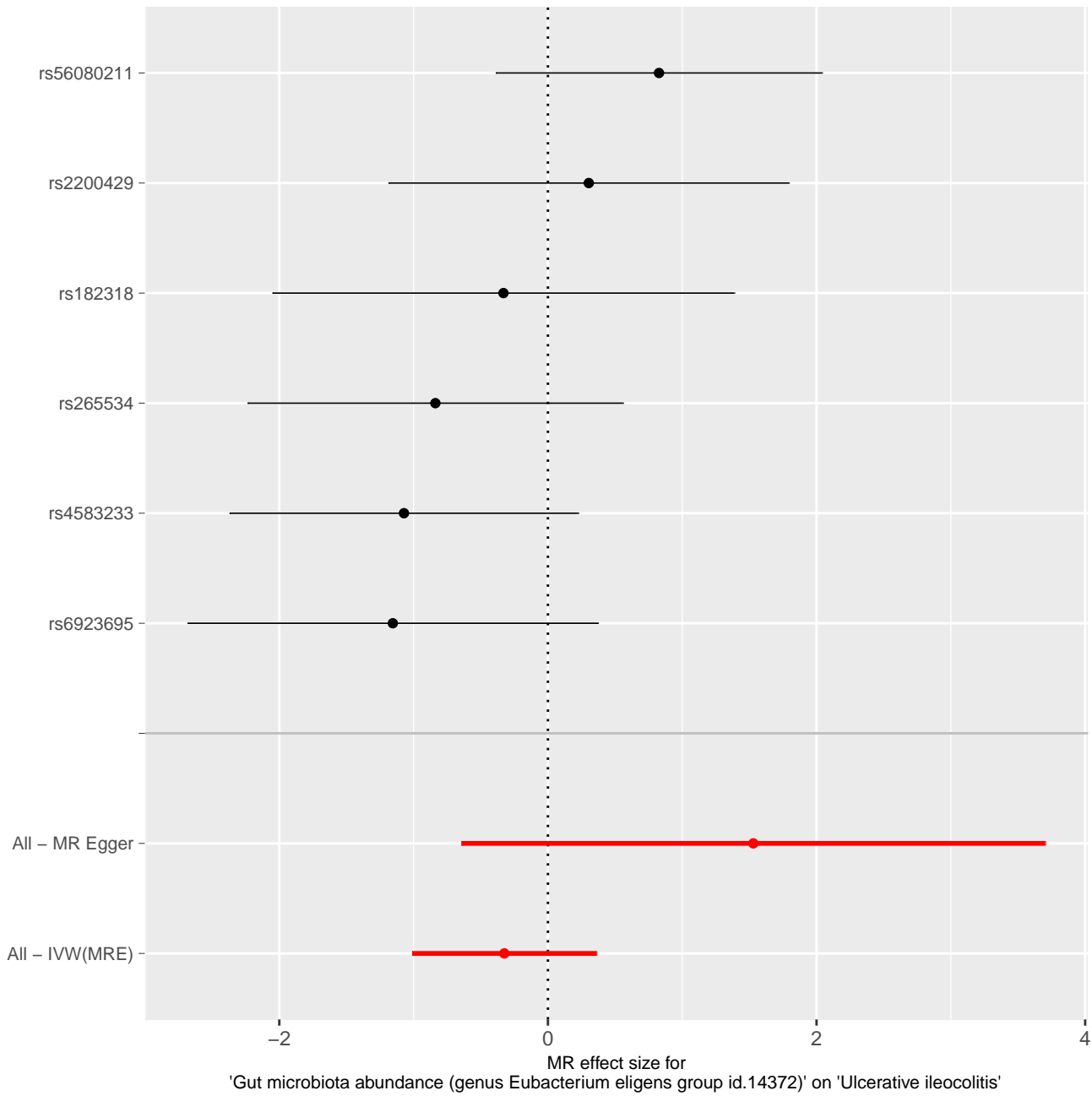


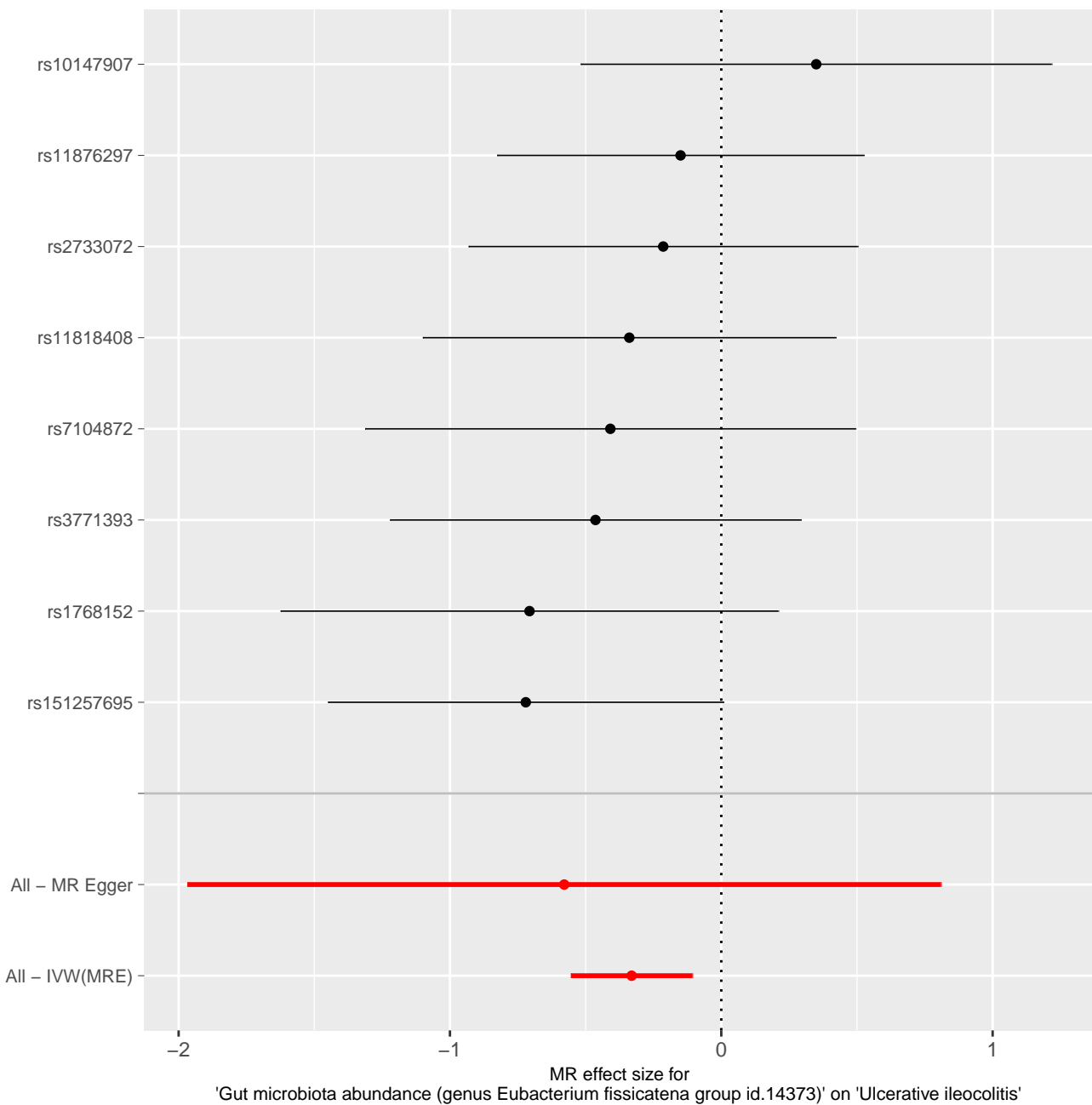


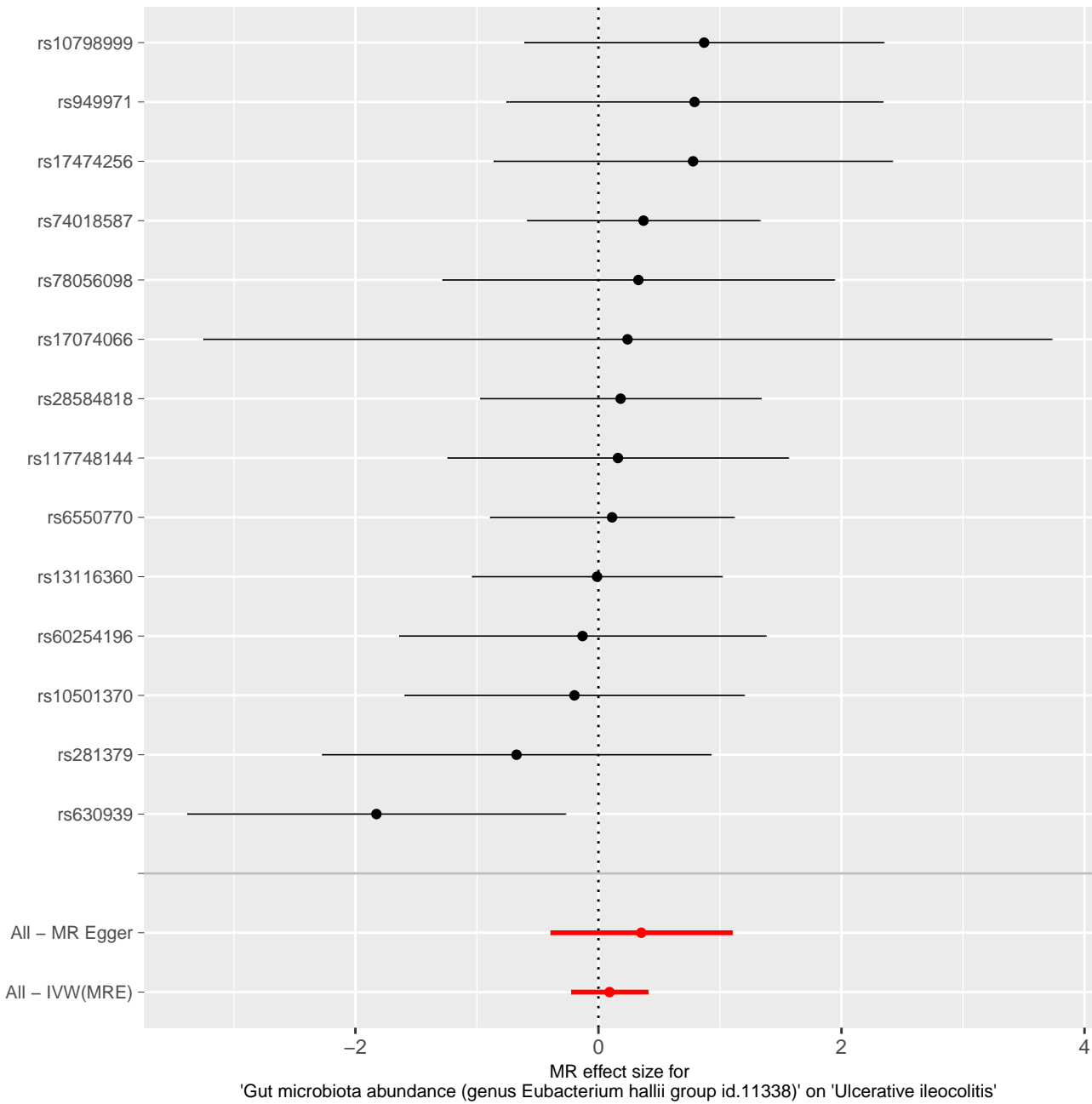


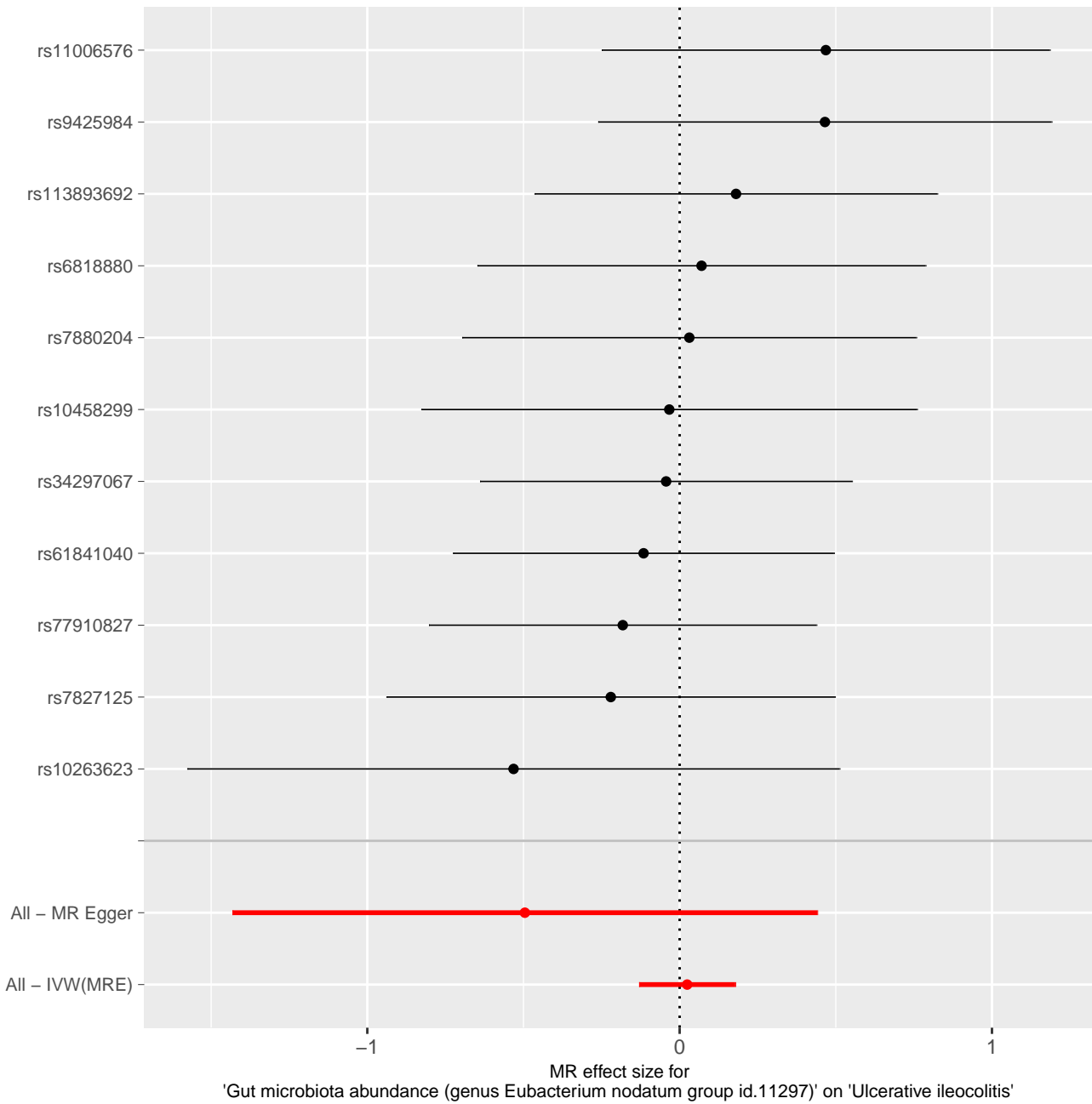


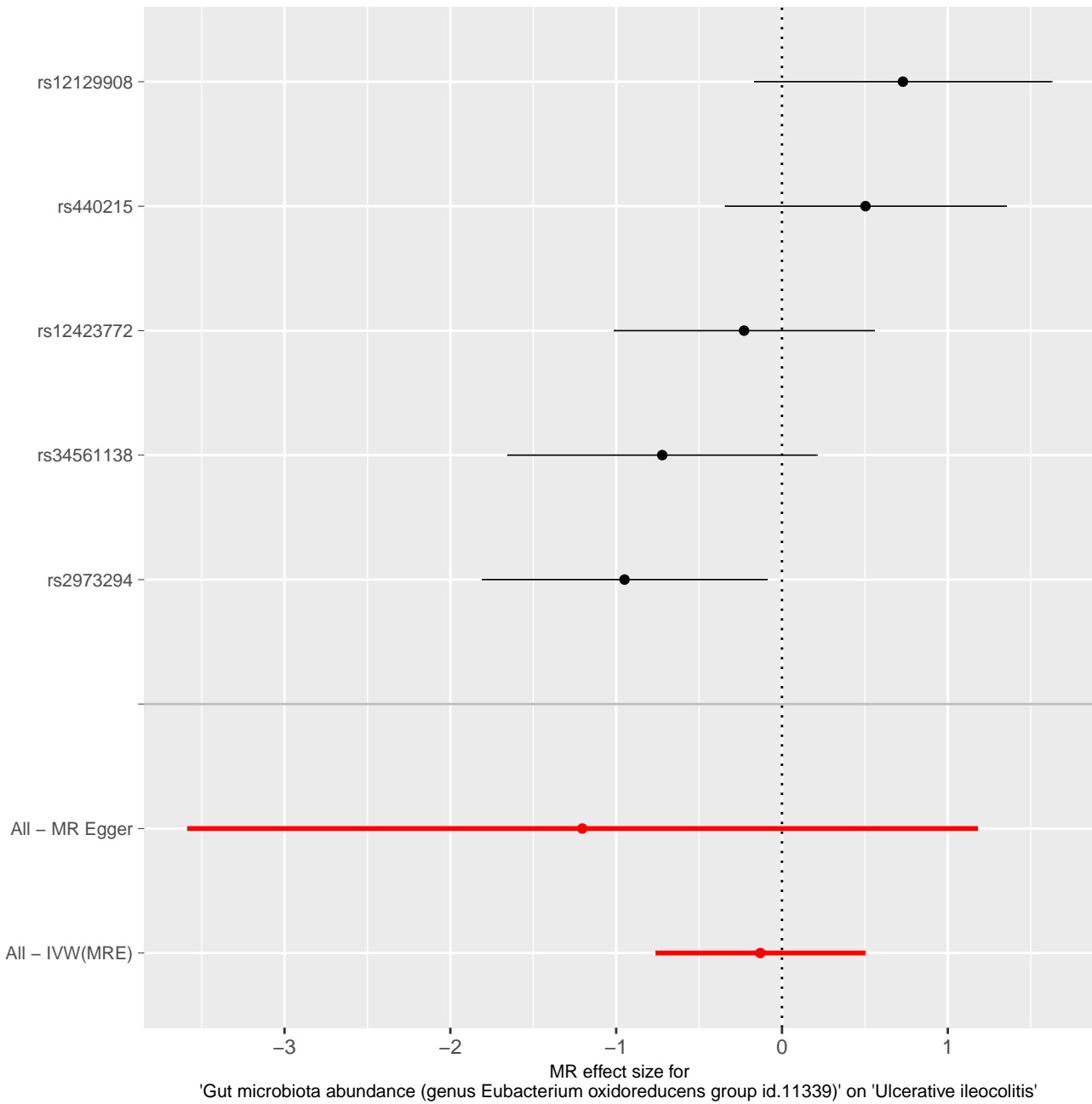


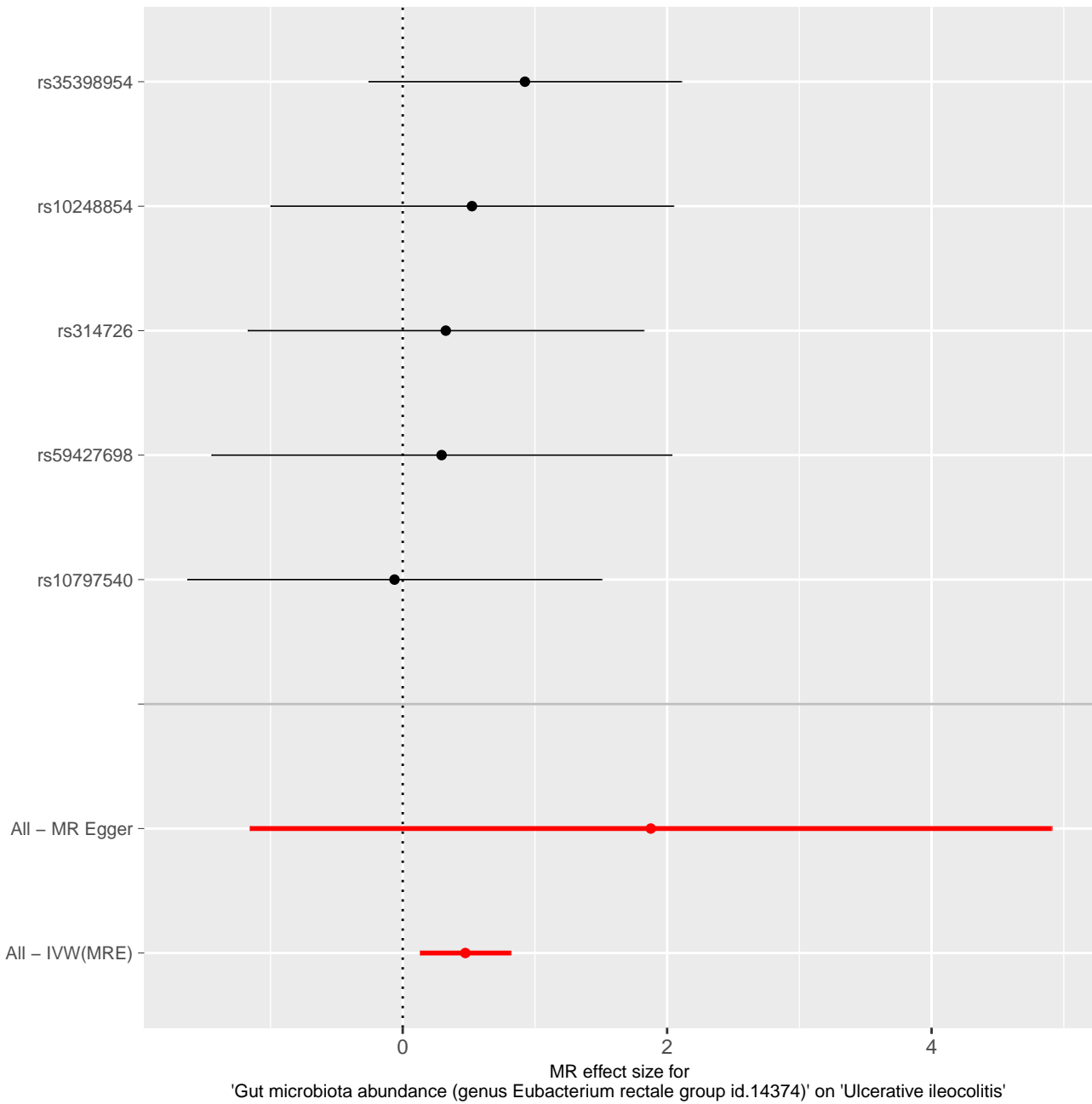


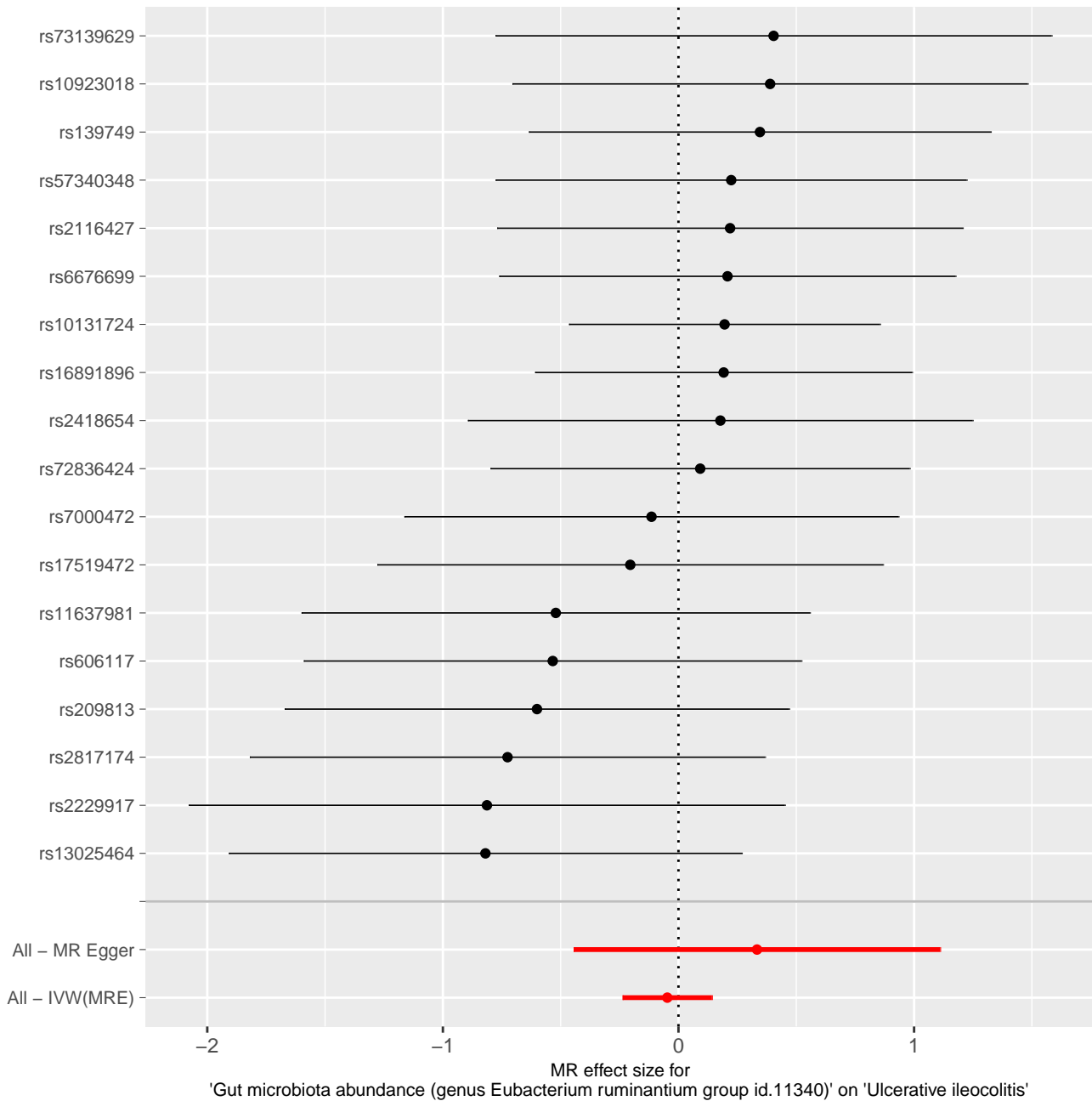


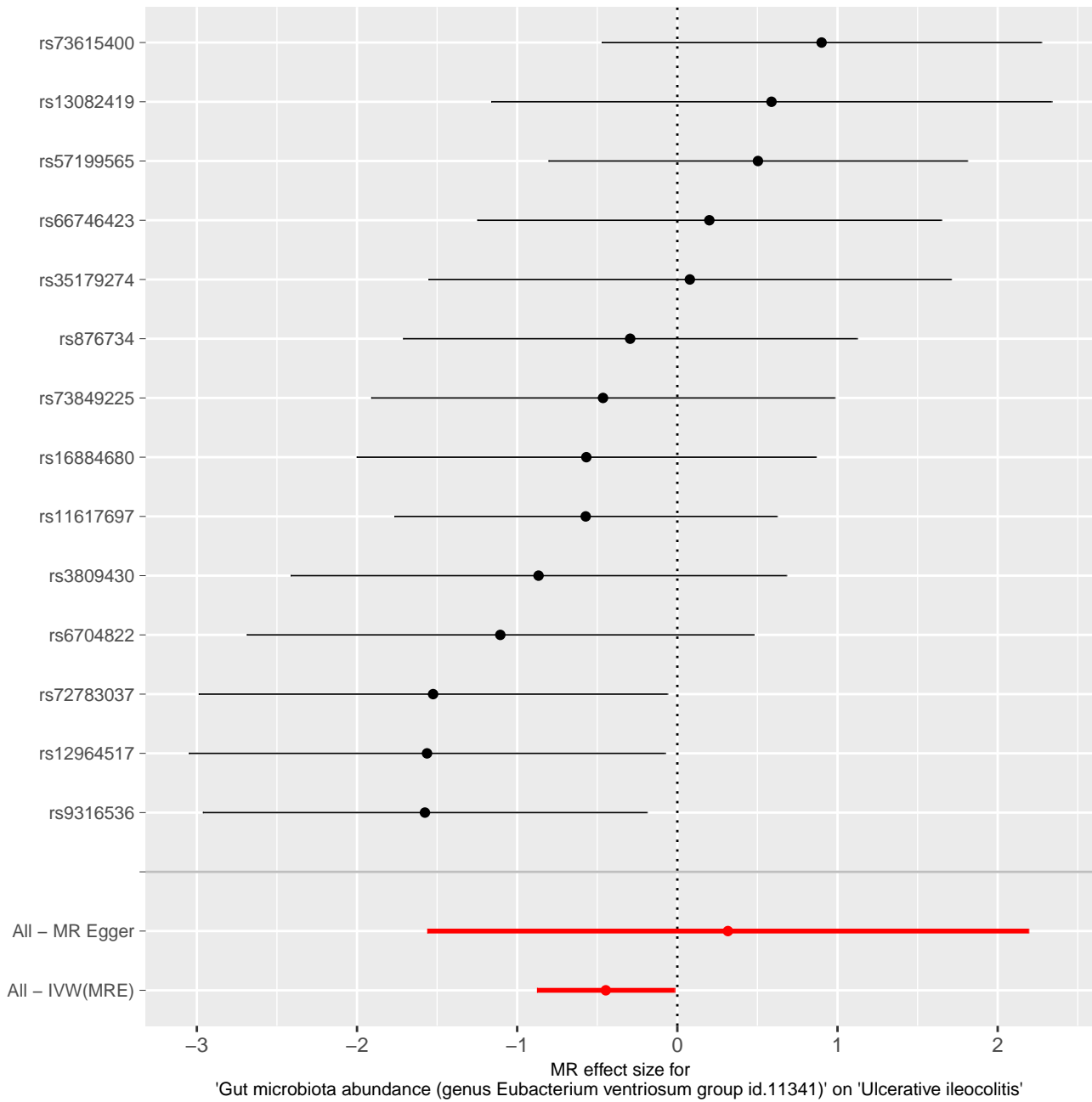


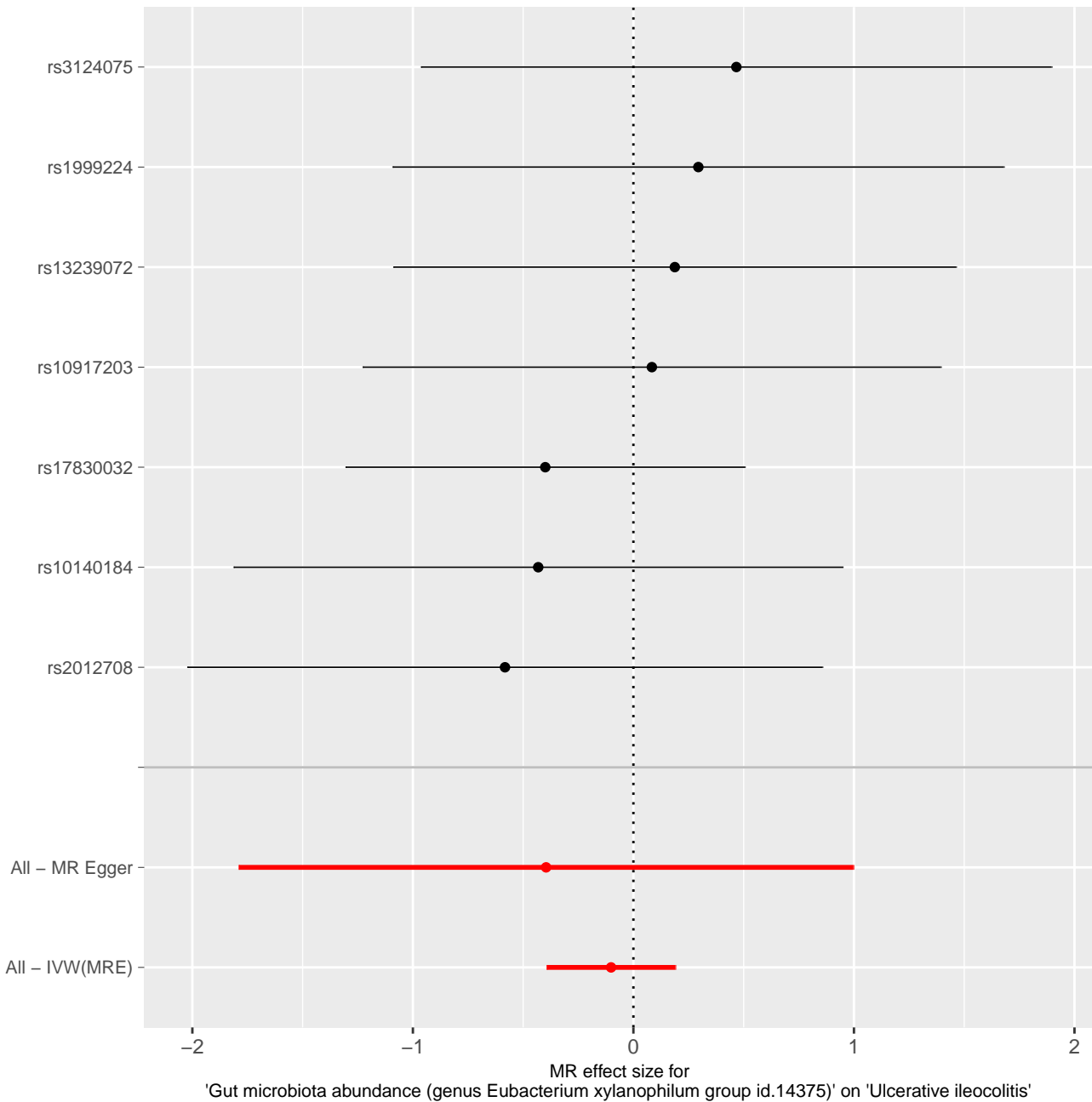




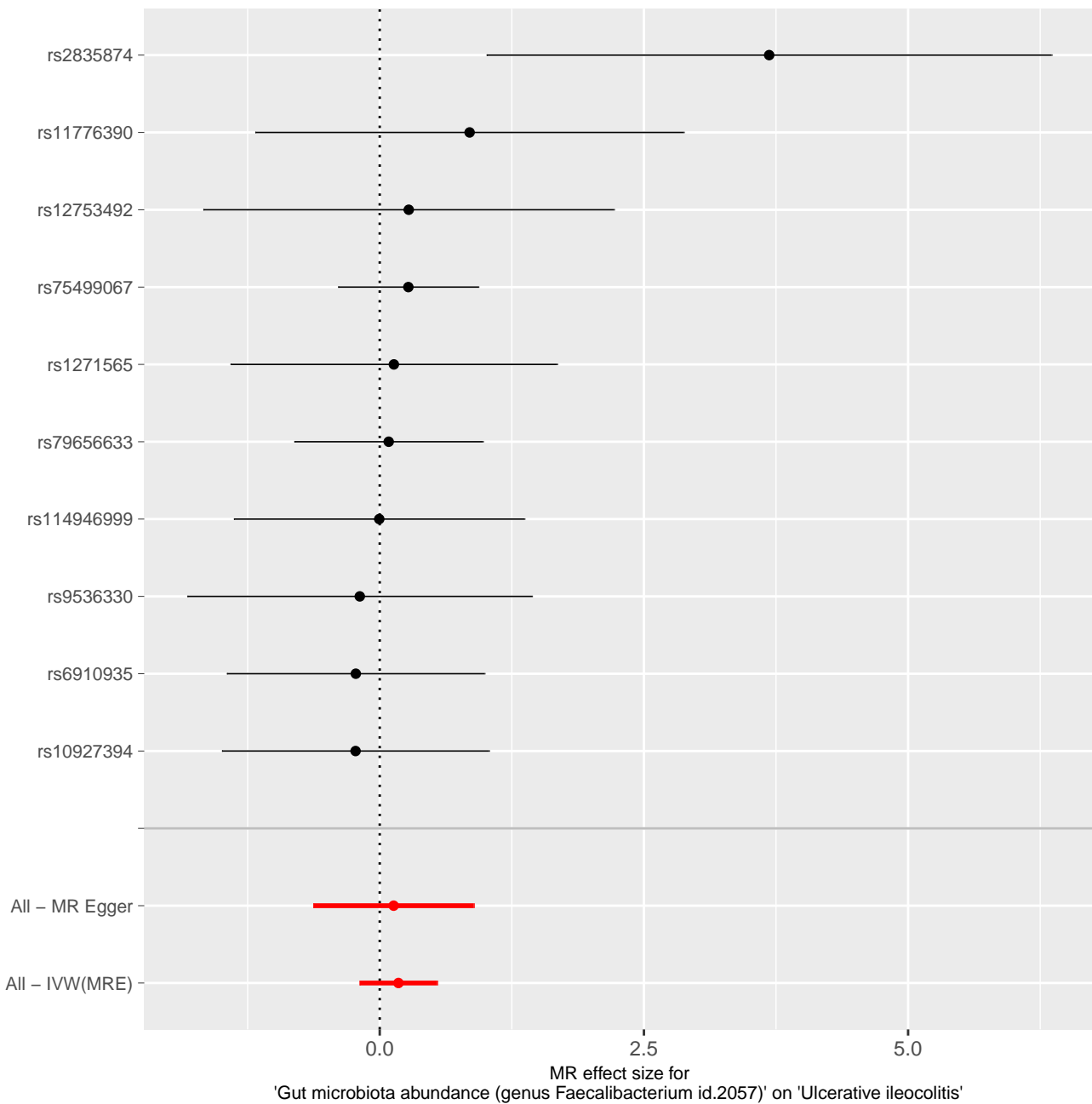


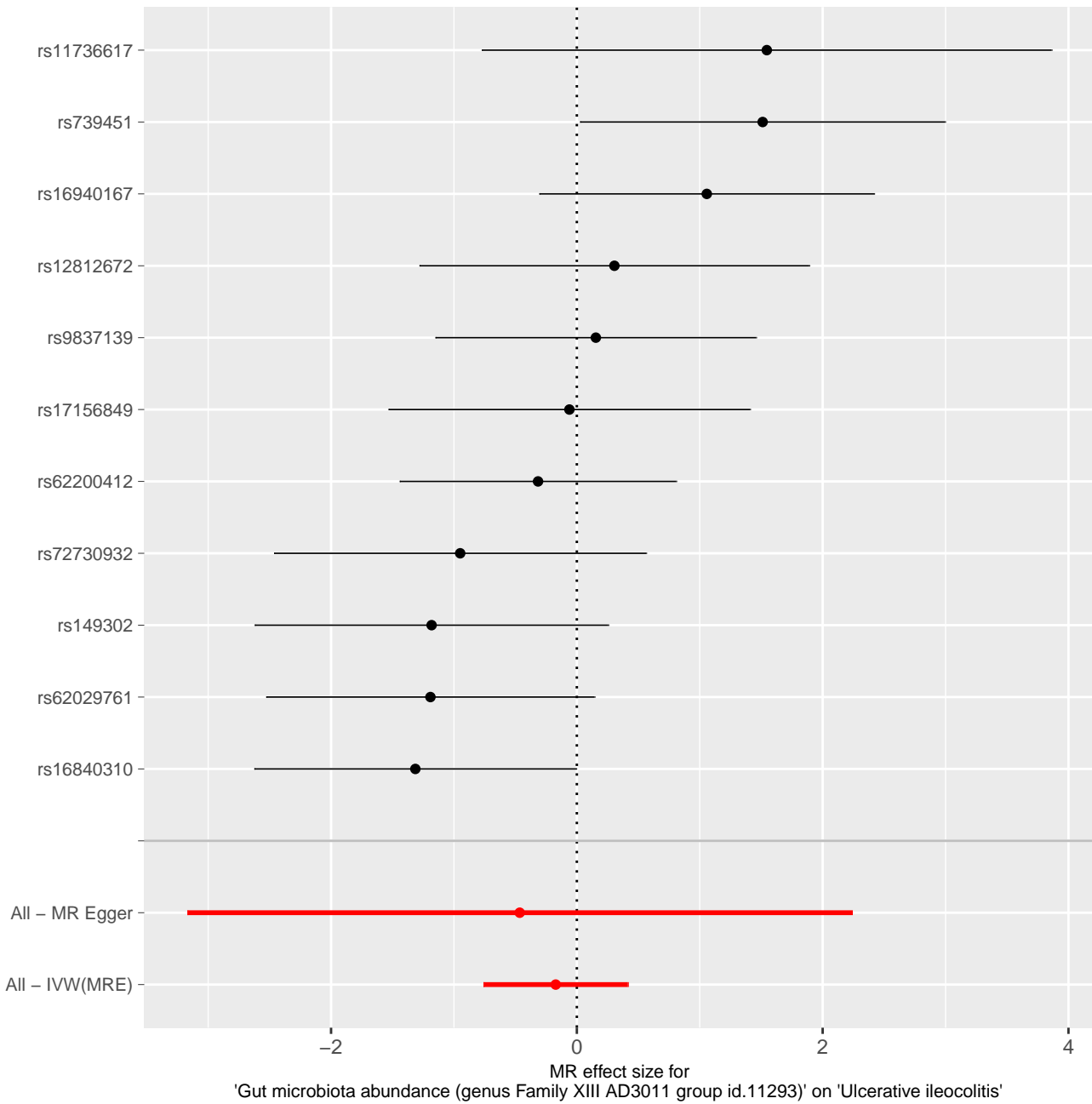


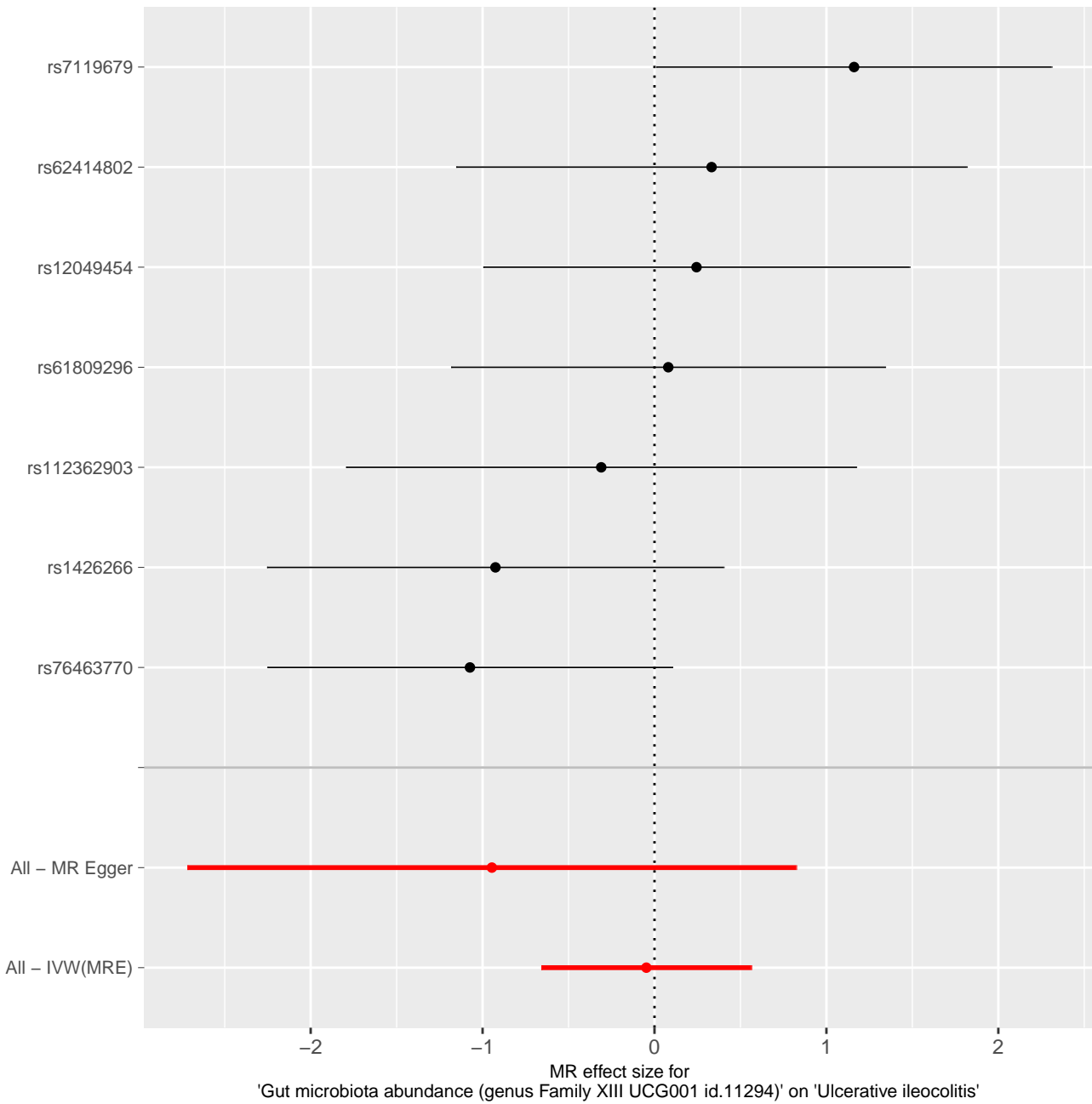


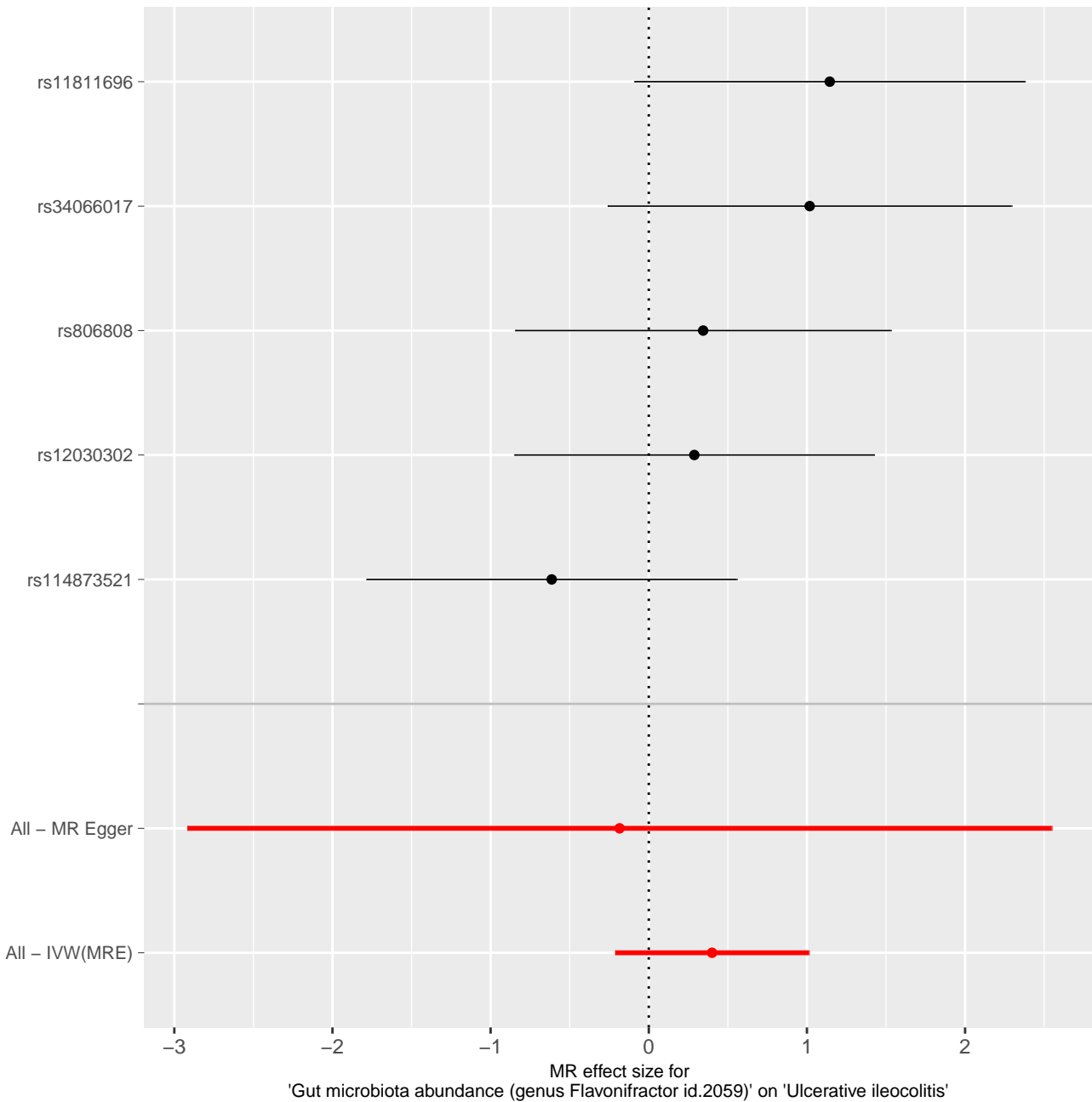


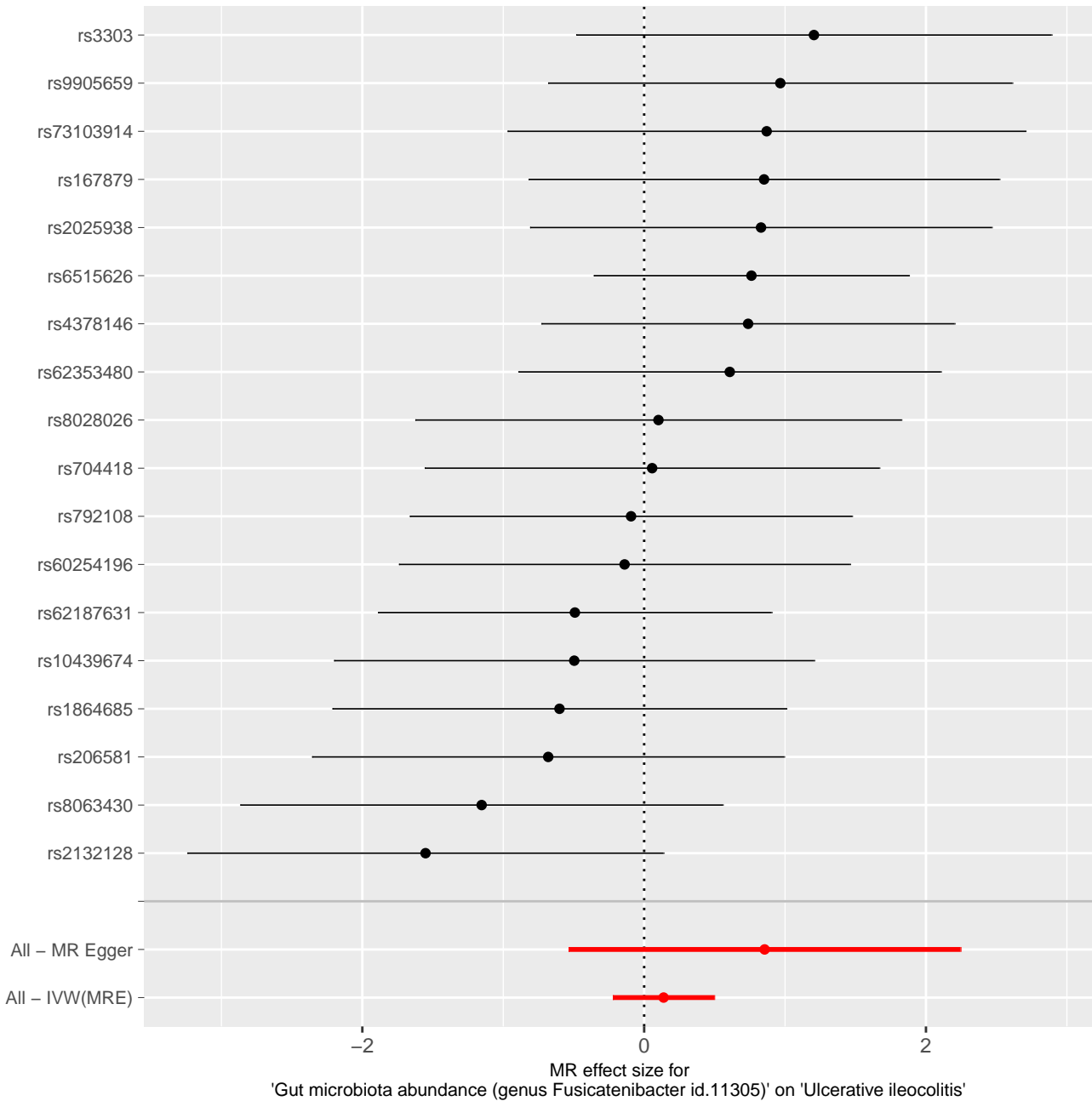
MR effect size for
'Gut microbiota abundance (genus Eubacterium xylanophilum group id.14375)' on 'Ulcerative ileocolitis'

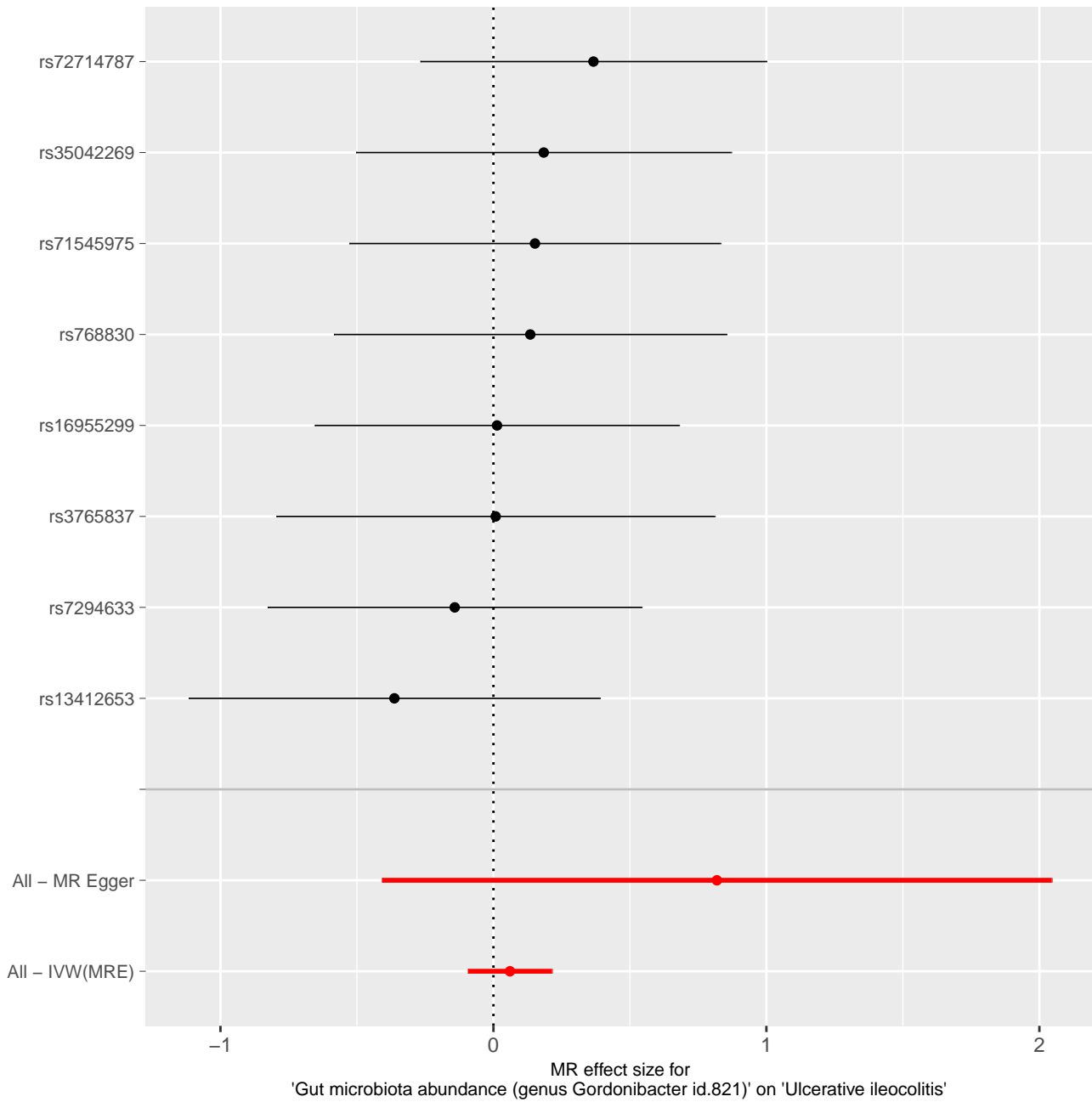


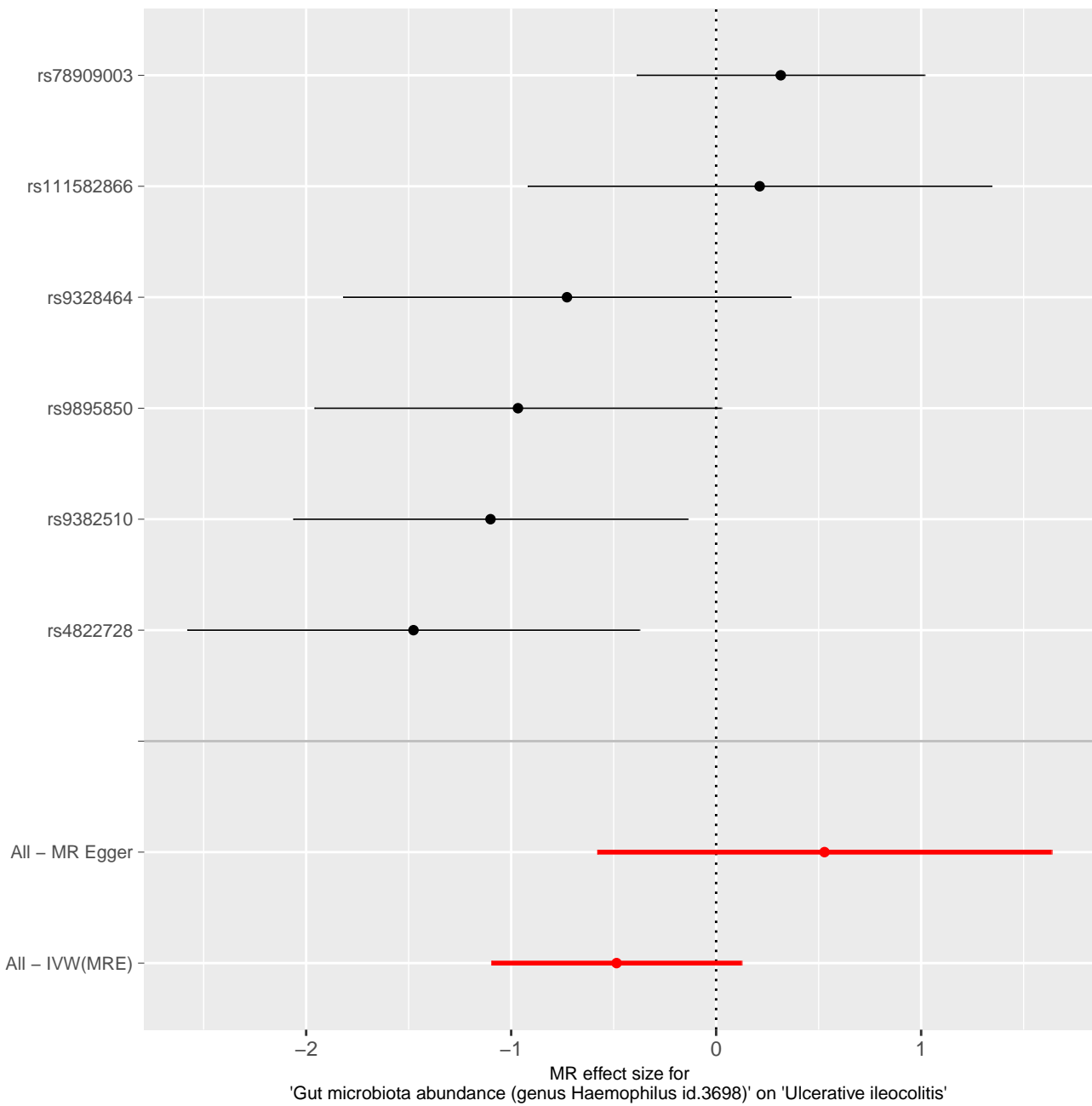


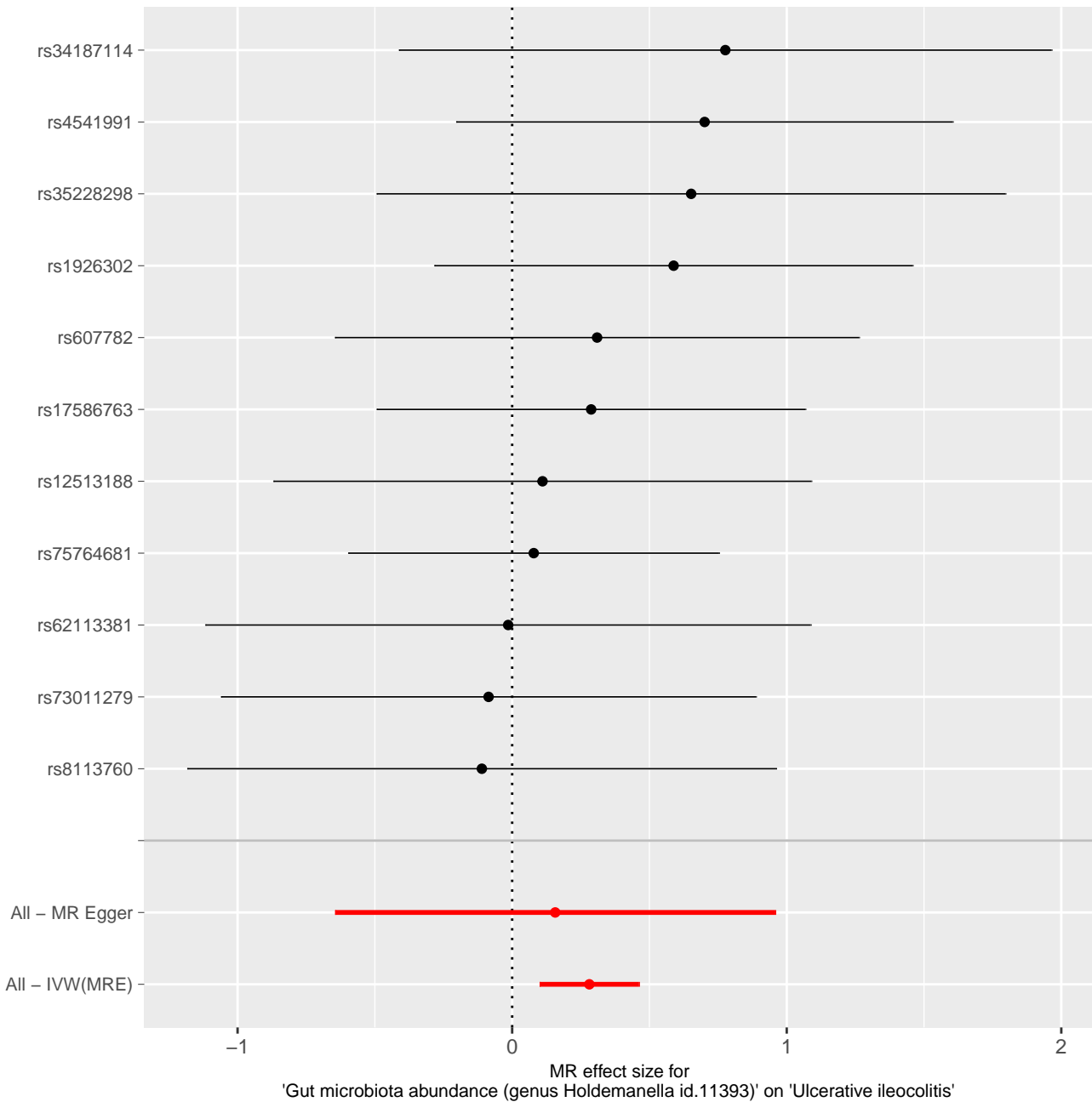


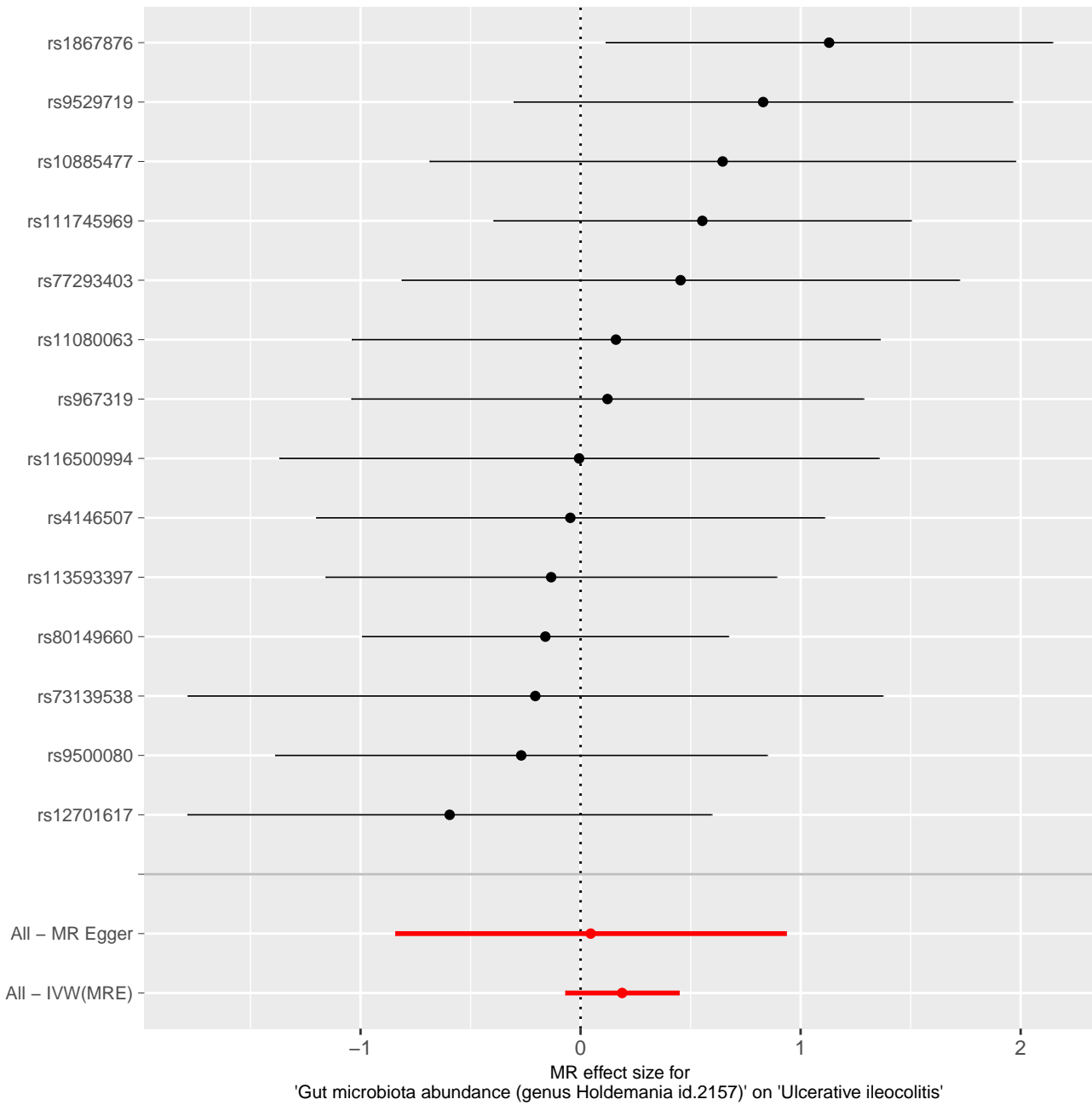


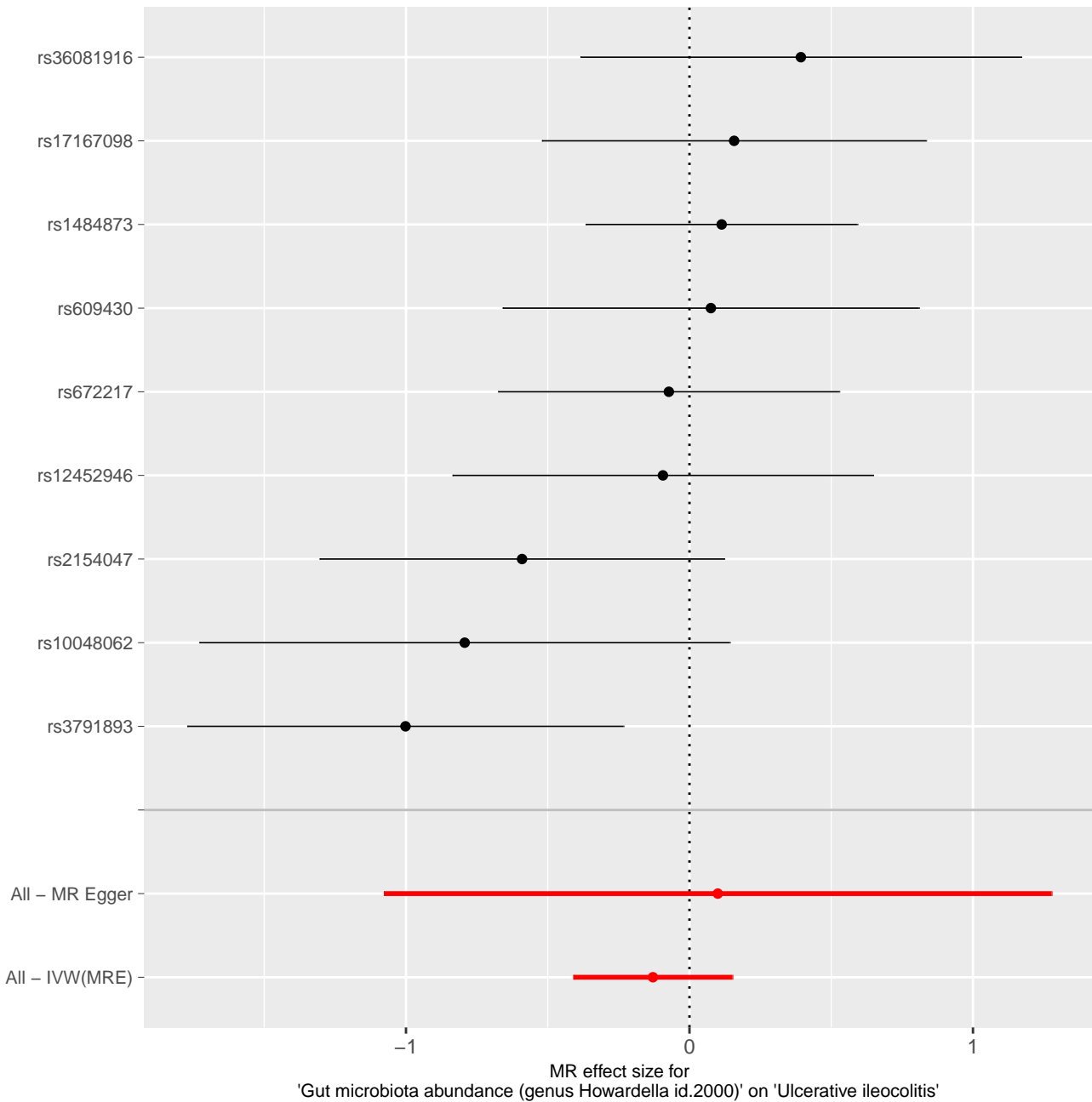


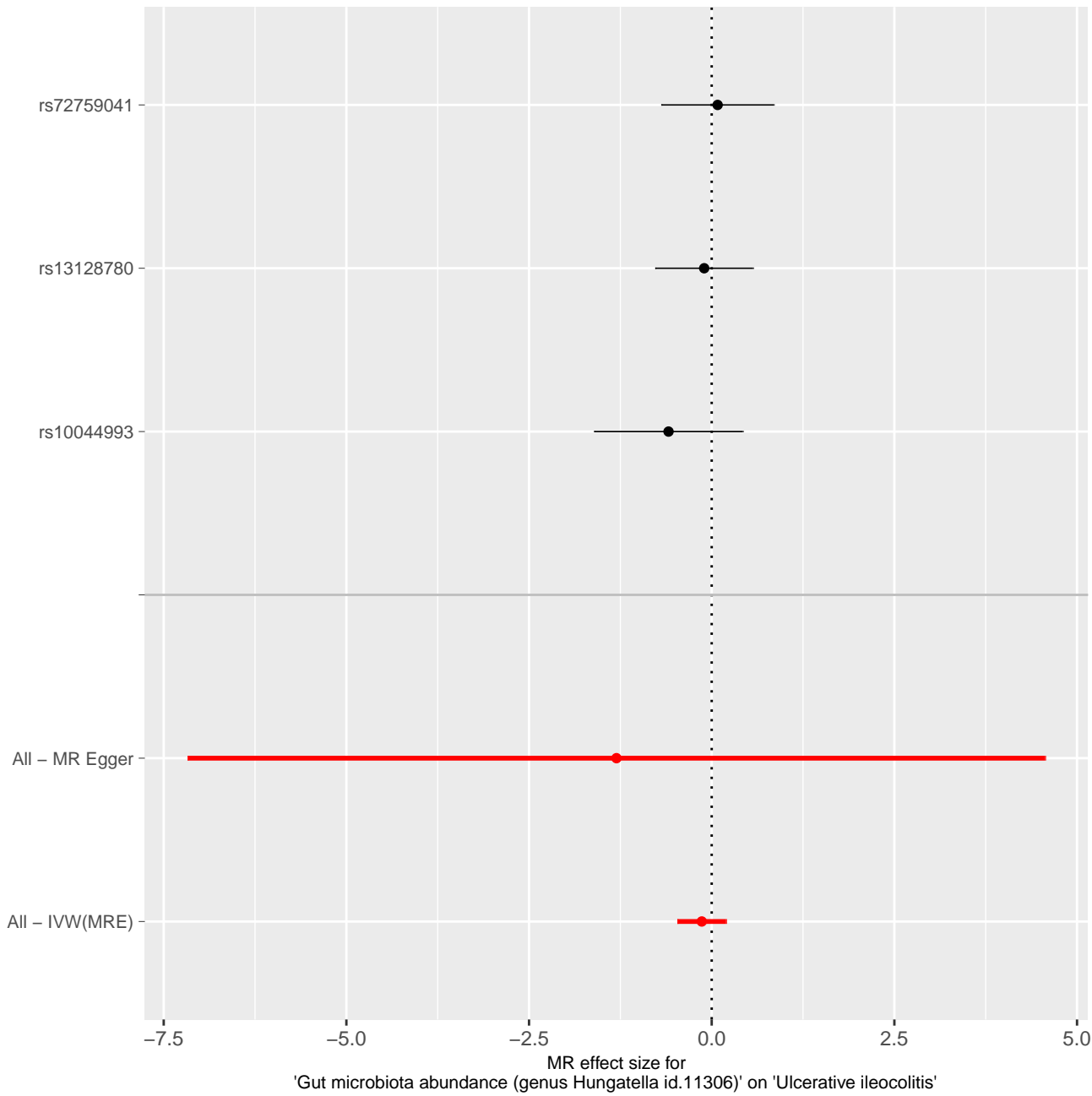




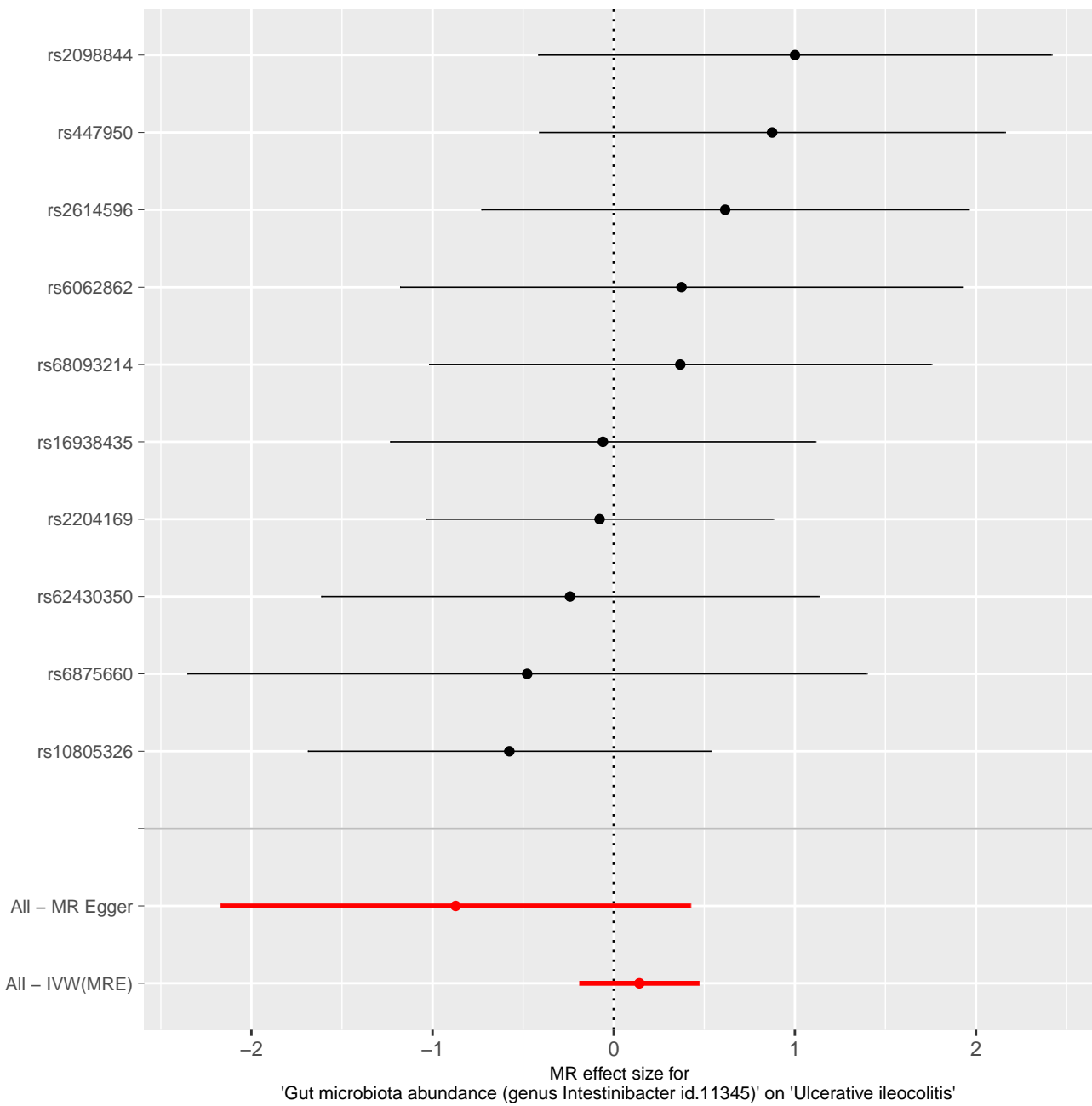


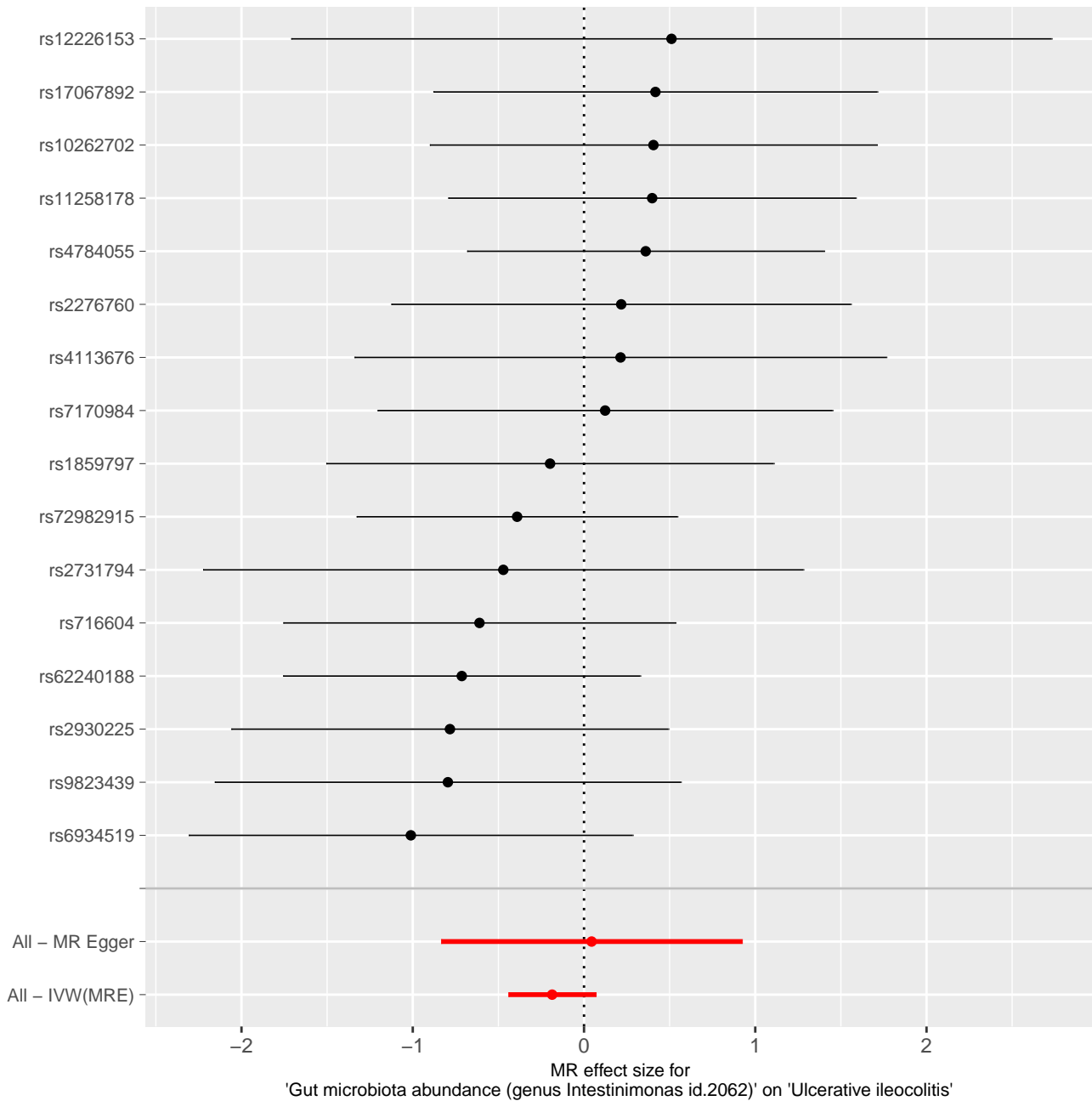


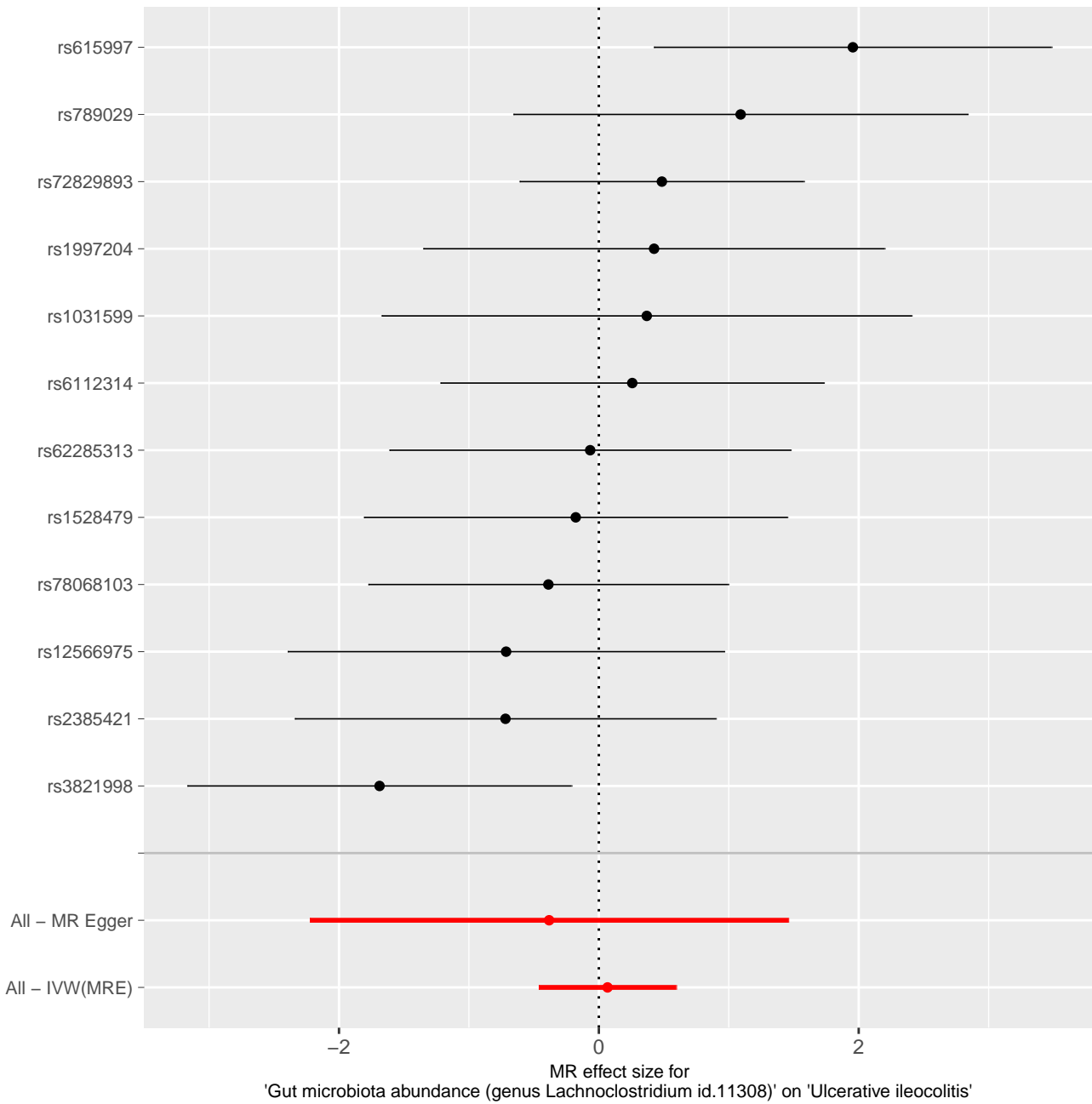


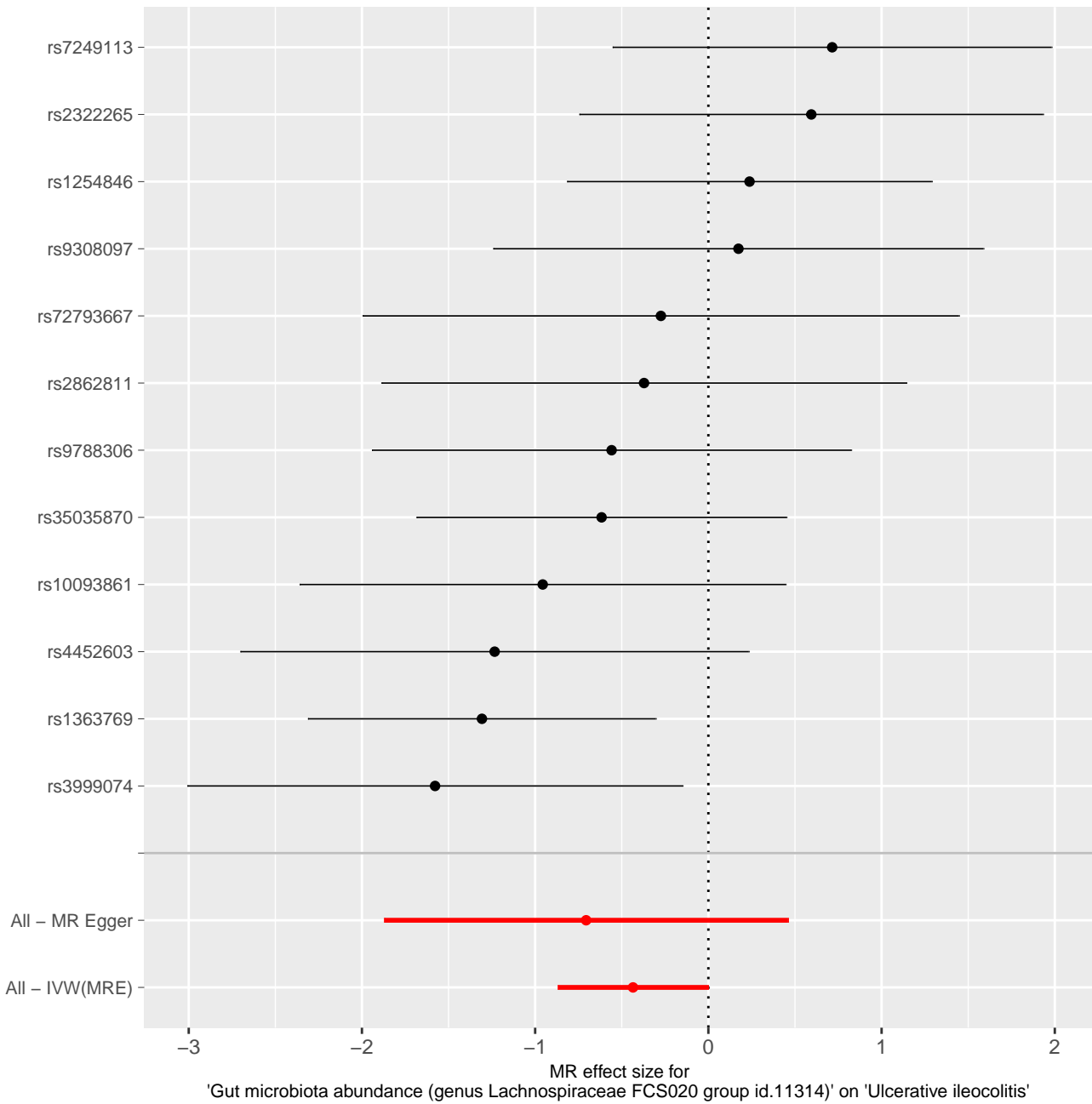


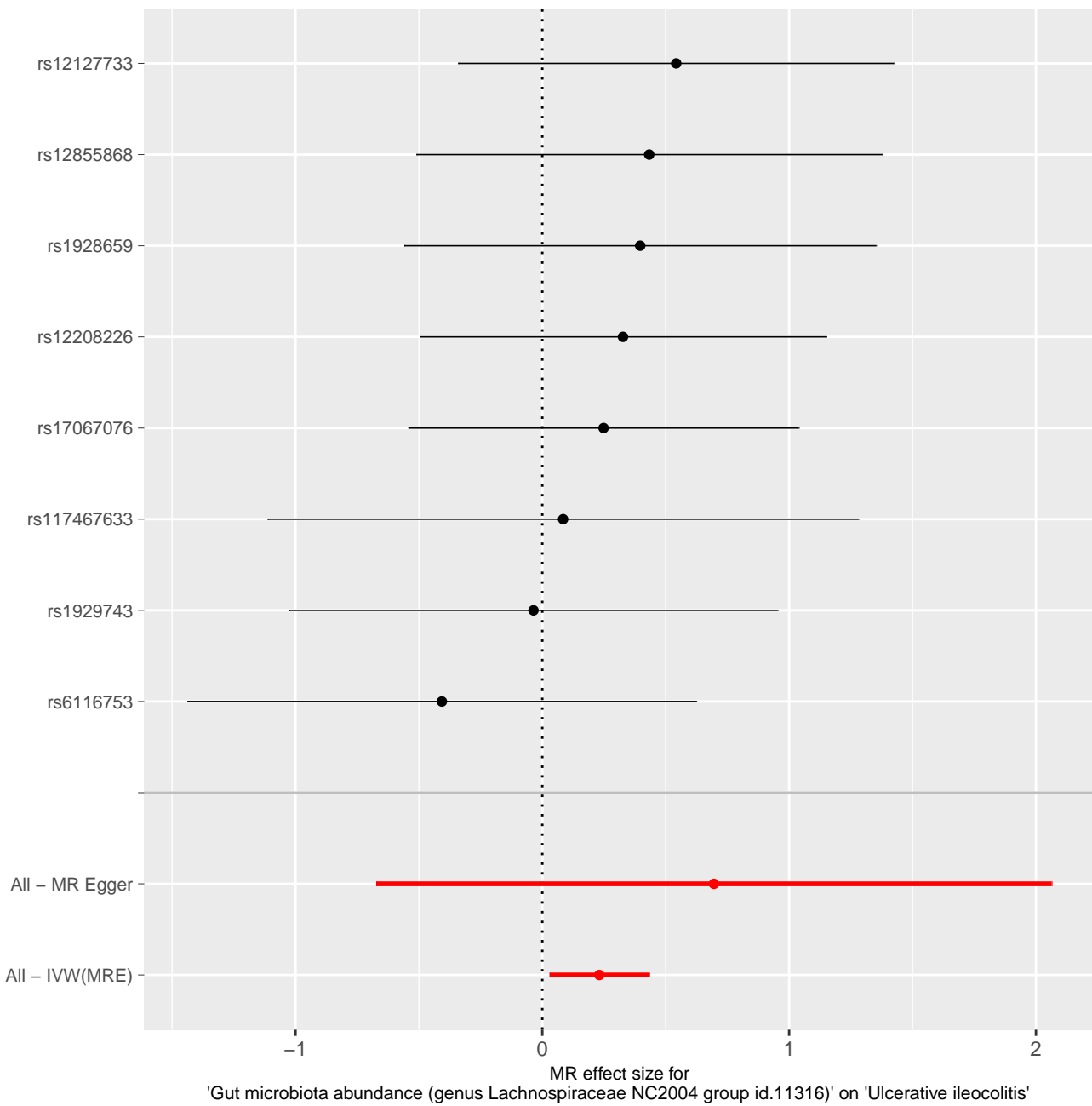
Batch 1142 : Gut microbiota abundance (genus Intestinibacter id.11345) on Ulcerative ileocolitis

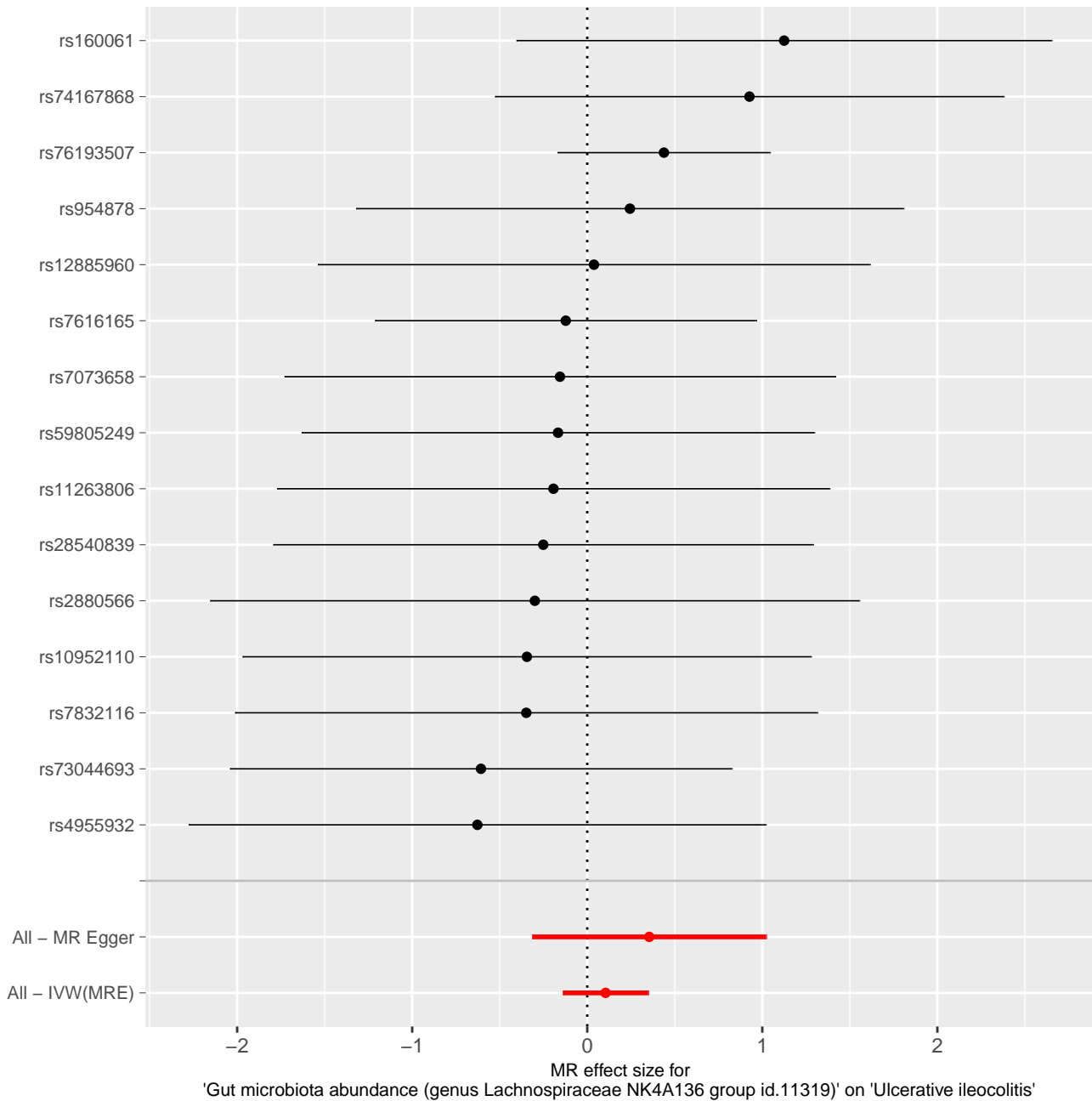


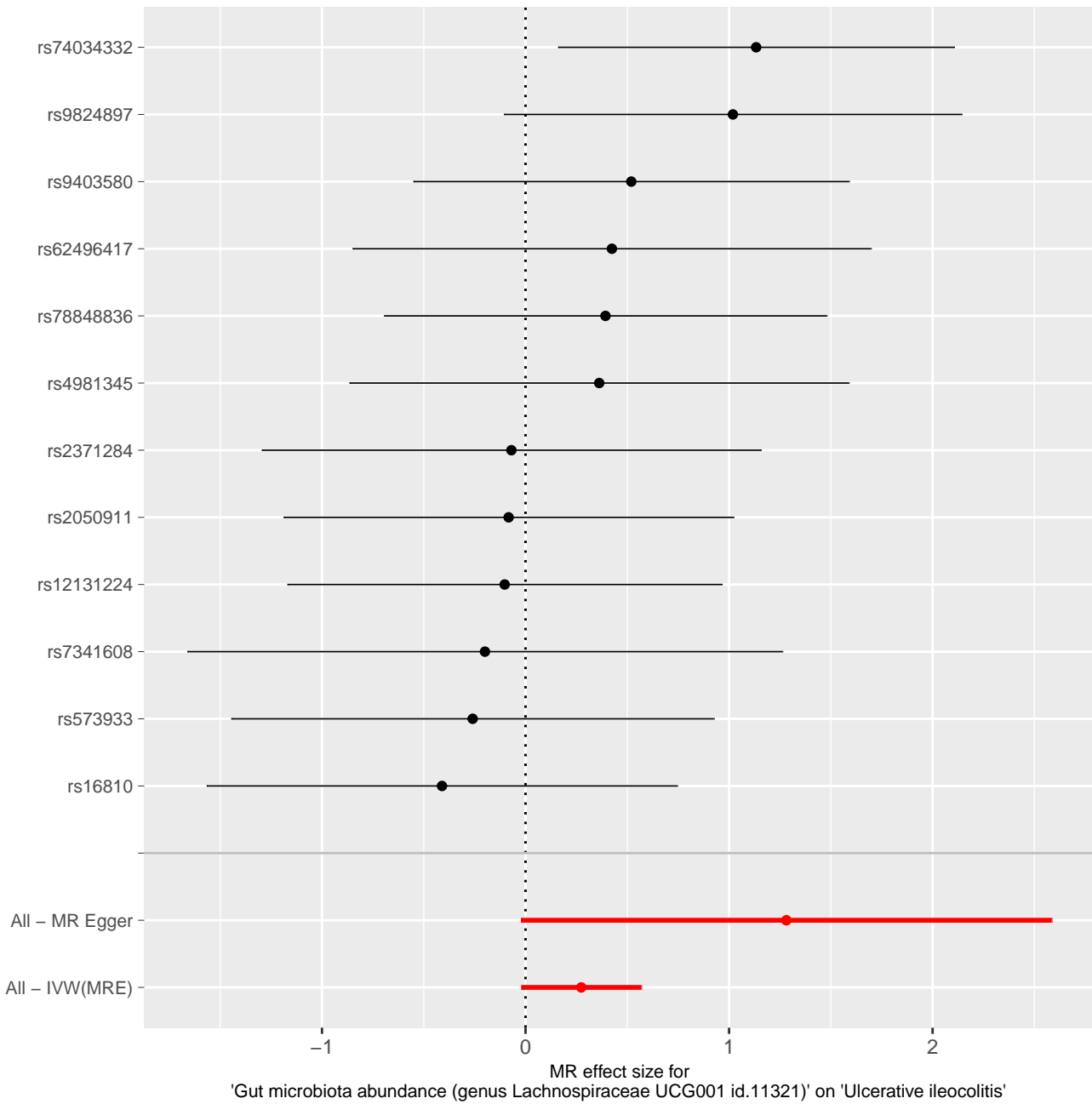


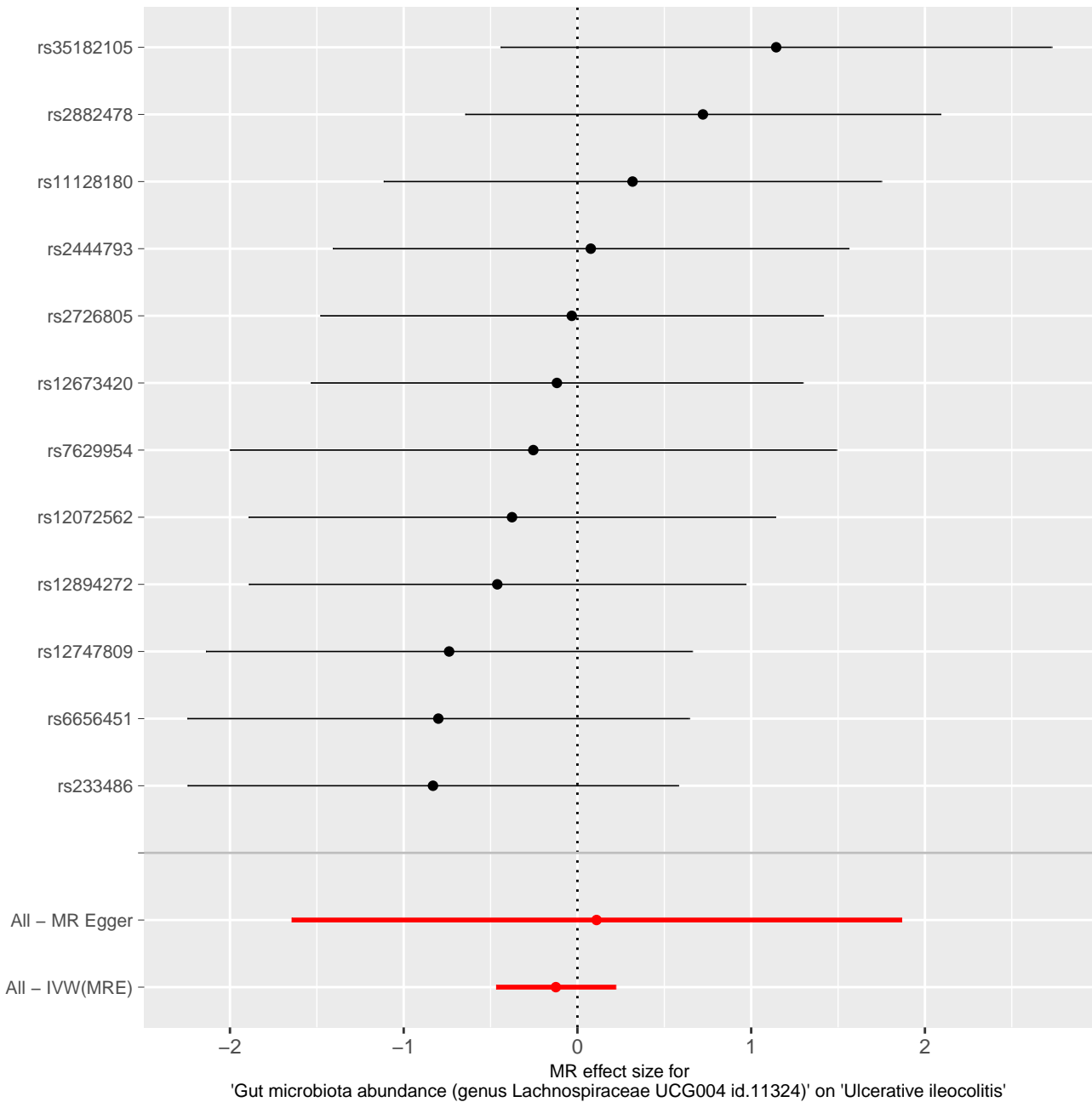


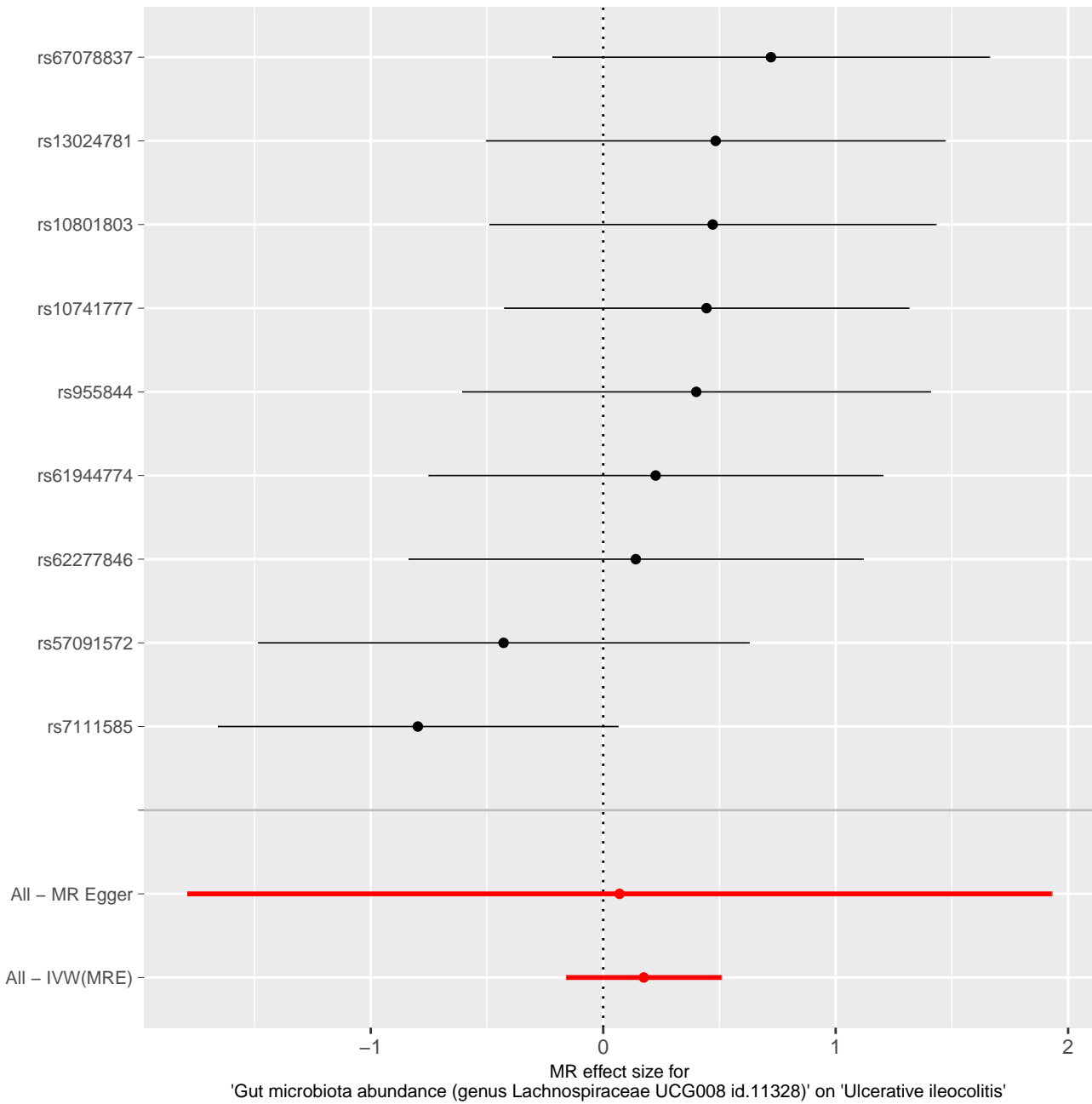


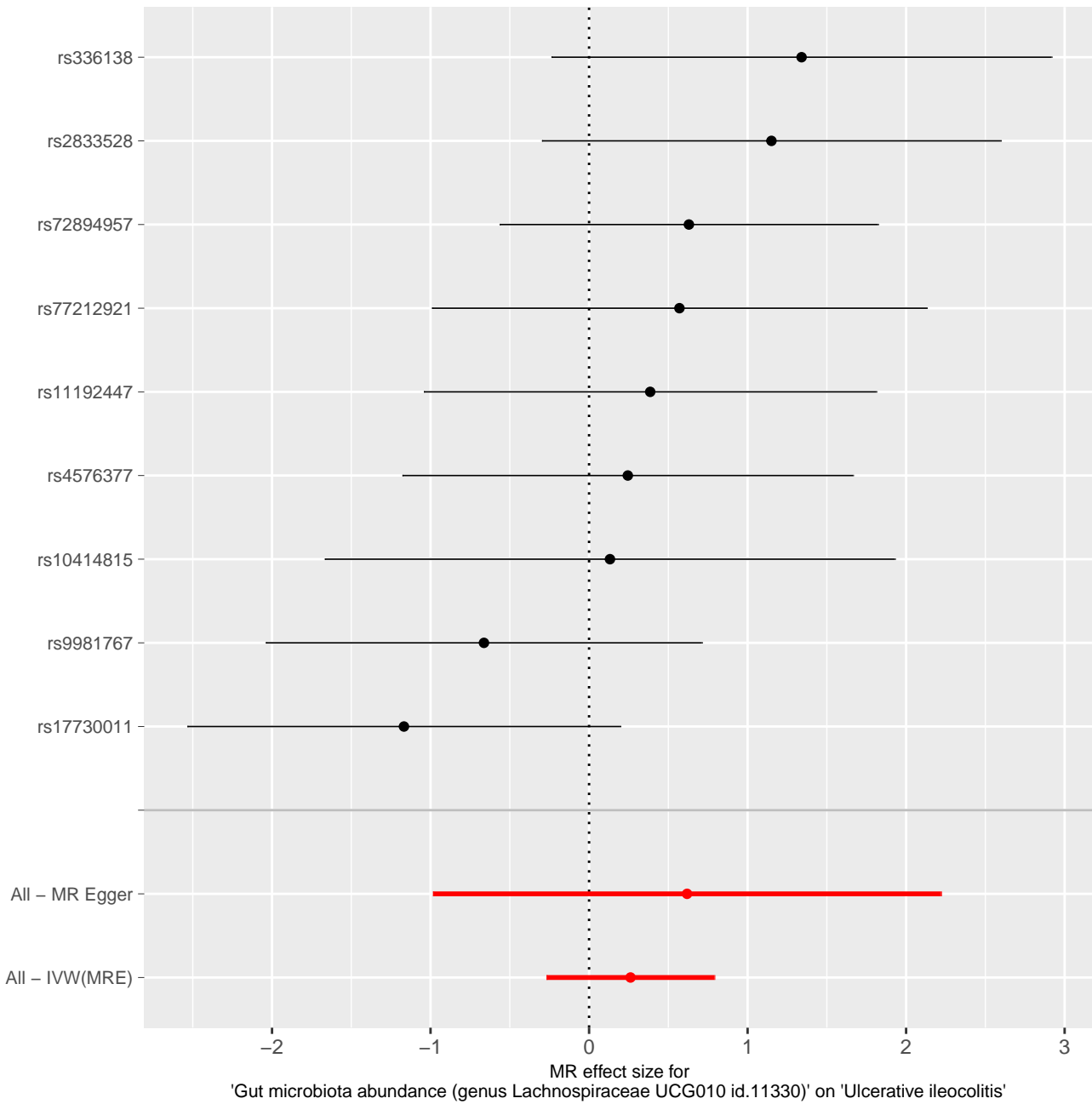


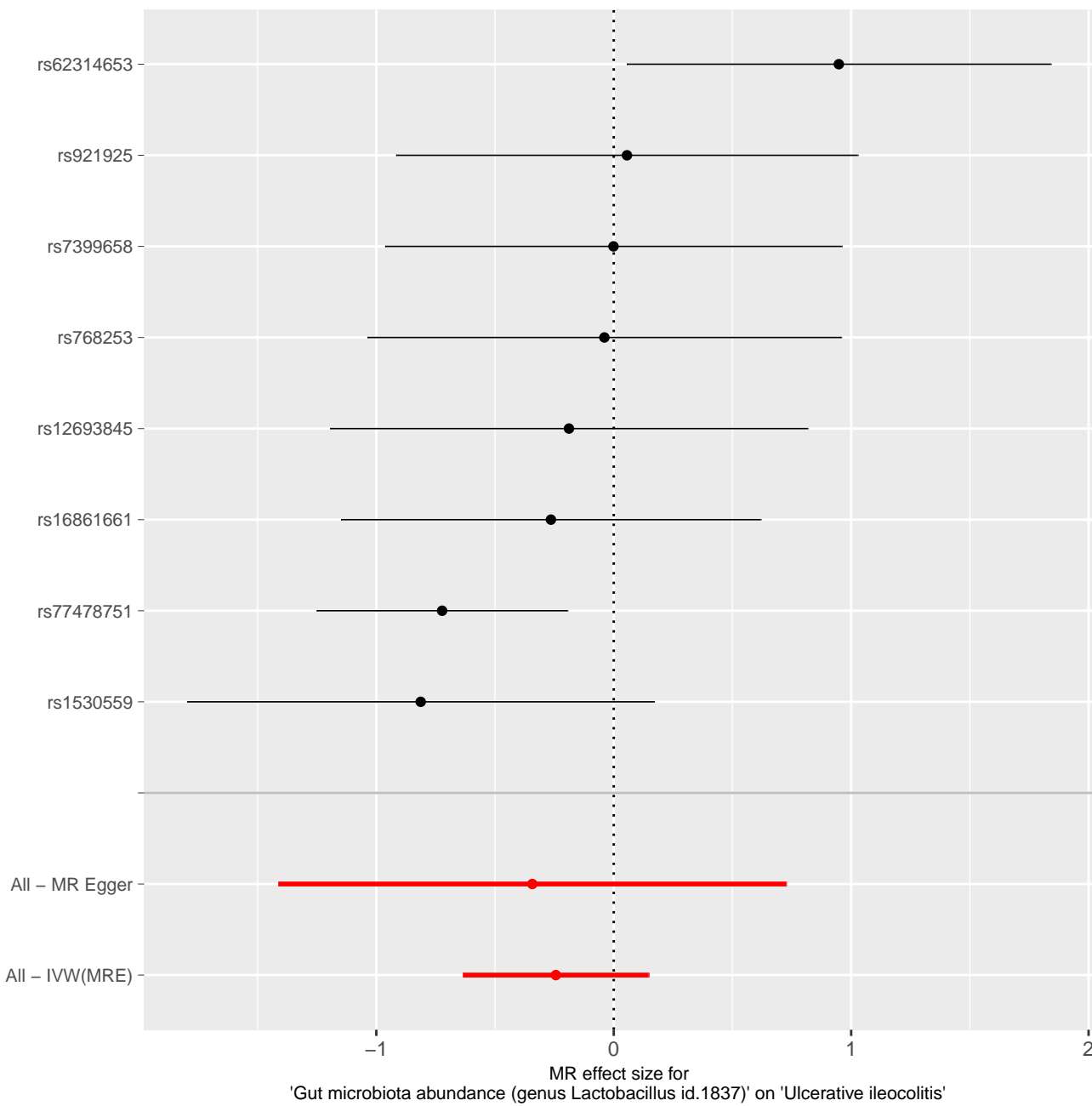


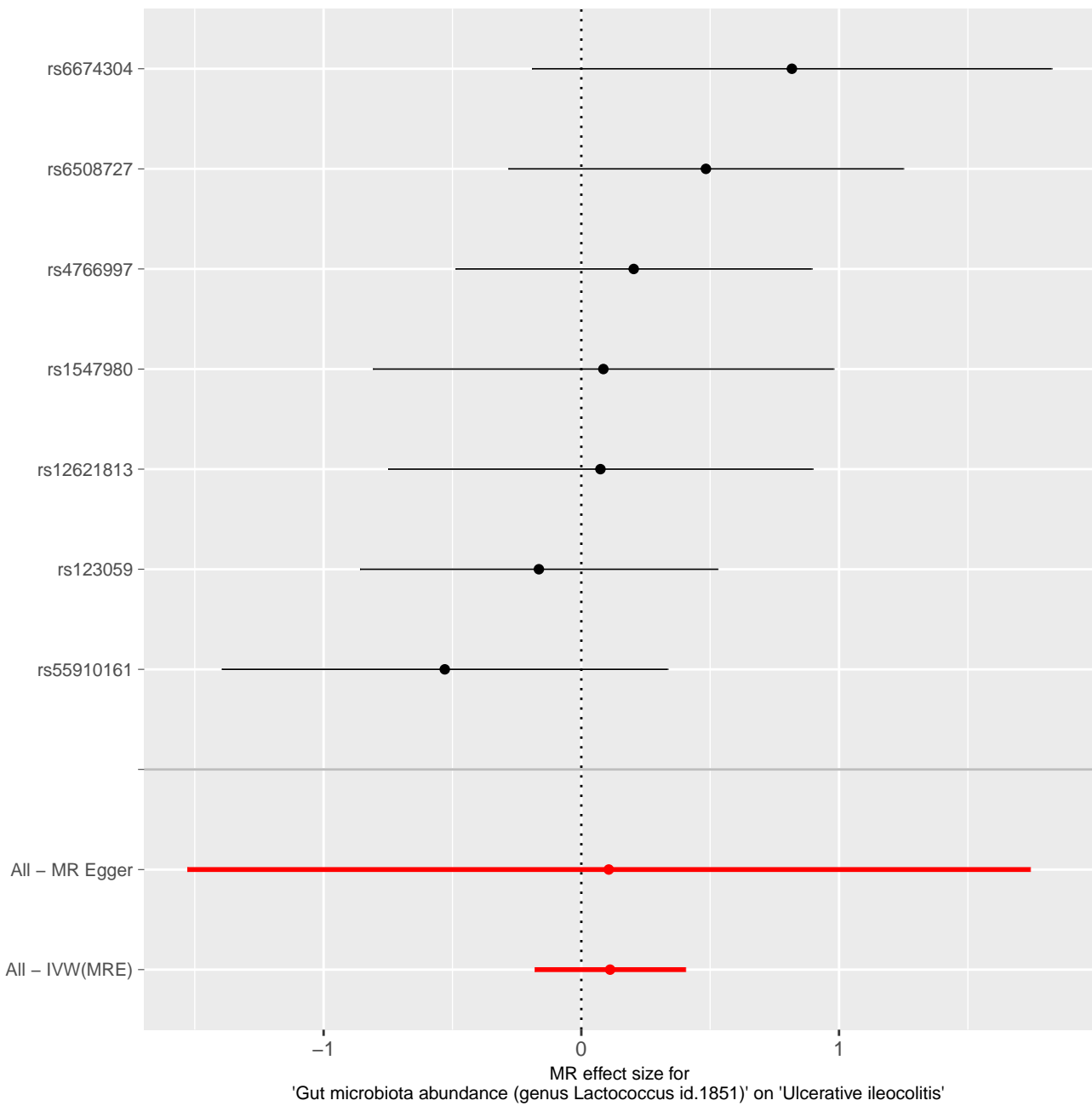


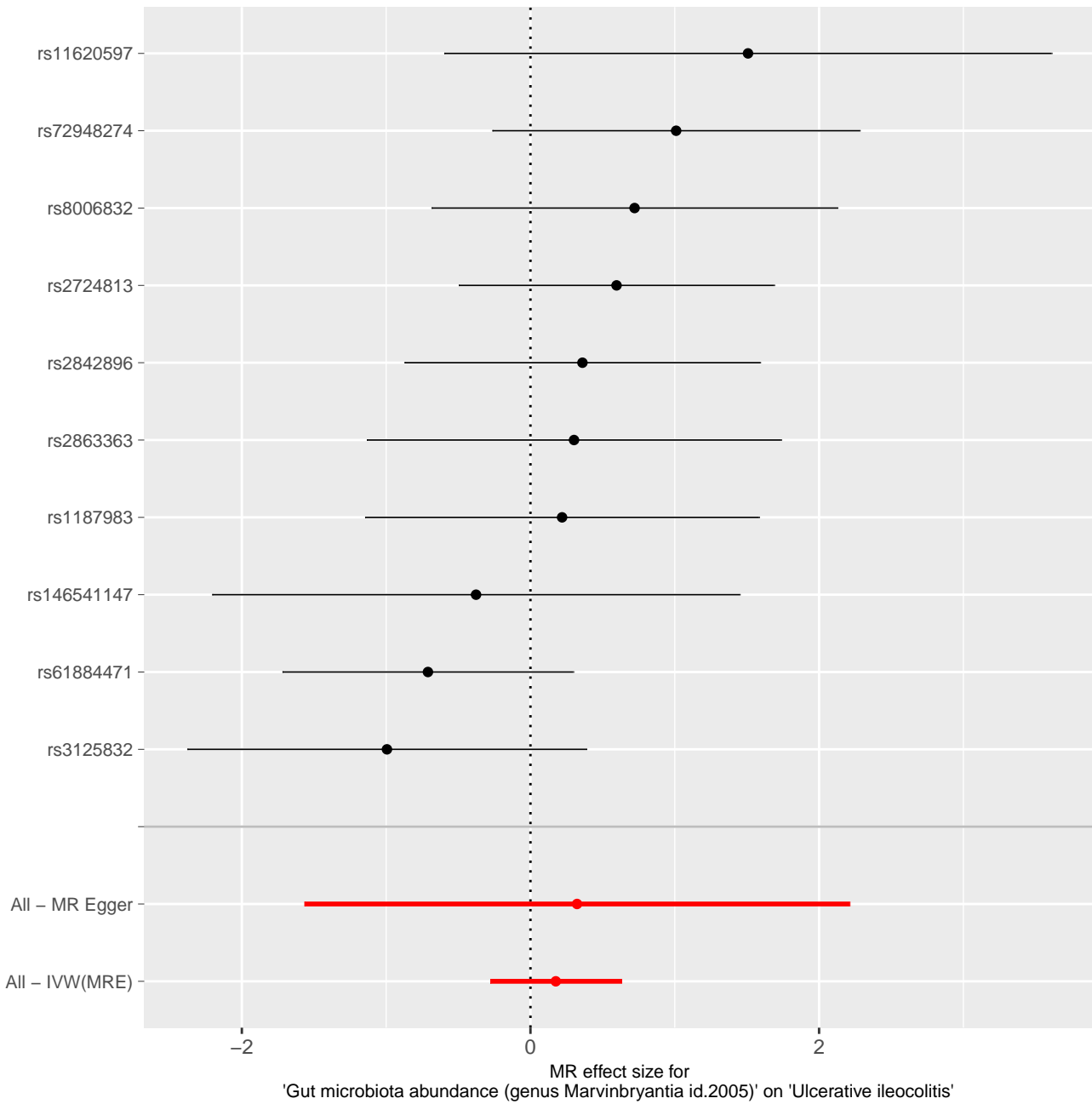


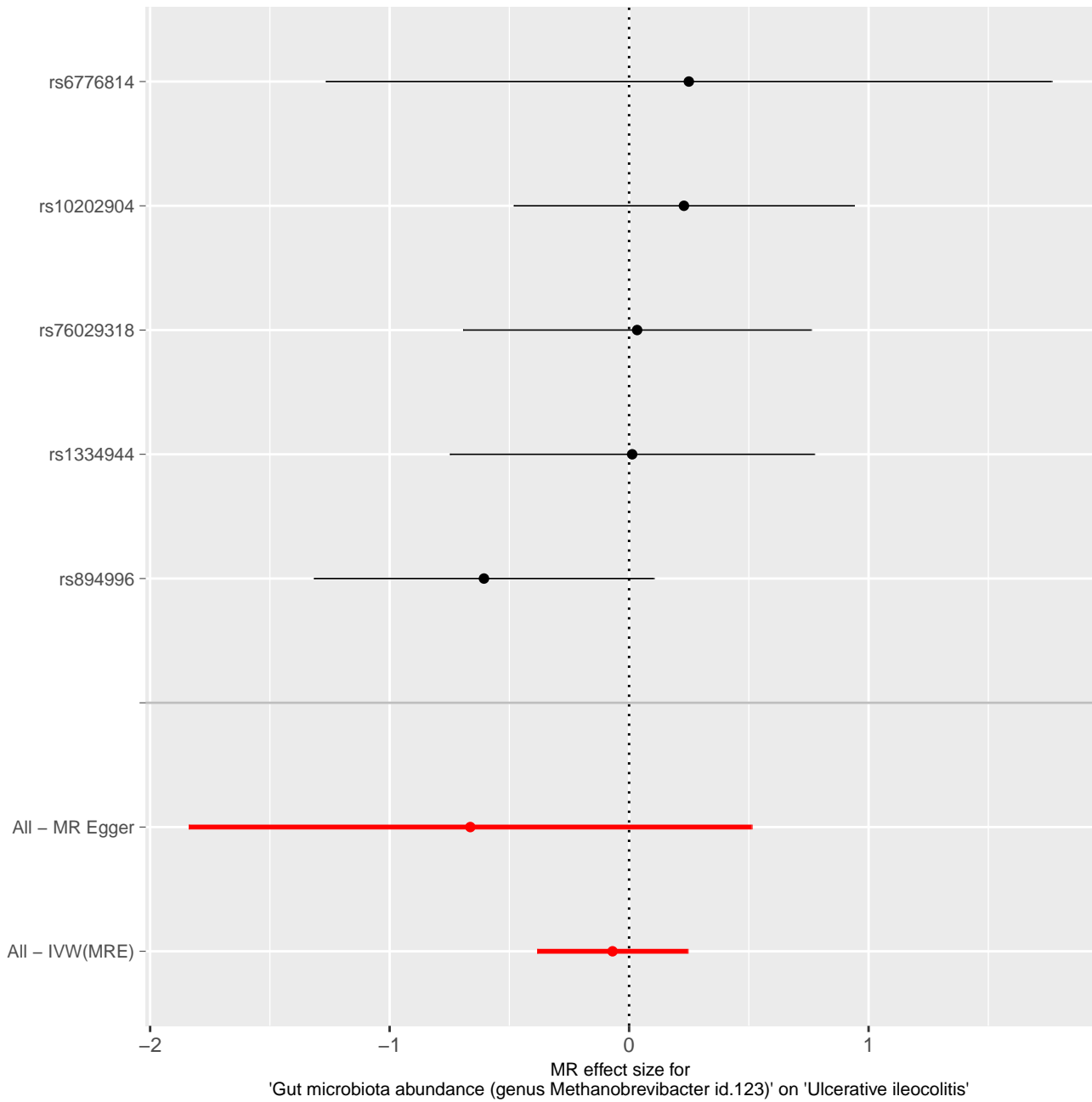


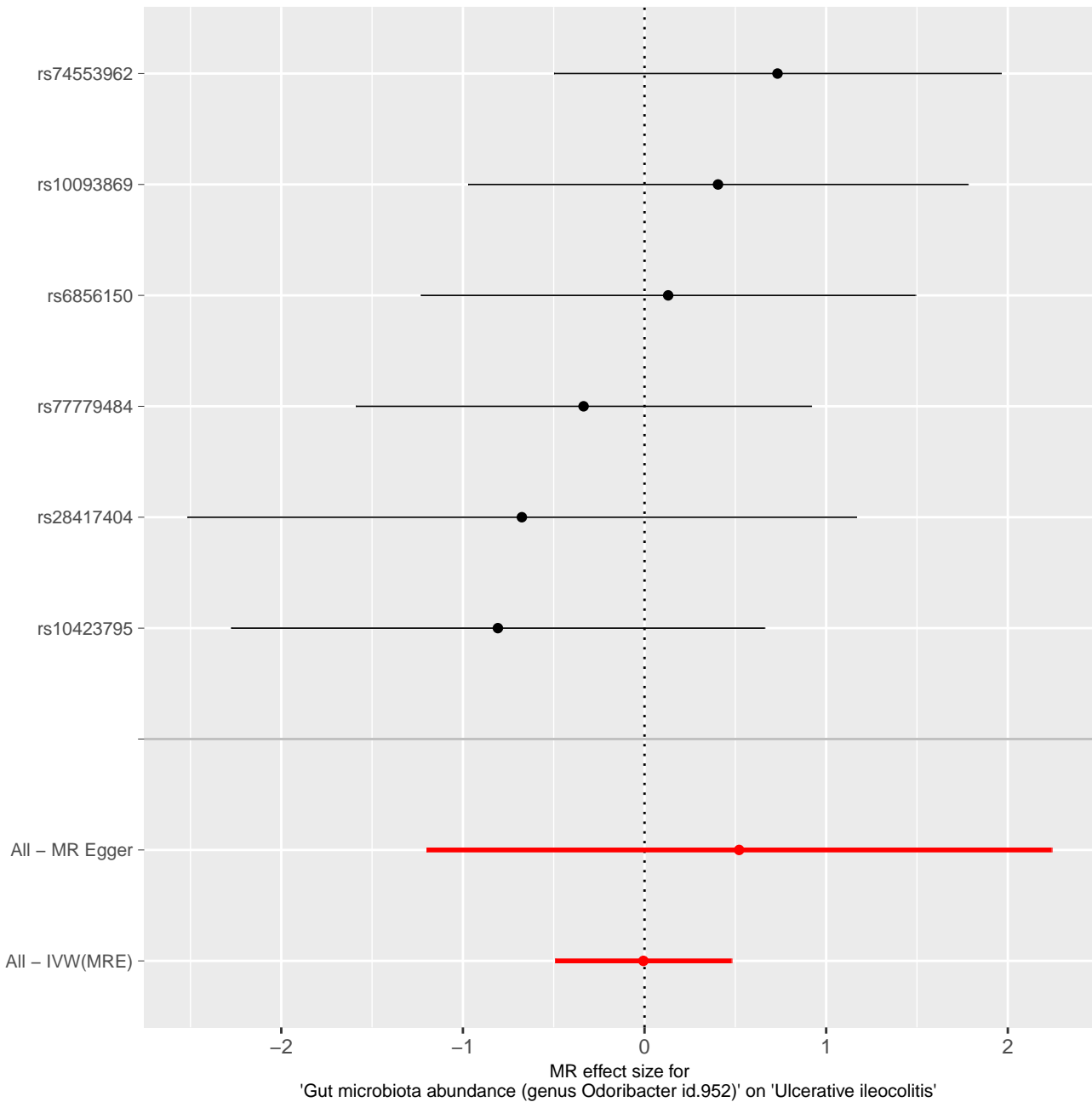


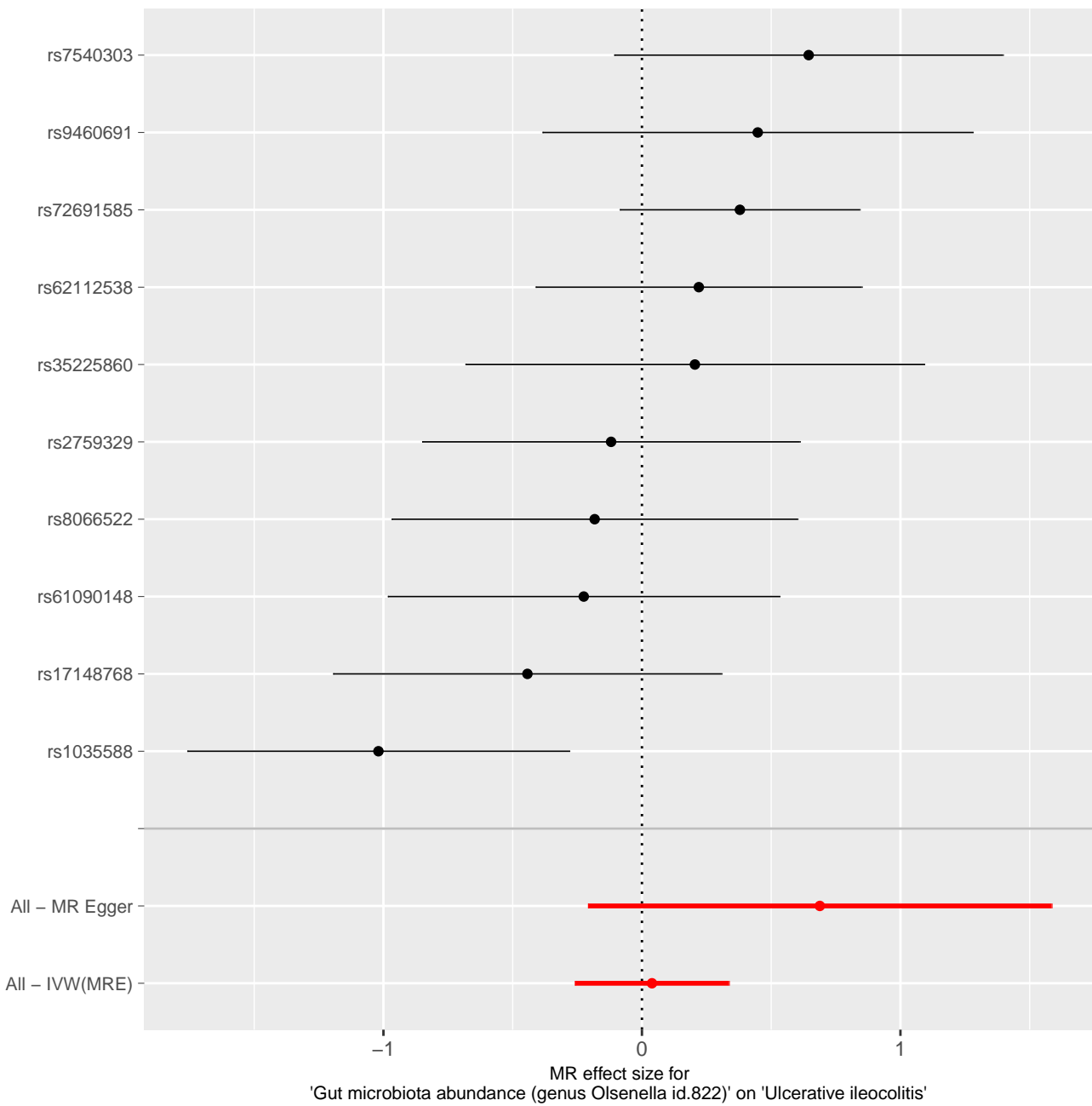




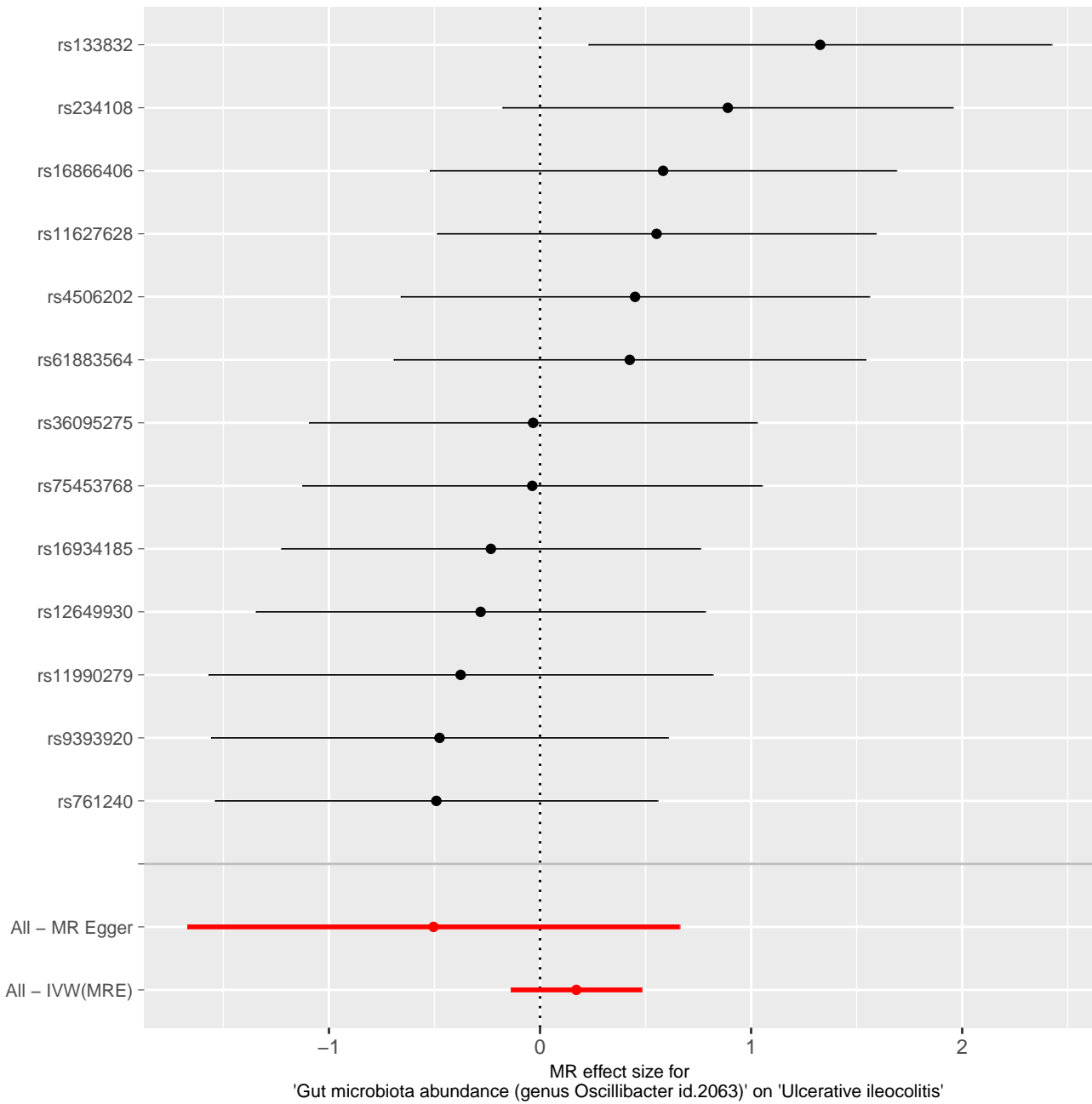


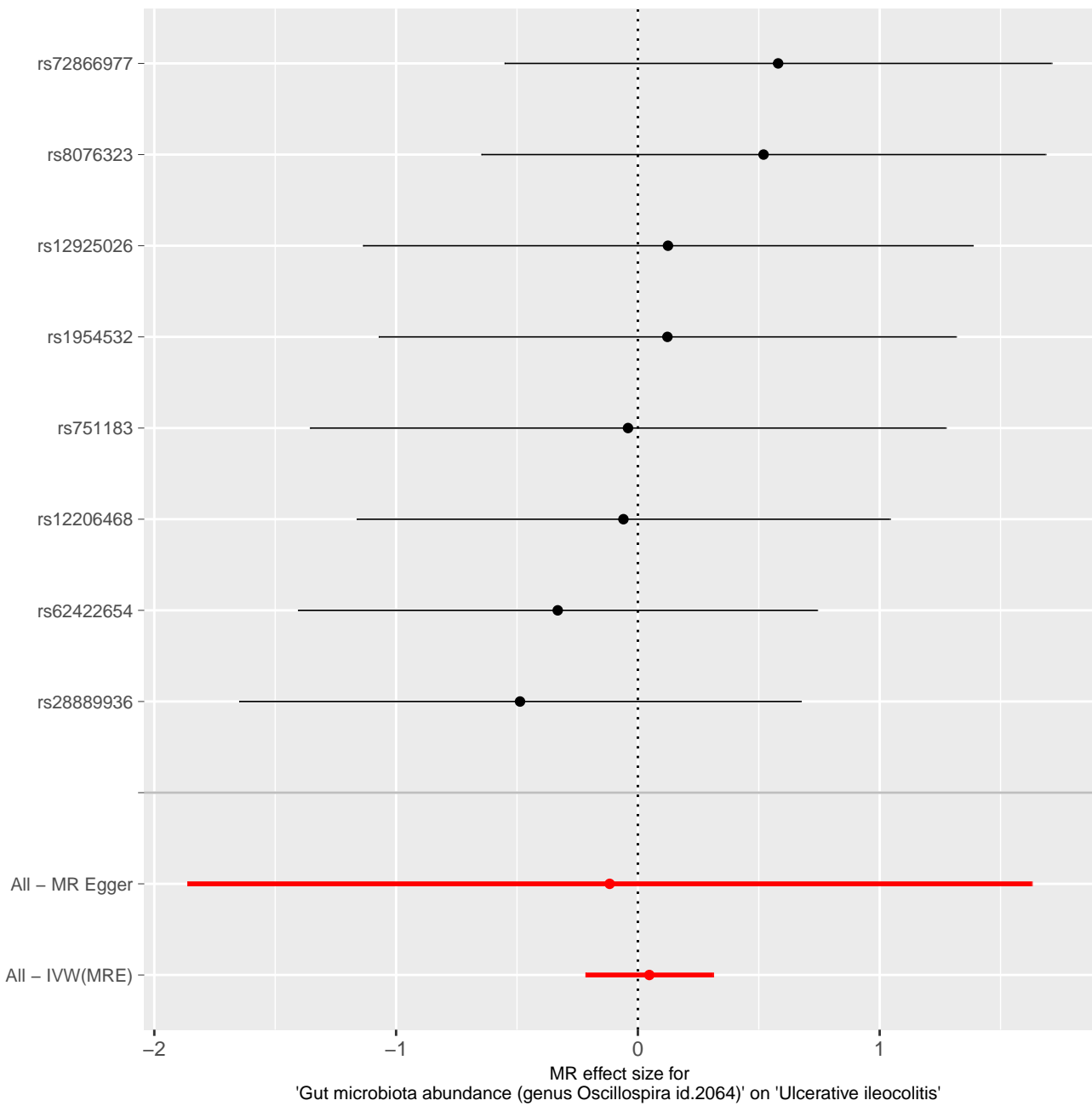


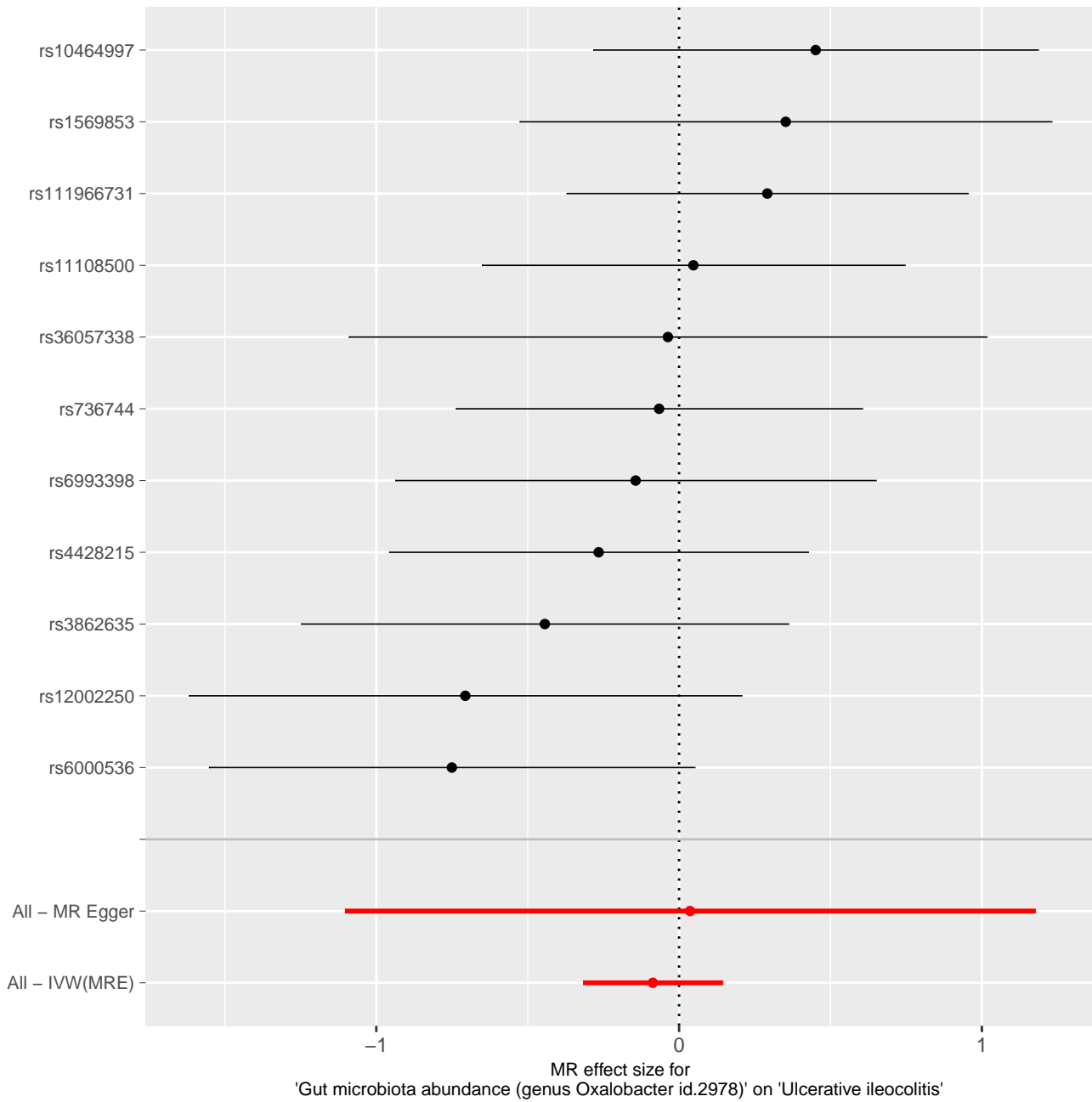


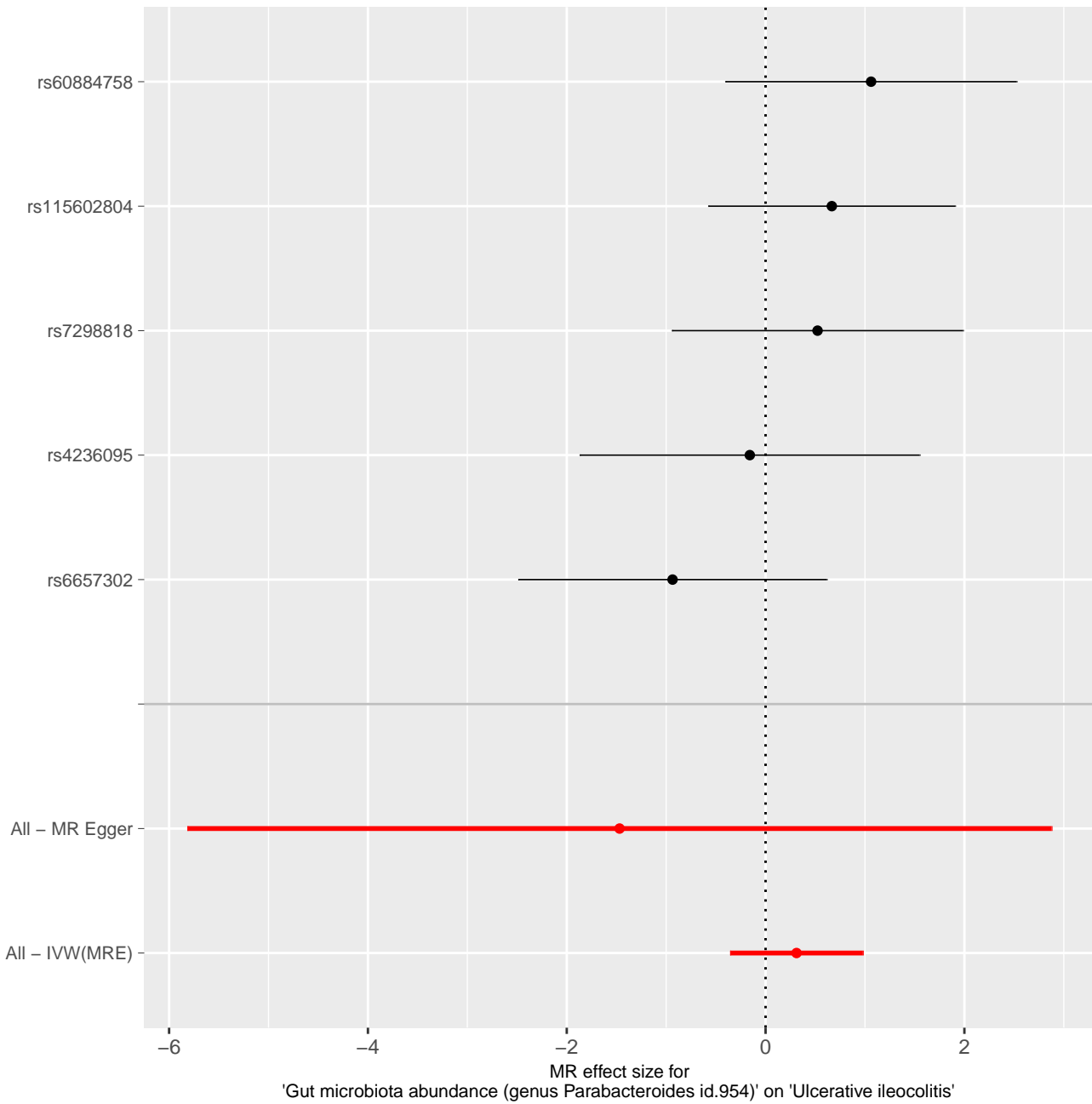


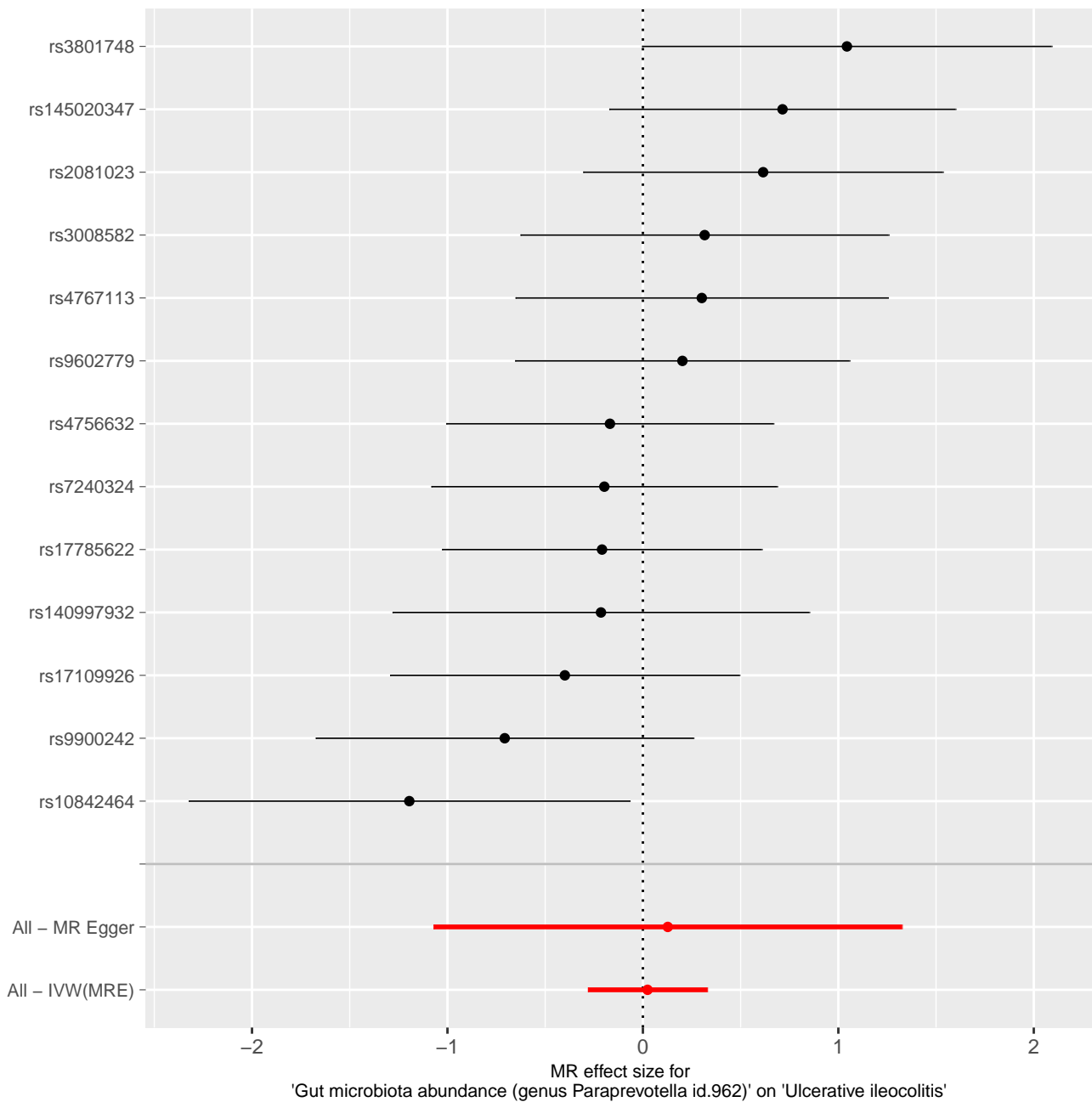
Batch 1160 : Gut microbiota abundance (genus Oscillibacter id.2063) on Ulcerative ileocolitis

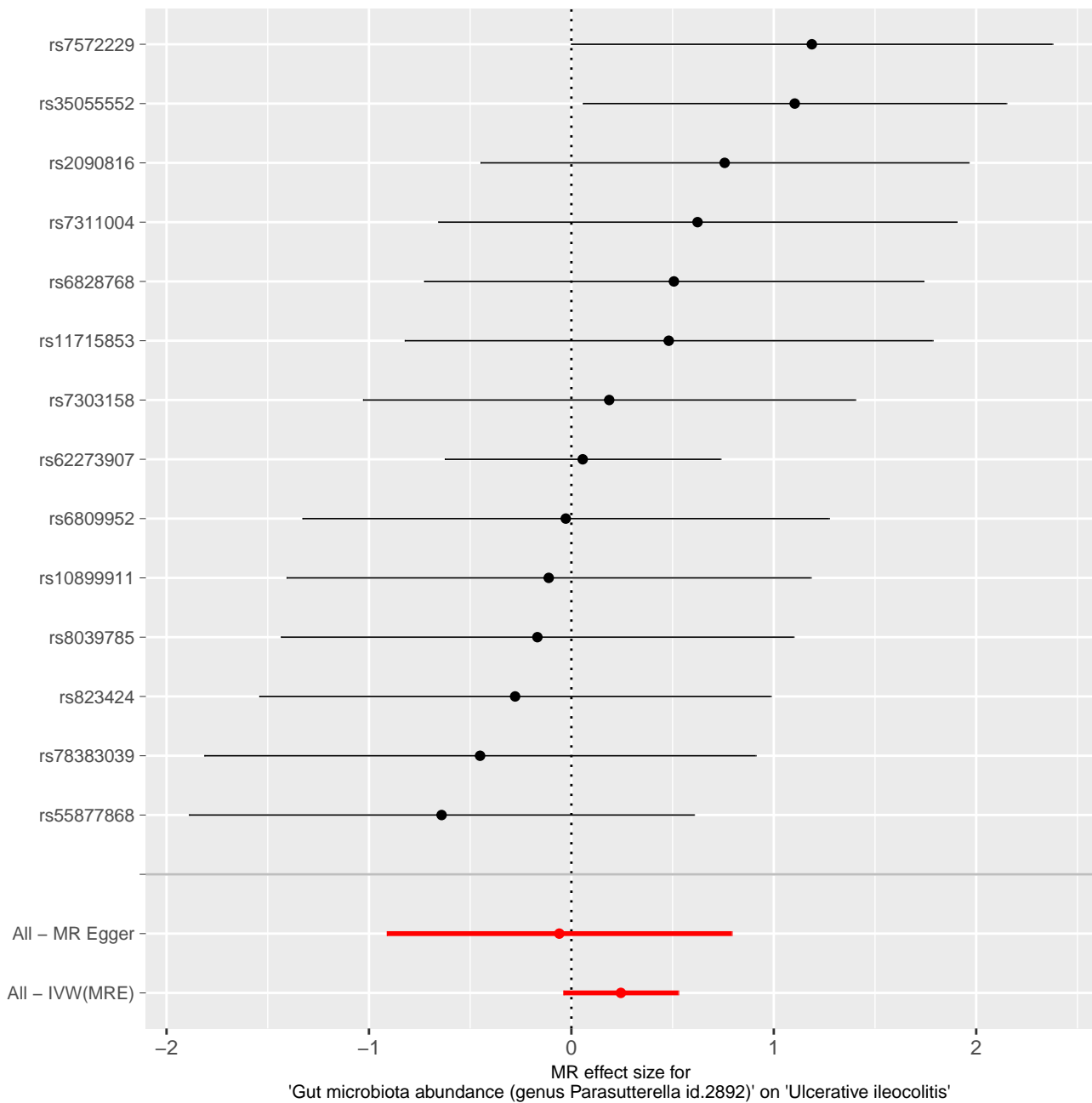


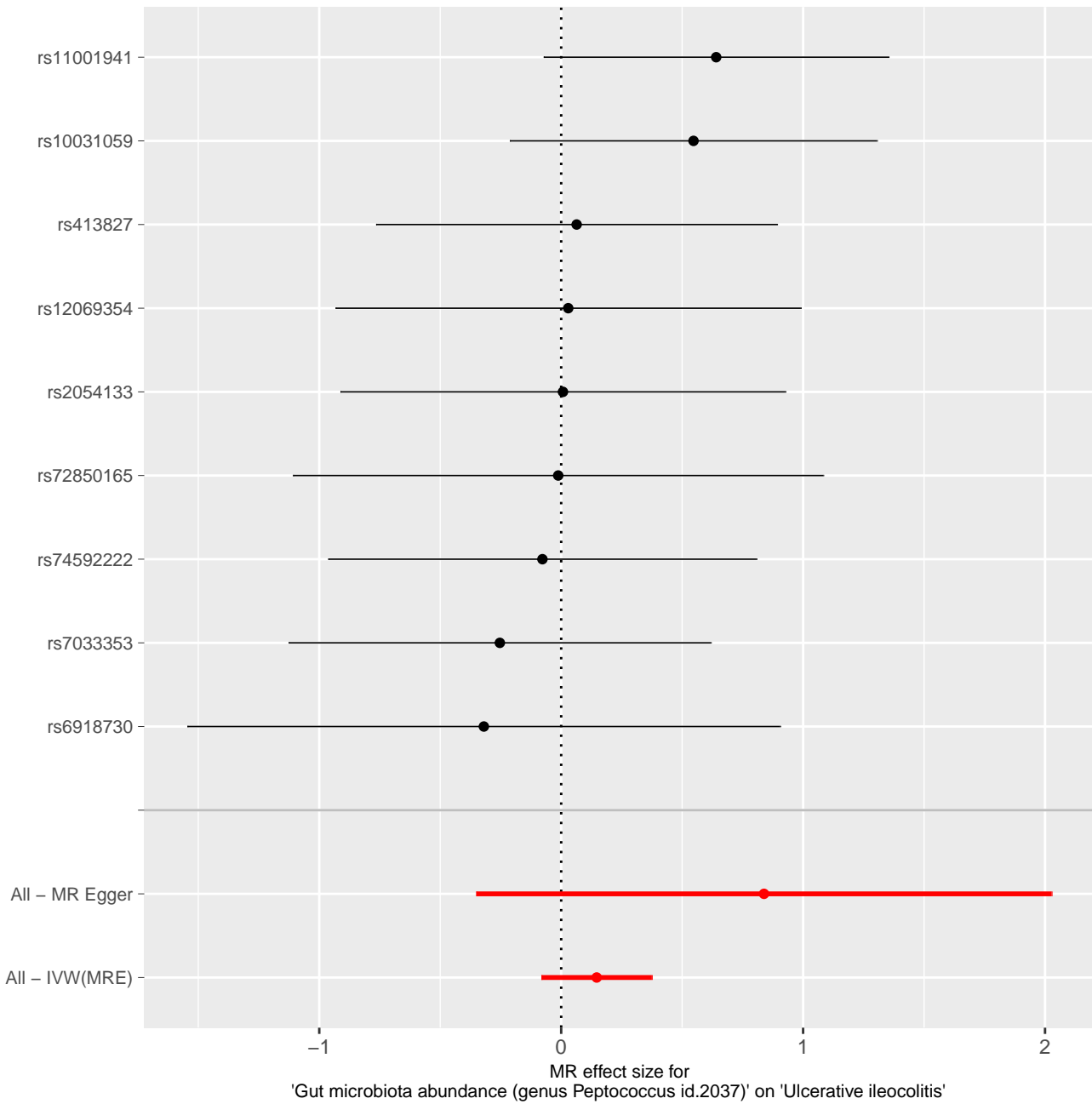


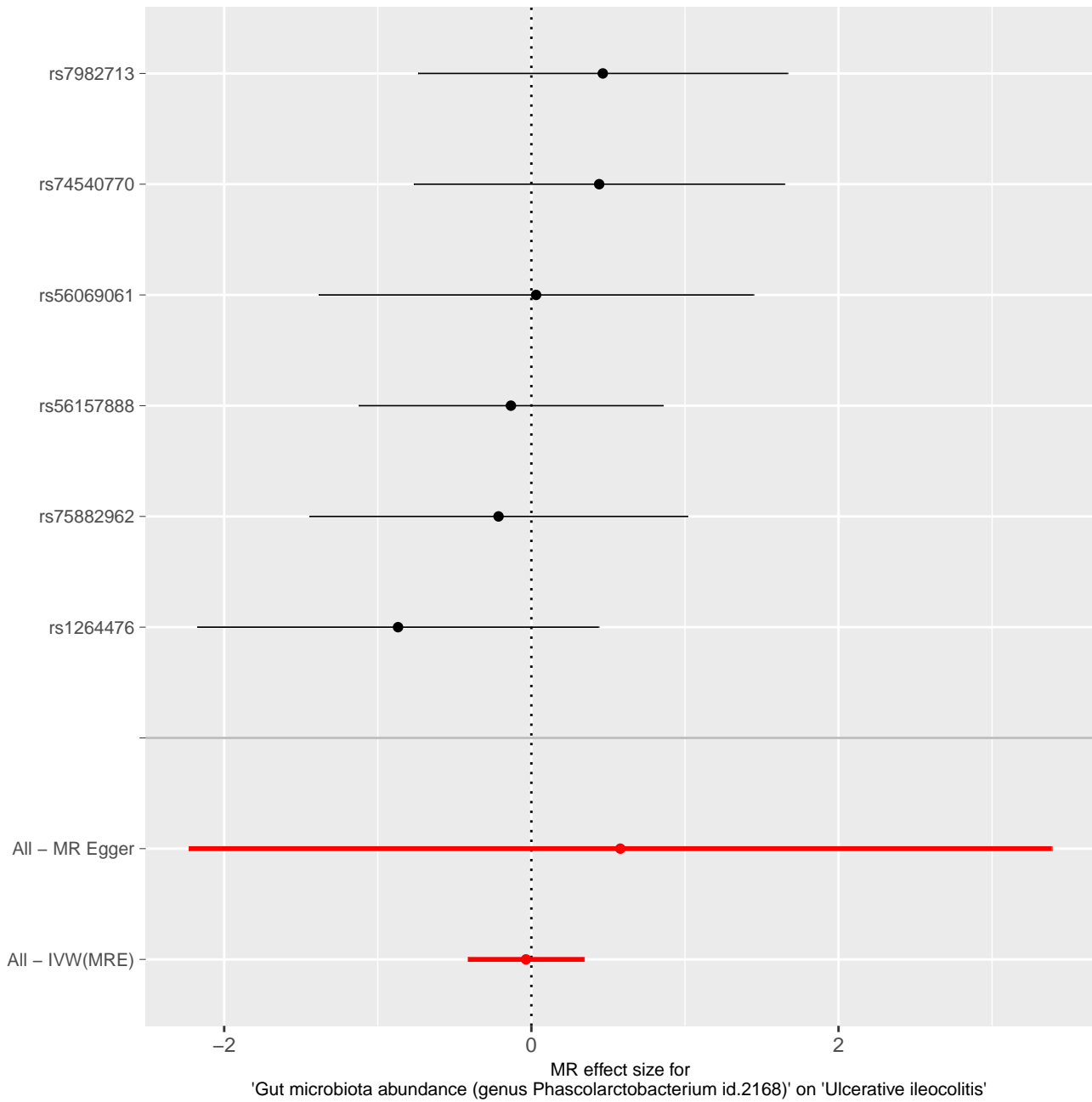


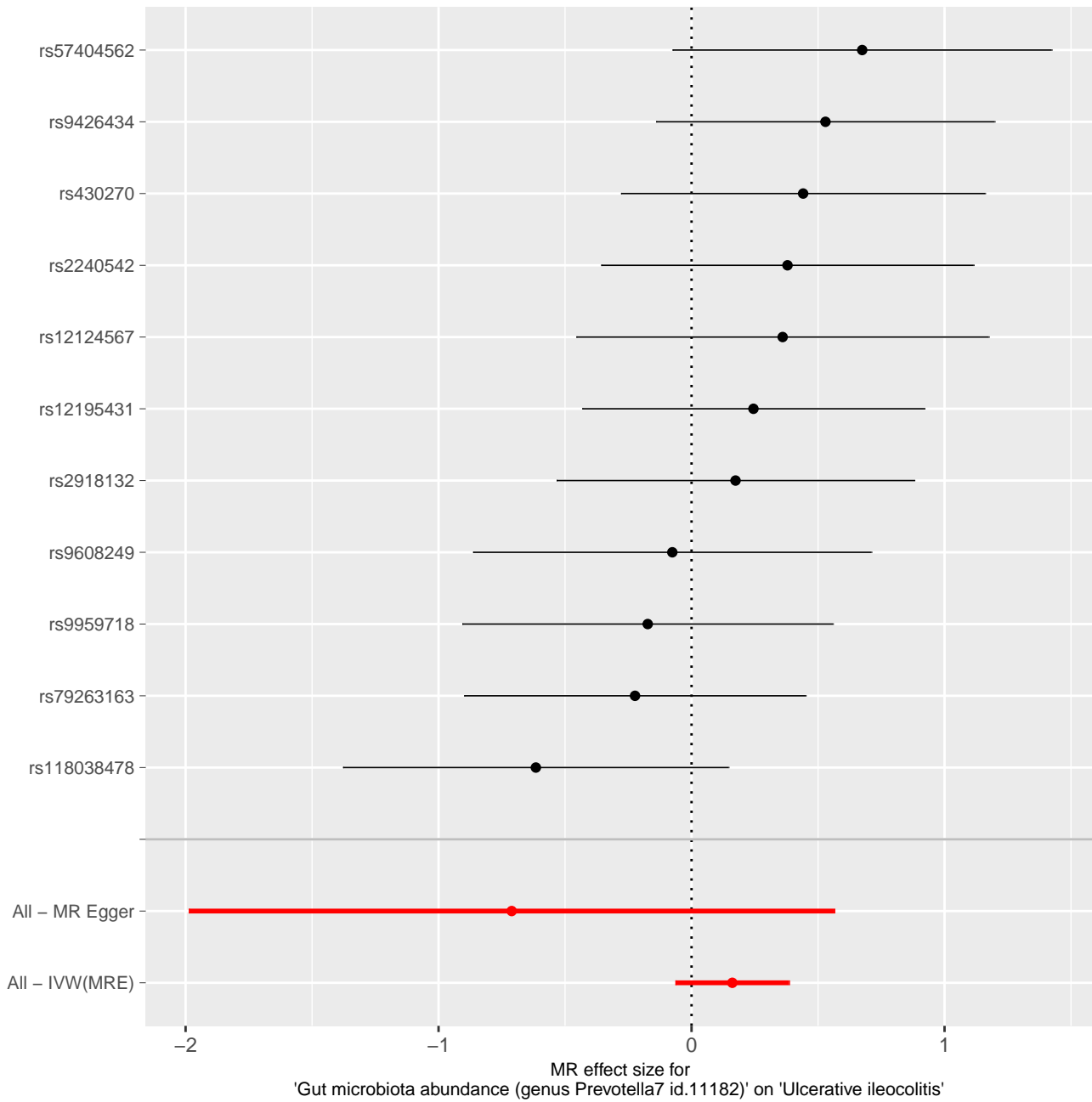




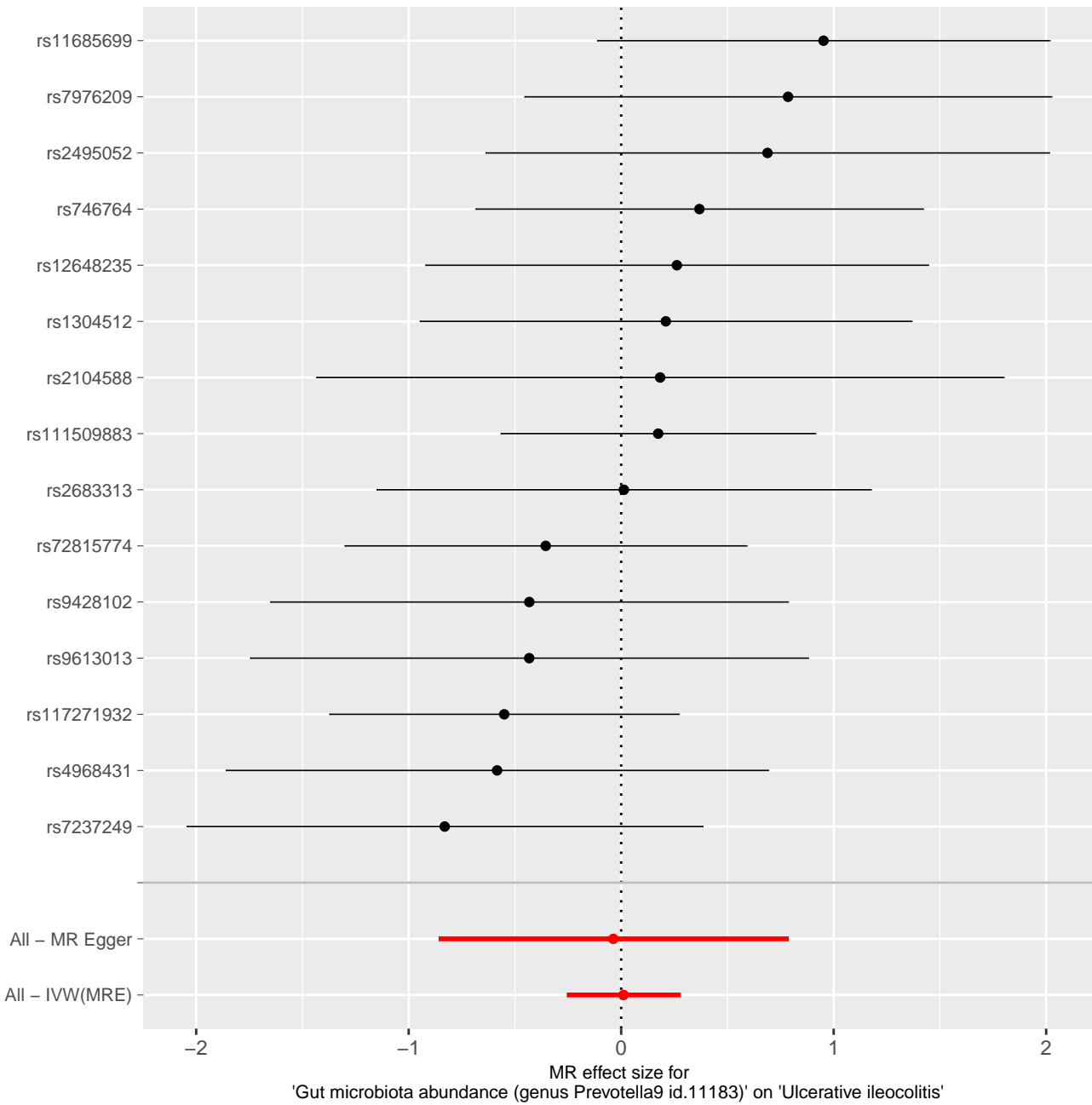


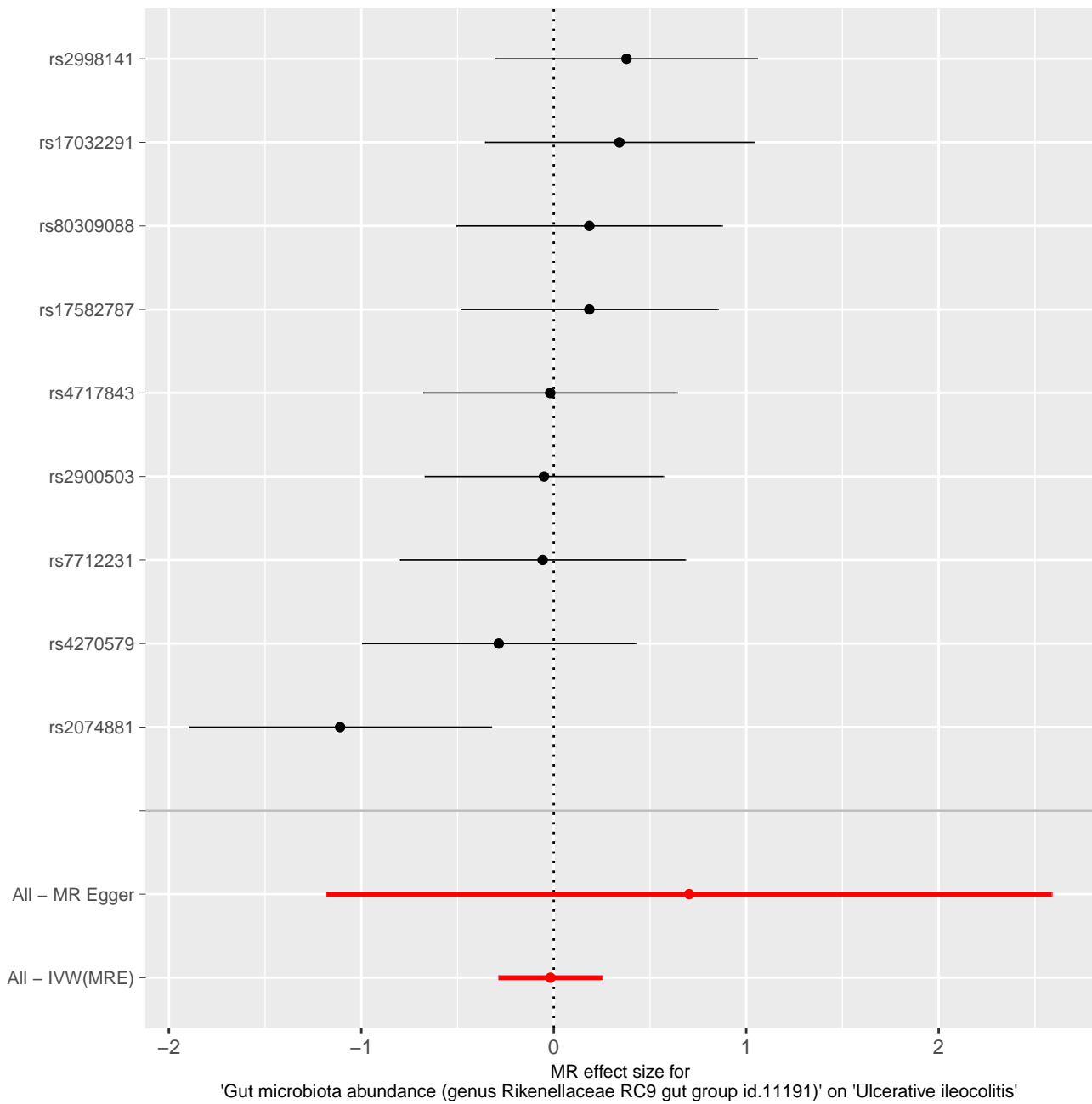


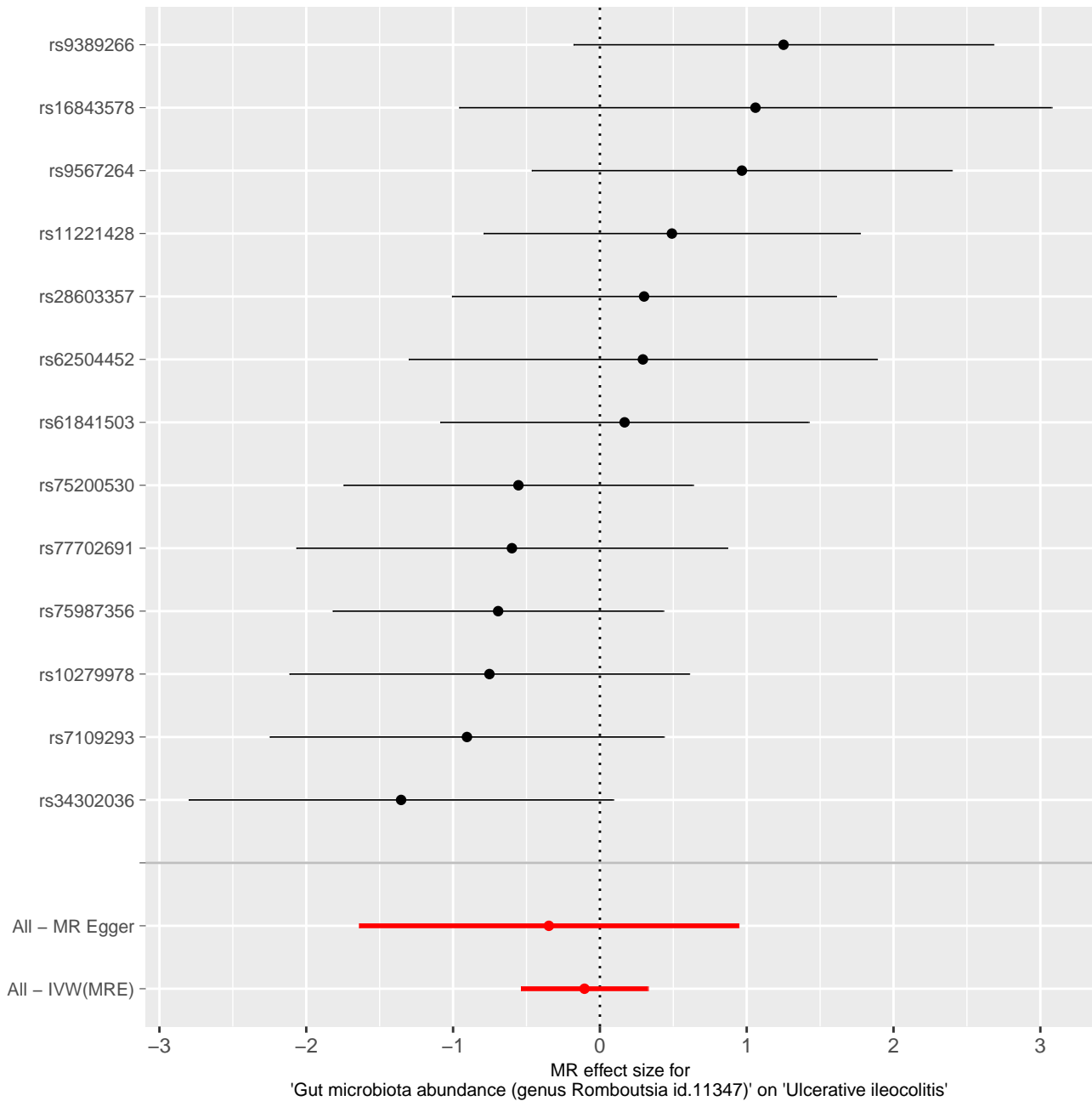


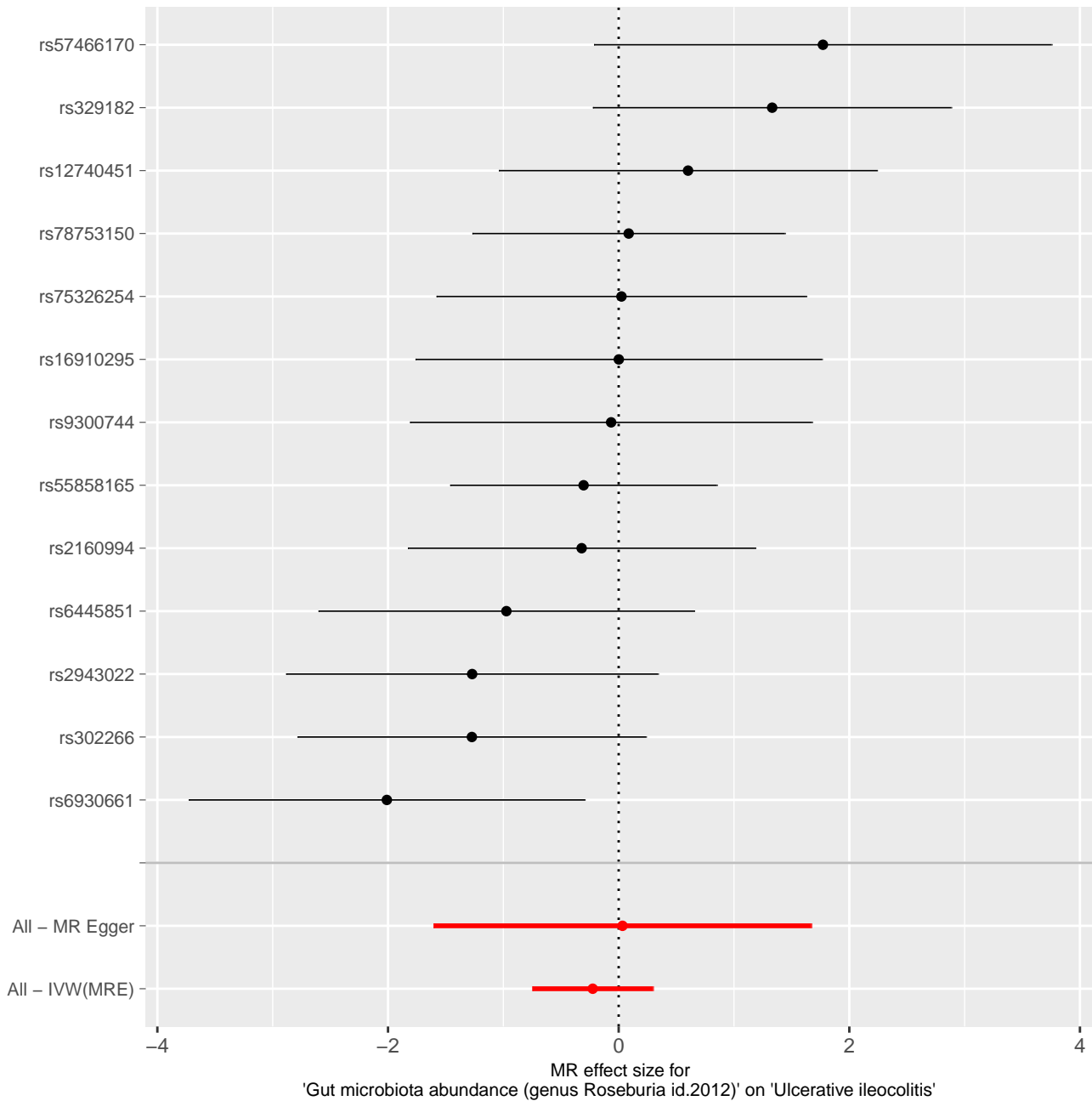


Batch 1169 : Gut microbiota abundance (genus Prevotella9 id.11183) on Ulcerative ileocolitis

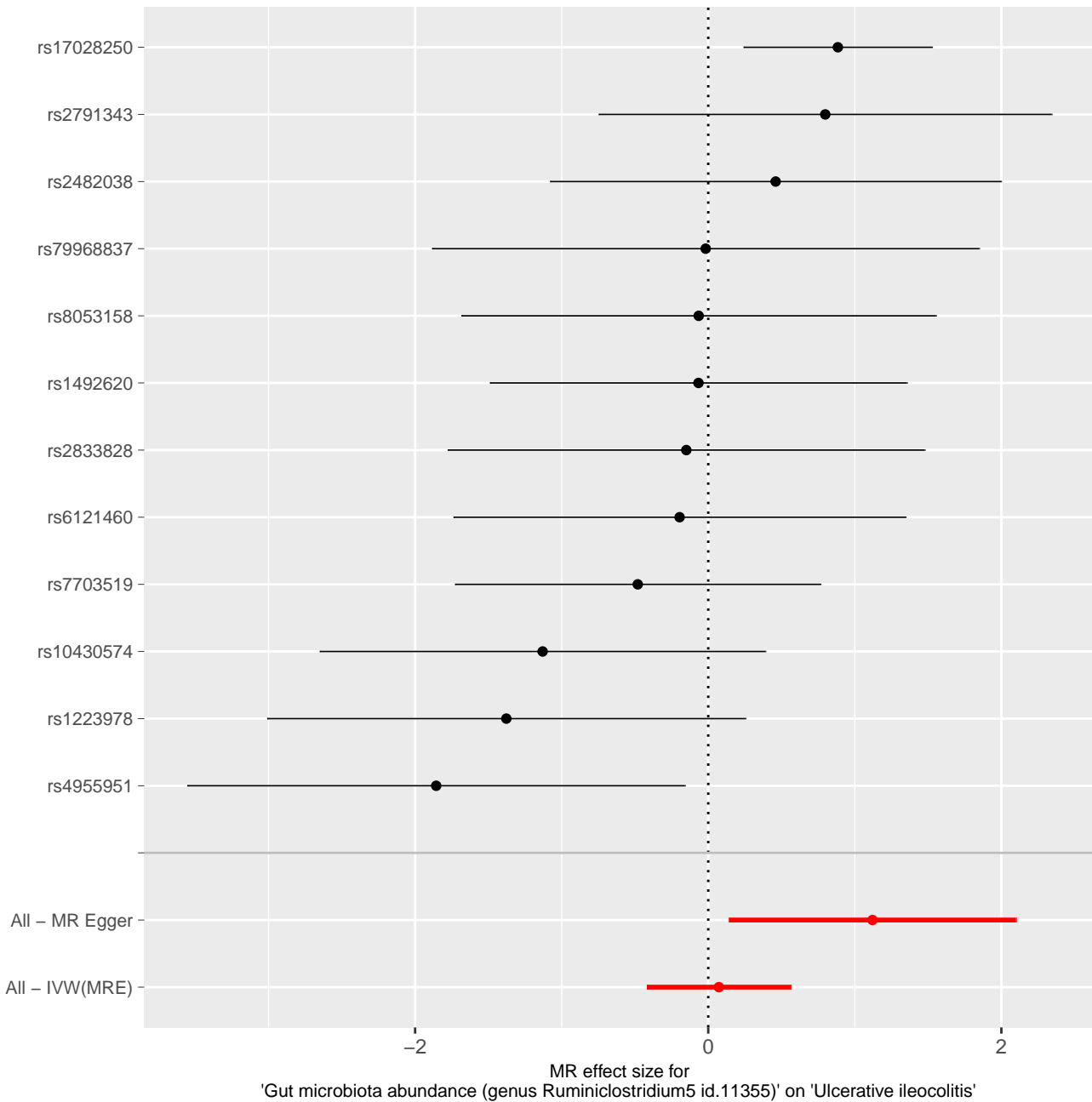


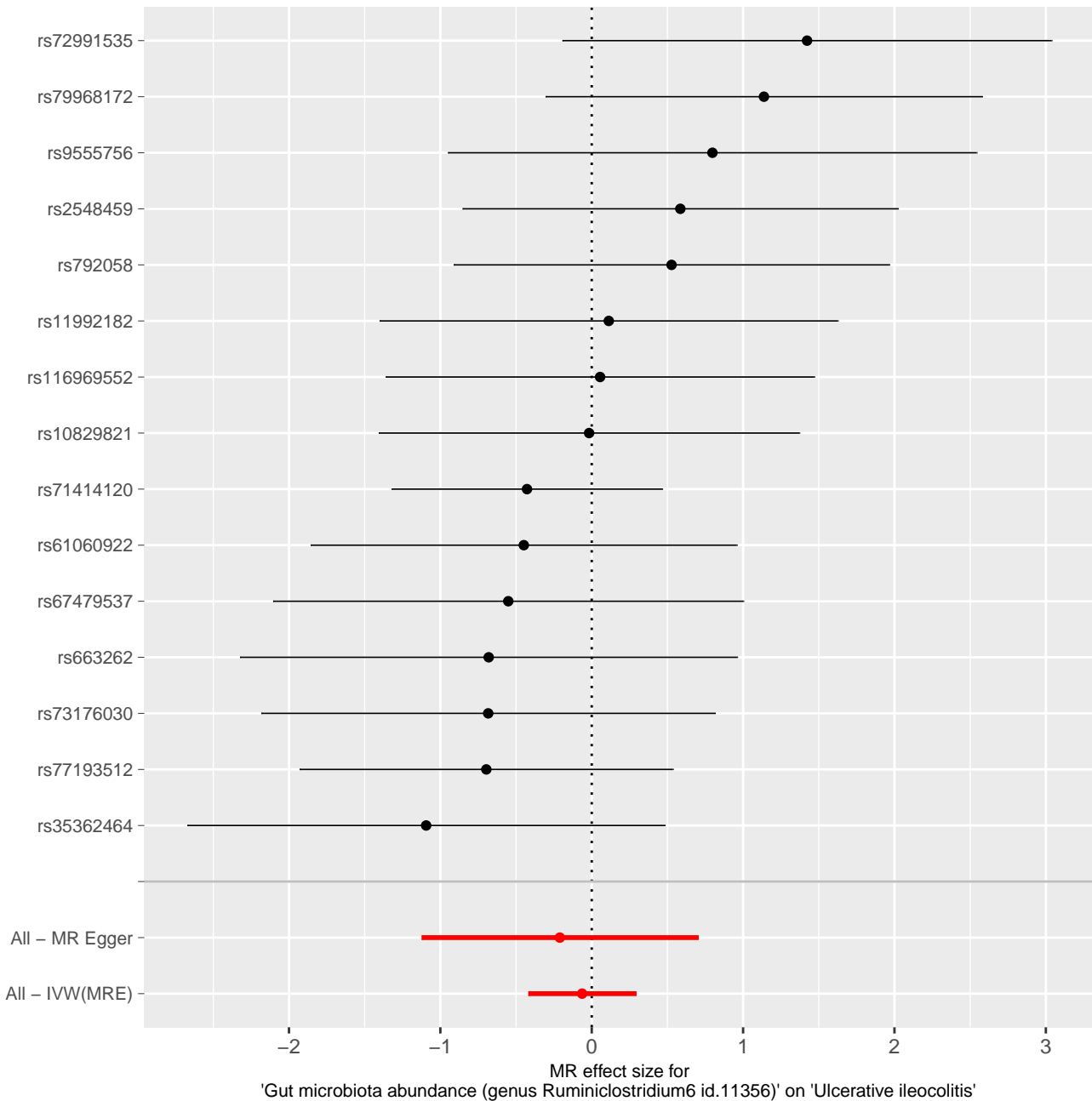


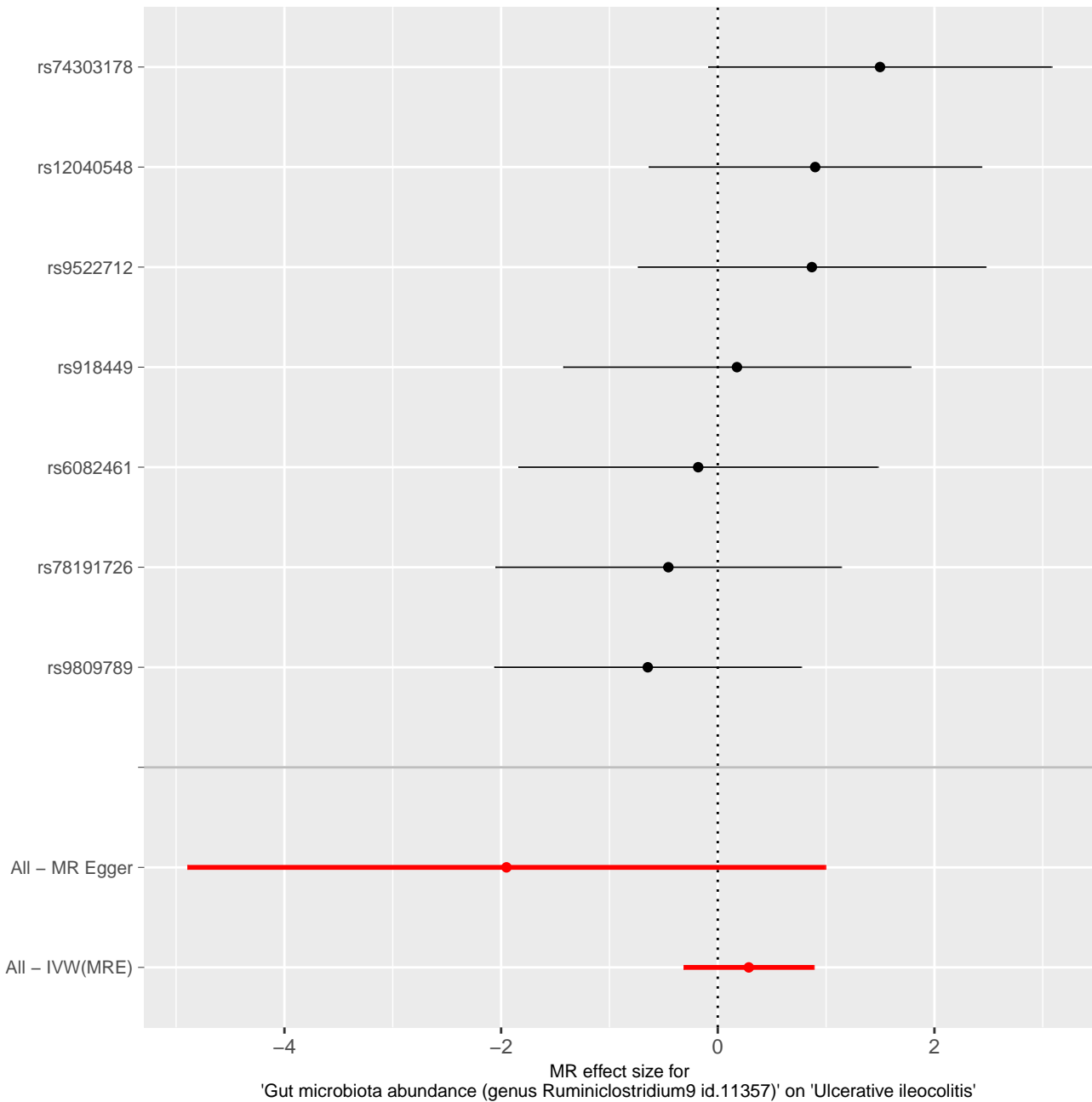


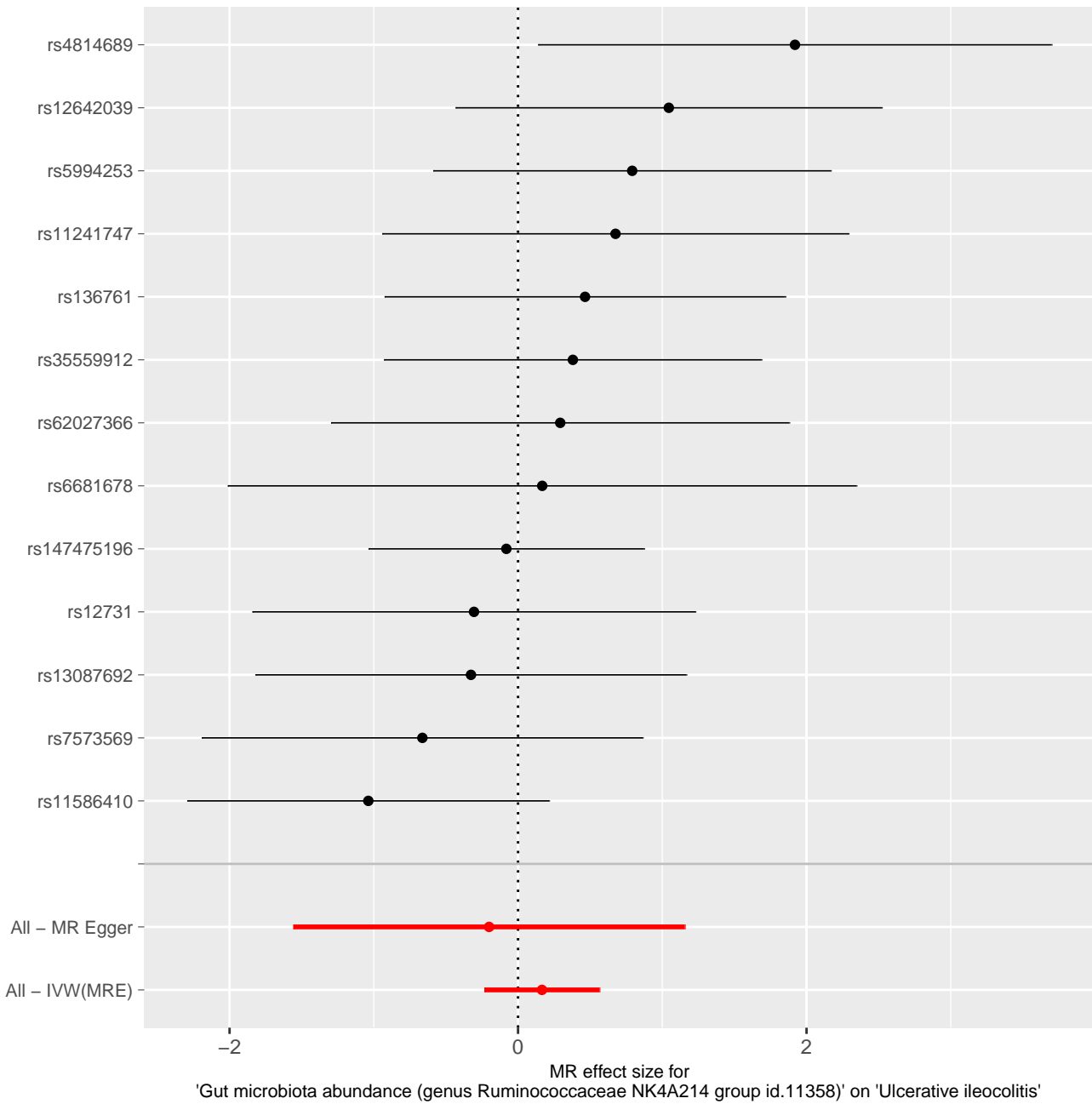


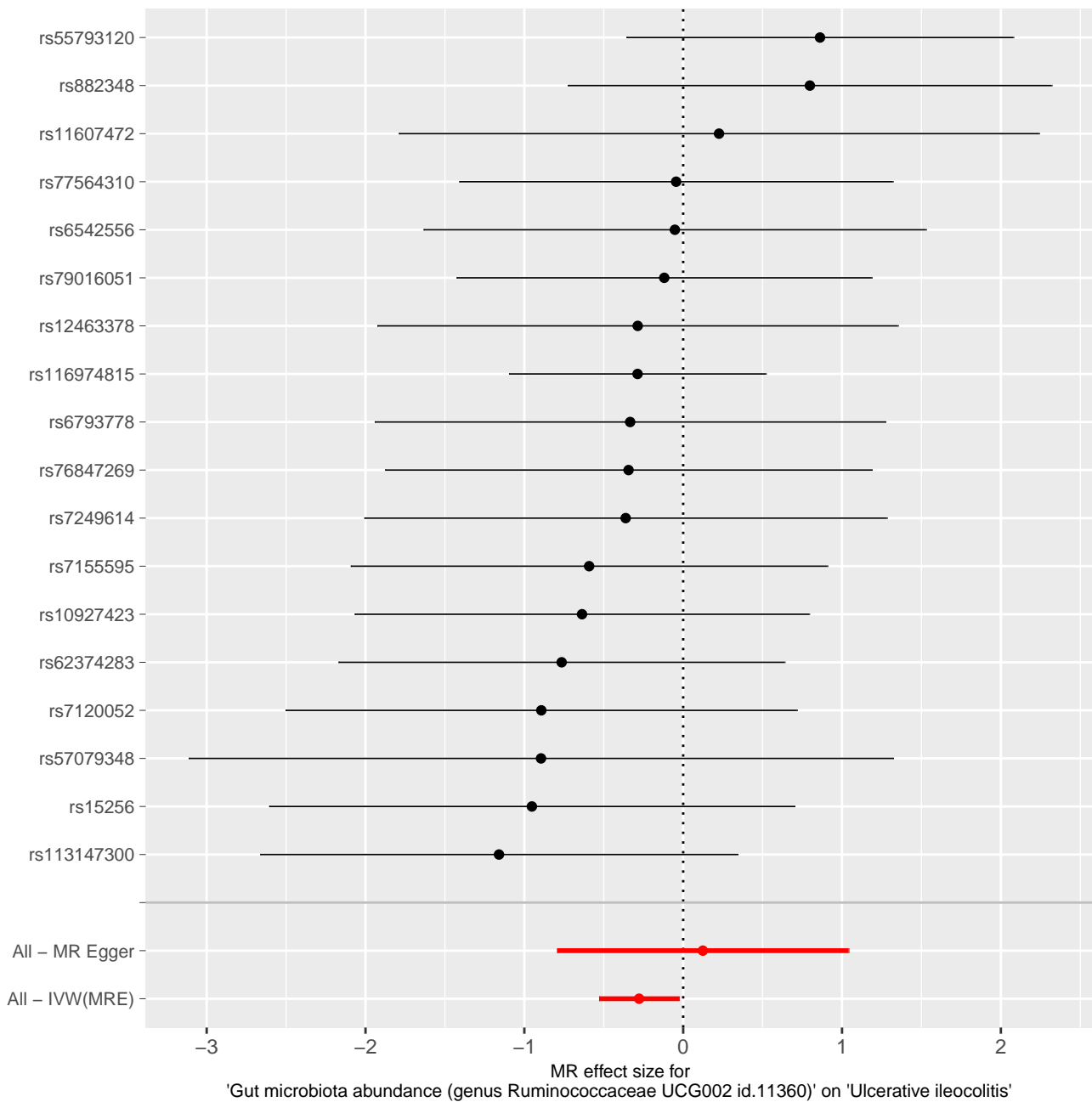
Batch 1173 : Gut microbiota abundance (genus Ruminiclostridium5 id.11355) on Ulcerative ileocolitis

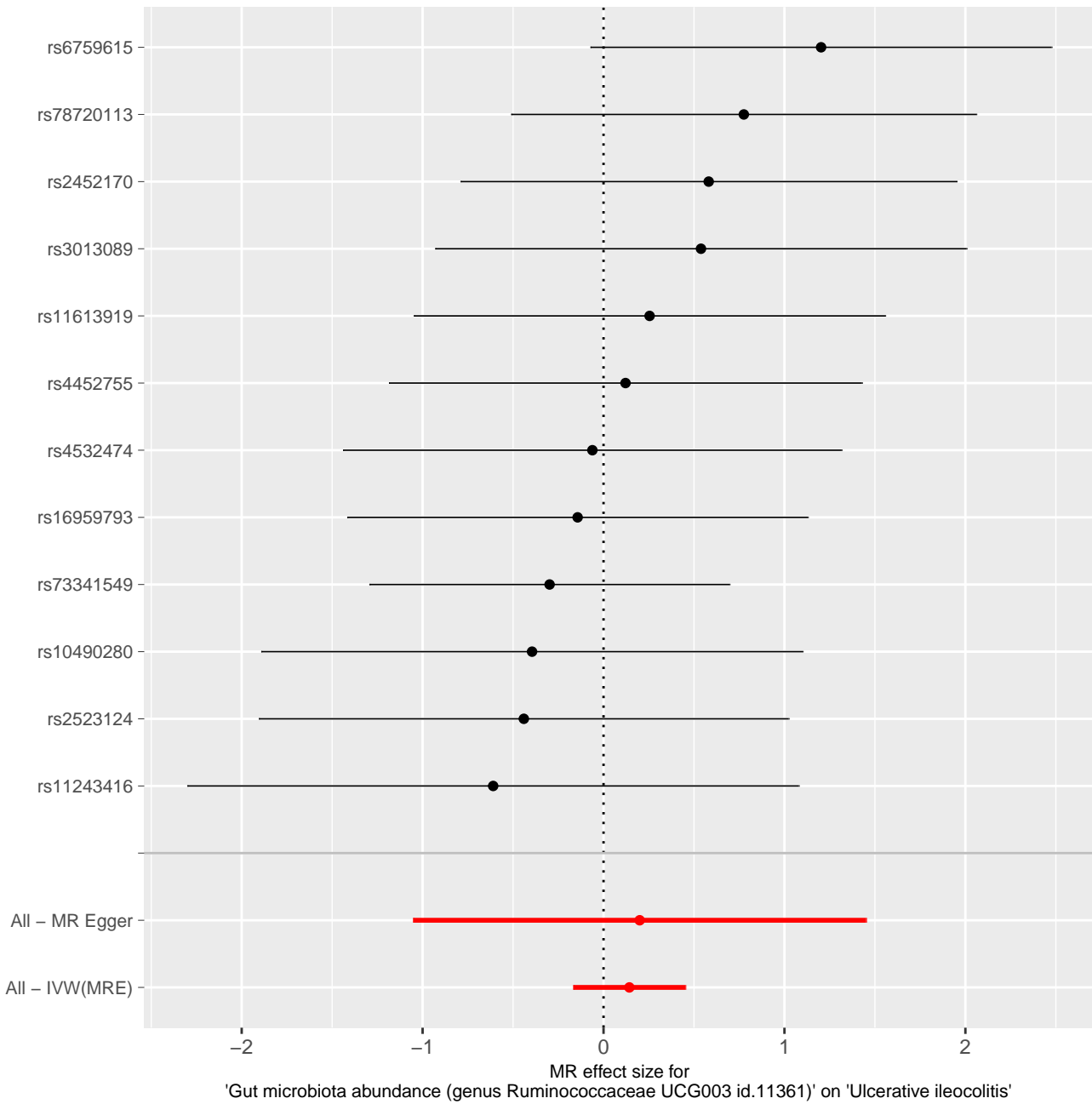


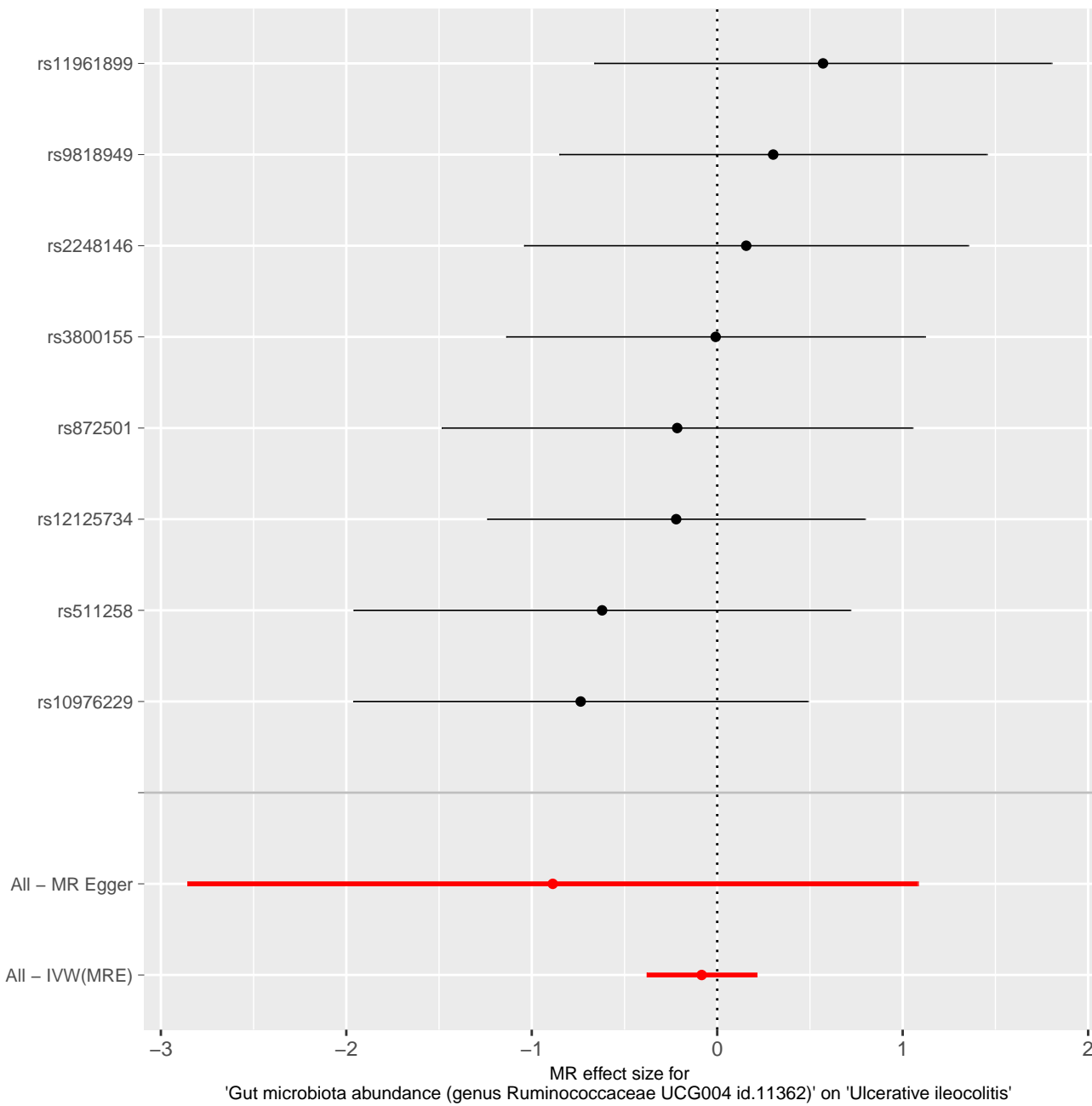


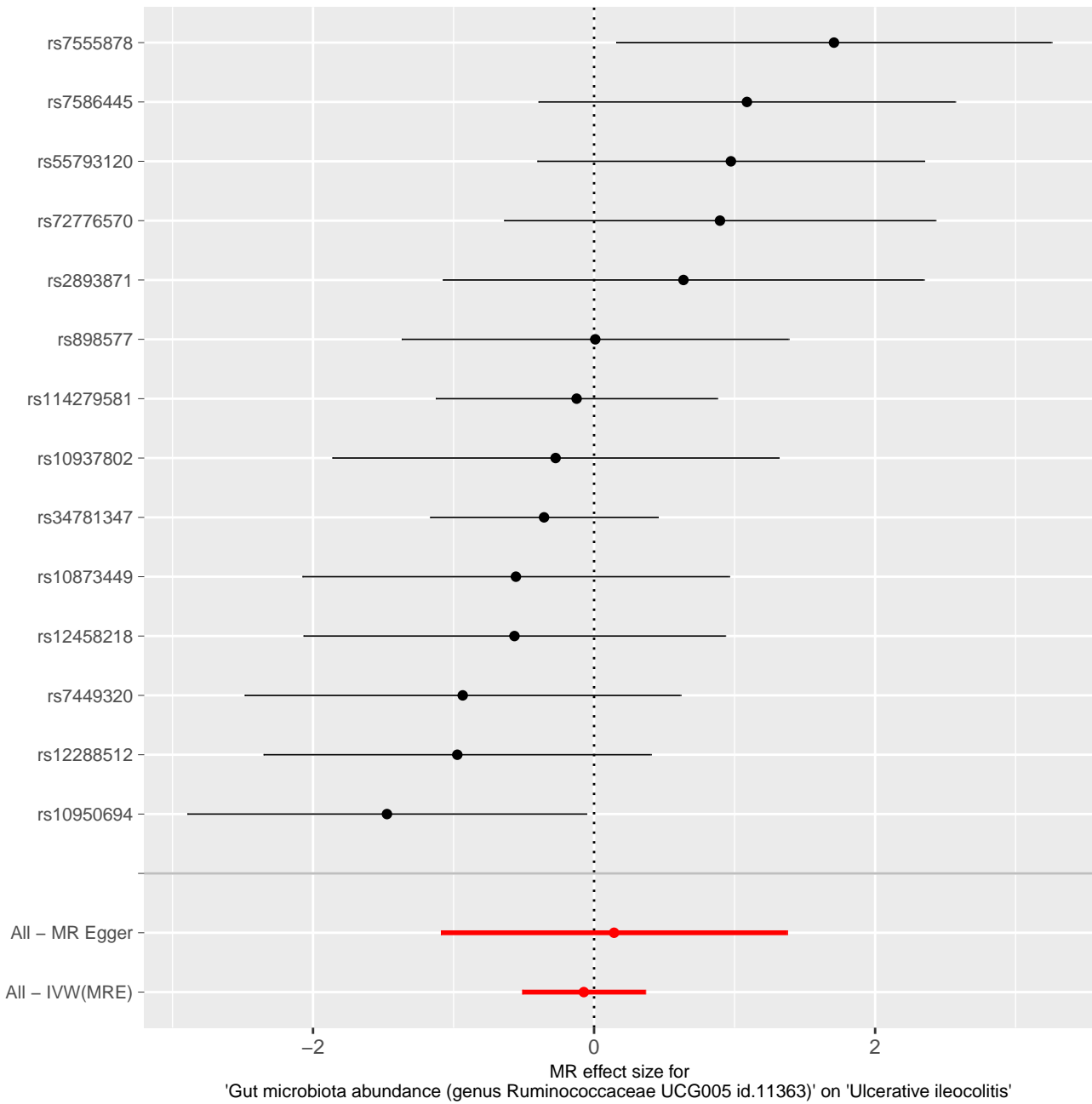


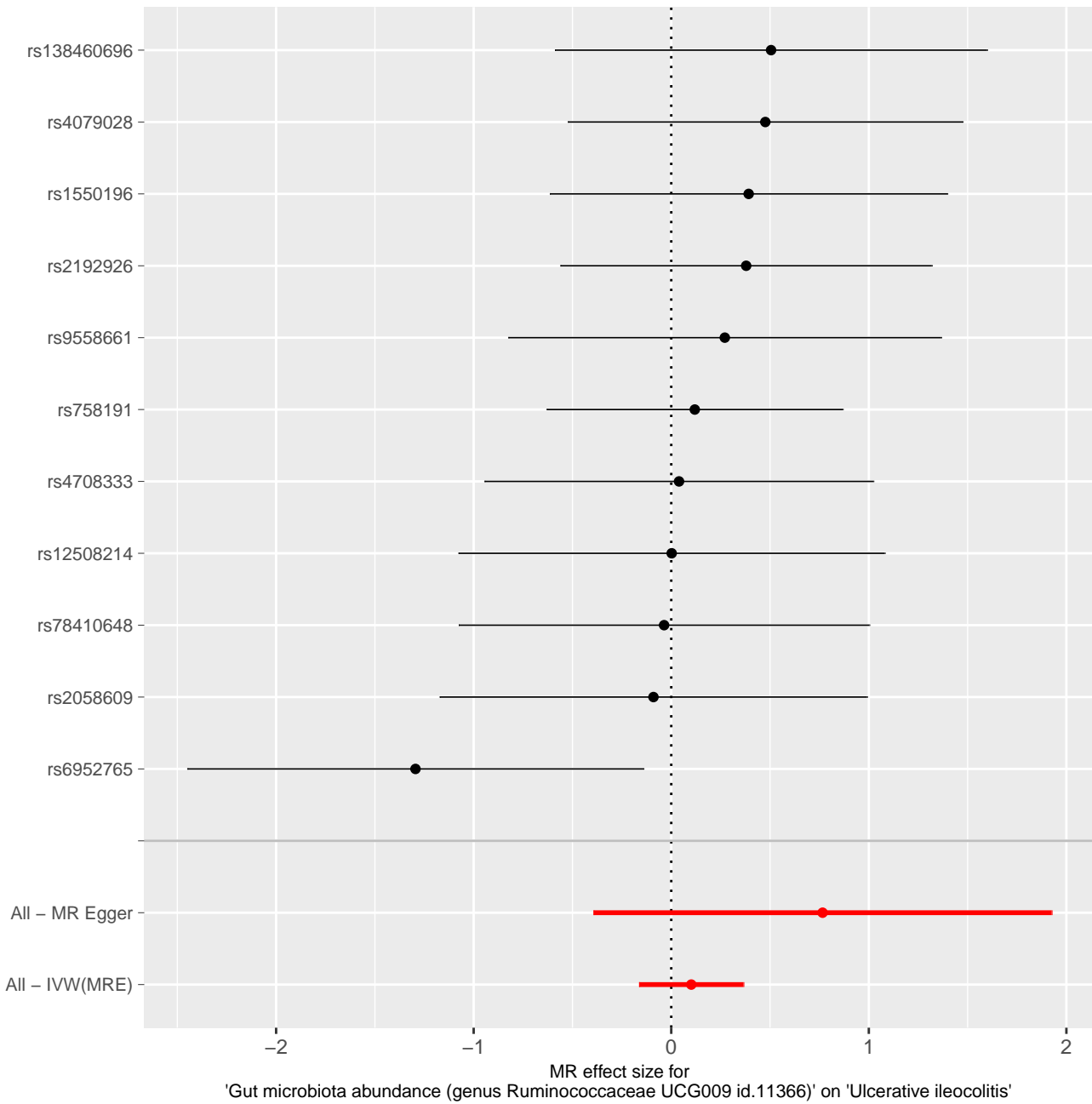


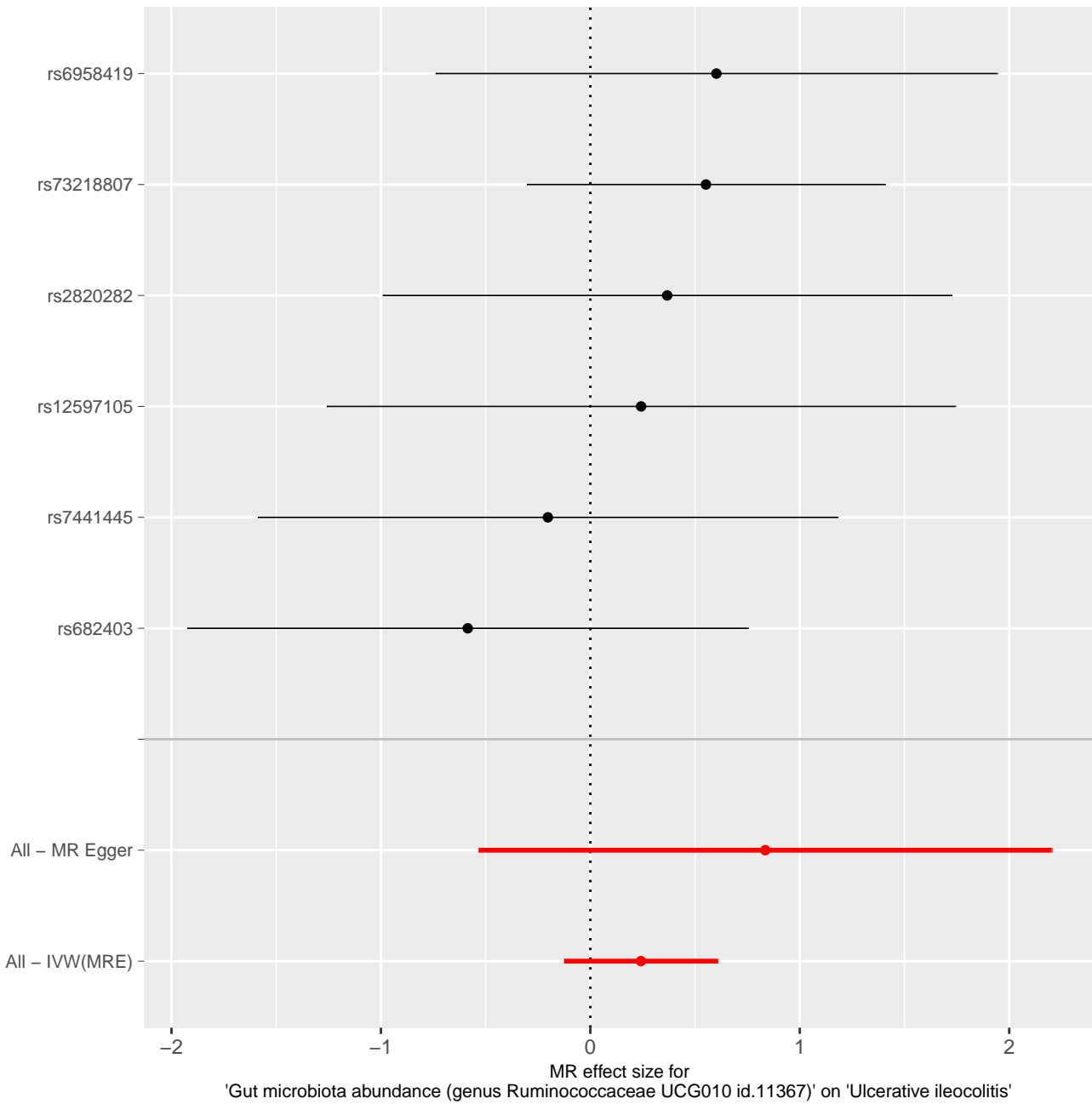


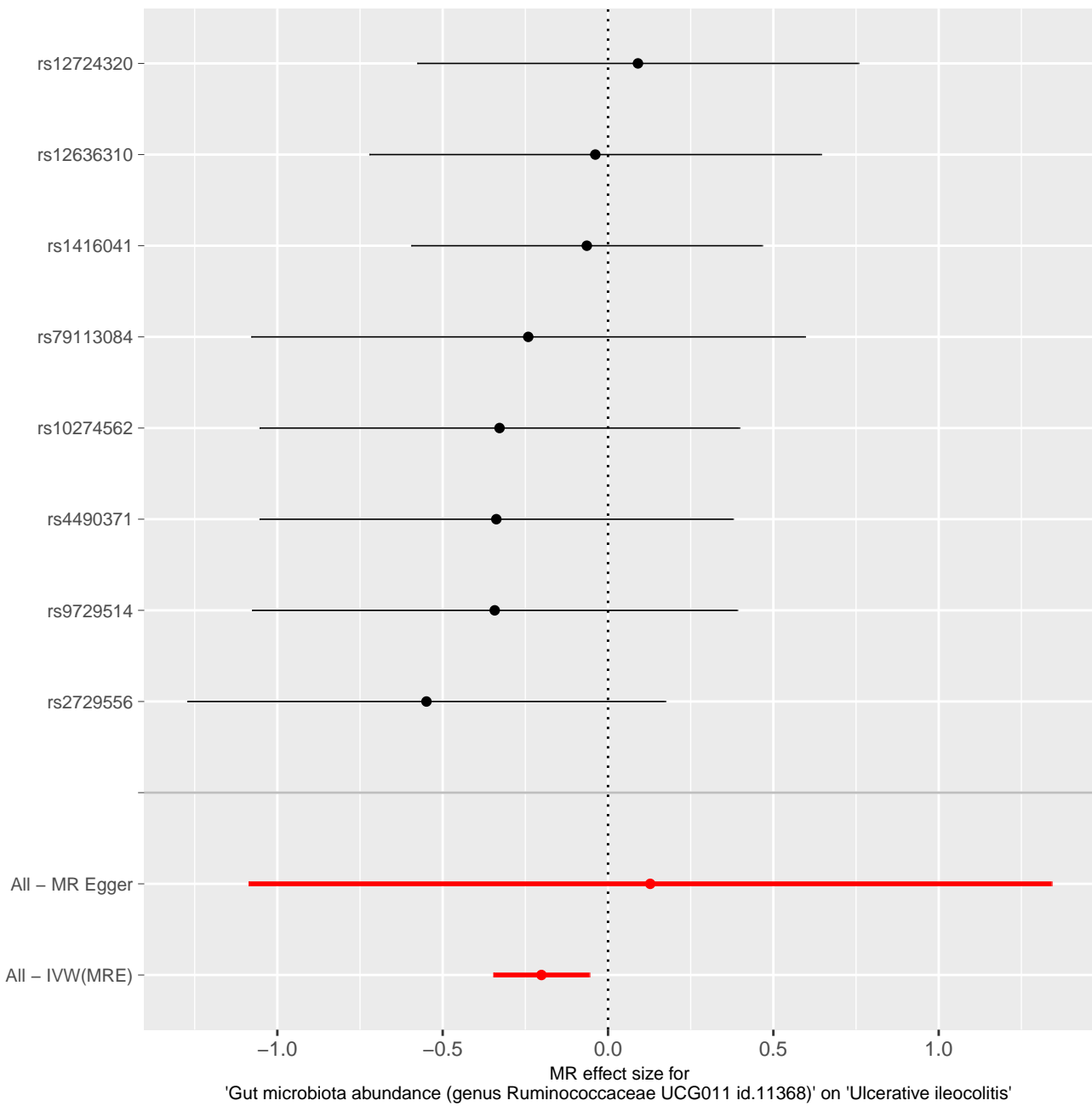


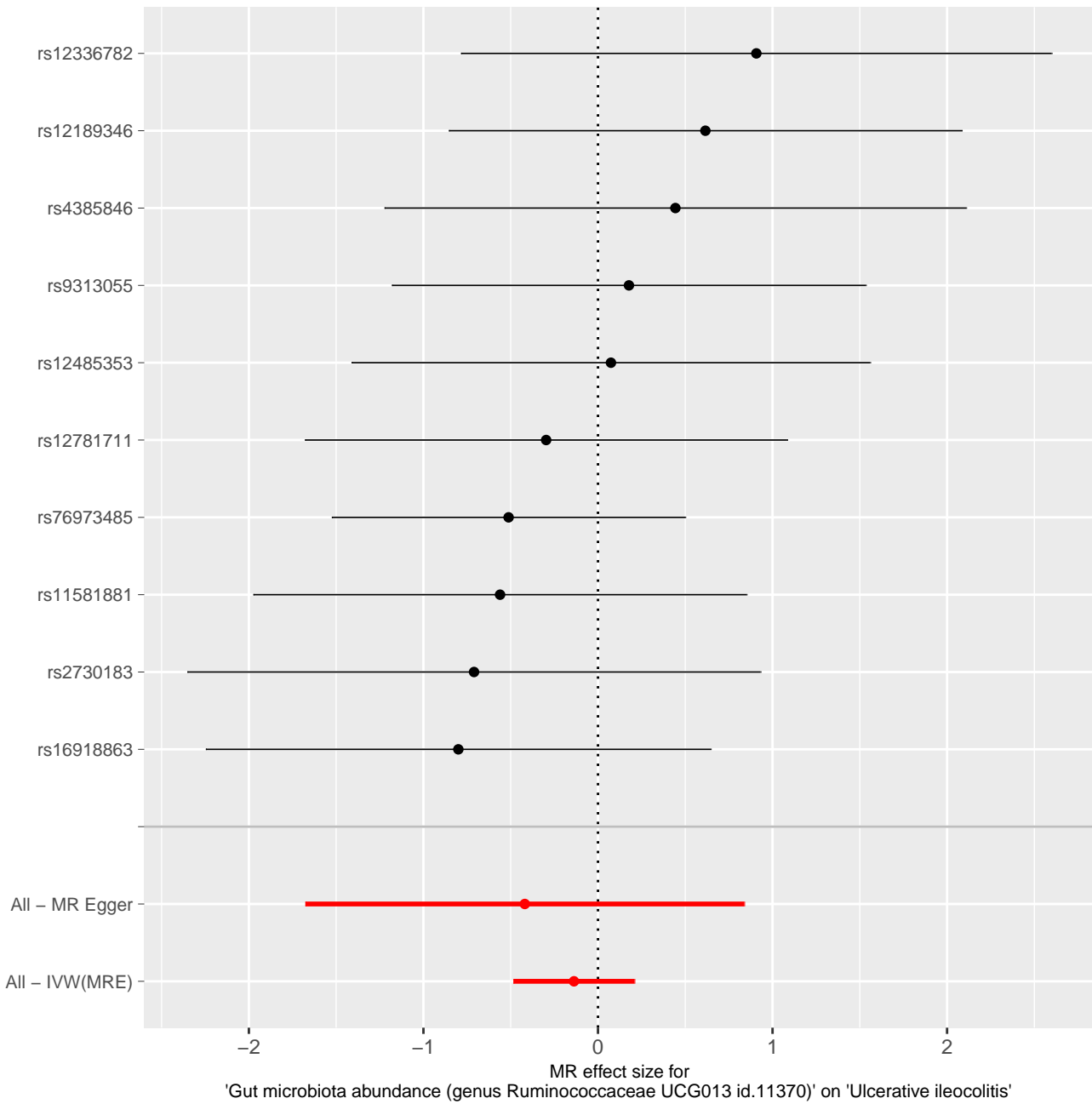


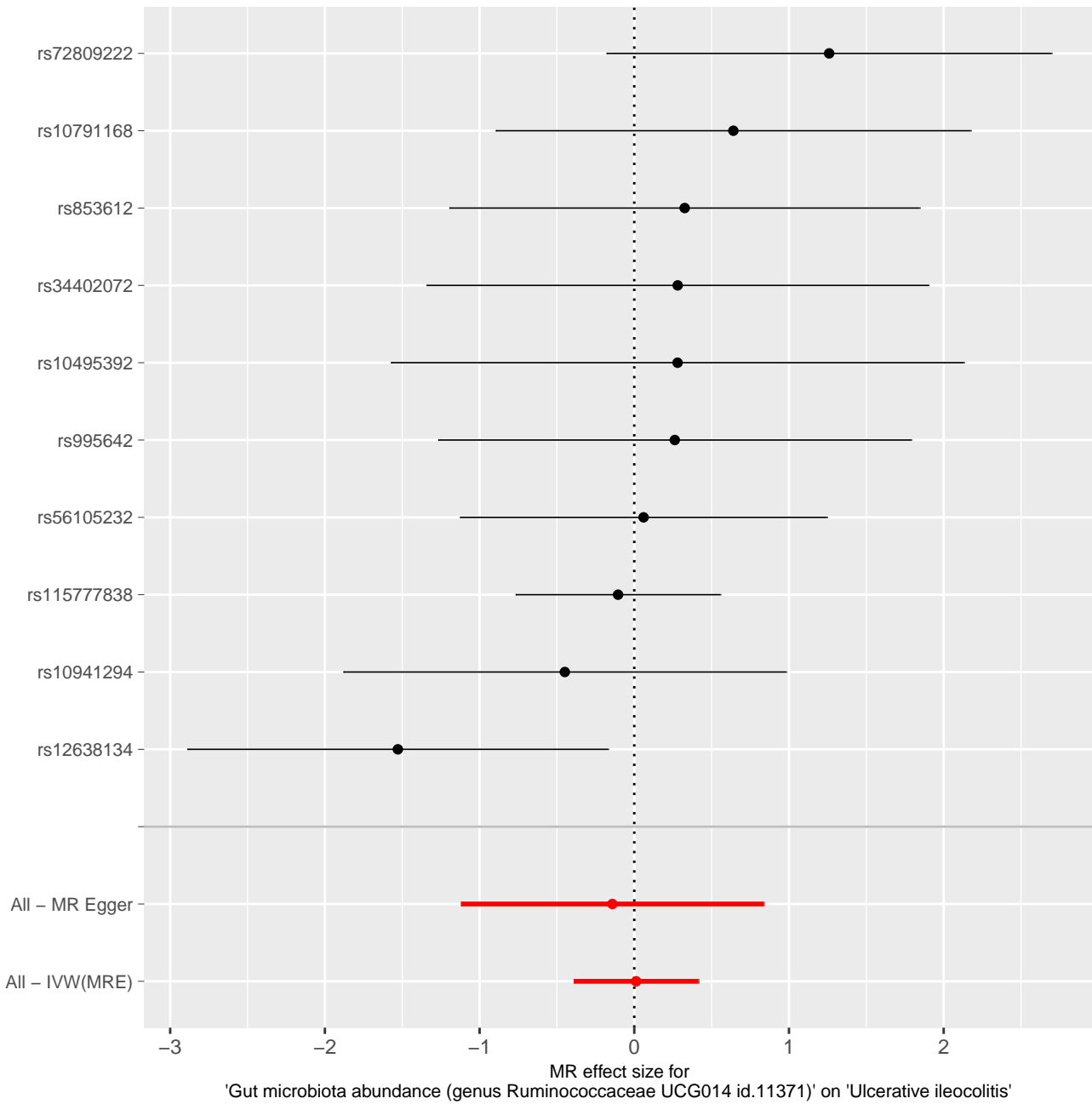


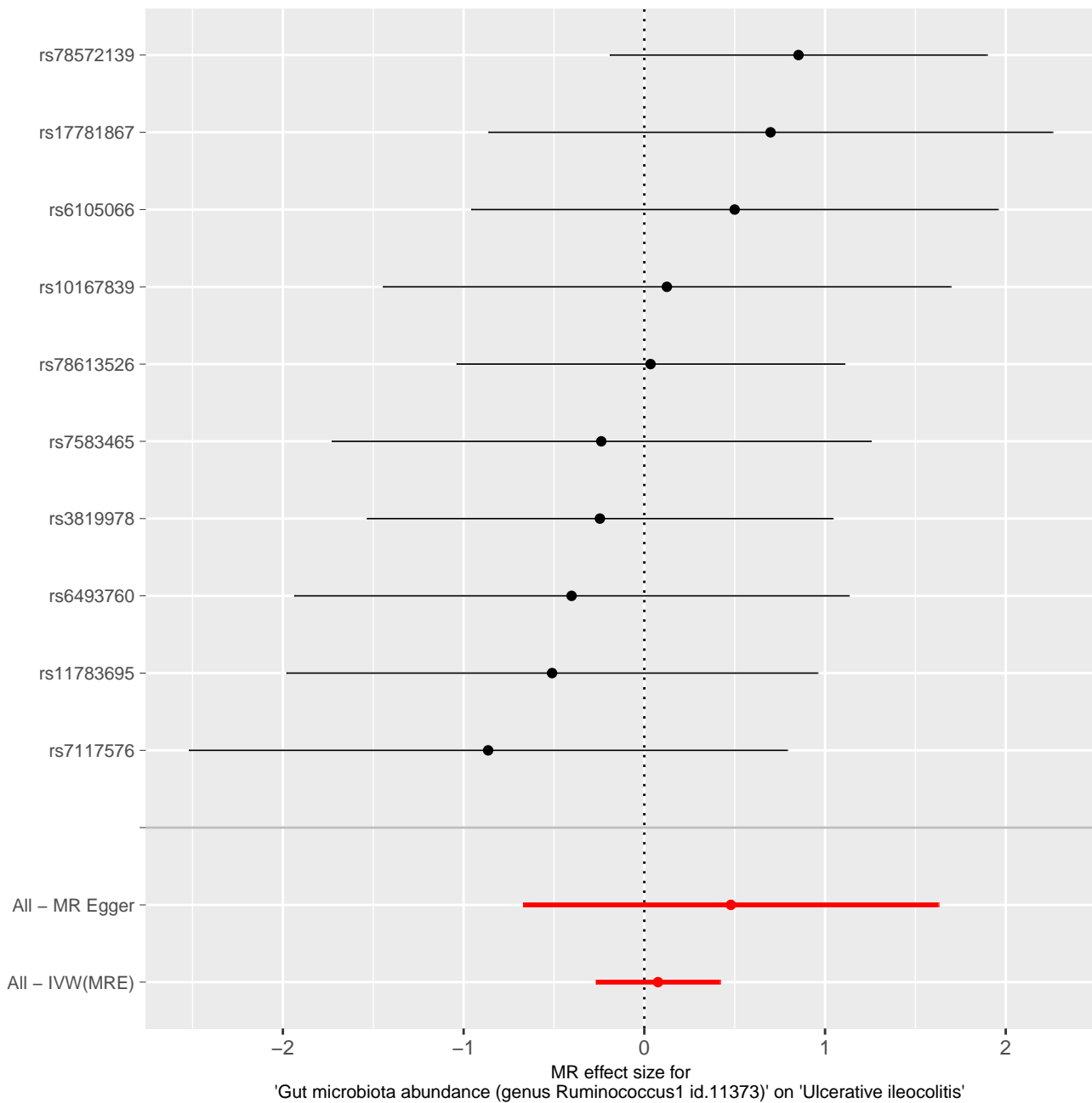


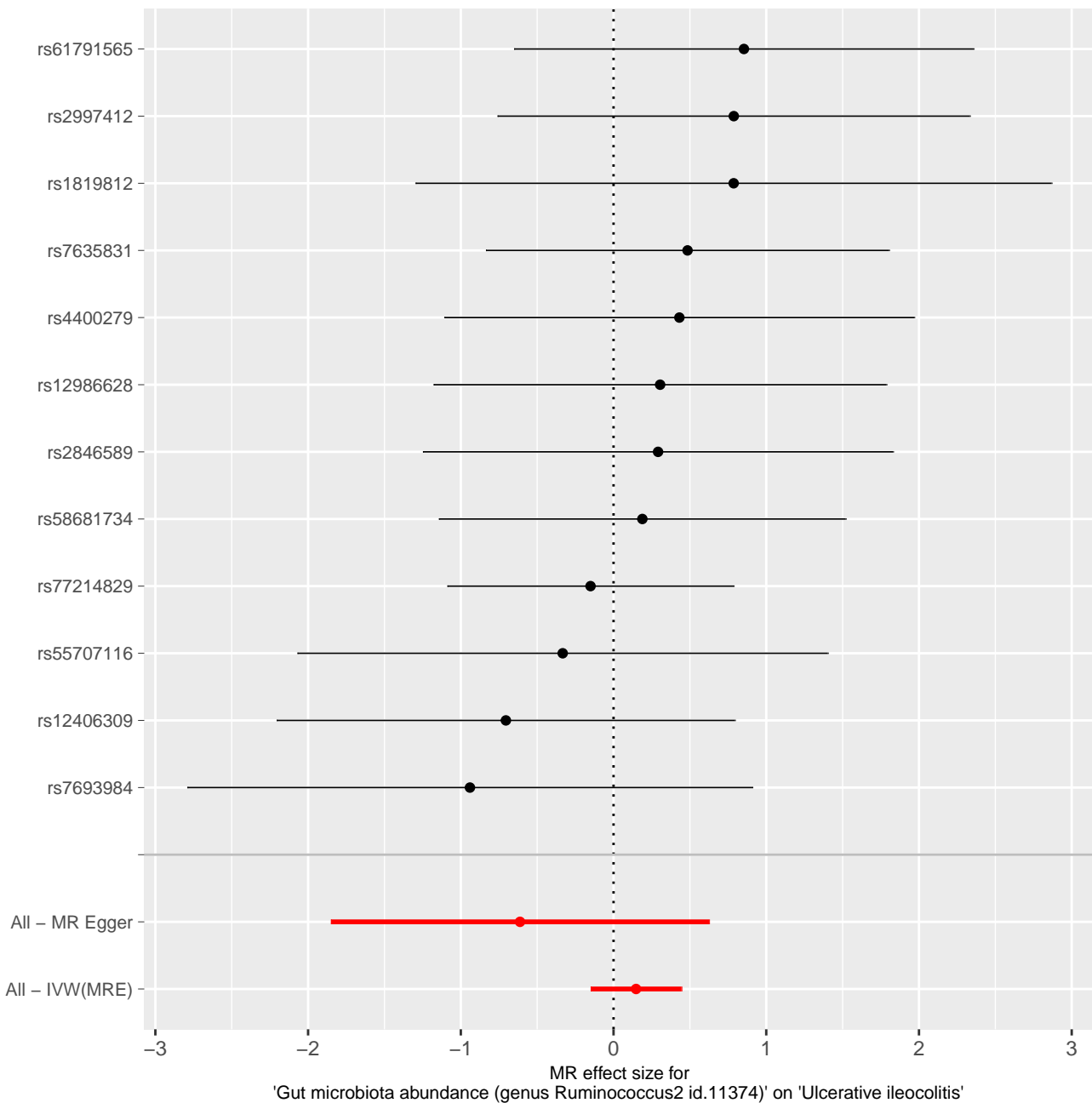


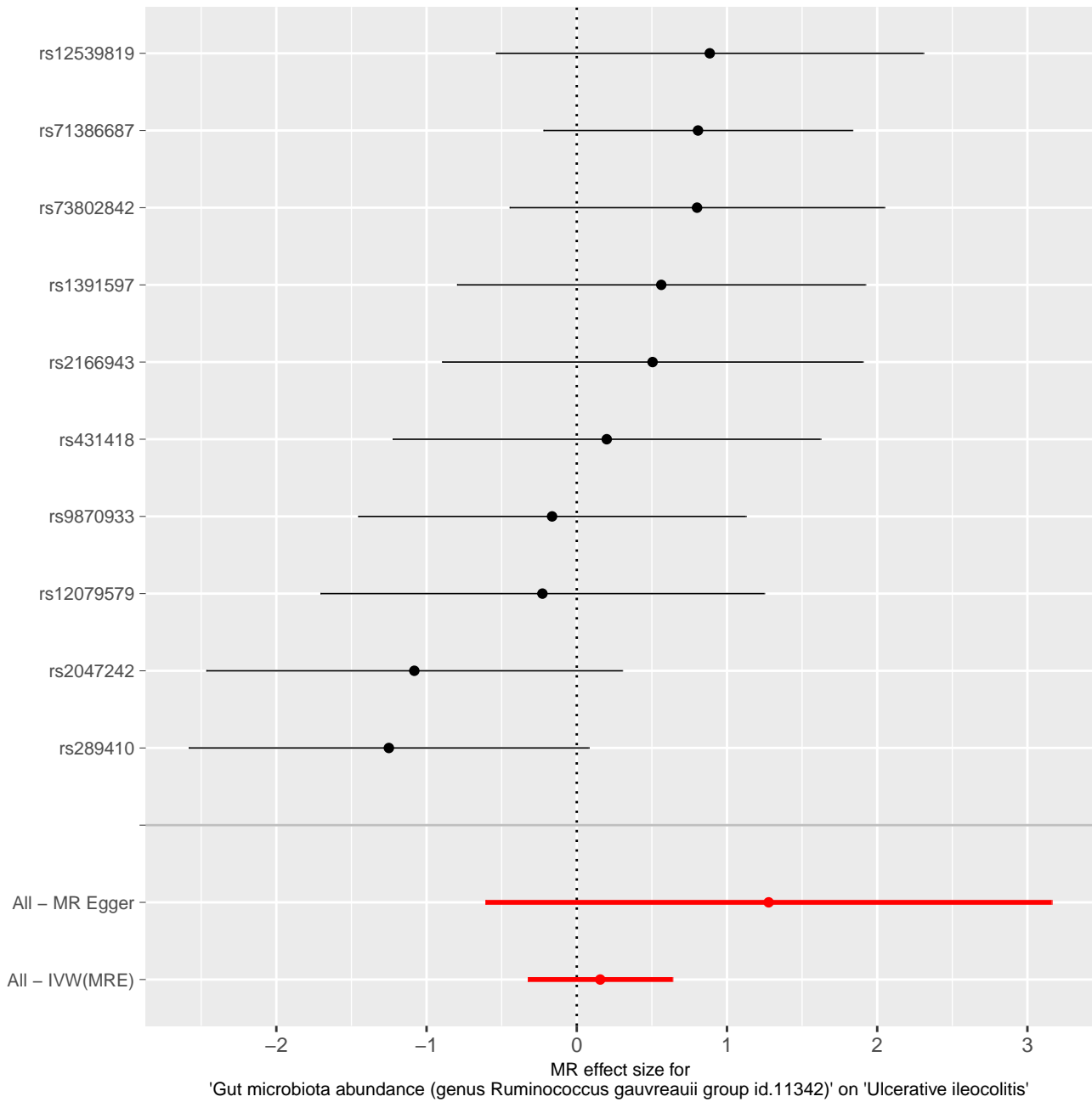


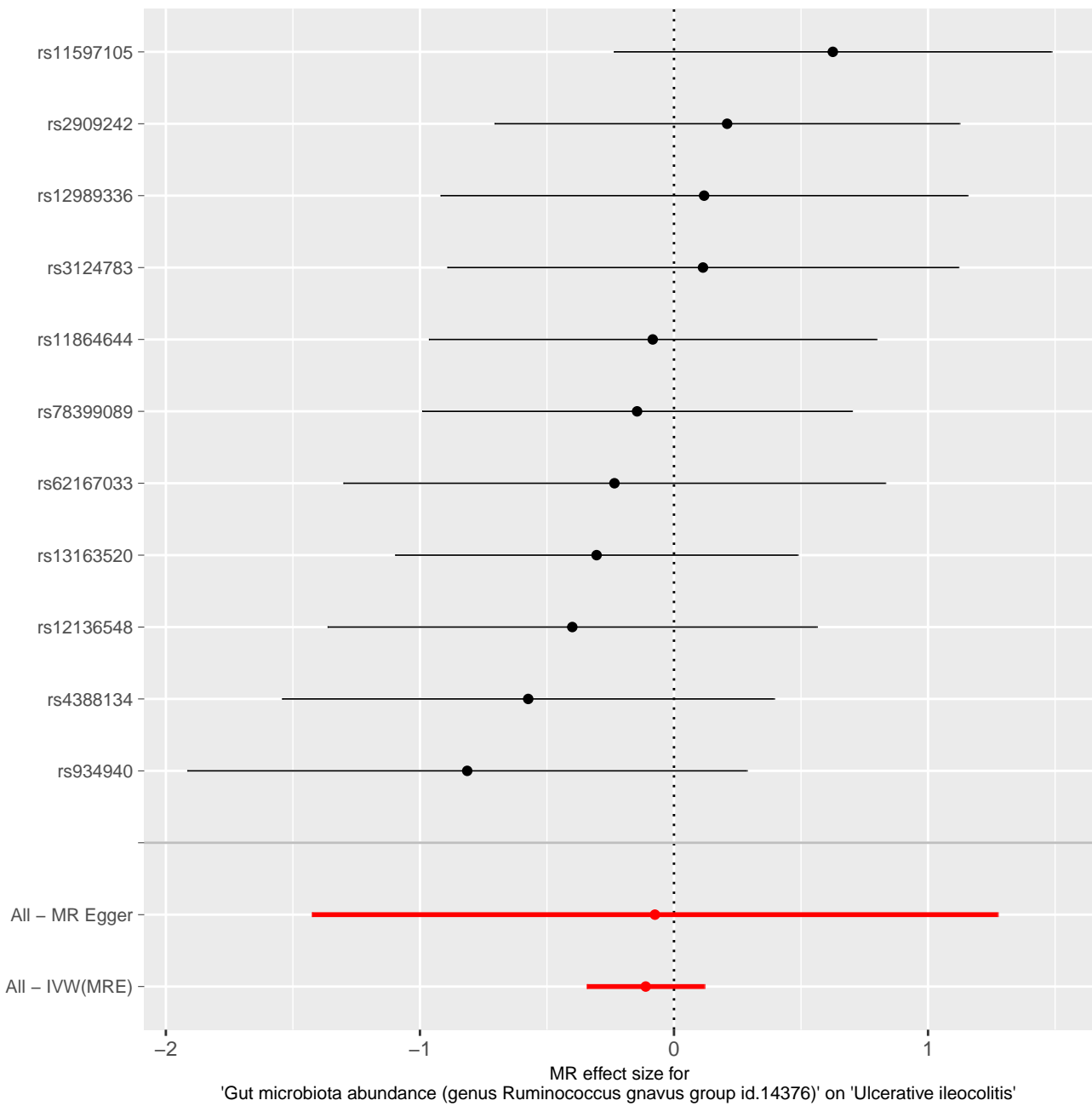


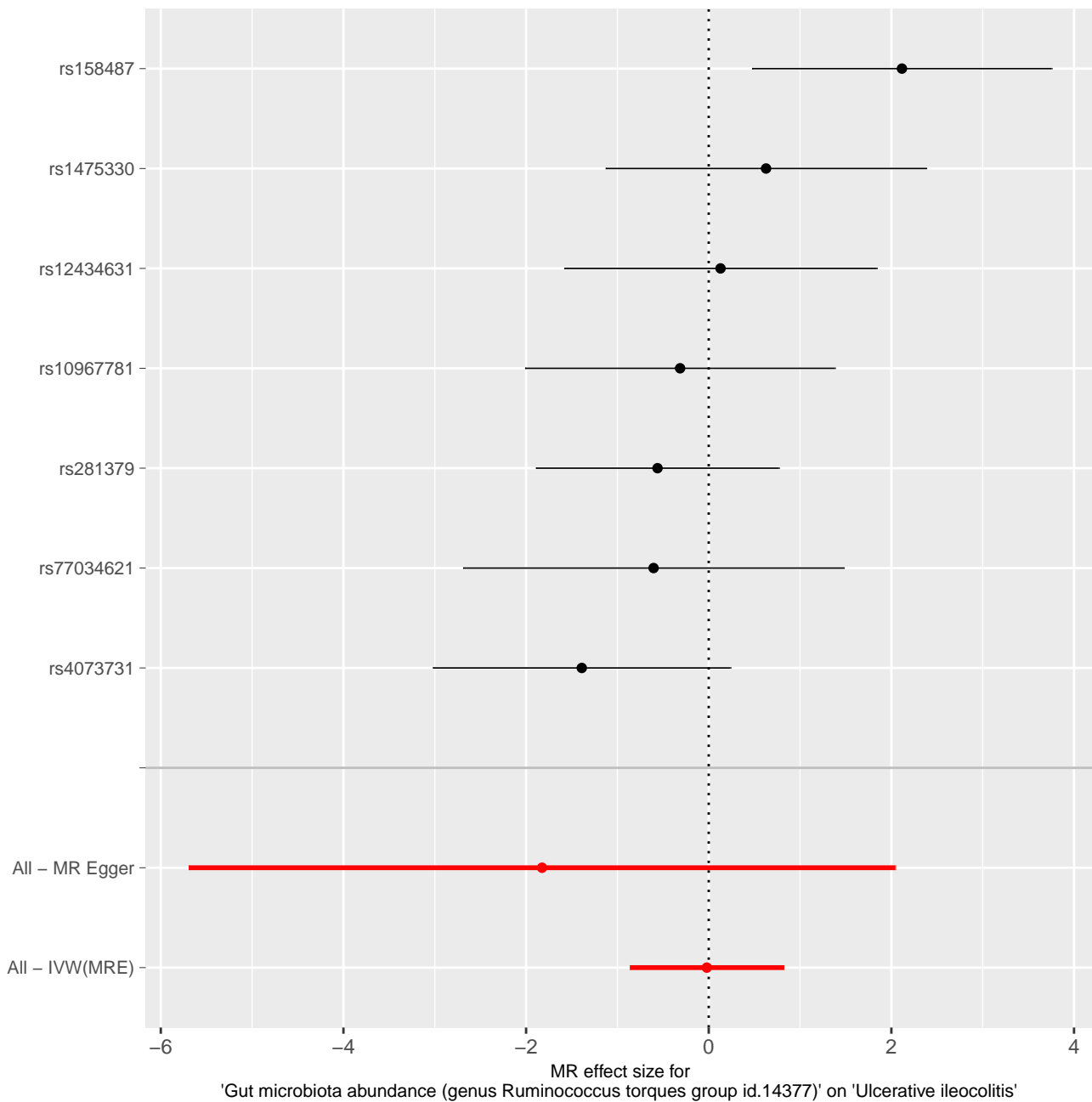


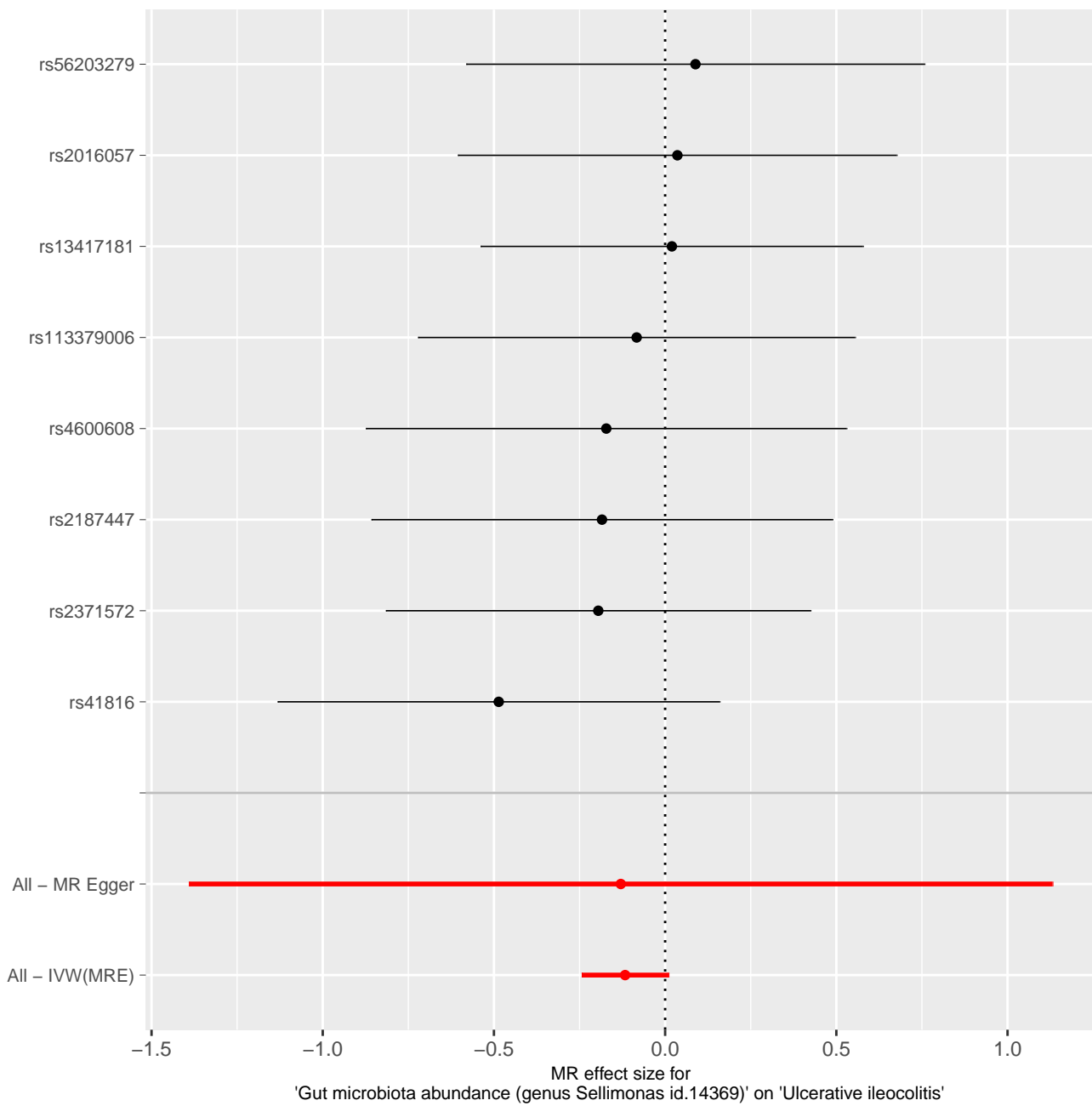


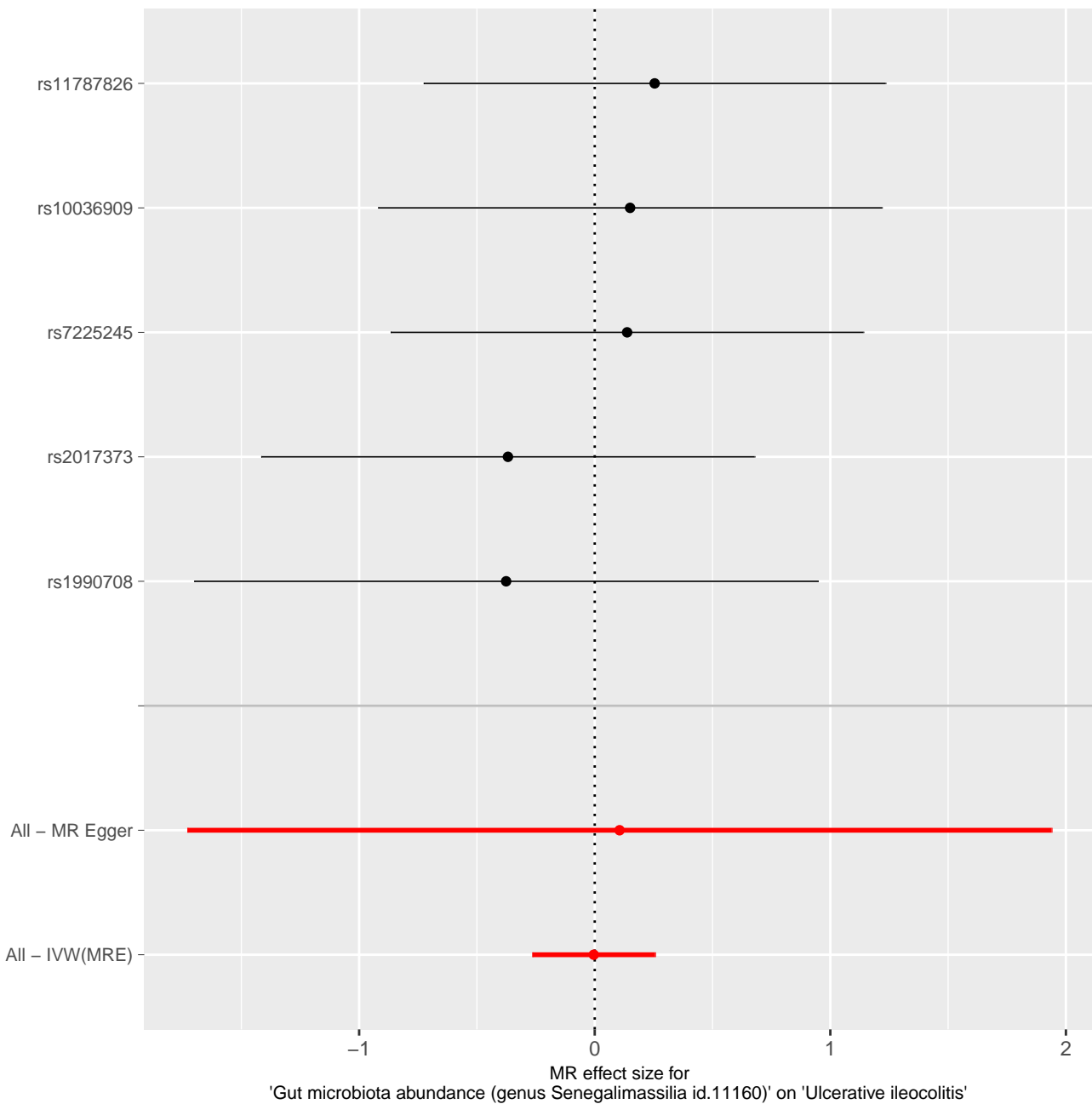


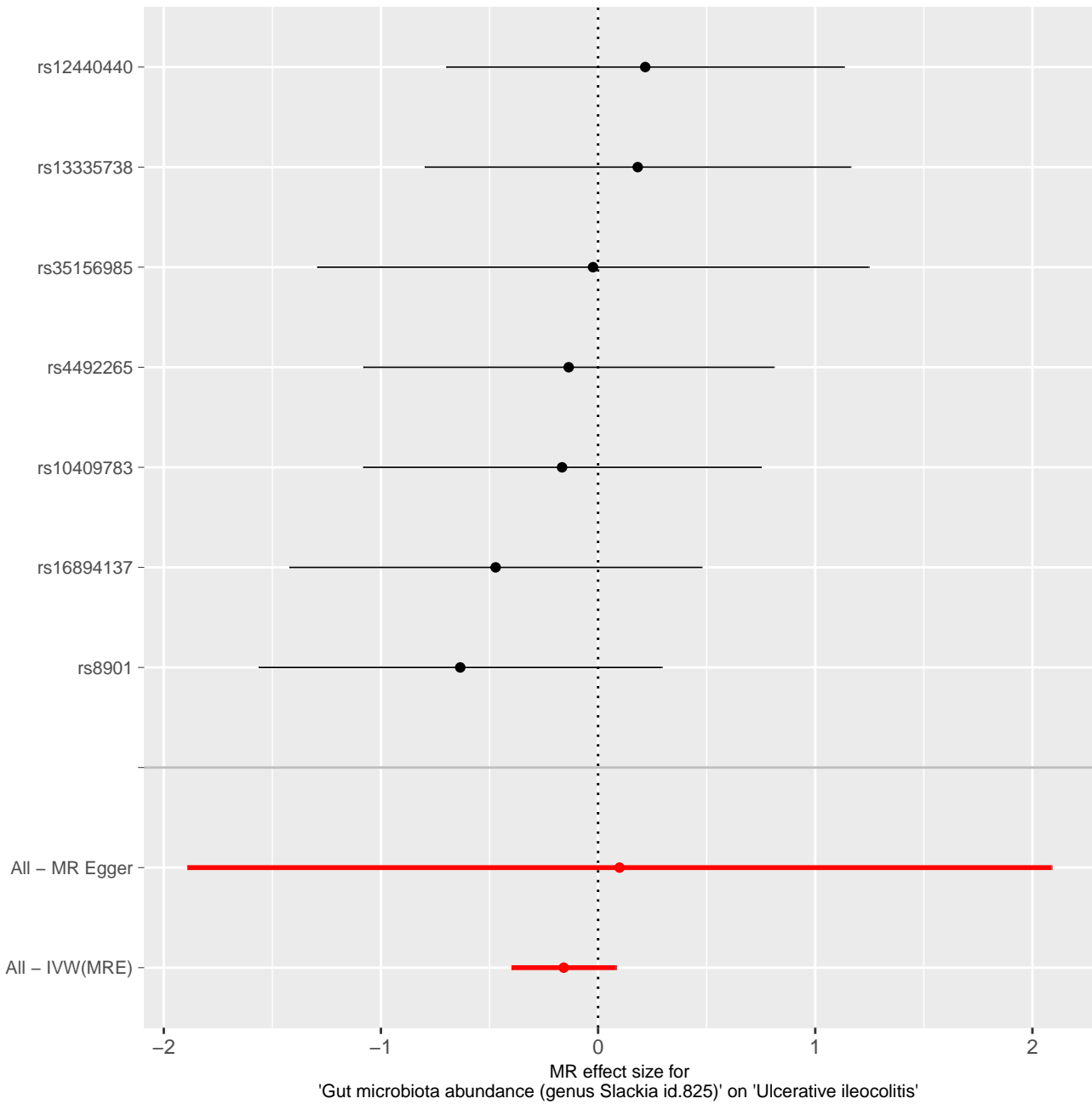


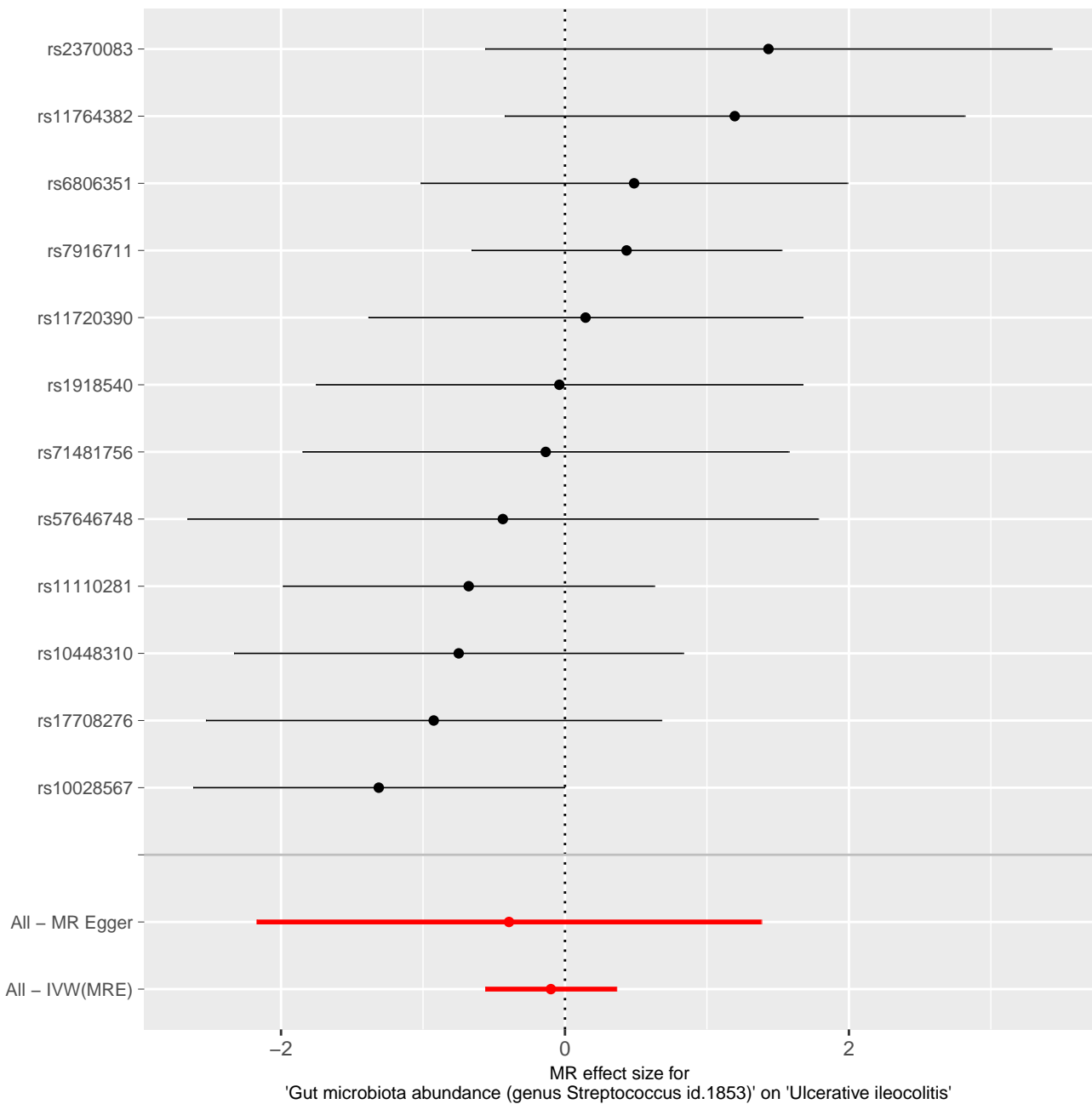


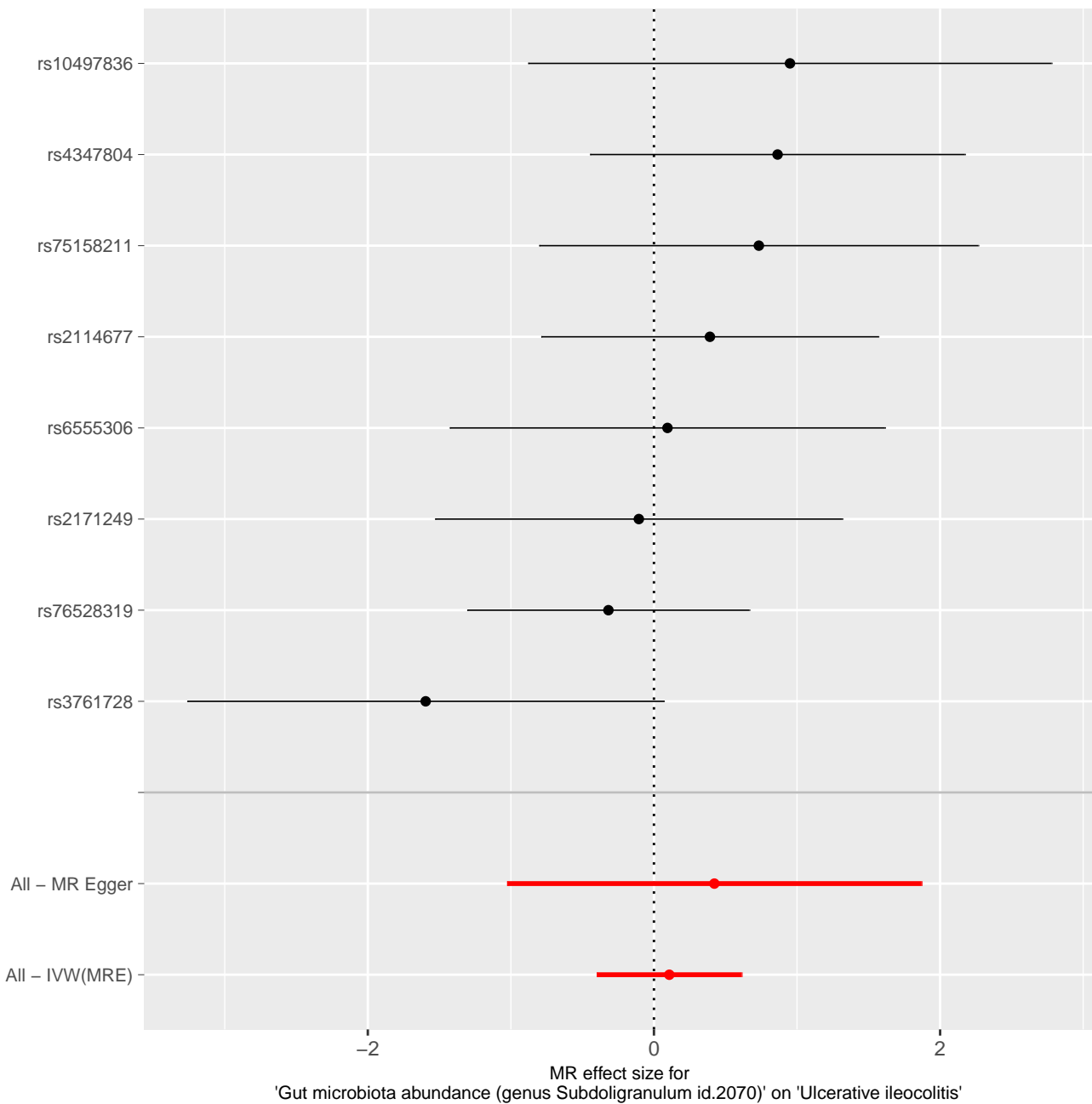


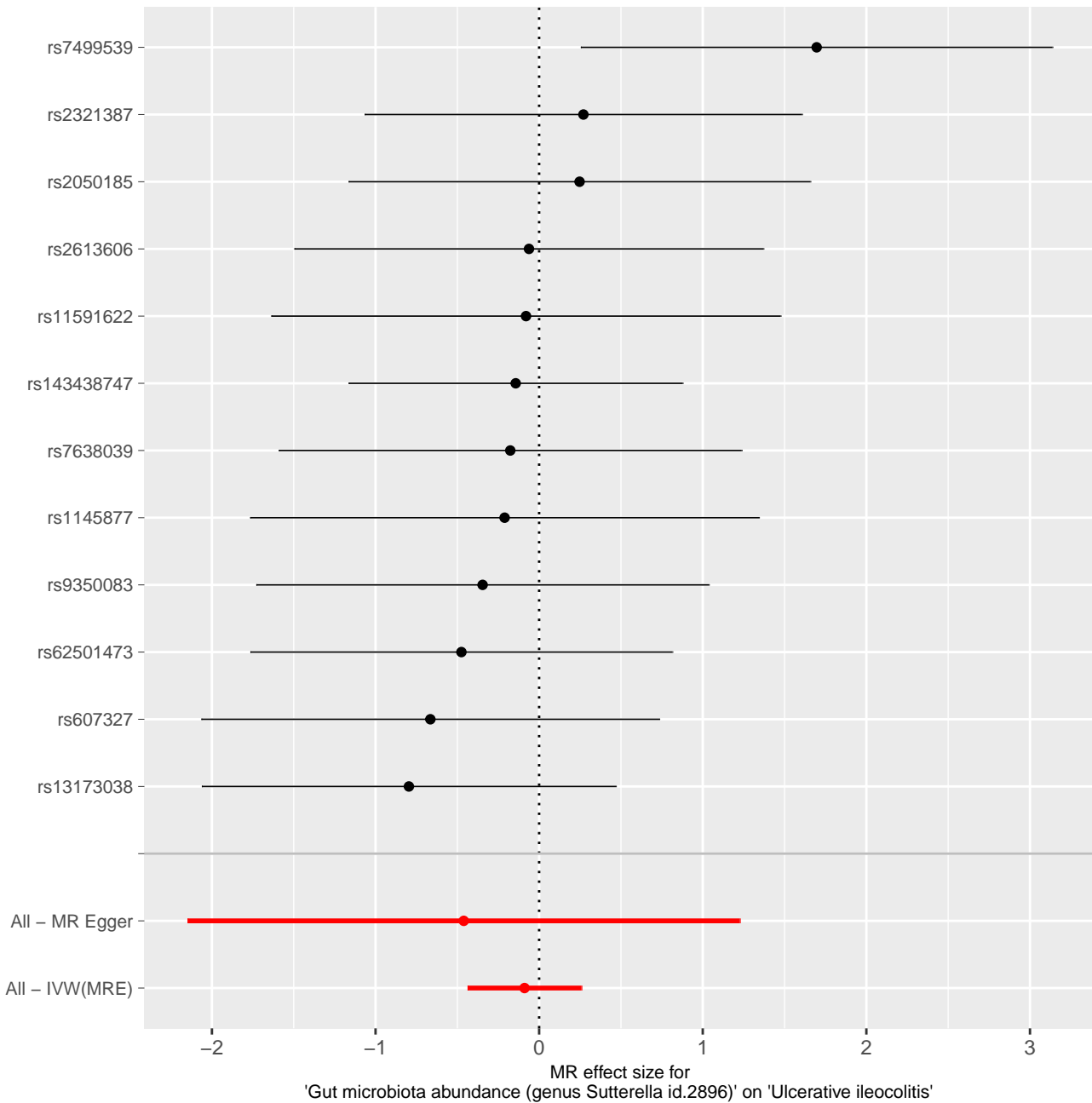


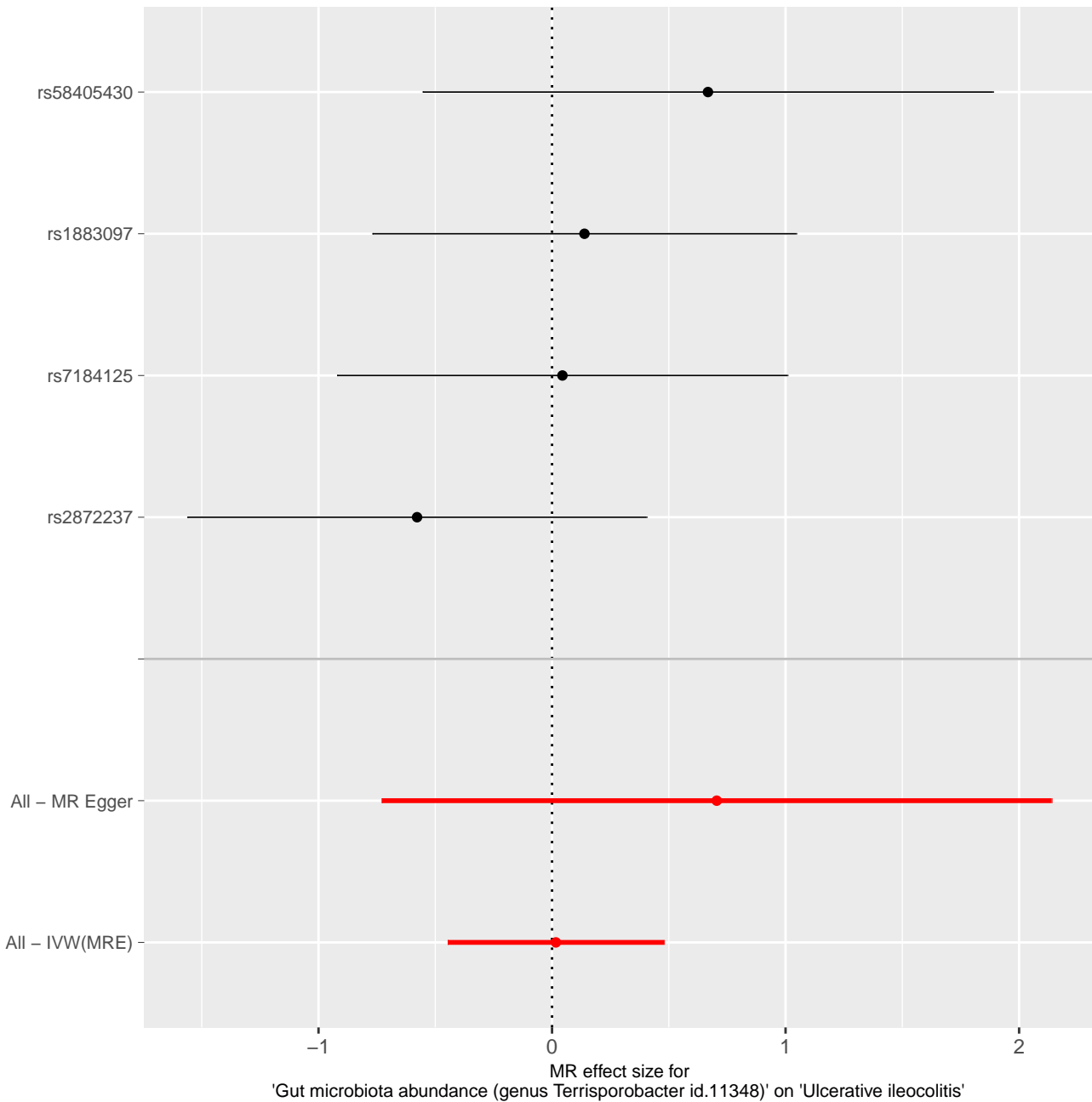


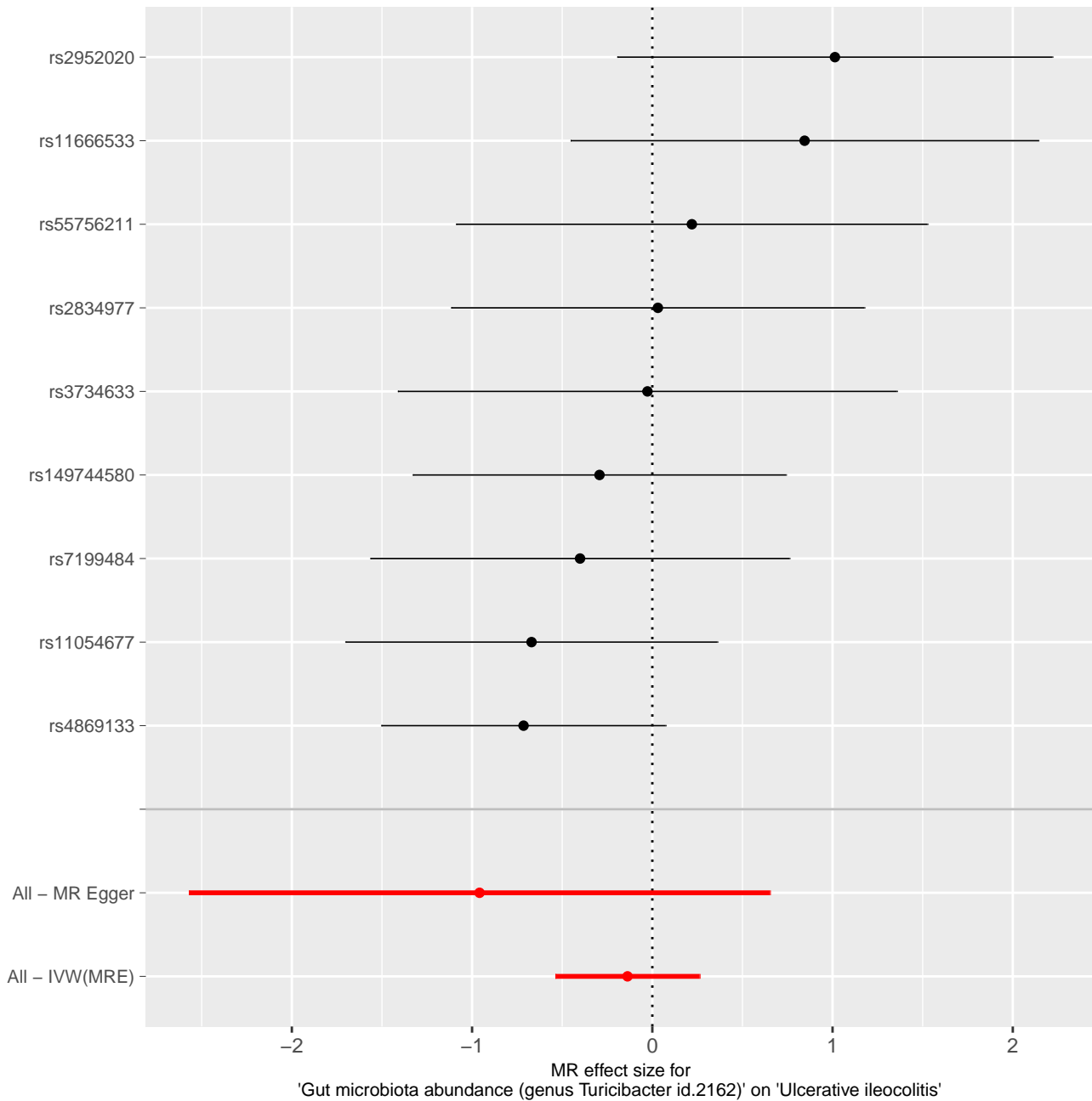


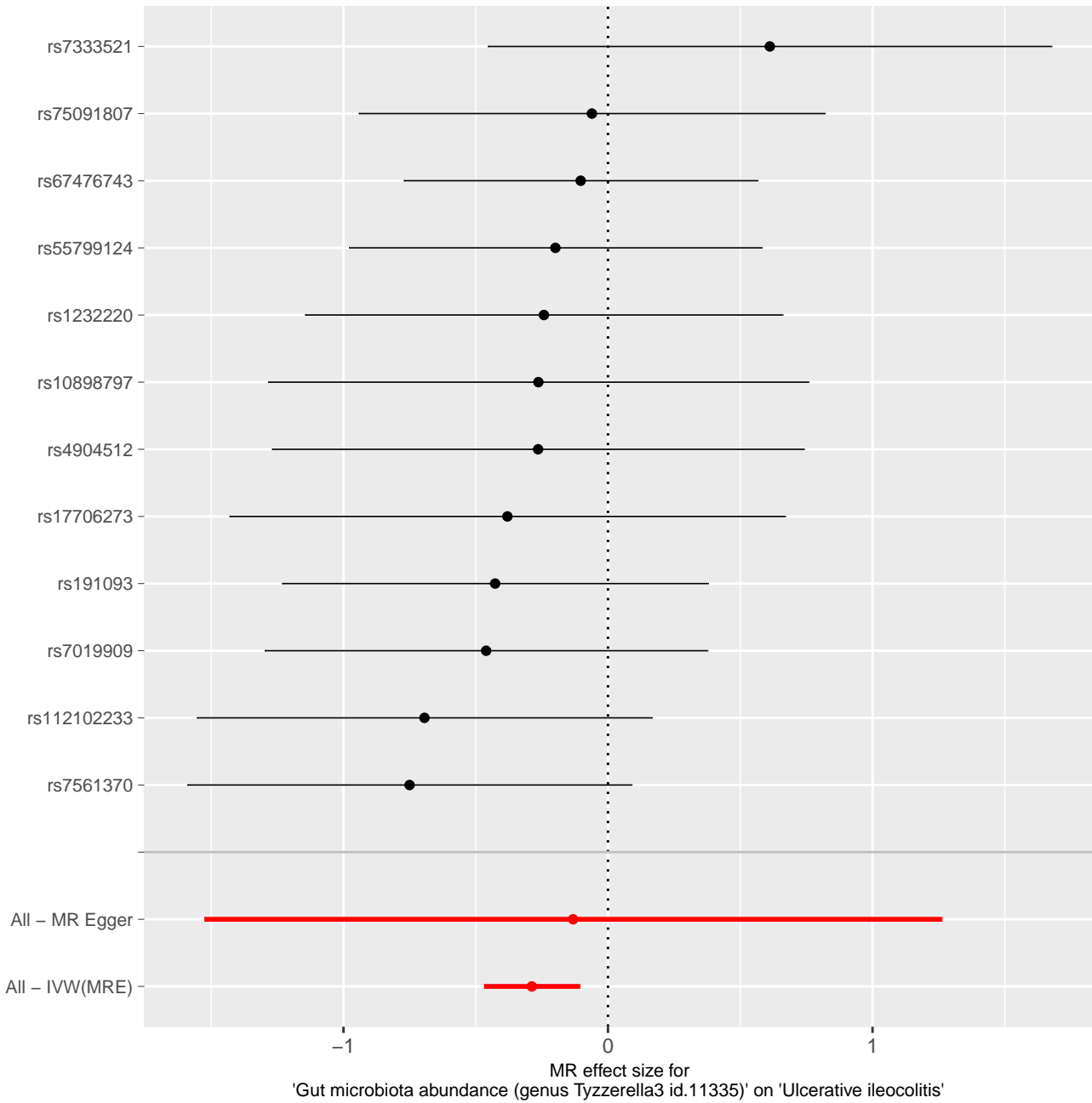


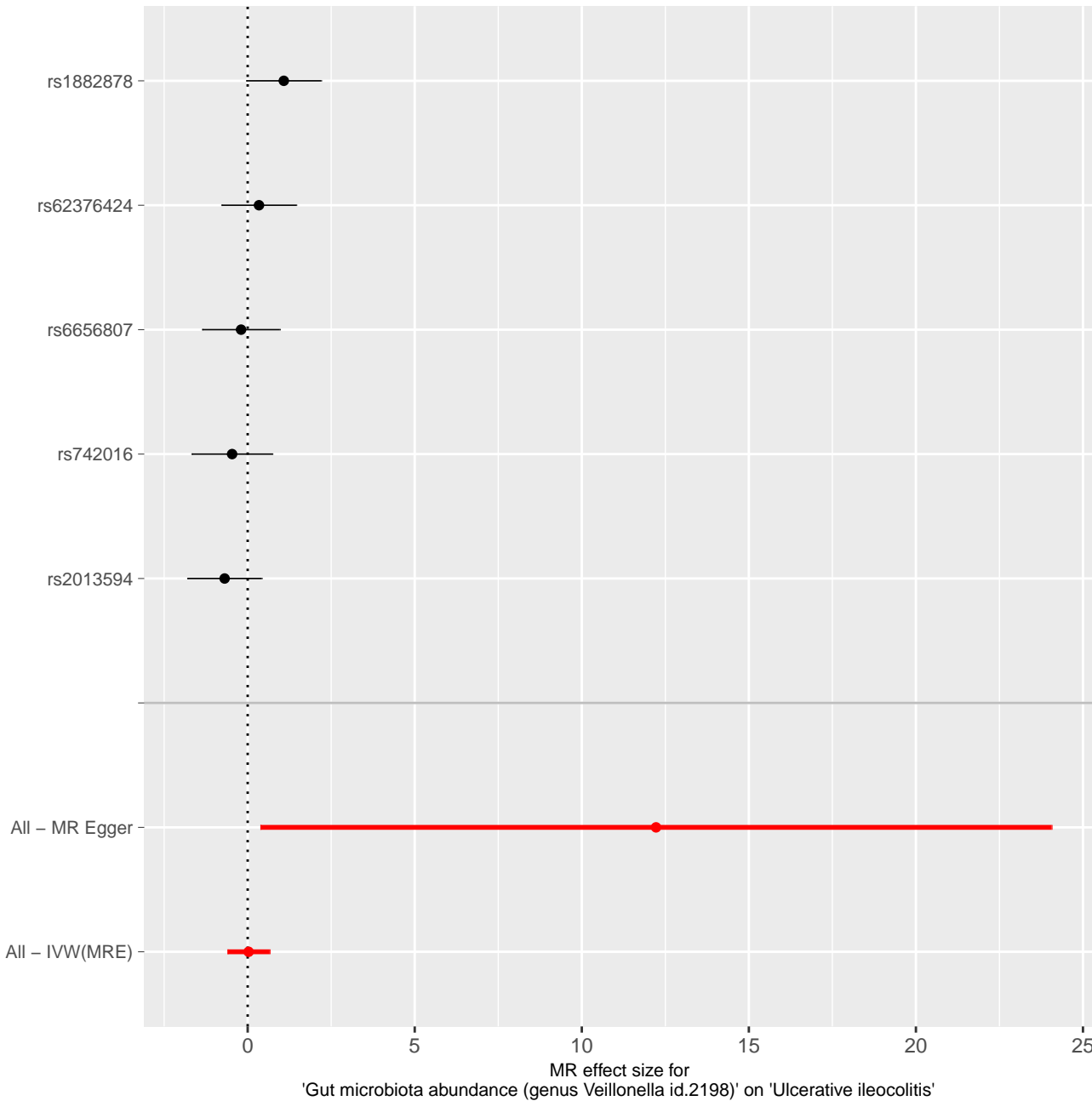


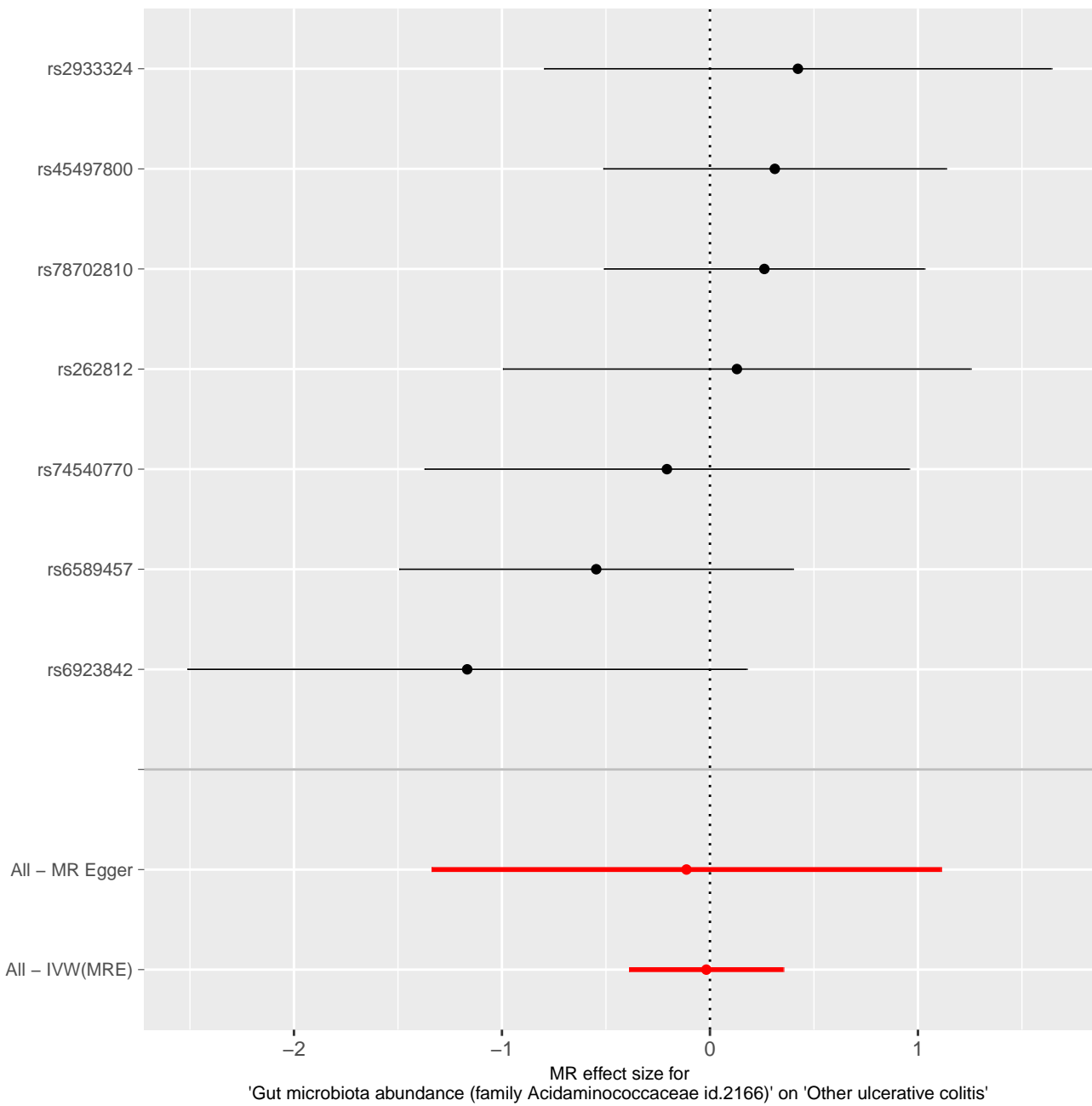


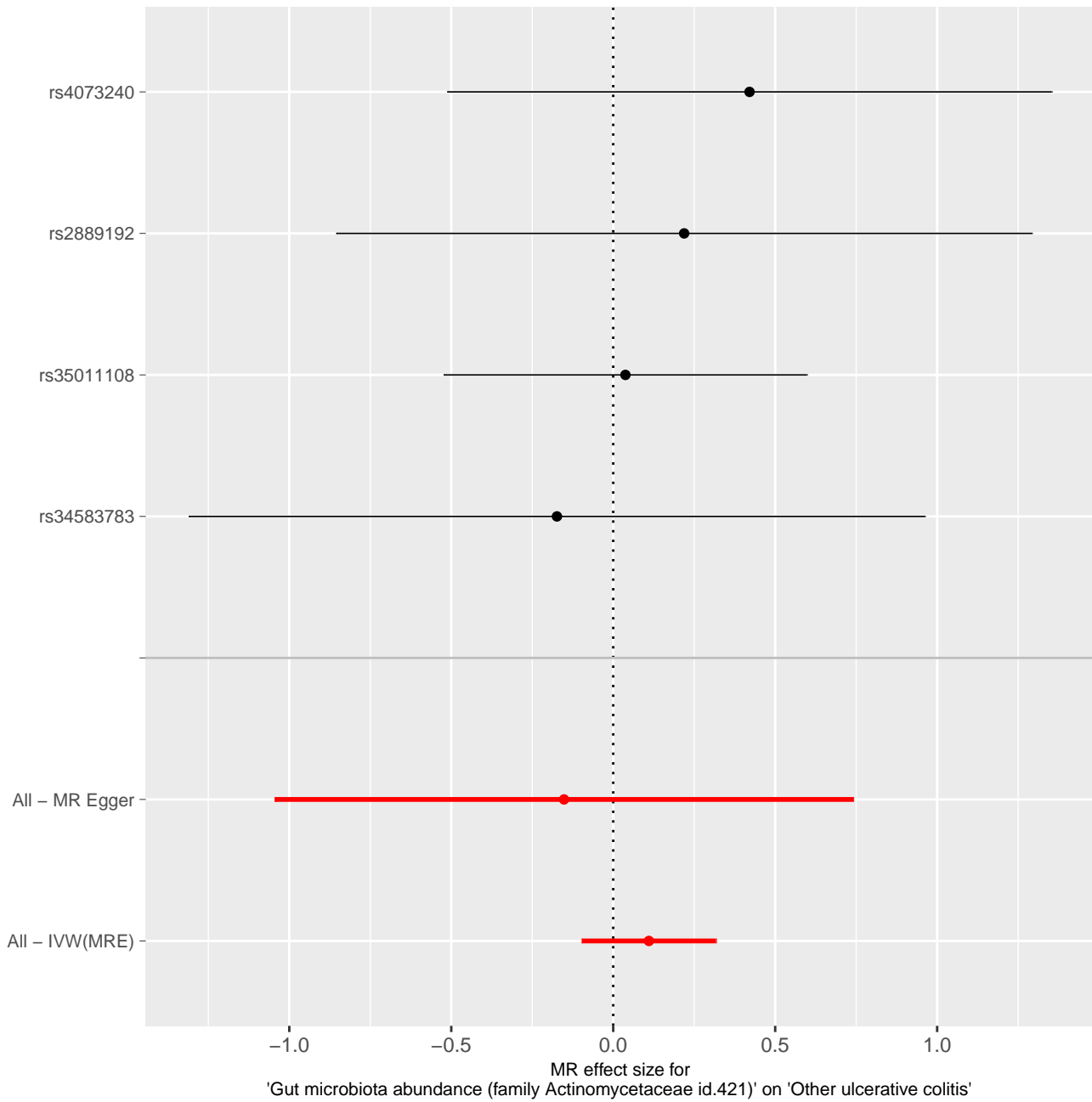


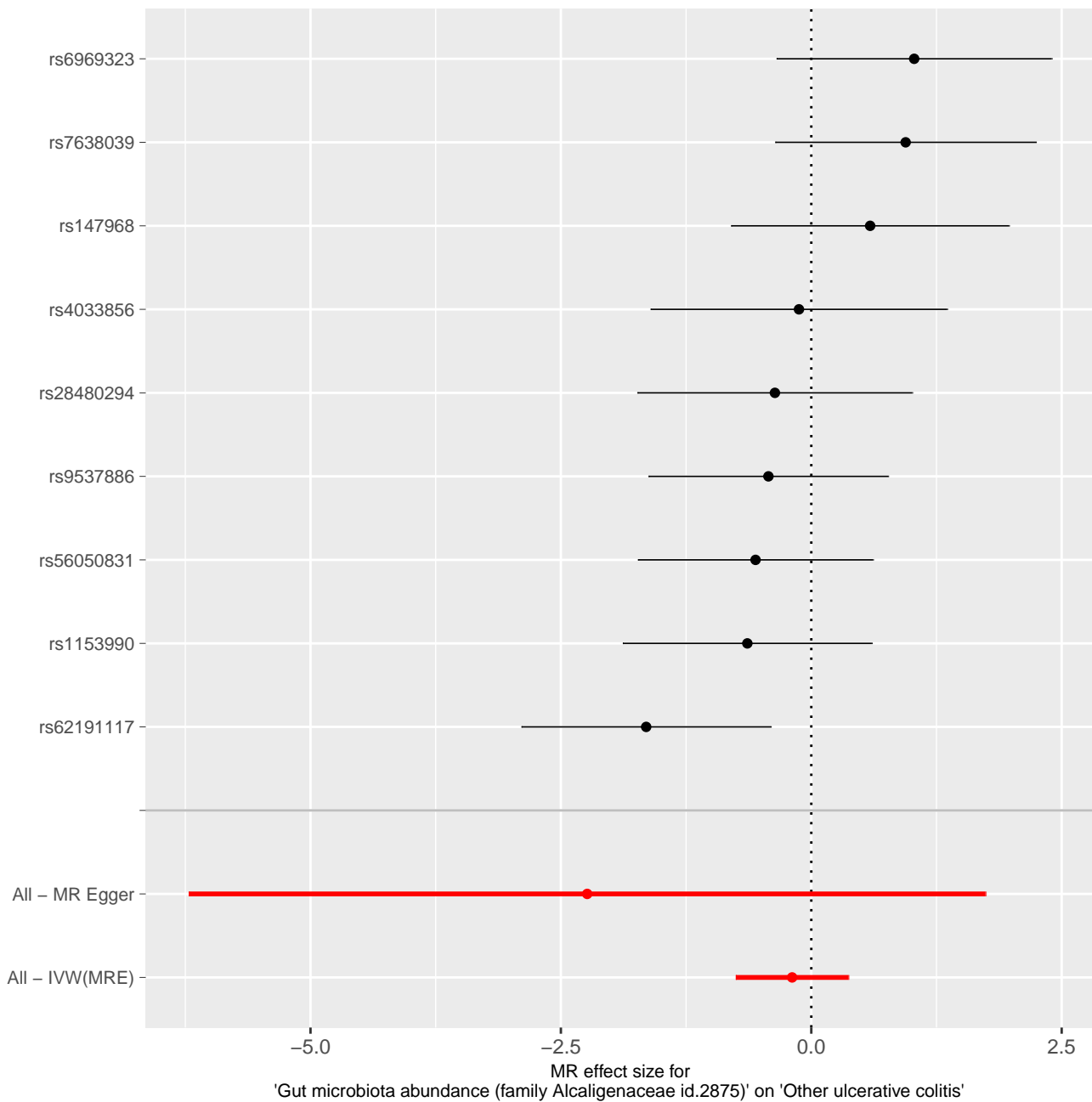


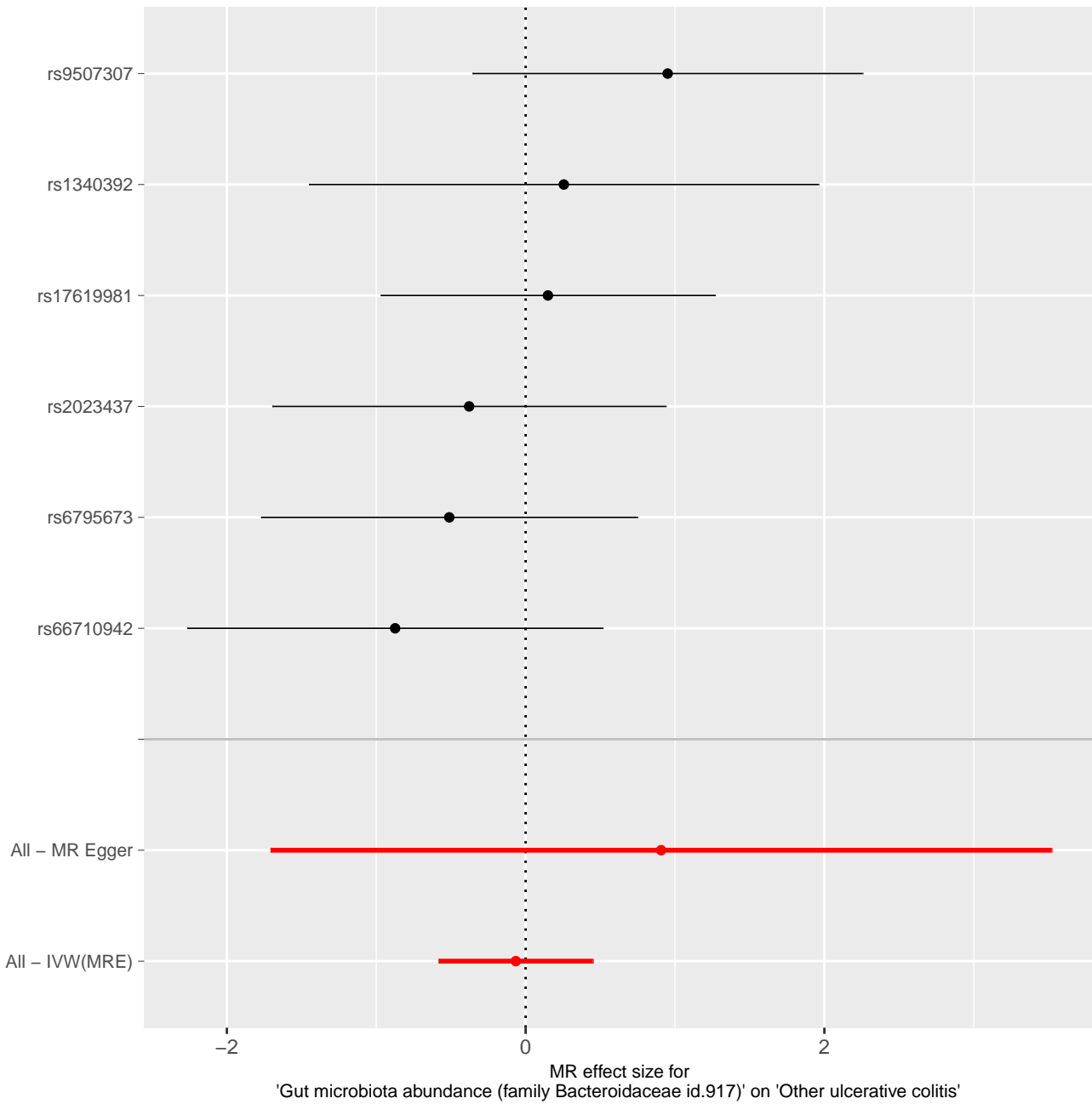


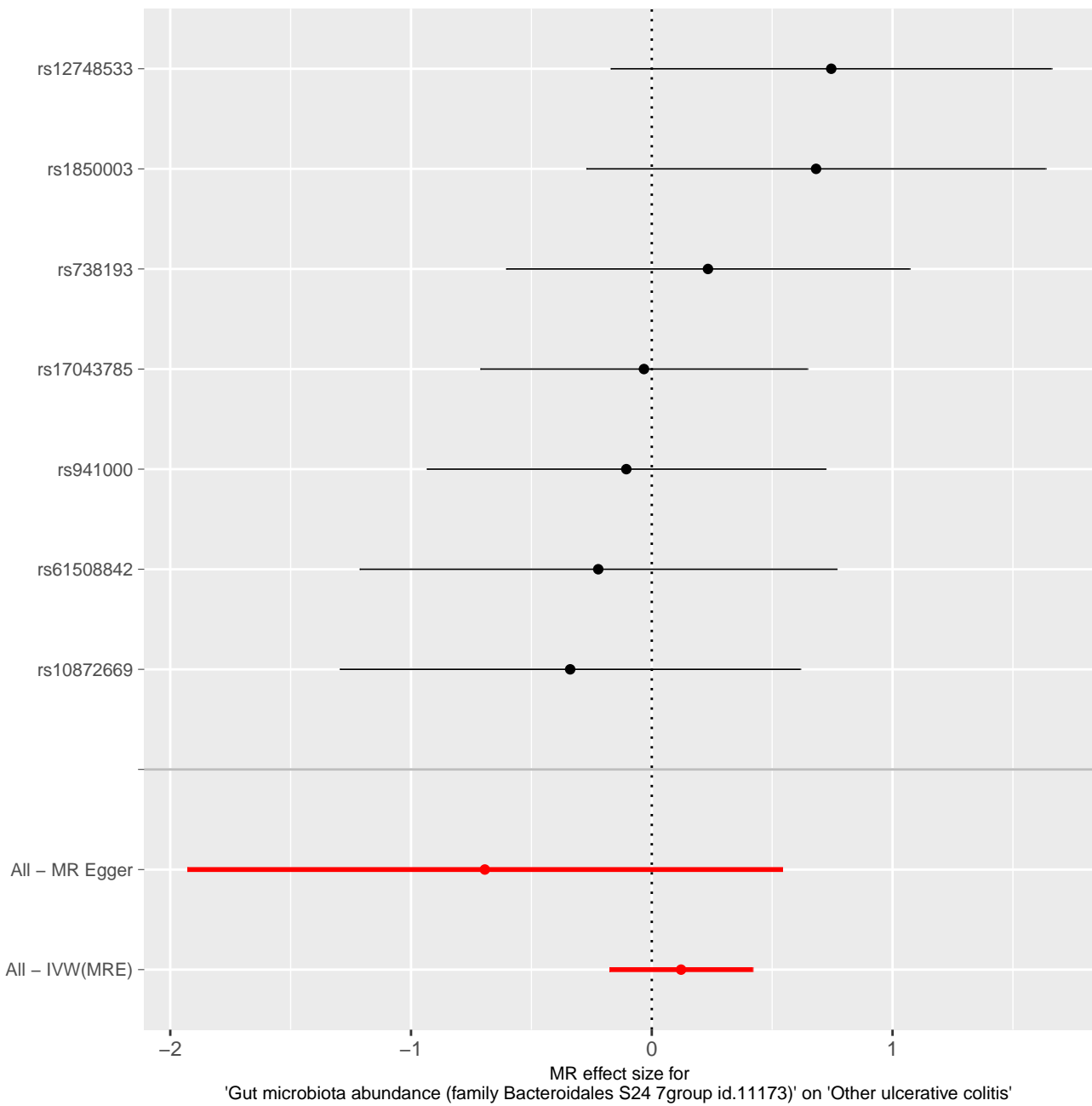


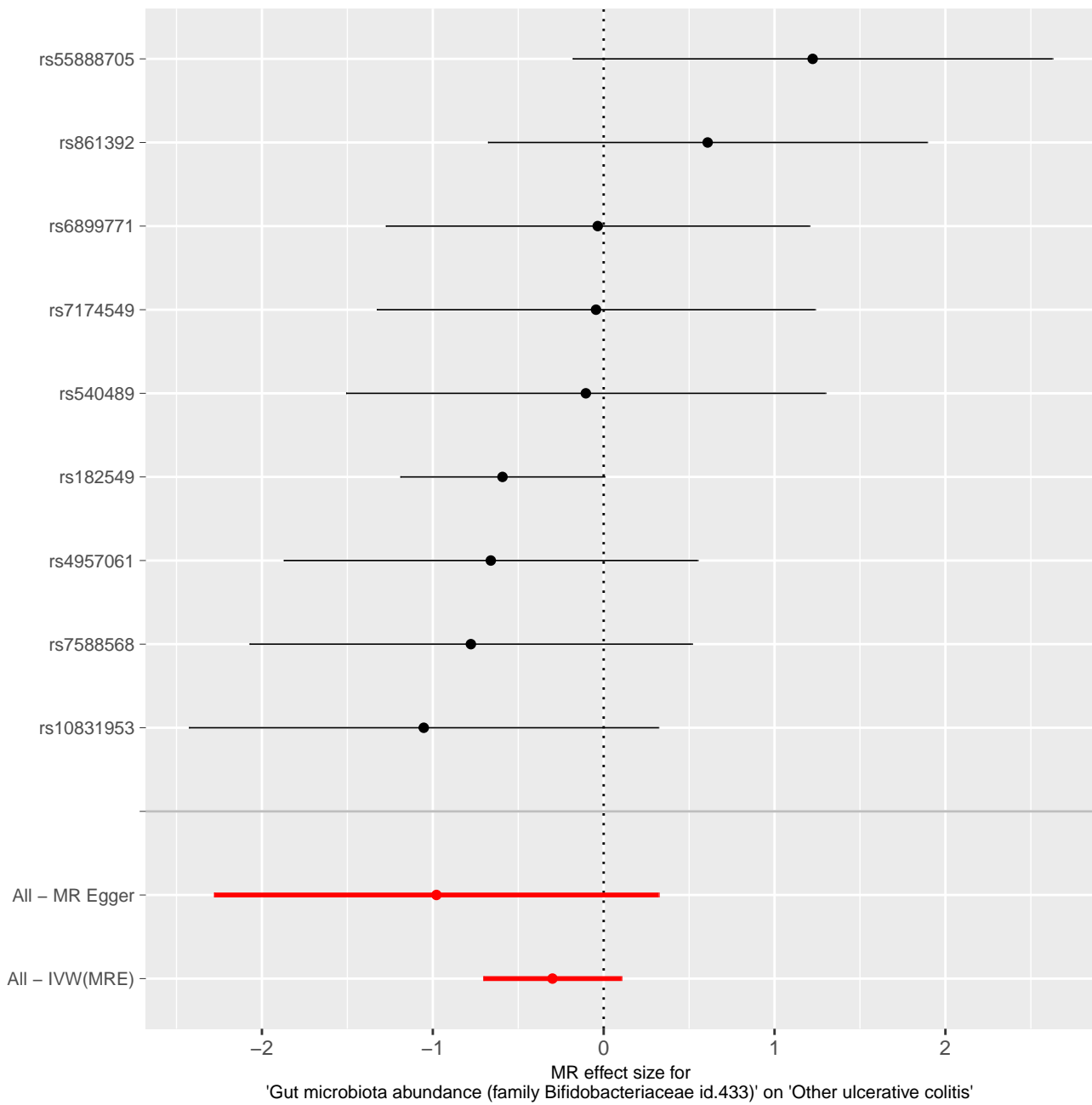


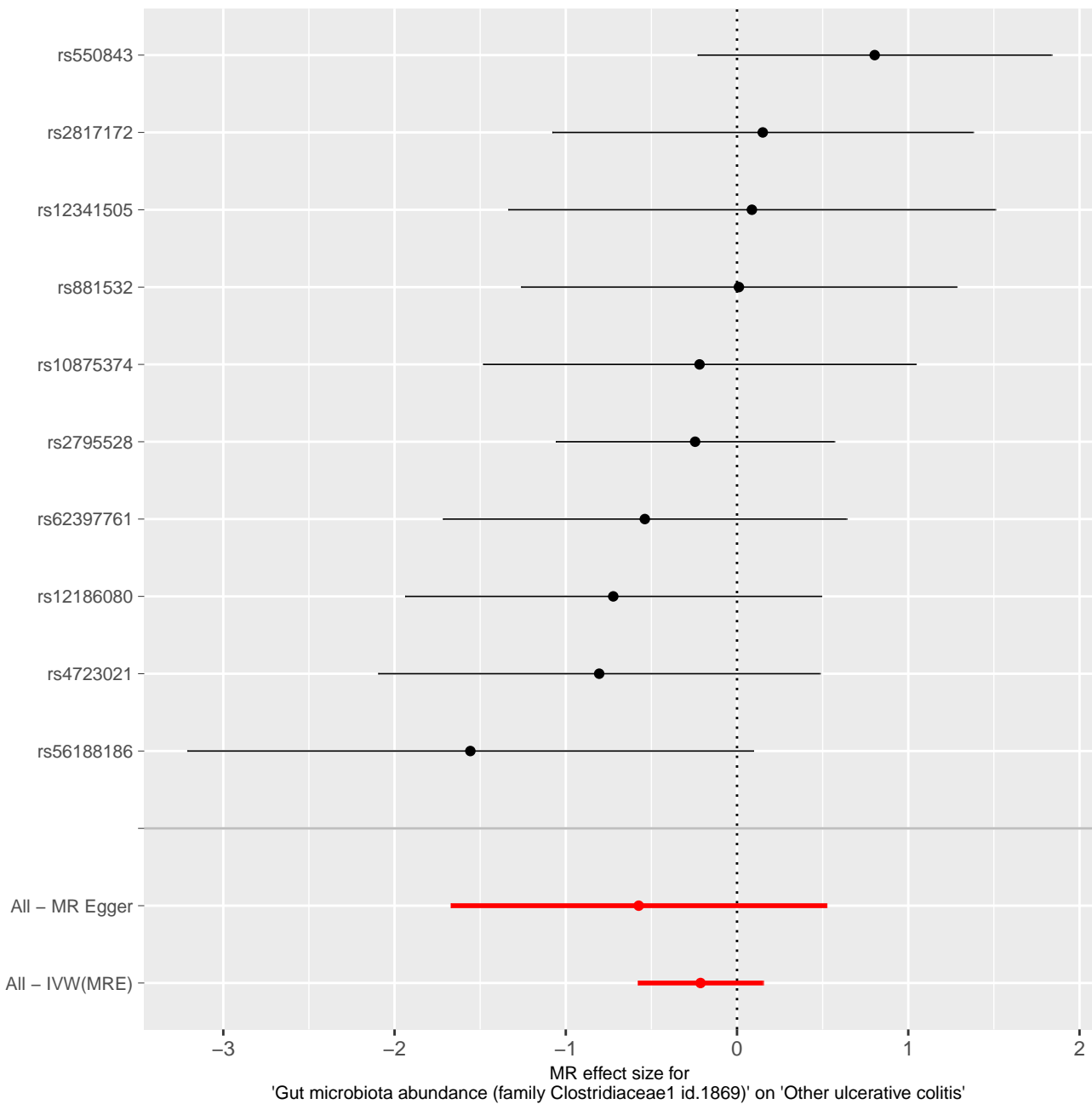


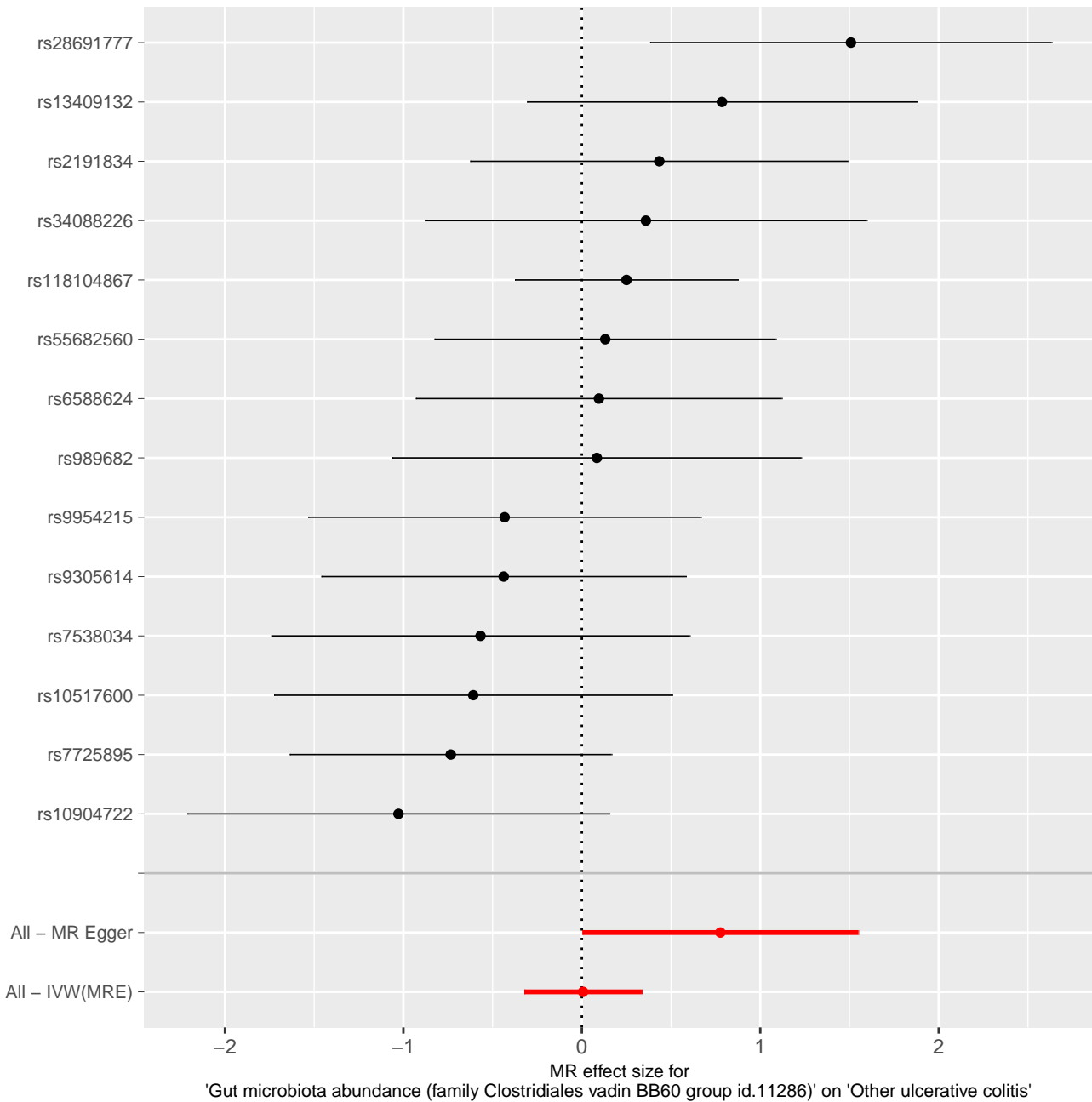


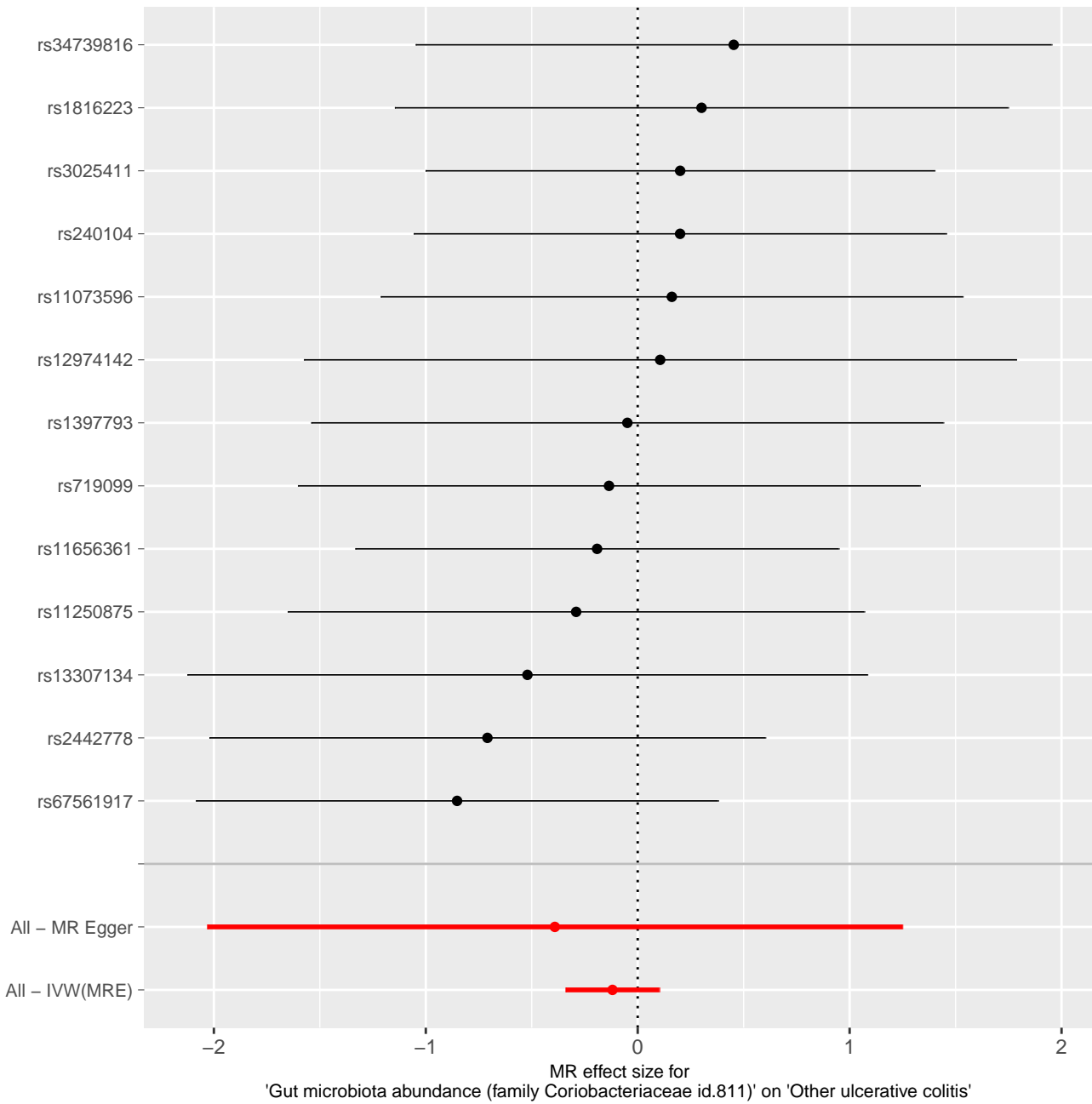


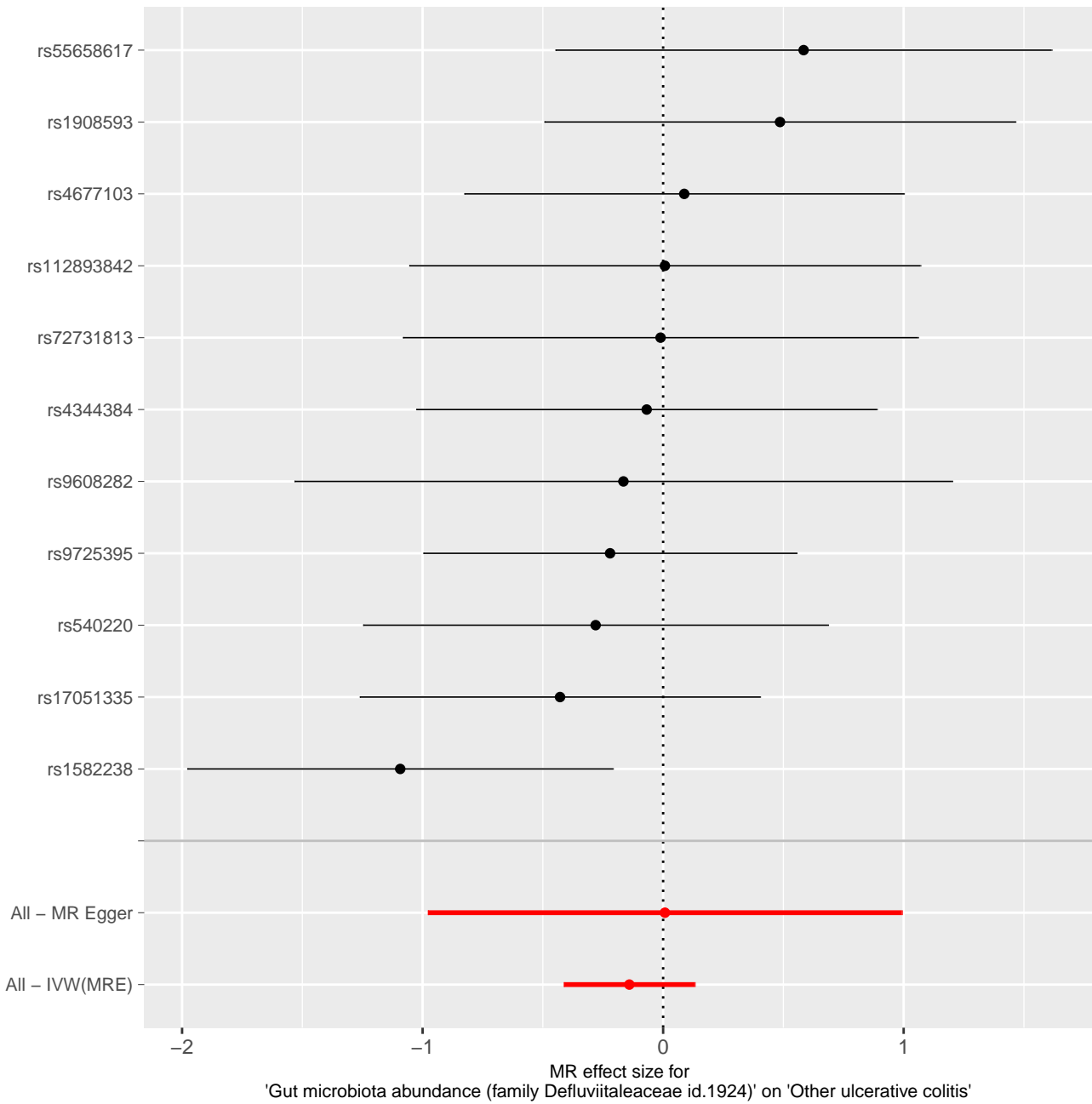


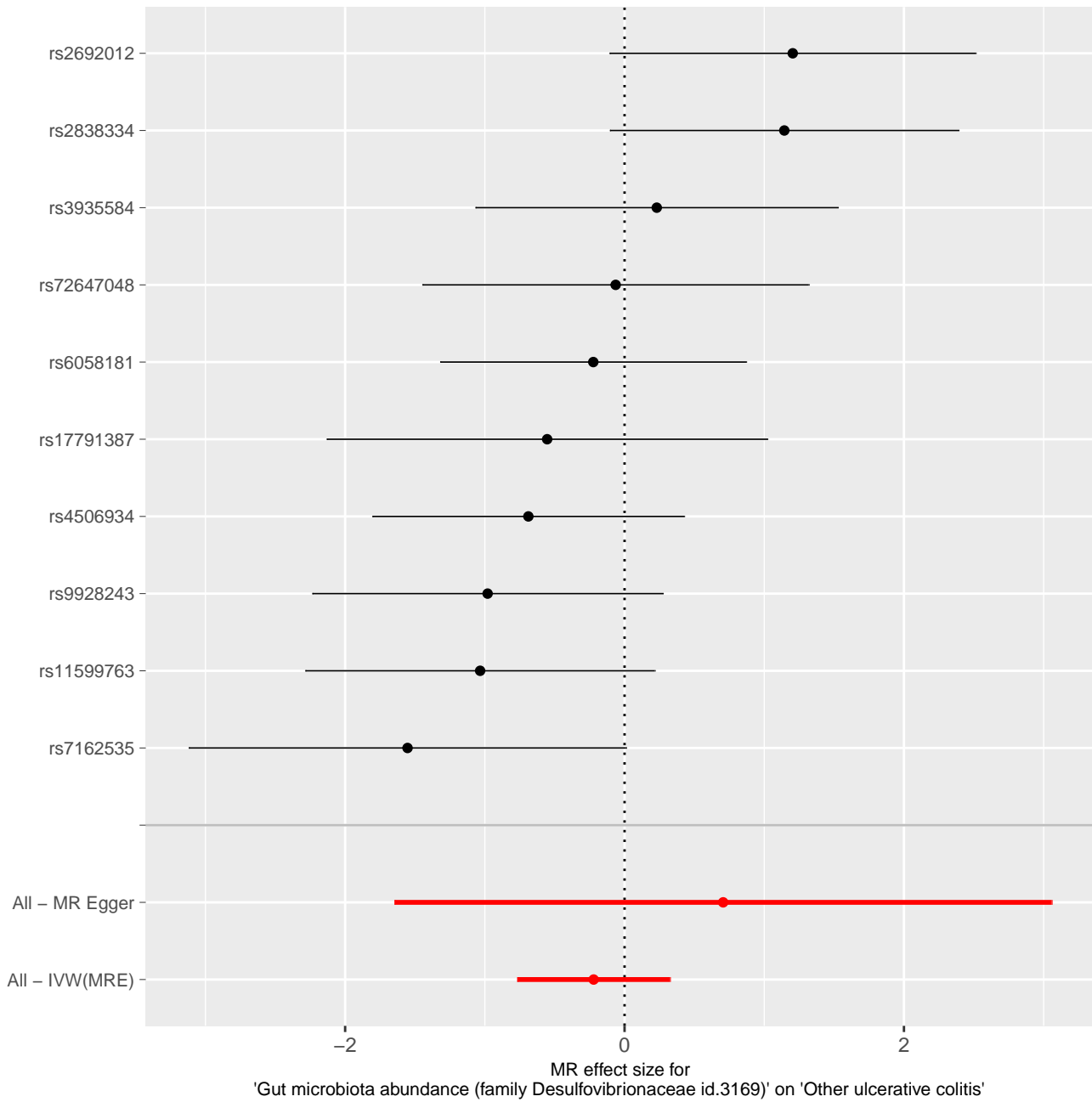


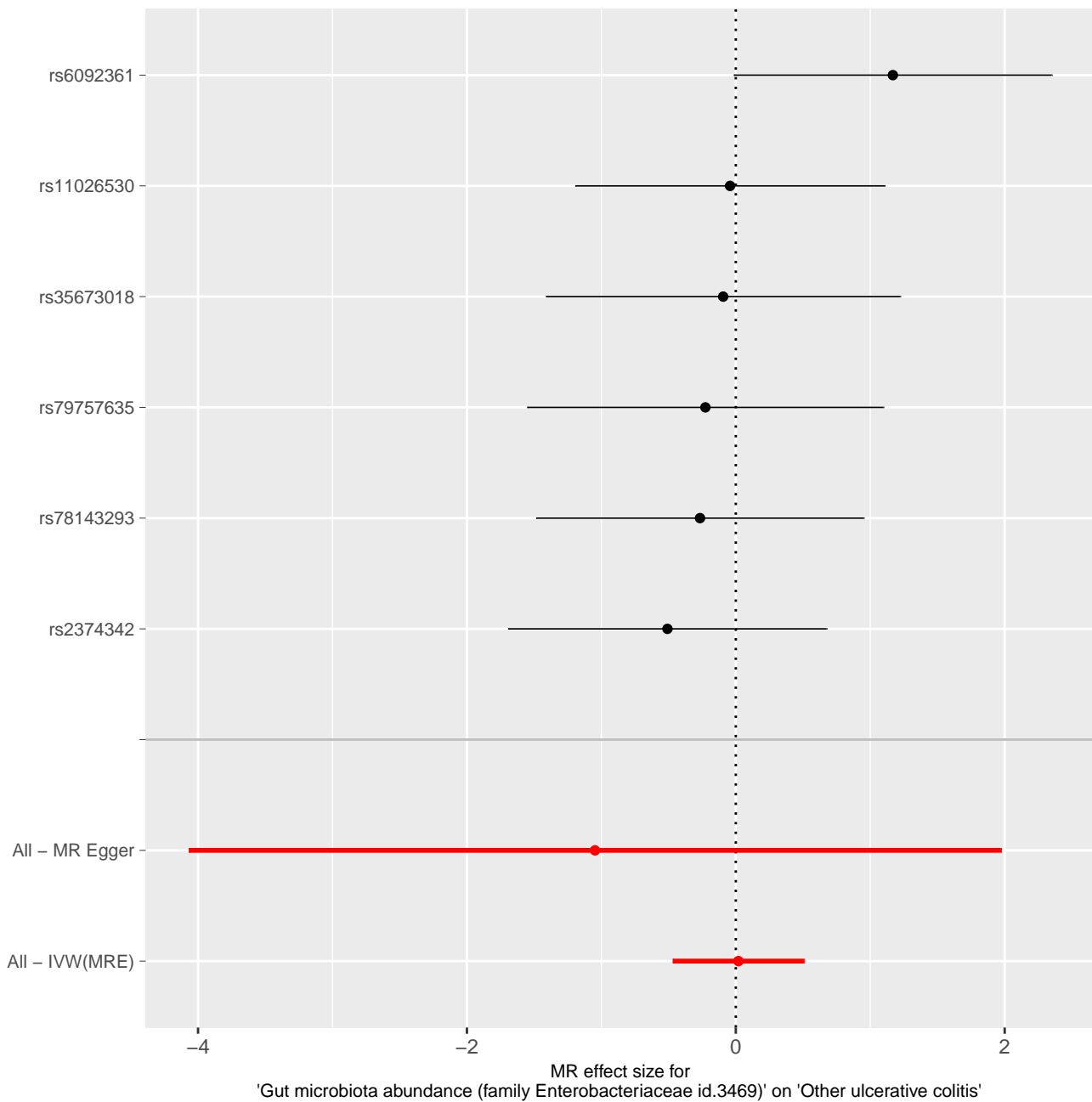


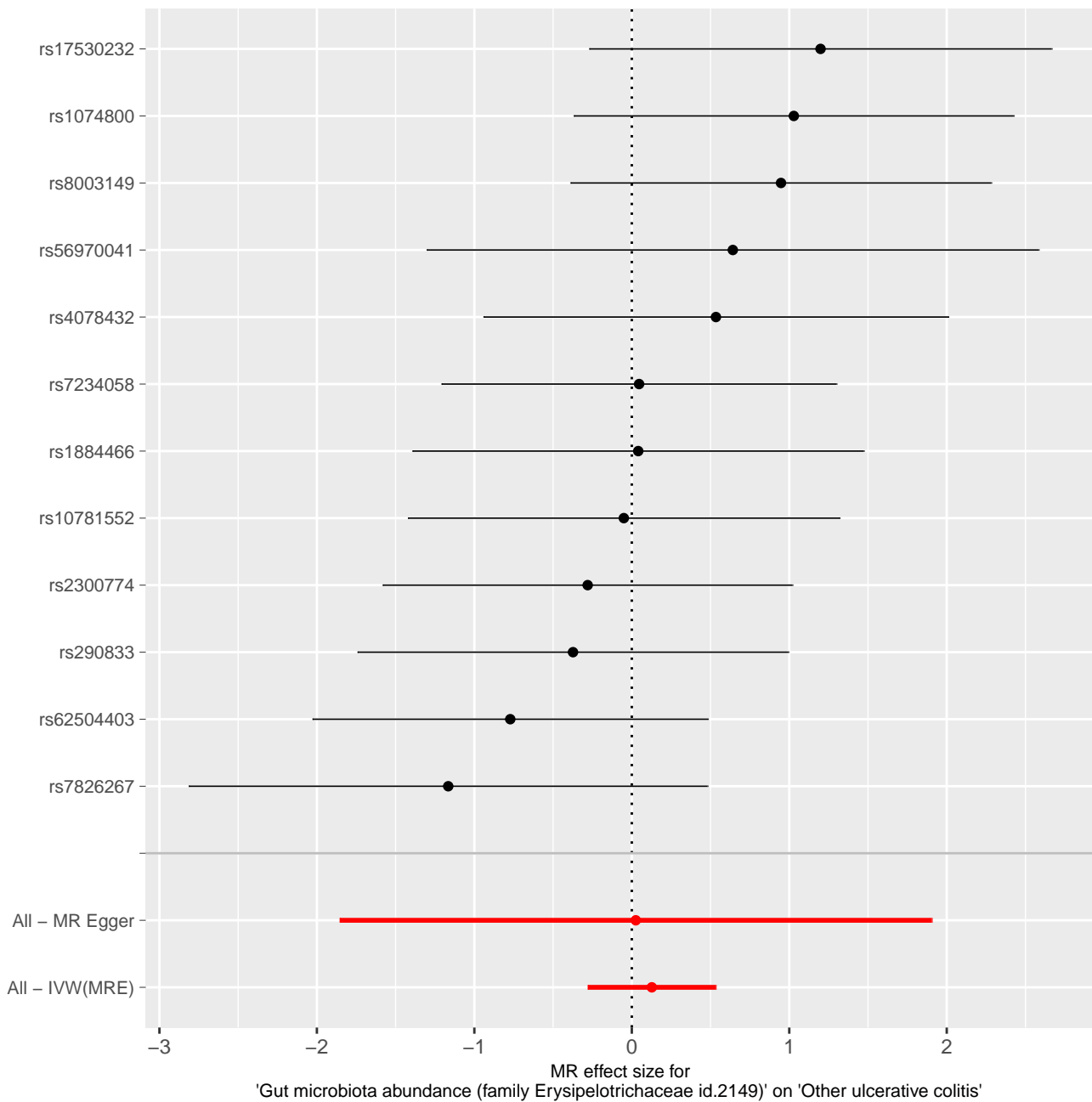


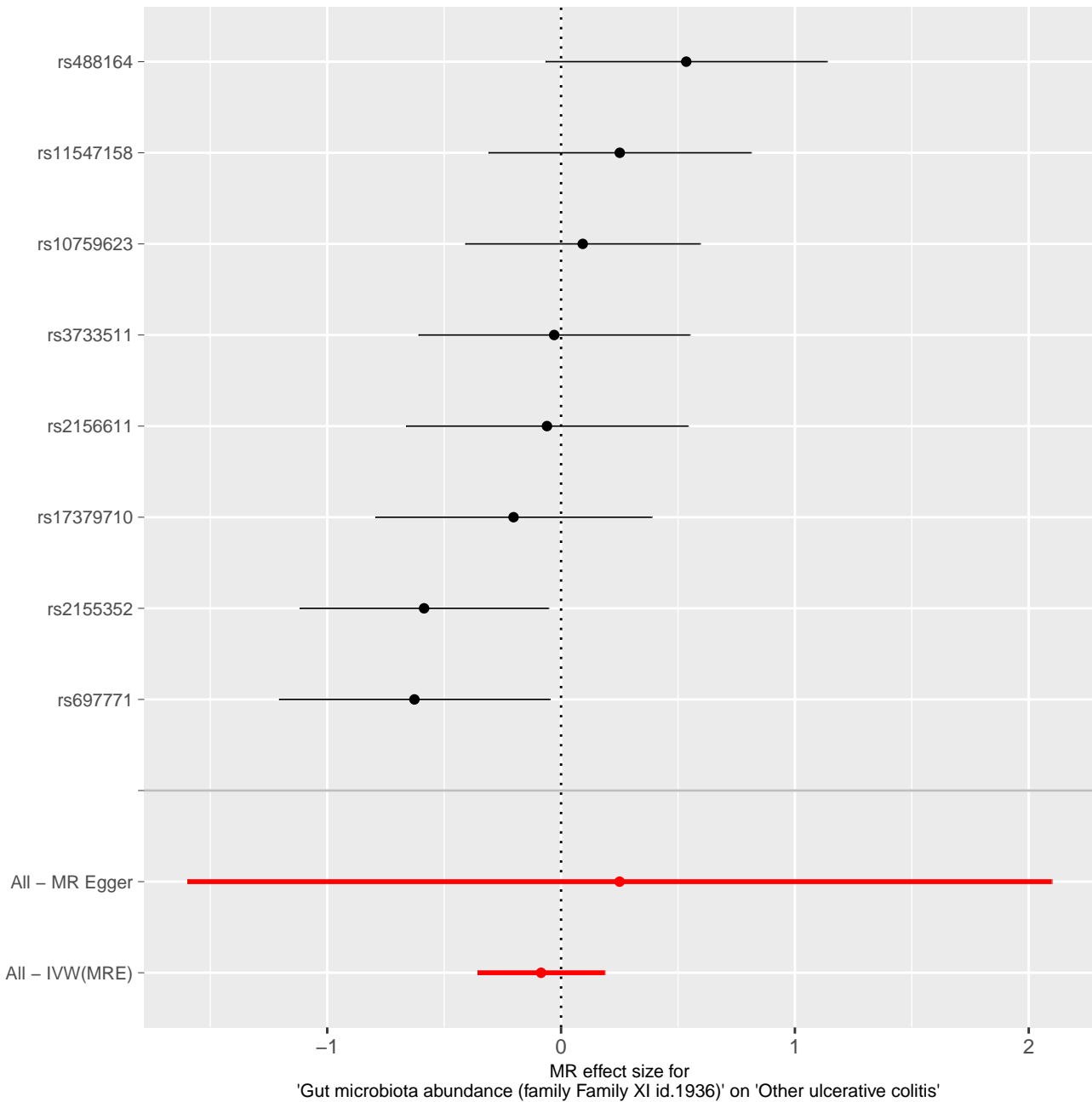


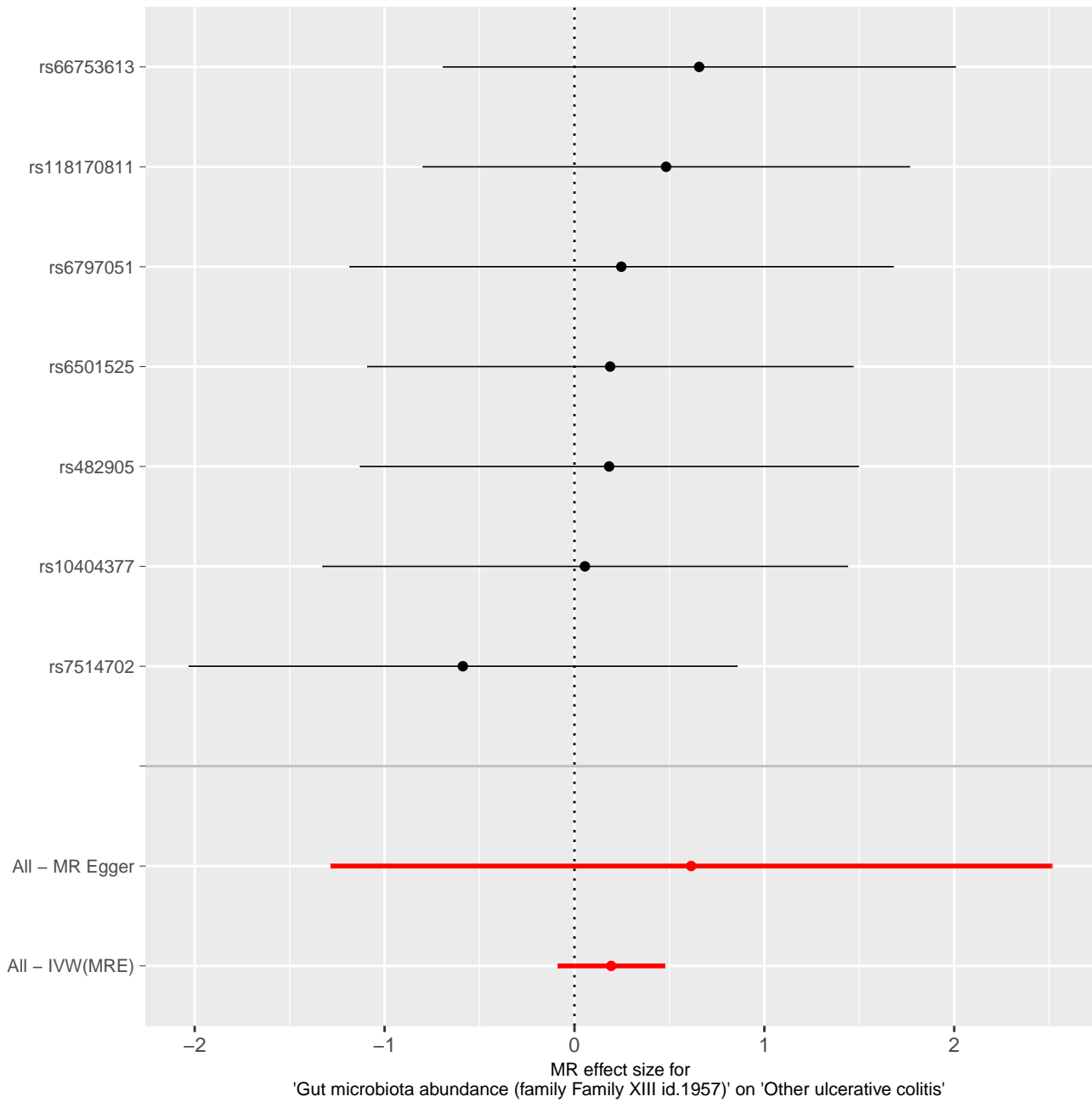


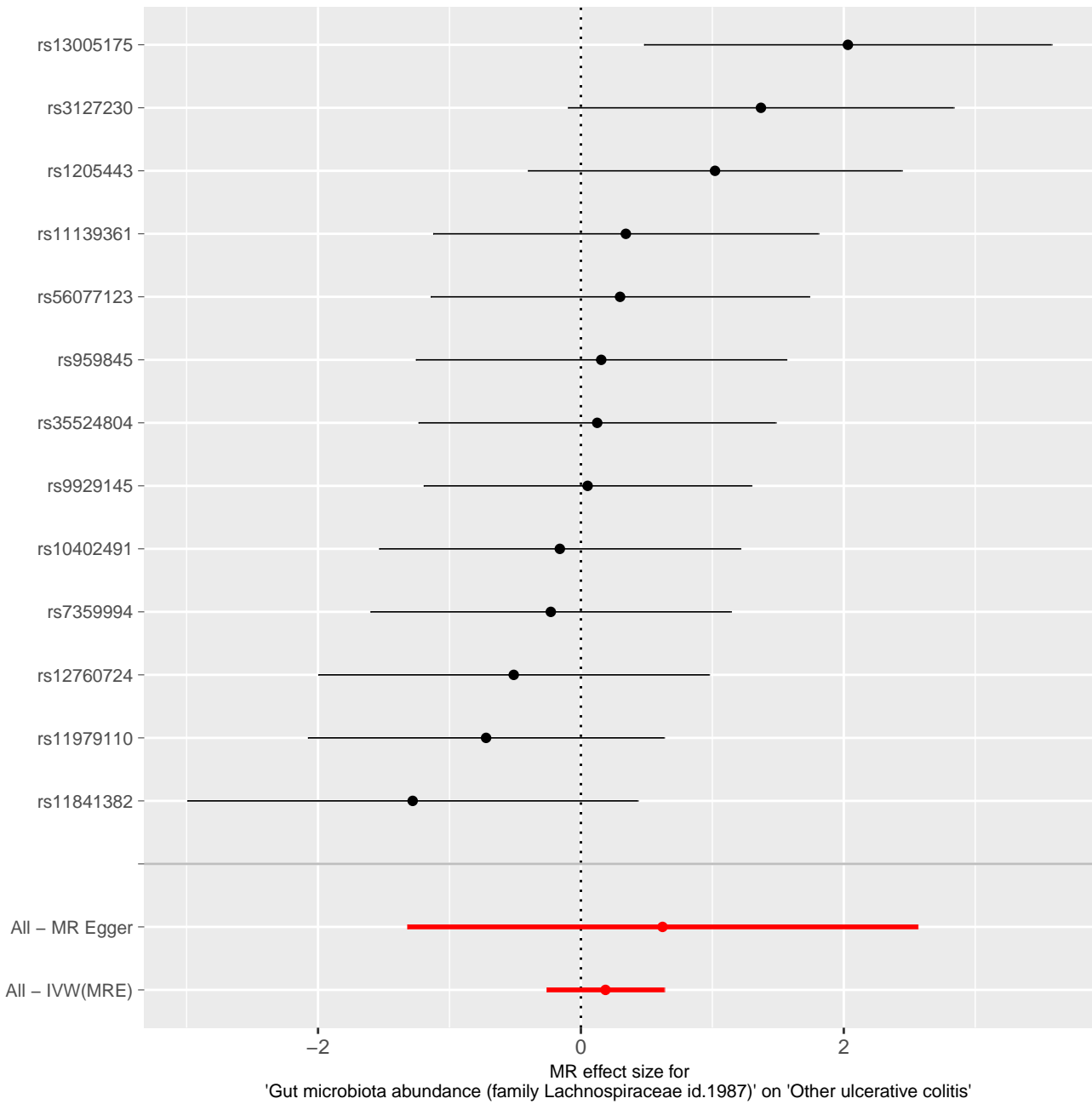


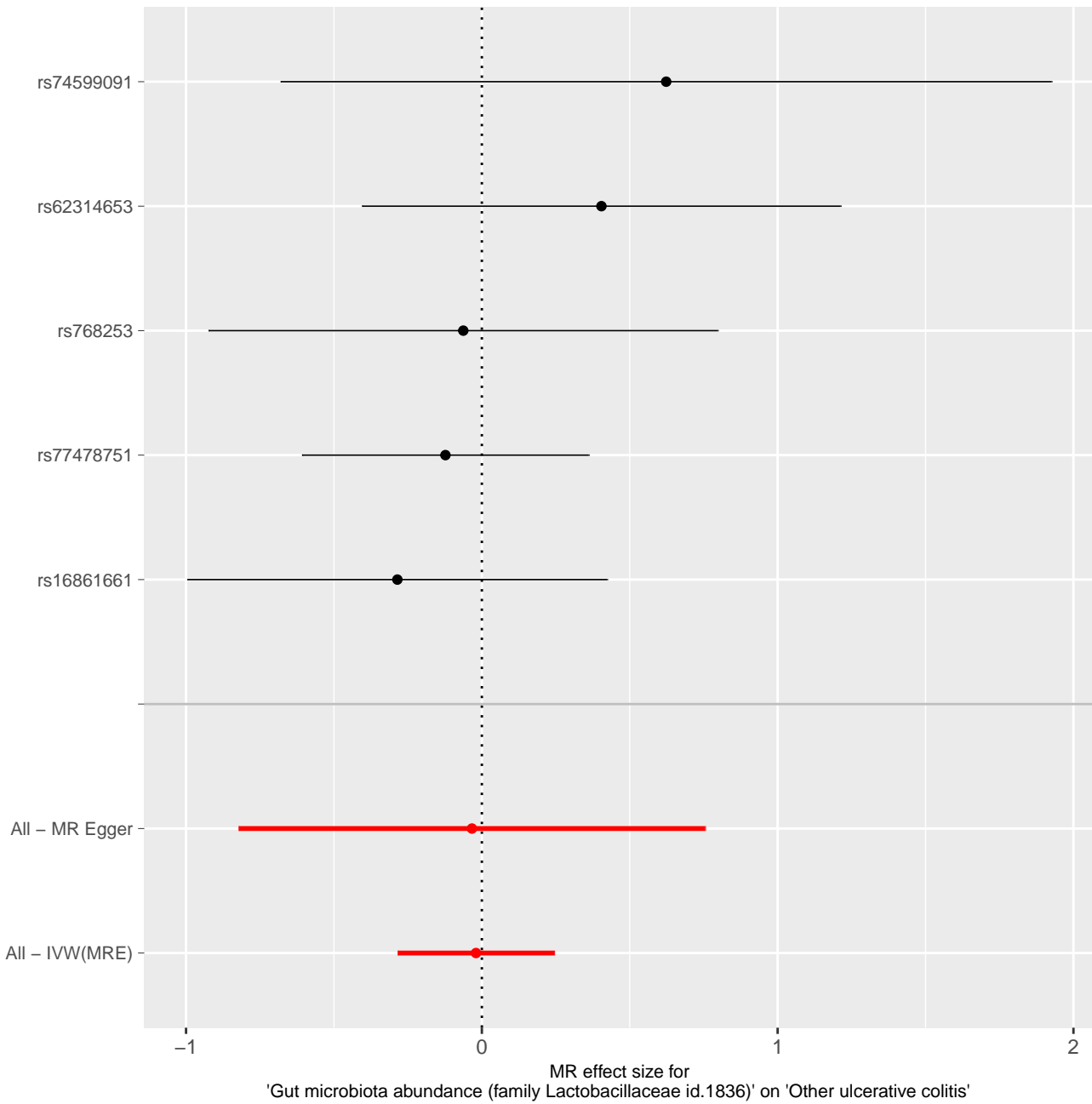


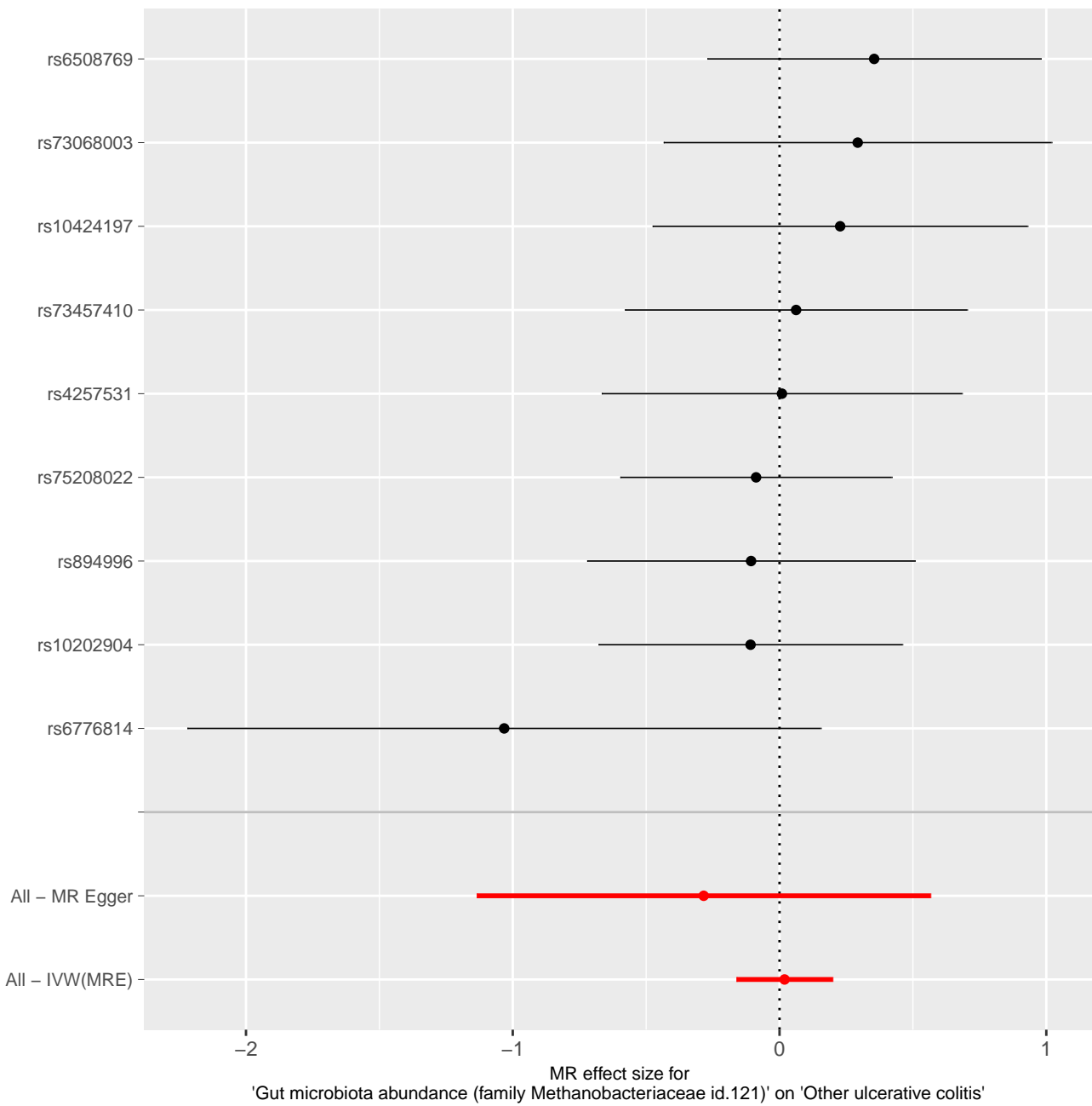


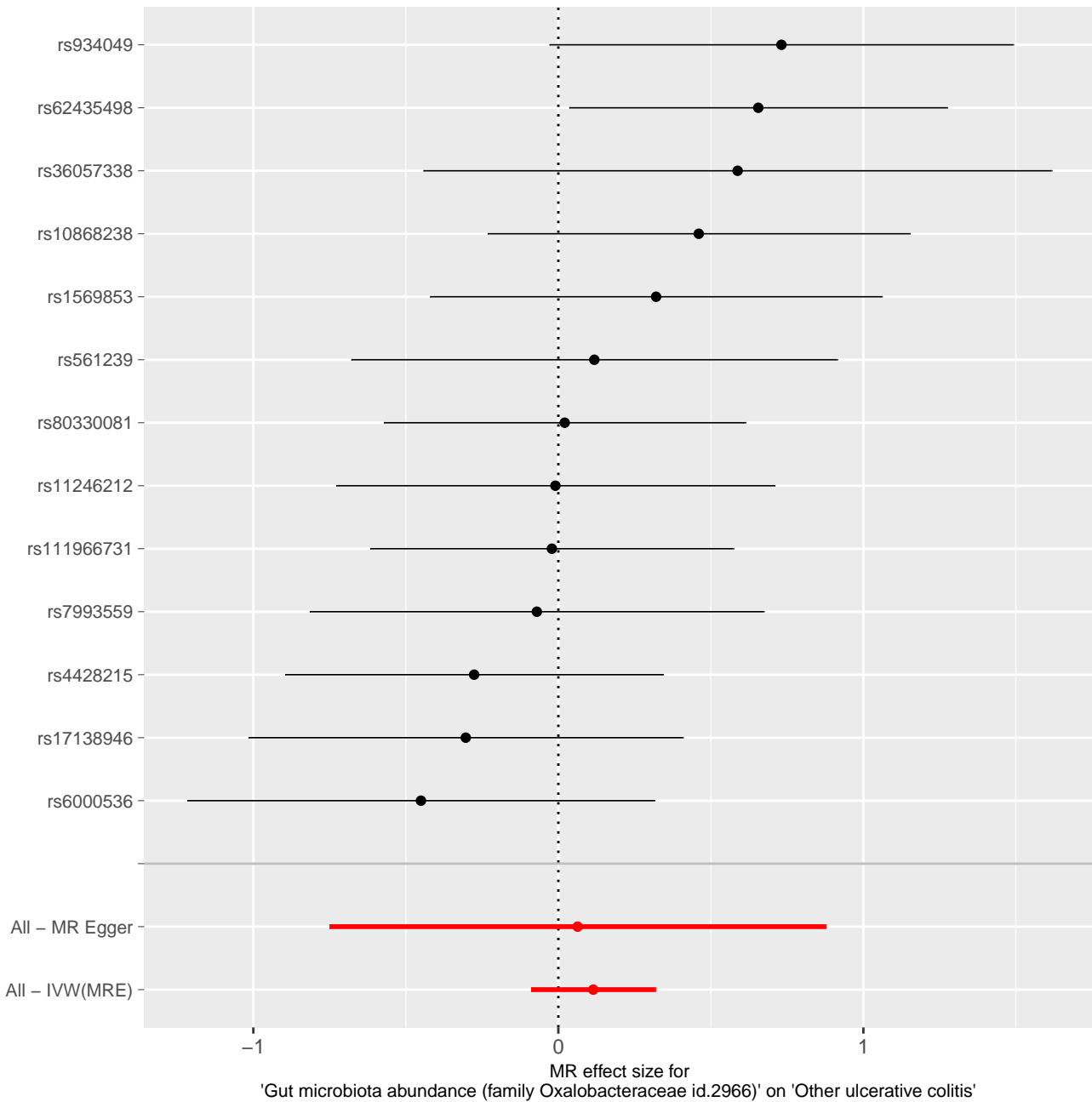


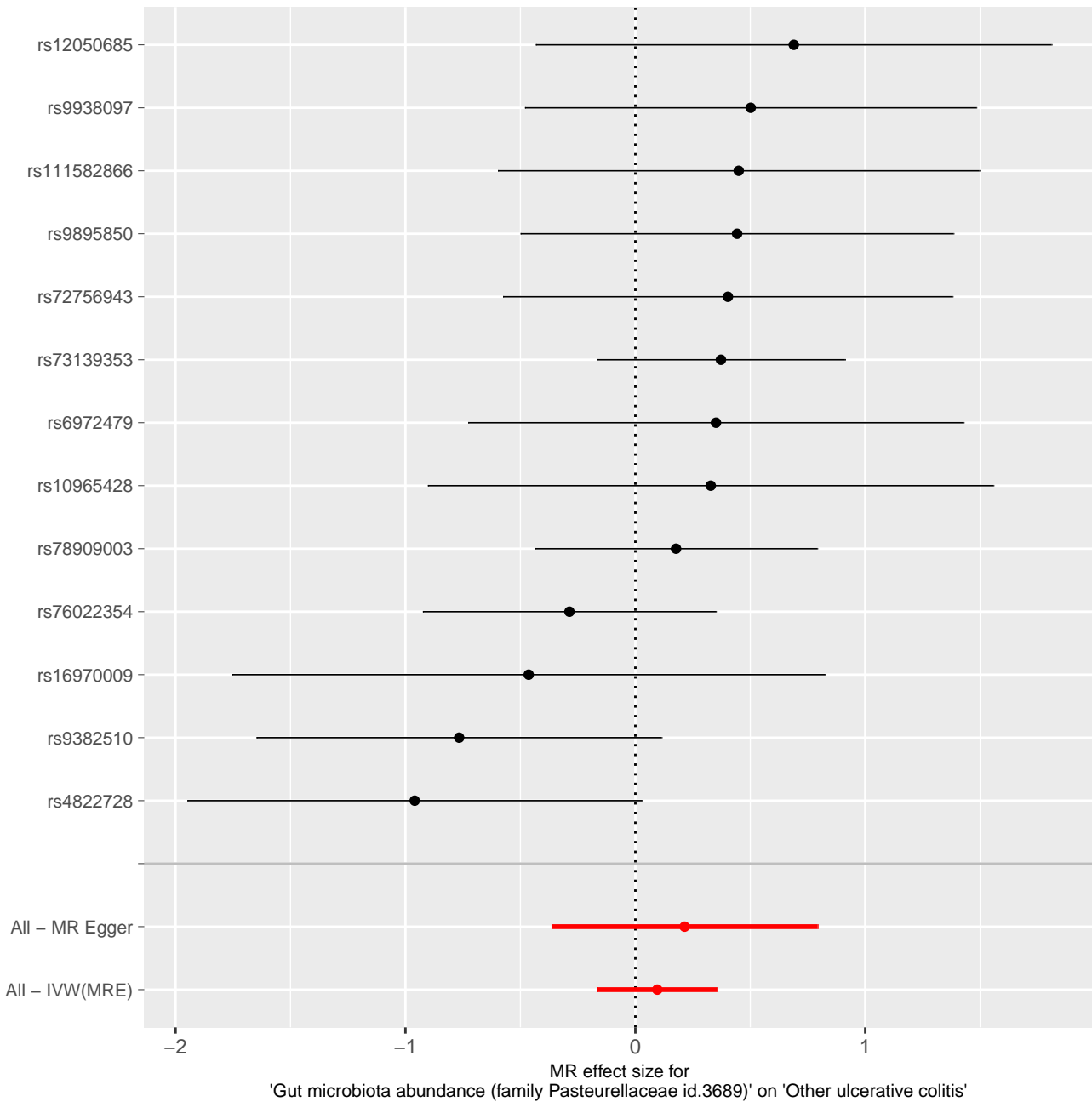


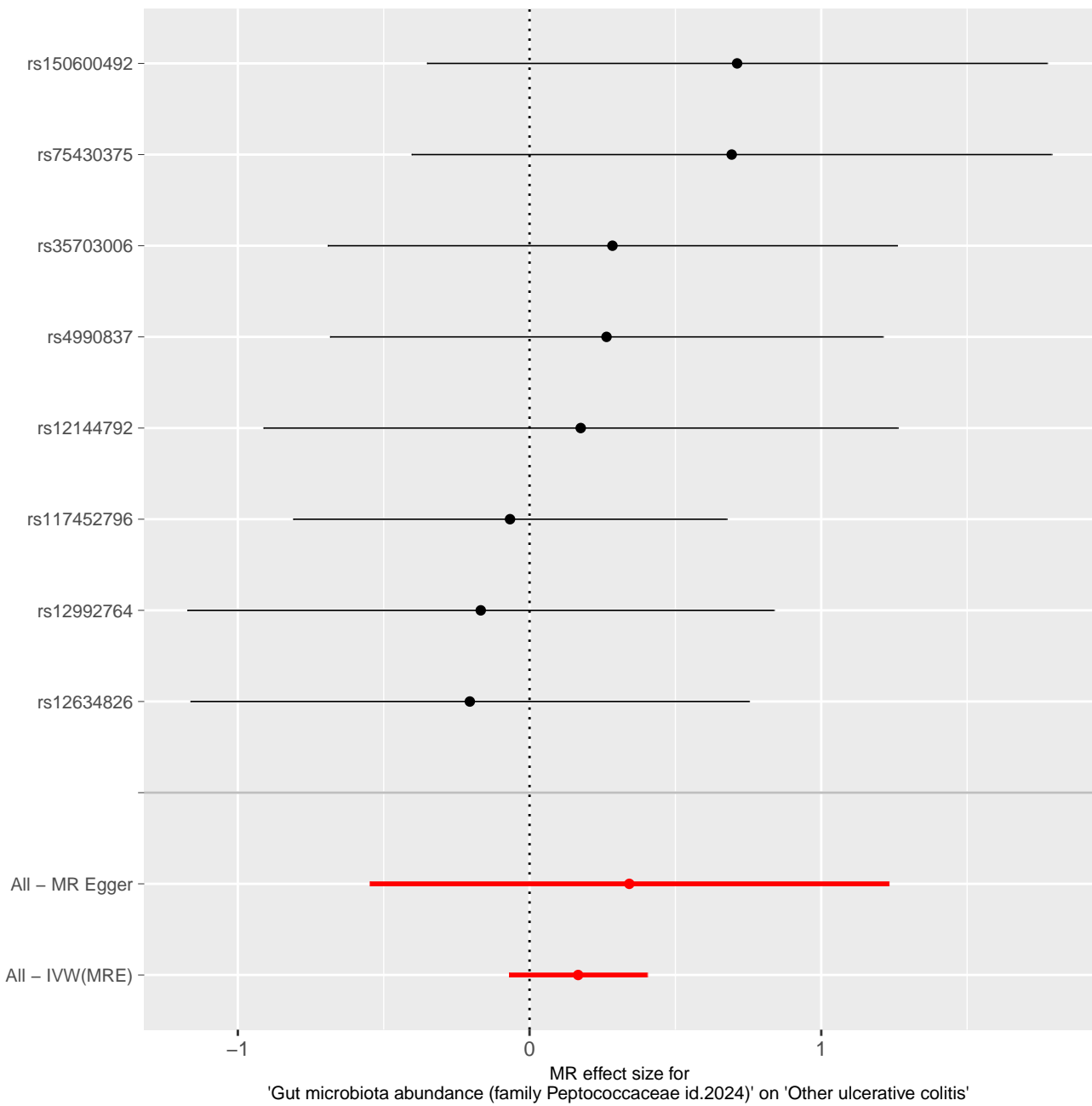


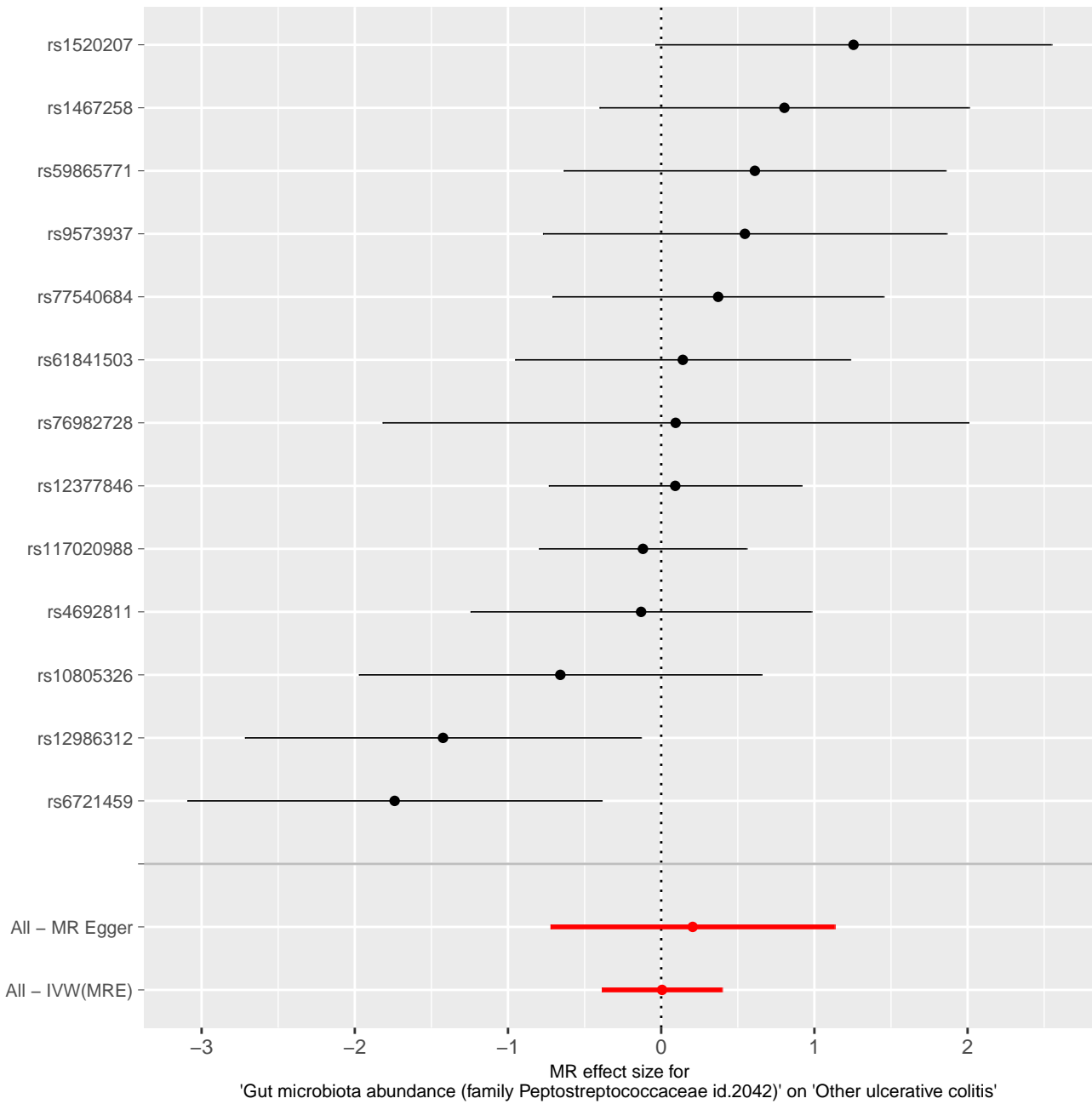


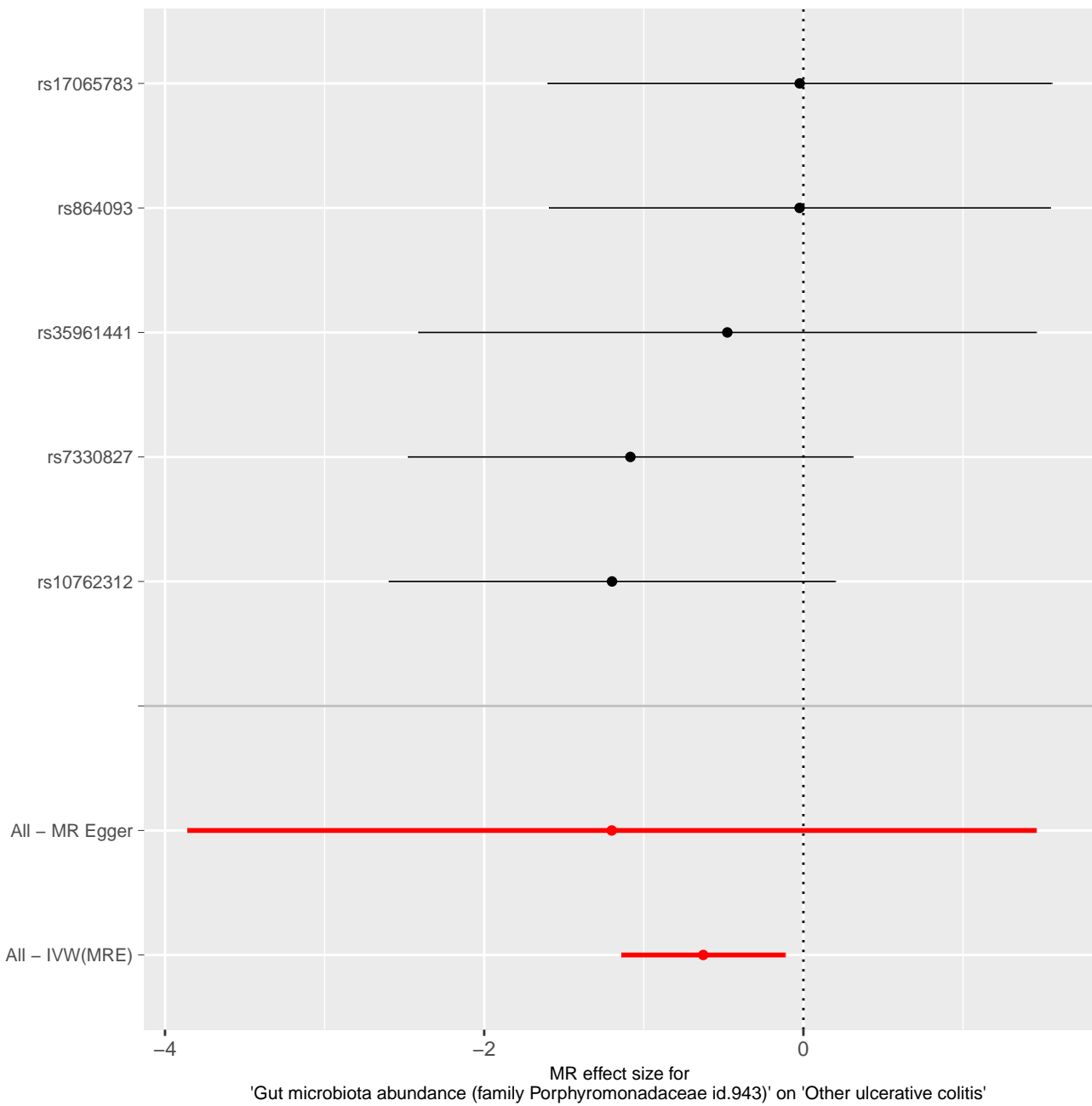


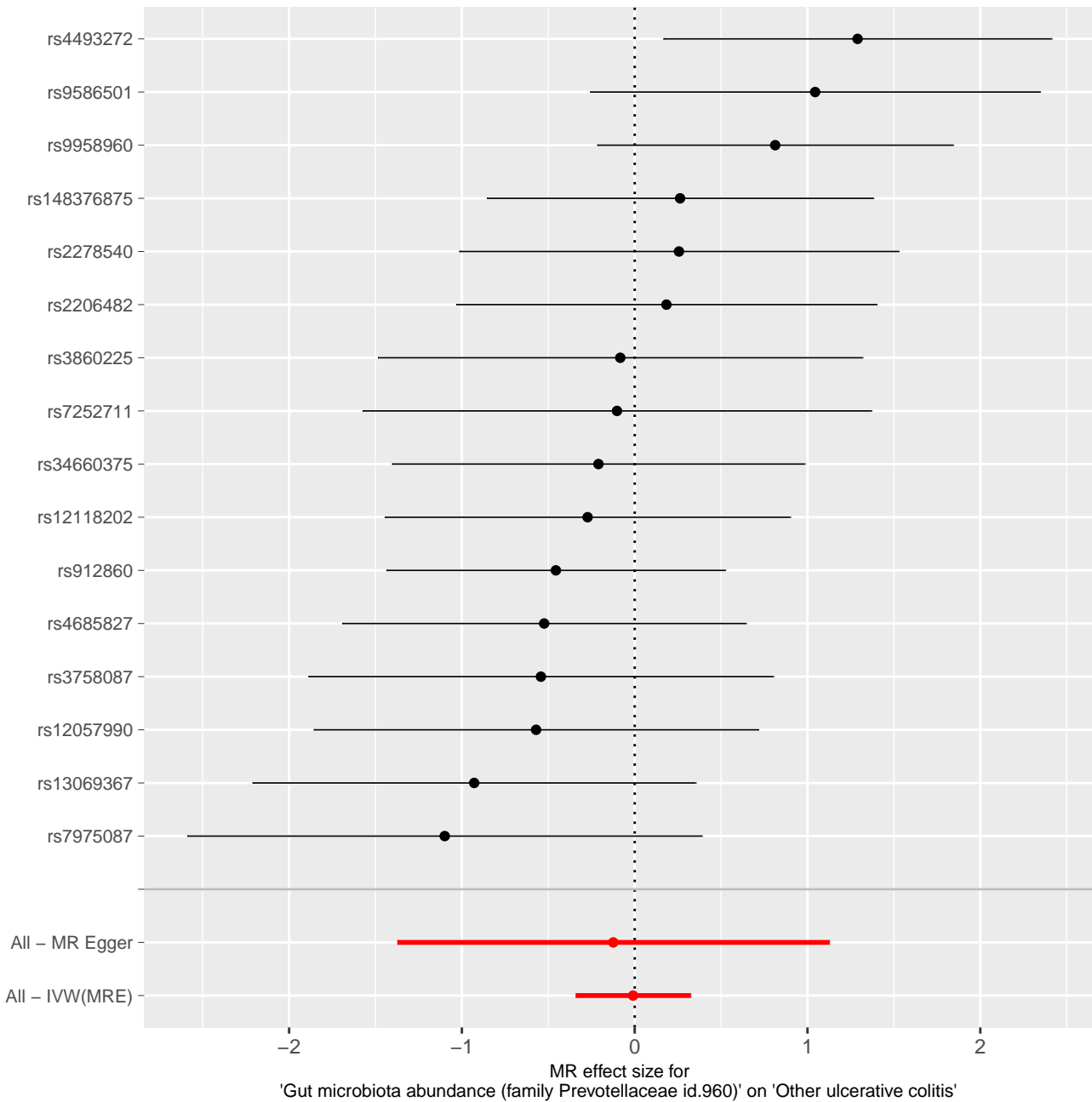


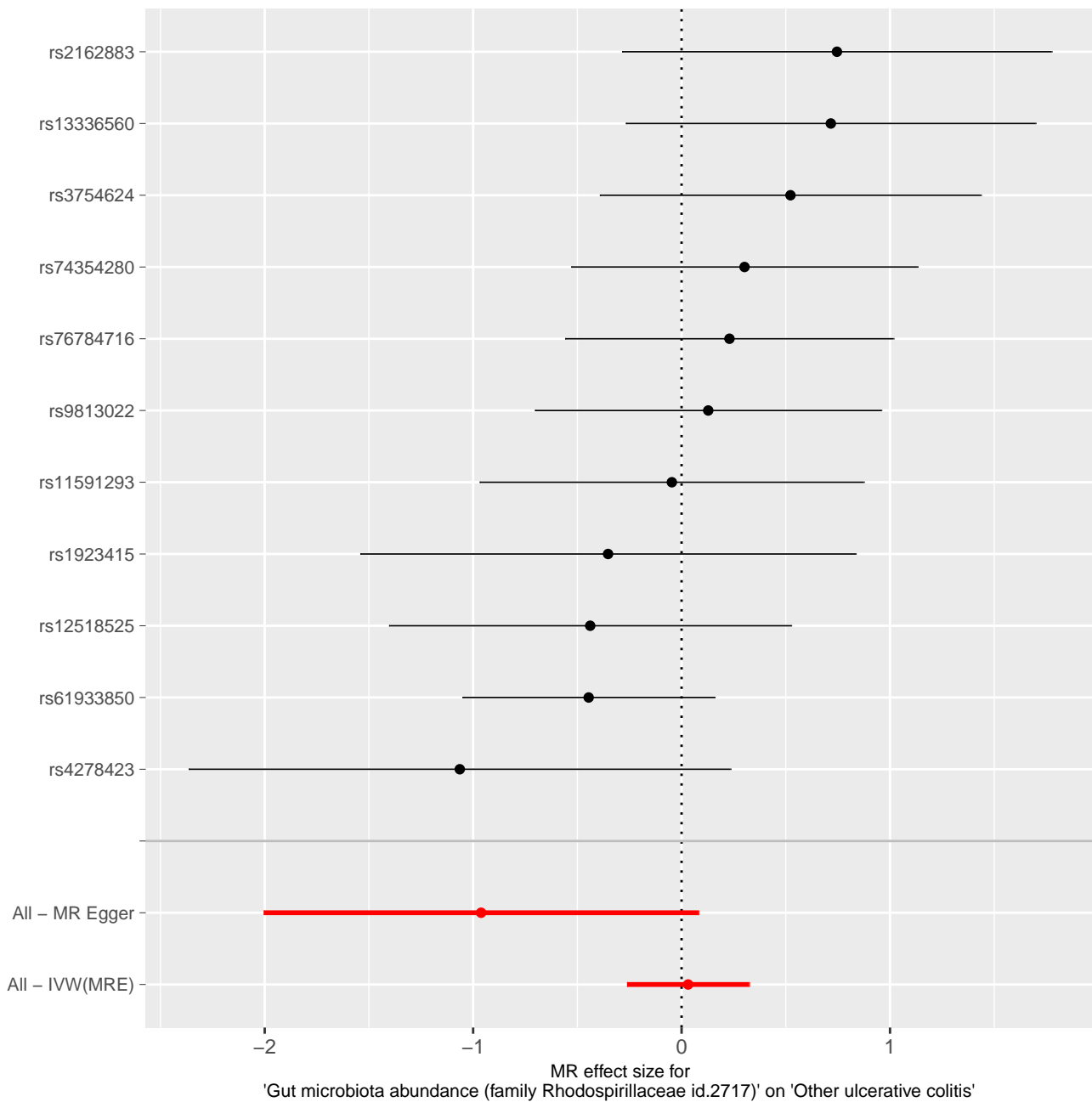


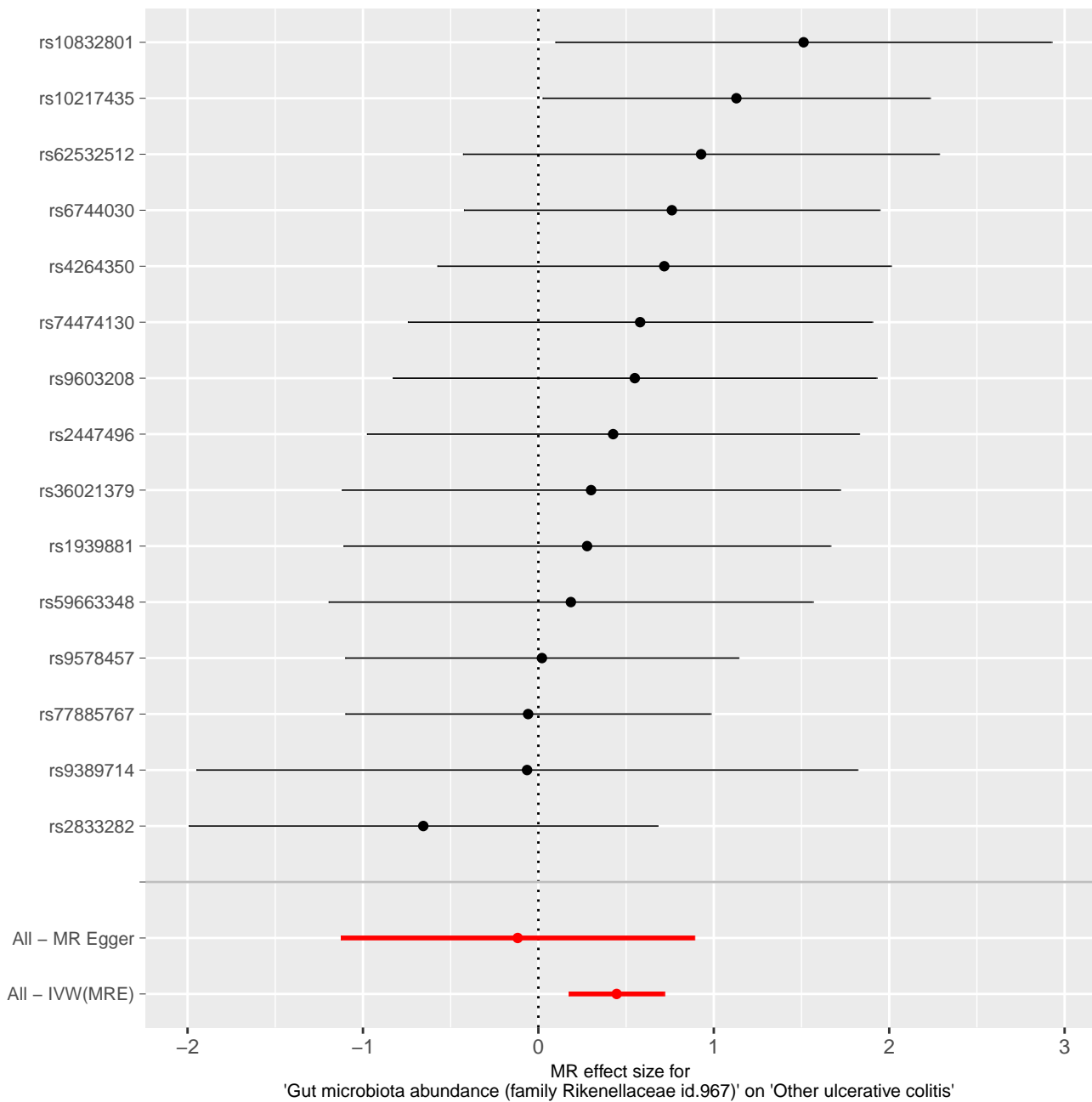


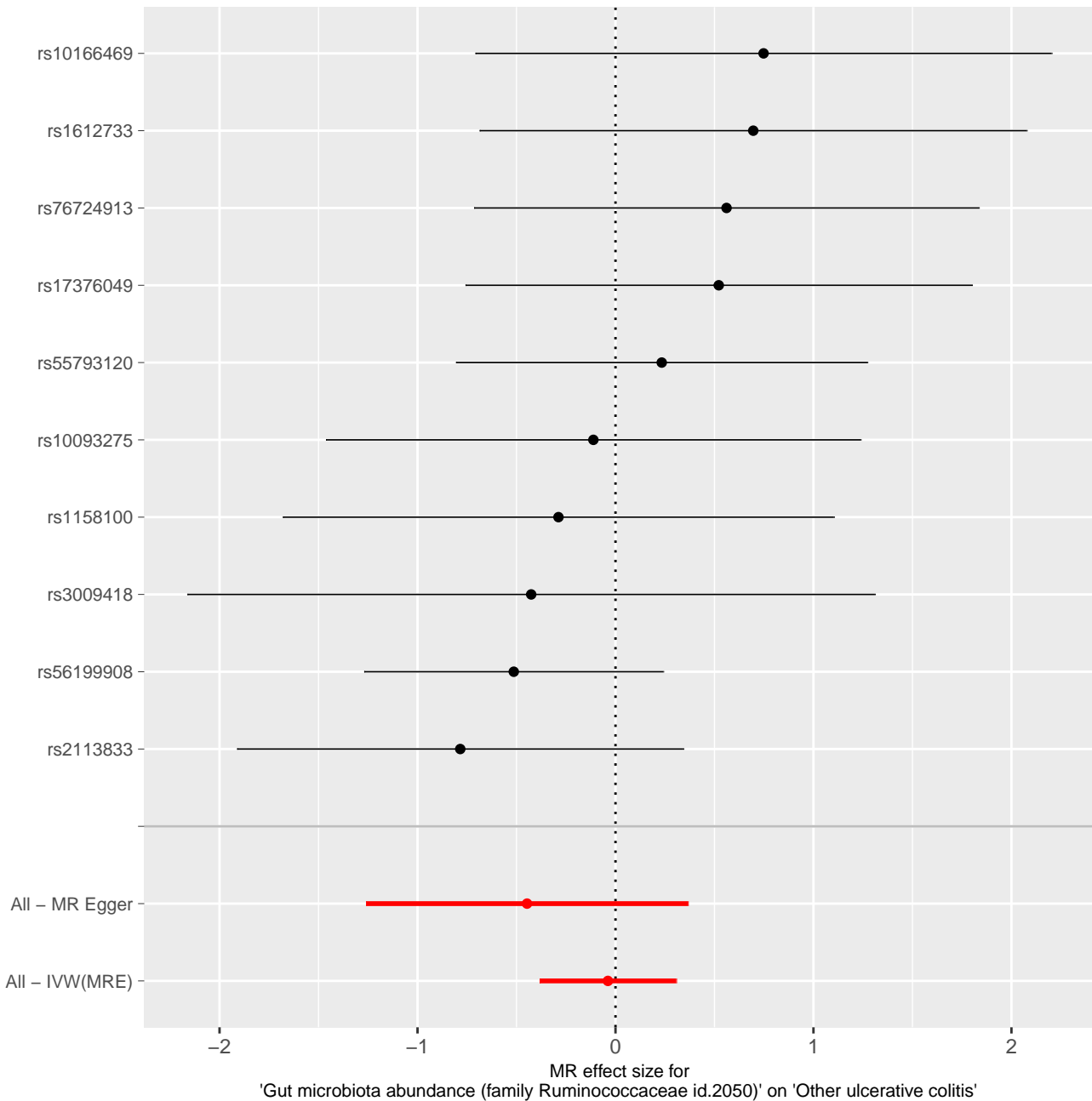


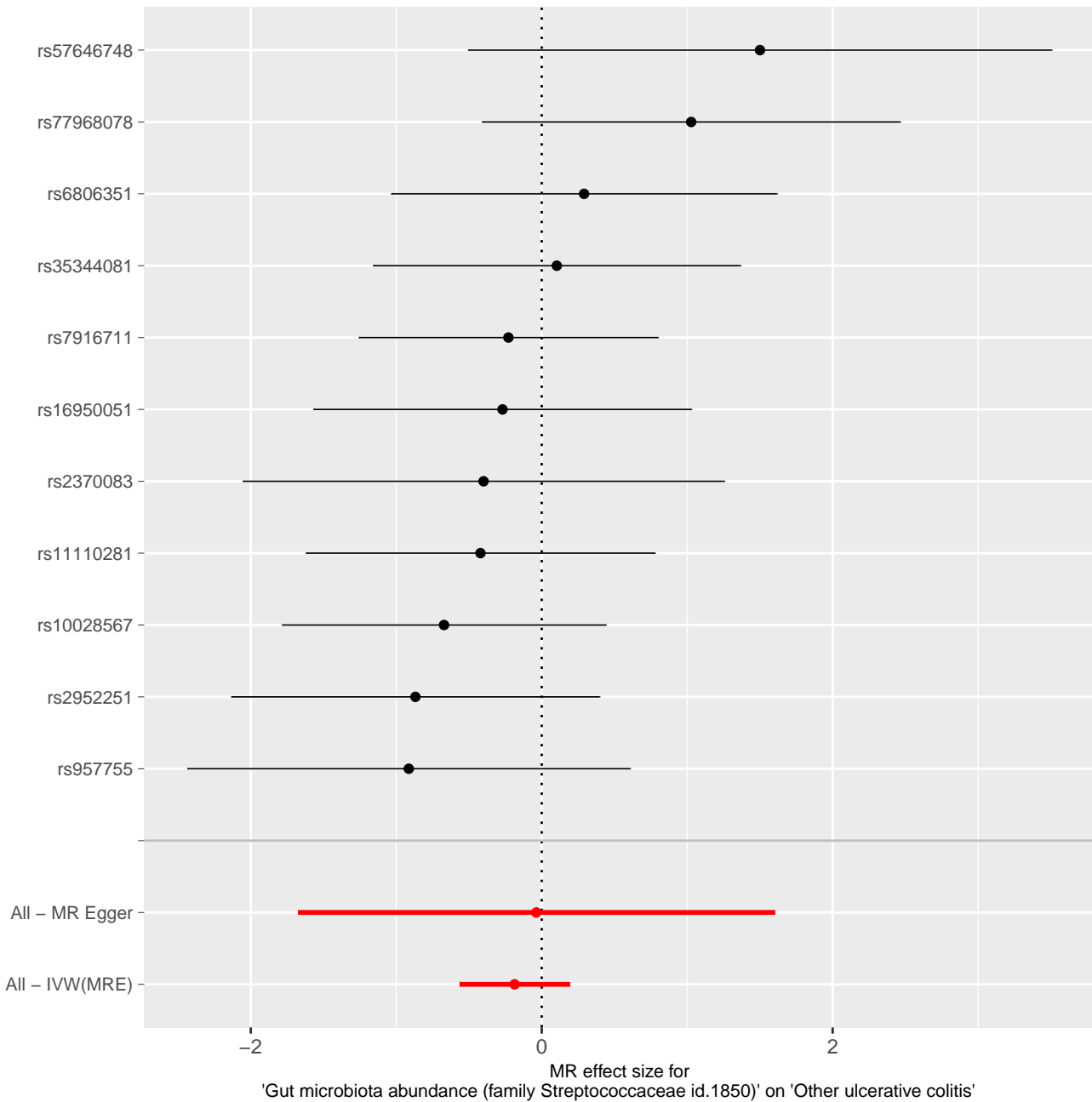


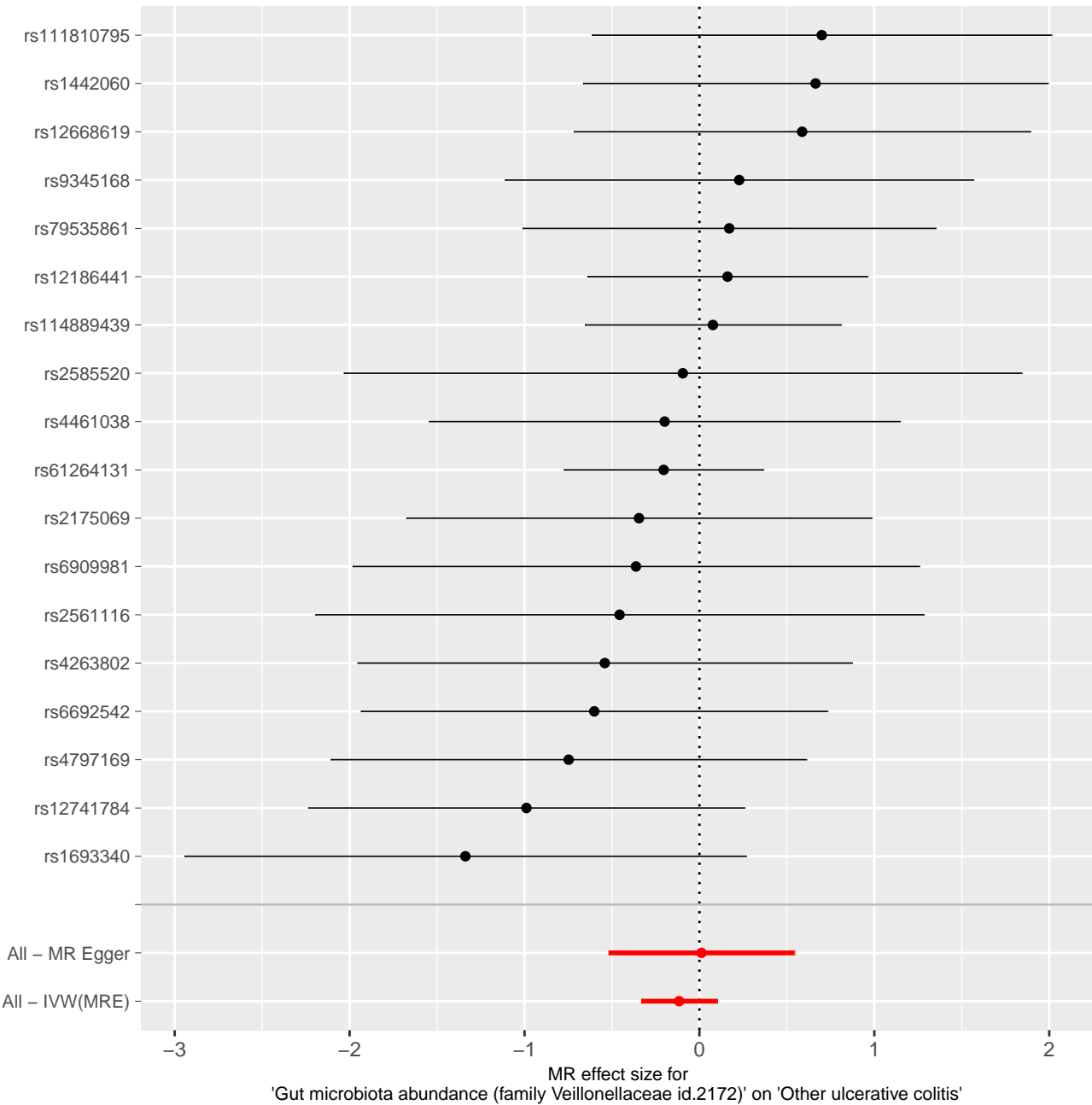


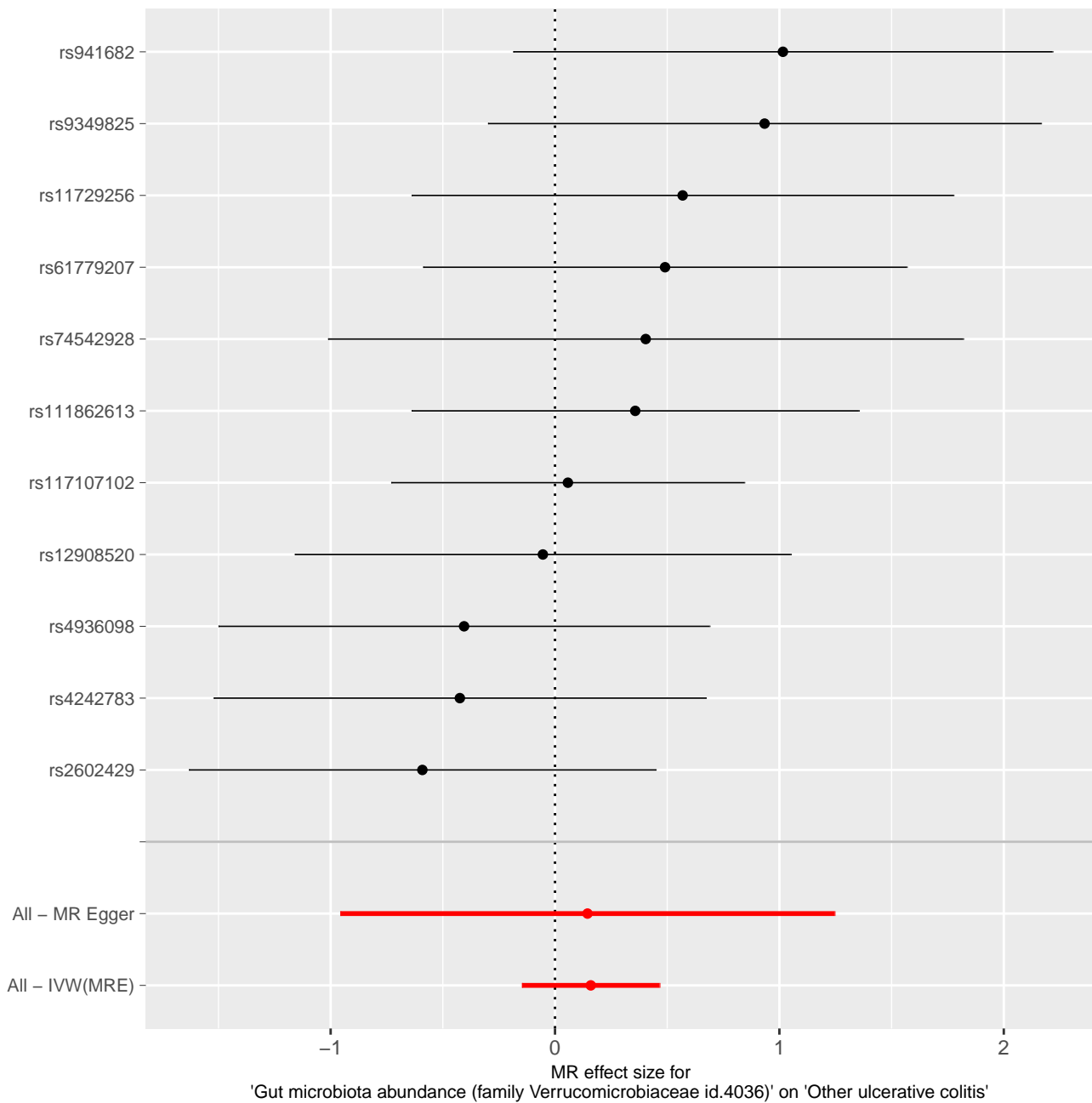


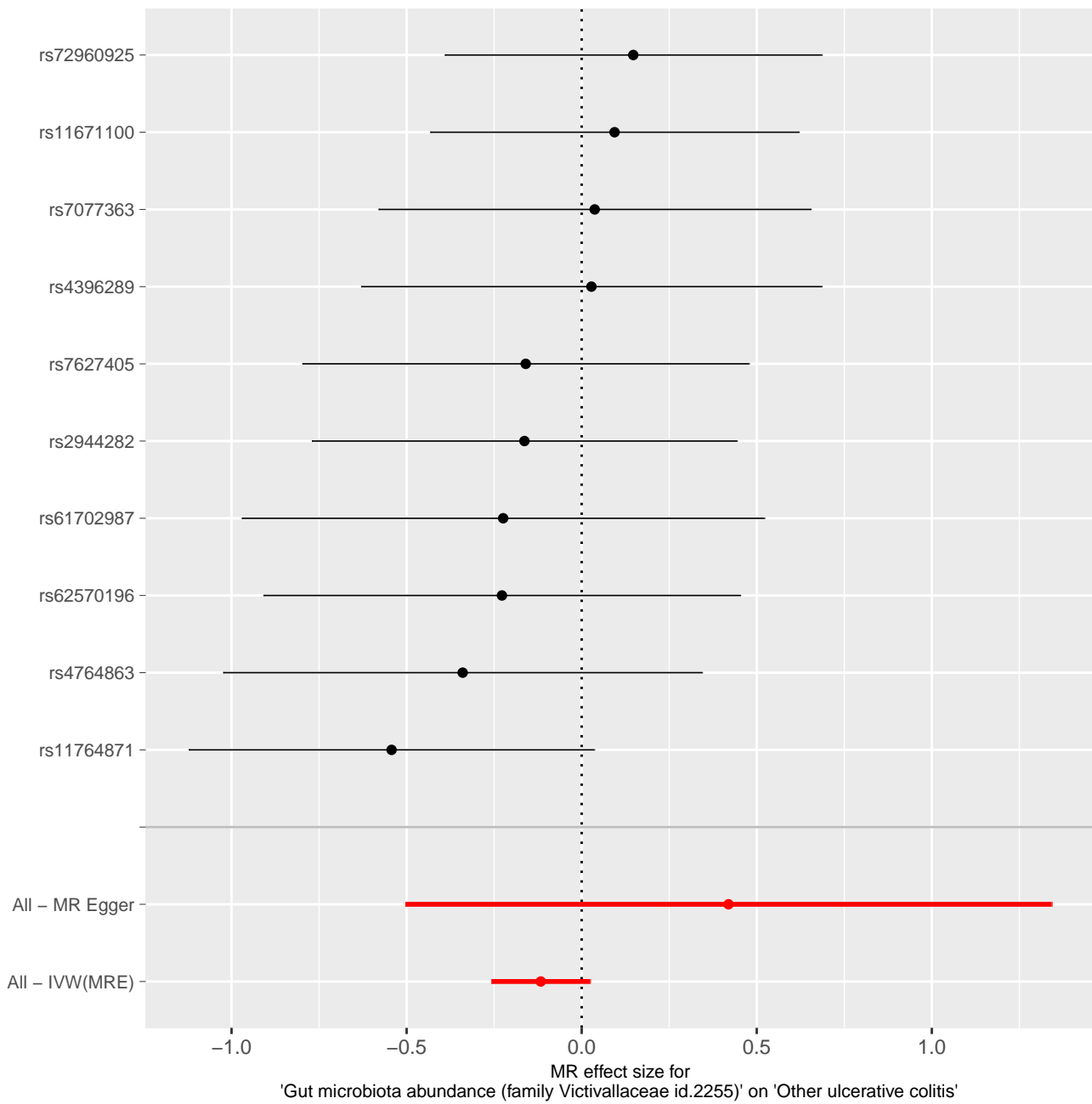


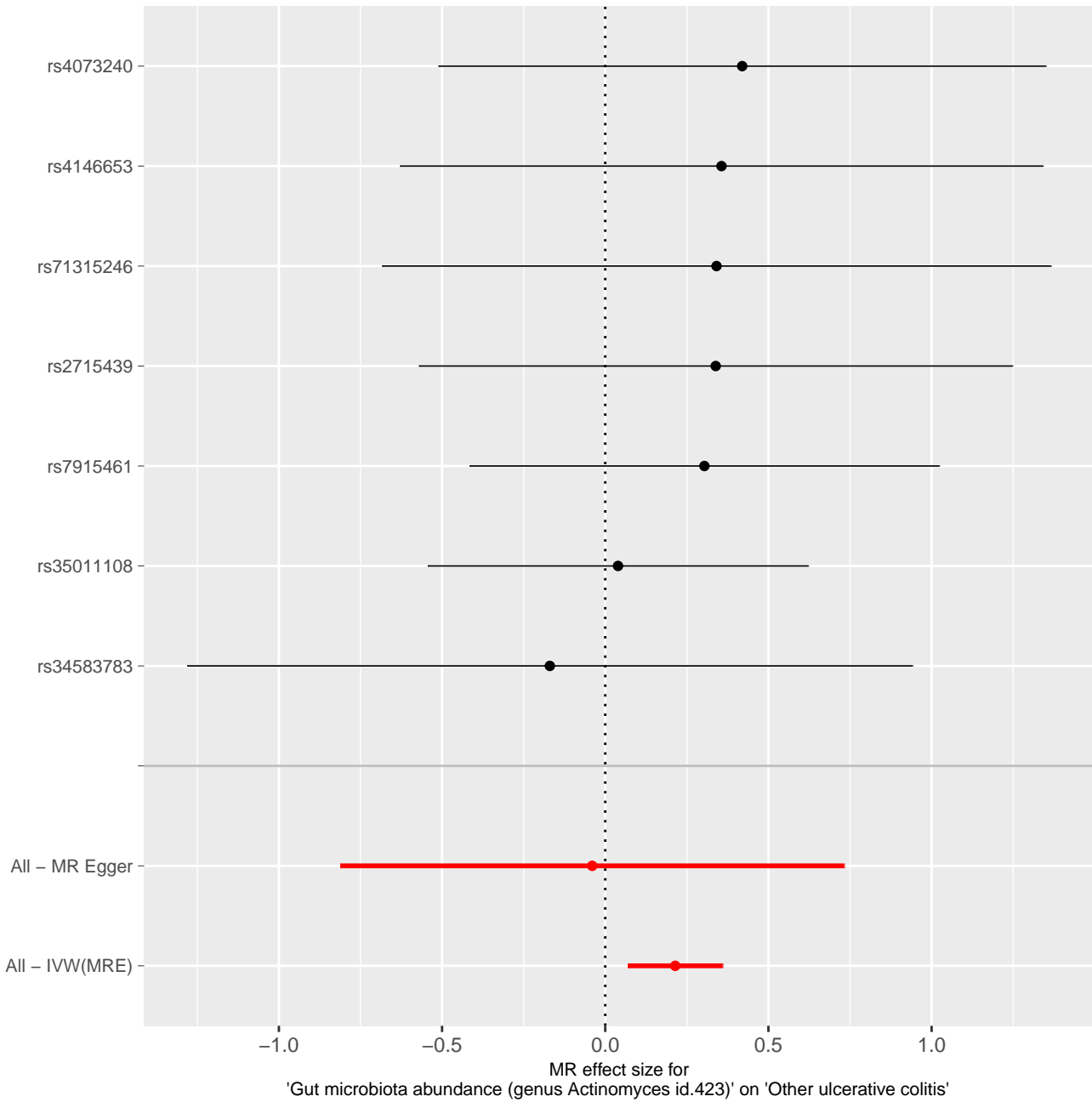


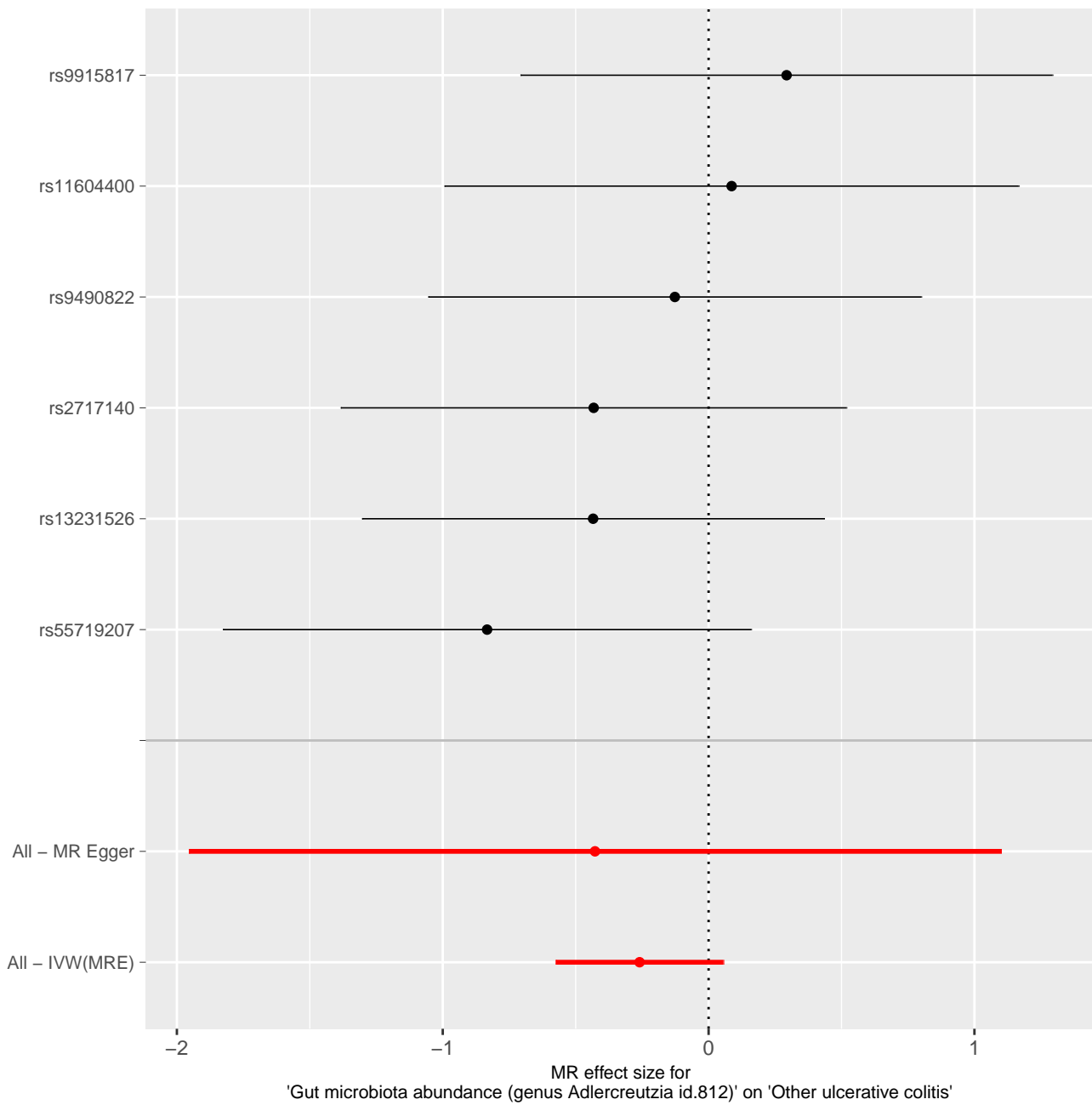




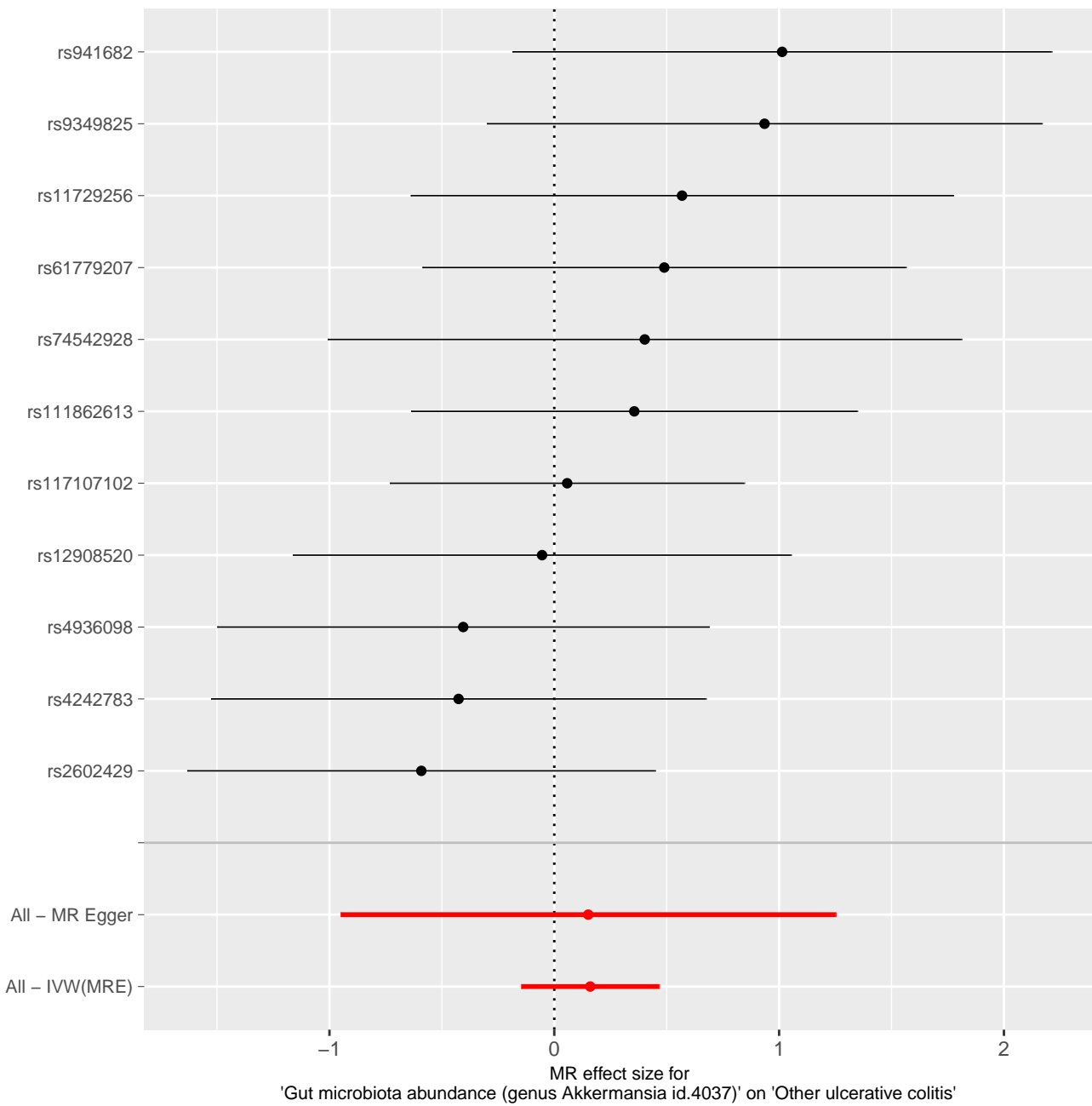


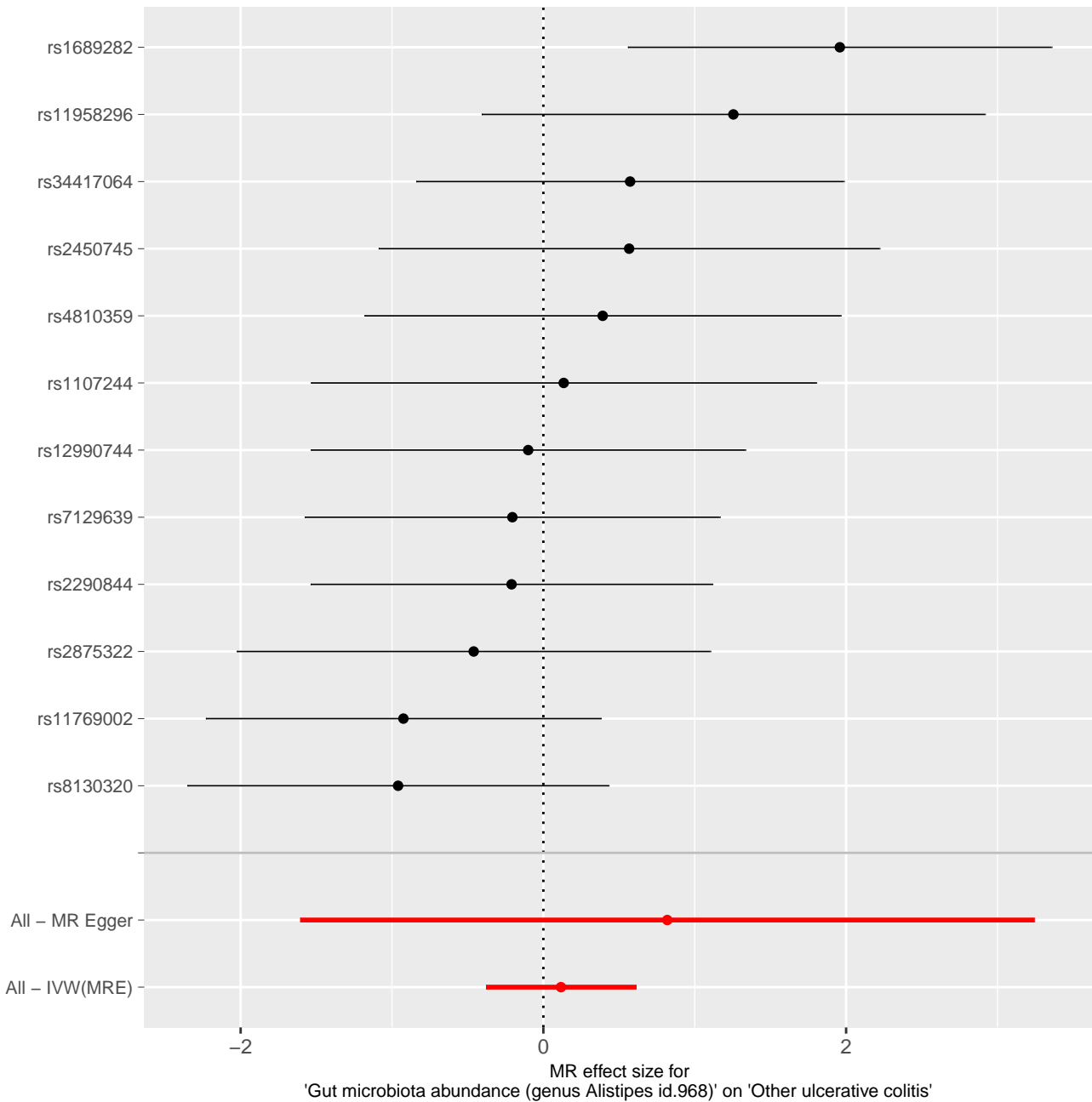


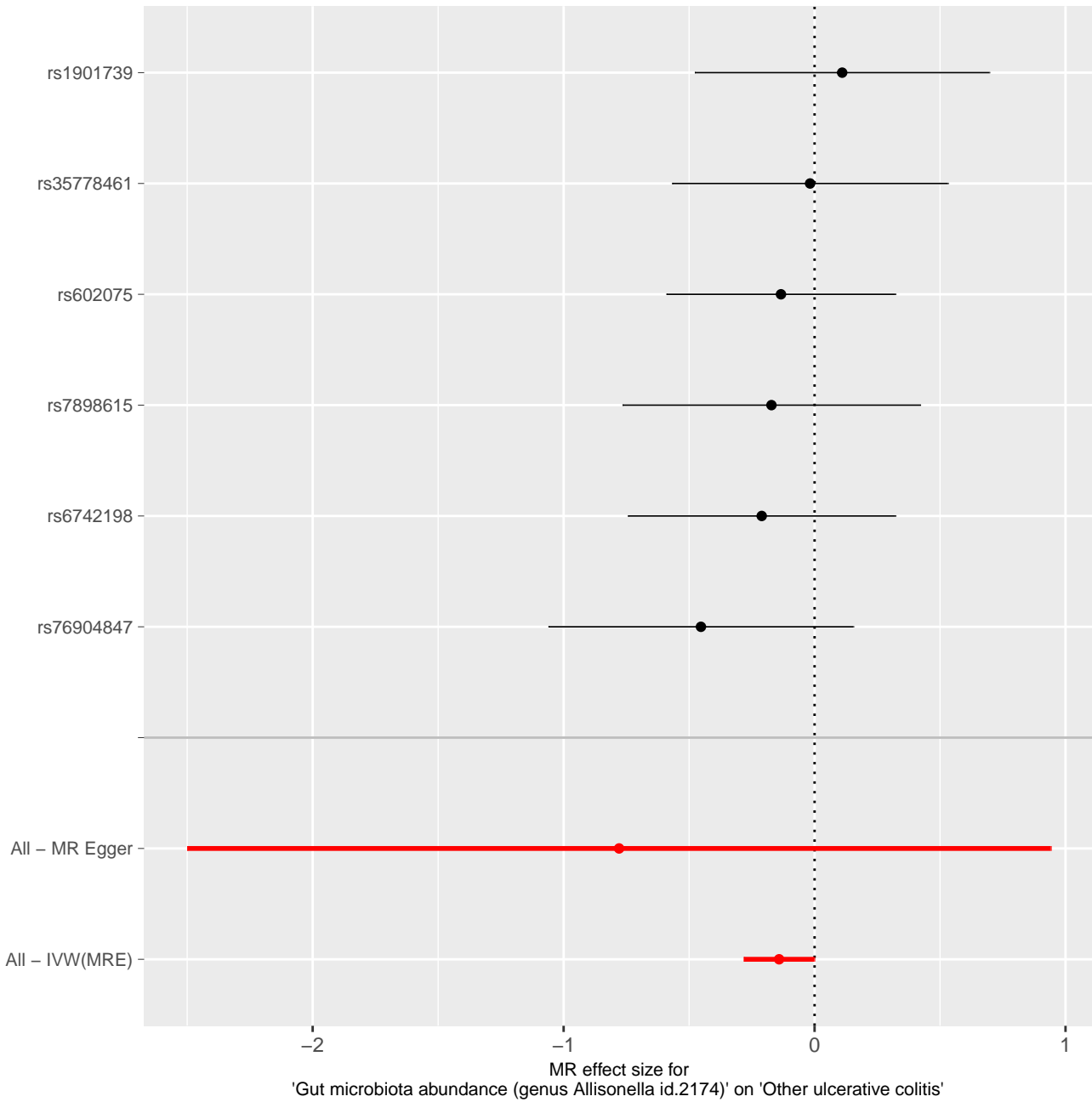


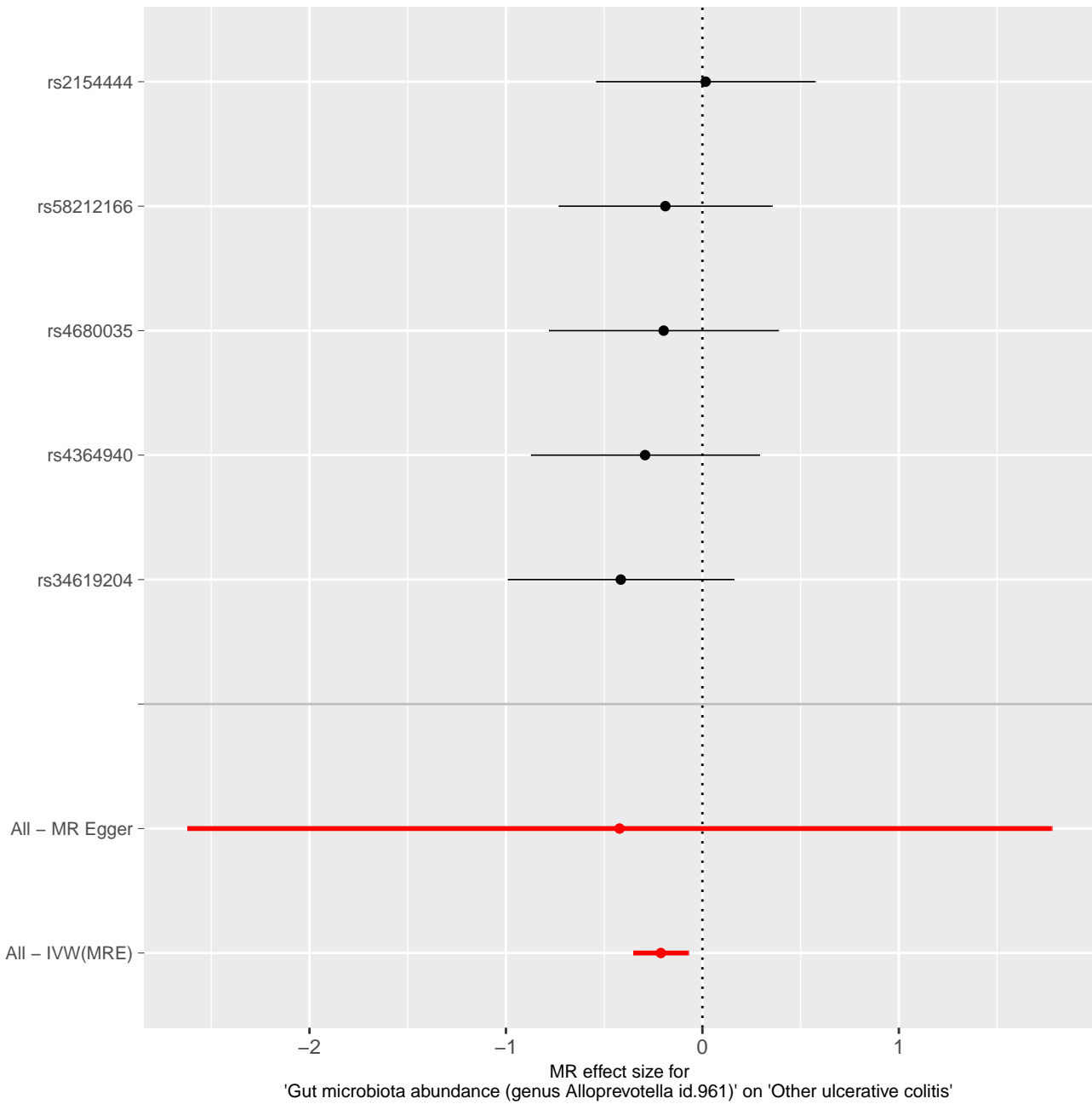


Batch 1235 : Gut microbiota abundance (genus Akkermansia id.4037) on Other ulcerative colitis

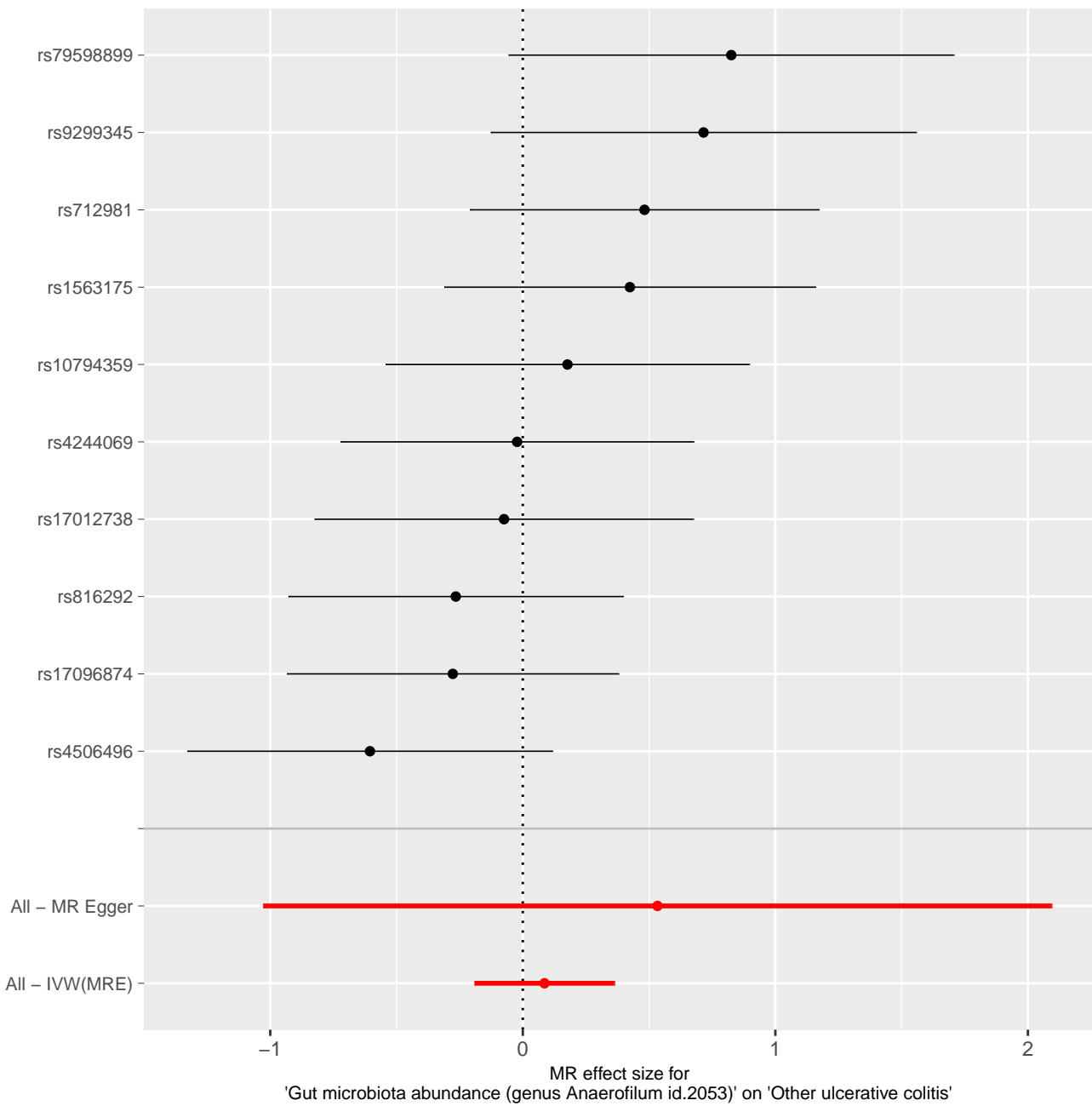




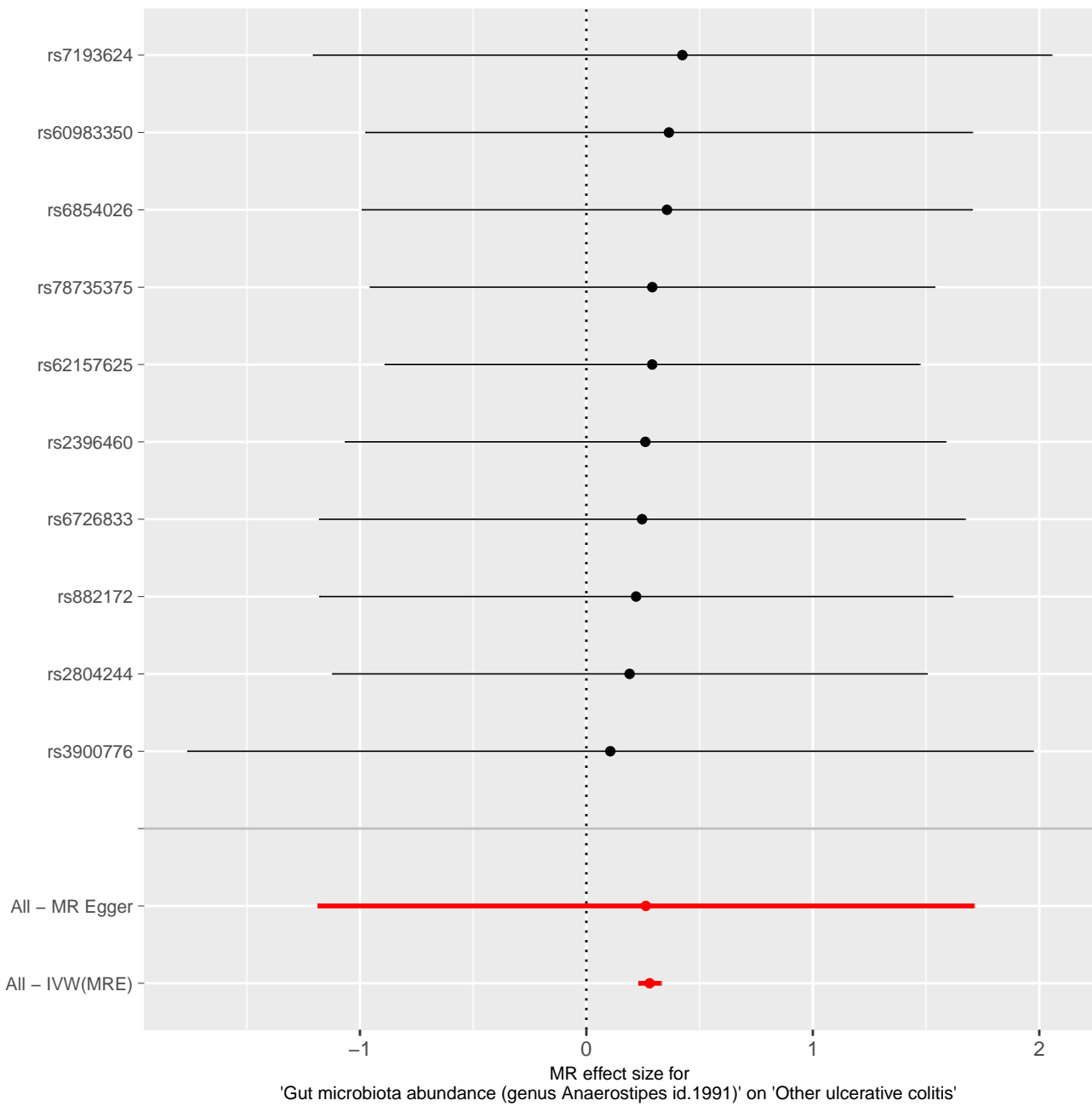


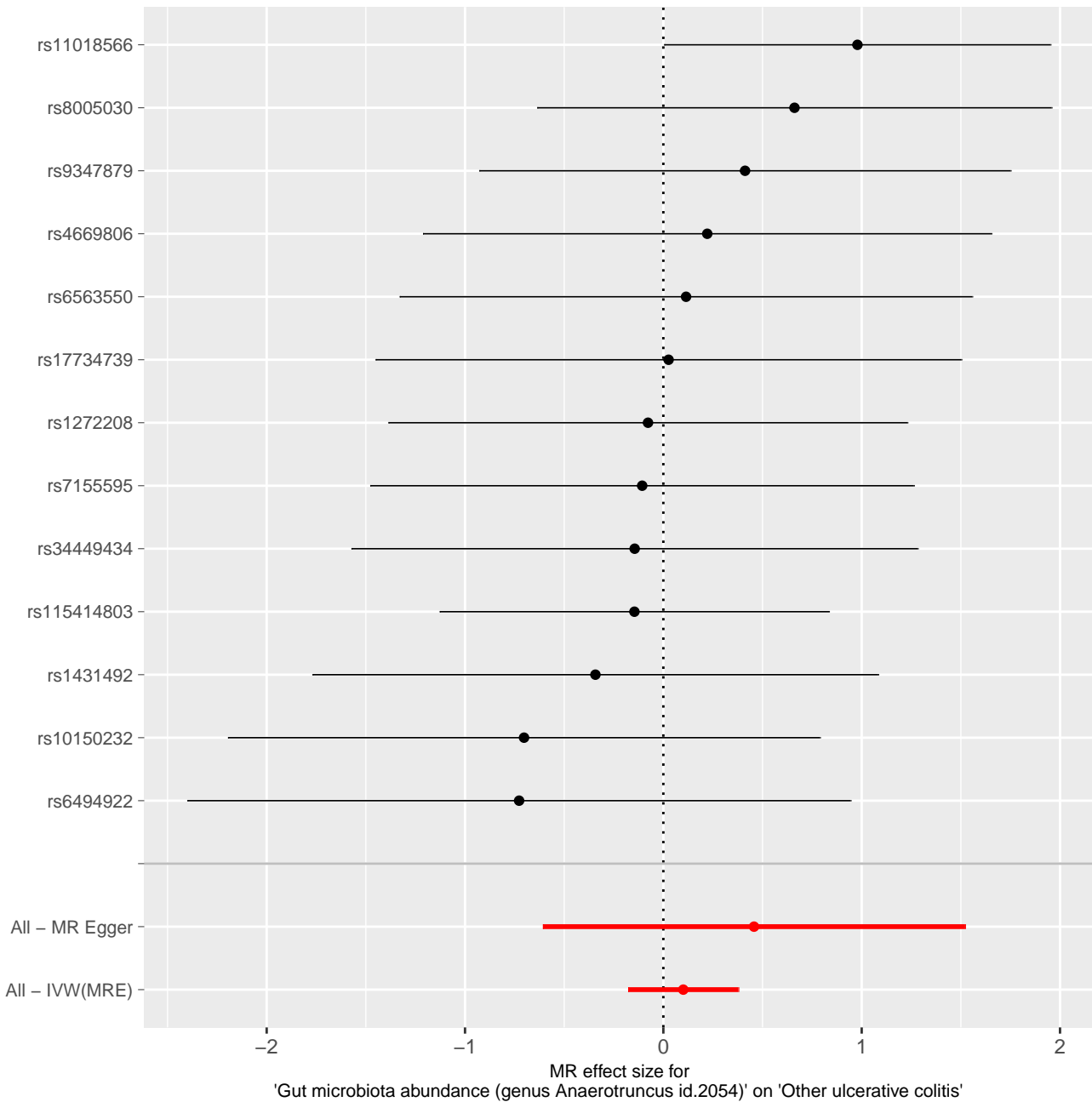


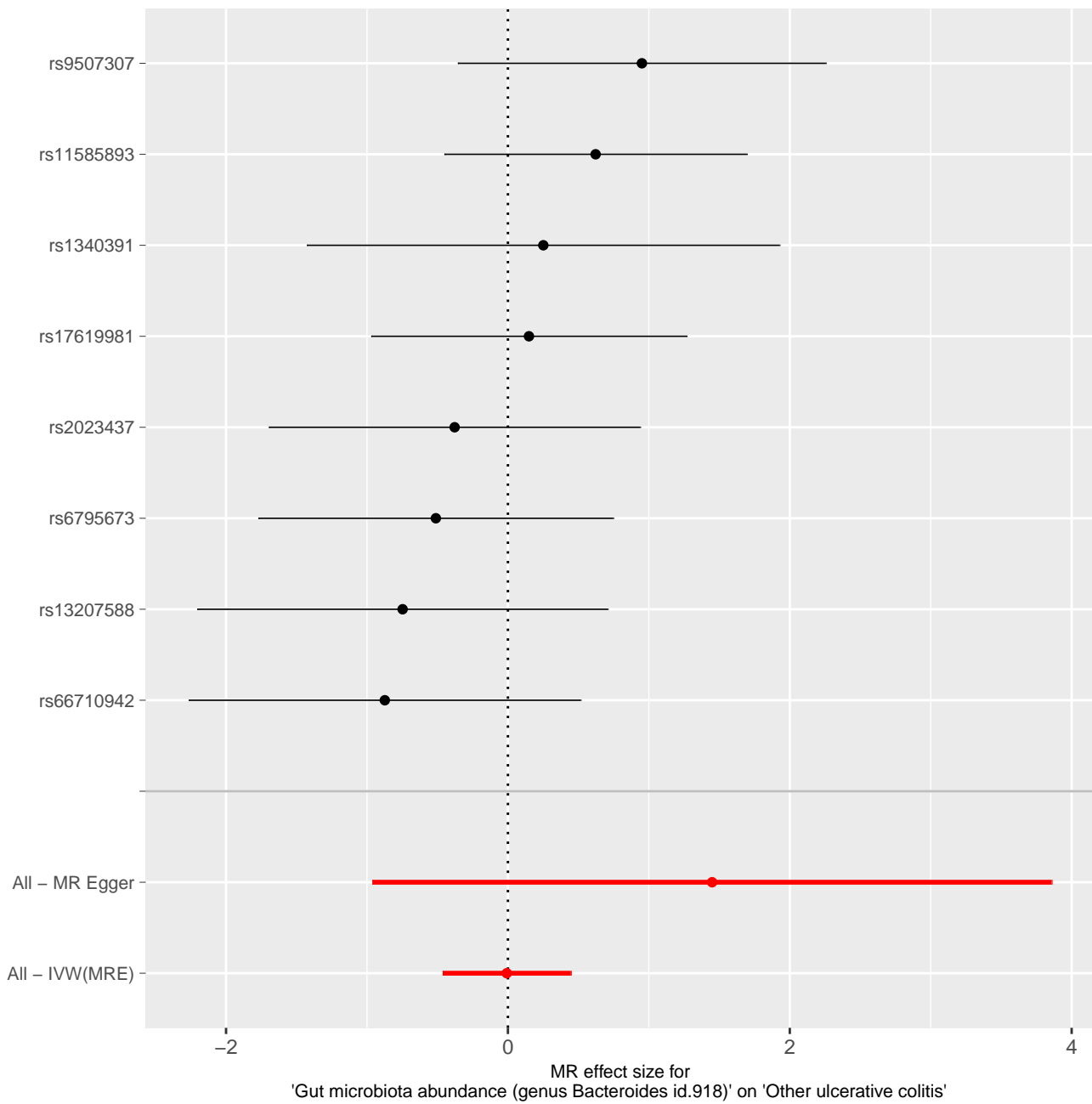
Batch 1239 : Gut microbiota abundance (genus Anaerofilum id.2053) on Other ulcerative colitis



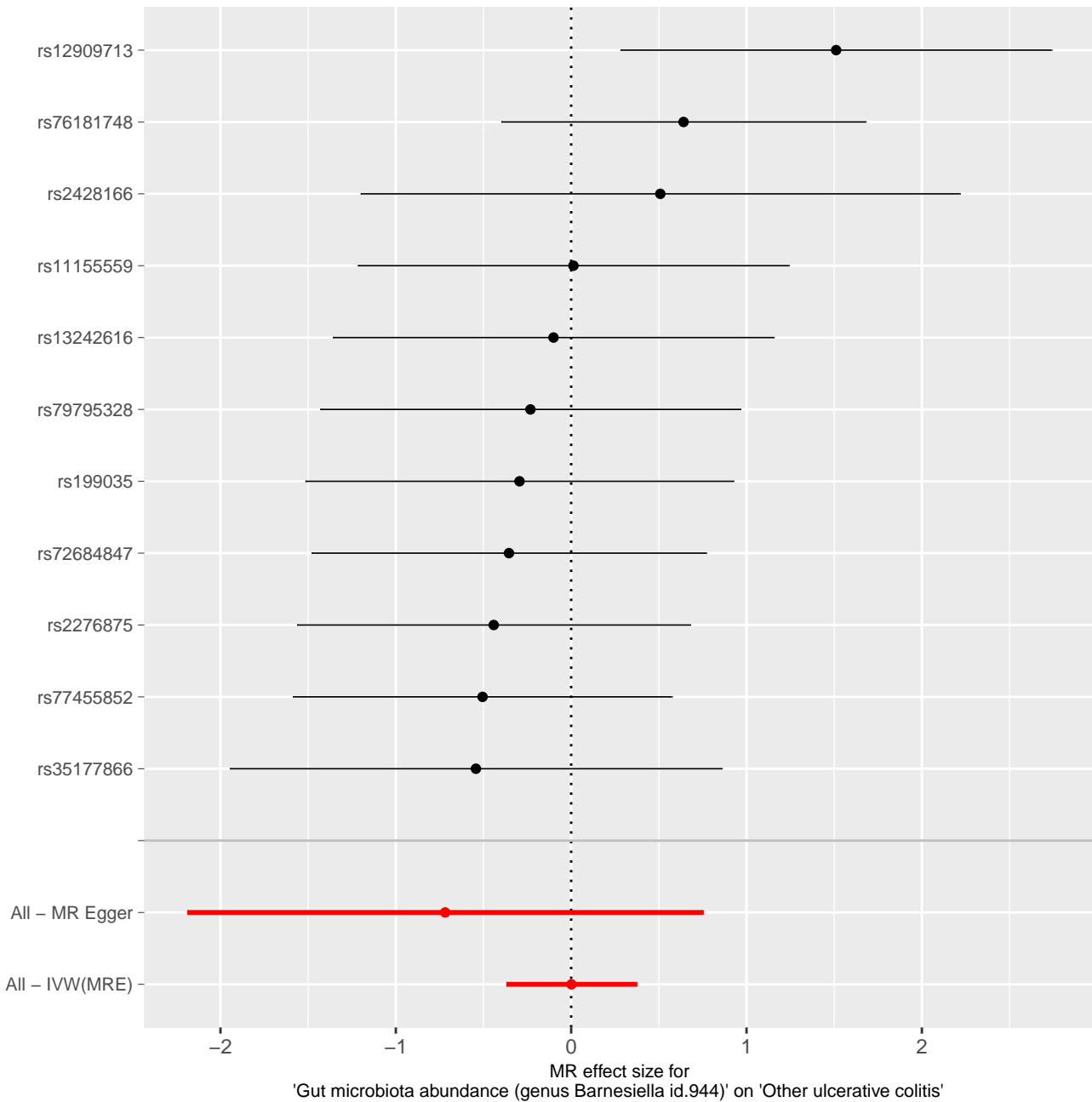
Batch 1240 : Gut microbiota abundance (genus Anaerostipes id.1991) on Other ulcerative colitis

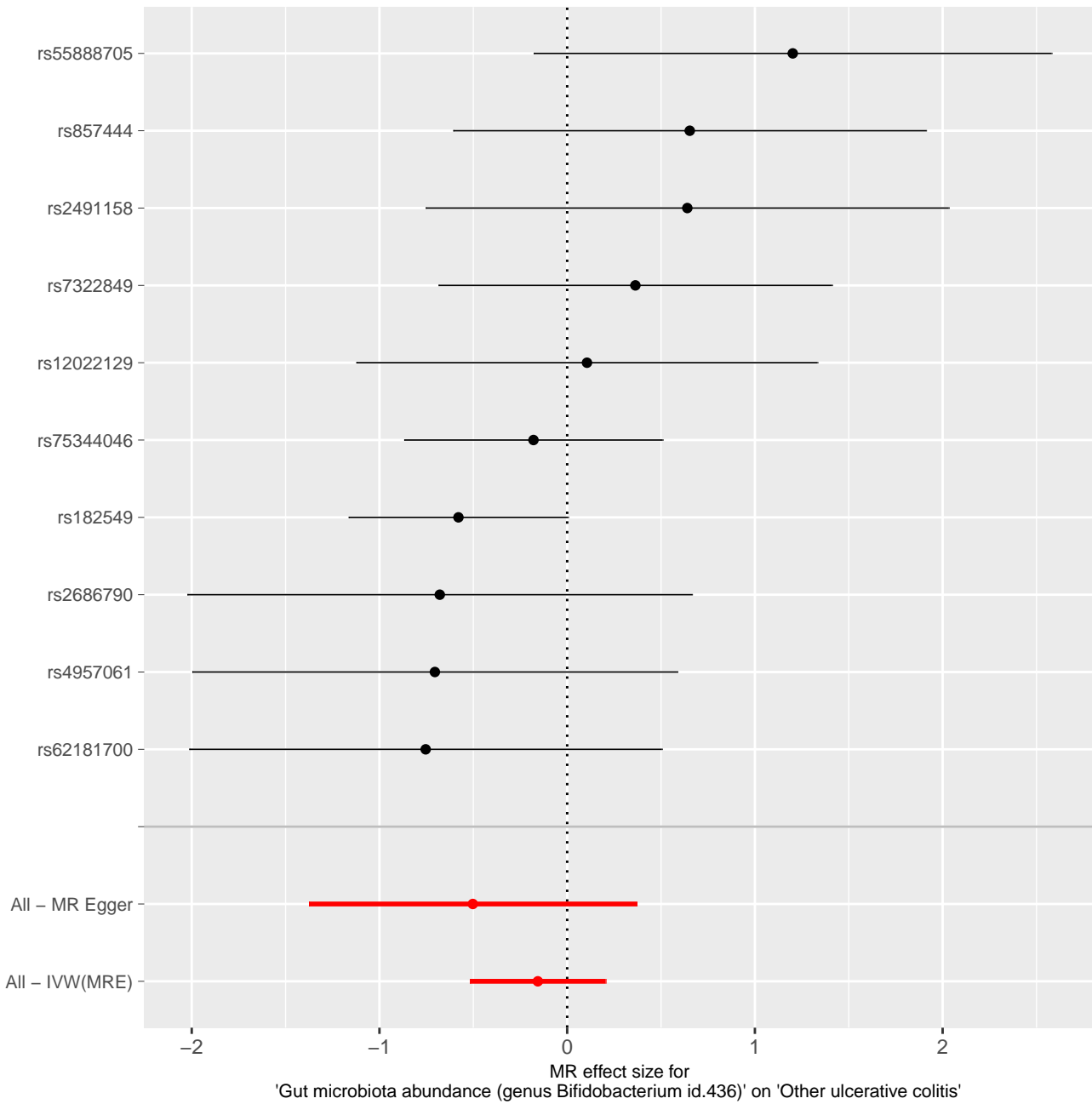




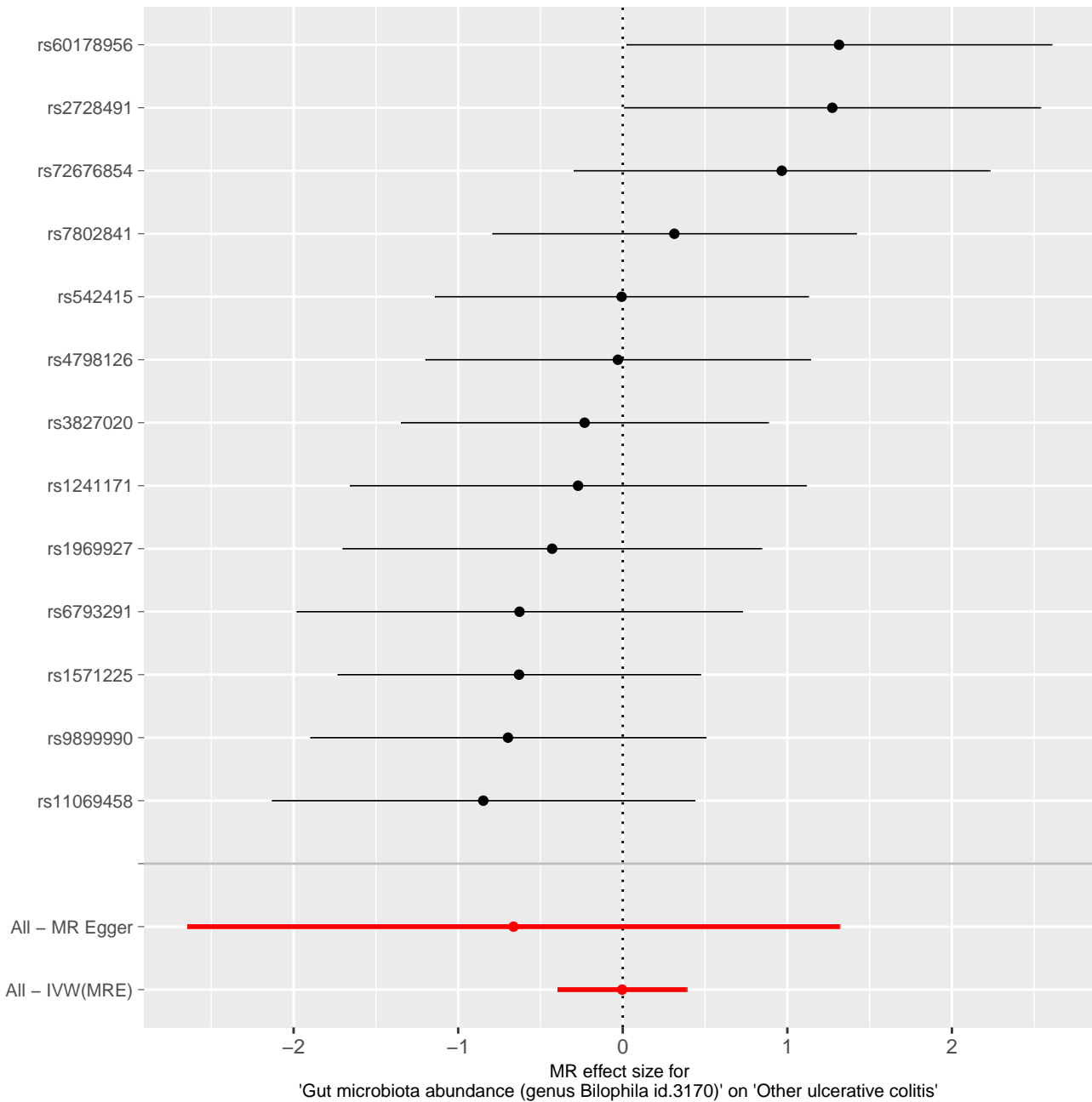


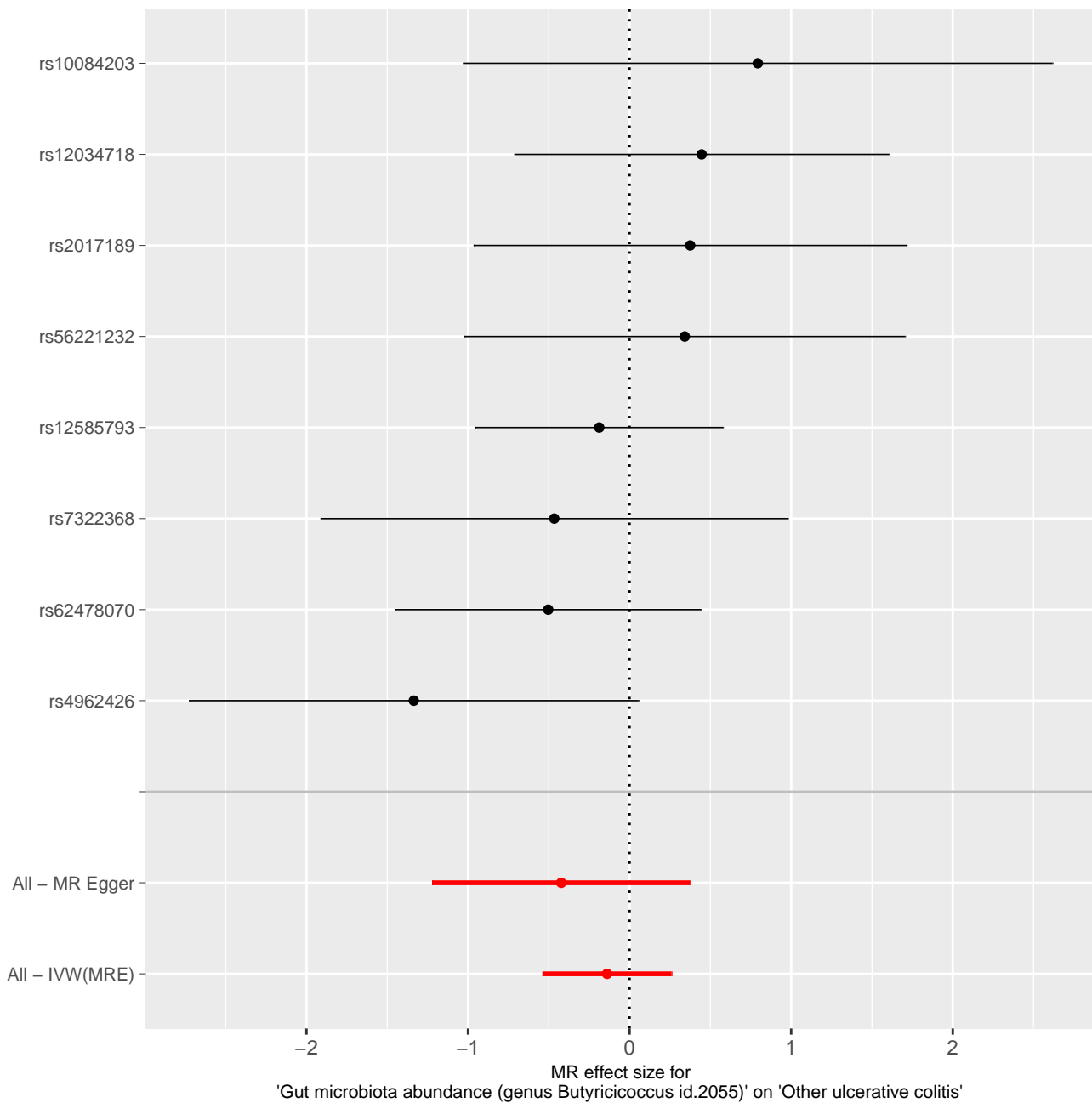
Batch 1243 : Gut microbiota abundance (genus *Barnesiella* id.944) on Other ulcerative colitis

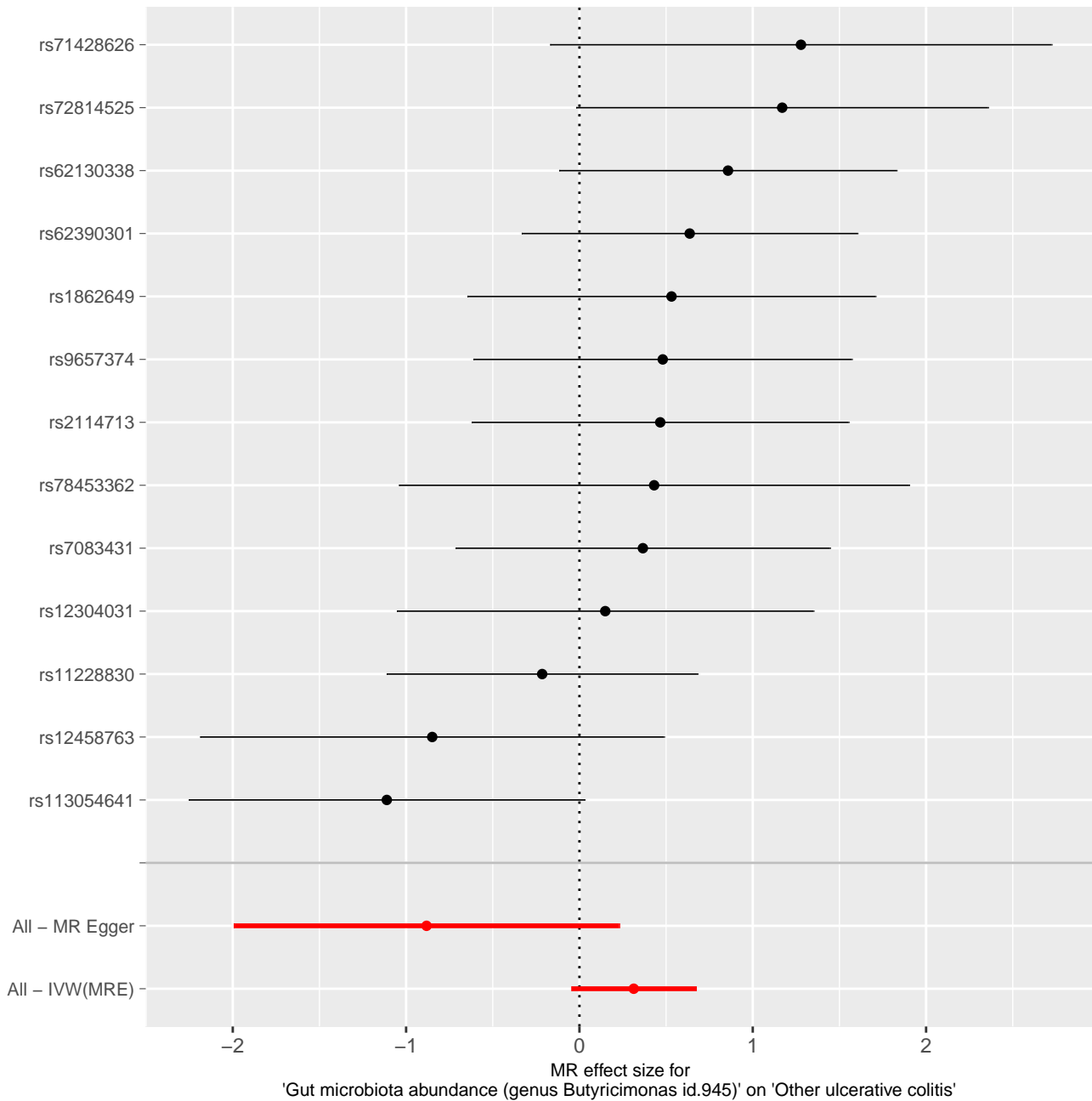


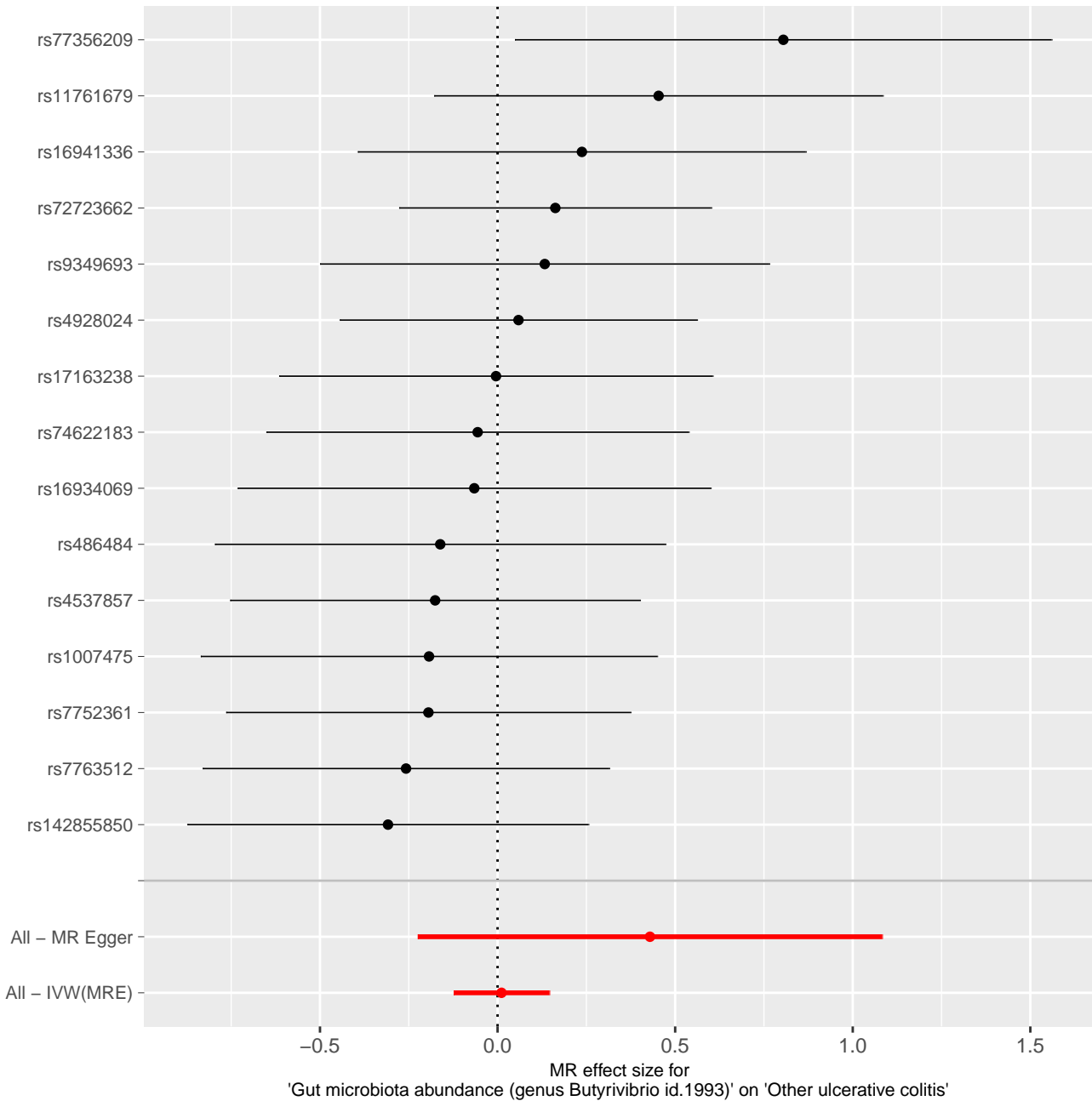


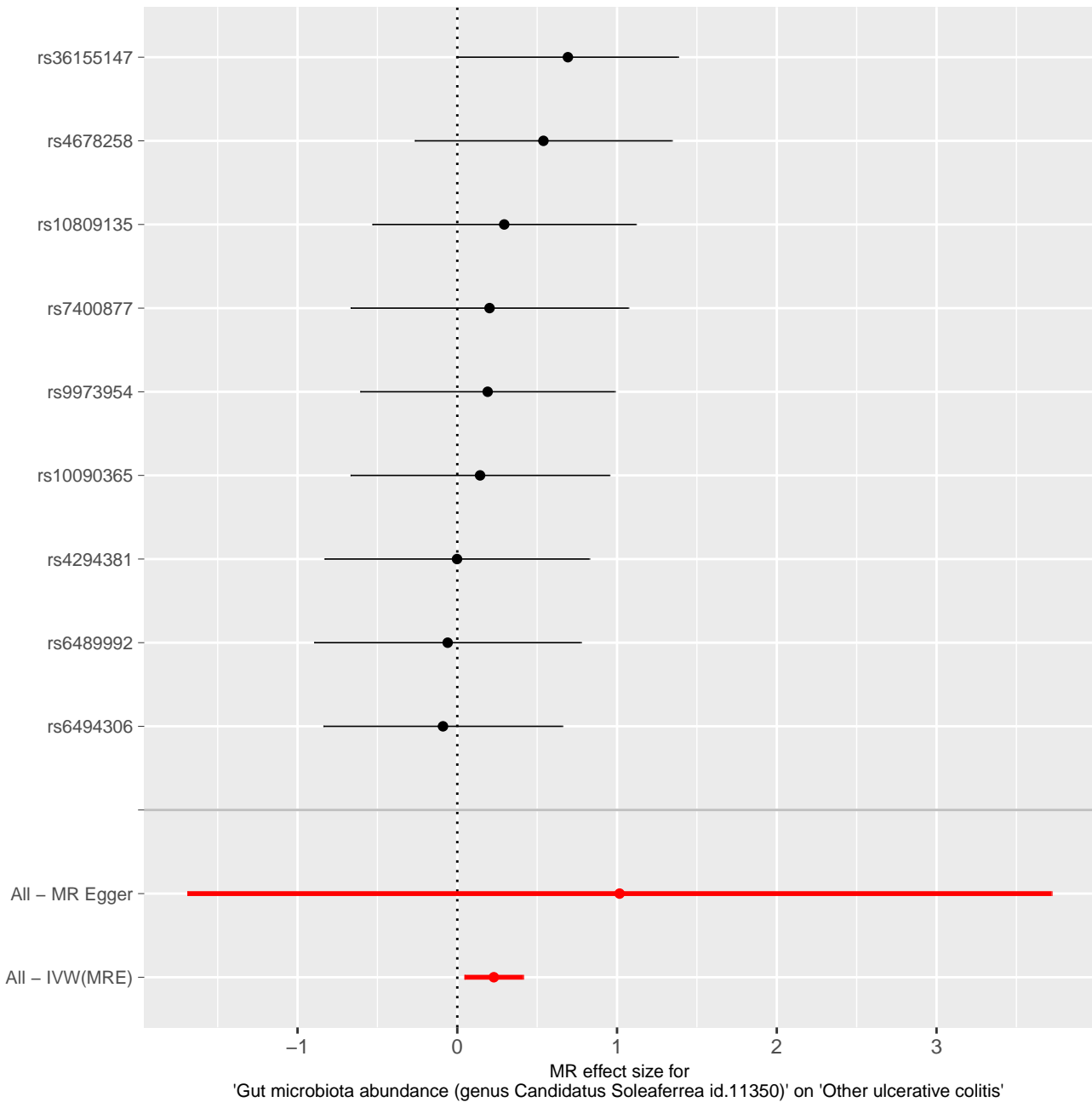
Batch 1245 : Gut microbiota abundance (genus Bilophila id.3170) on Other ulcerative colitis

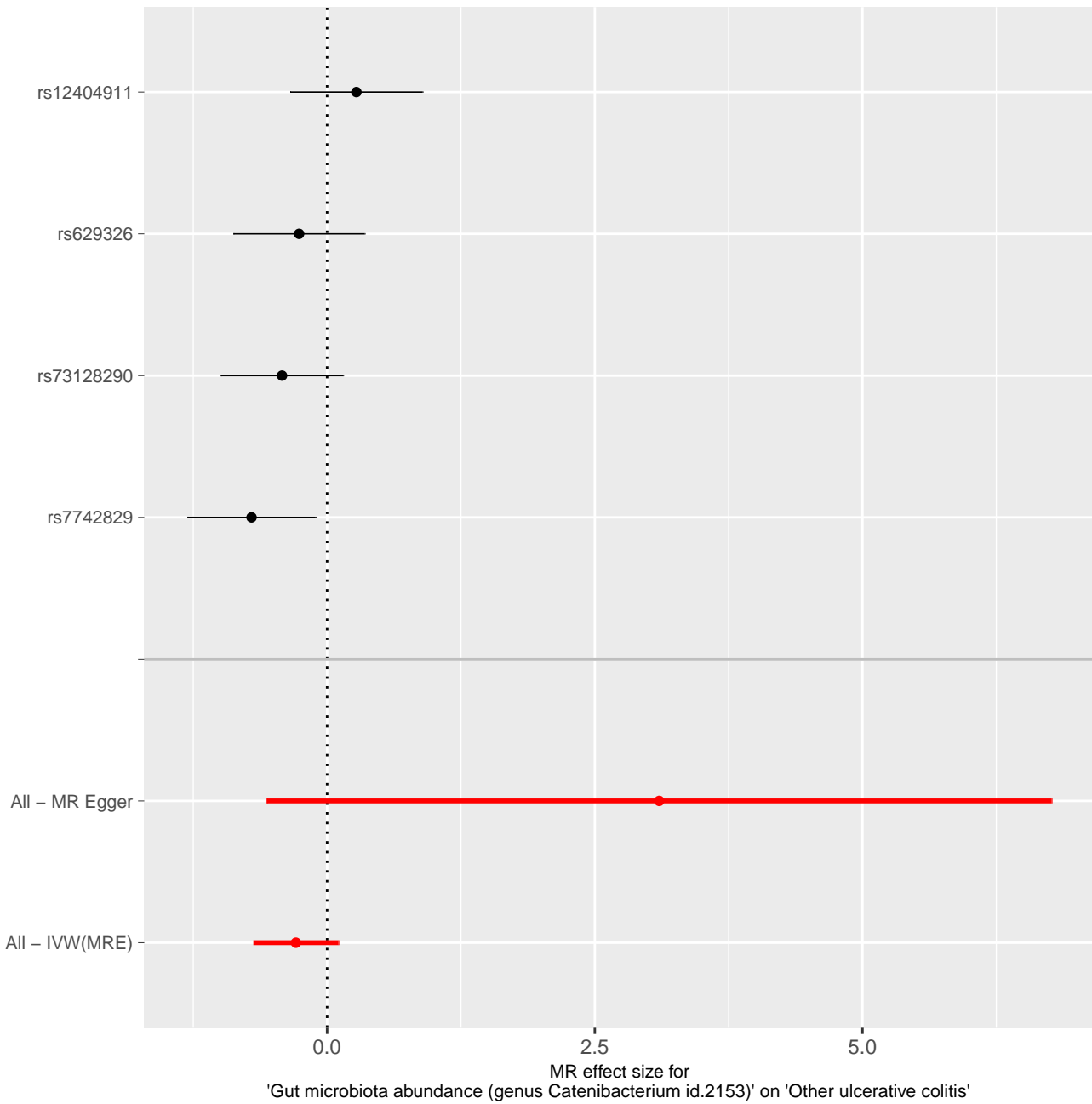


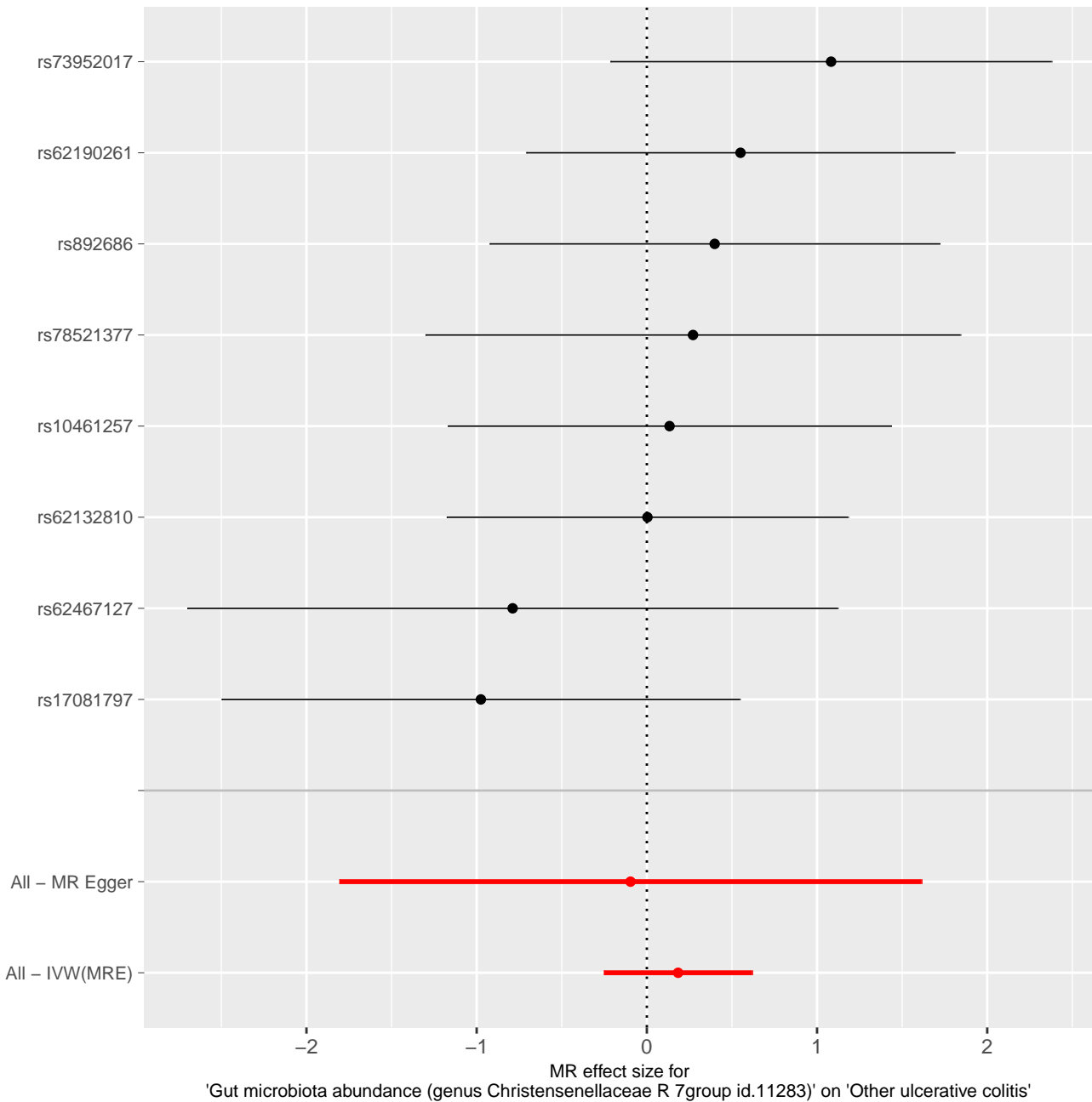


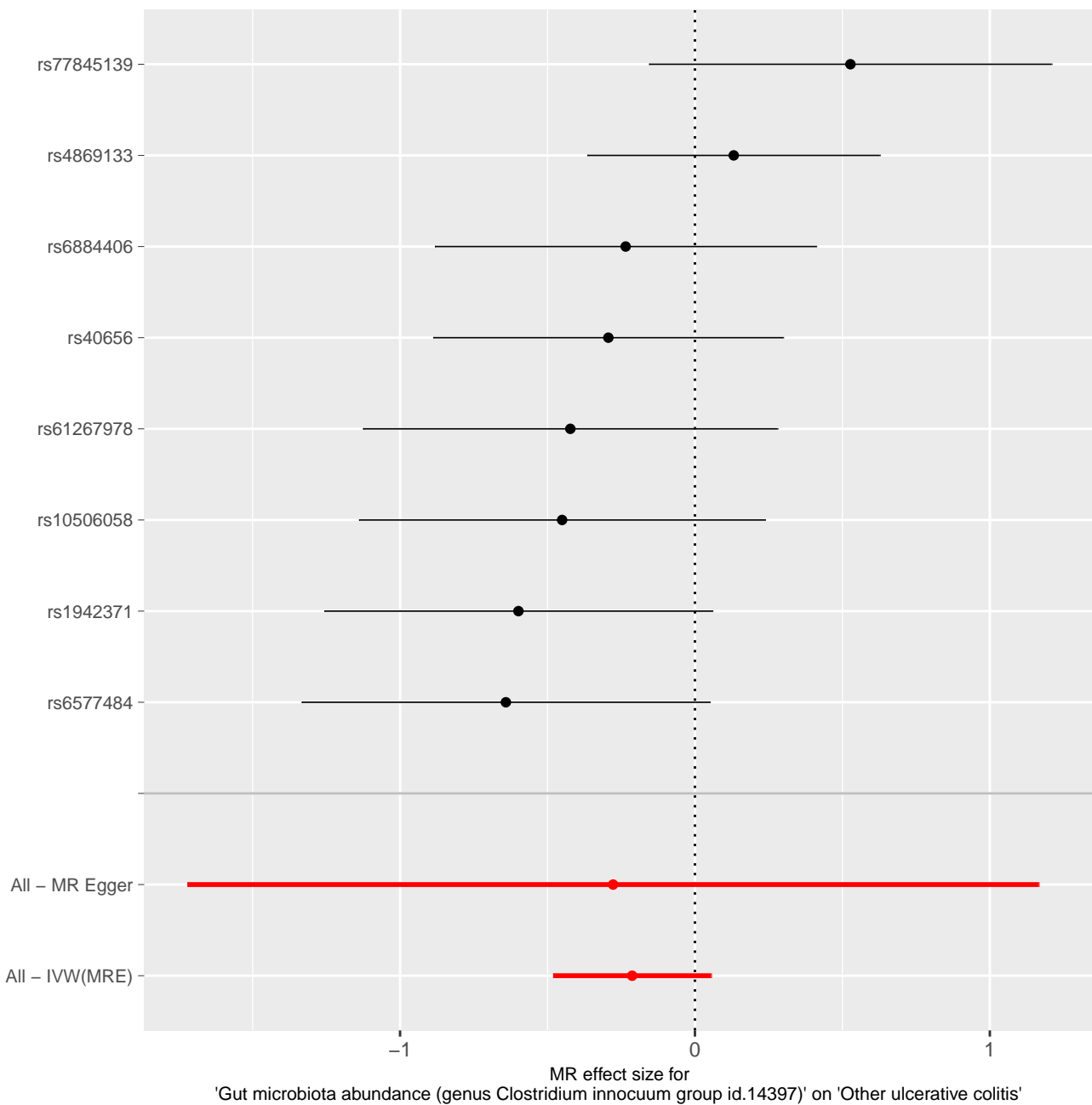


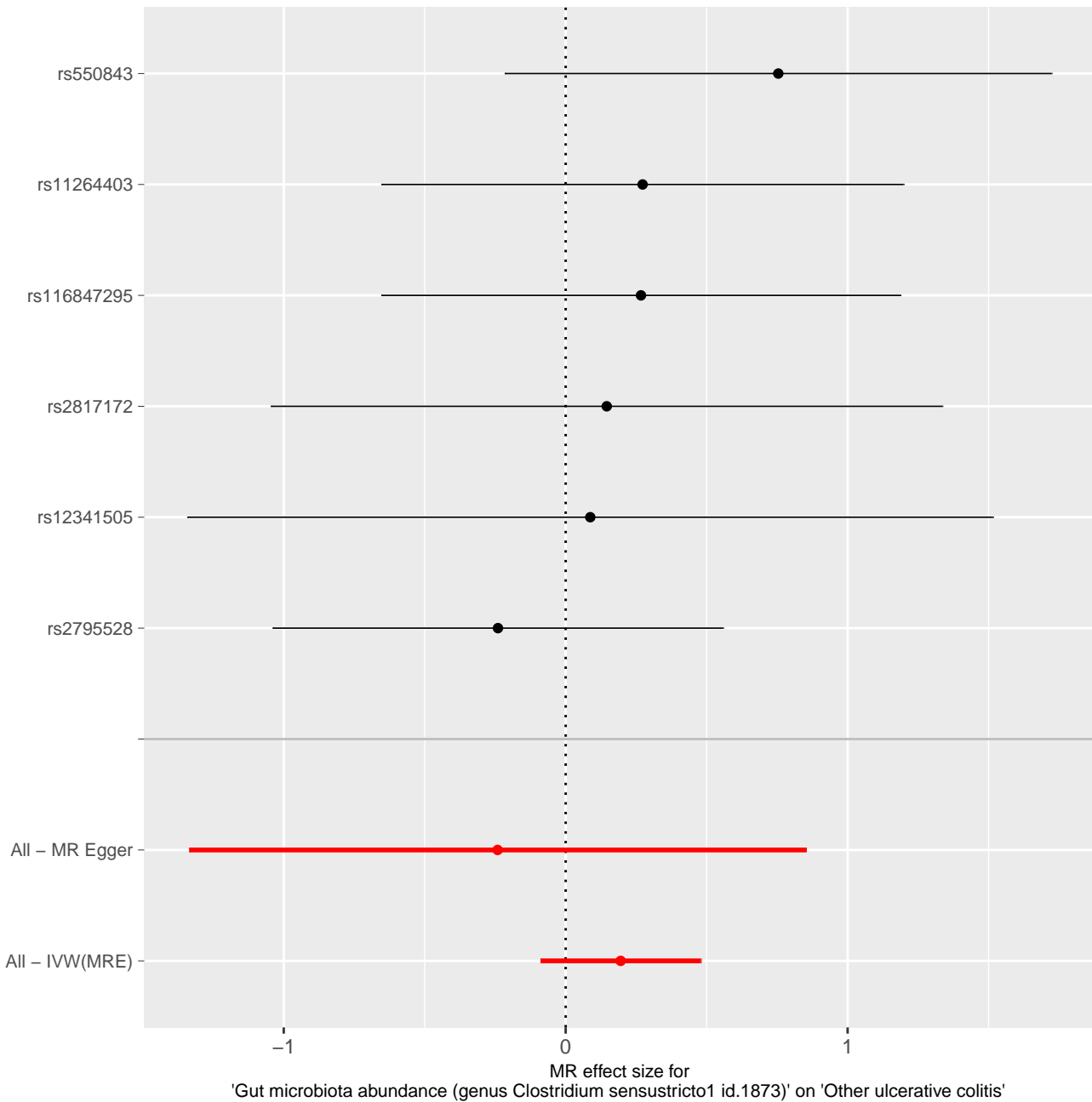


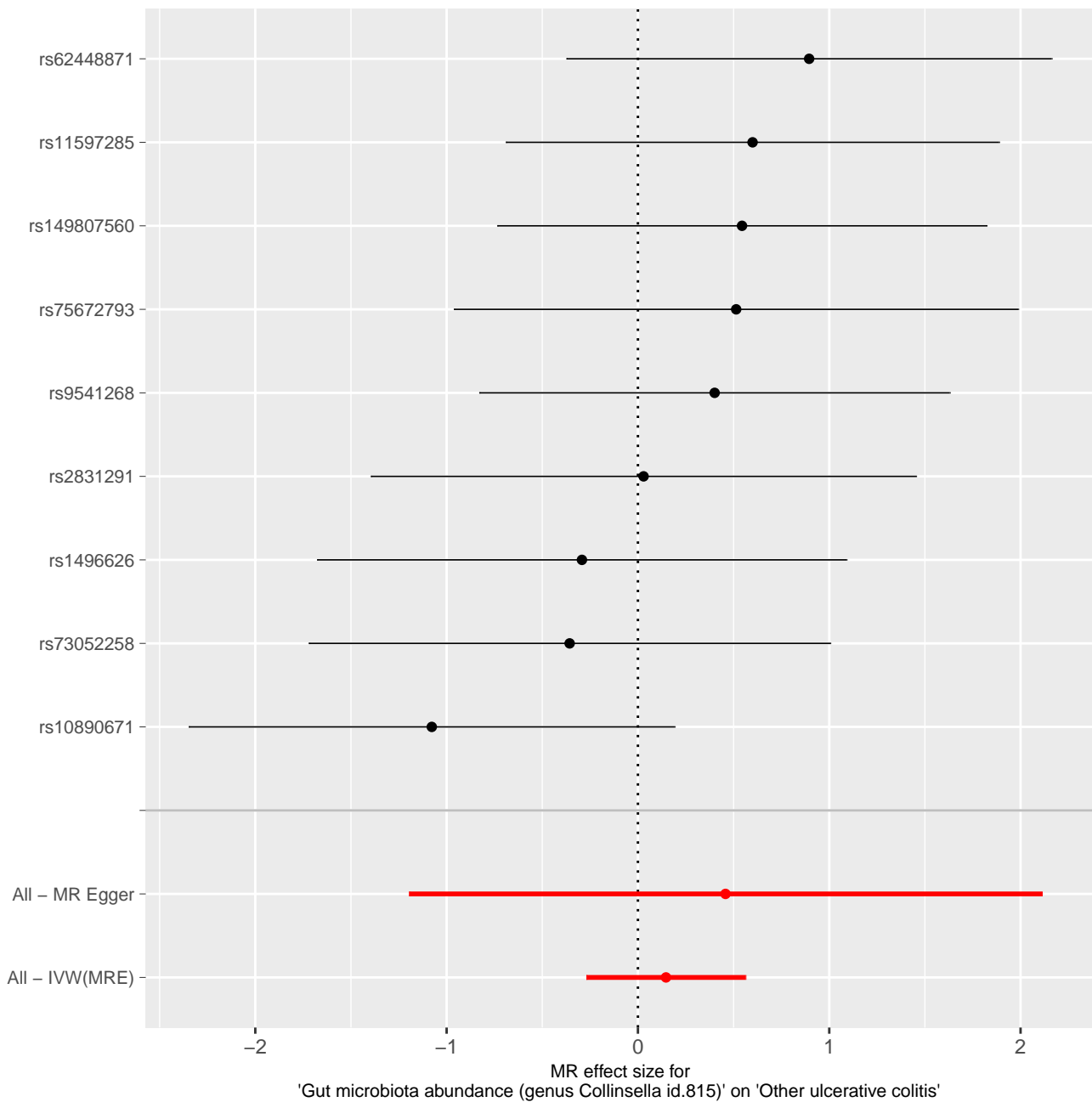


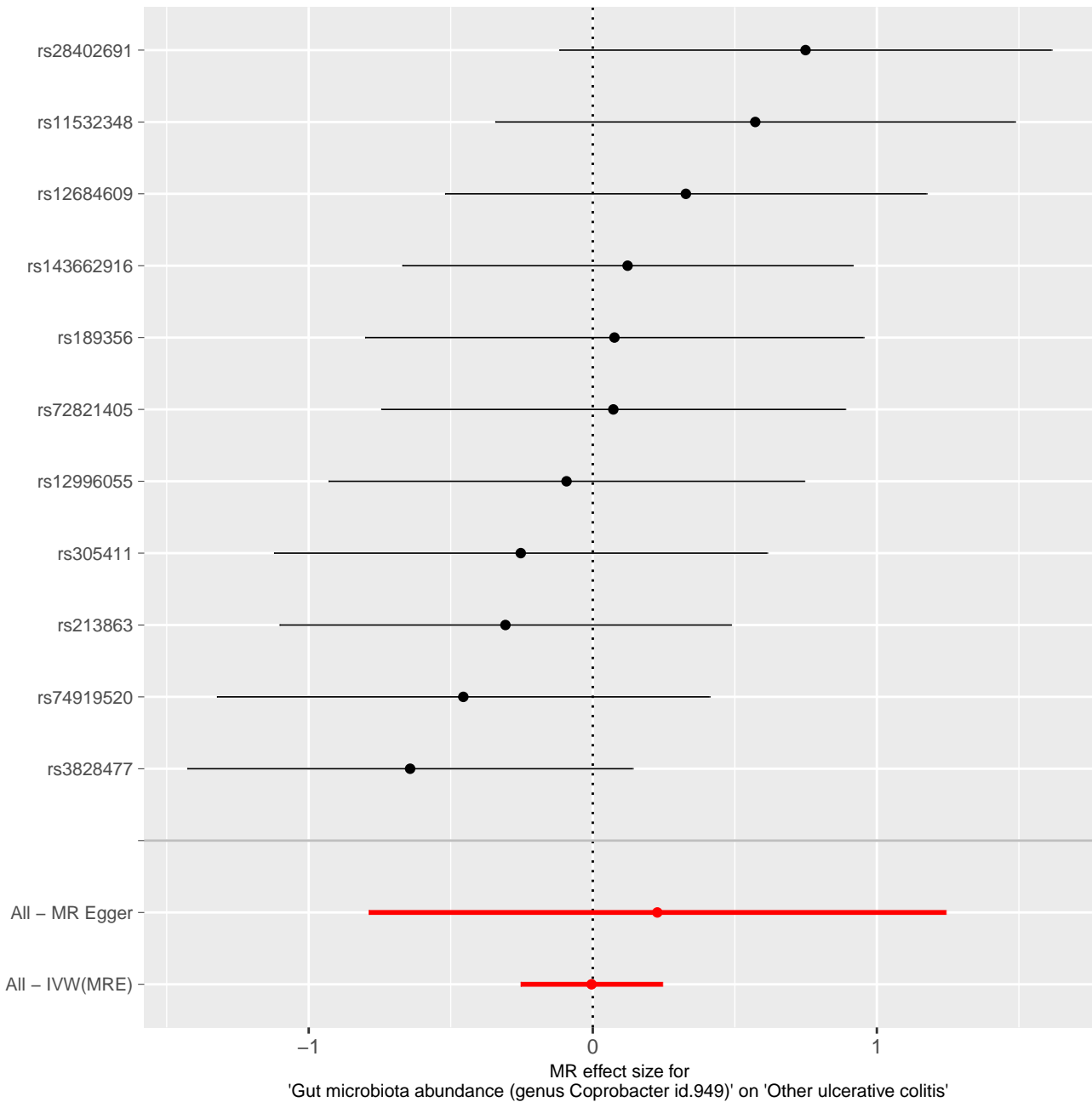


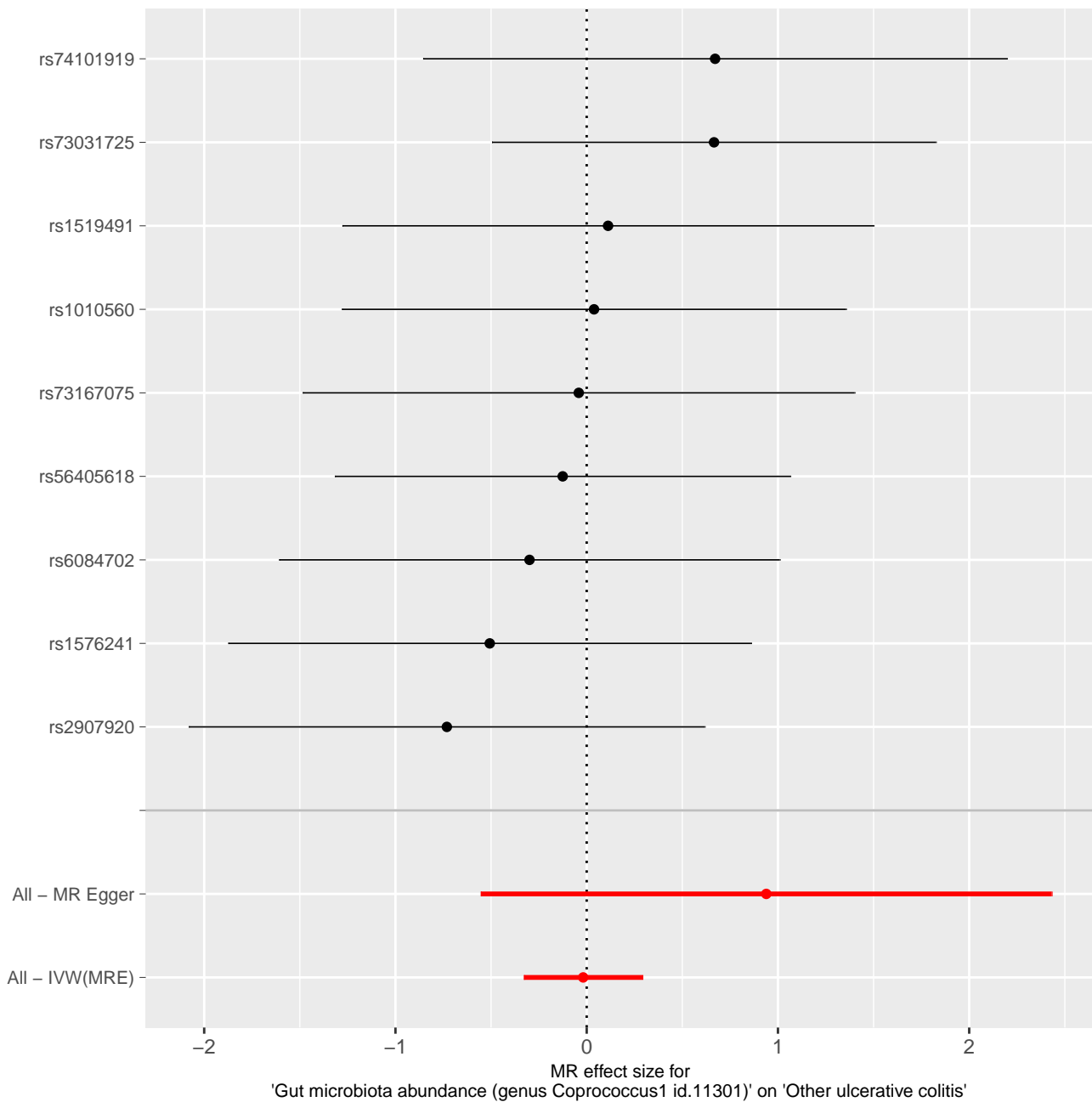


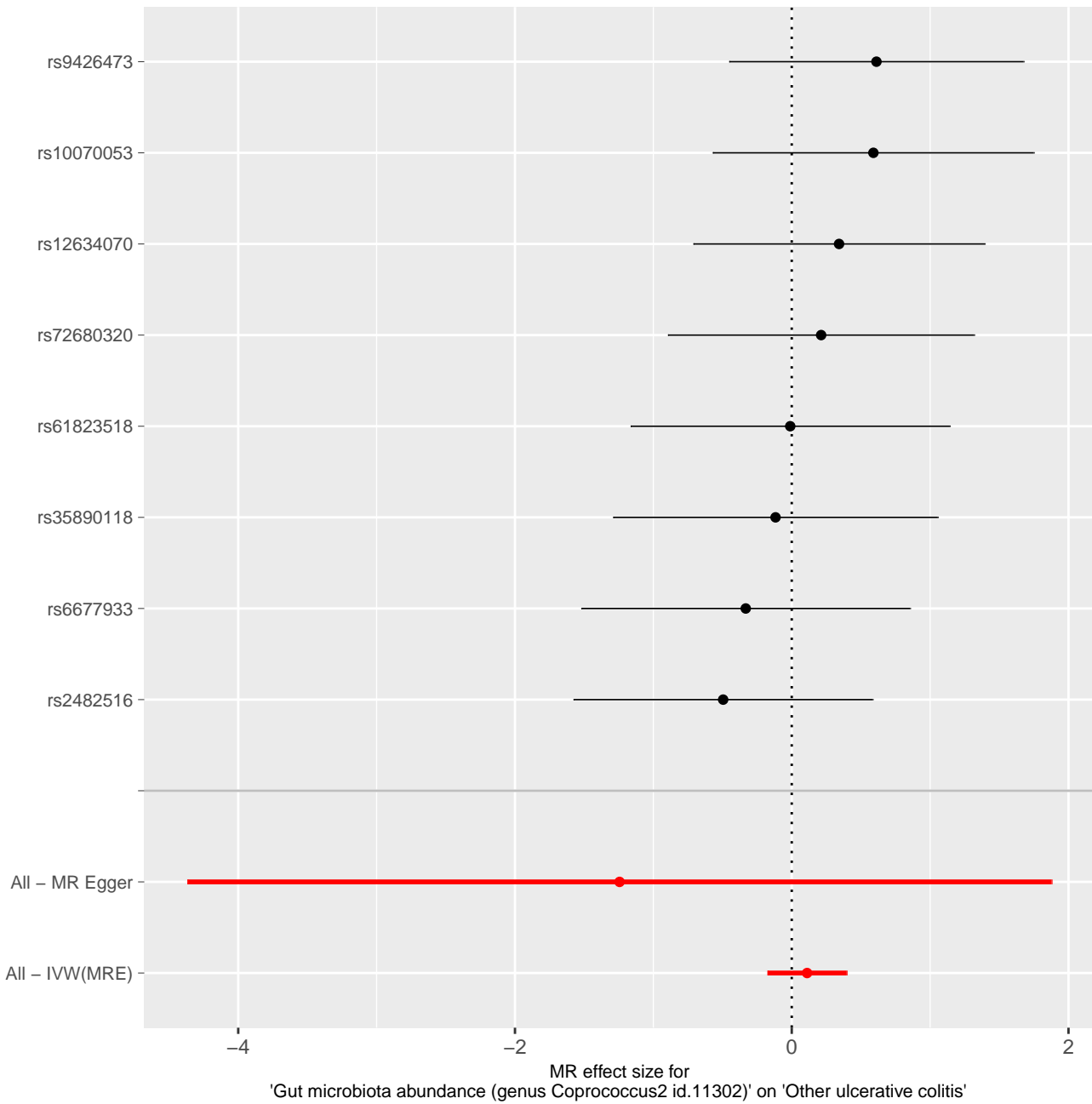


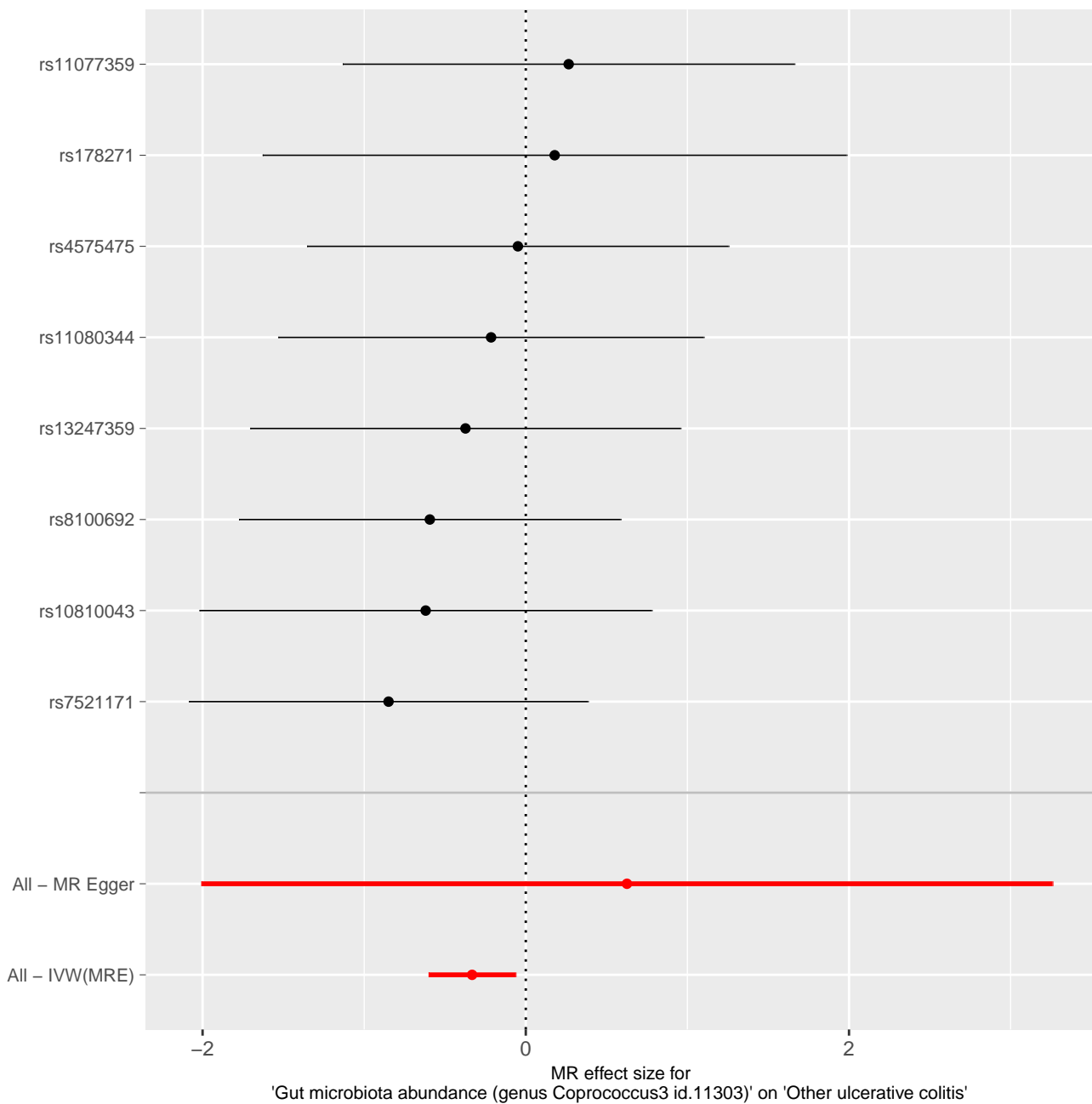


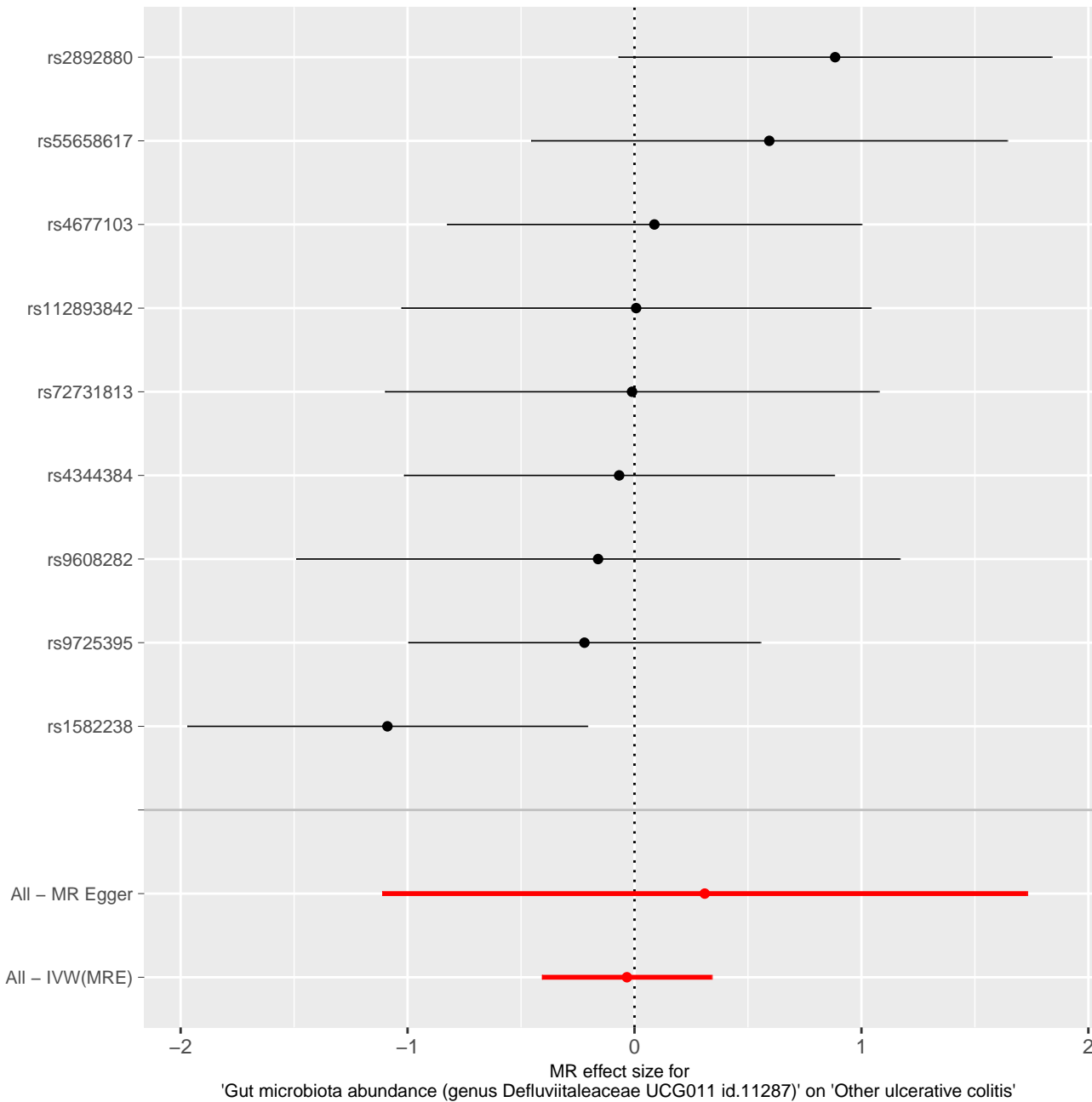


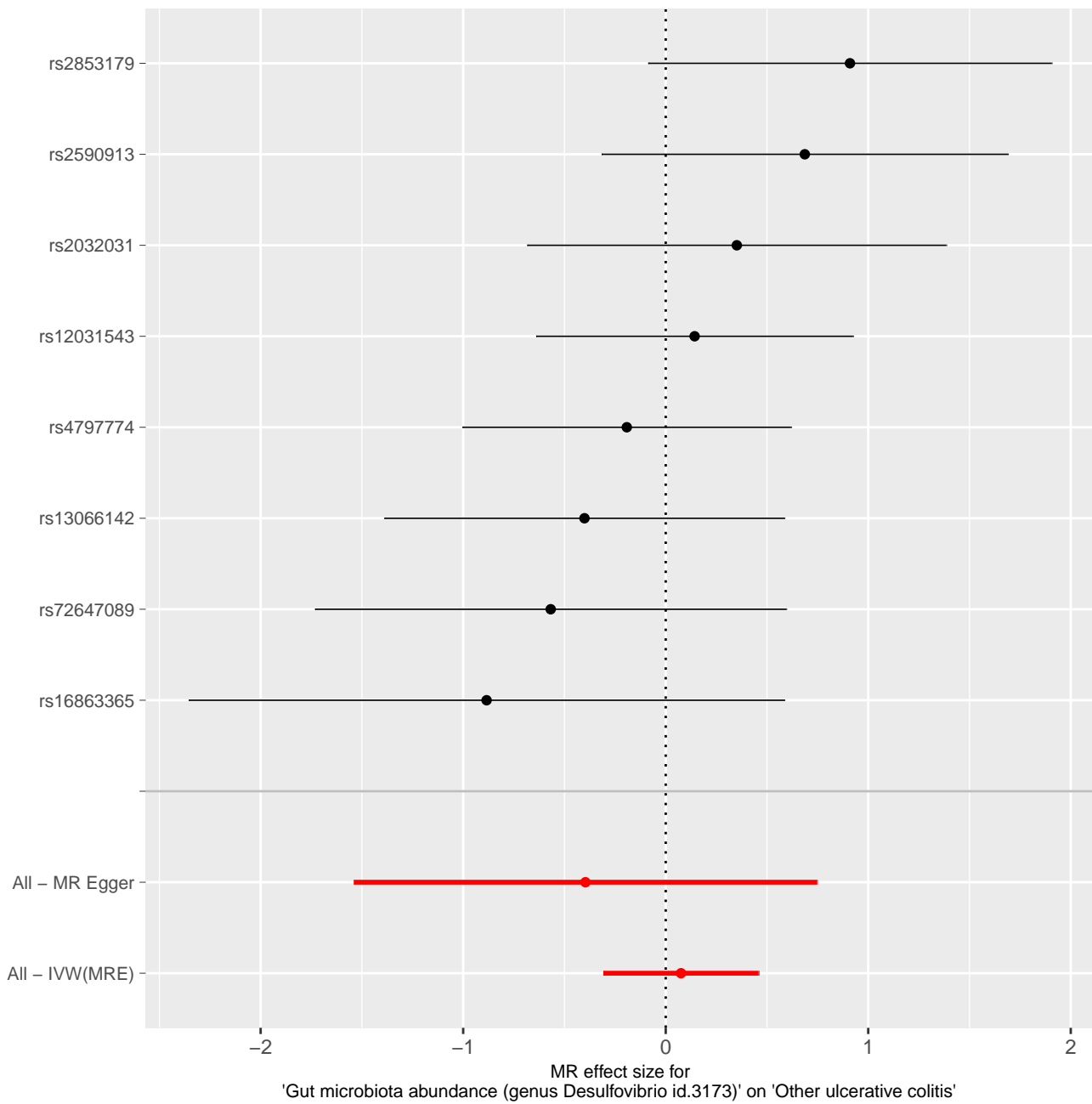


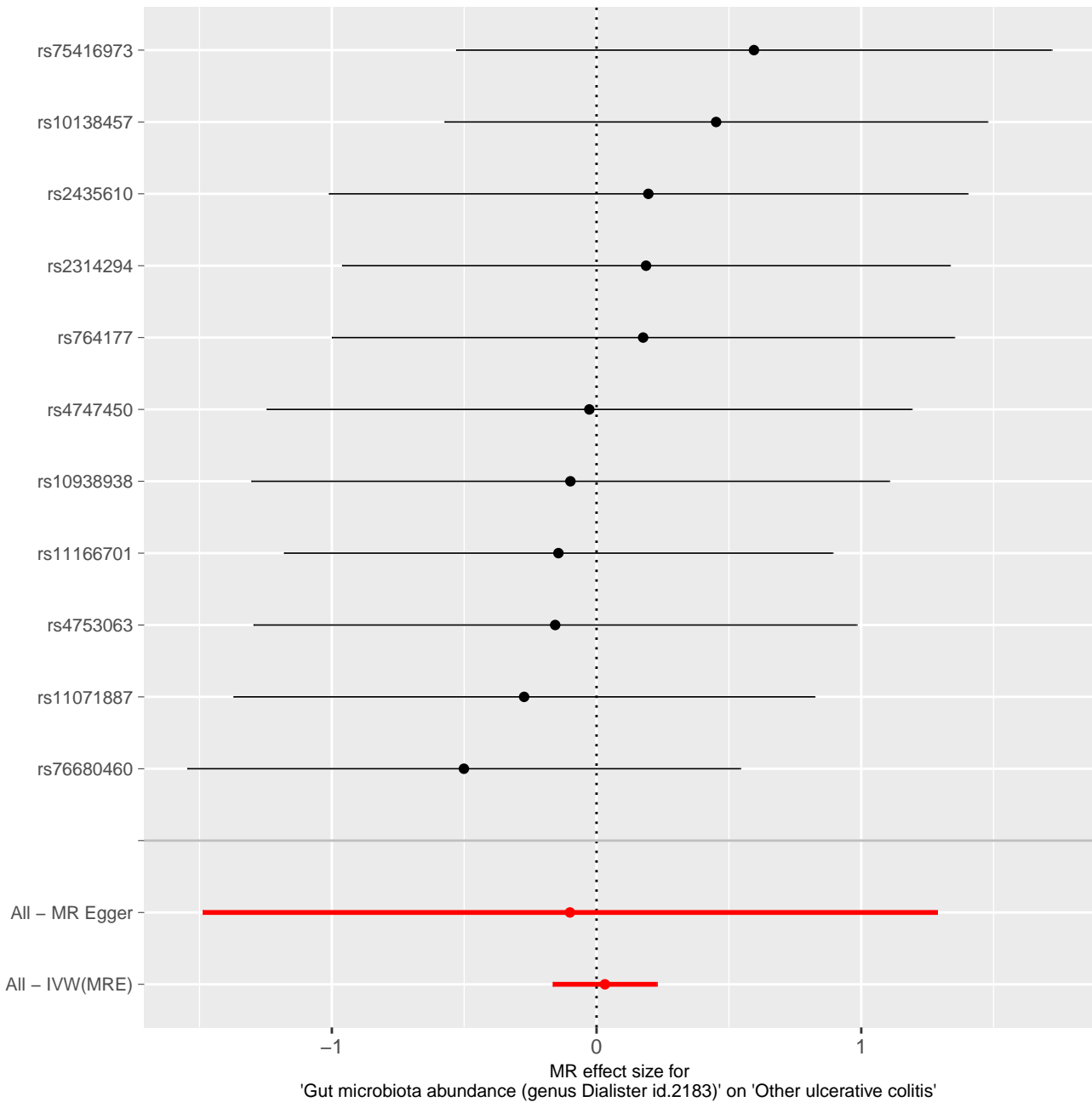


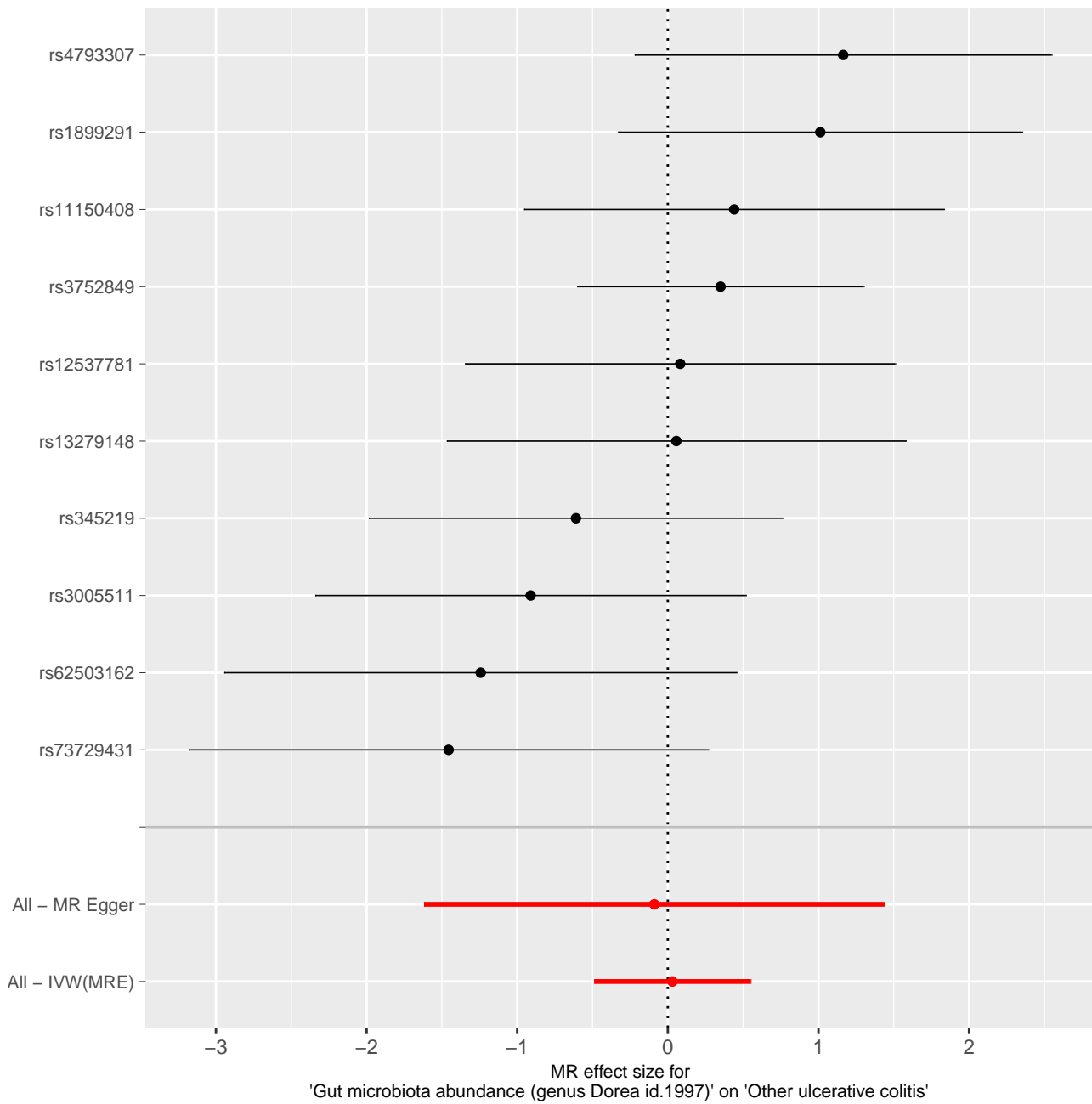


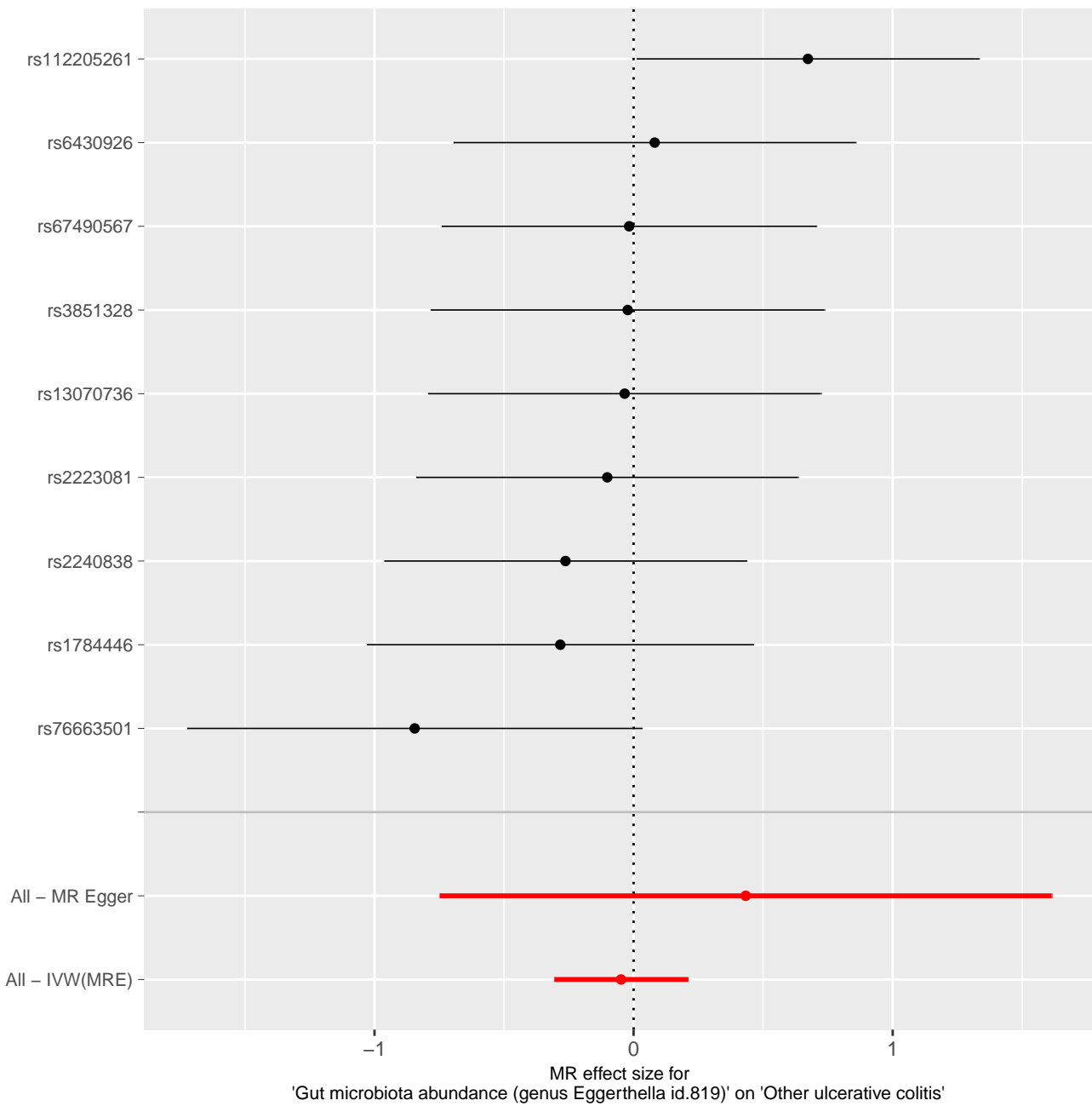


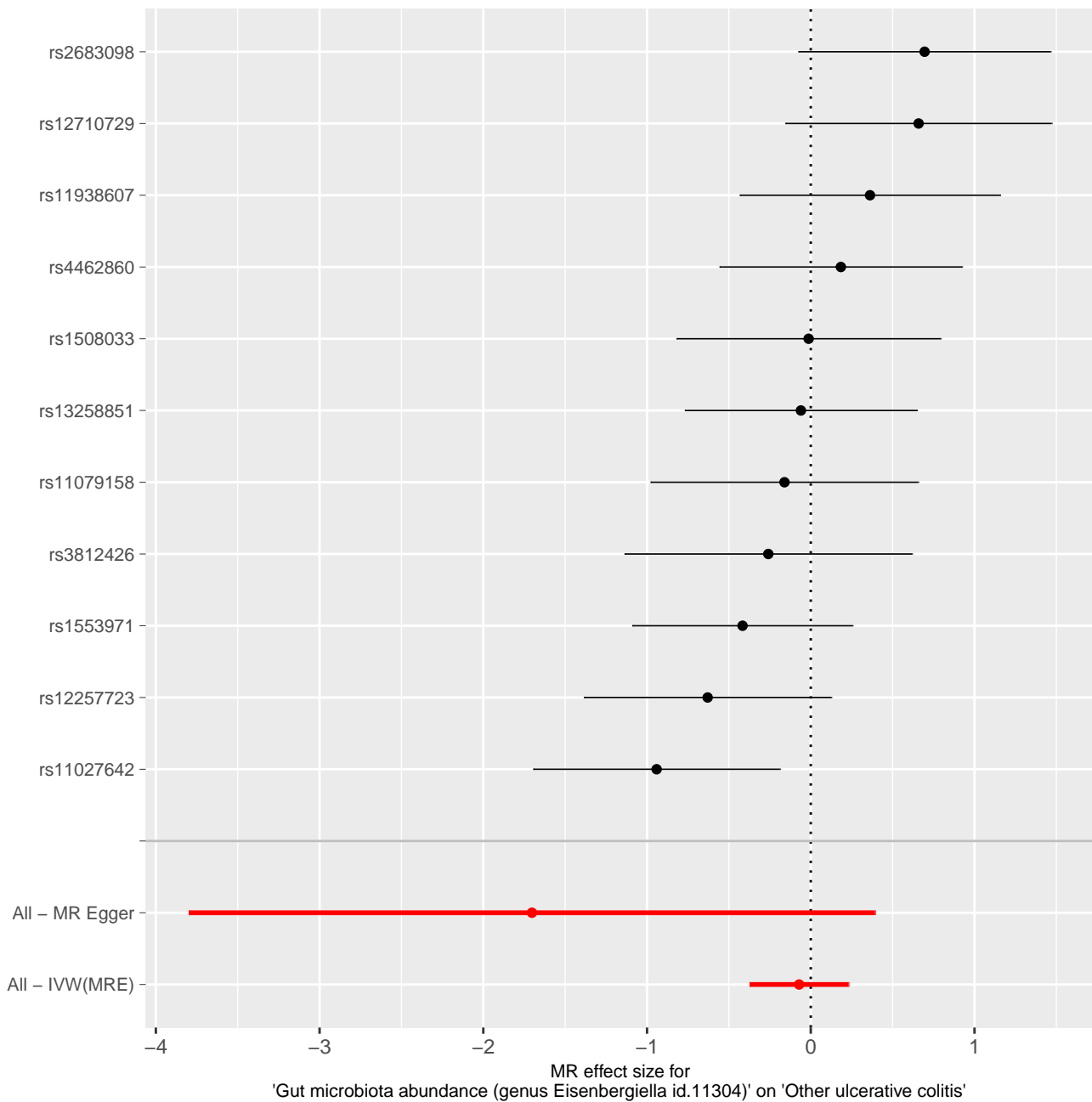


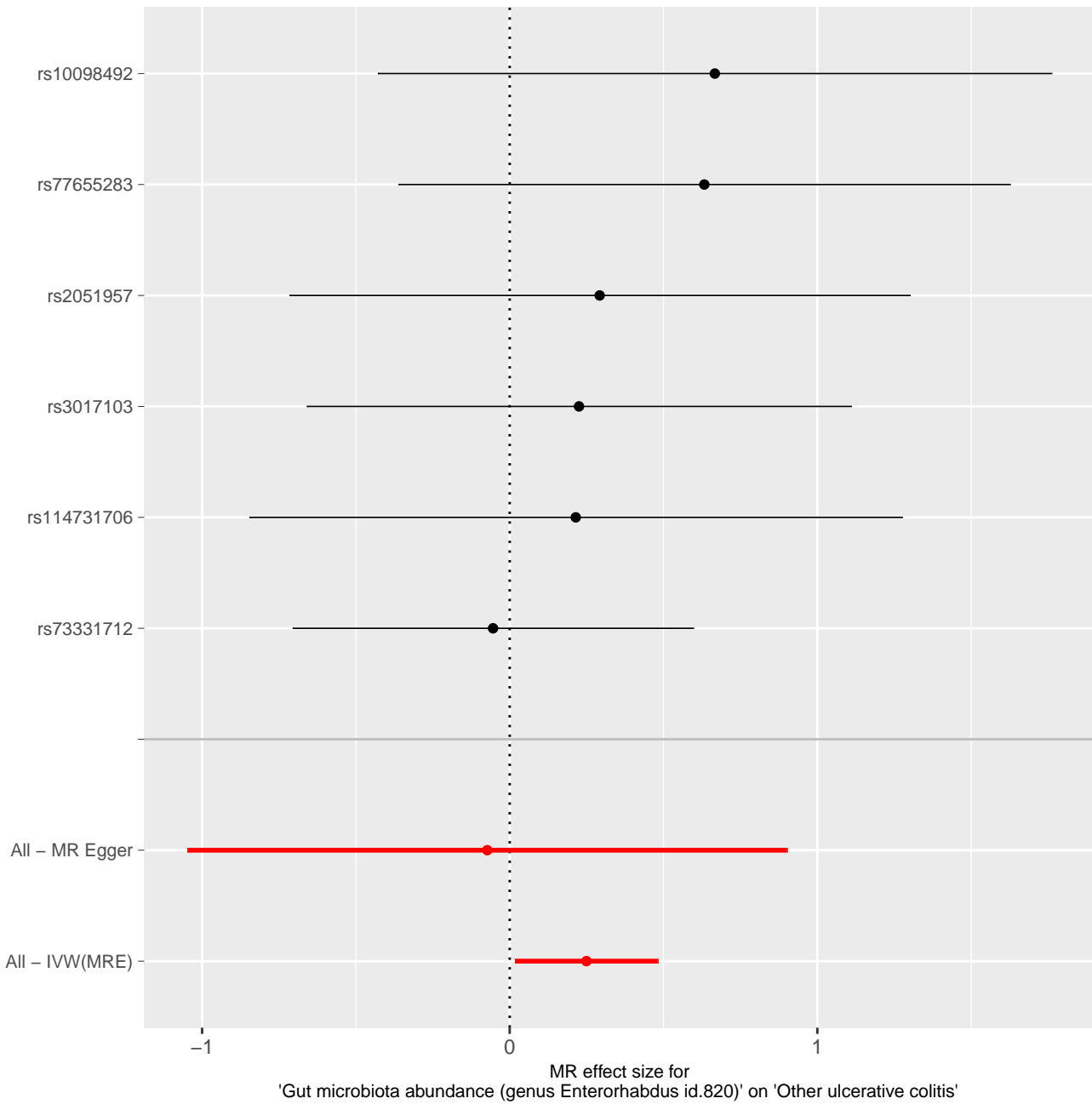


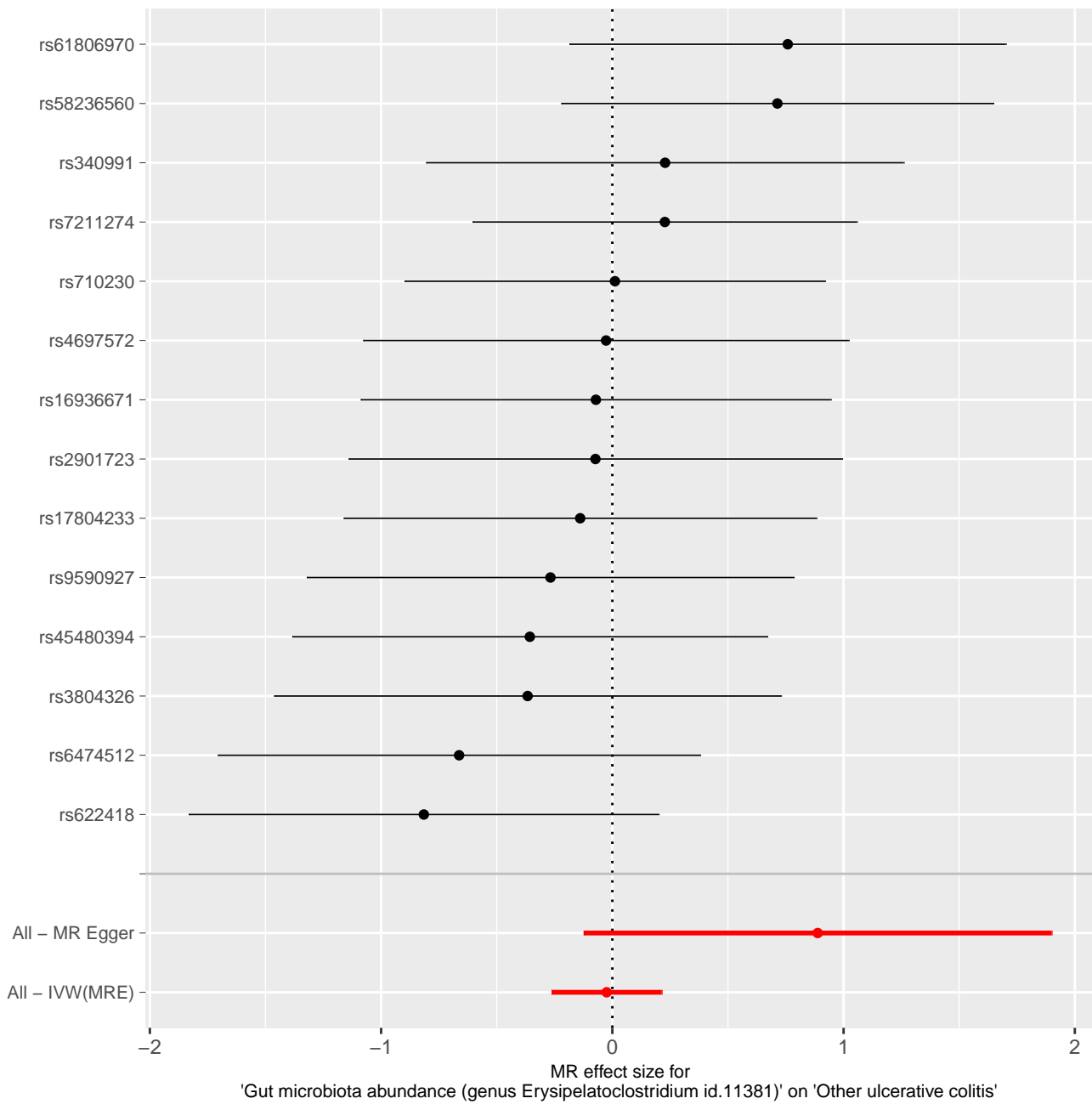


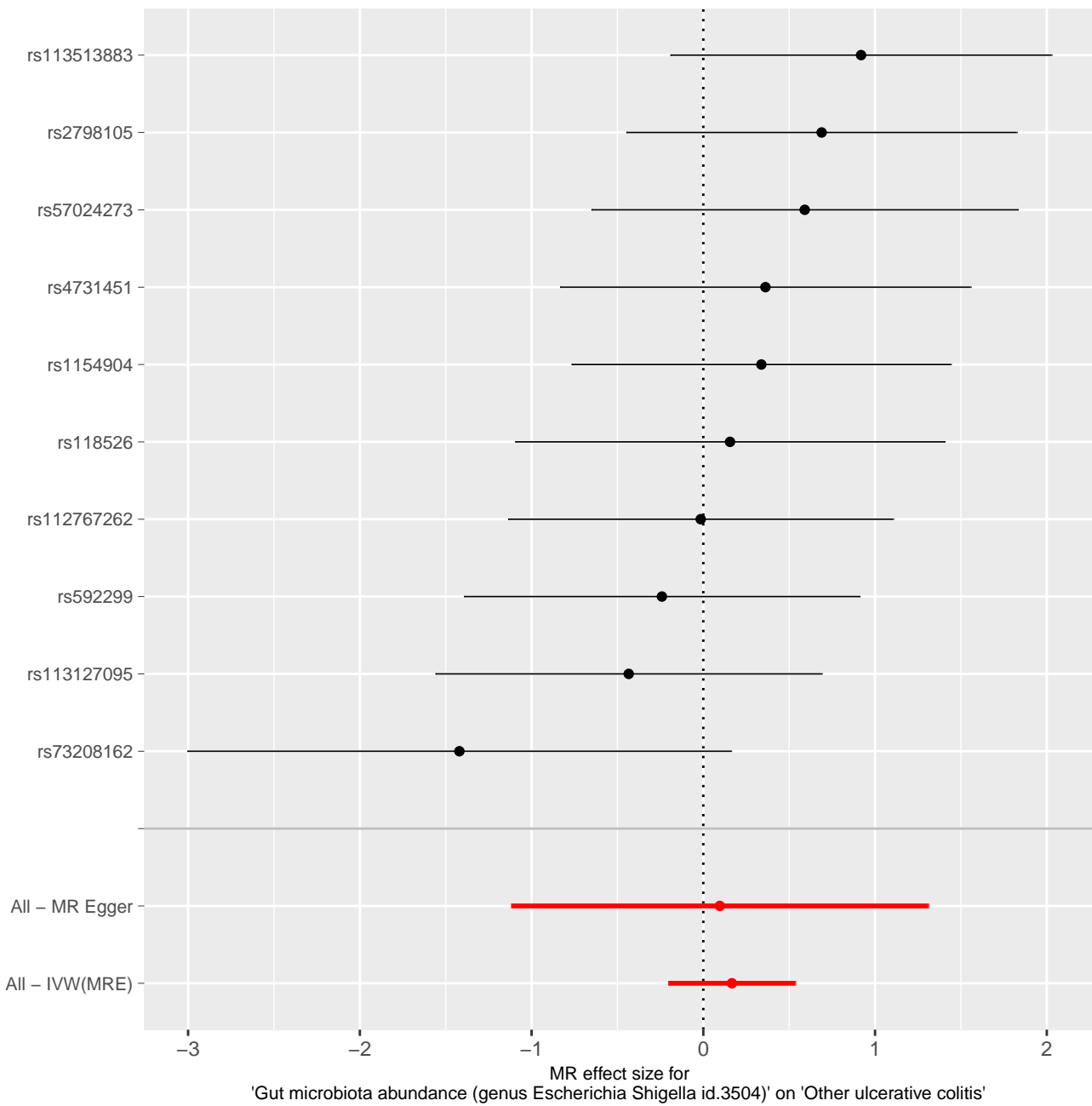


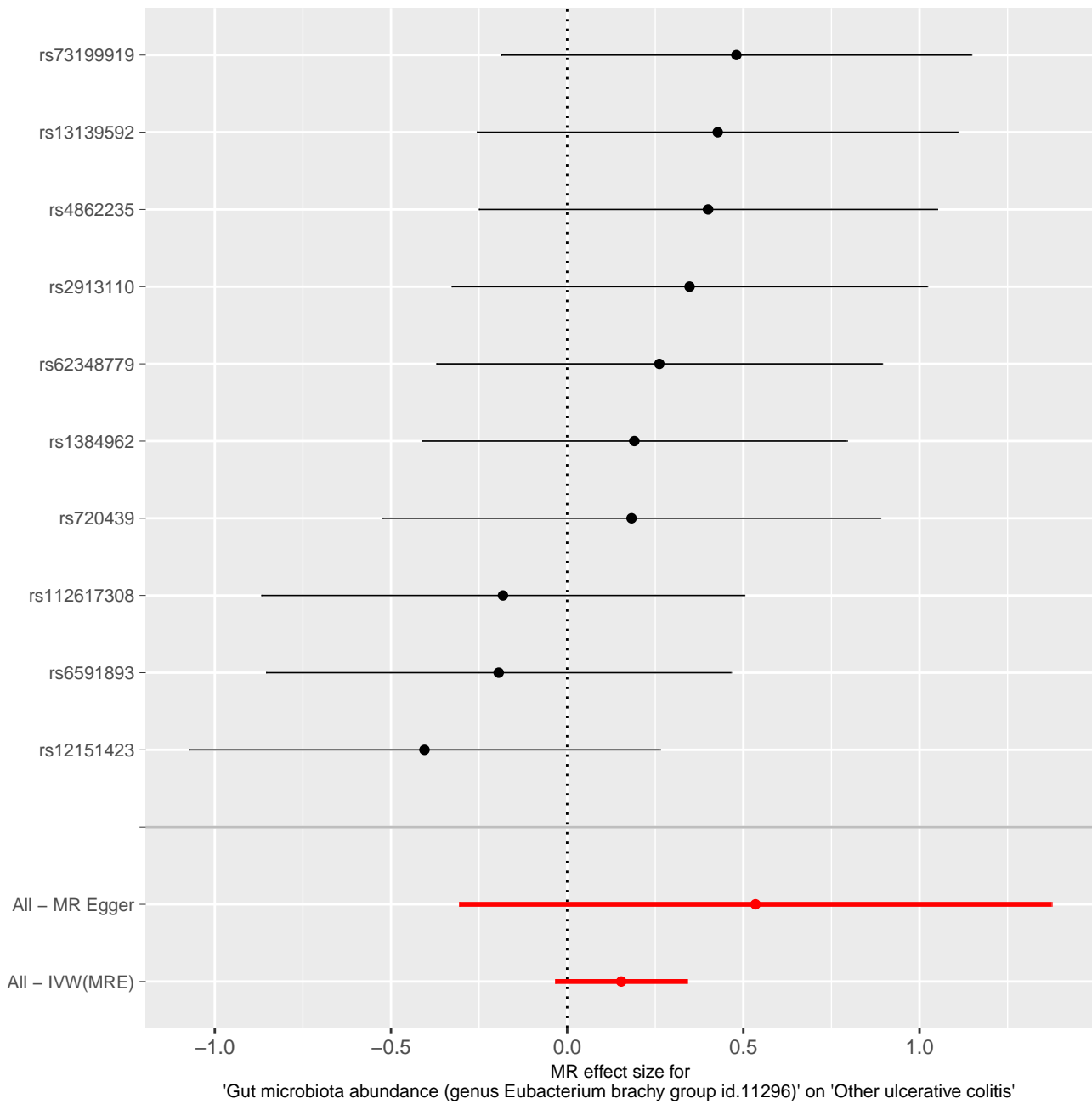




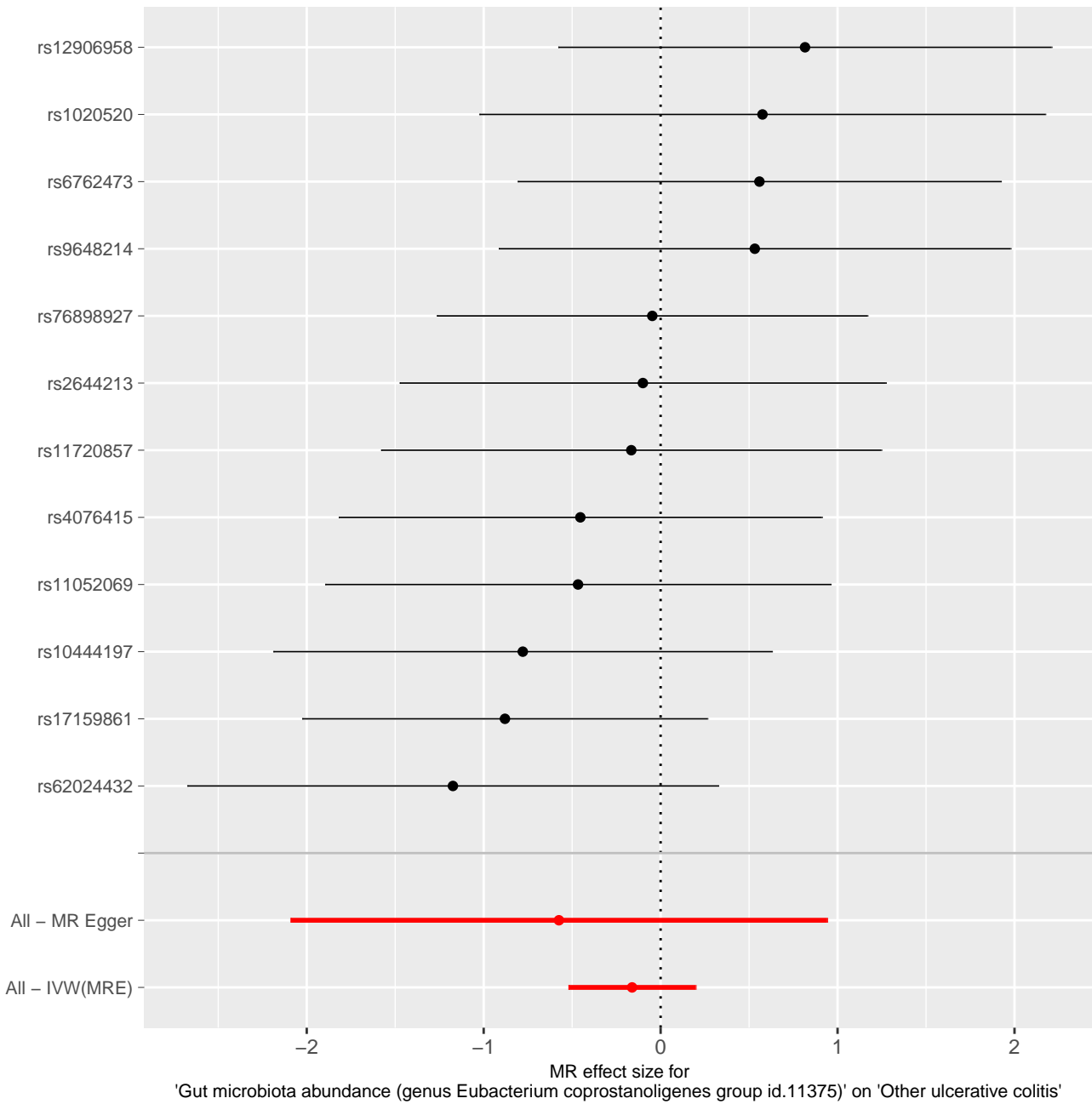


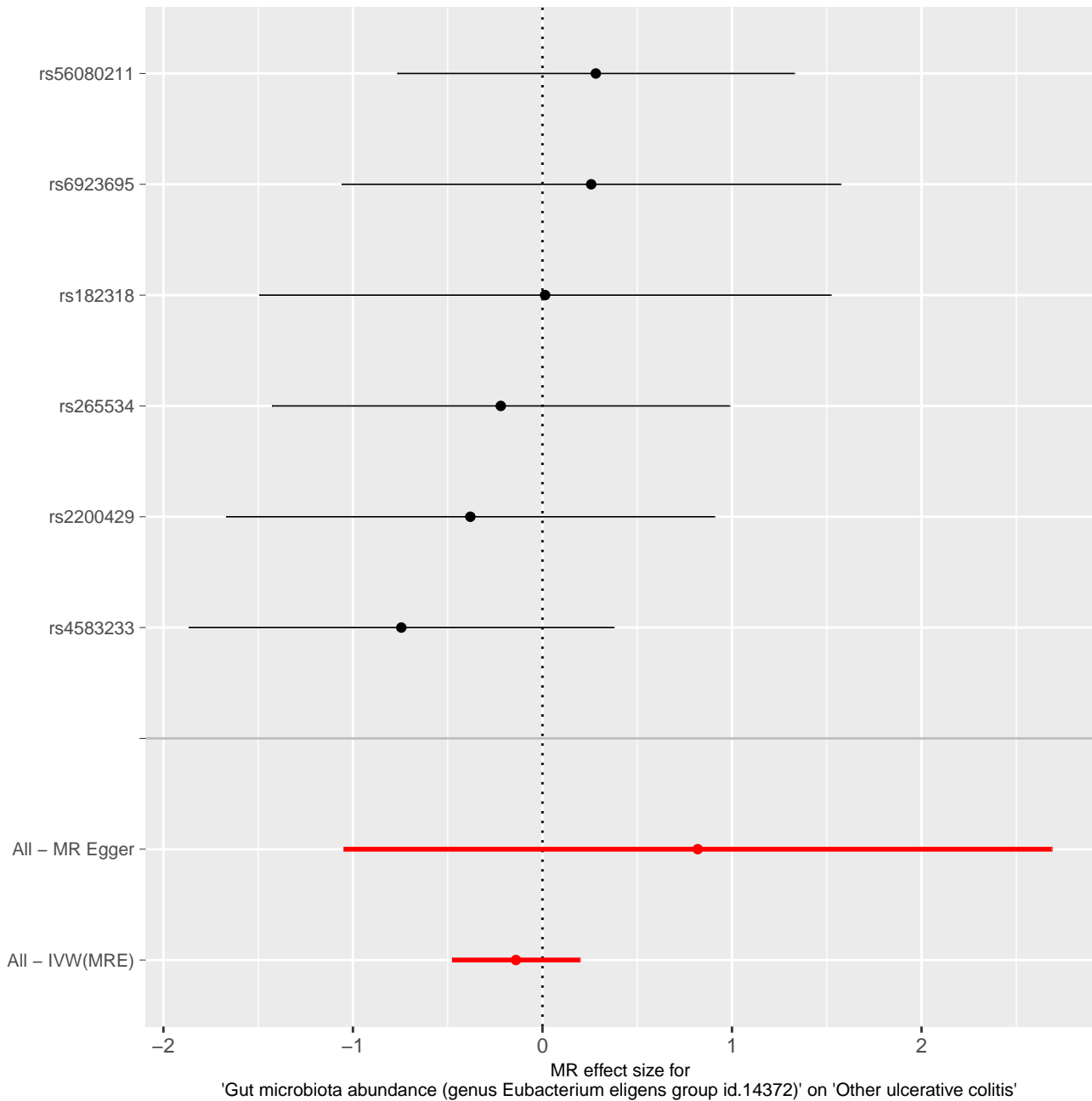


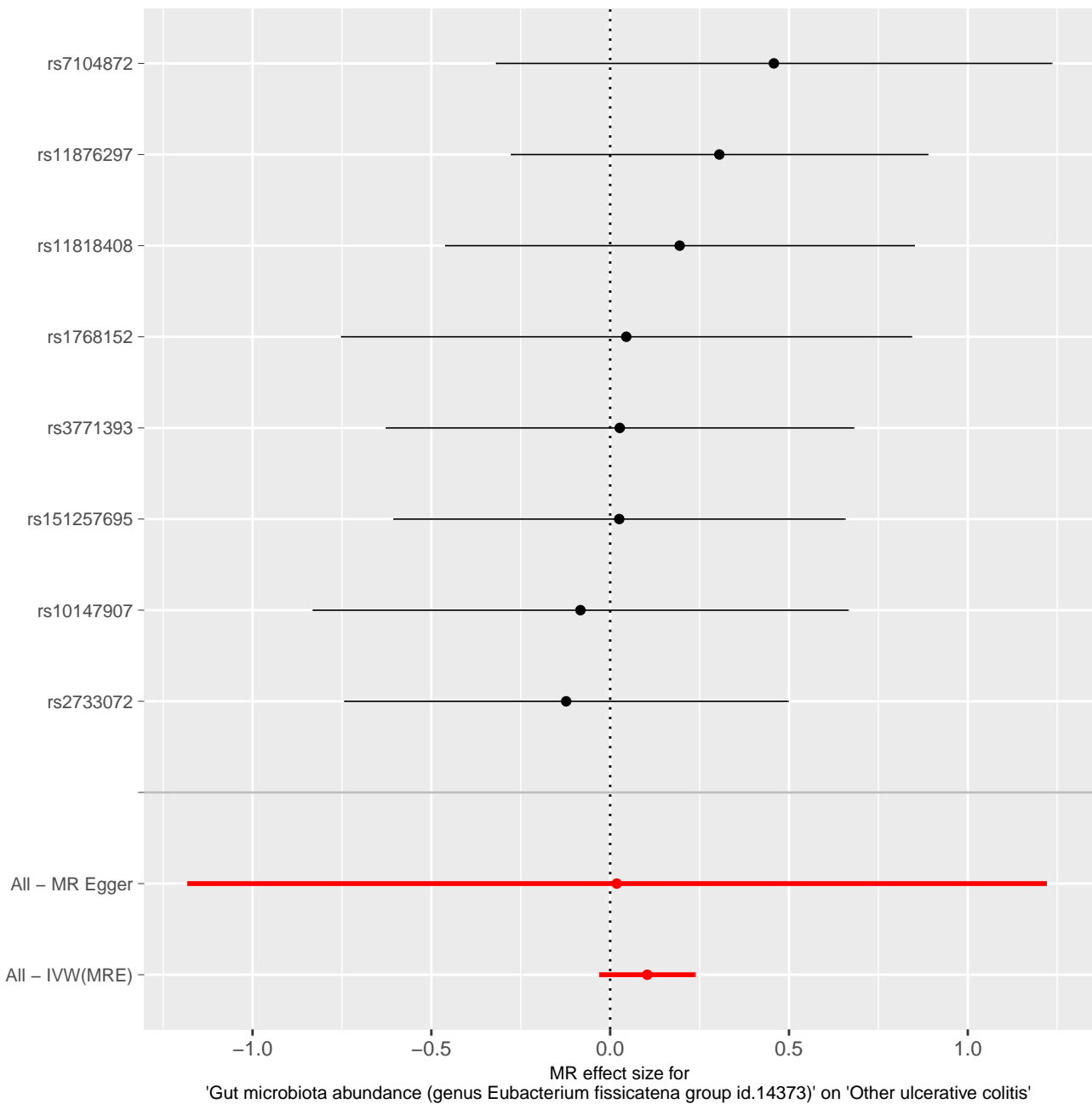


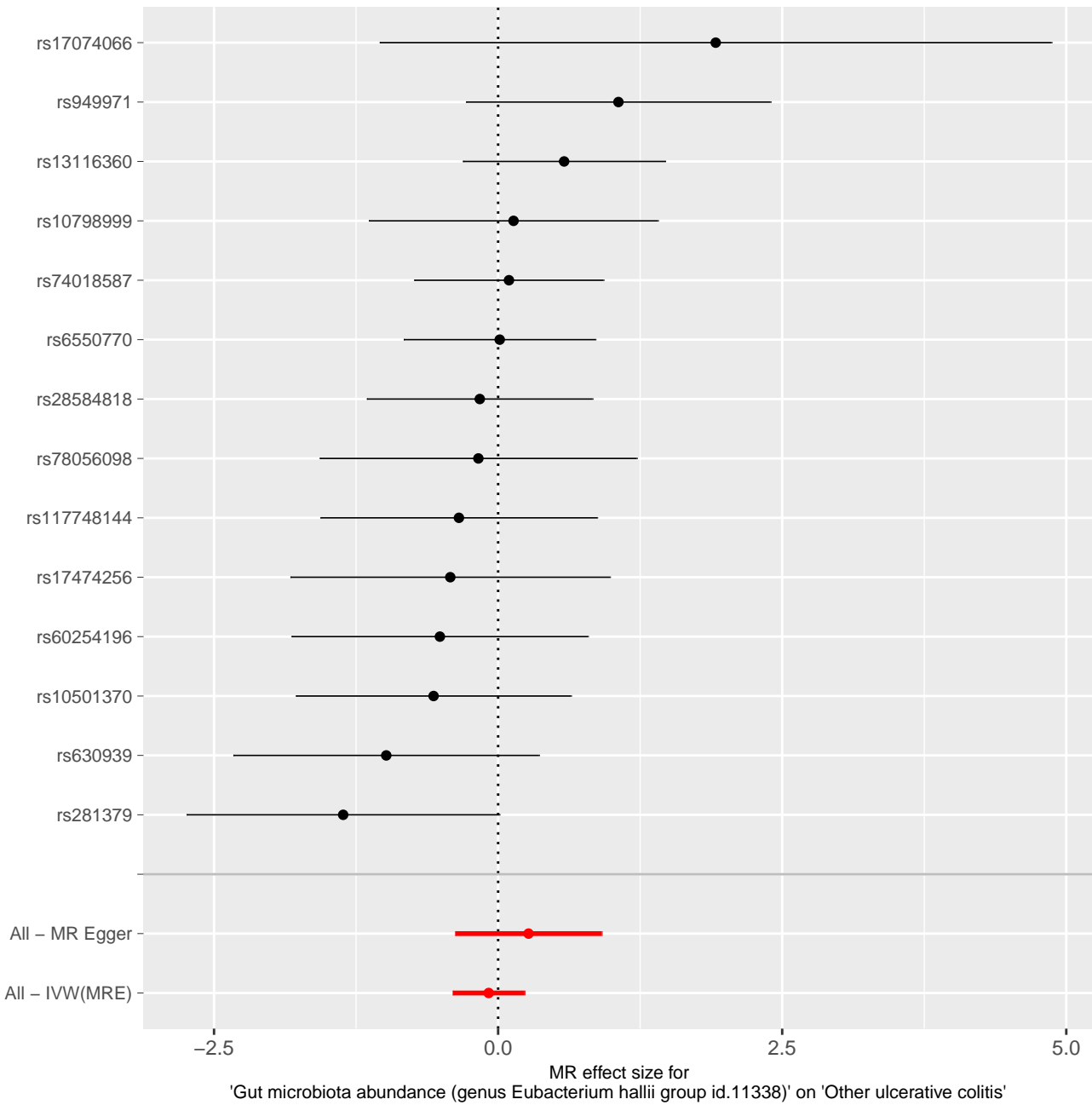


Batch 1271 : Gut microbiota abundance (genus Eubacterium coprostanoligenes group id.11375) on Other ulcerative colitis



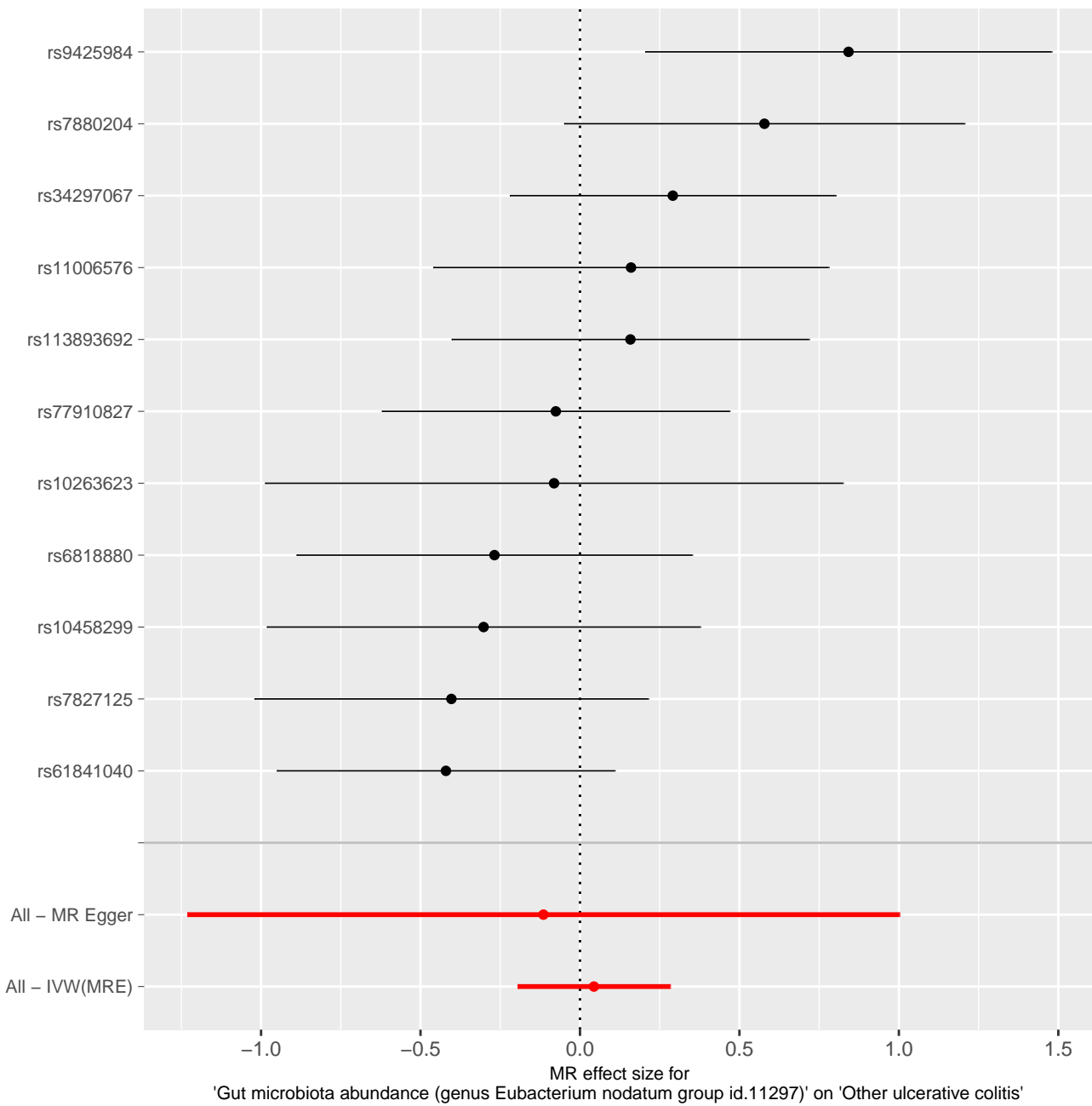


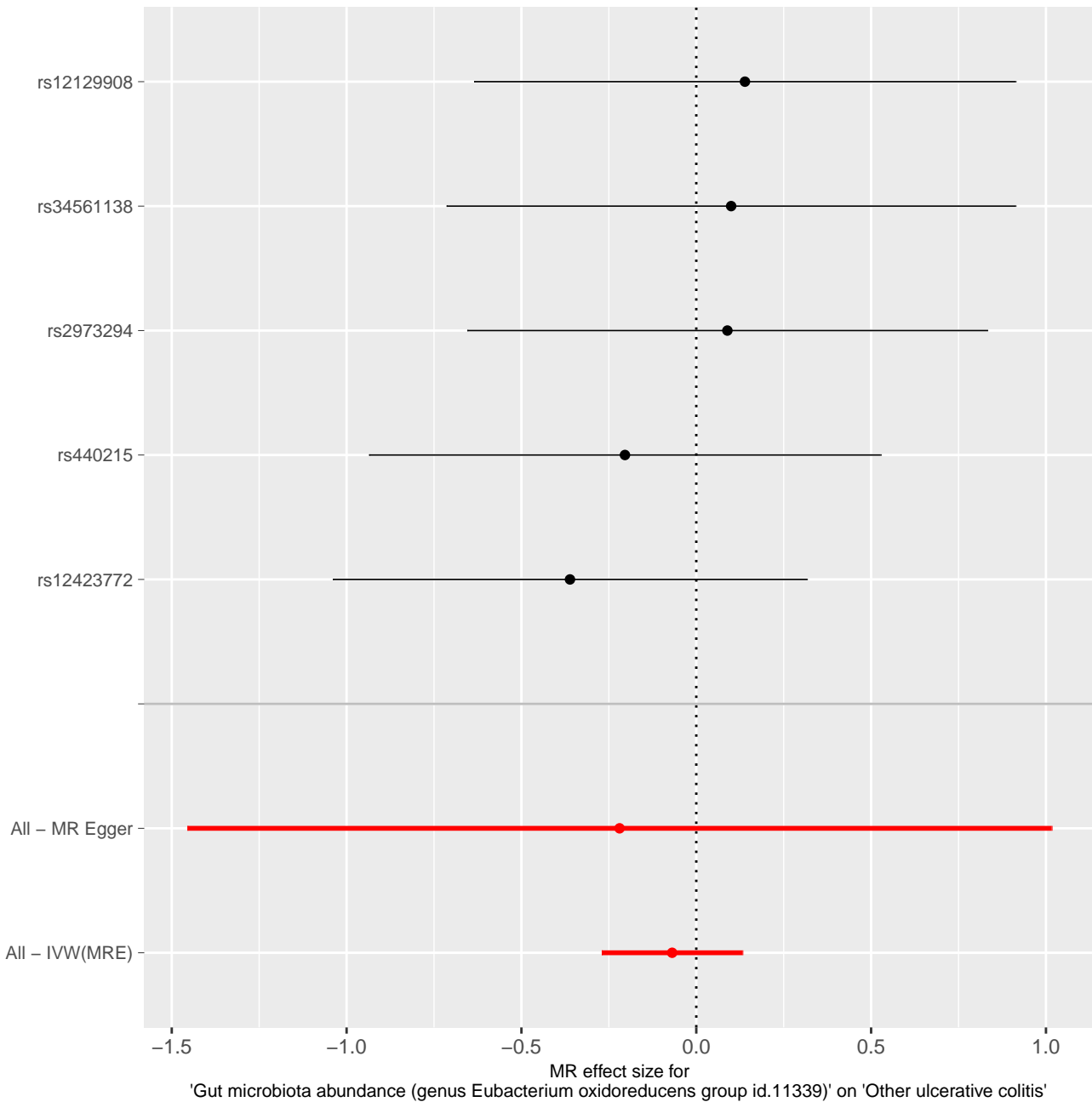


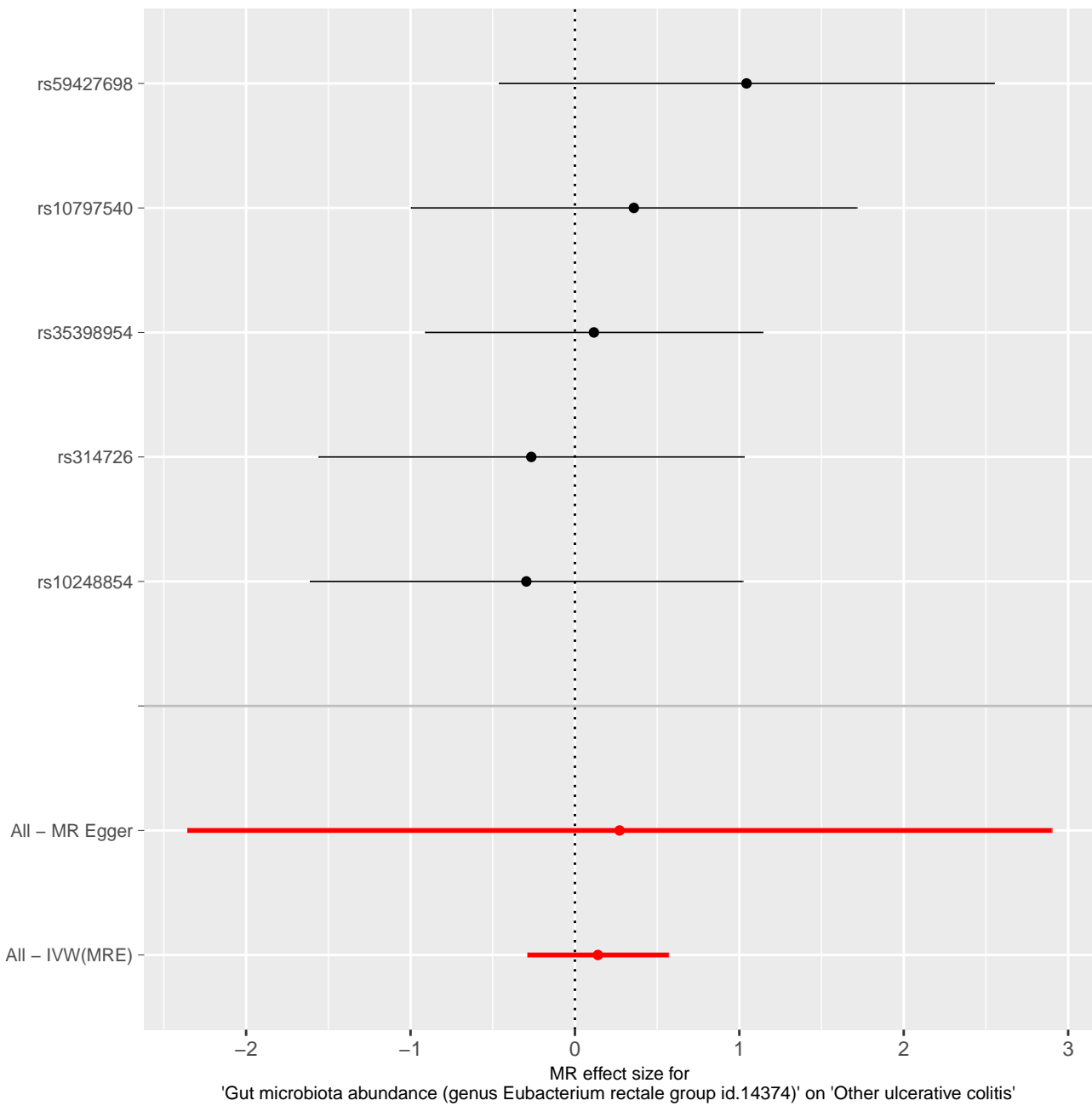


MR effect size for 'Gut microbiota abundance (genus Eubacterium hallii group id.11338) on 'Other ulcerative colitis'

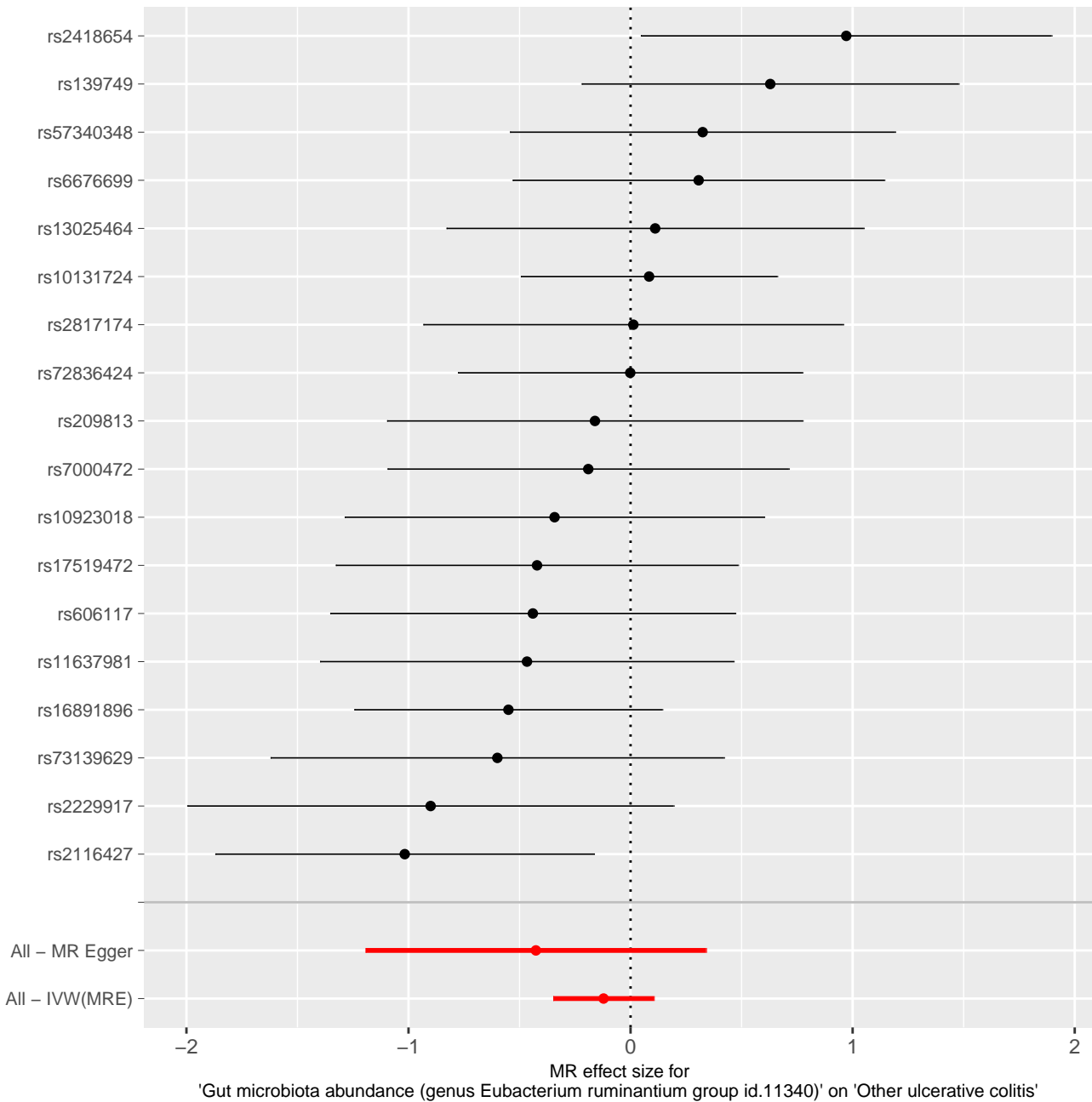
Batch 1275 : Gut microbiota abundance (genus Eubacterium nodatum group id.11297) on Other ulcerative colitis

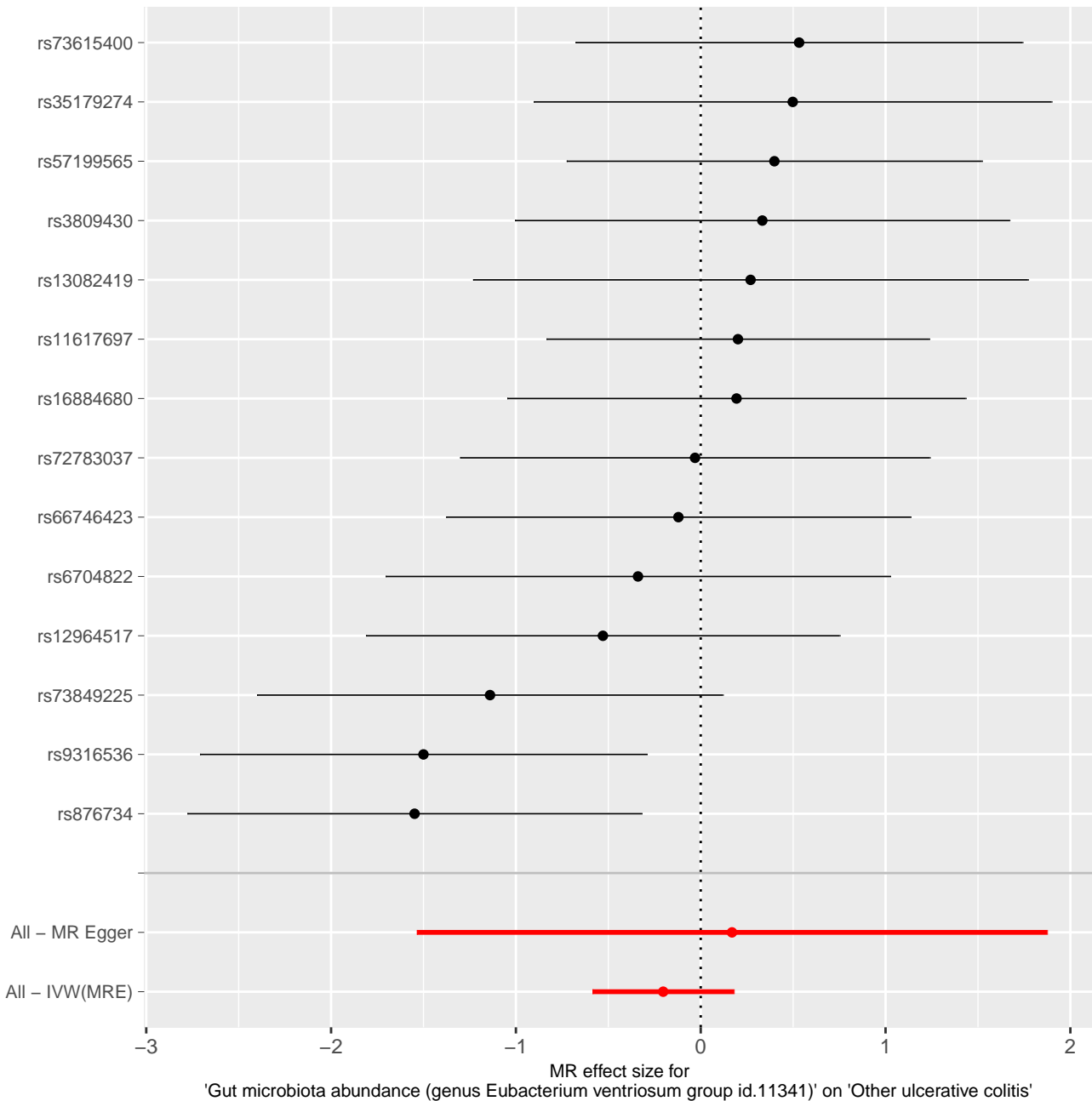


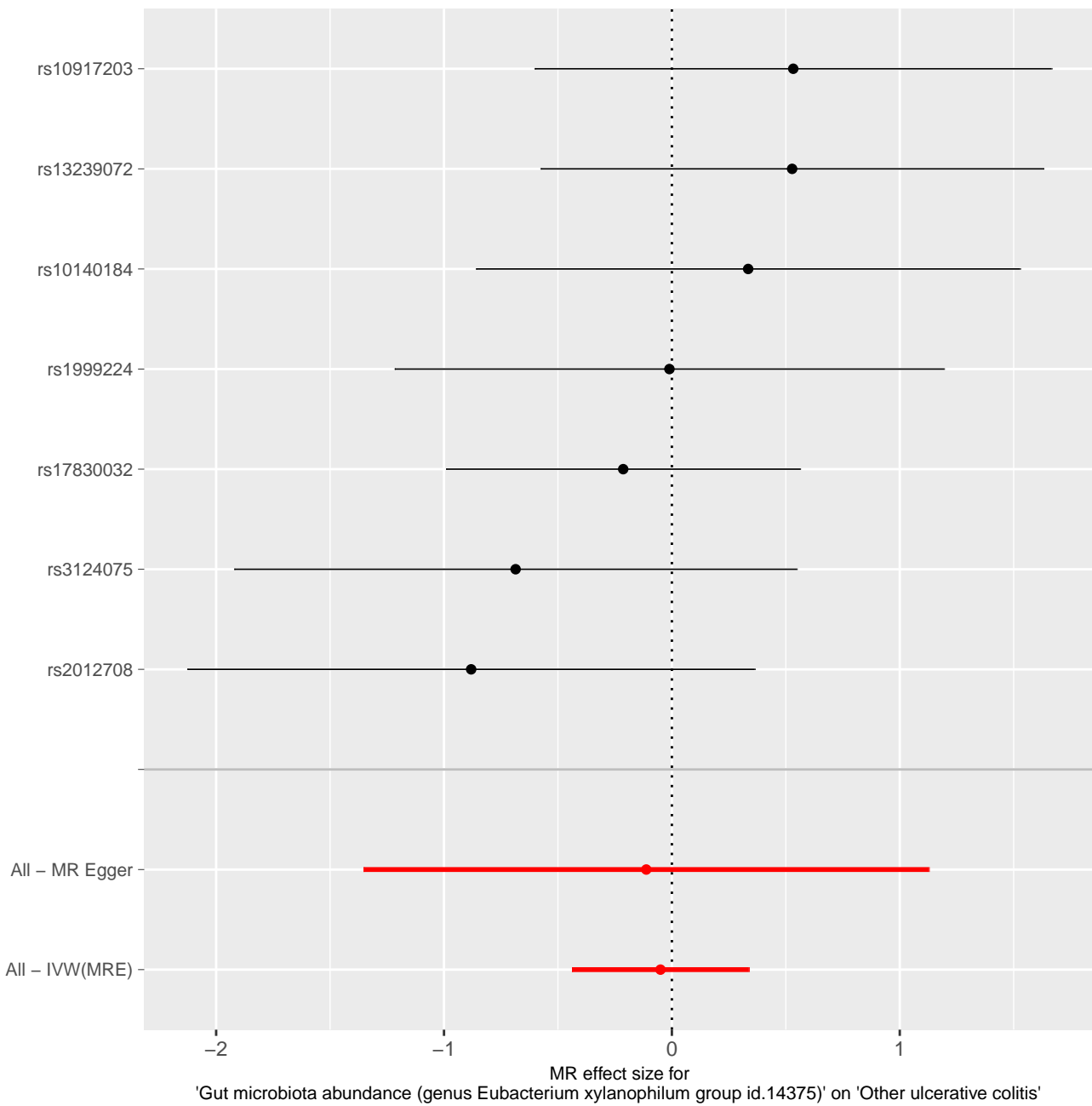




Batch 1278 : Gut microbiota abundance (genus Eubacterium ruminantium group id.11340) on Other ulcerative colitis

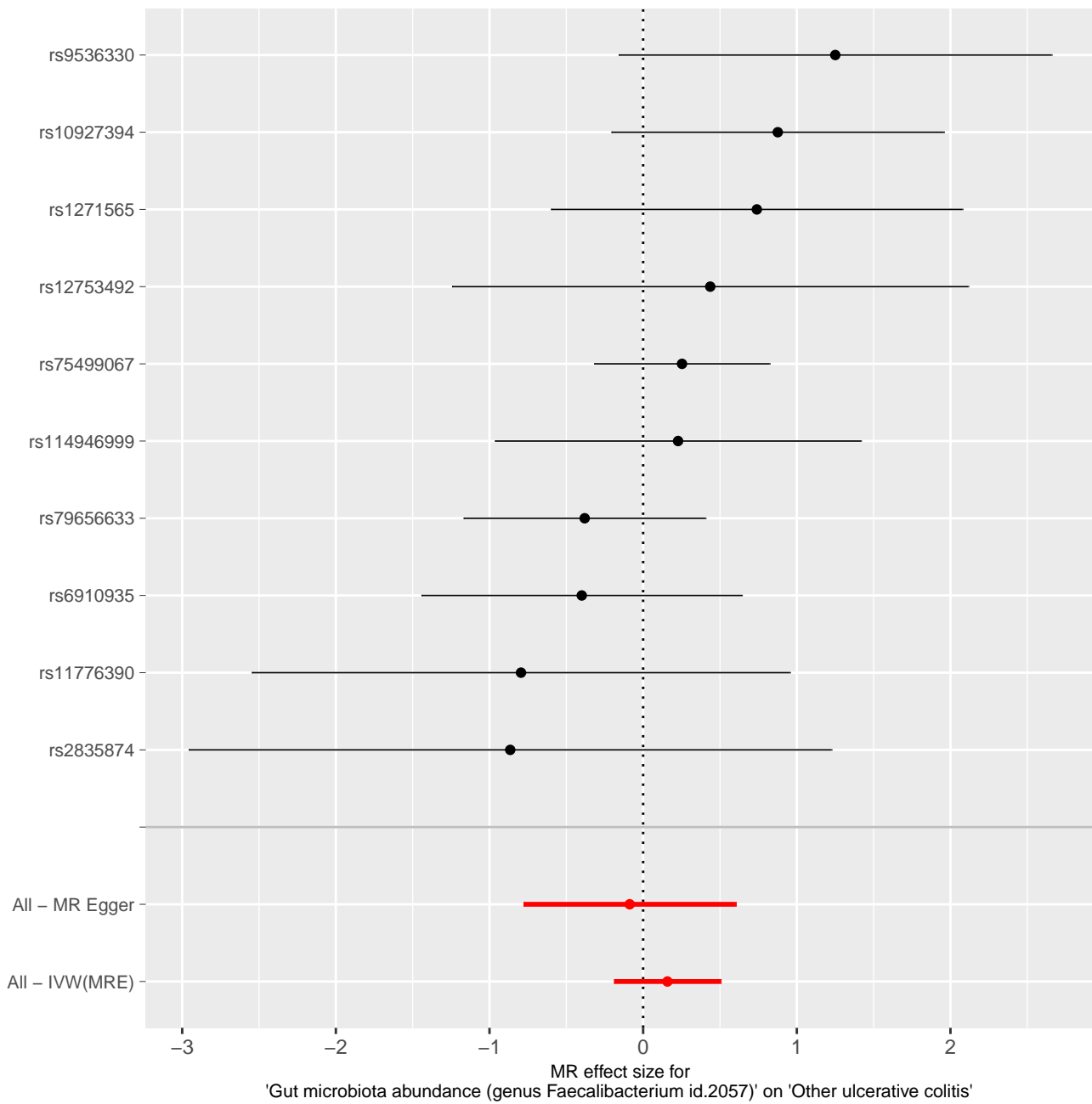


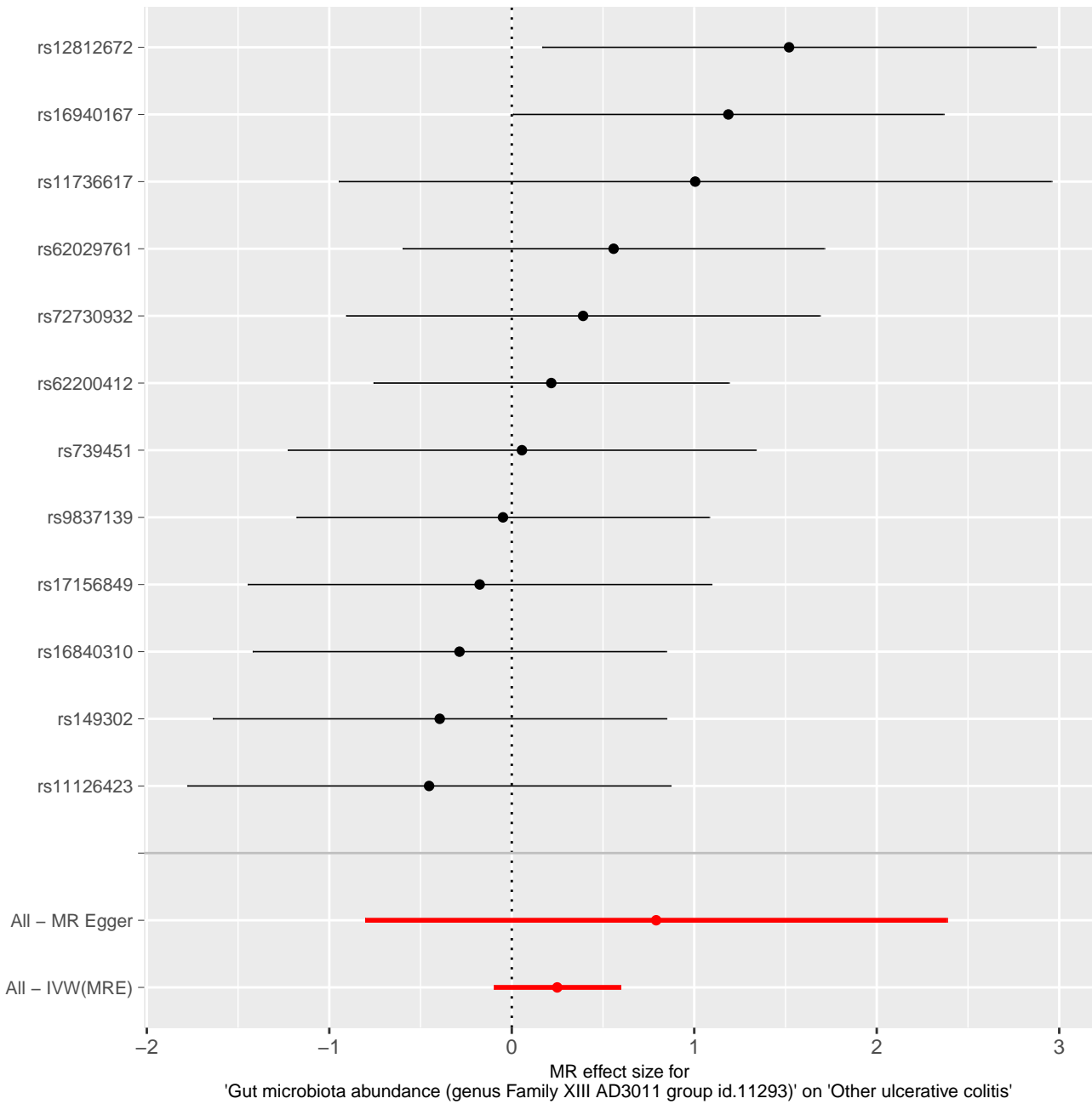


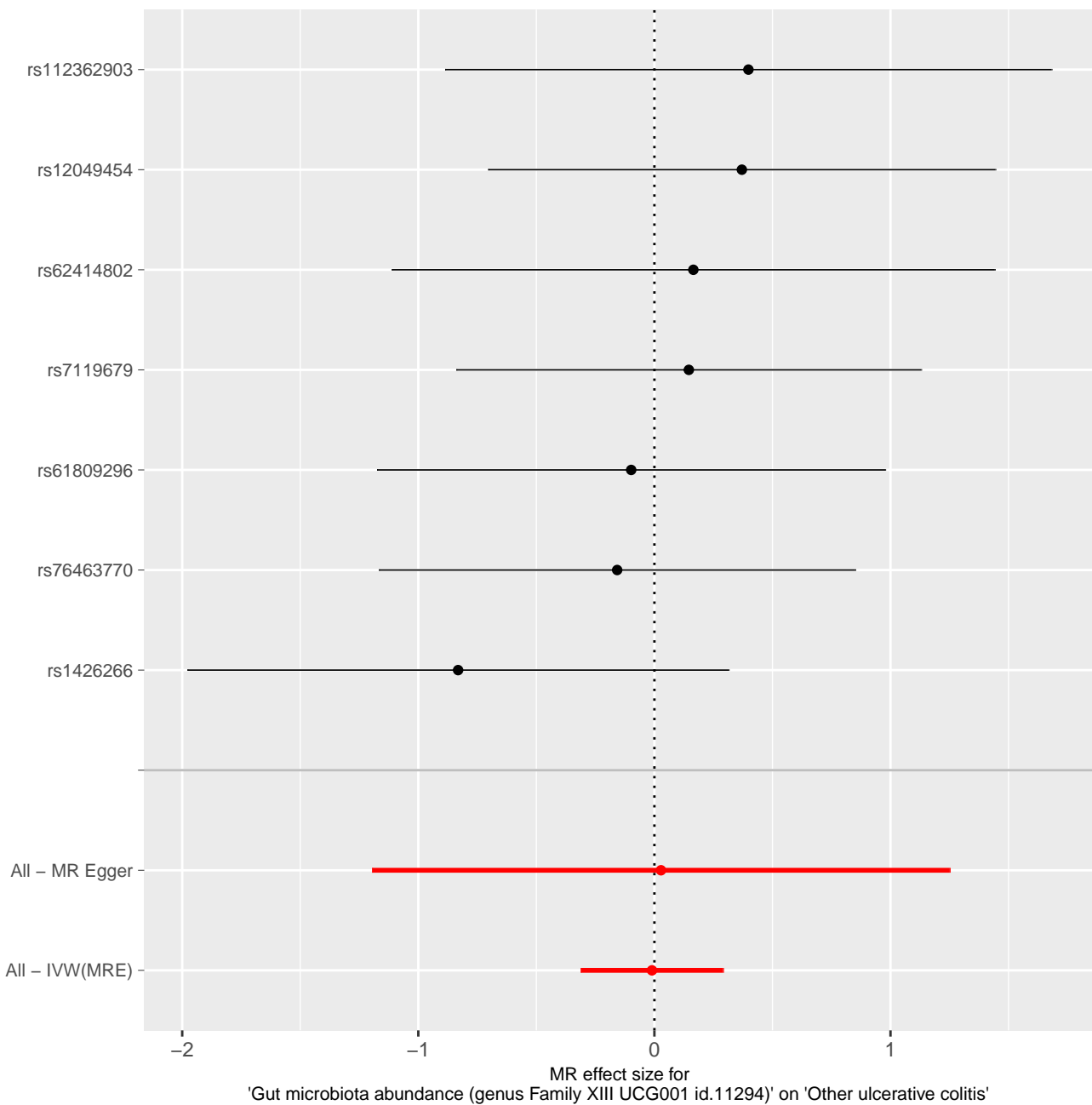


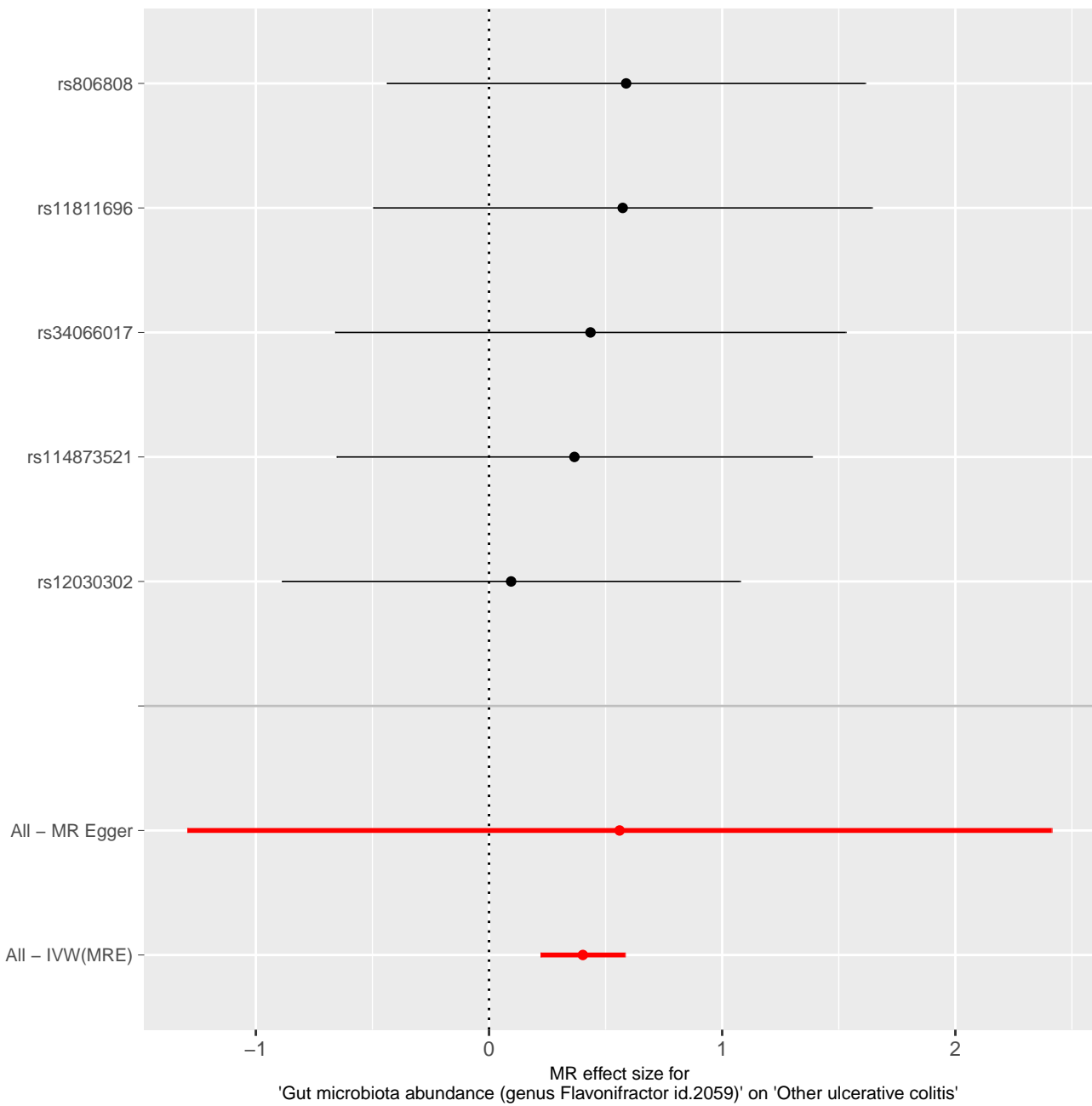
MR effect size for
'Gut microbiota abundance (genus Eubacterium xylanophilum group id.14375)' on 'Other ulcerative colitis'

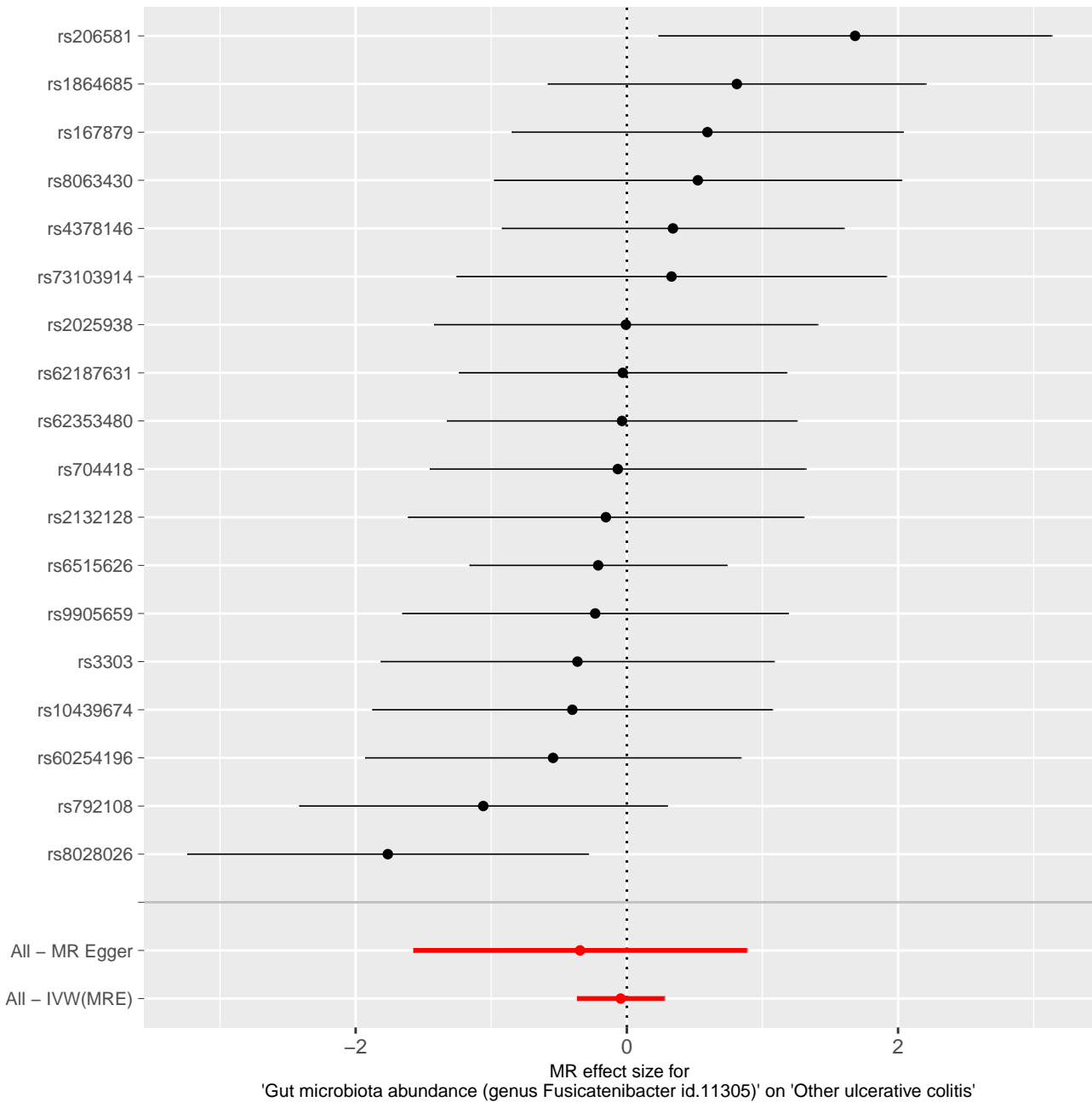
Batch 1281 : Gut microbiota abundance (genus Faecalibacterium id.2057) on Other ulcerative colitis

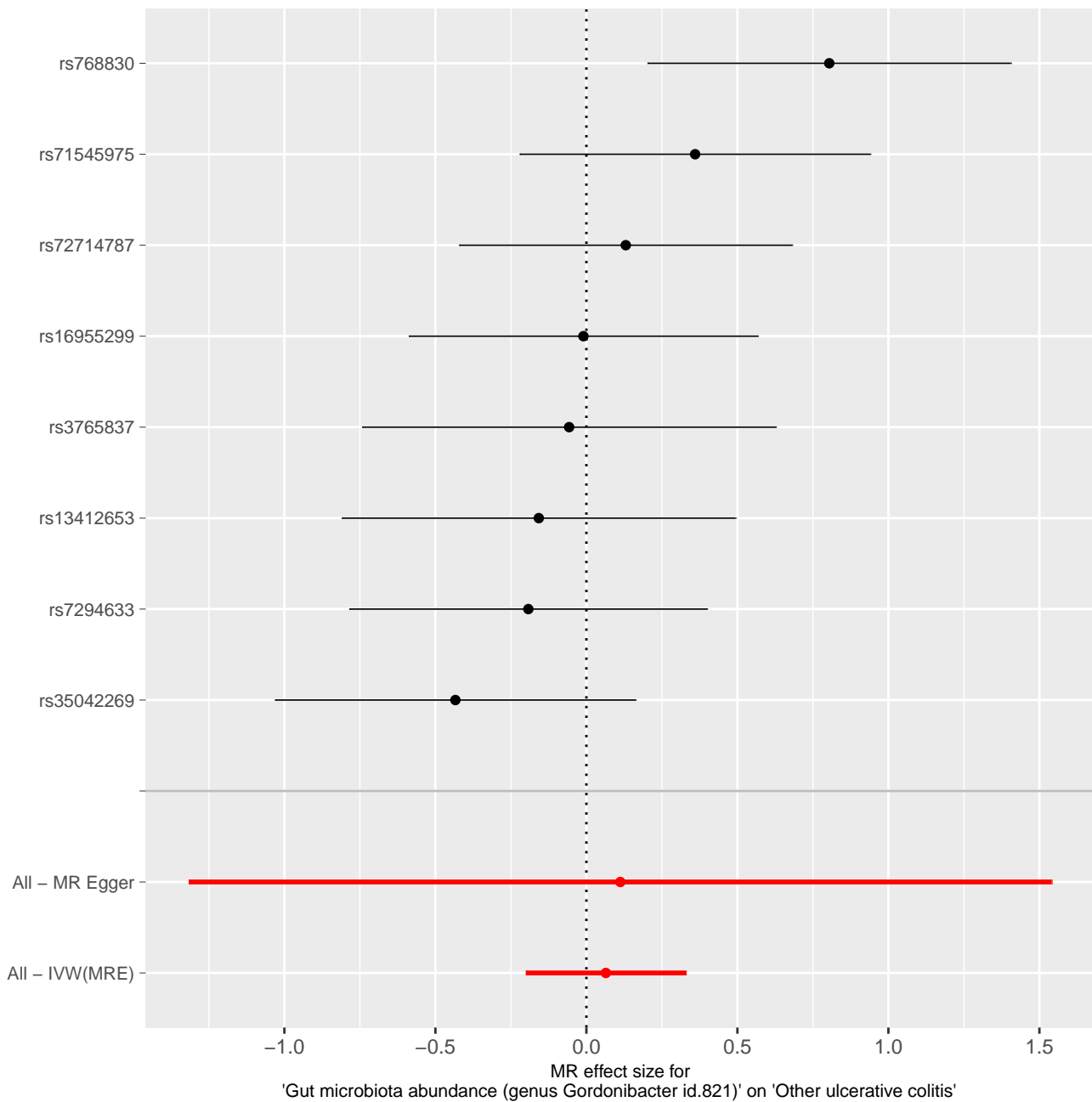


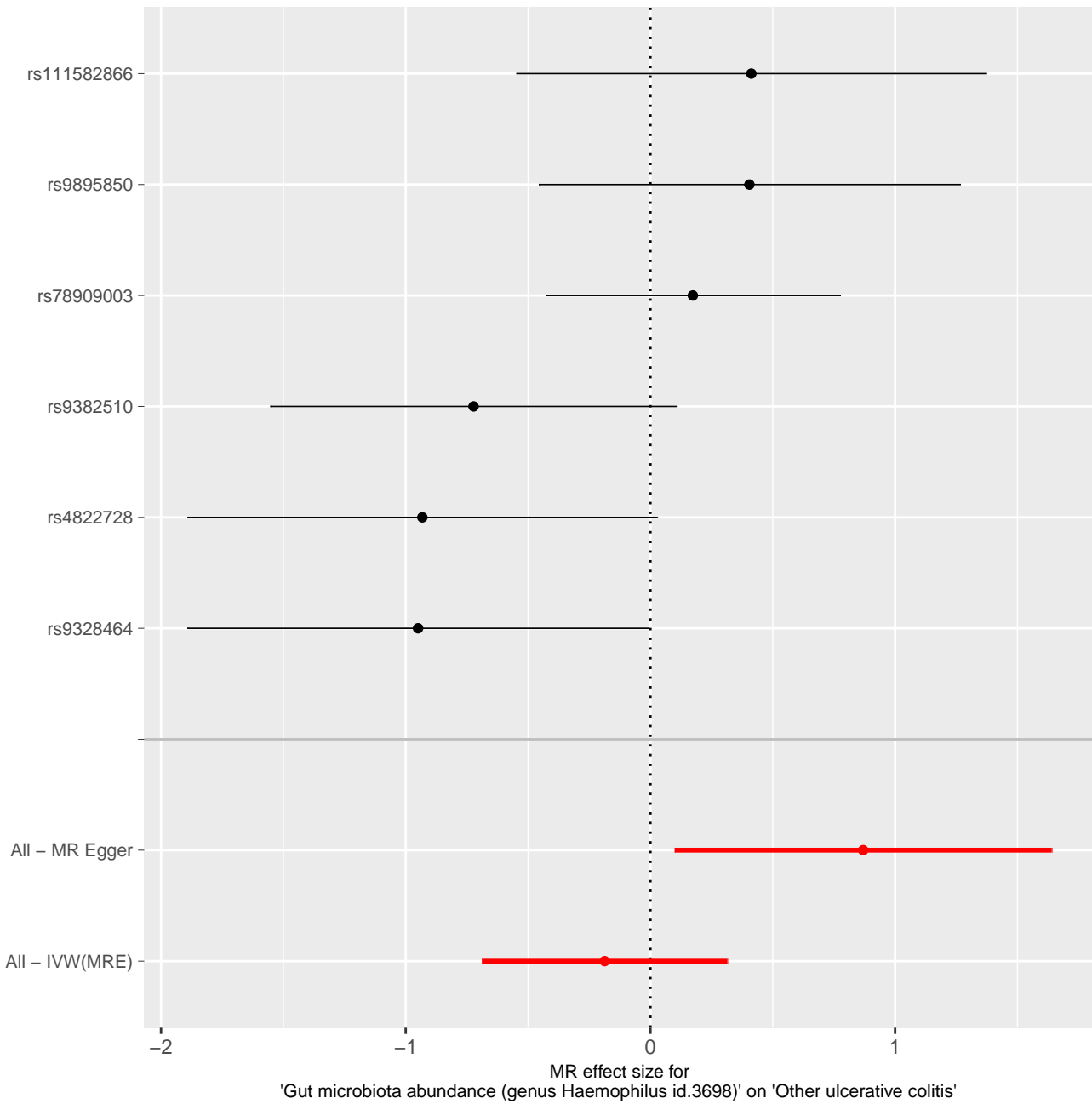


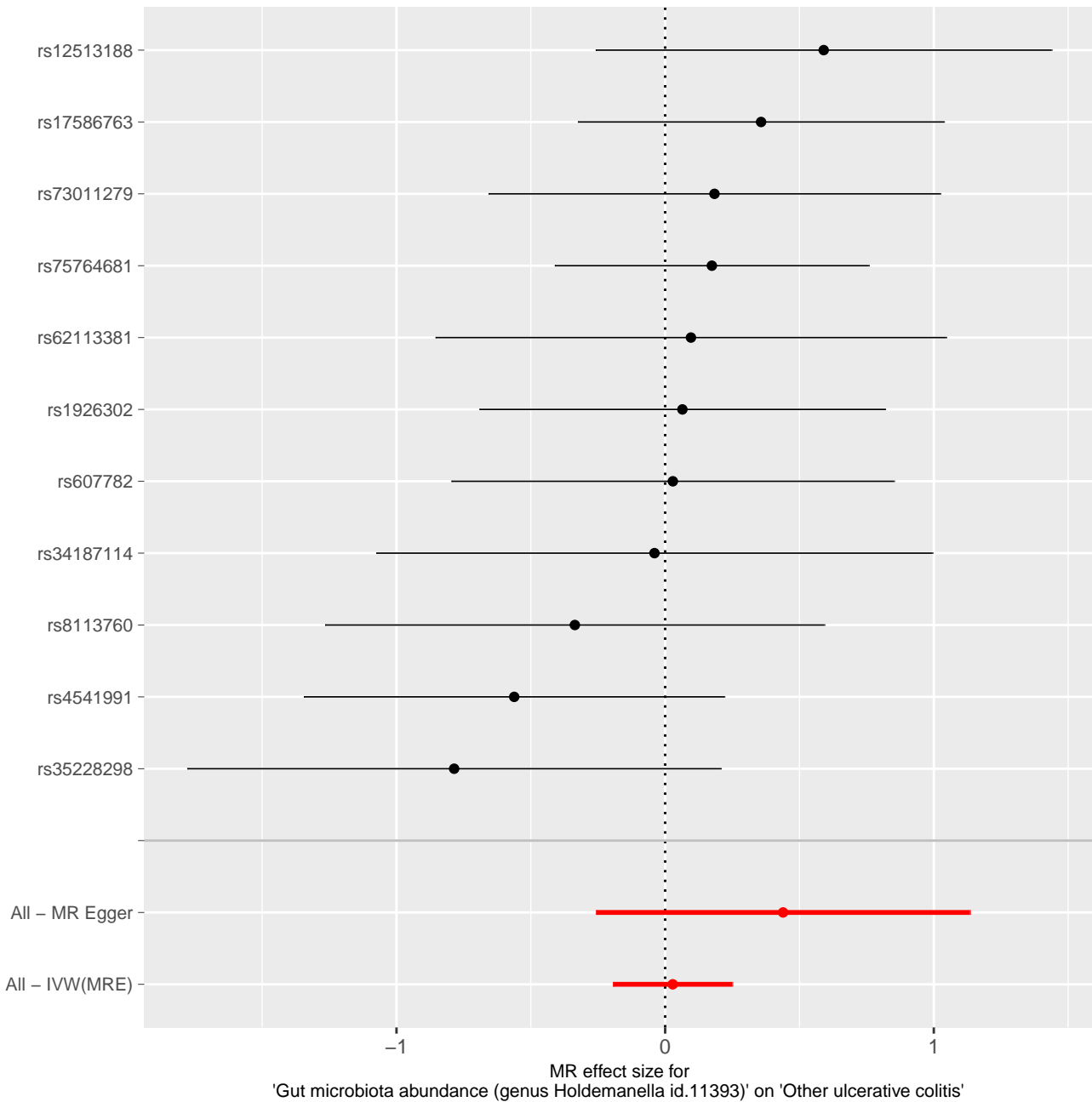


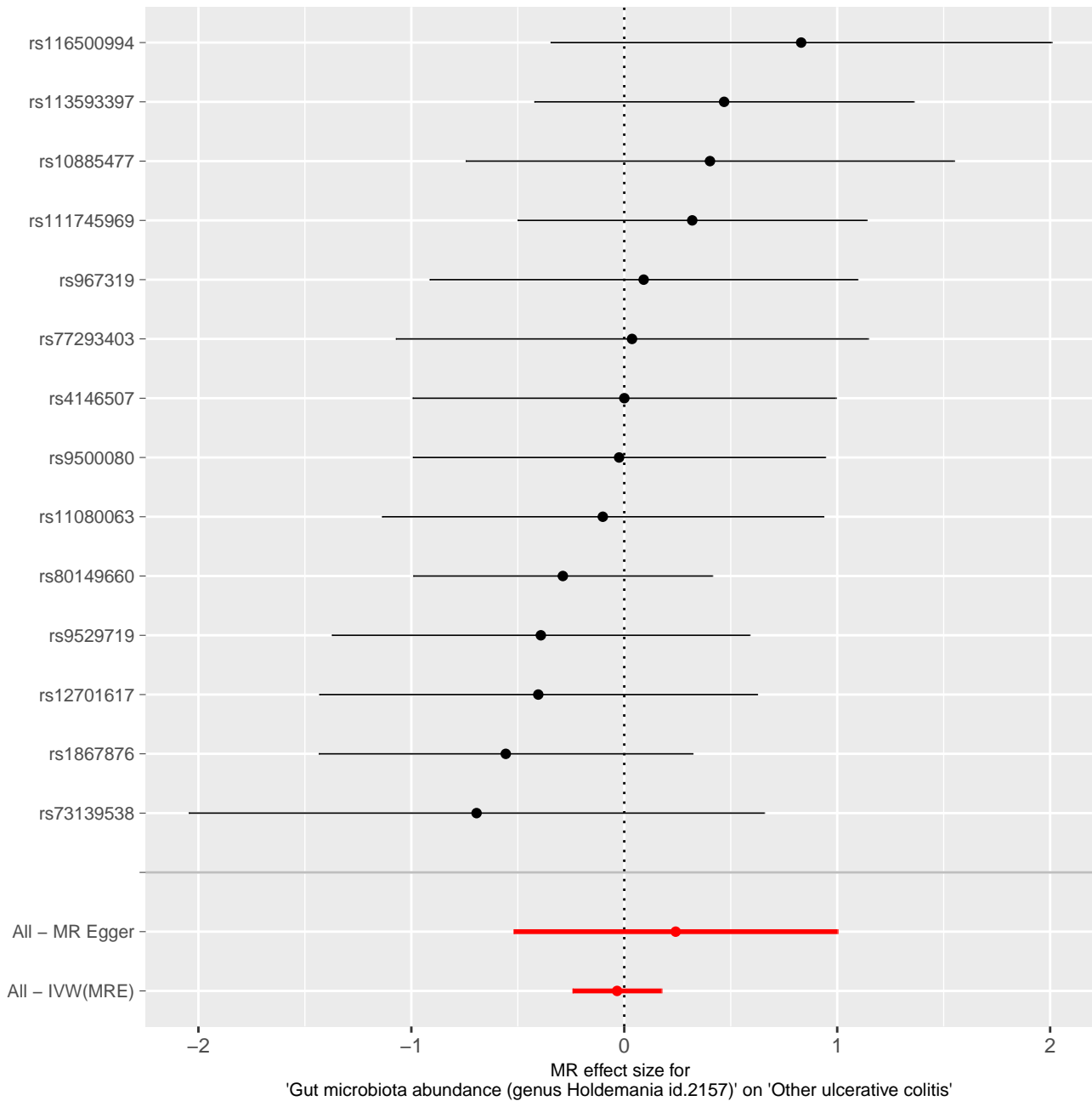


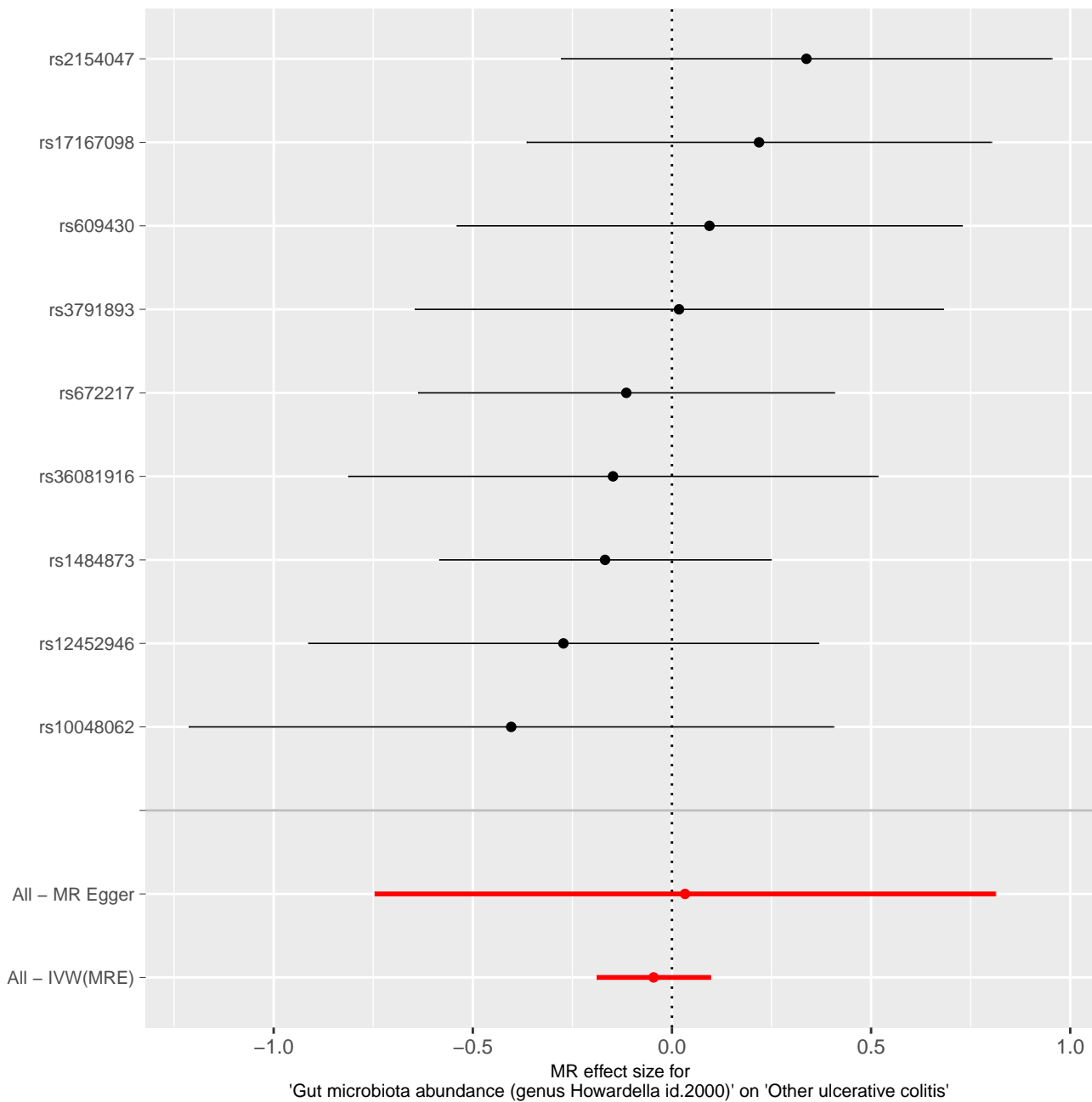


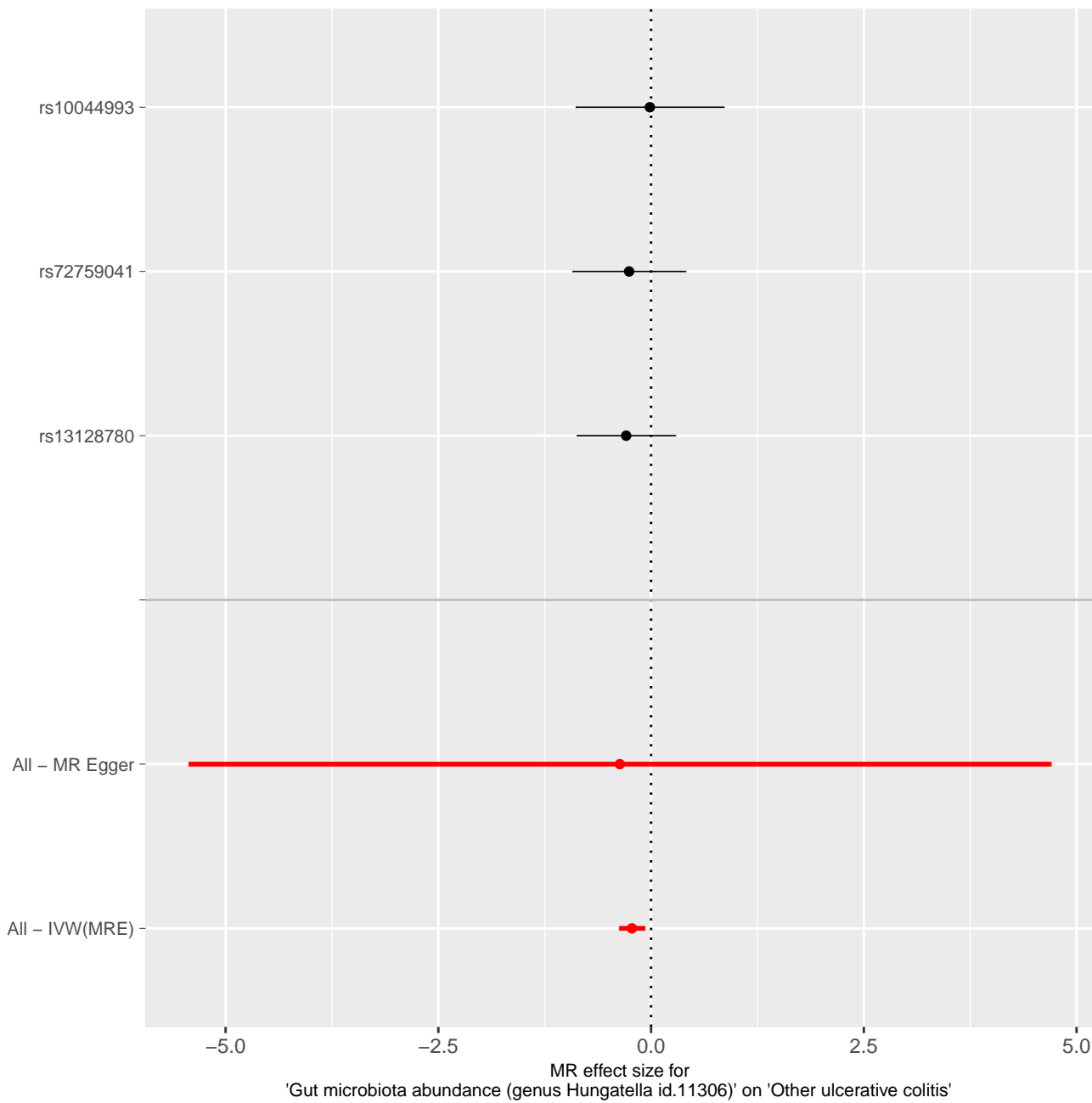




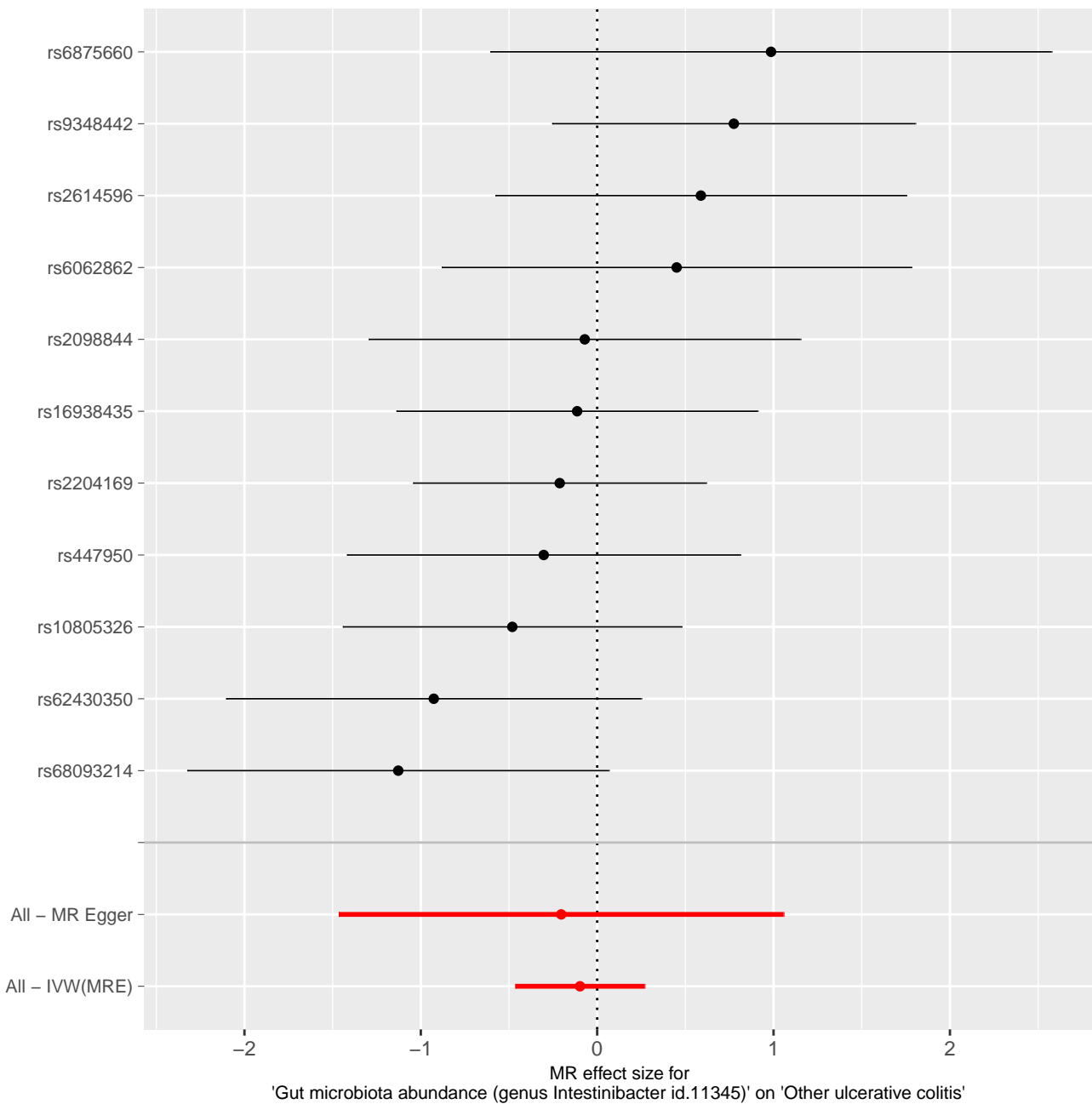


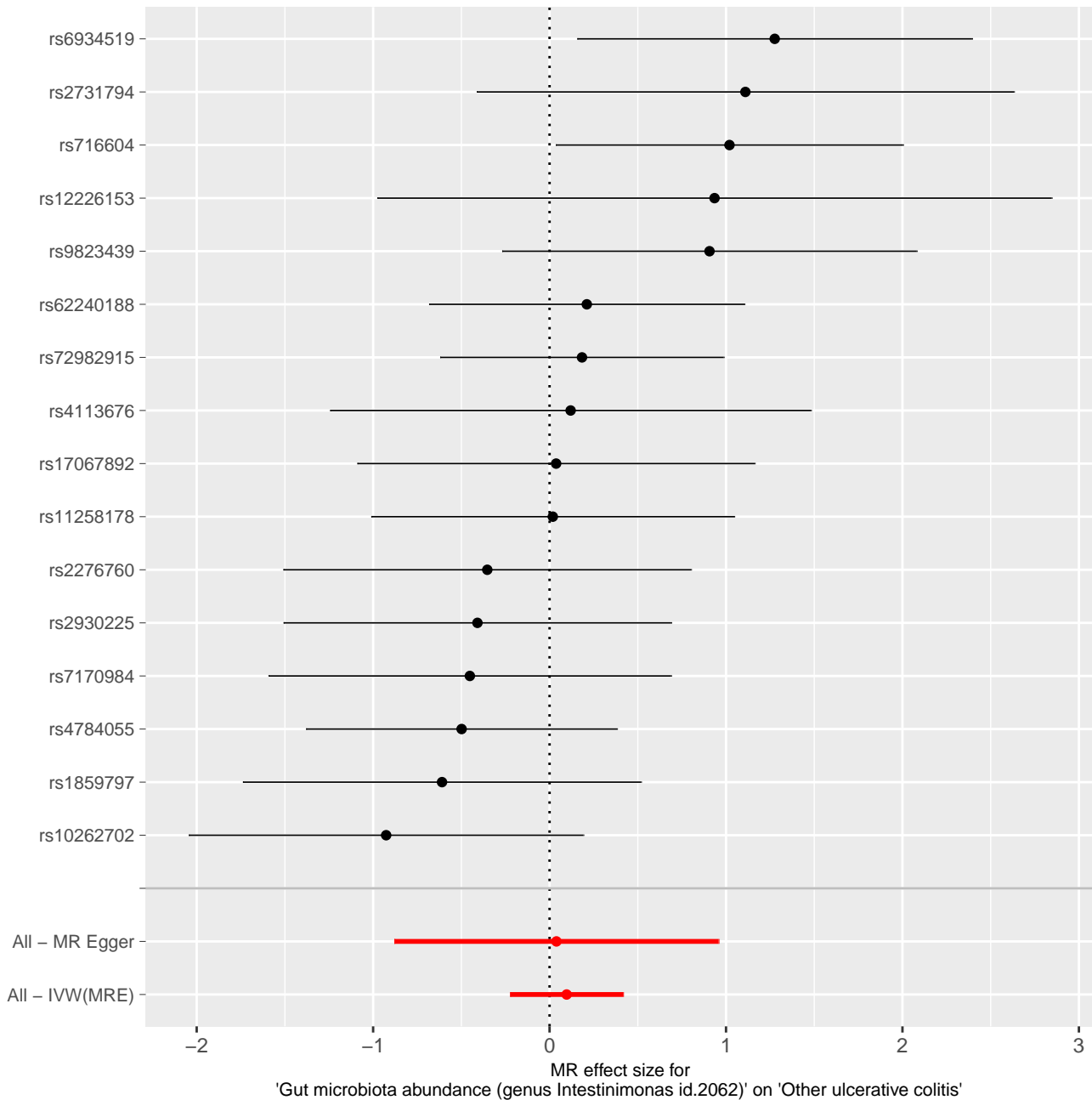


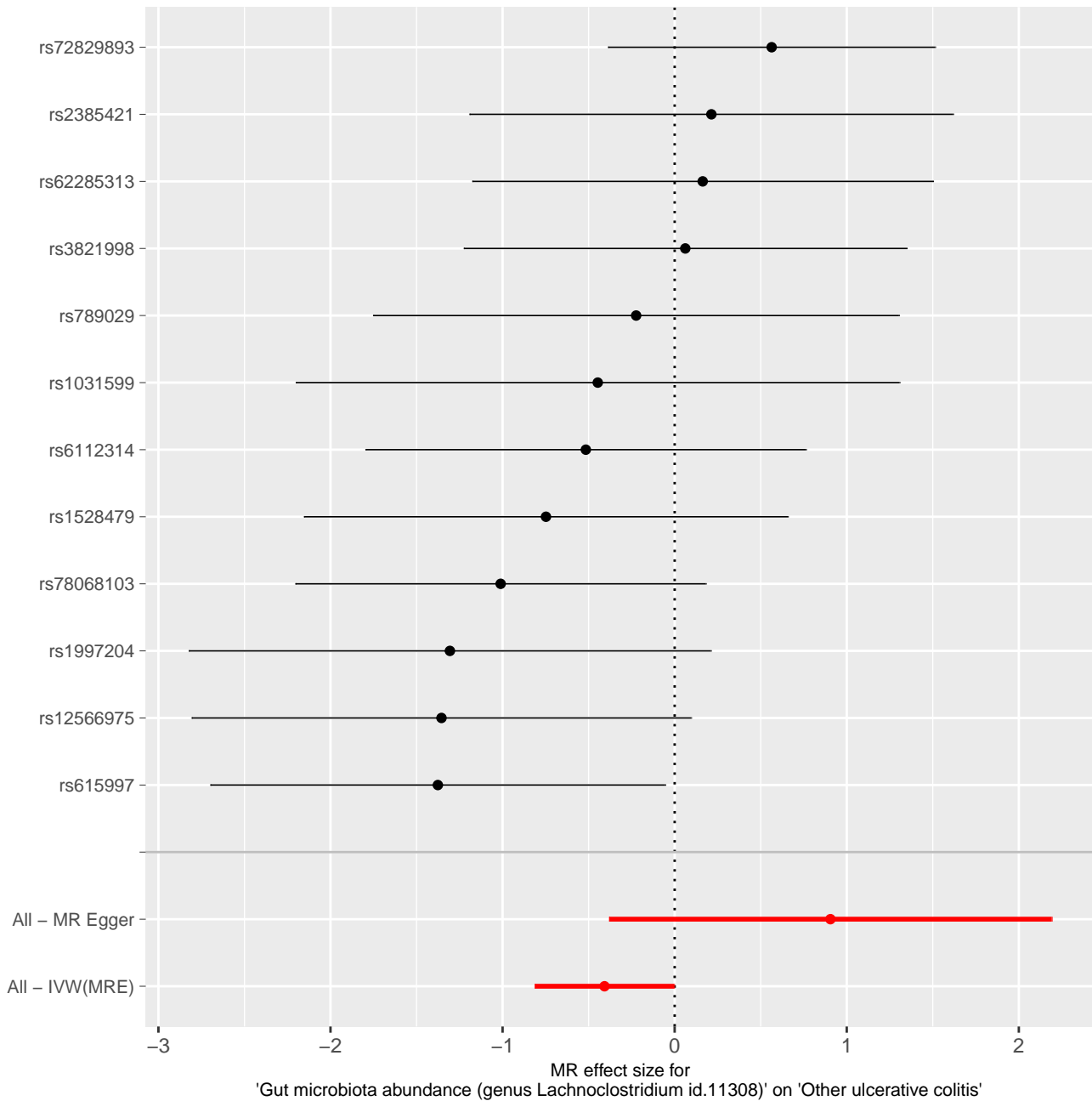


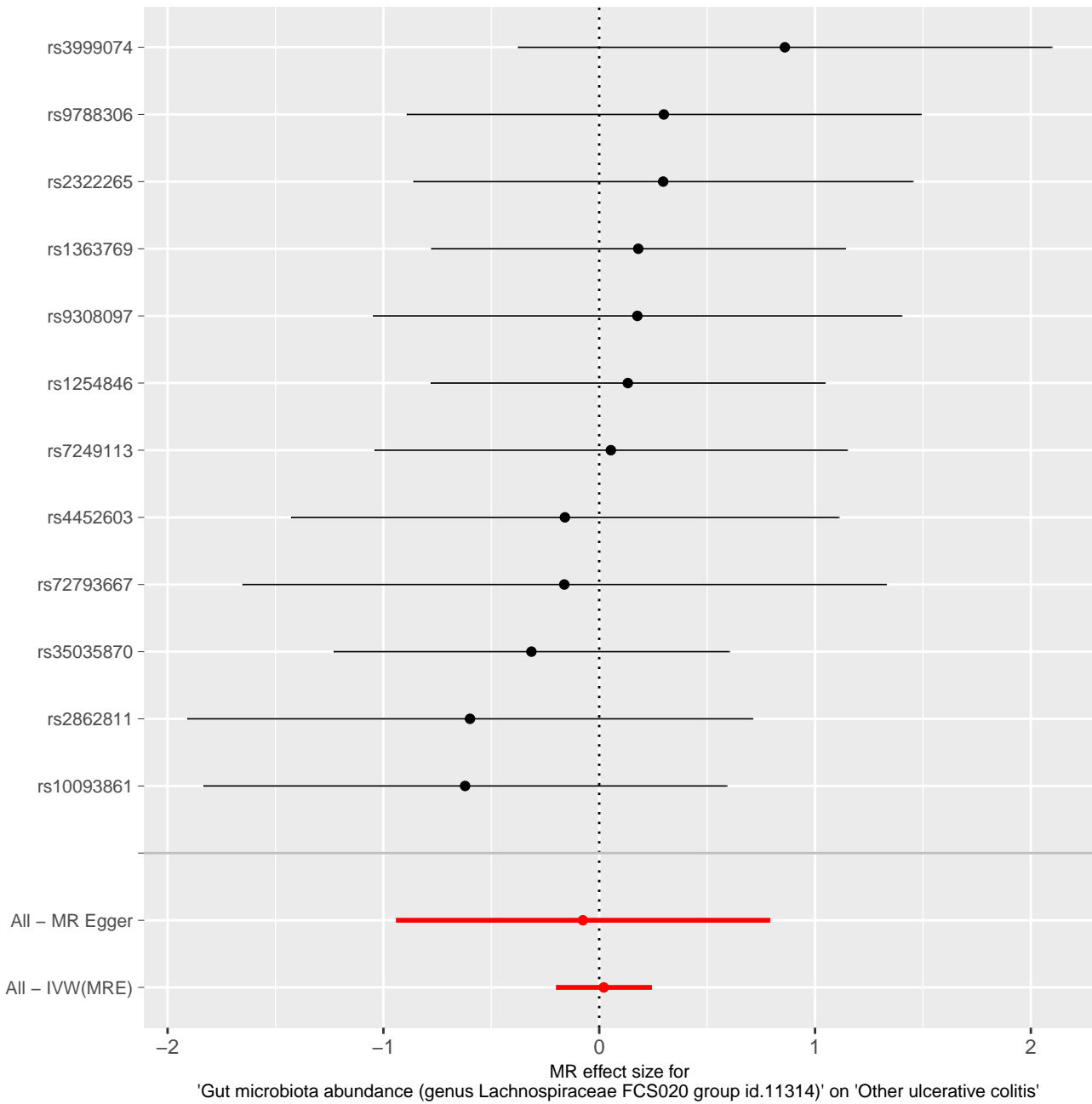


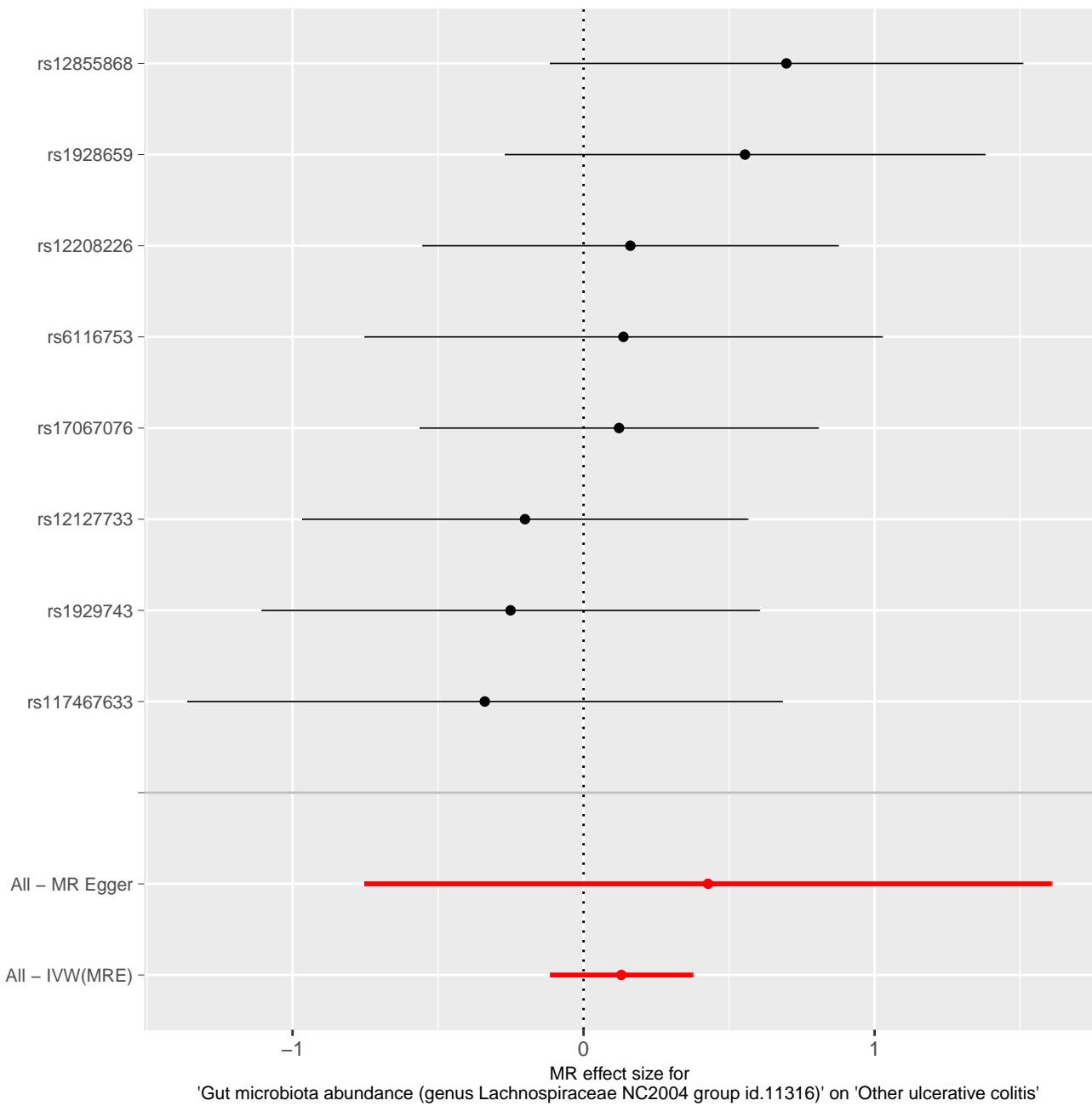
Batch 1292 : Gut microbiota abundance (genus Intestinibacter id.11345) on Other ulcerative colitis



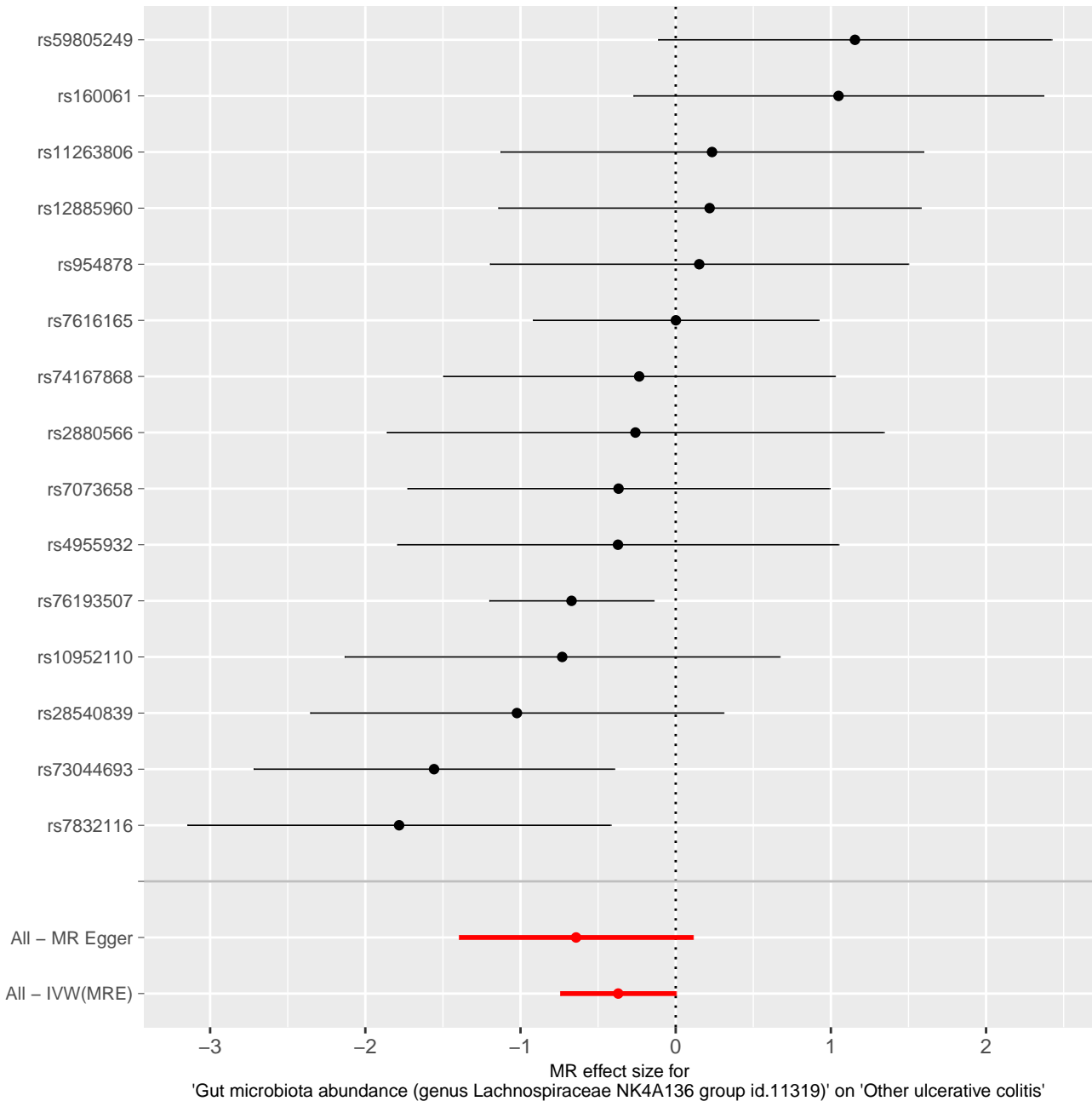


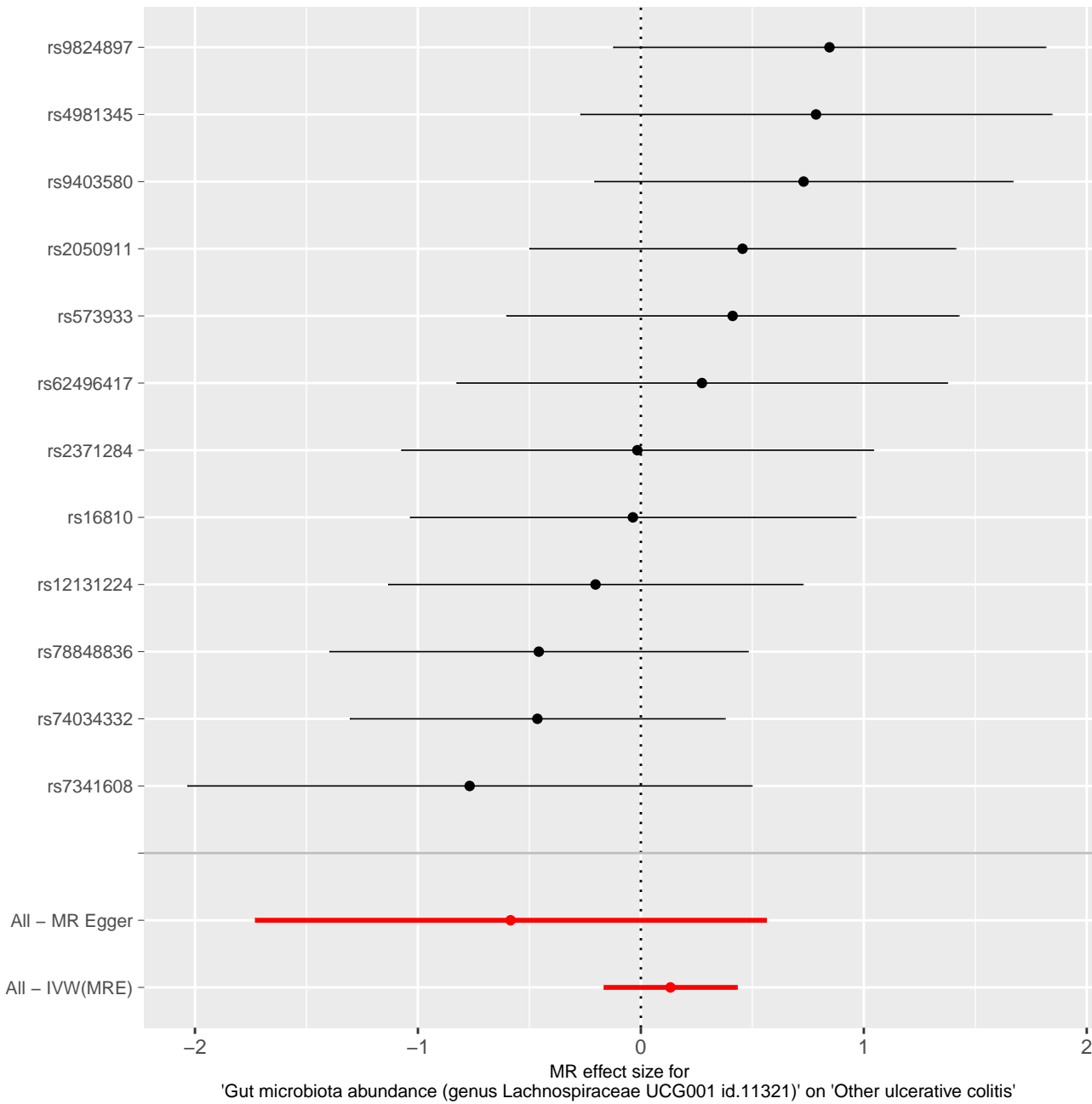


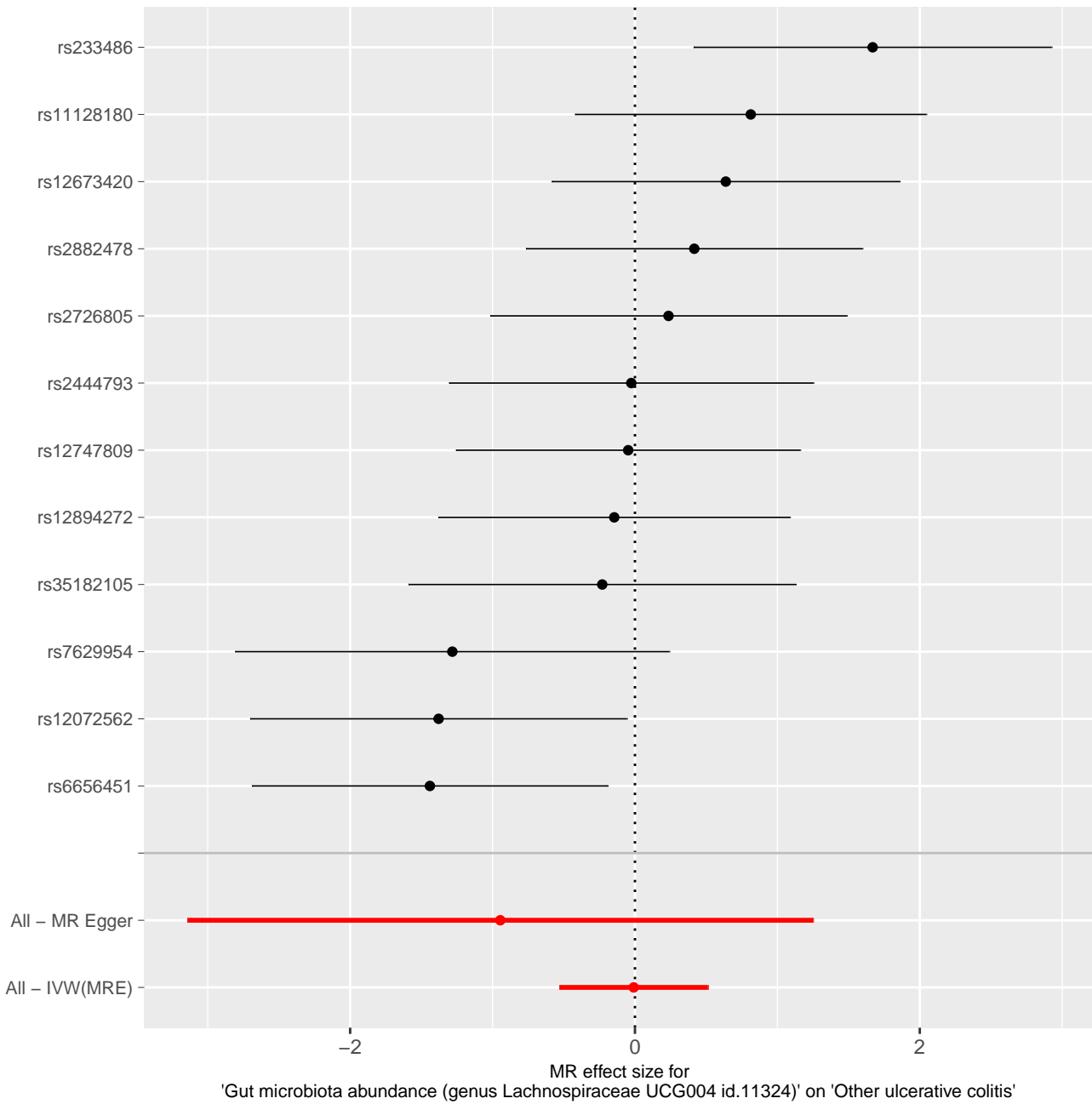


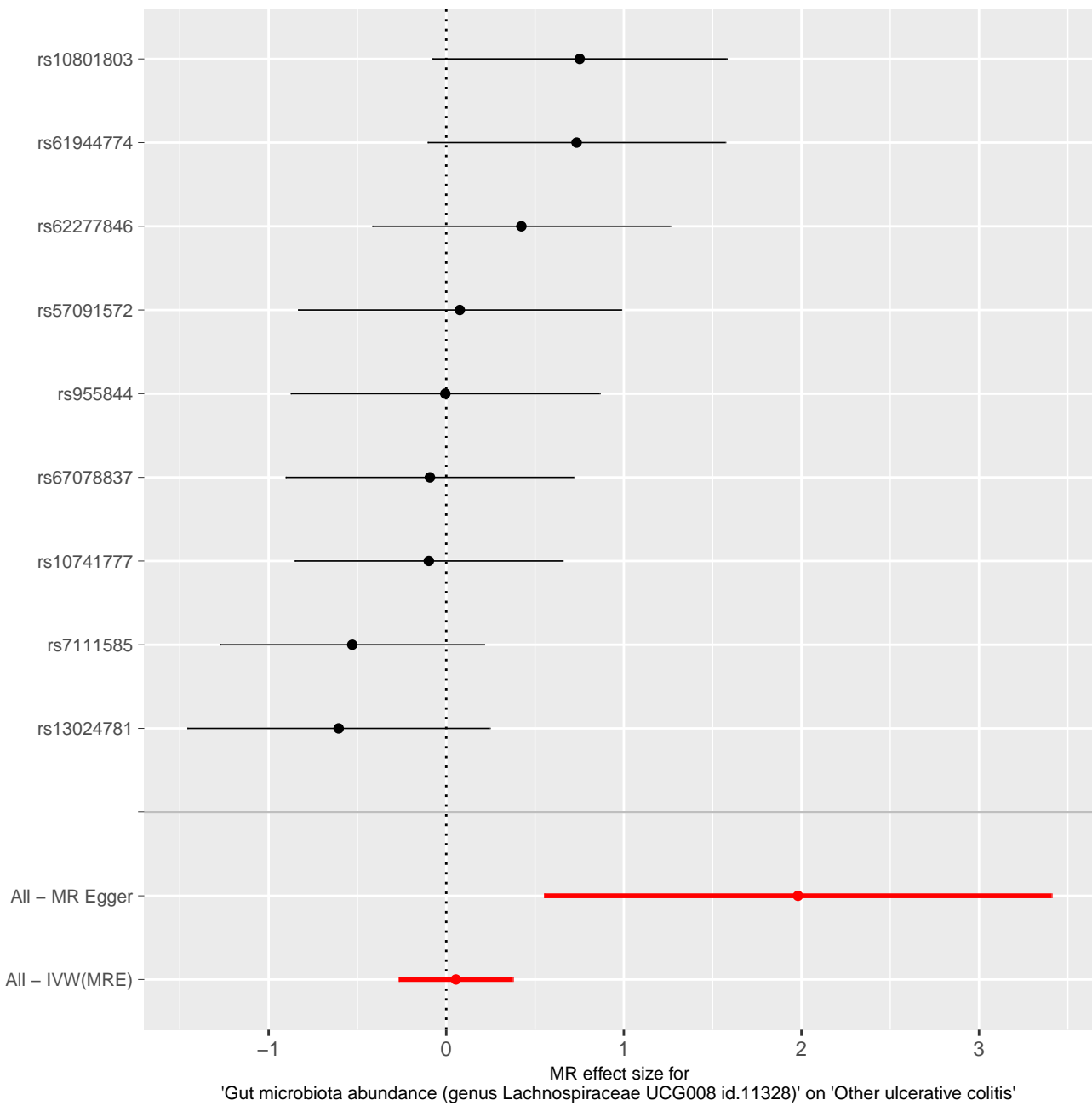


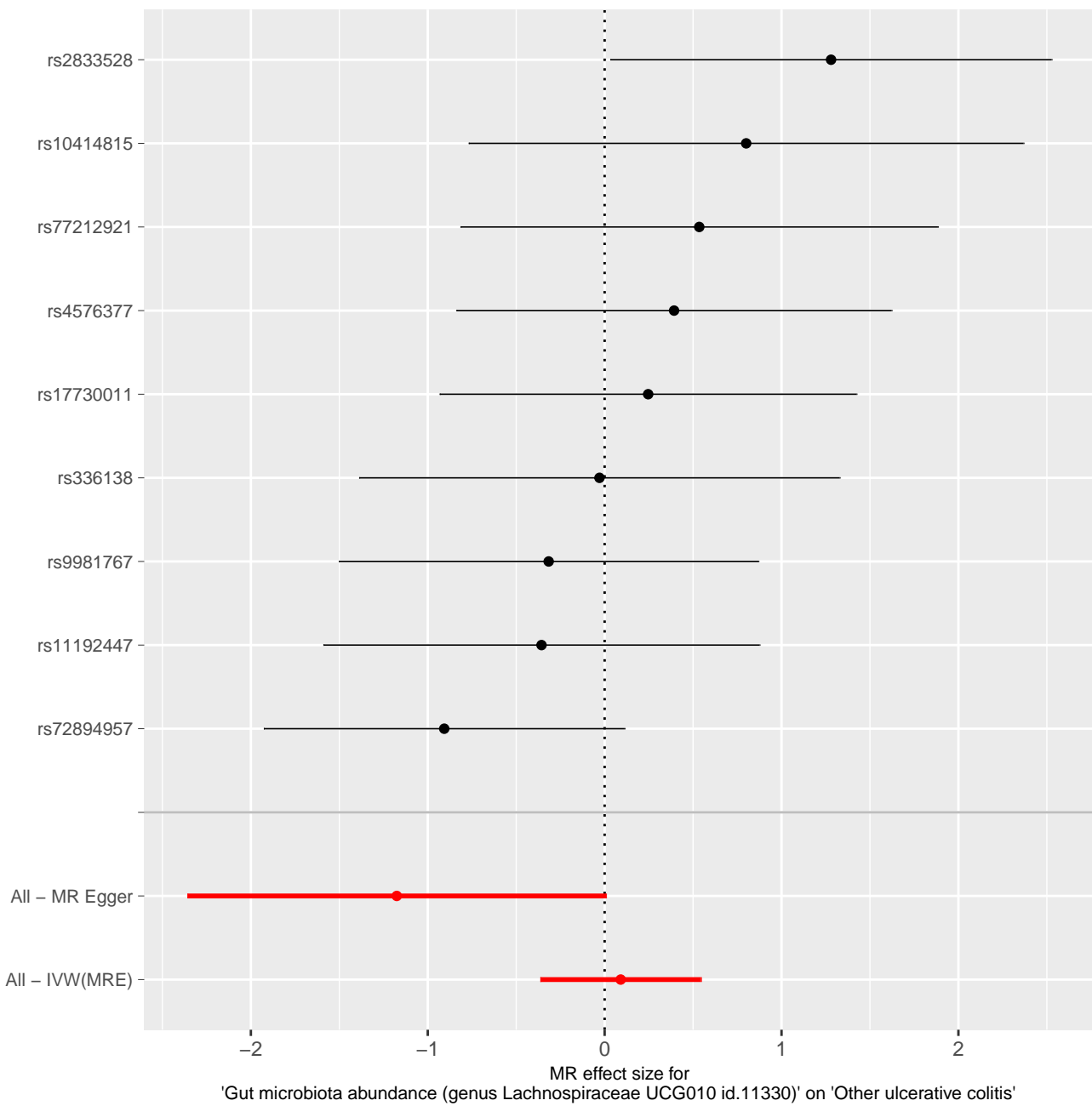
MR effect size for 'Gut microbiota abundance (genus Lachnospiraceae NC2004 group id.11316)' on 'Other ulcerative colitis'

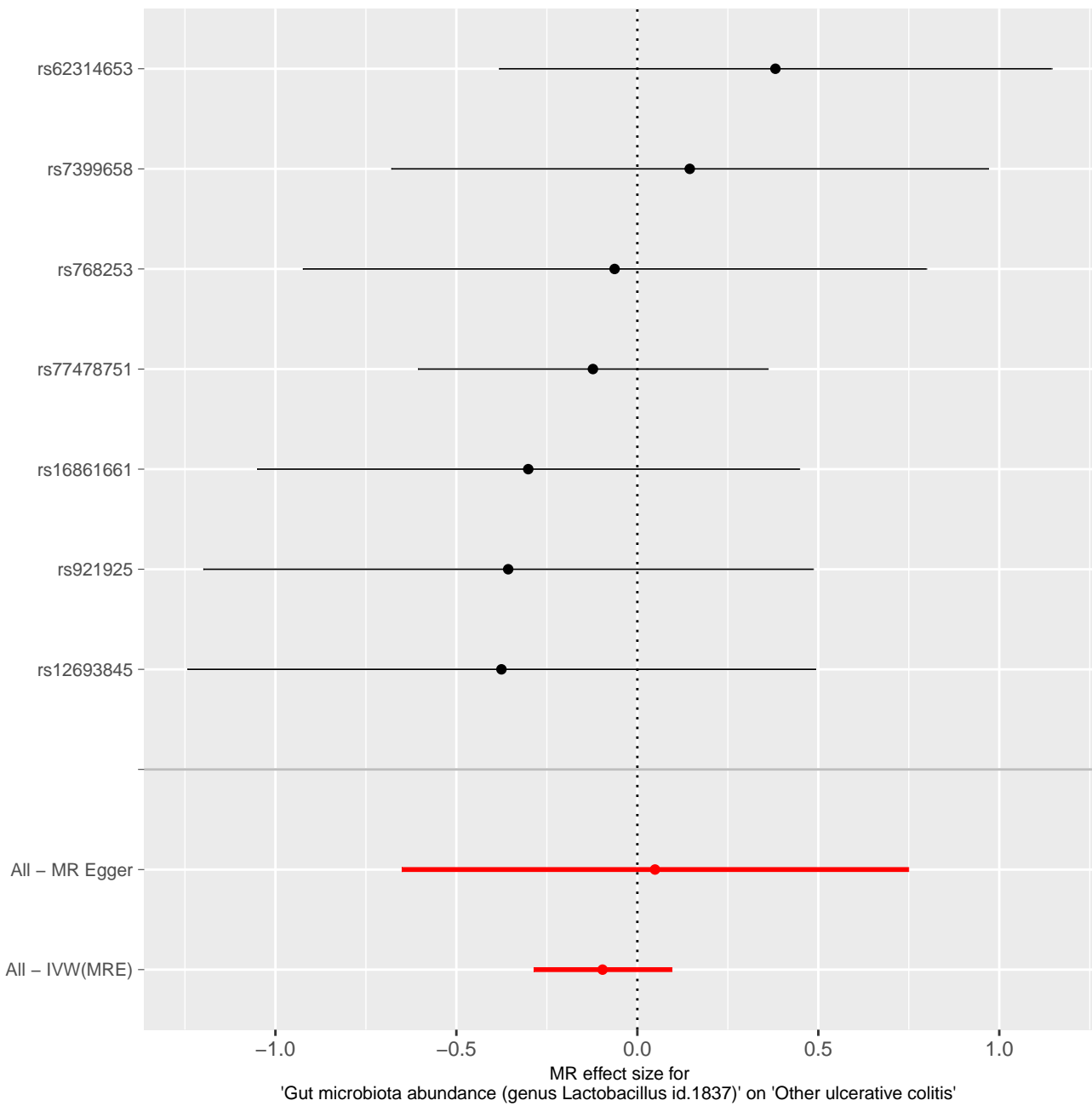


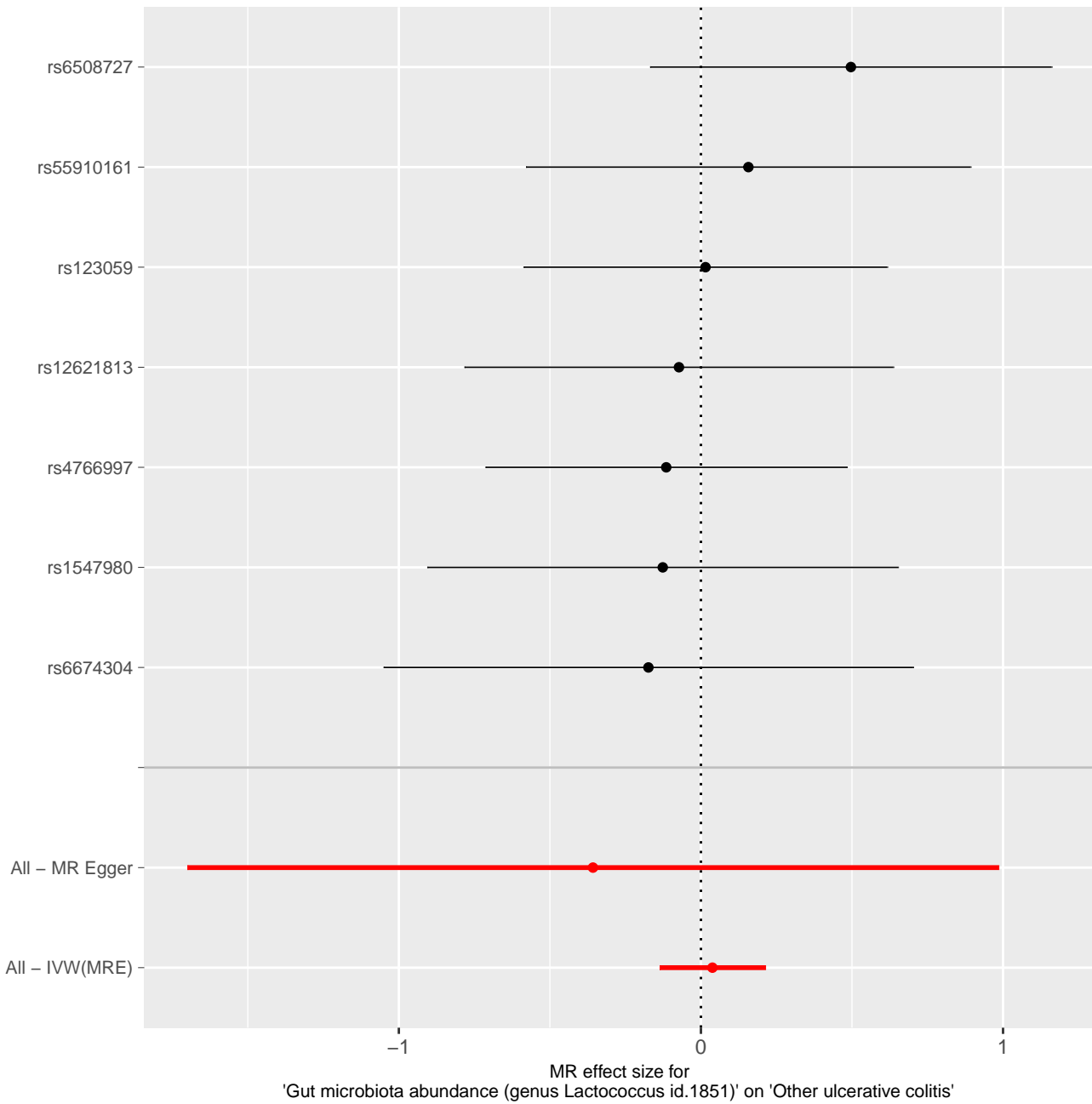


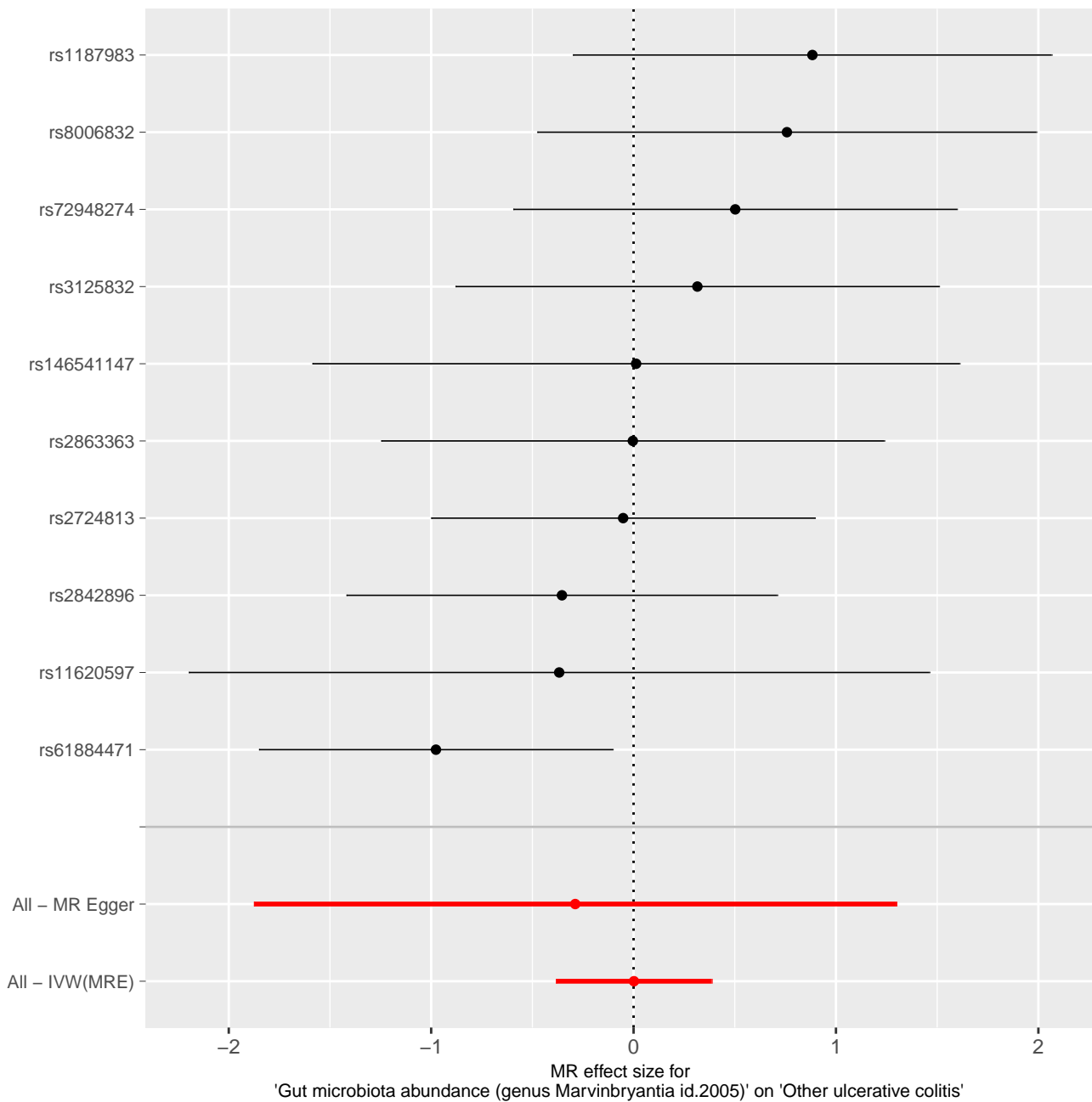


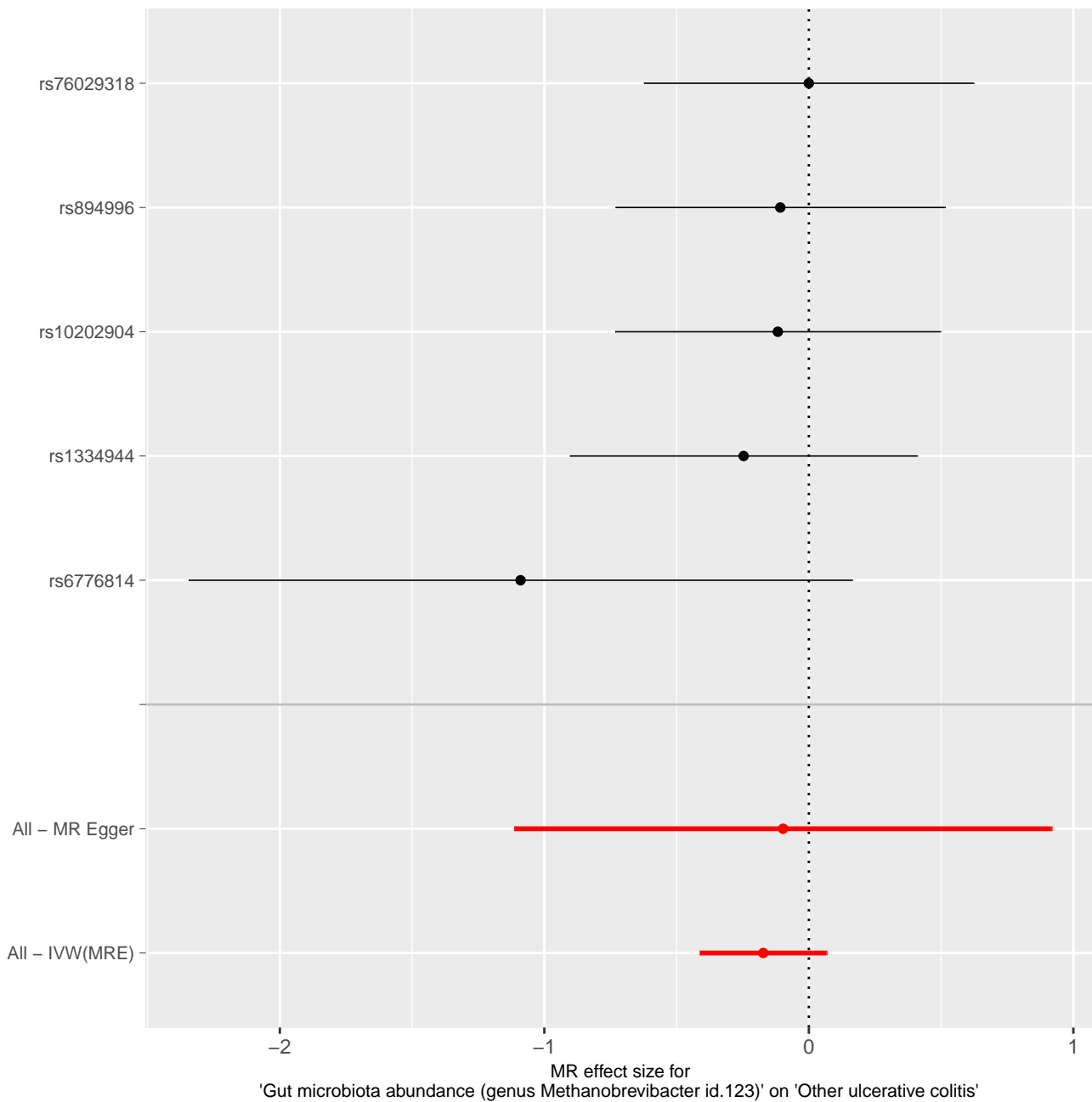




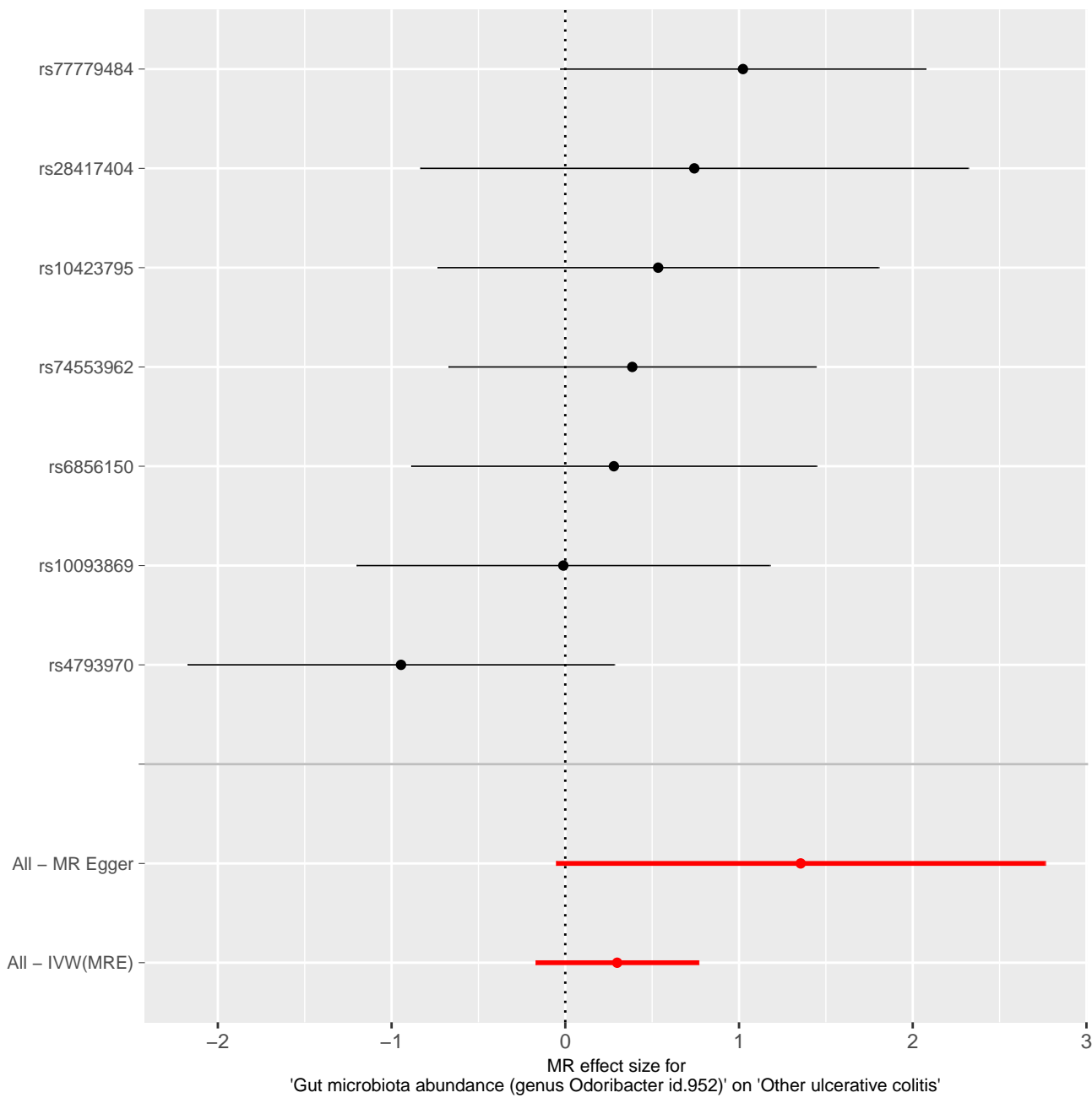


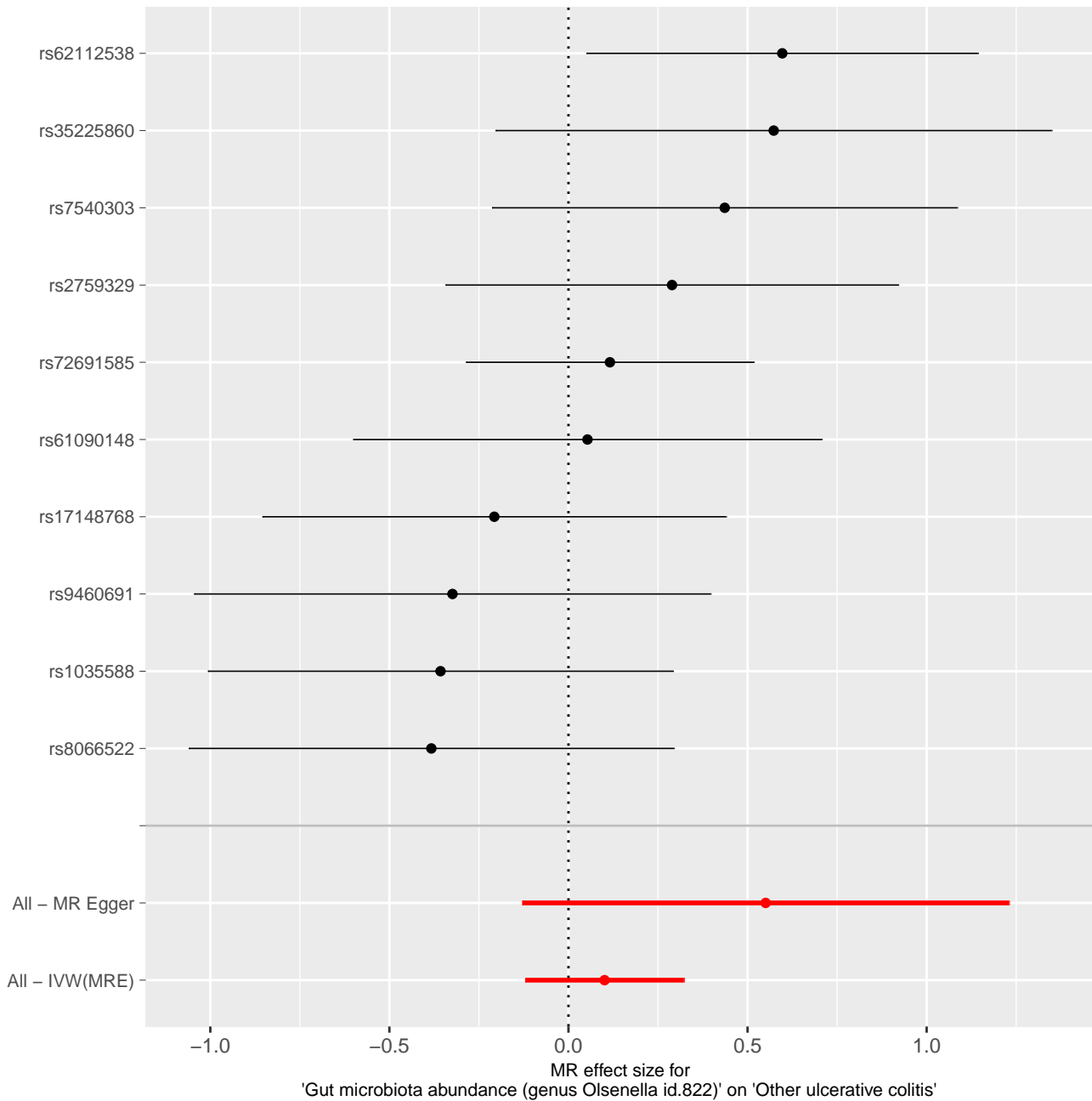


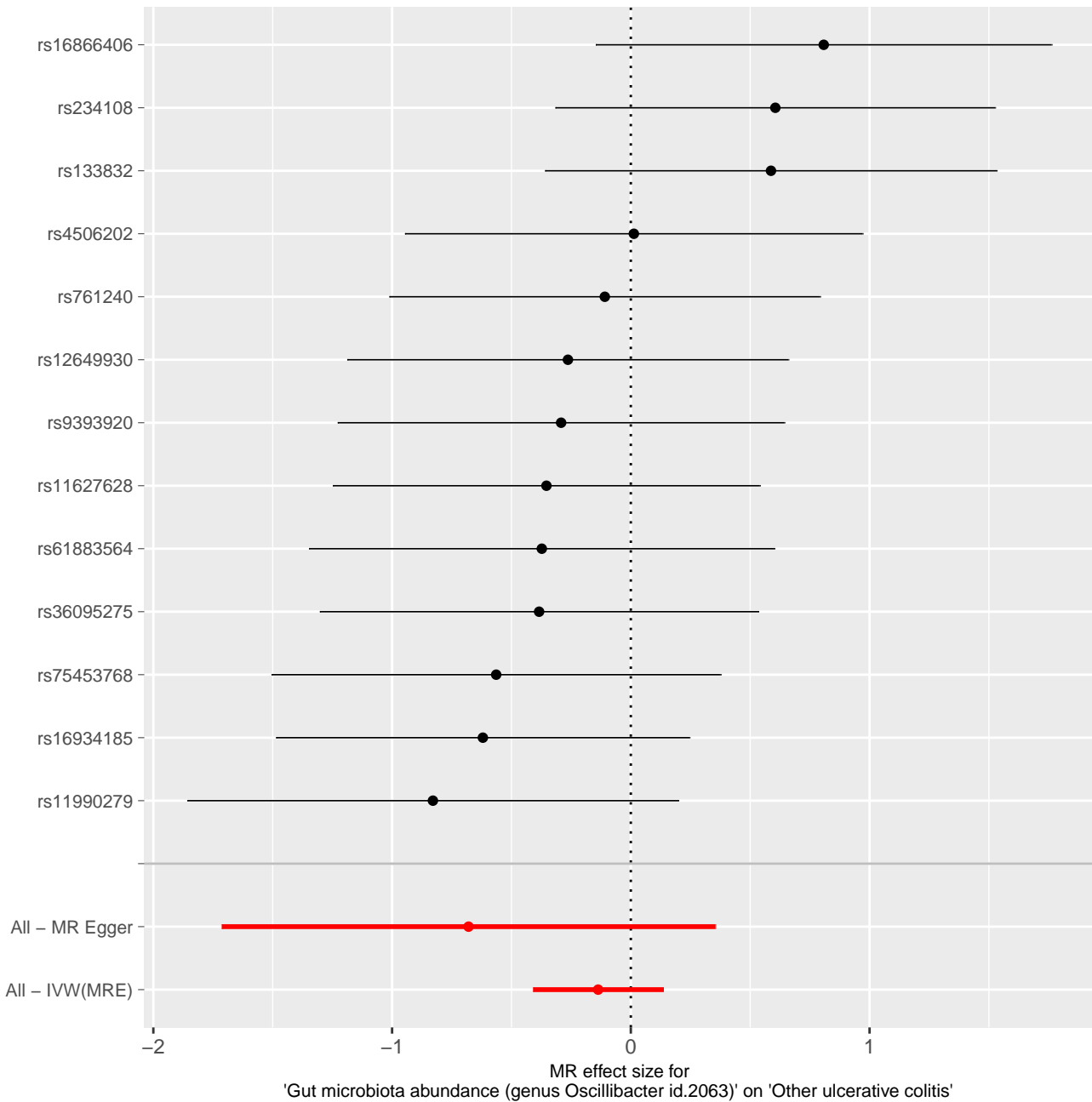


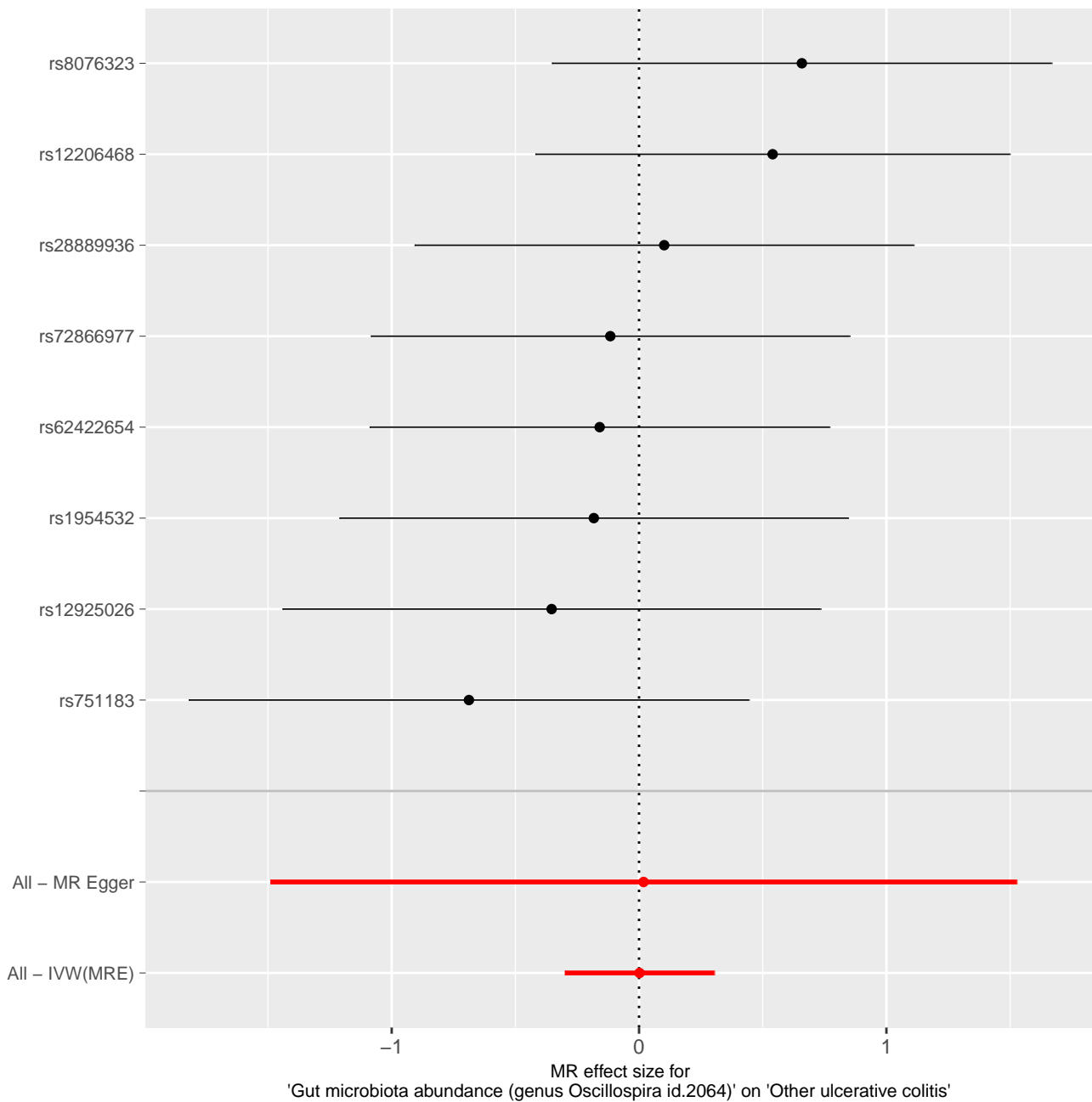


Batch 1308 : Gut microbiota abundance (genus Odoribacter id.952) on Other ulcerative colitis

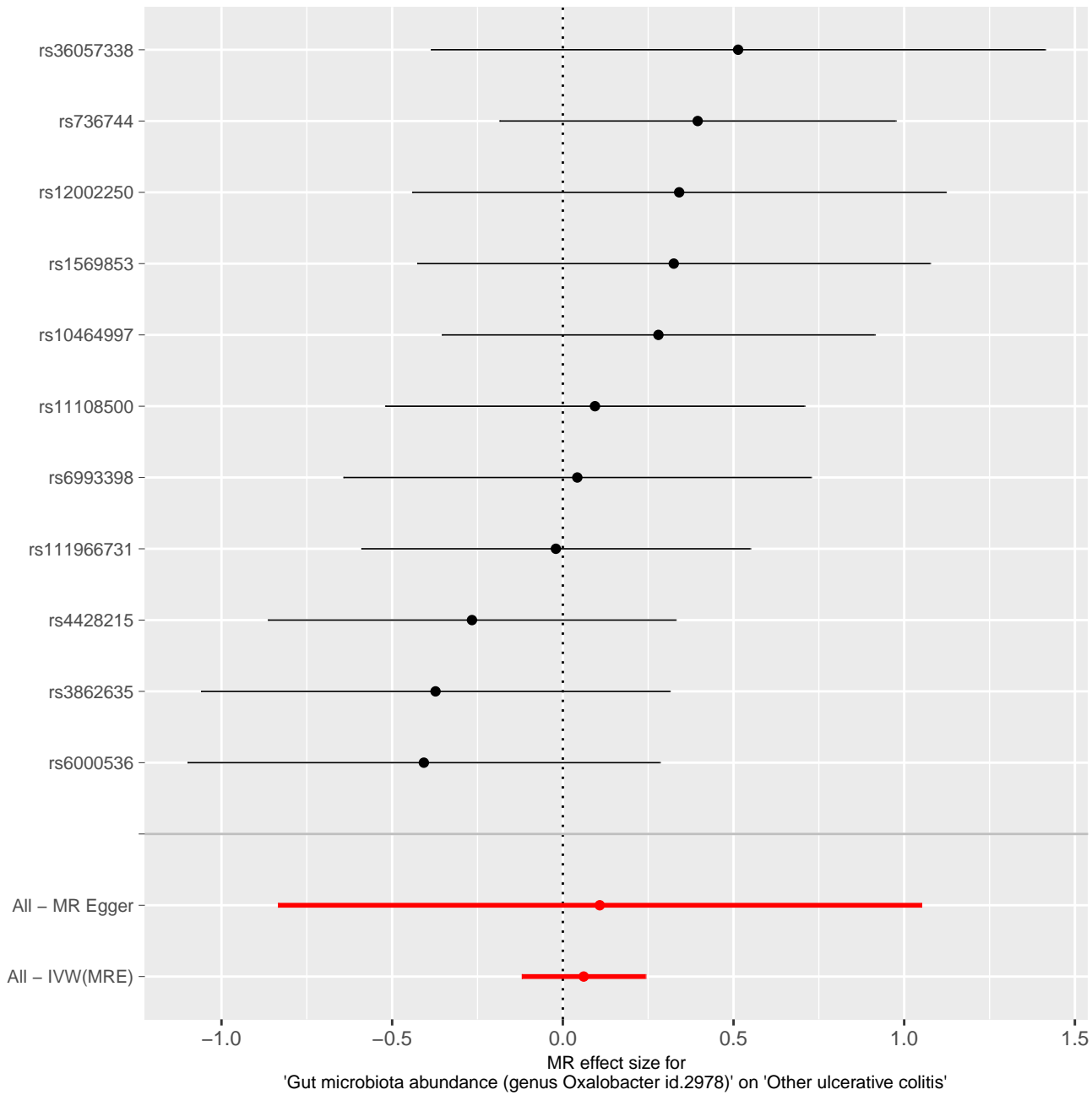


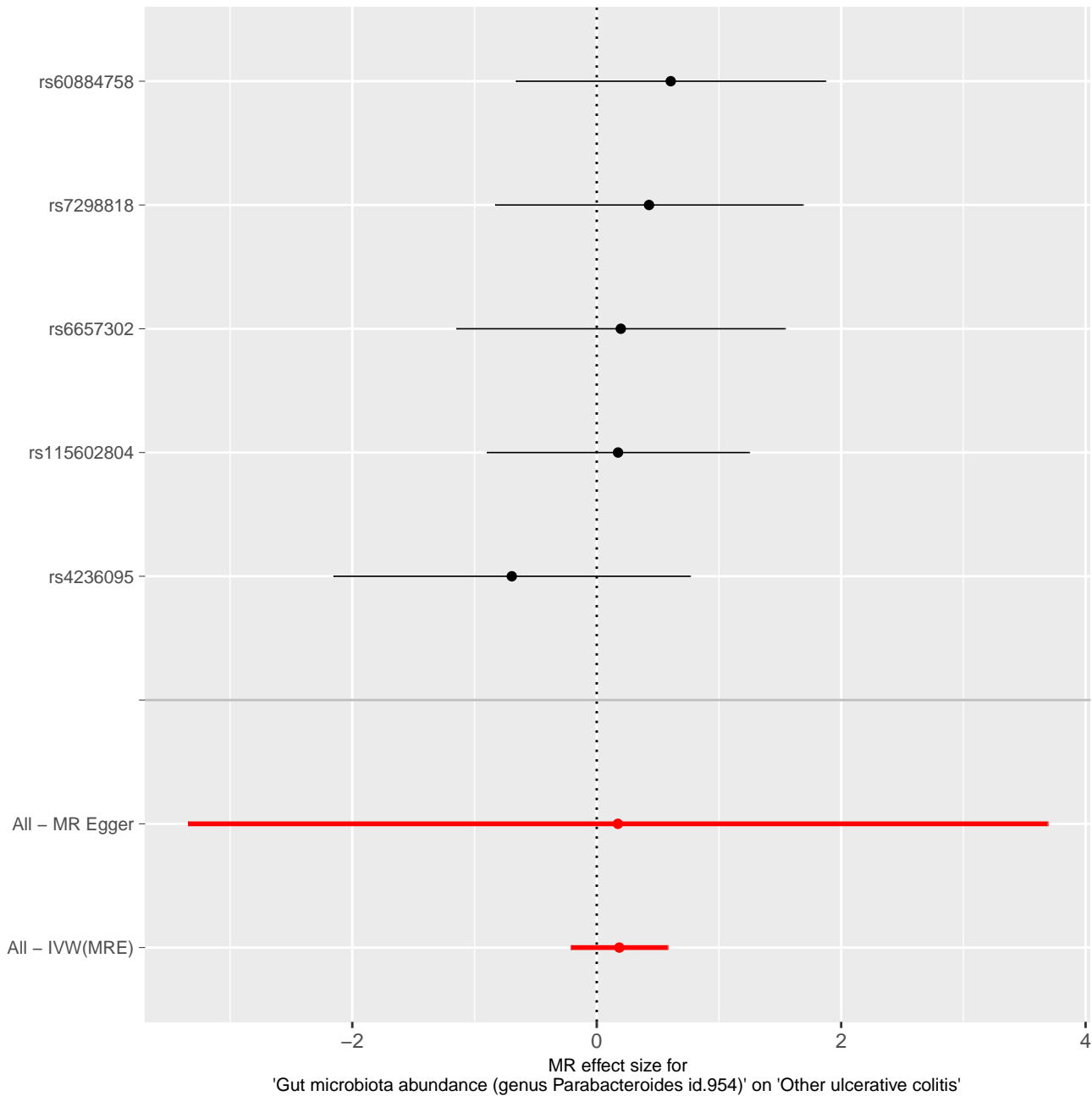


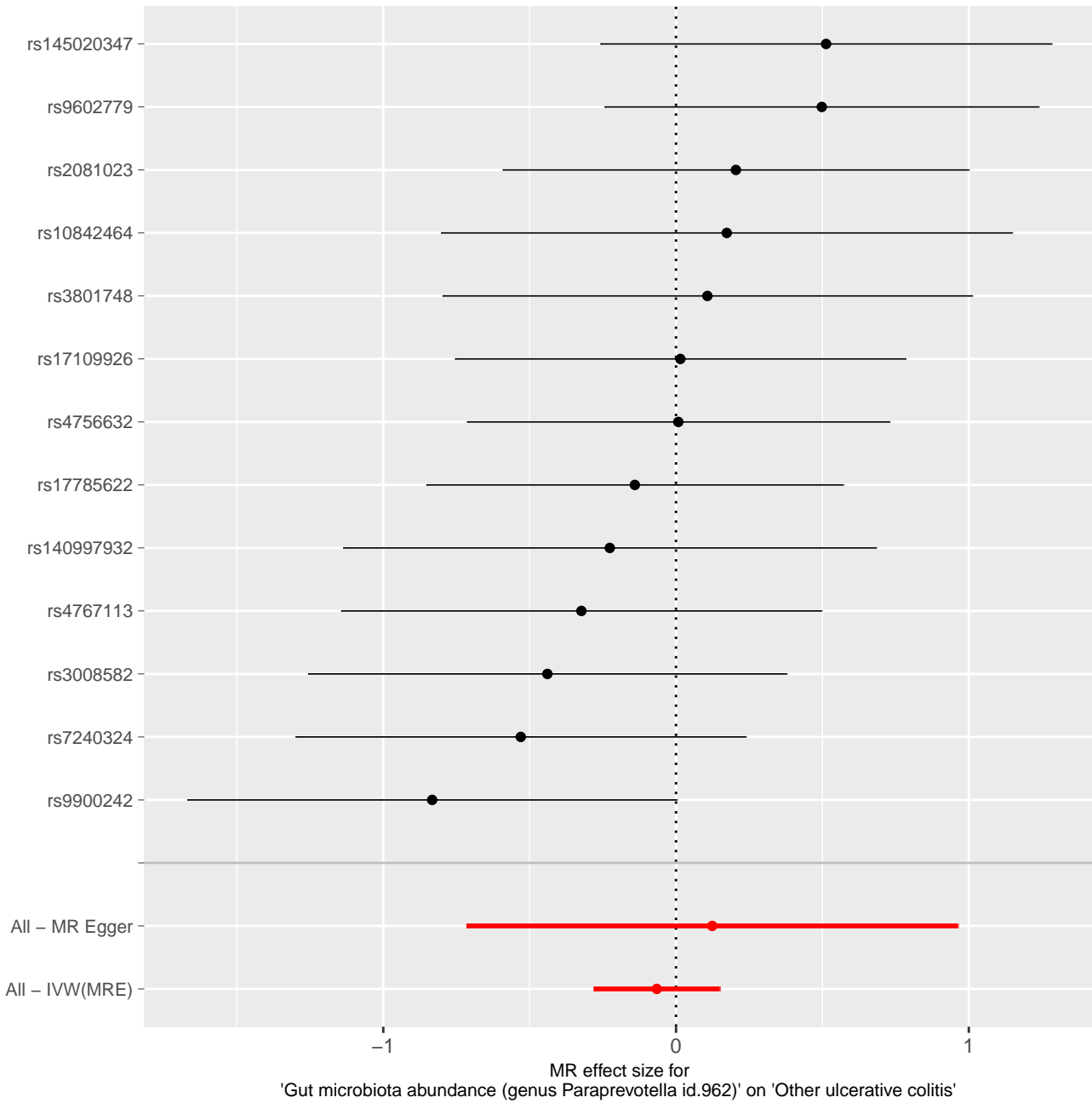


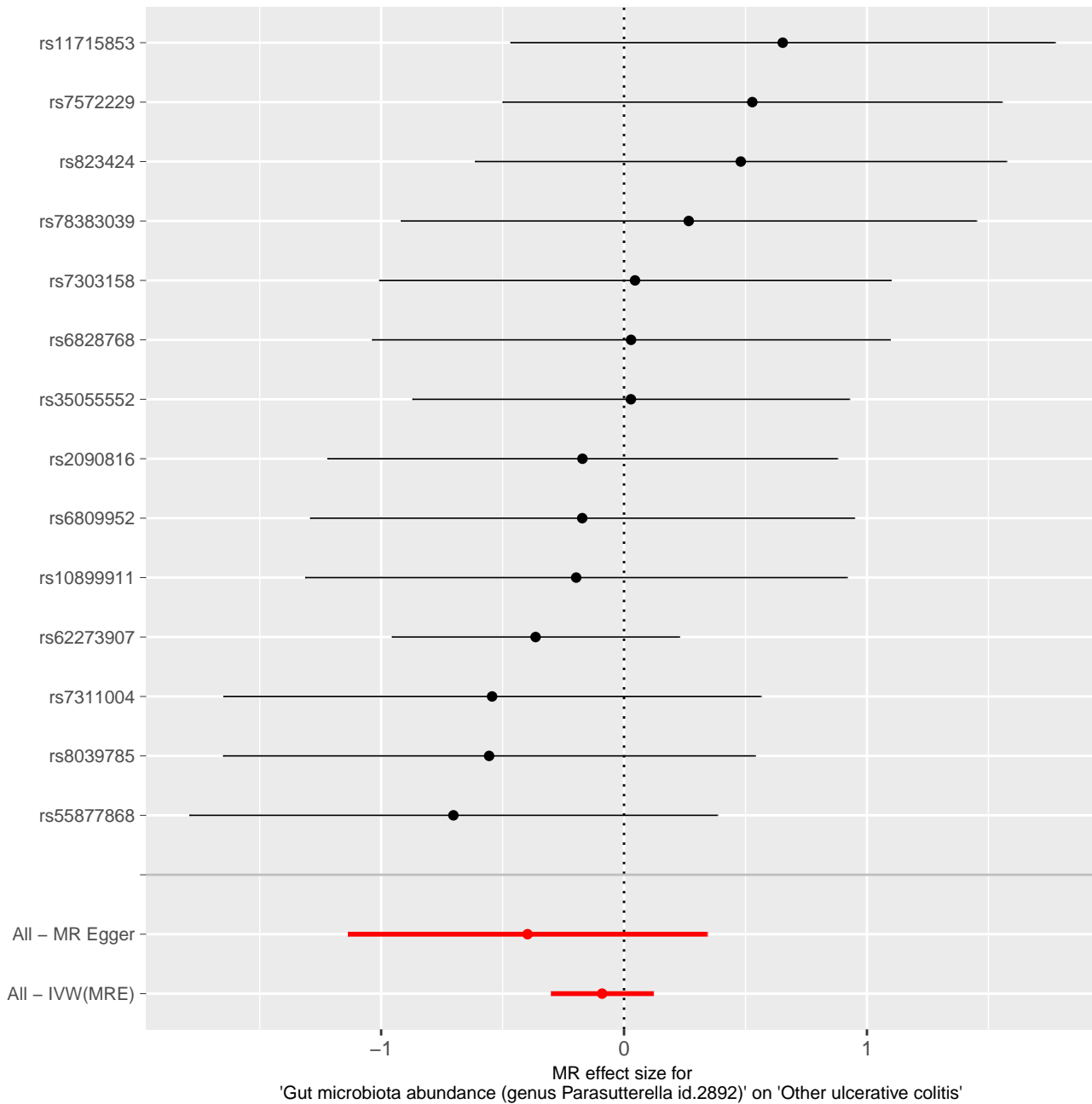


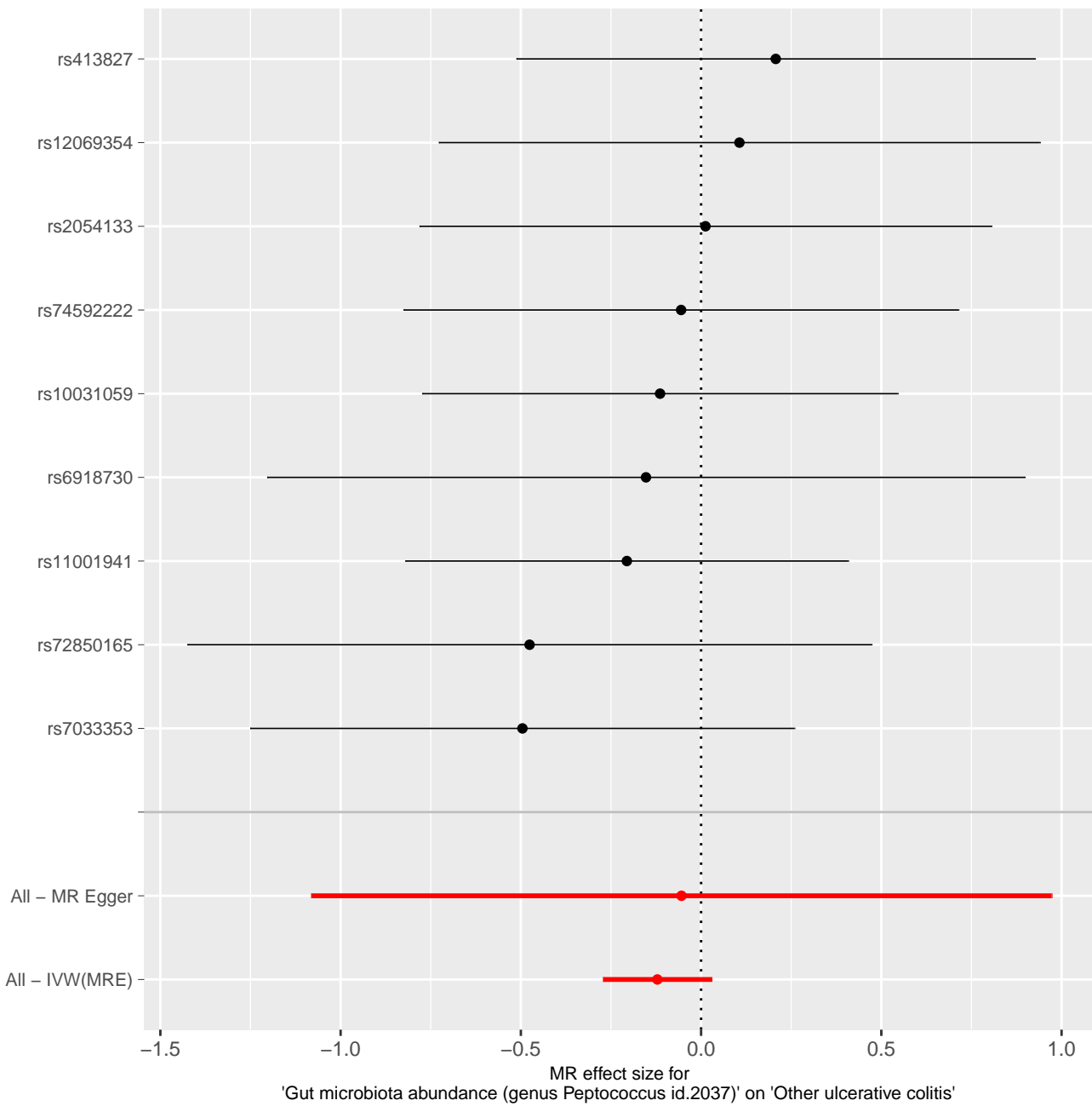
Batch 1312 : Gut microbiota abundance (genus Oxalobacter id.2978) on Other ulcerative colitis

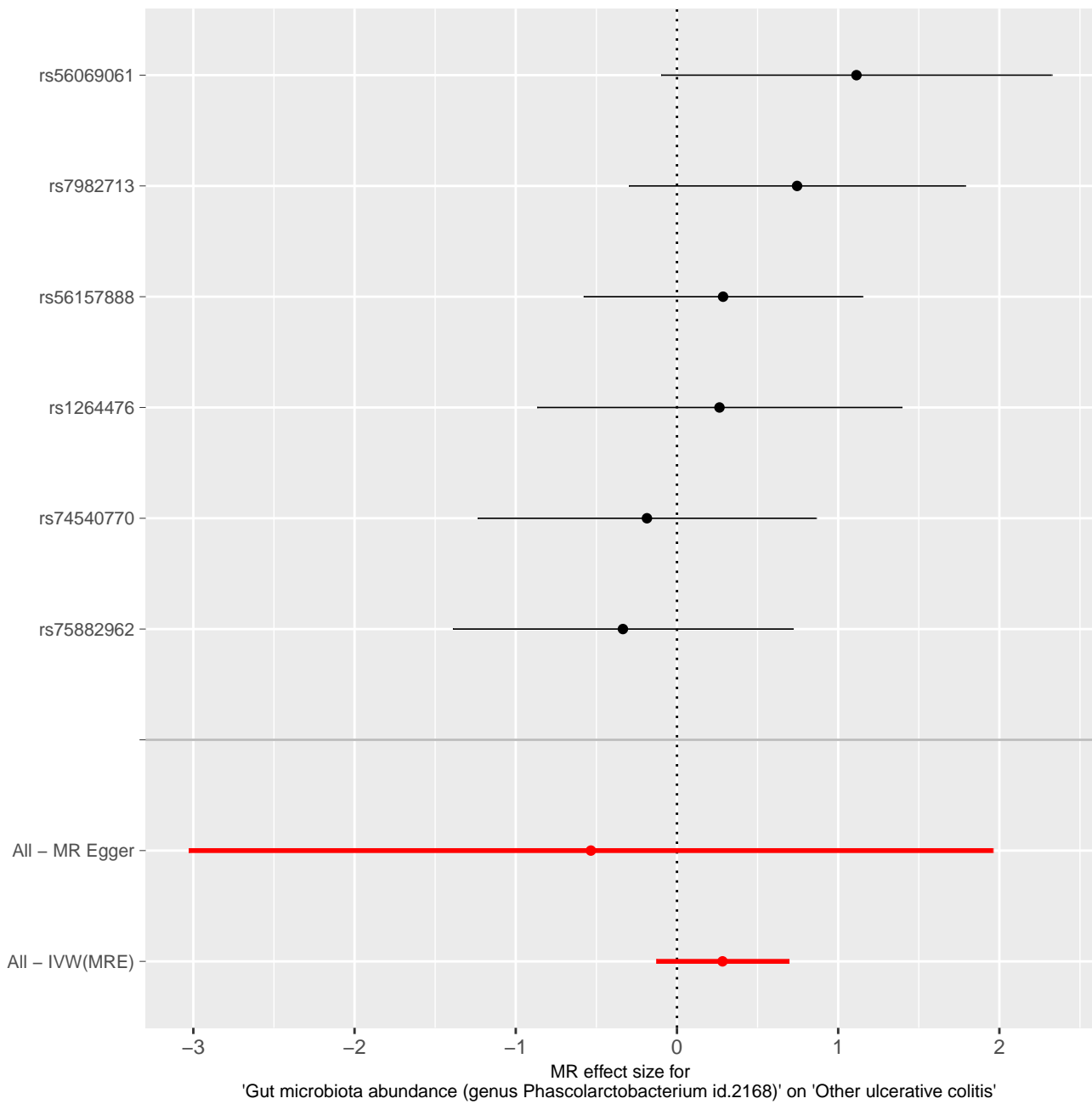


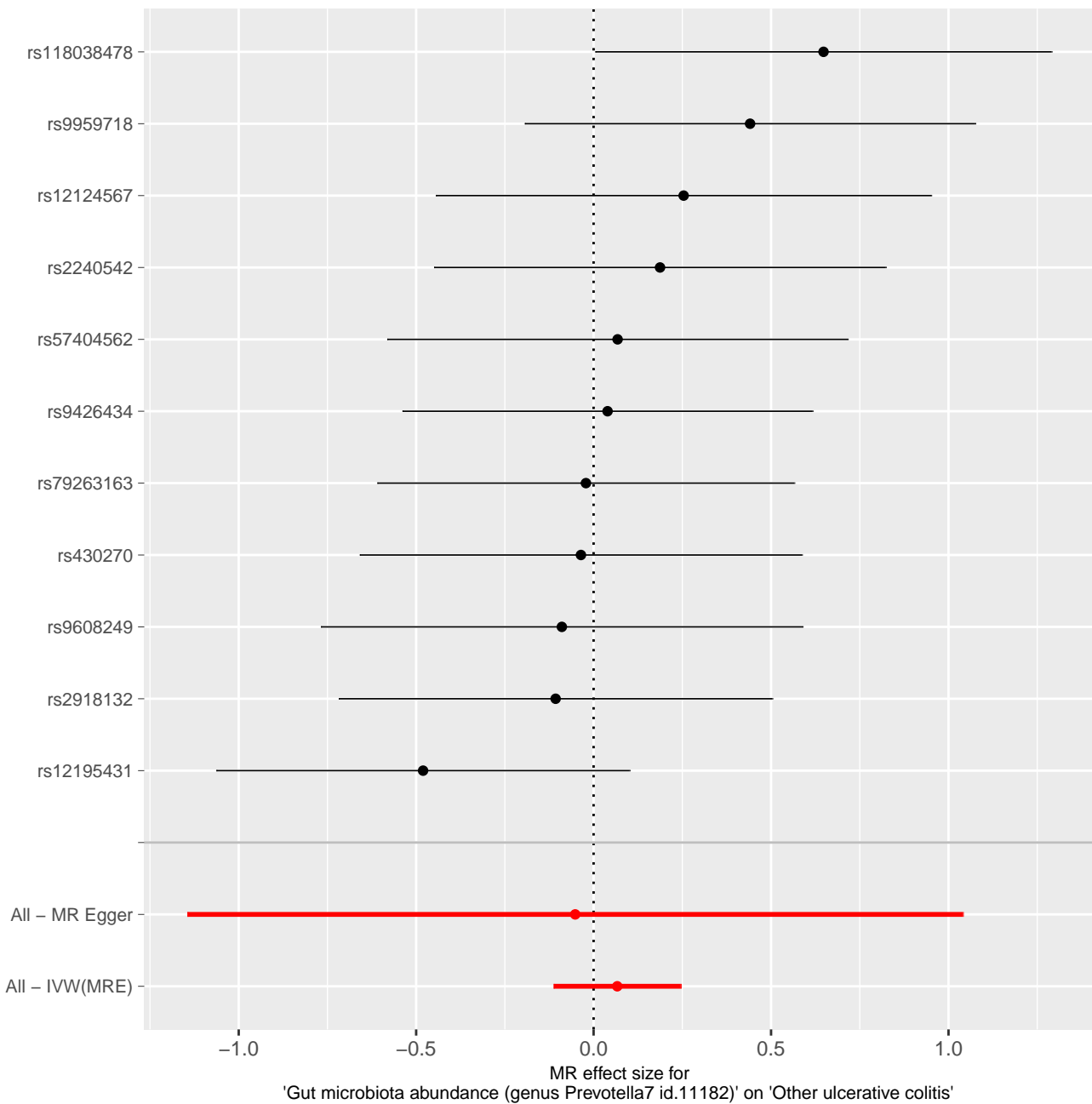


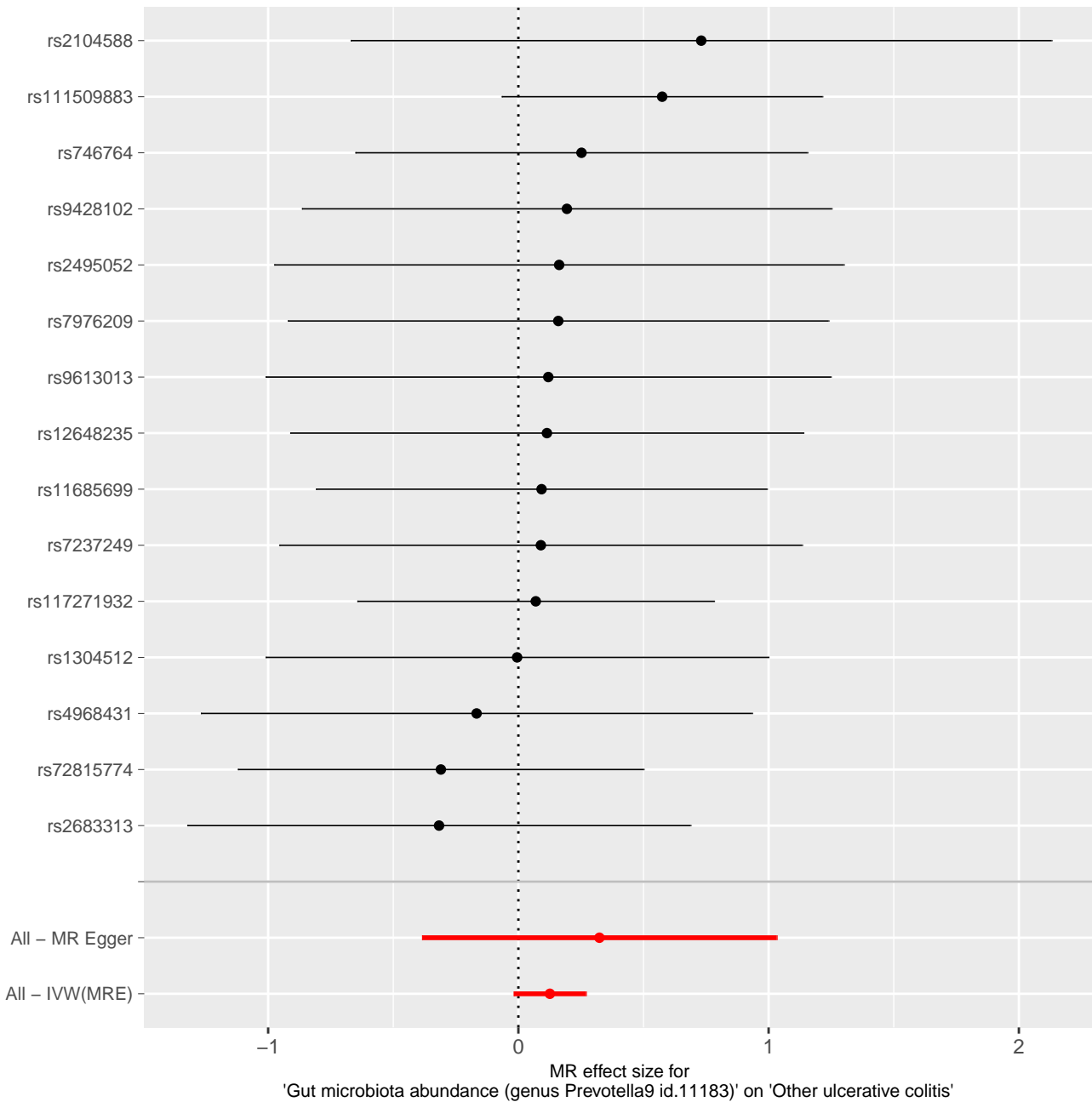


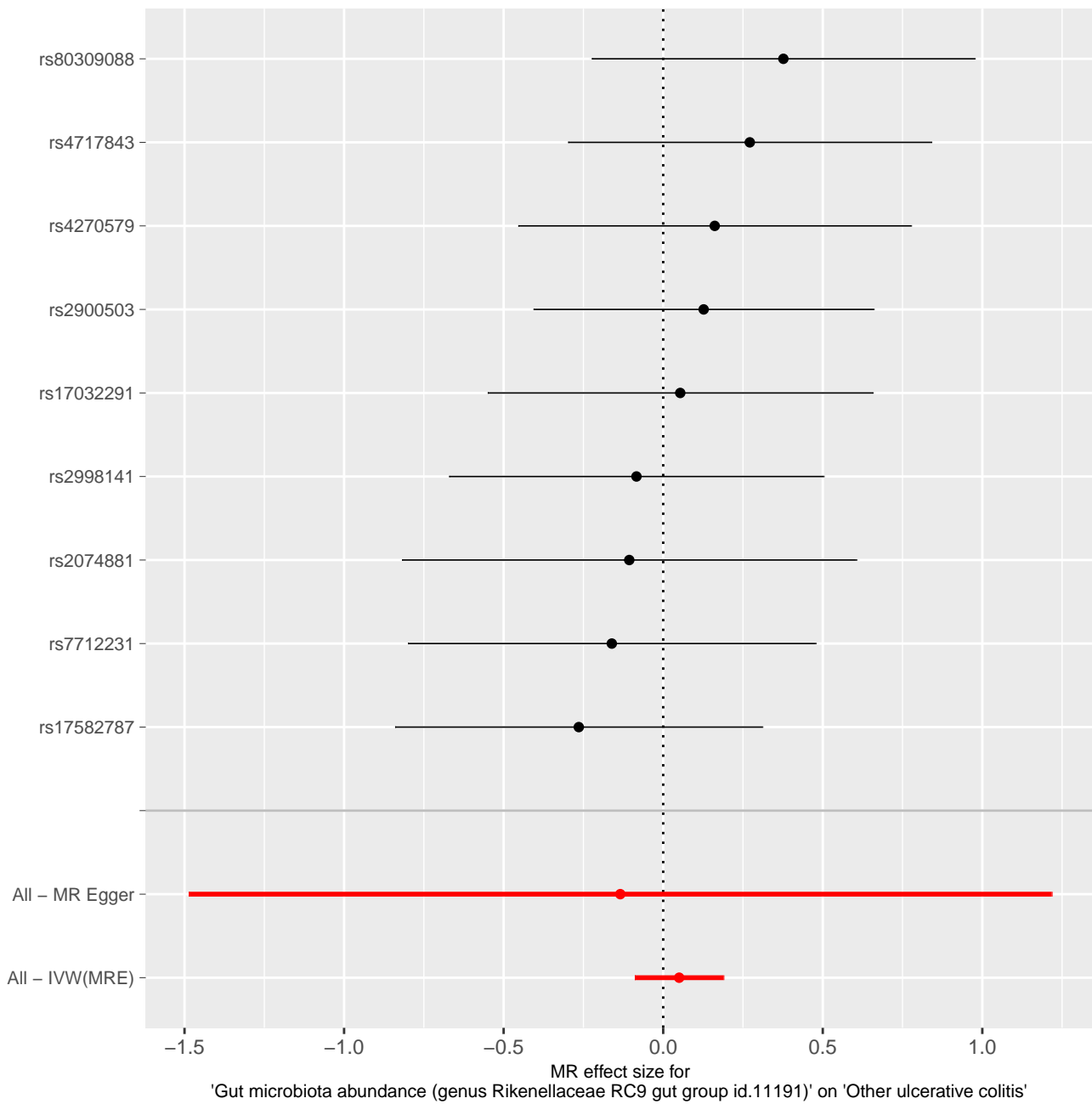


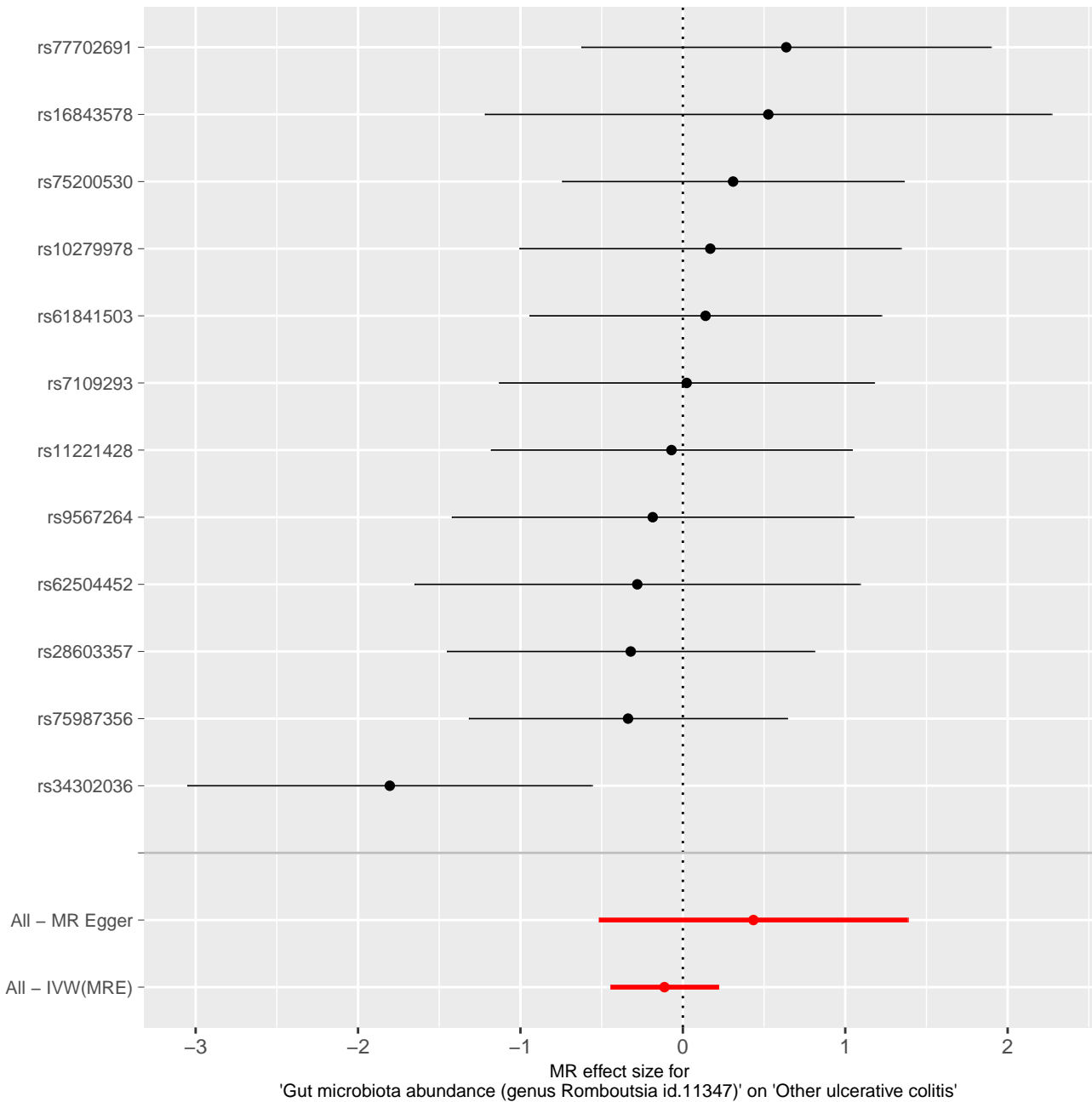


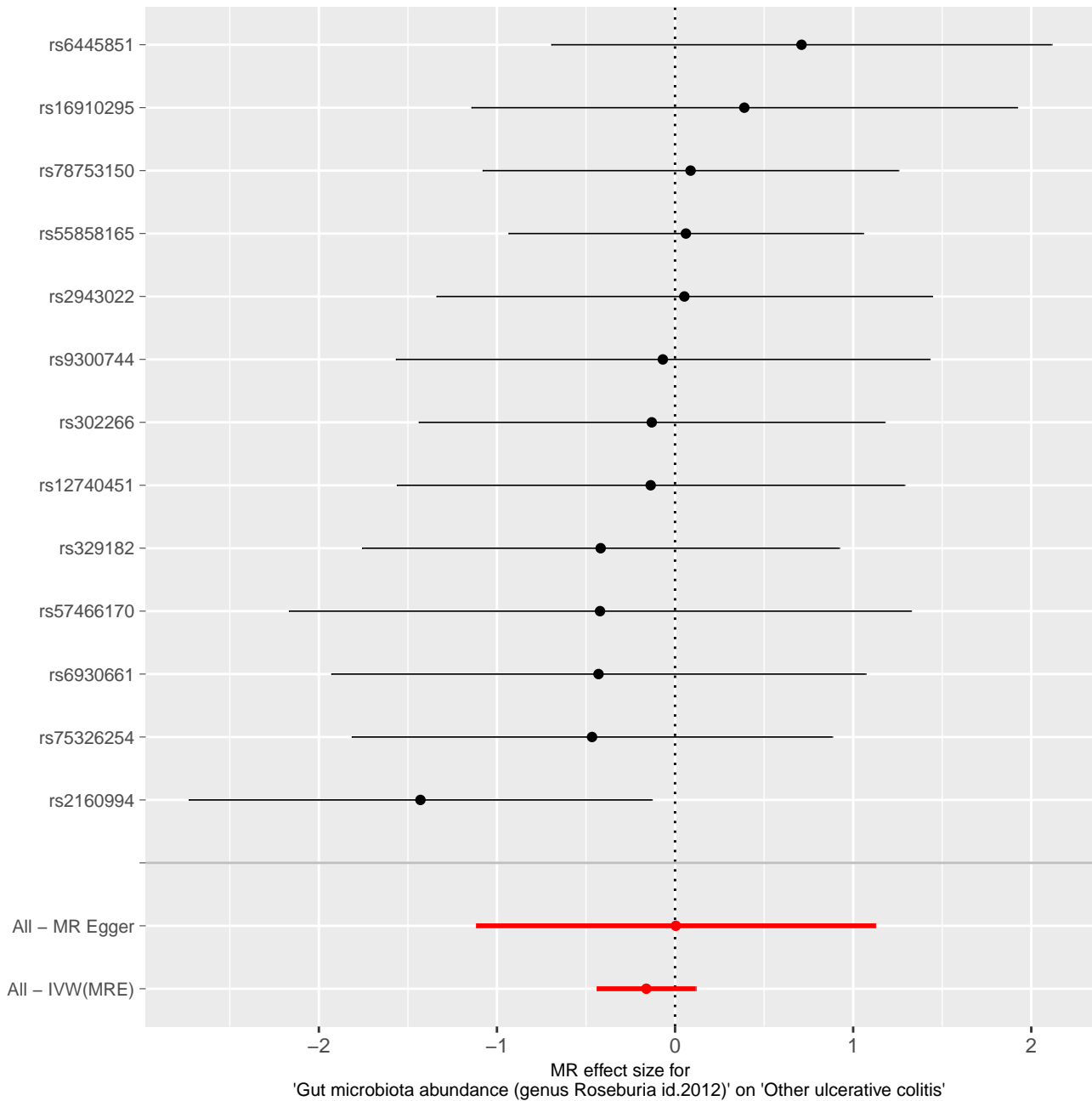


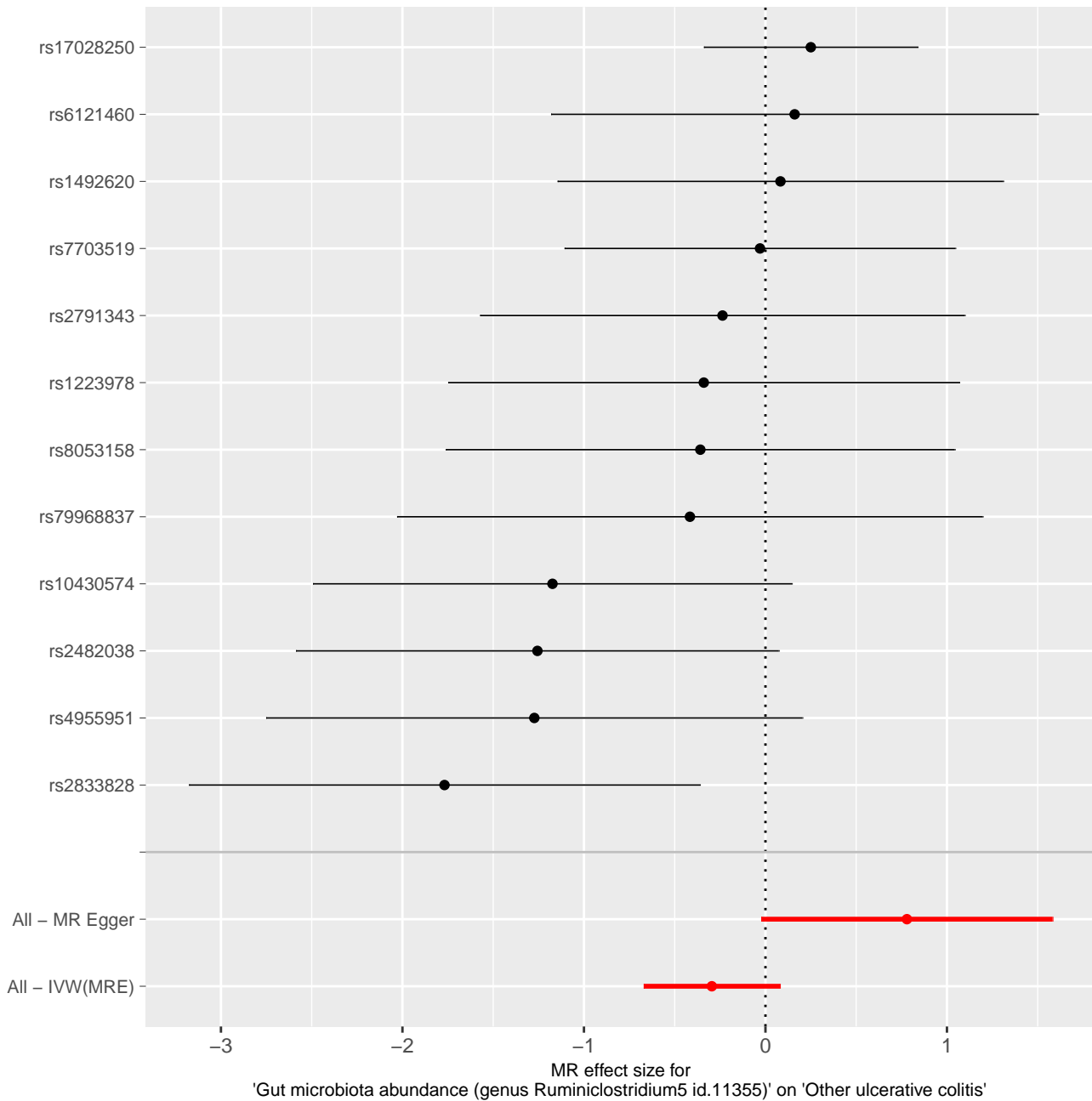


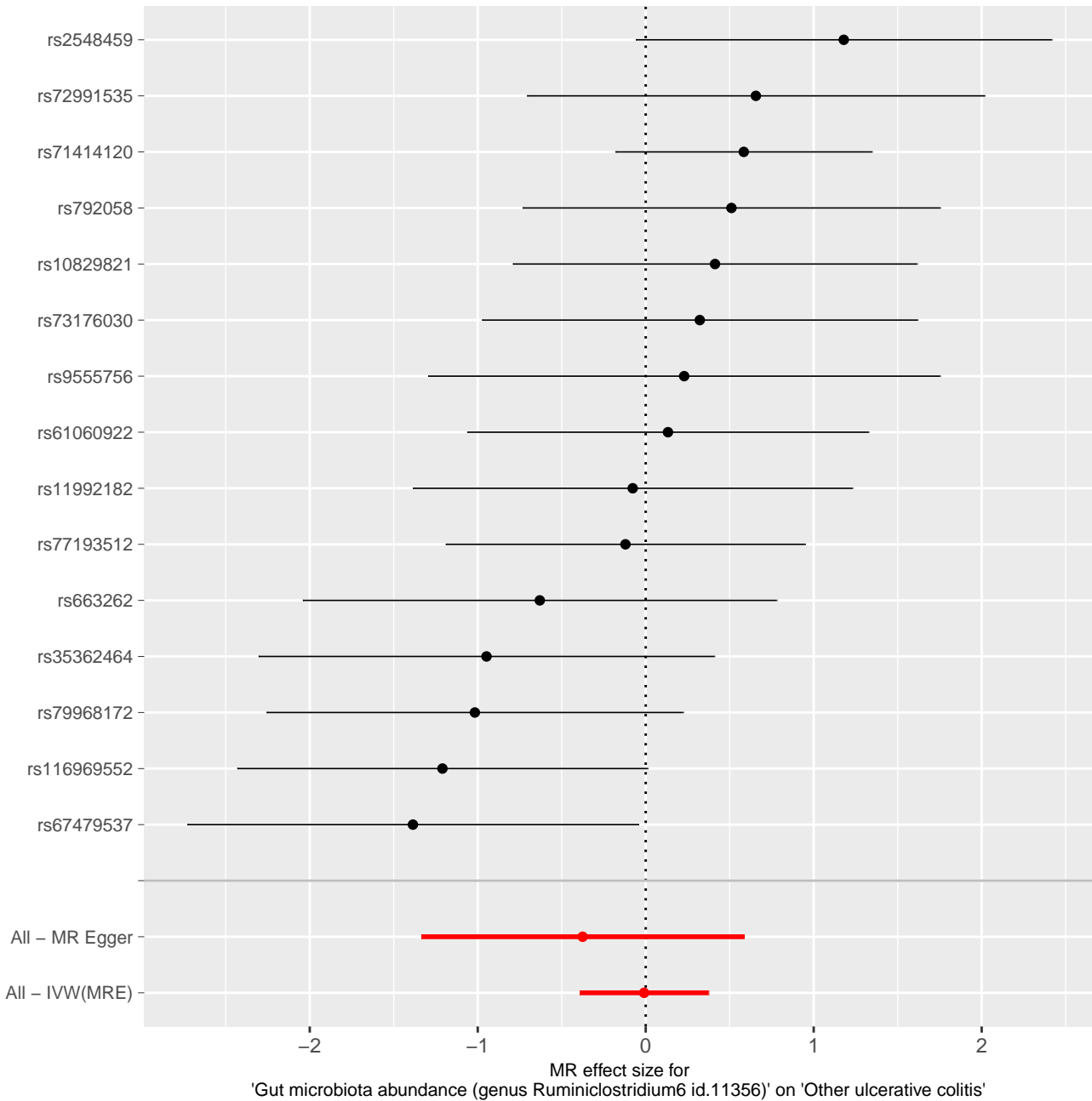


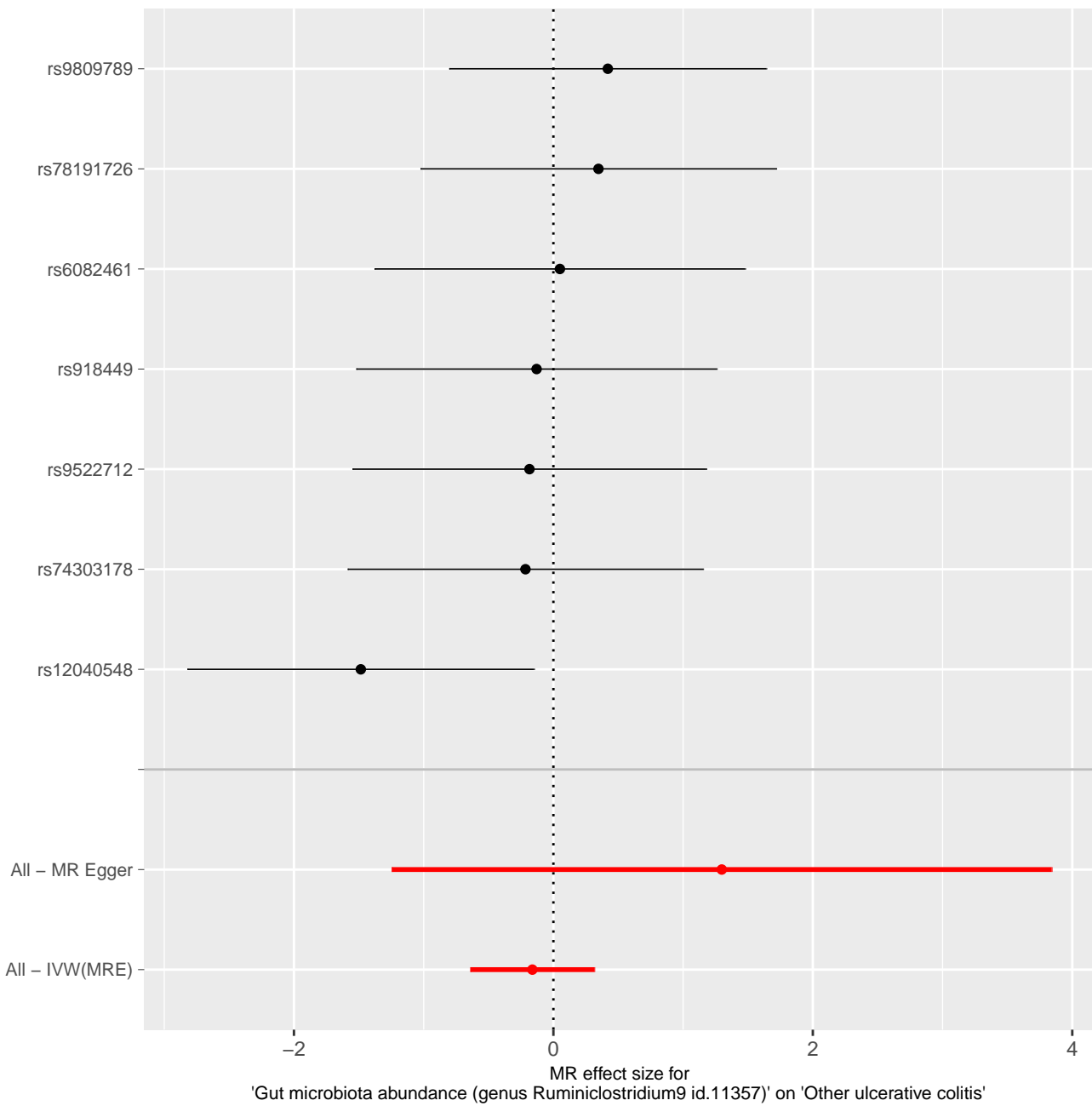


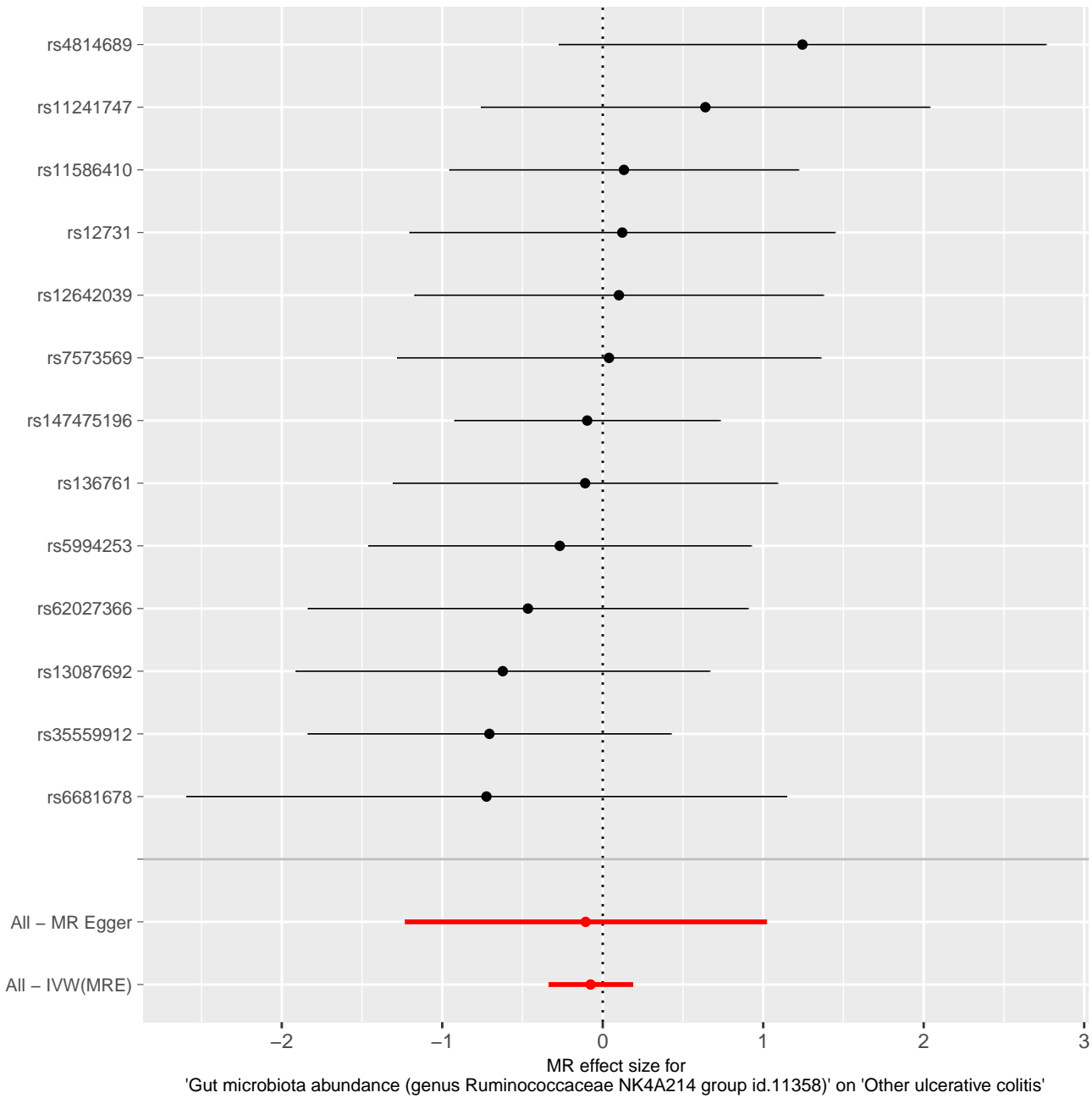


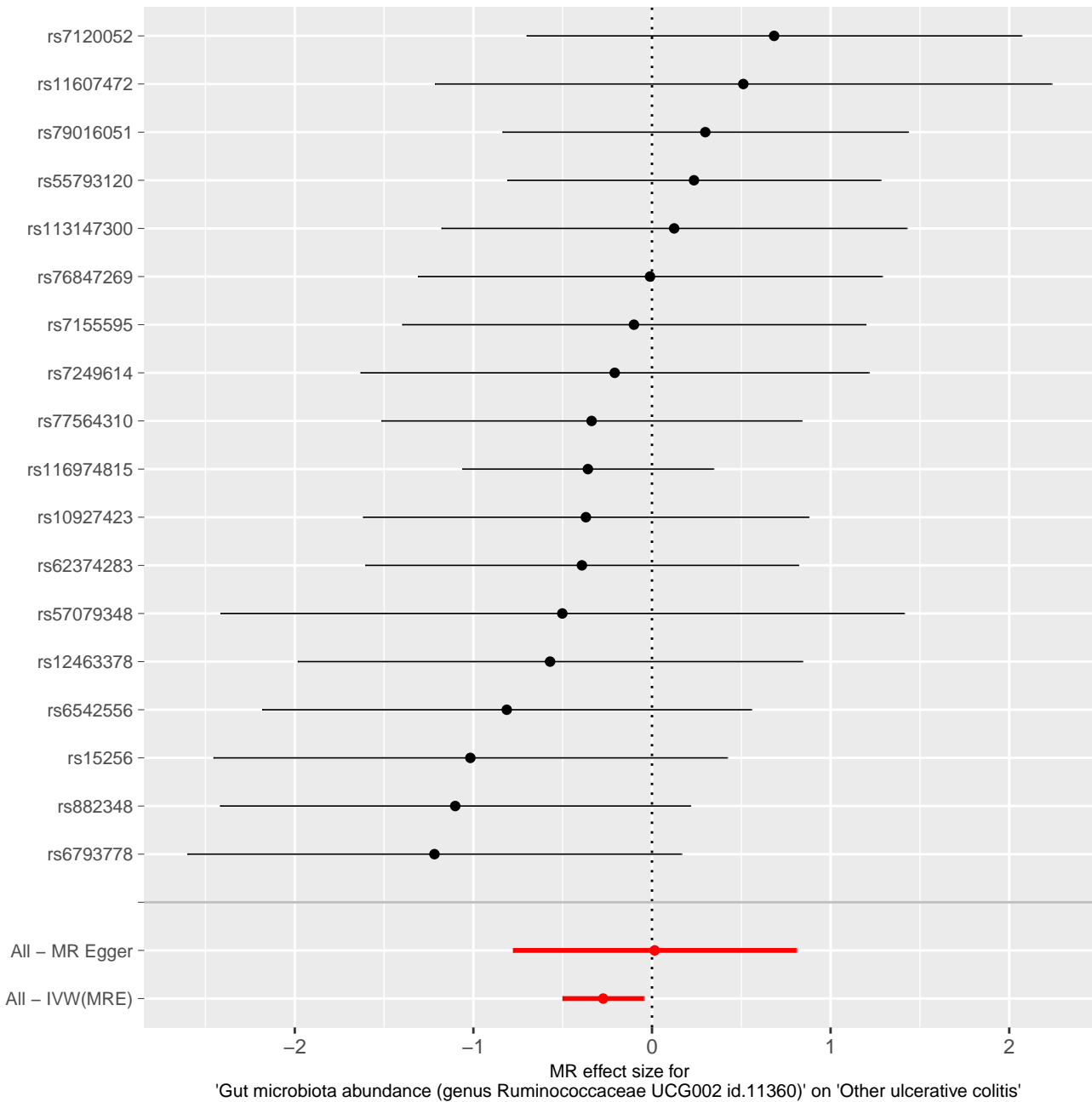


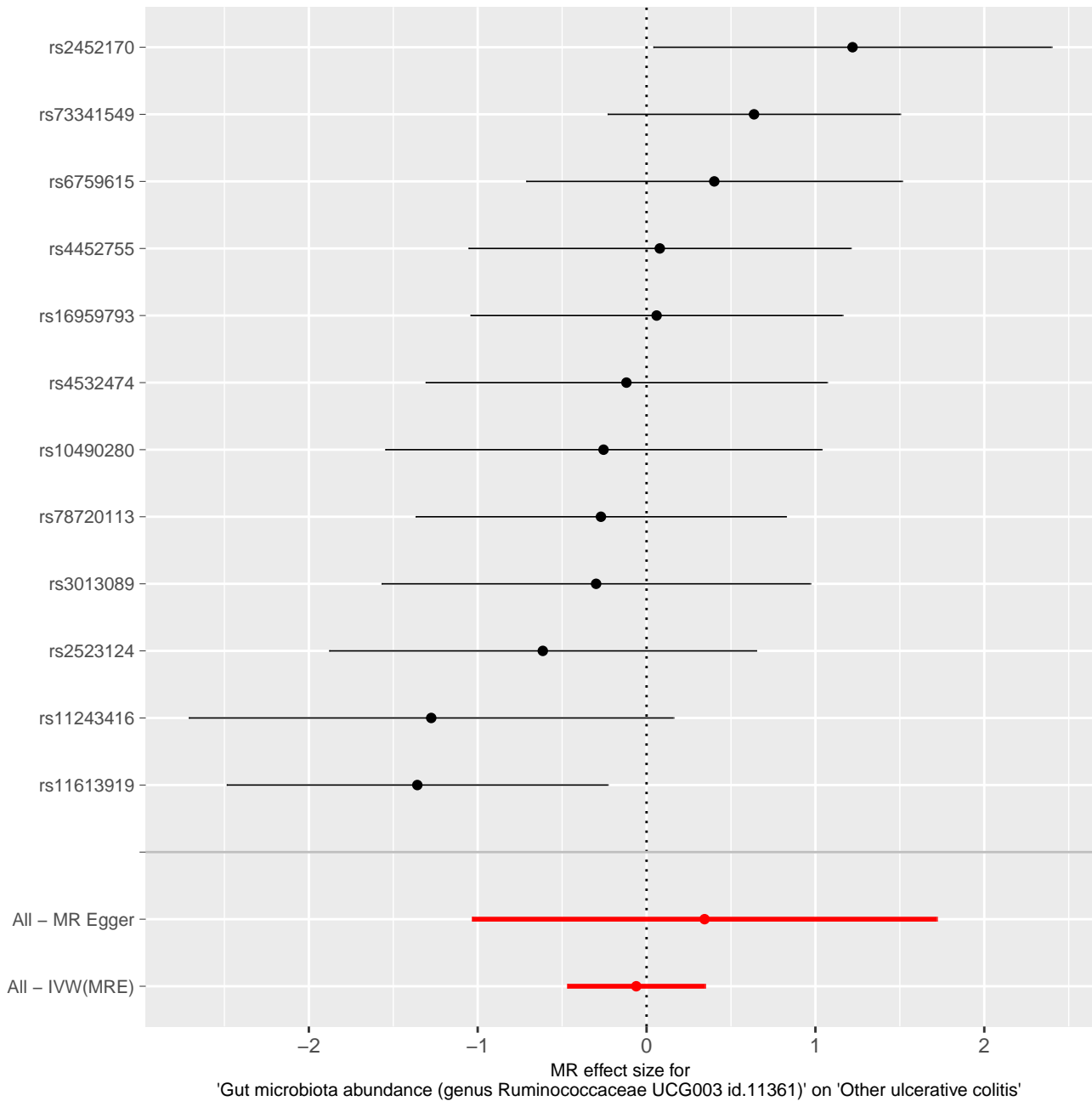


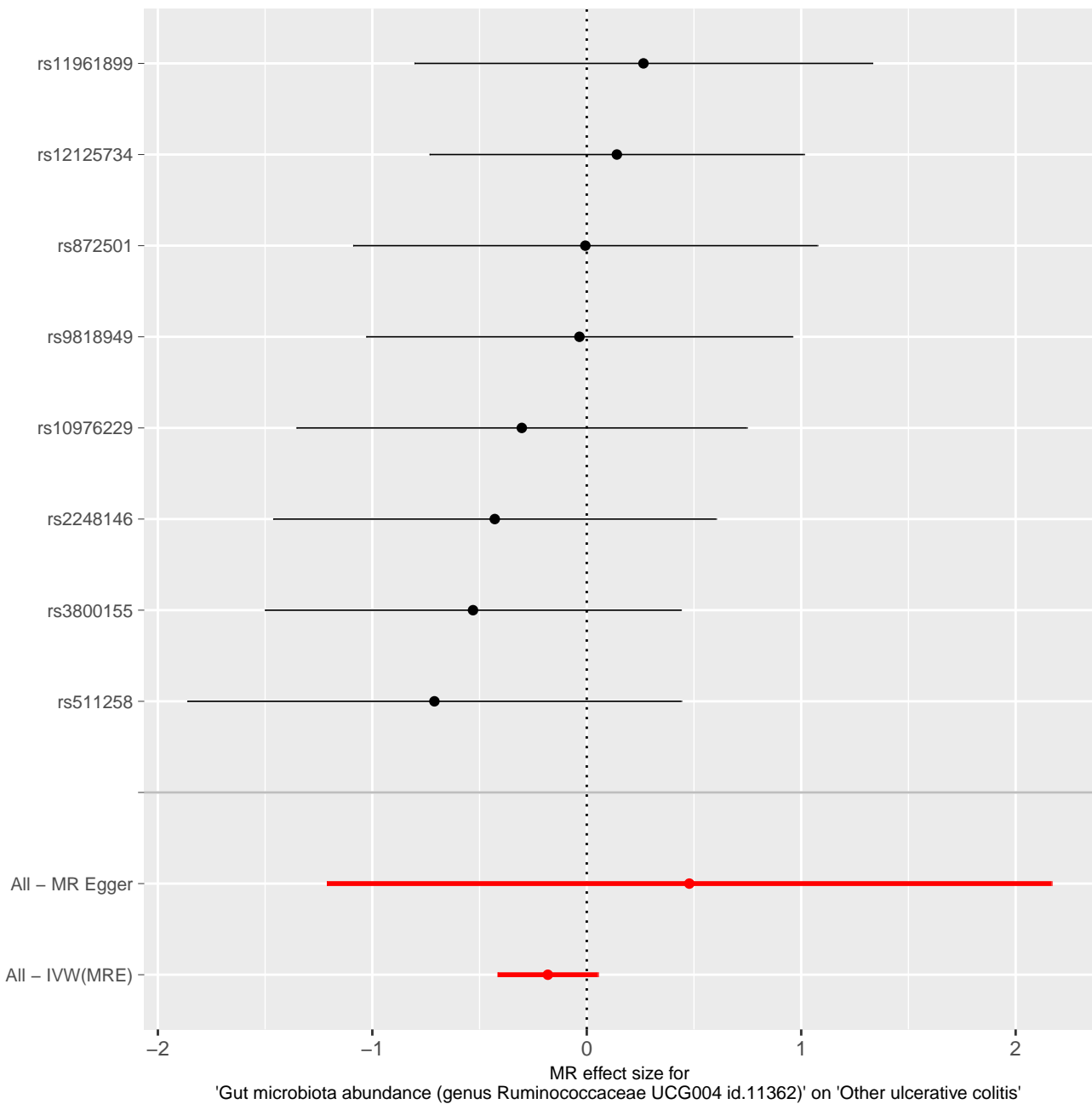


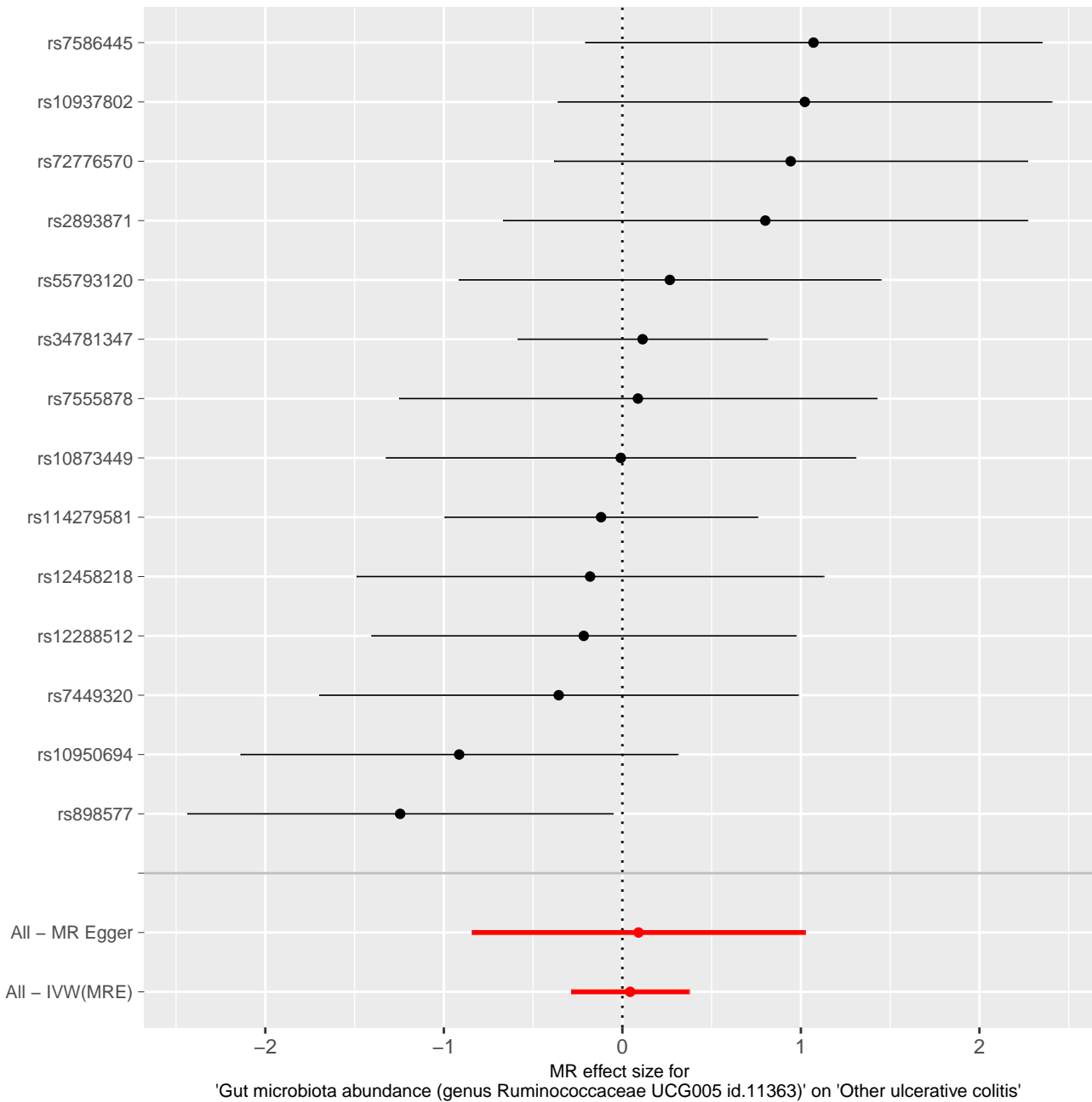


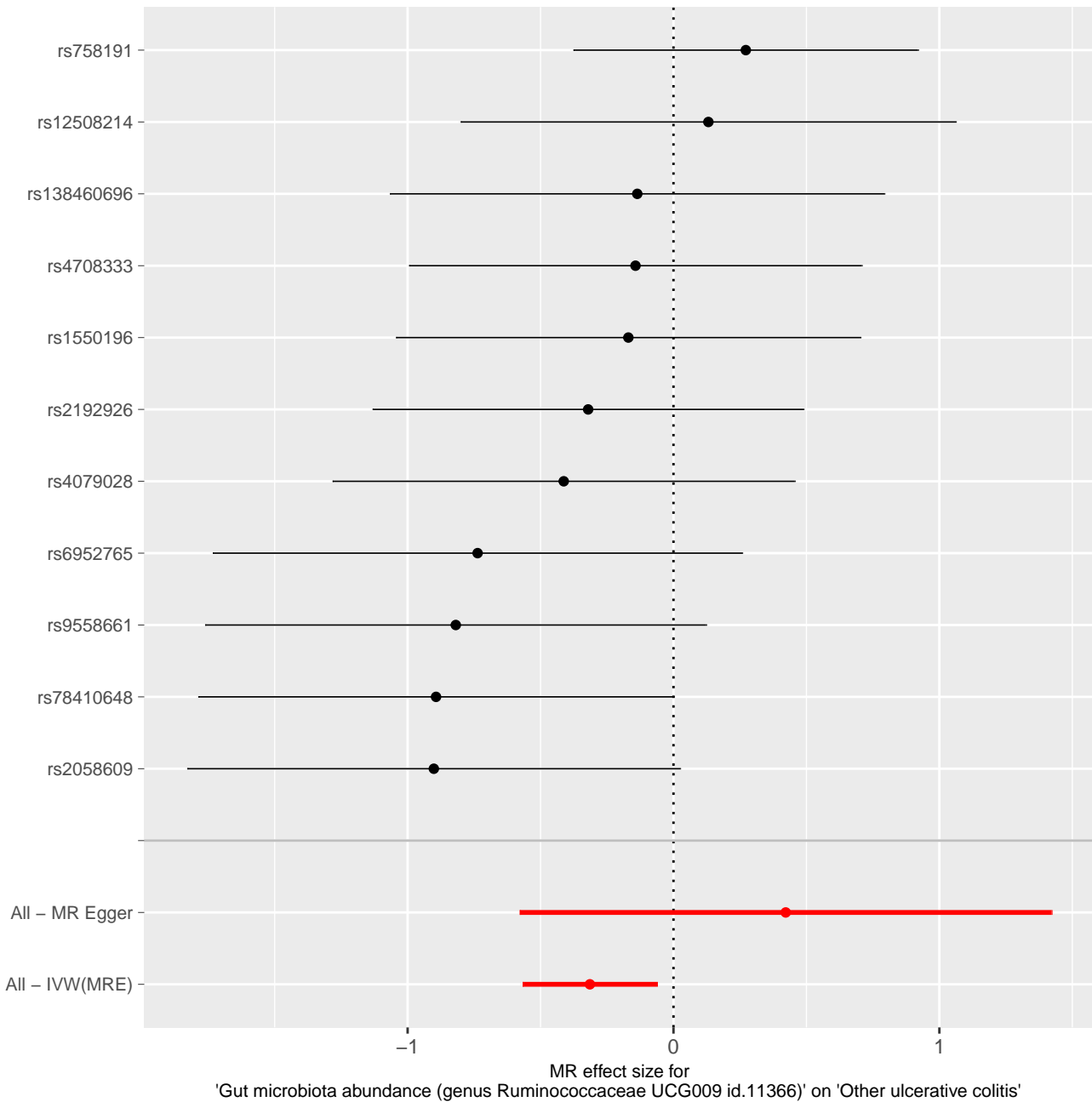


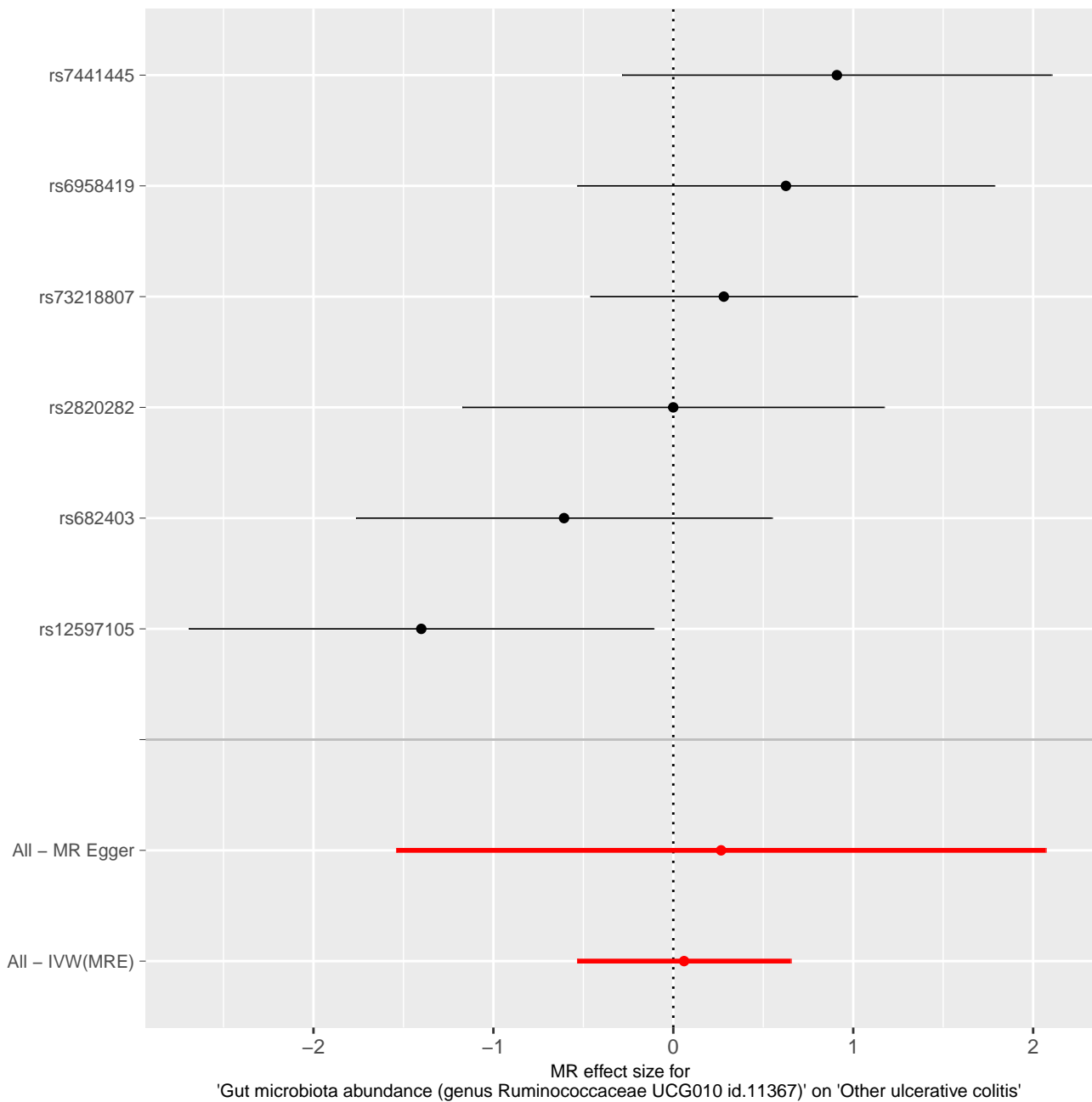


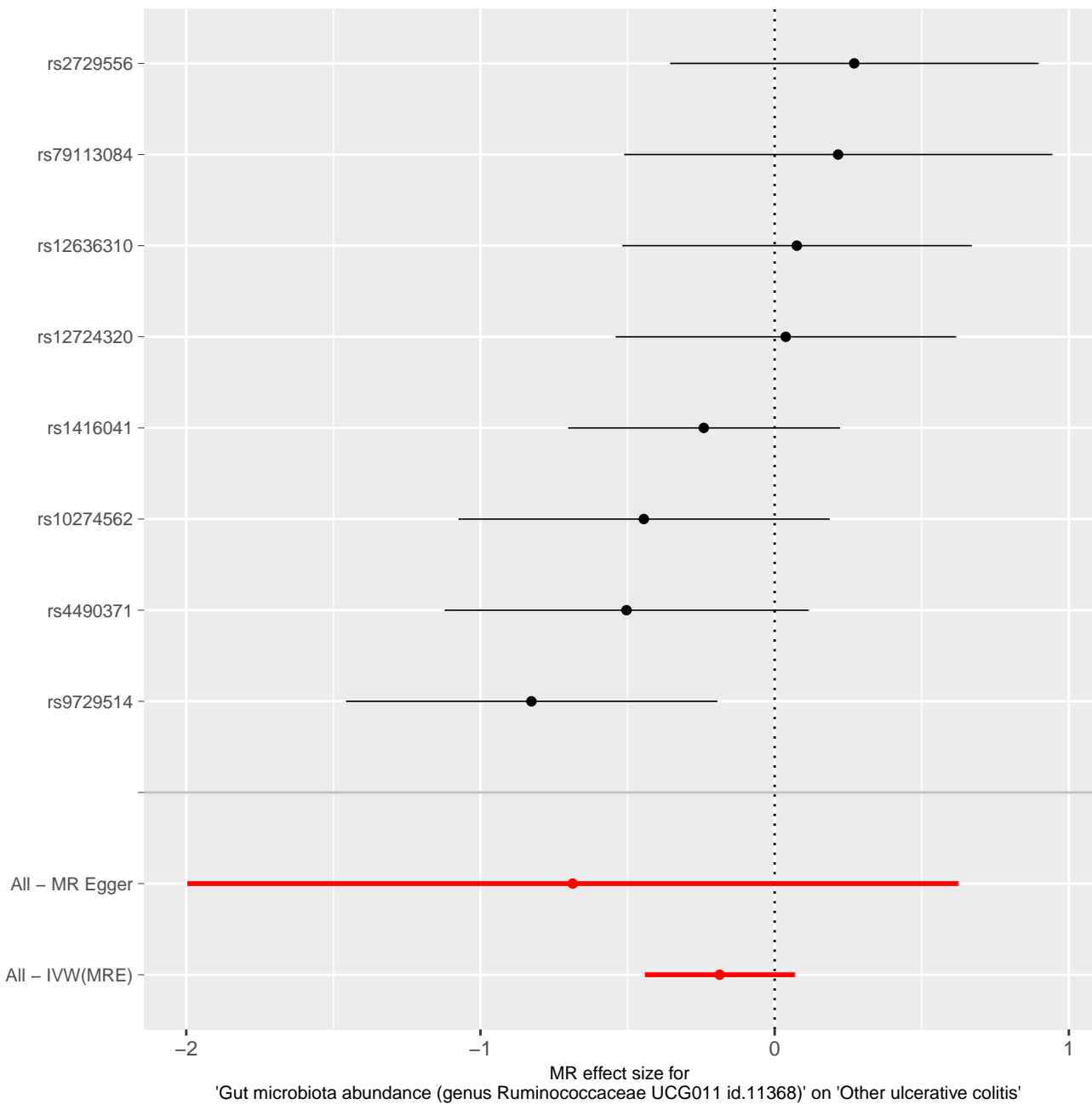


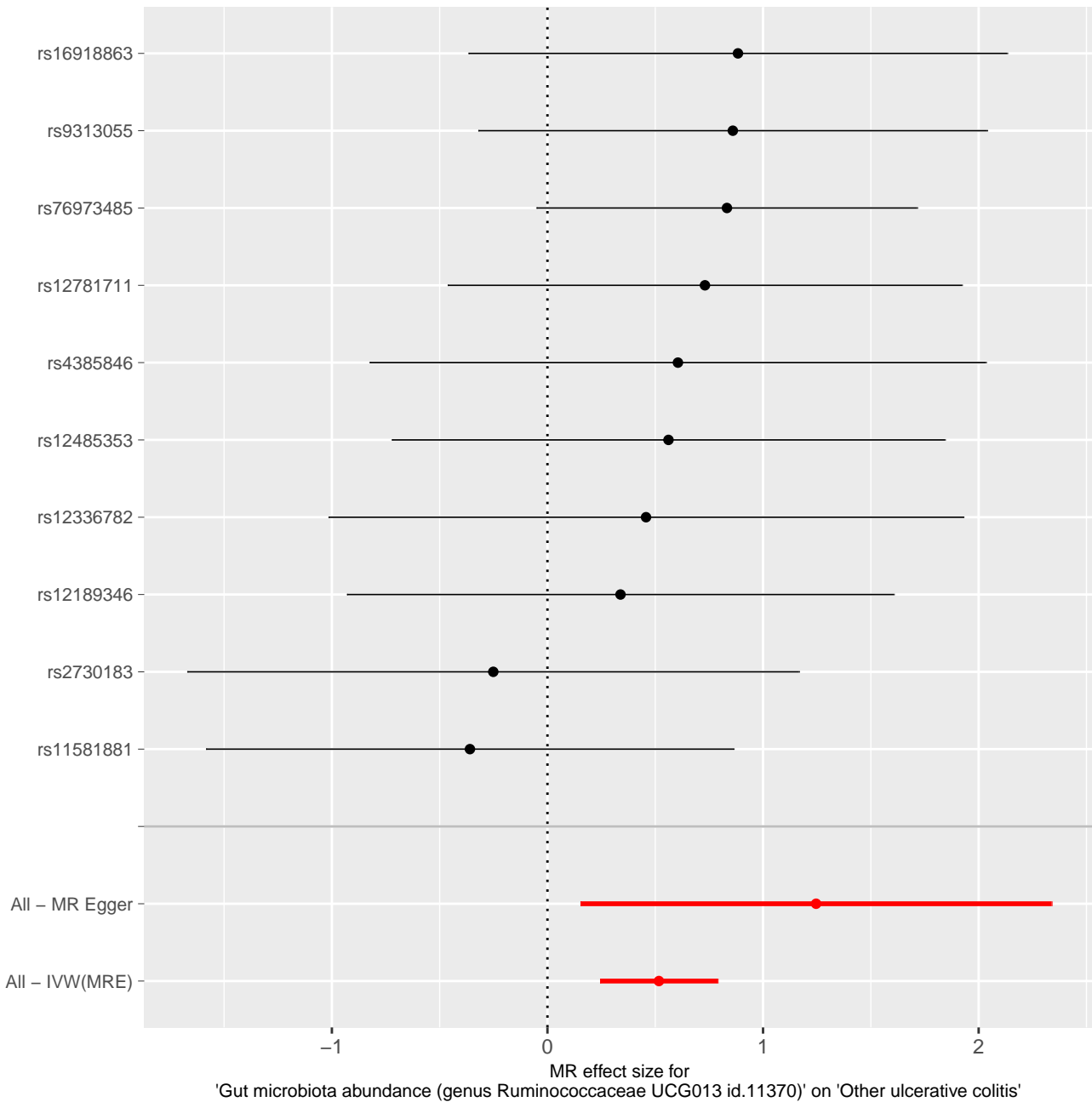


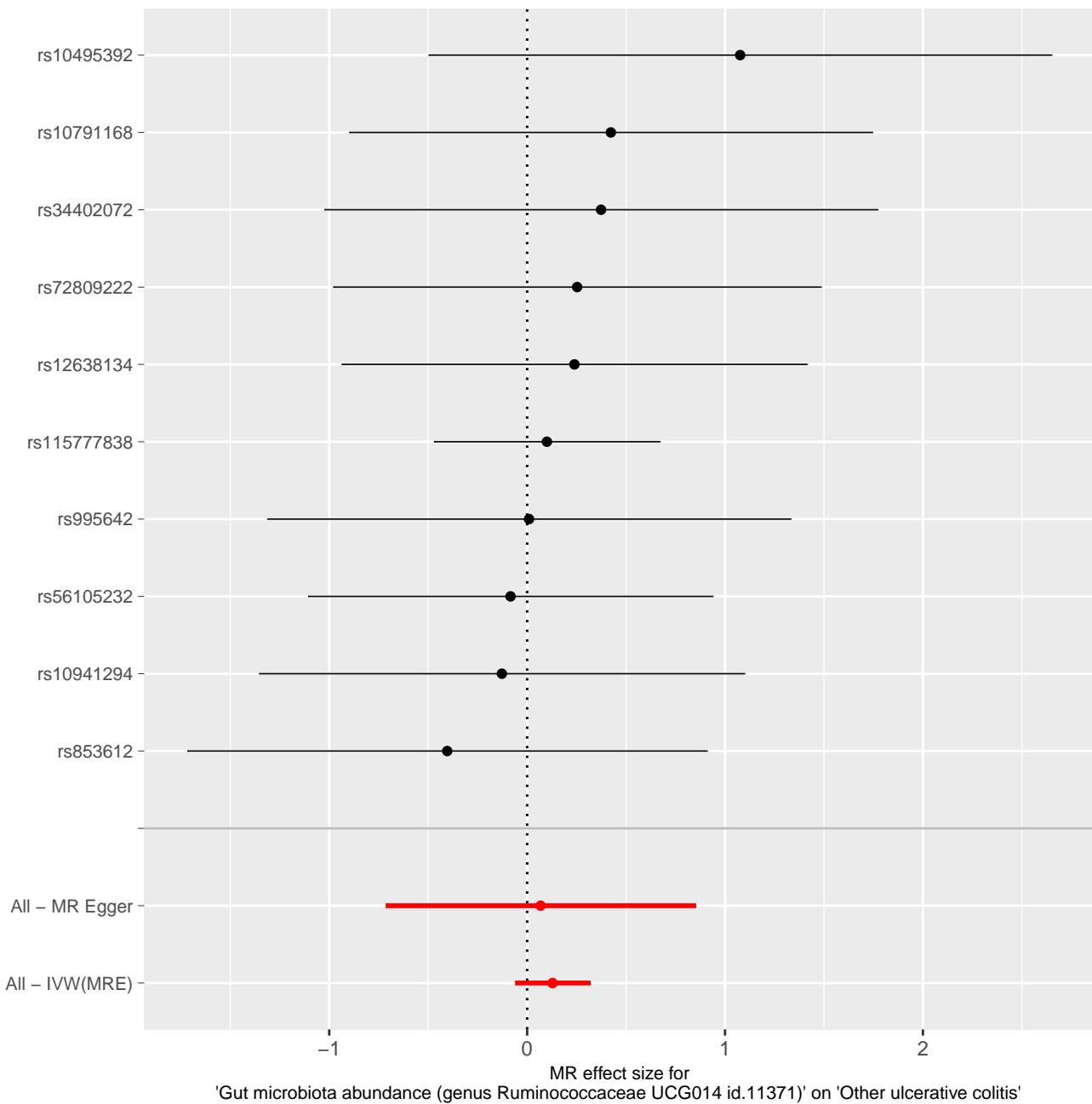


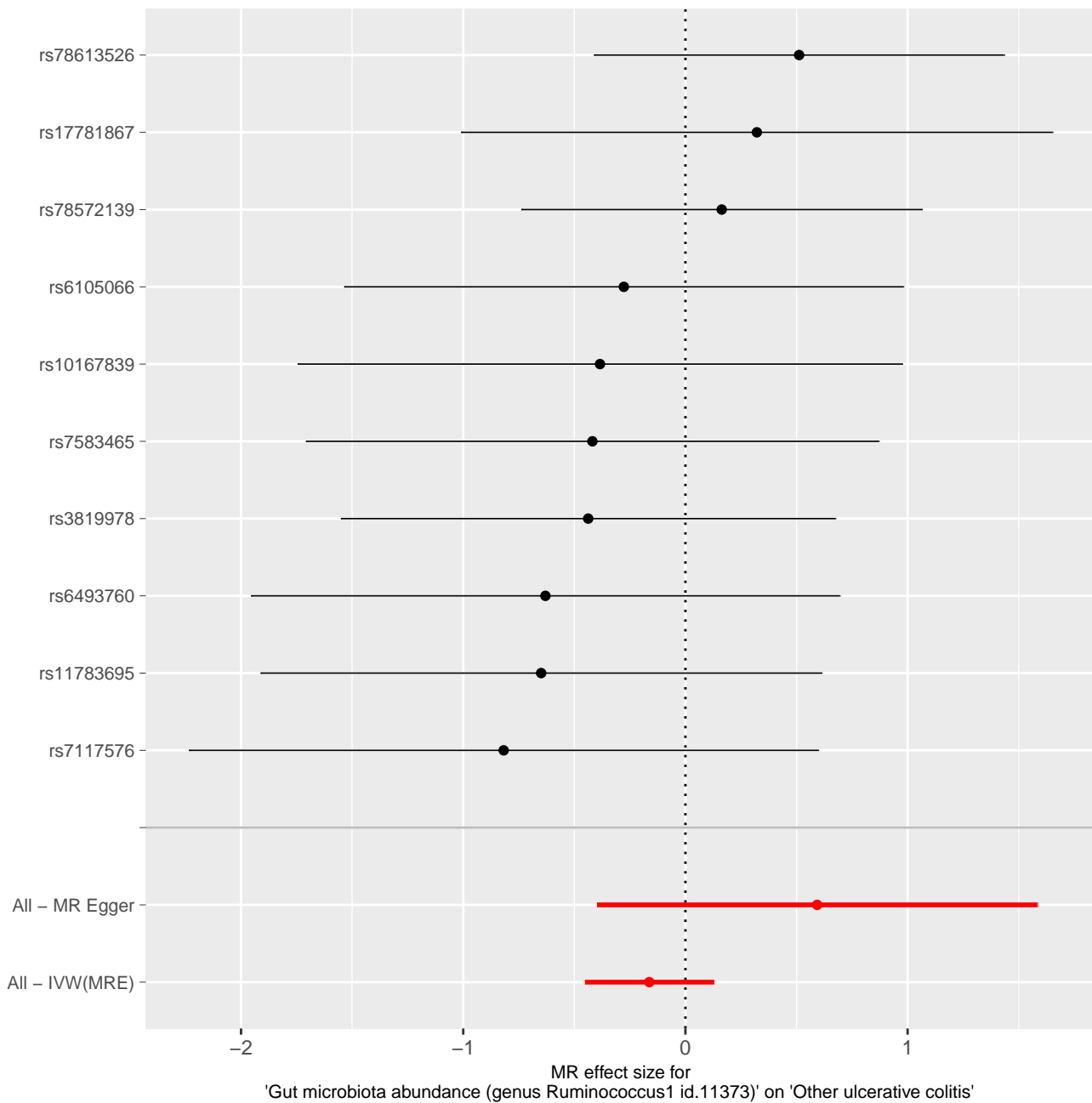


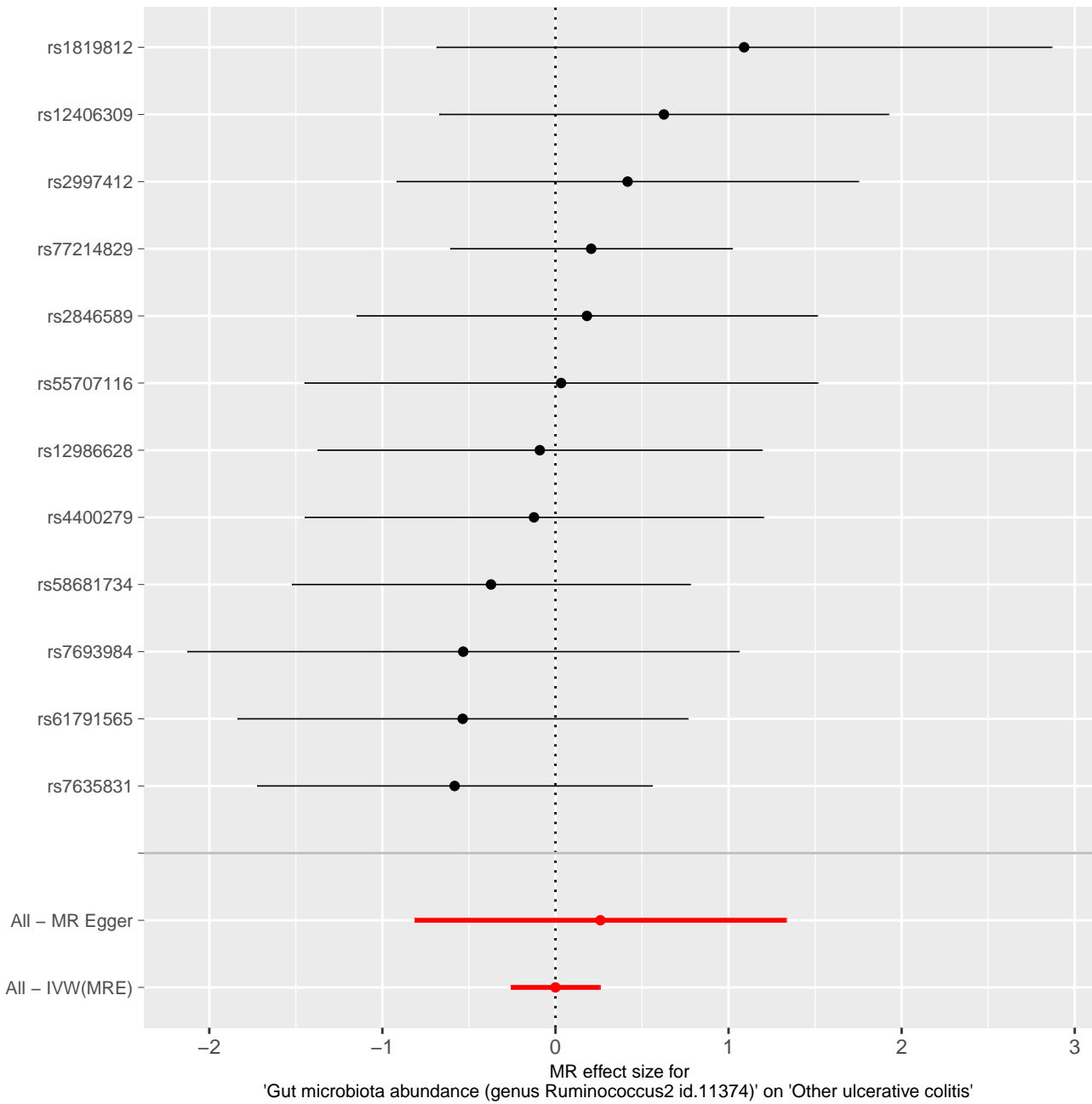


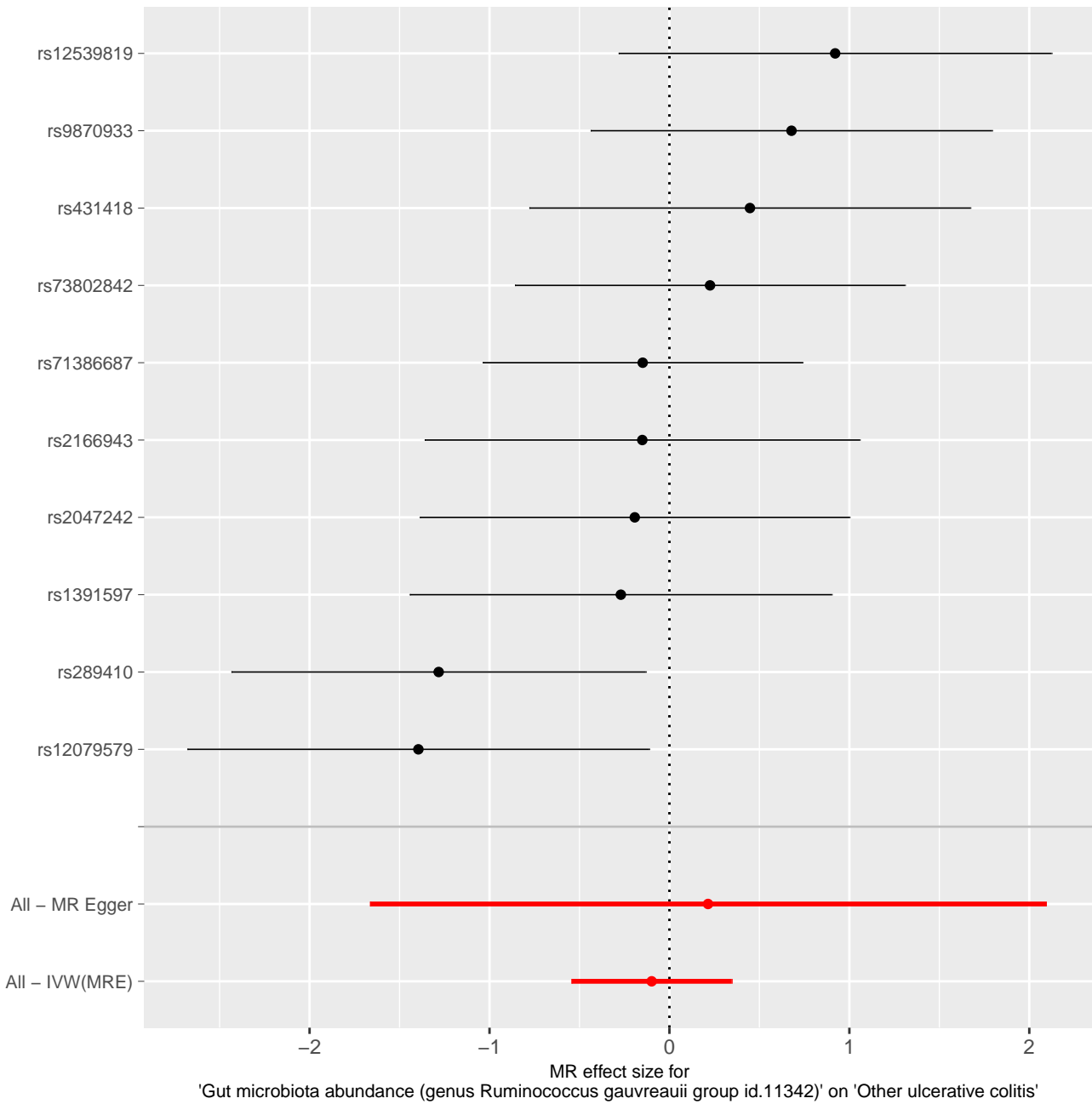


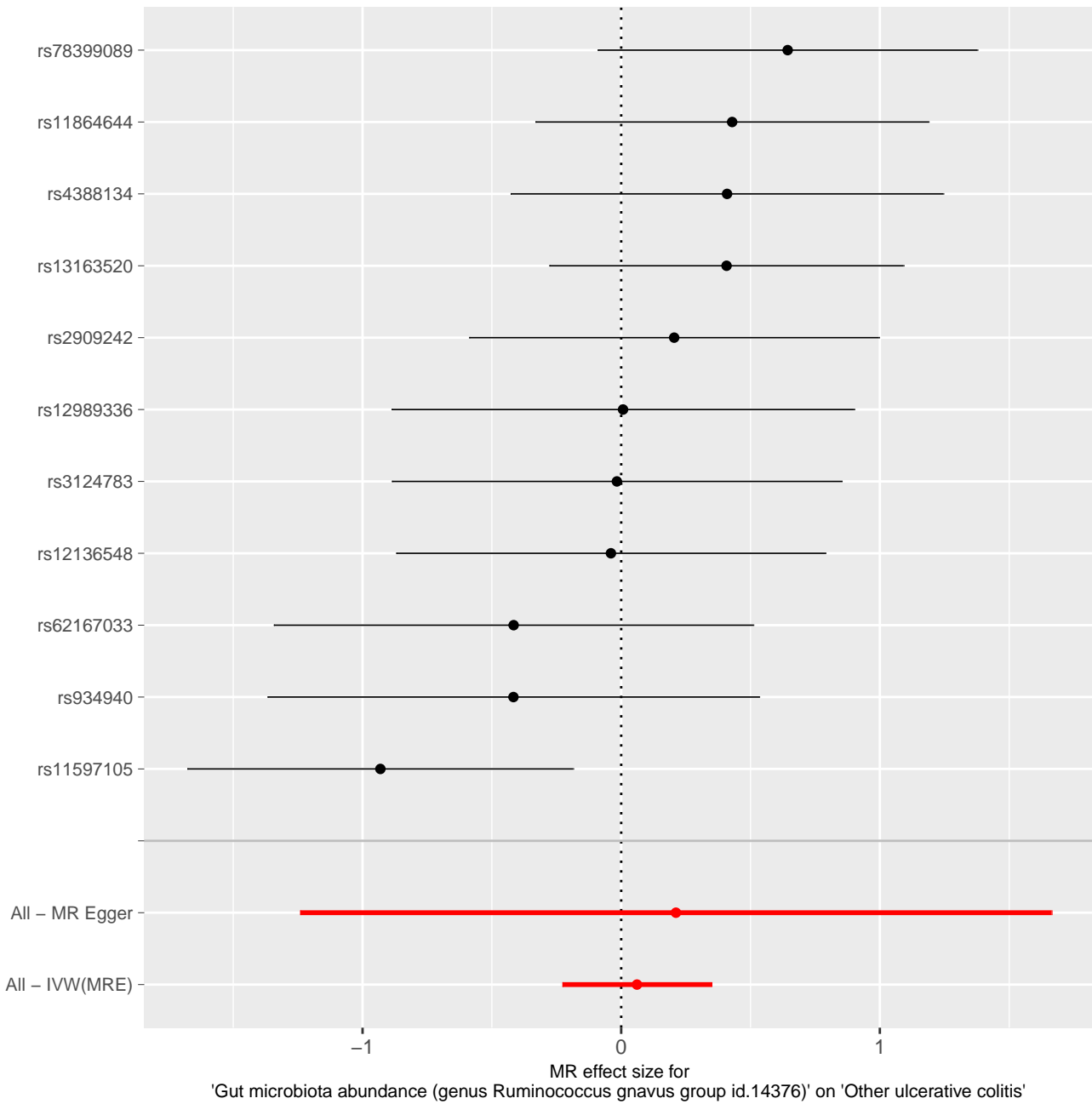


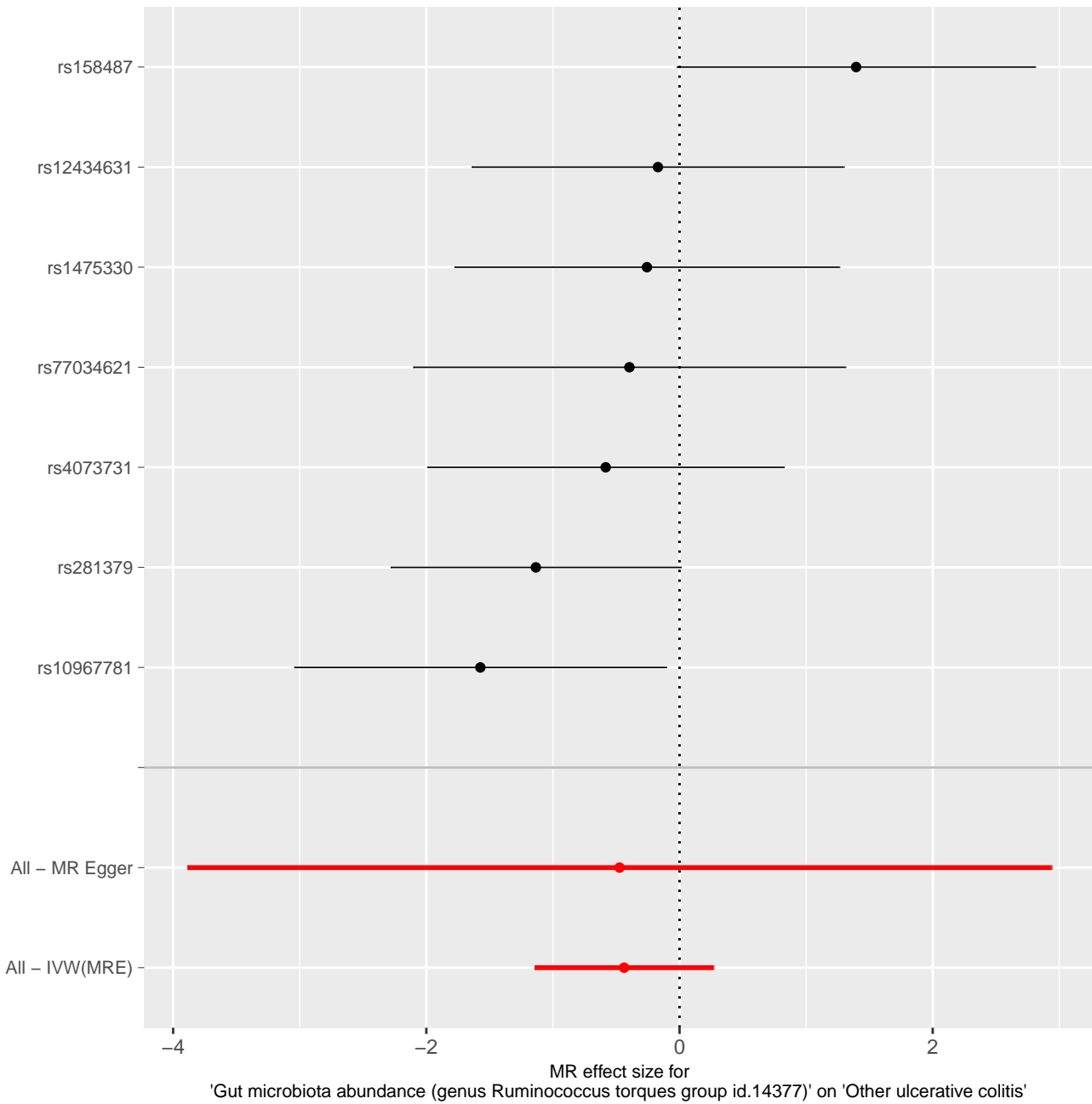


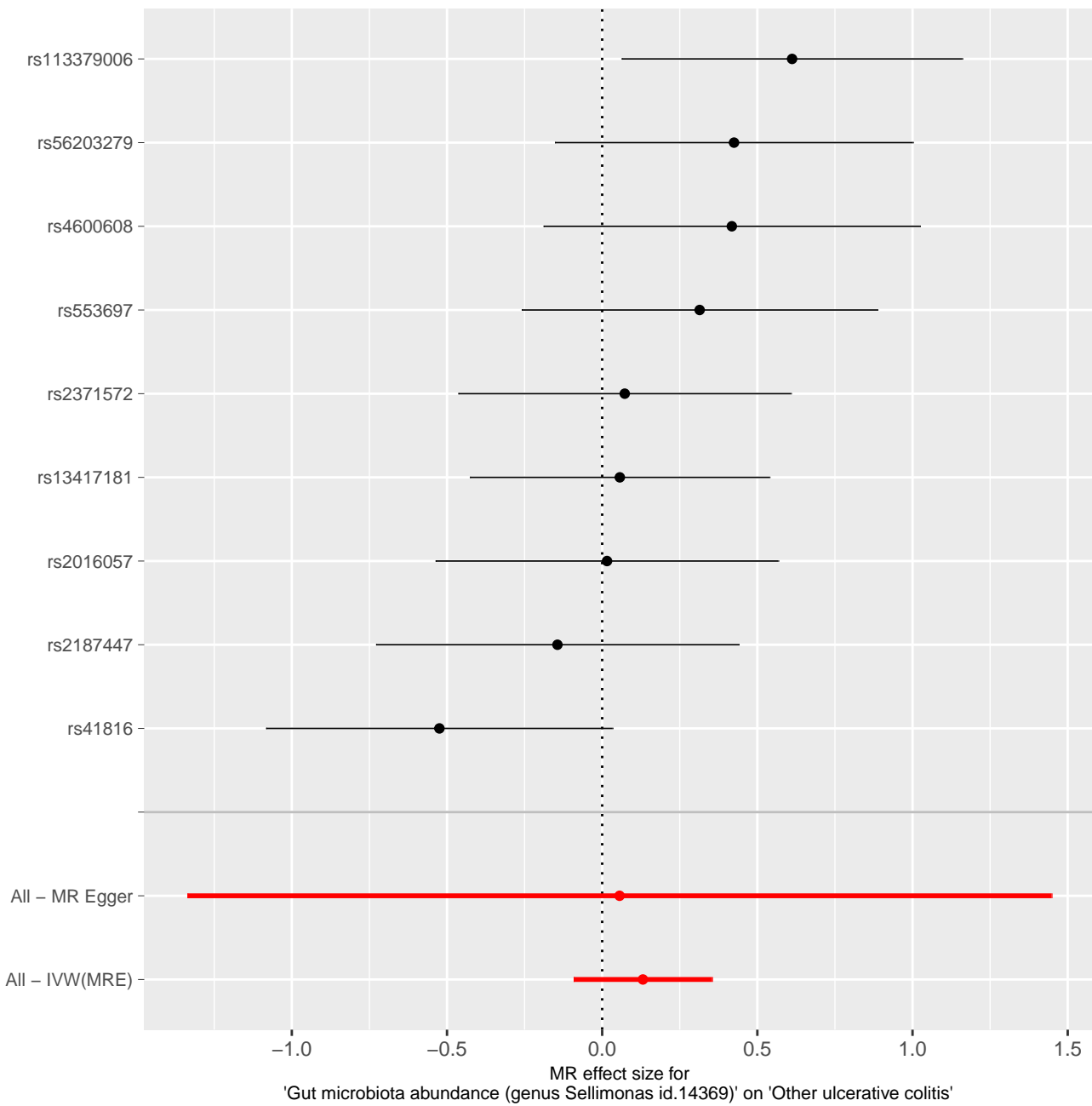


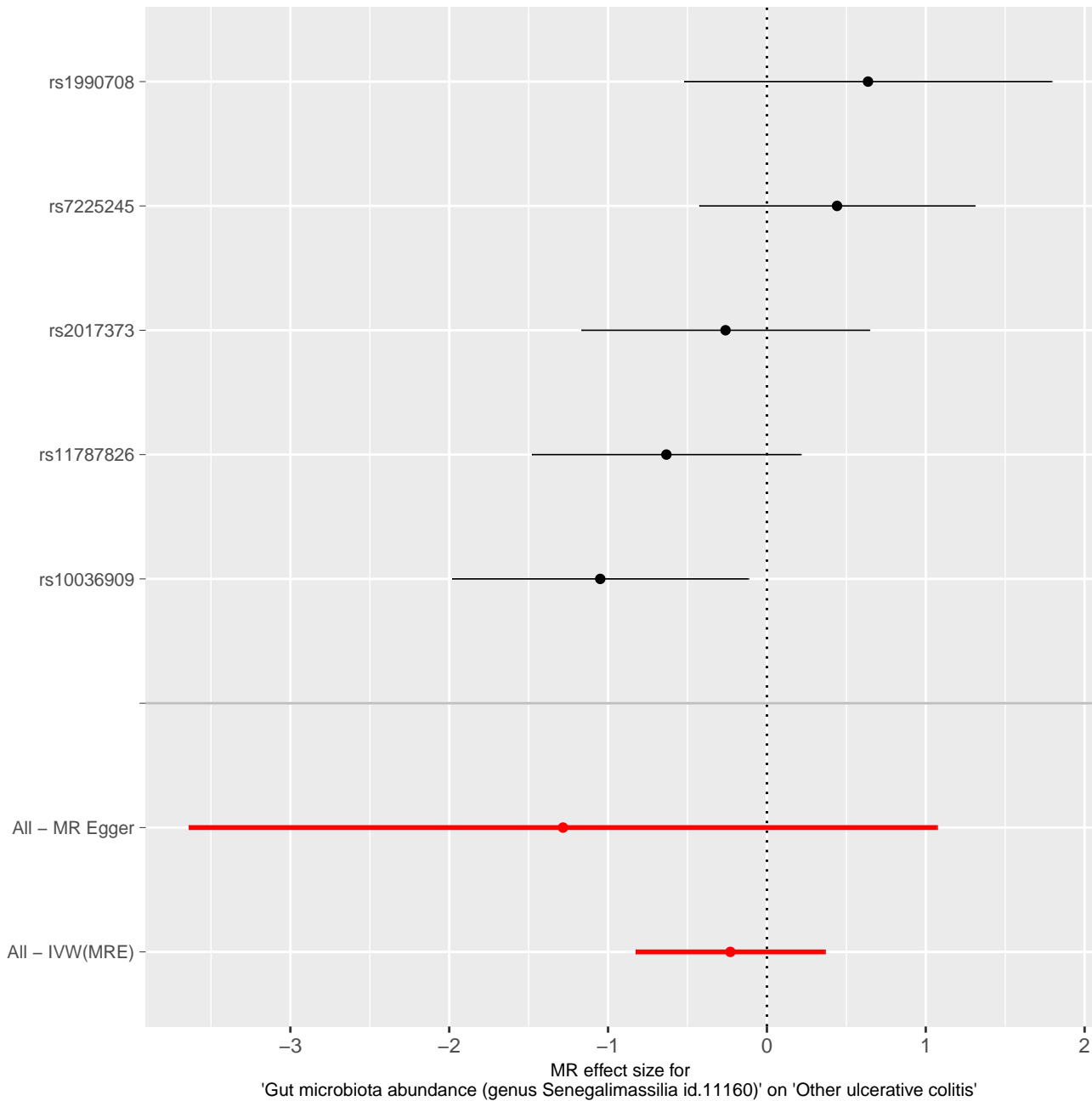


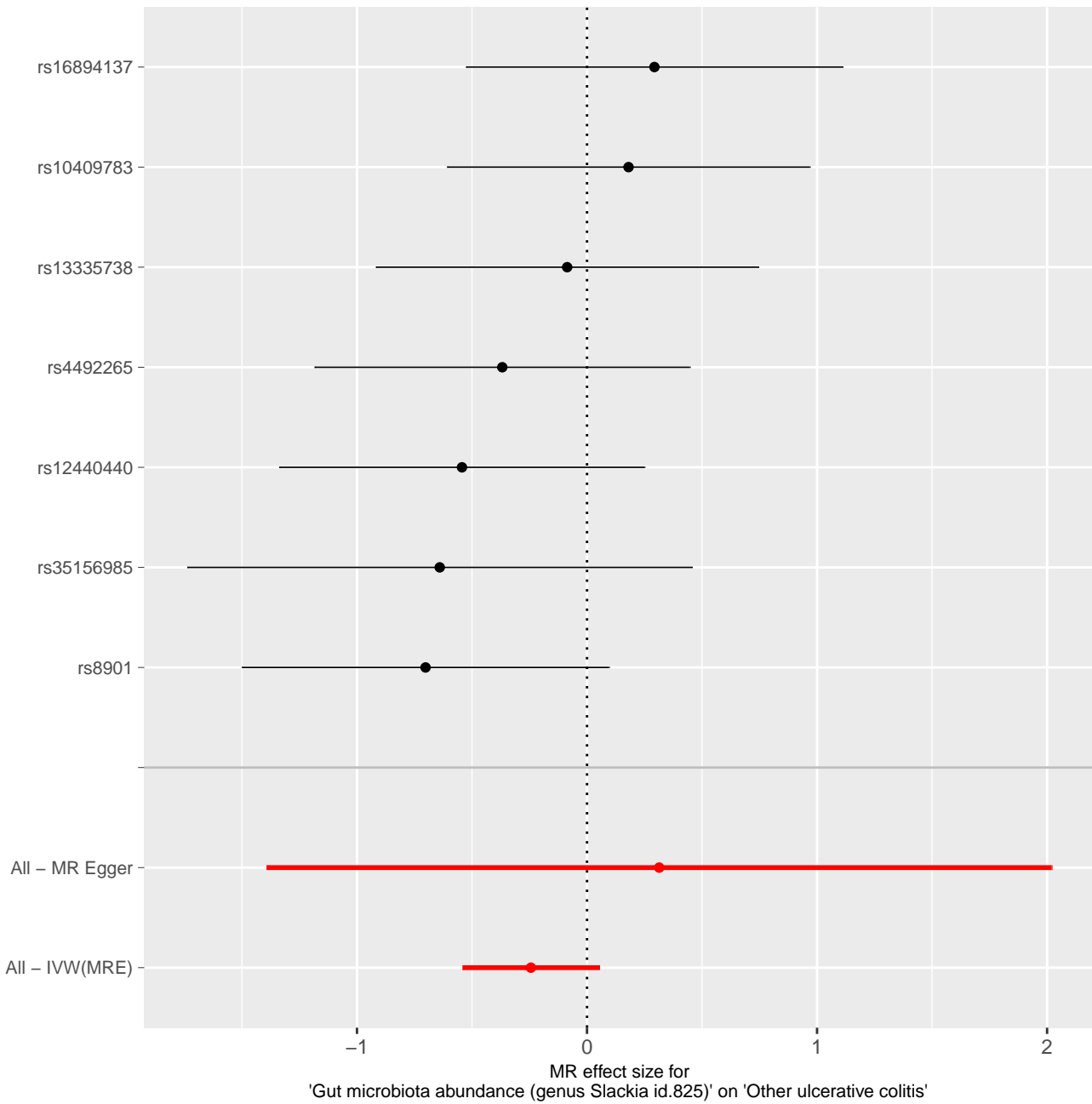


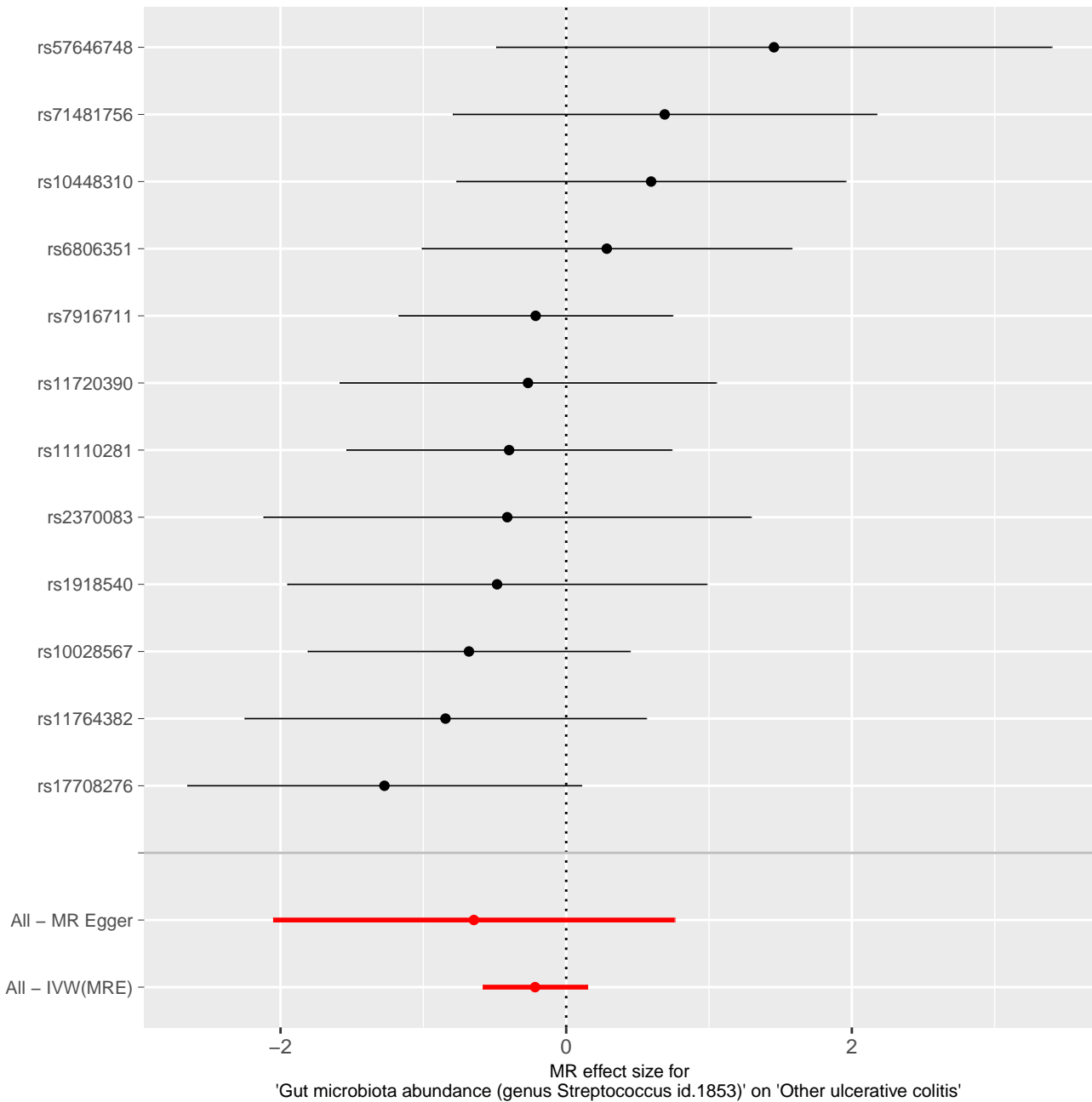


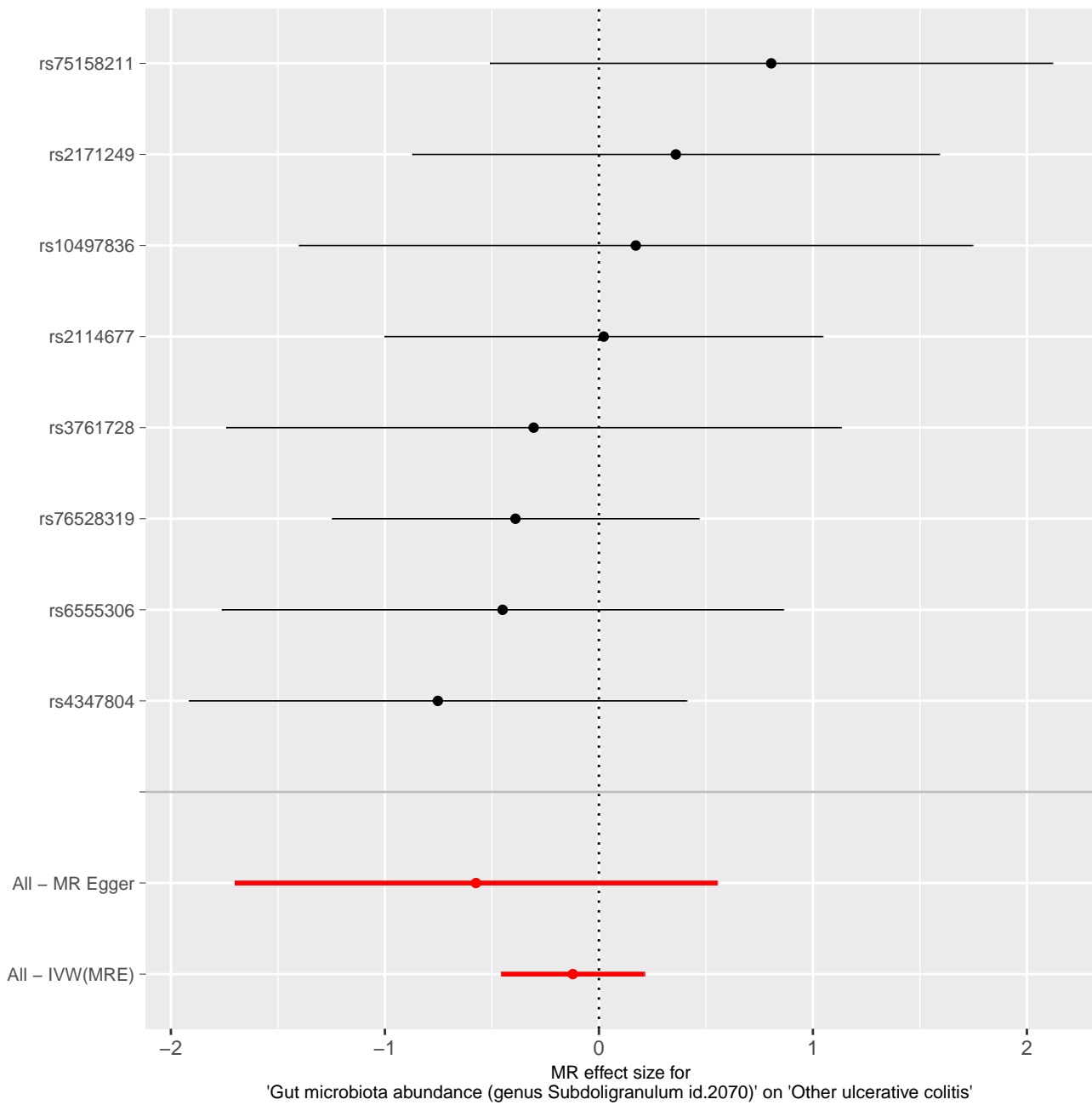


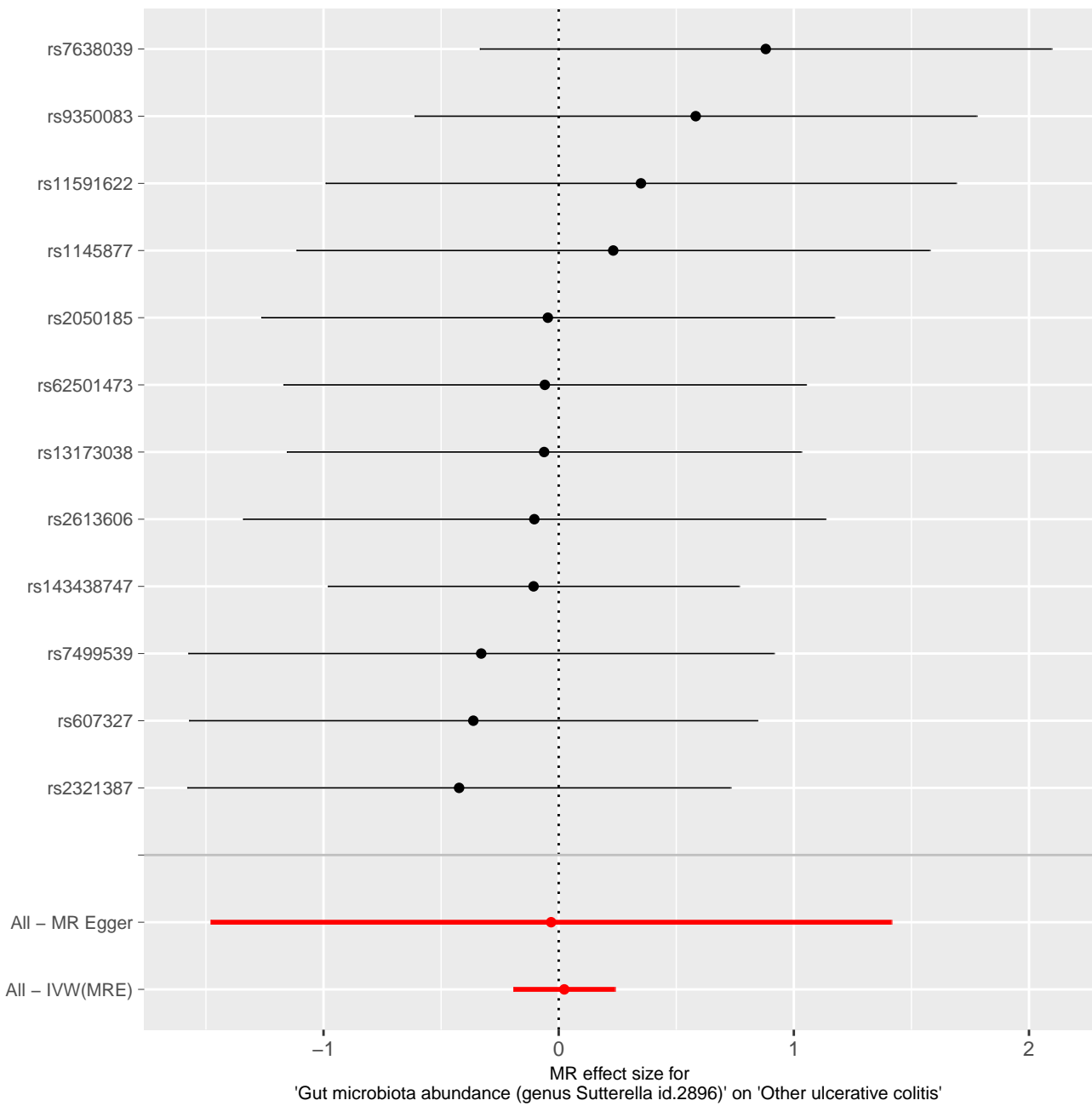


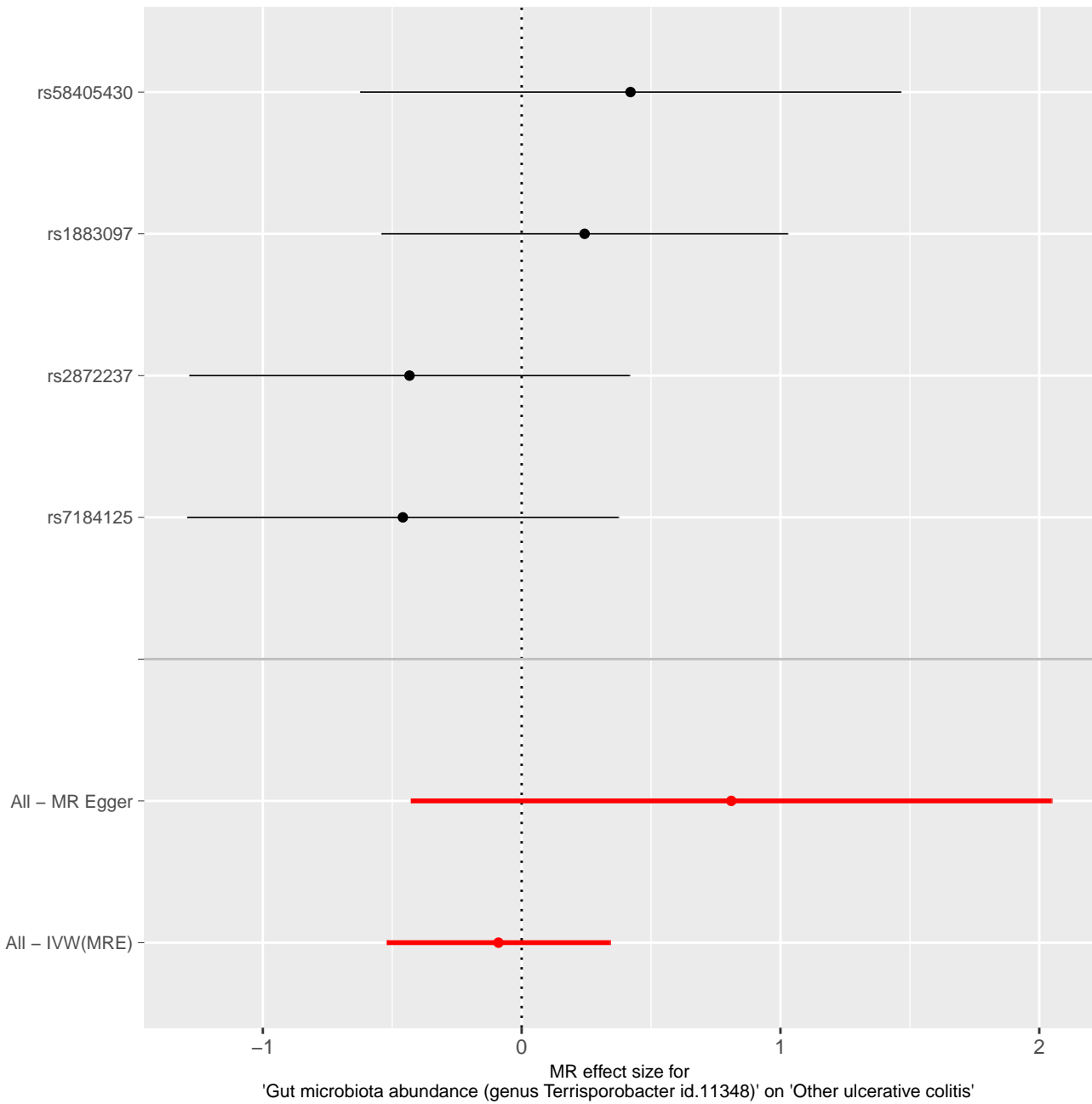


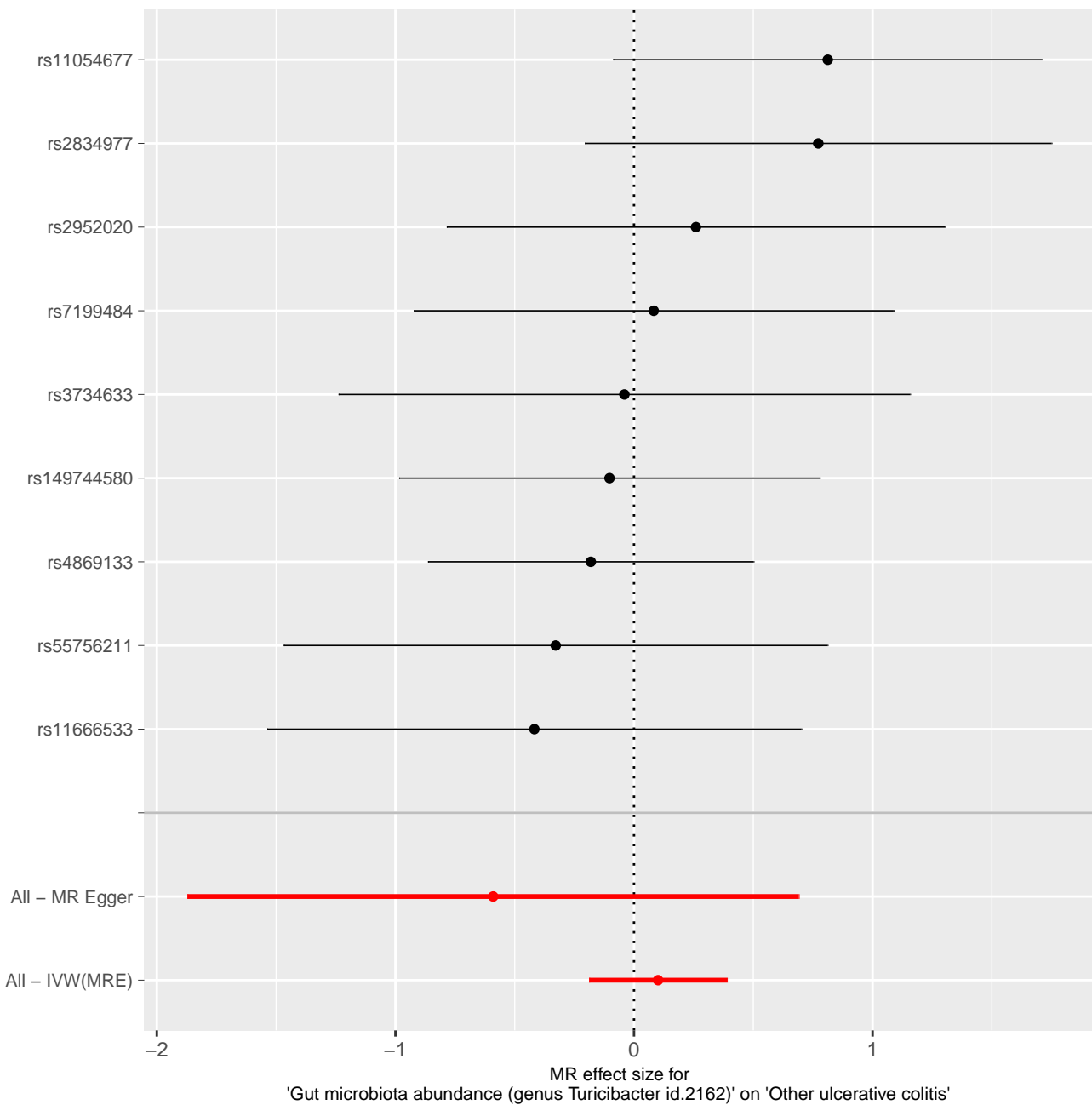


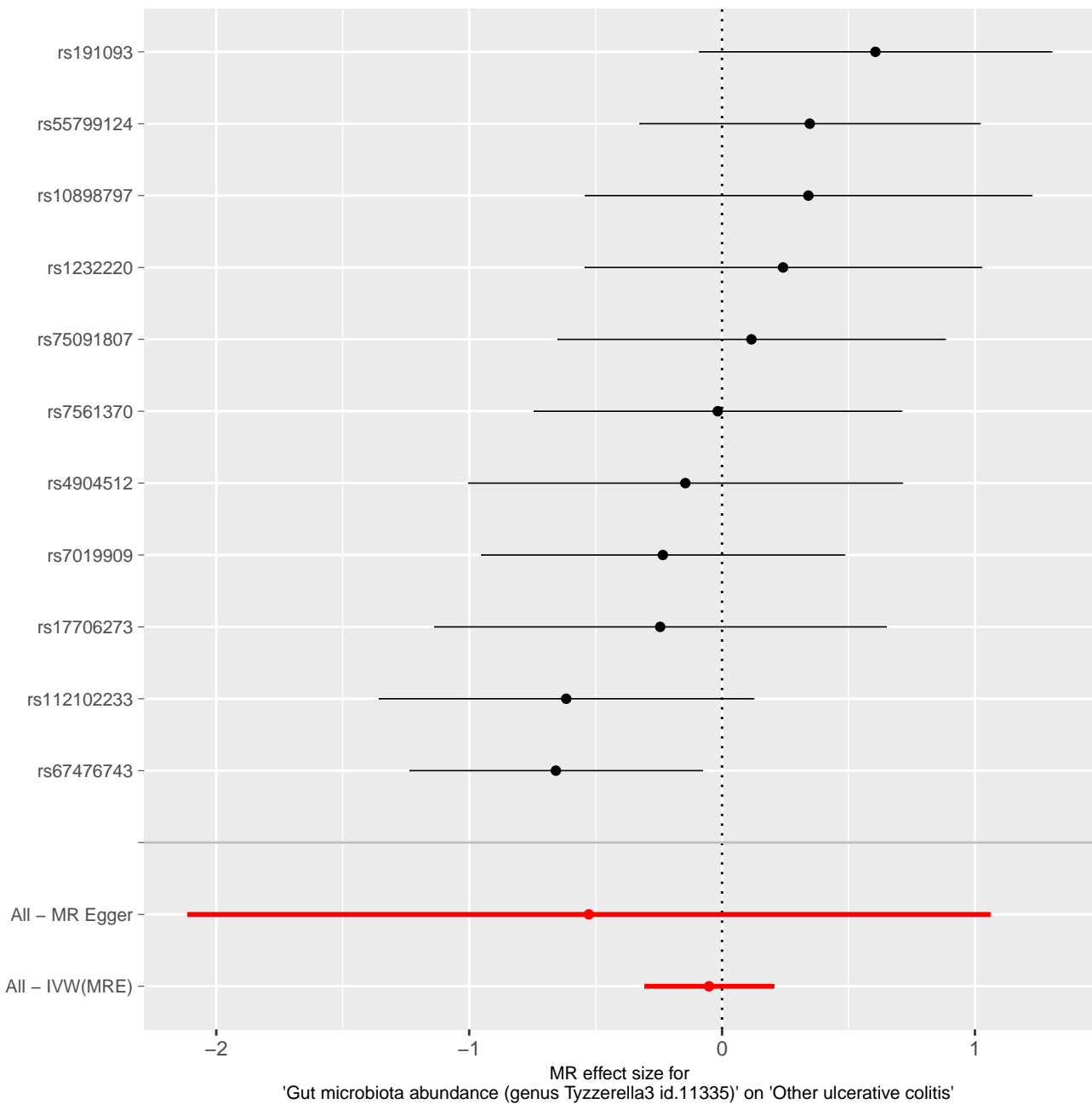


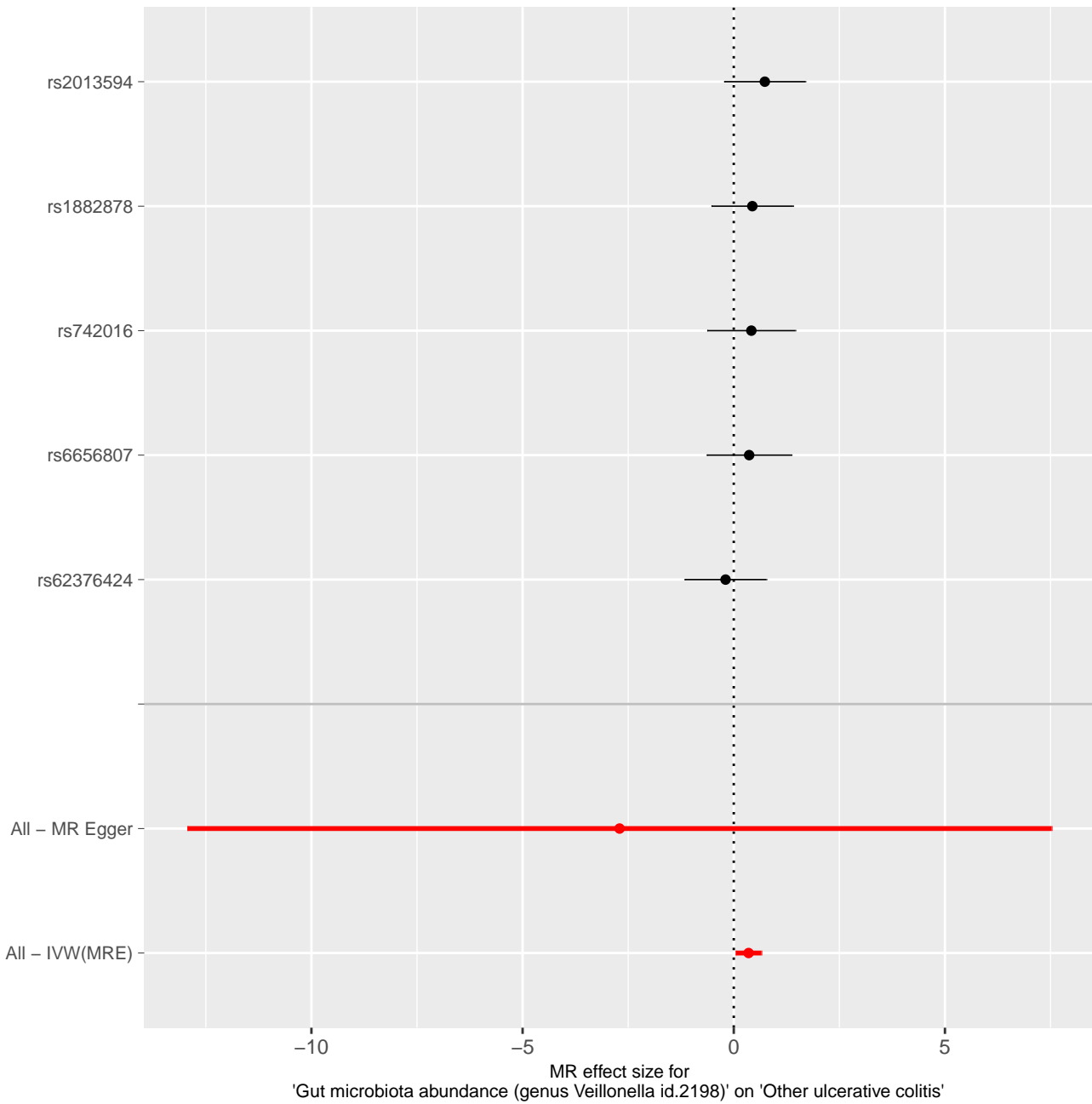


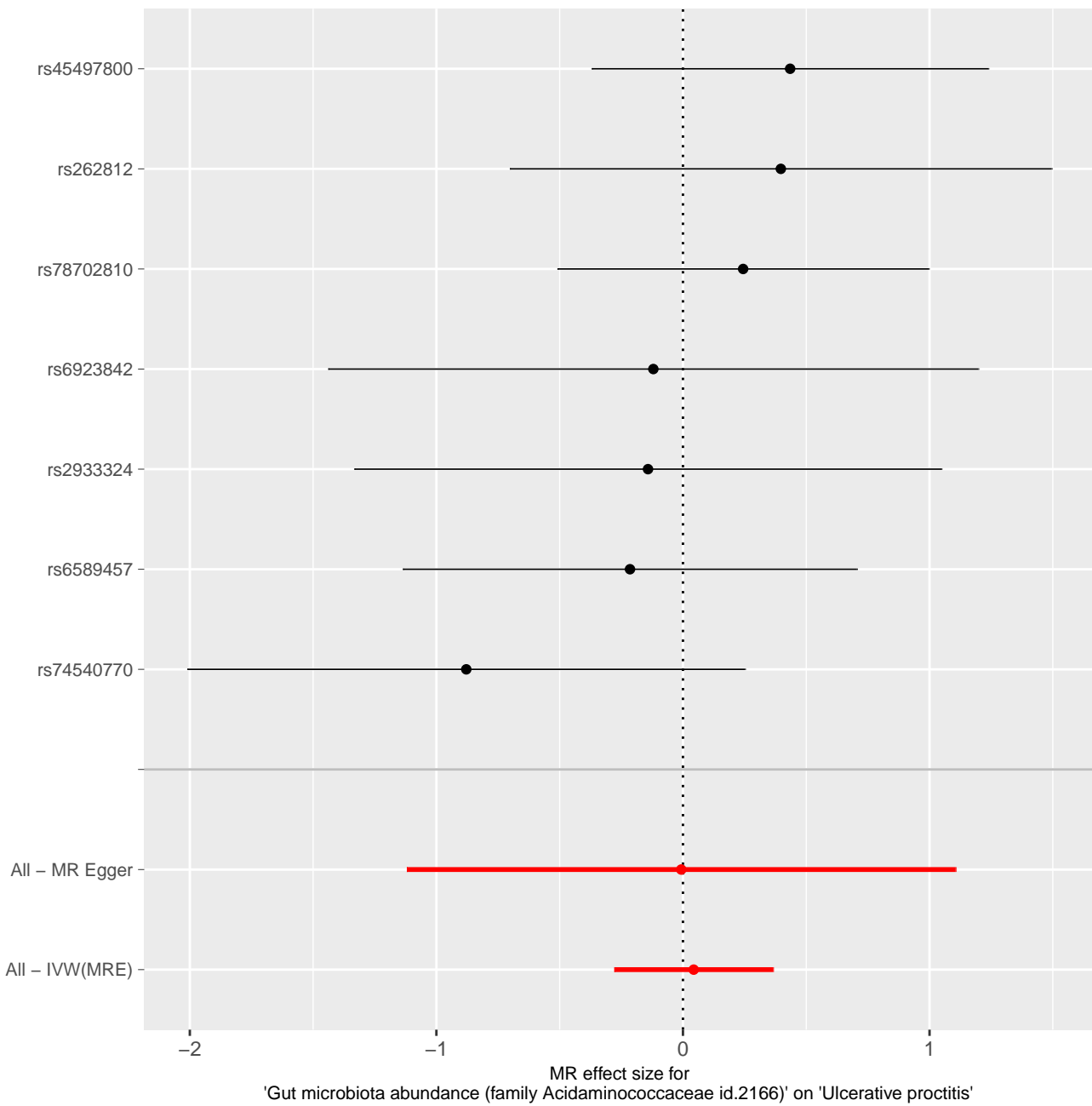


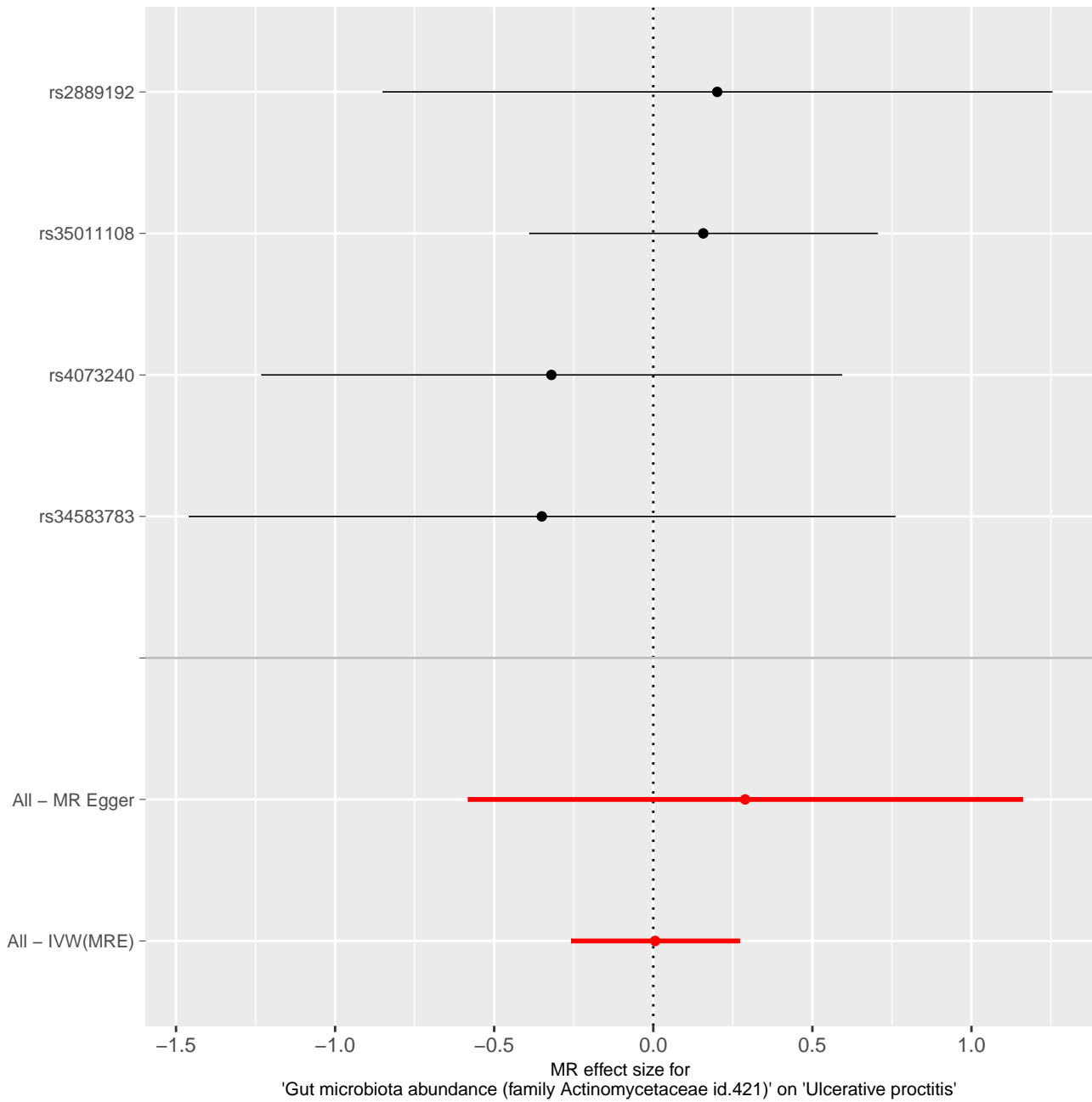


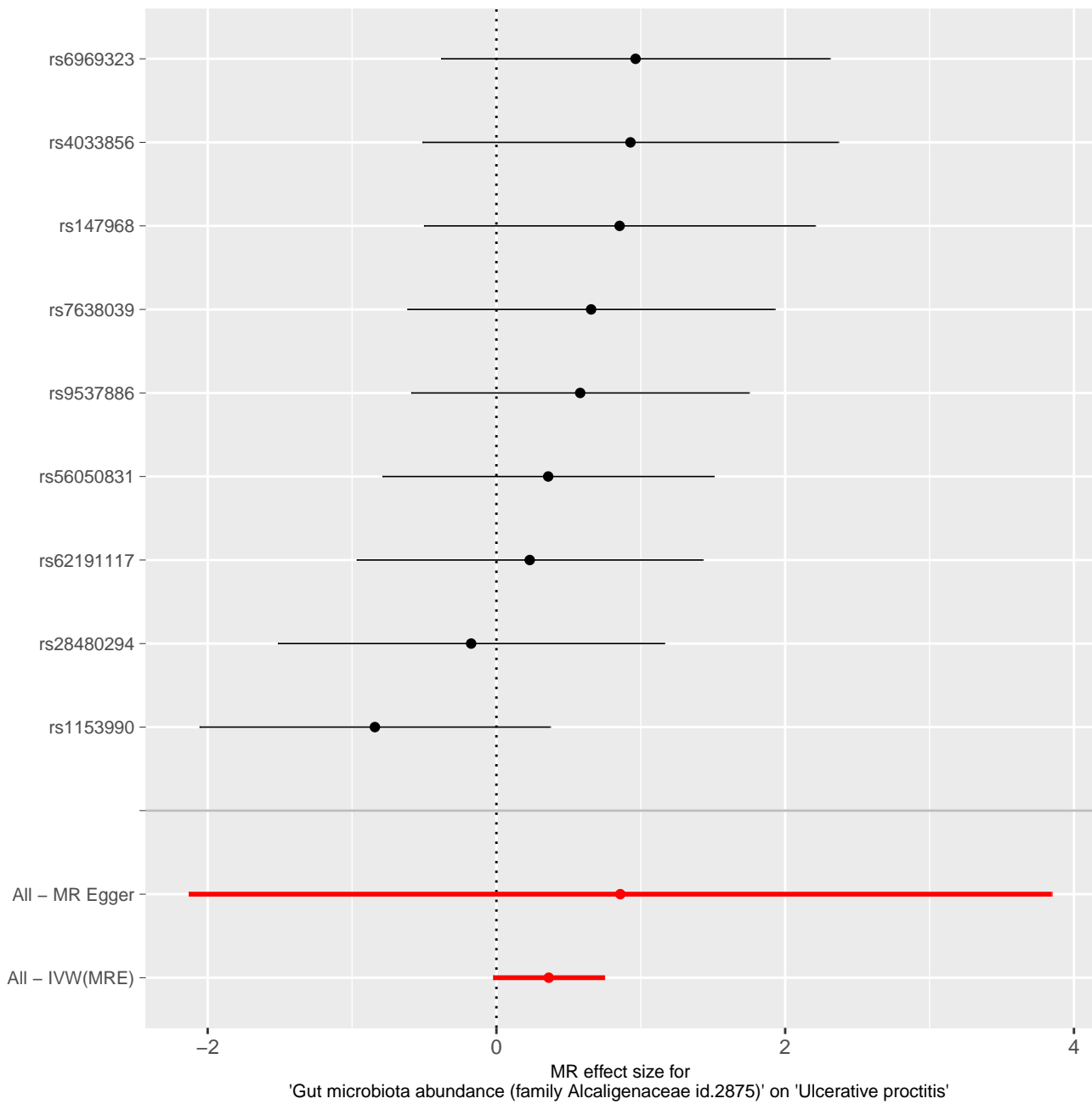


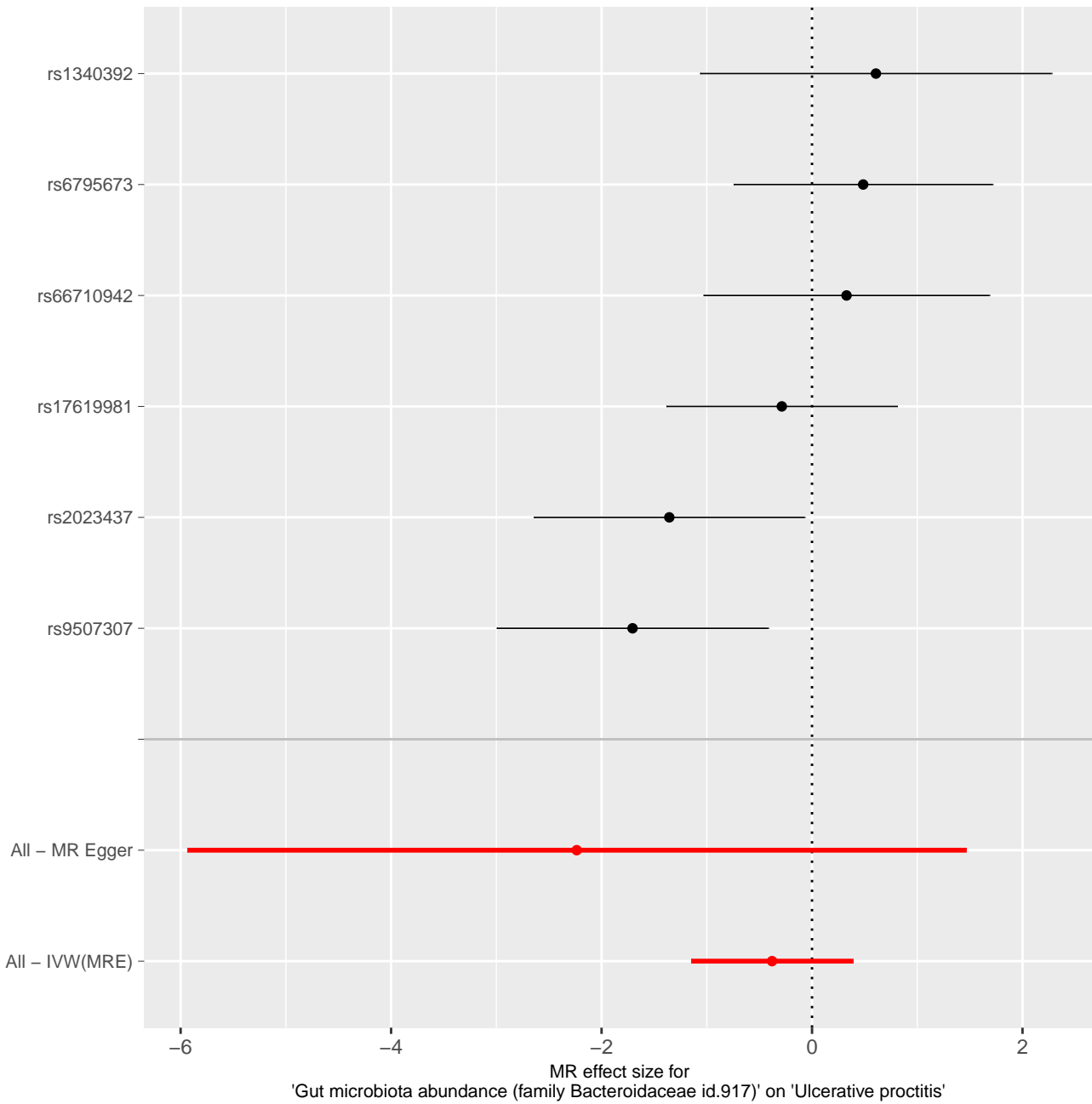


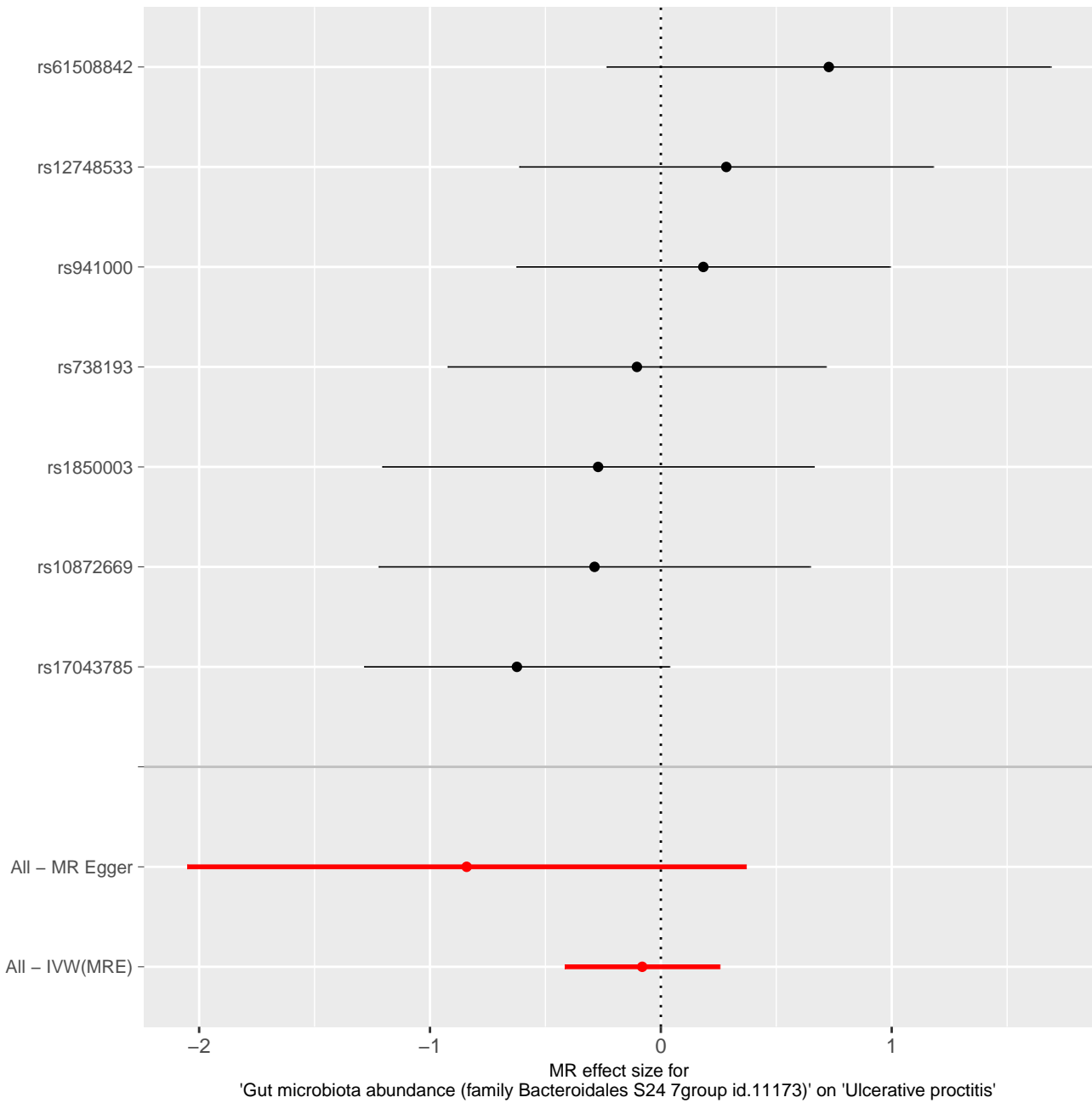


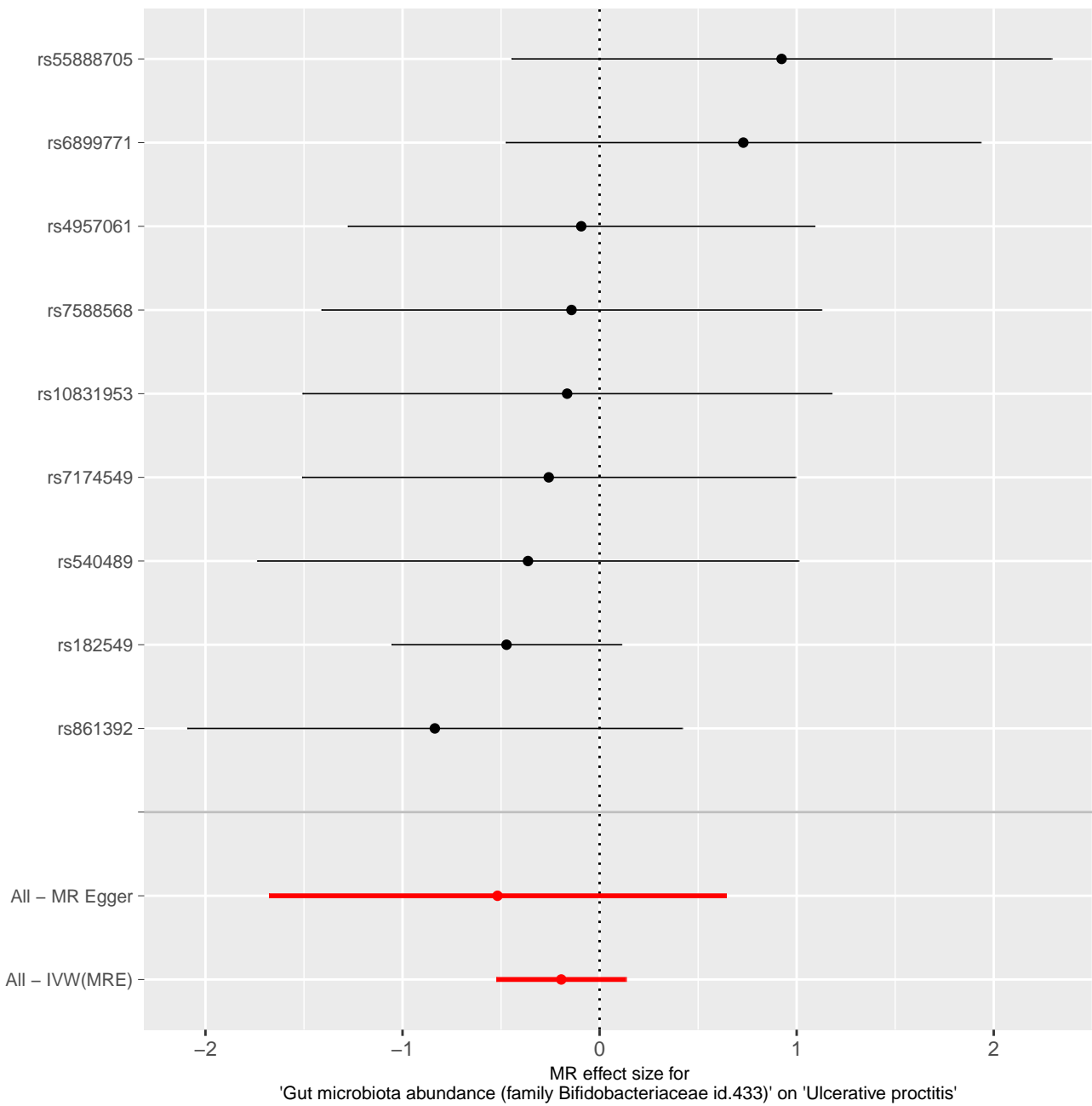


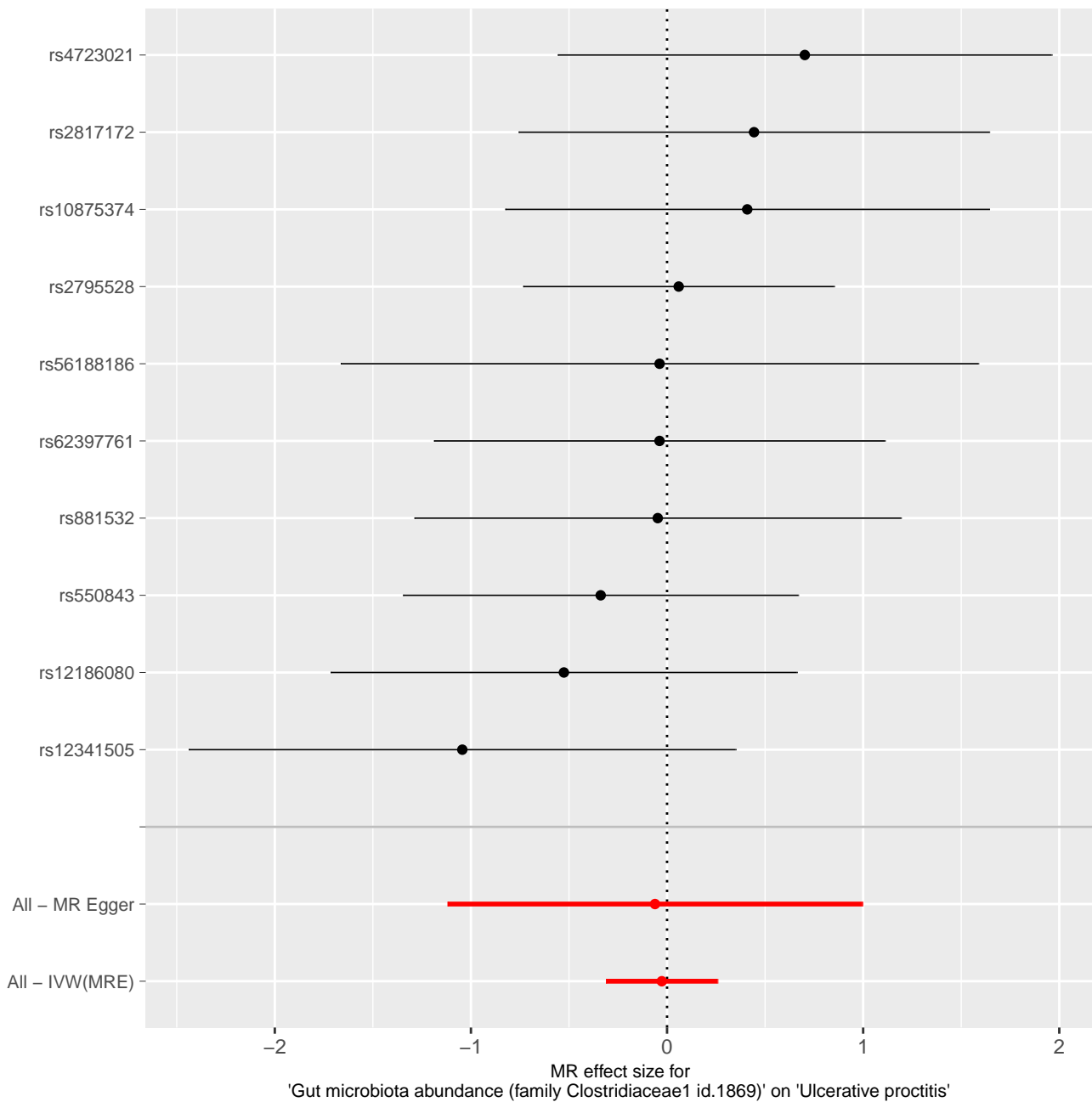


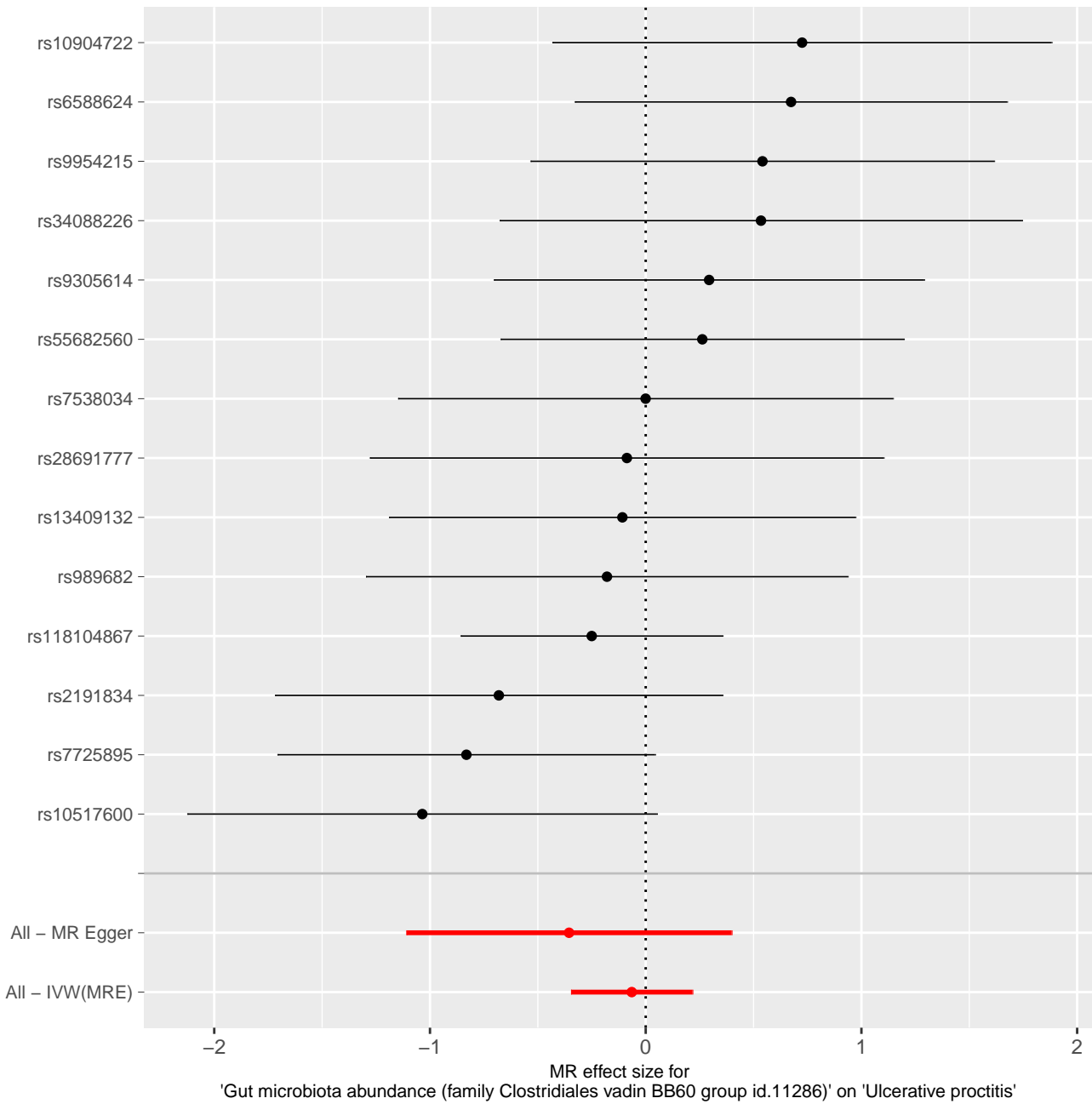


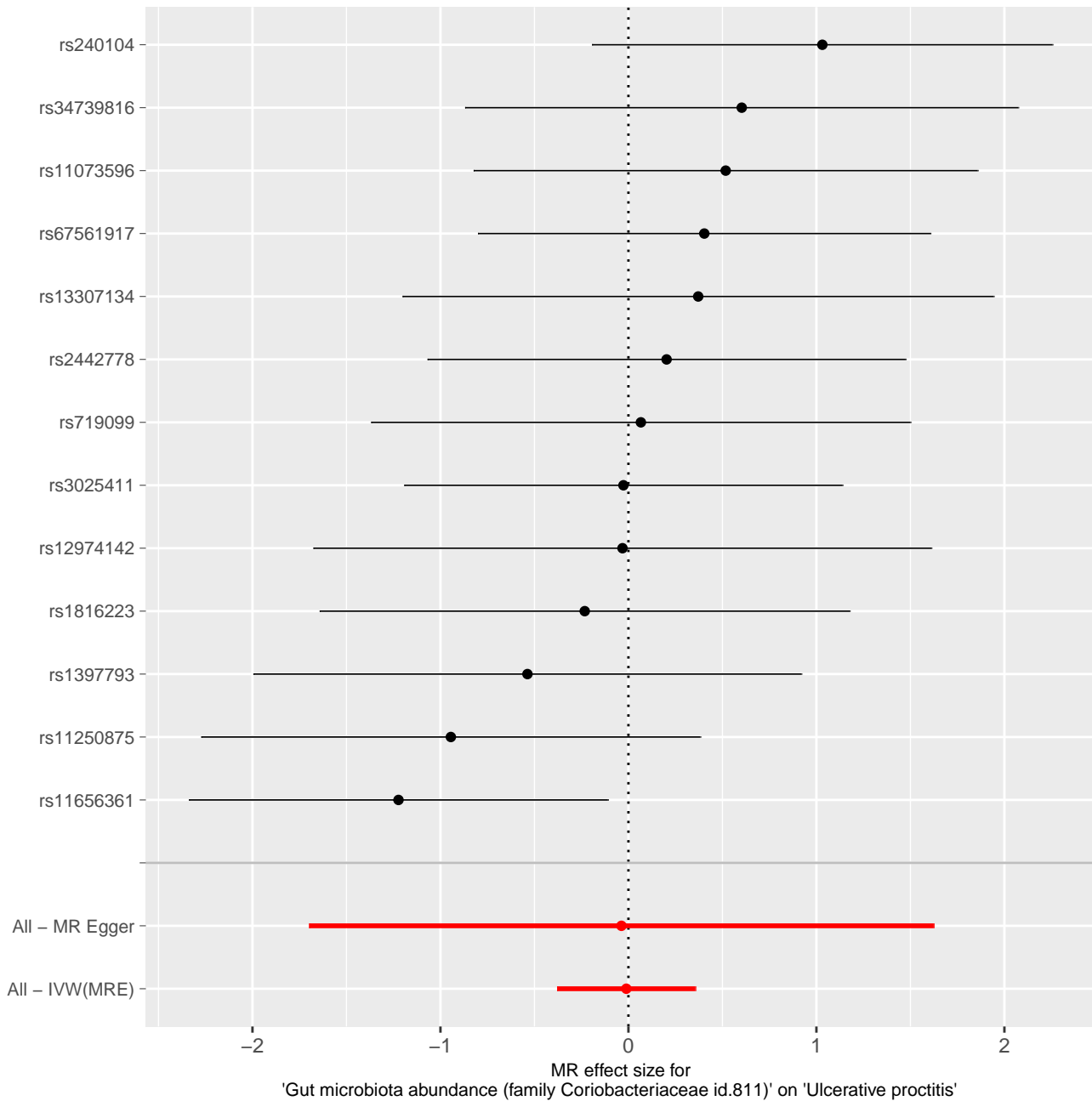


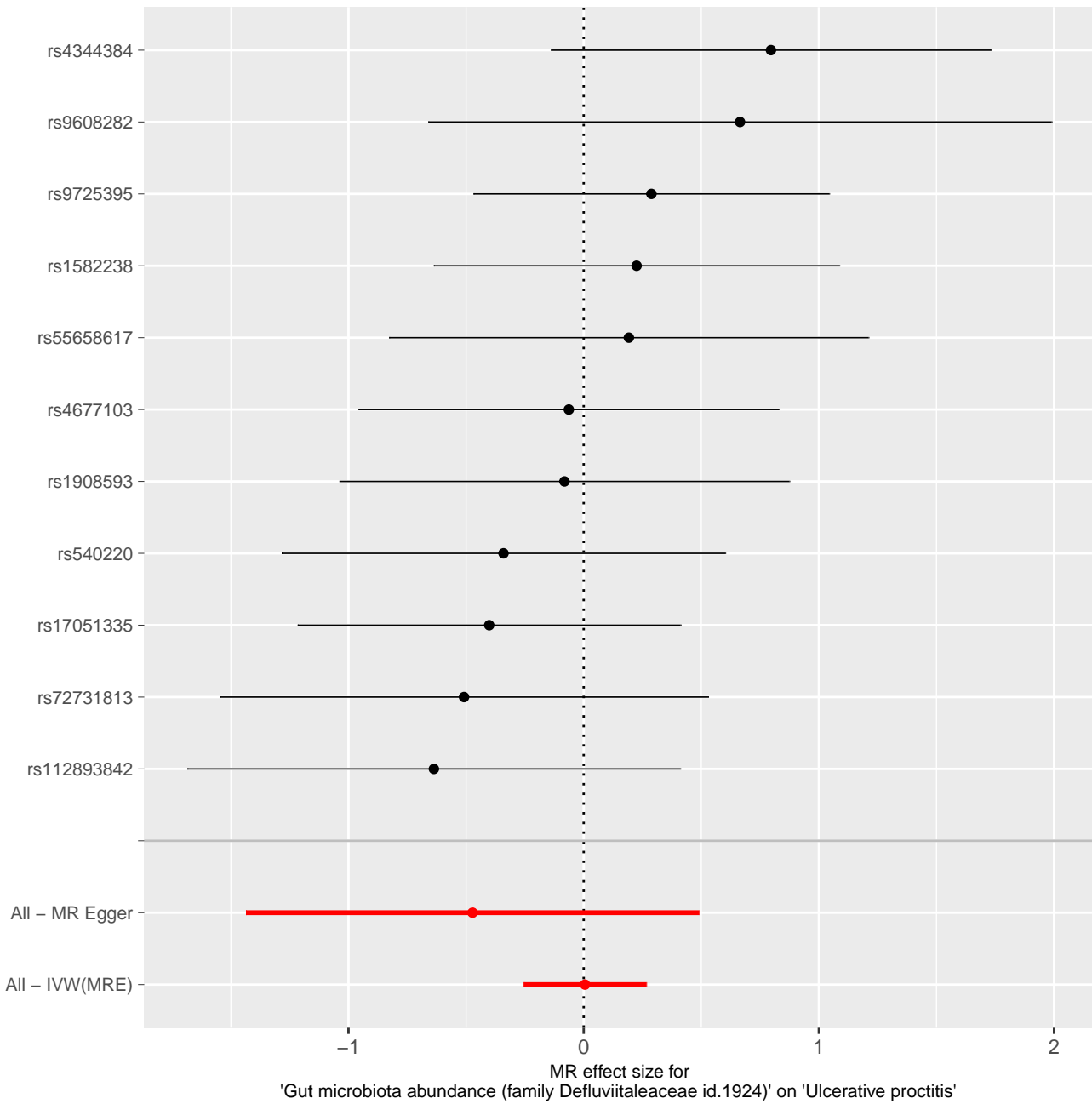


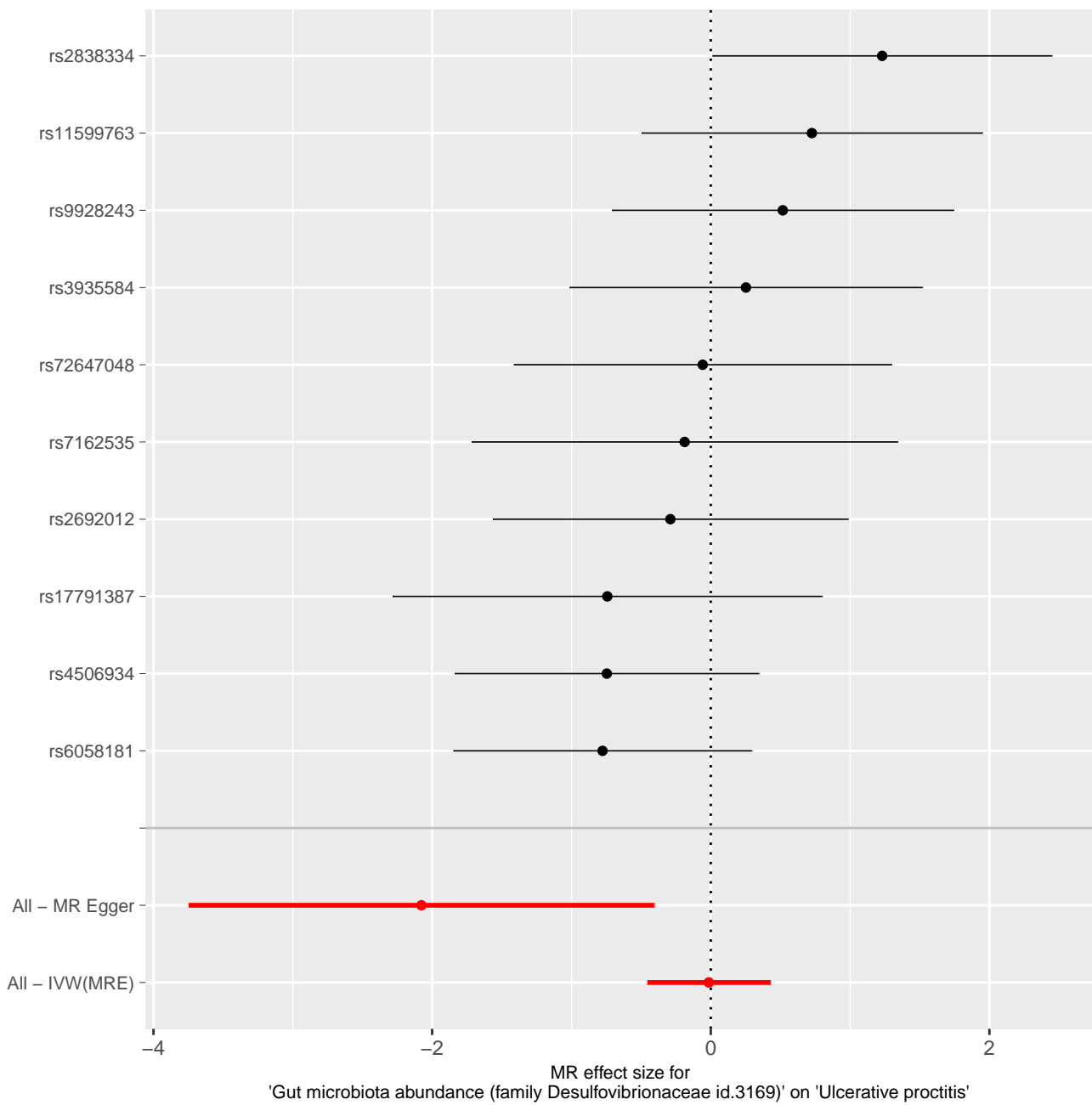


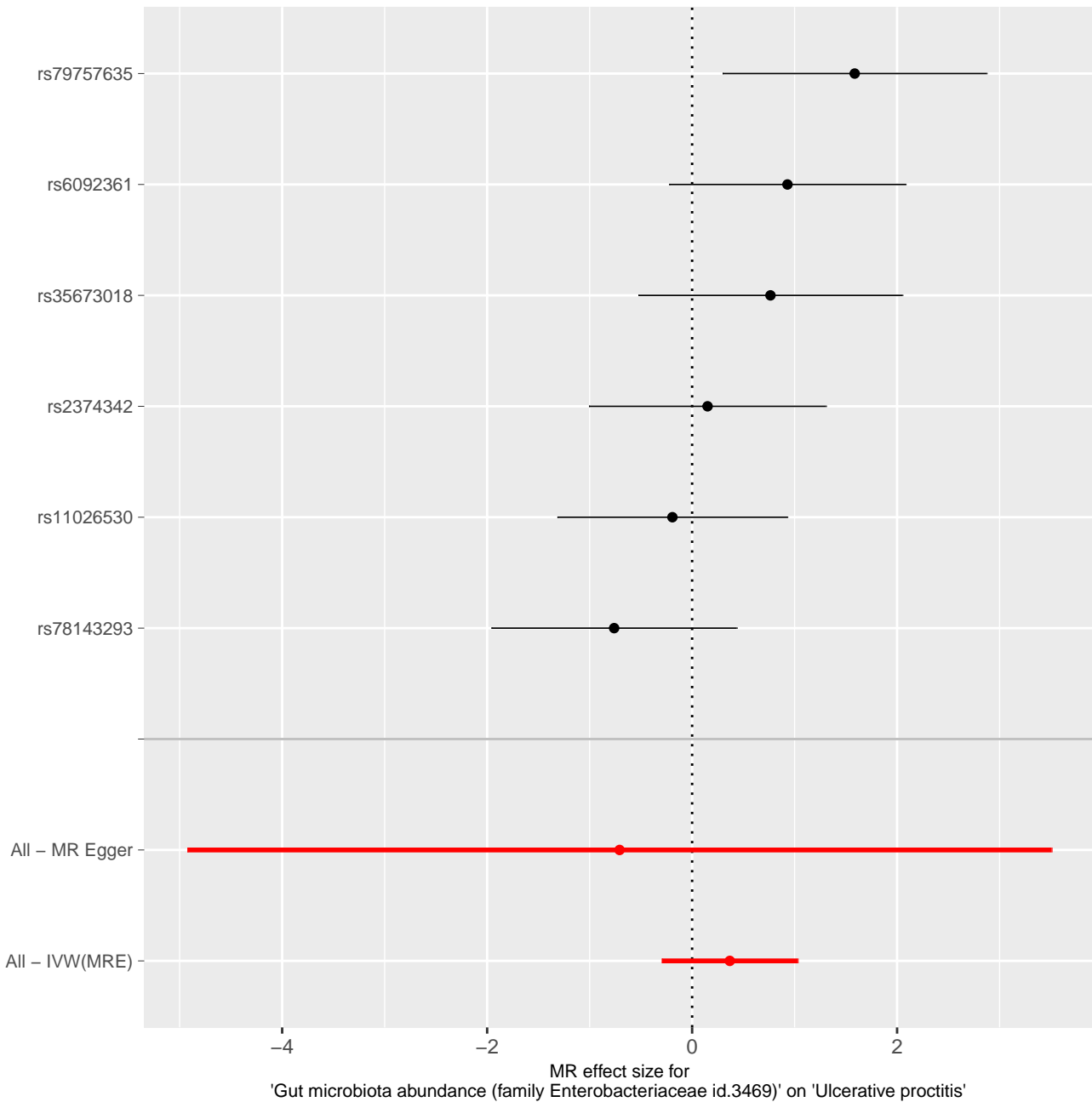


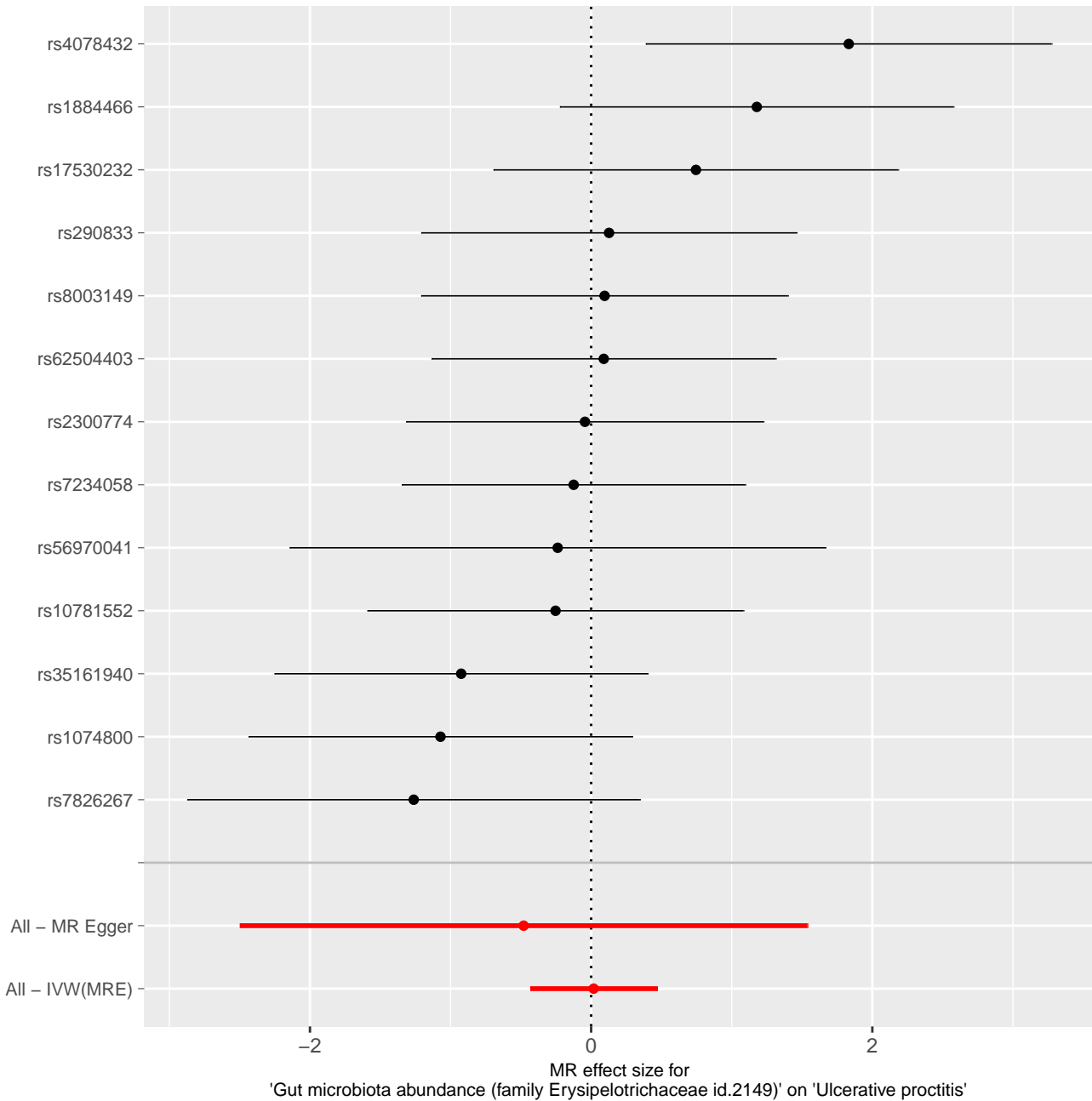


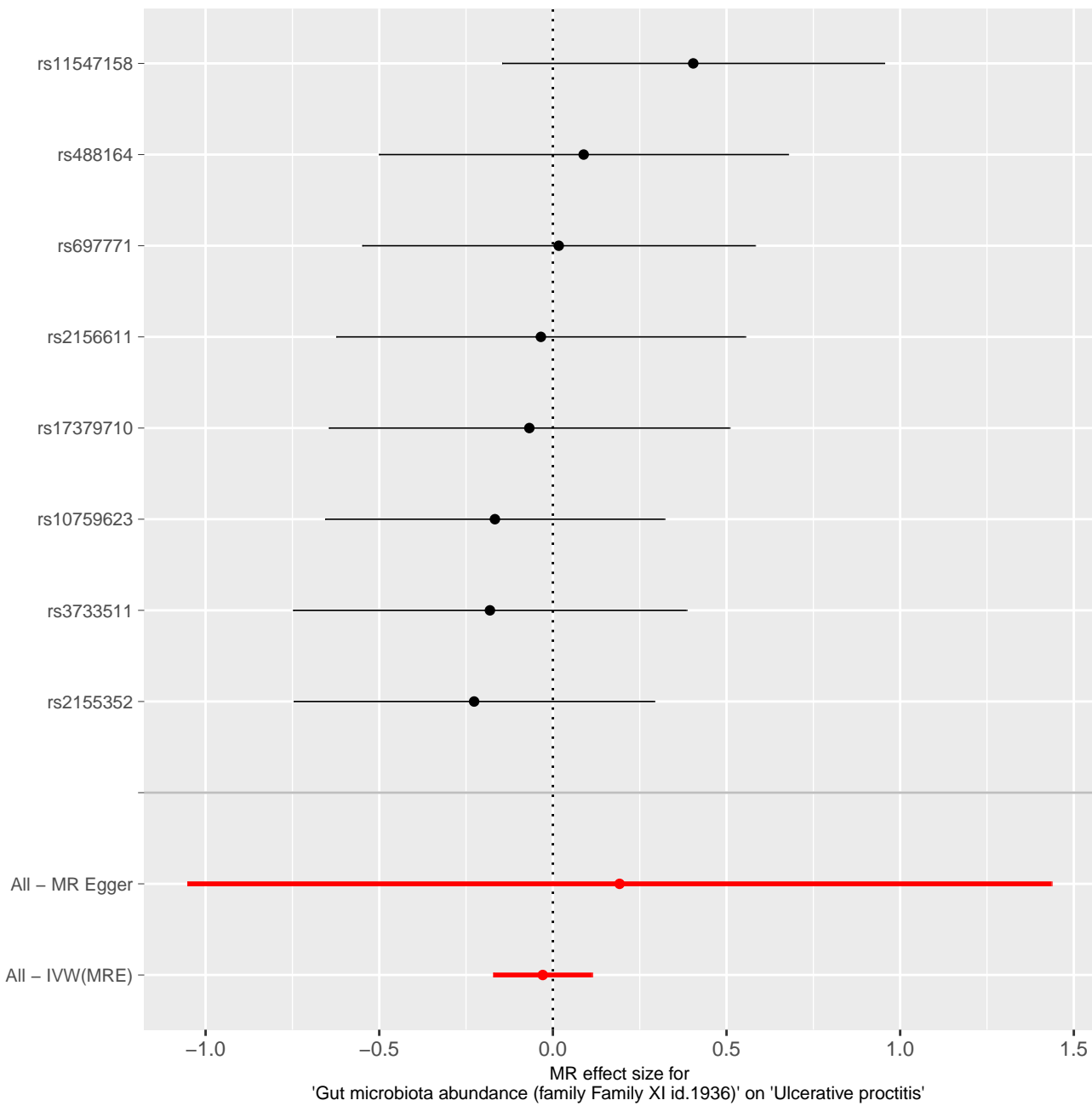


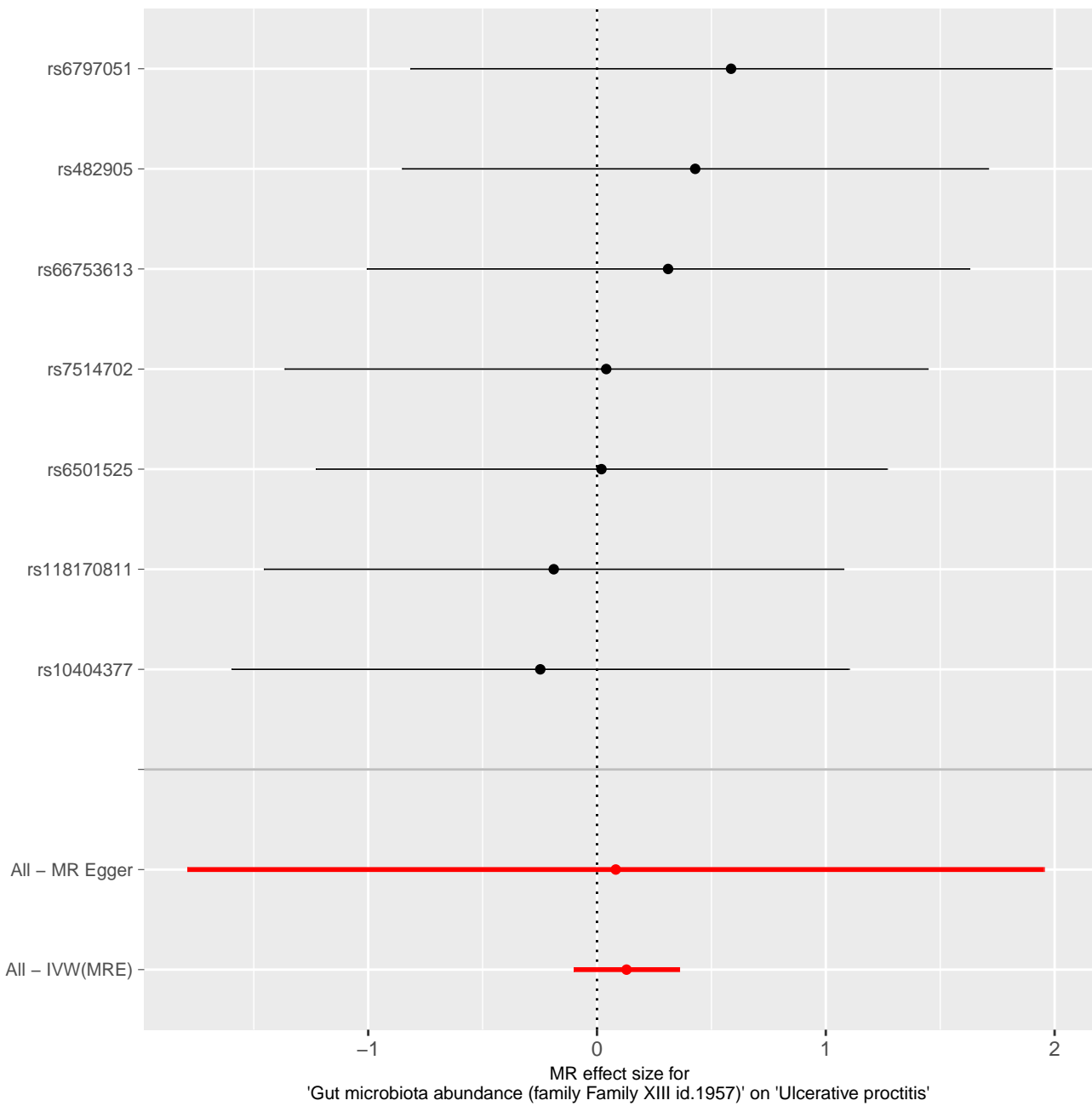


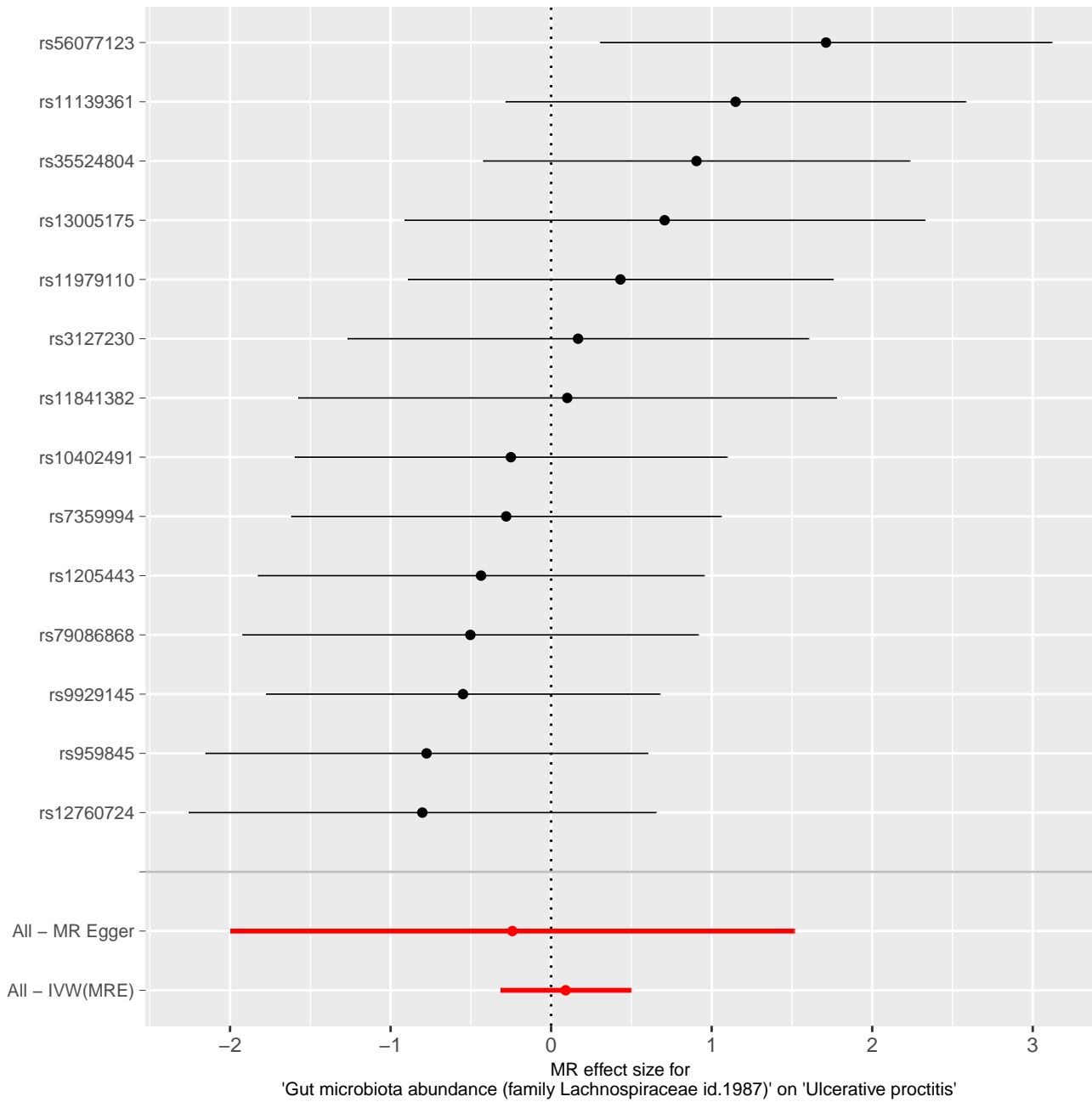


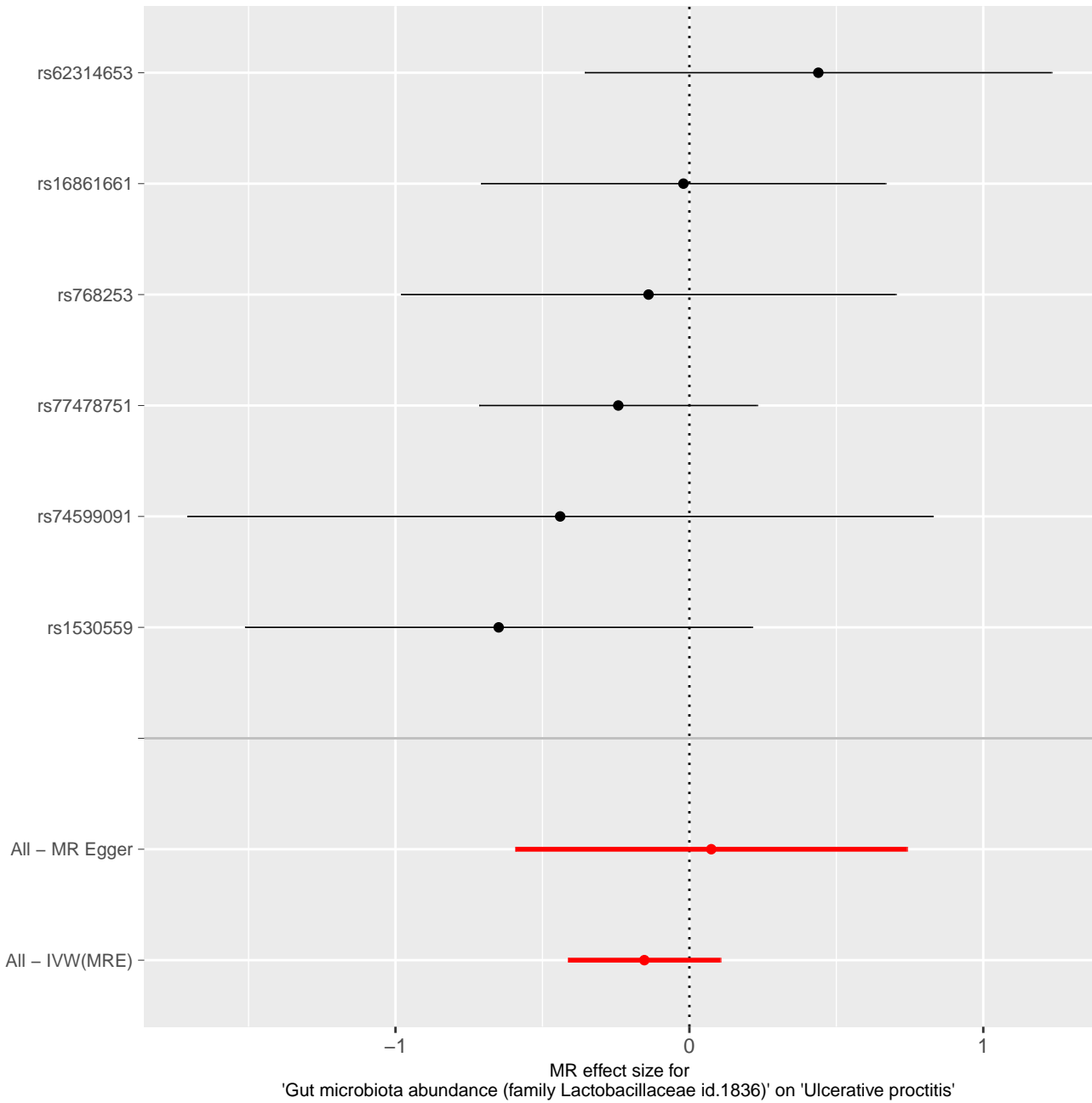


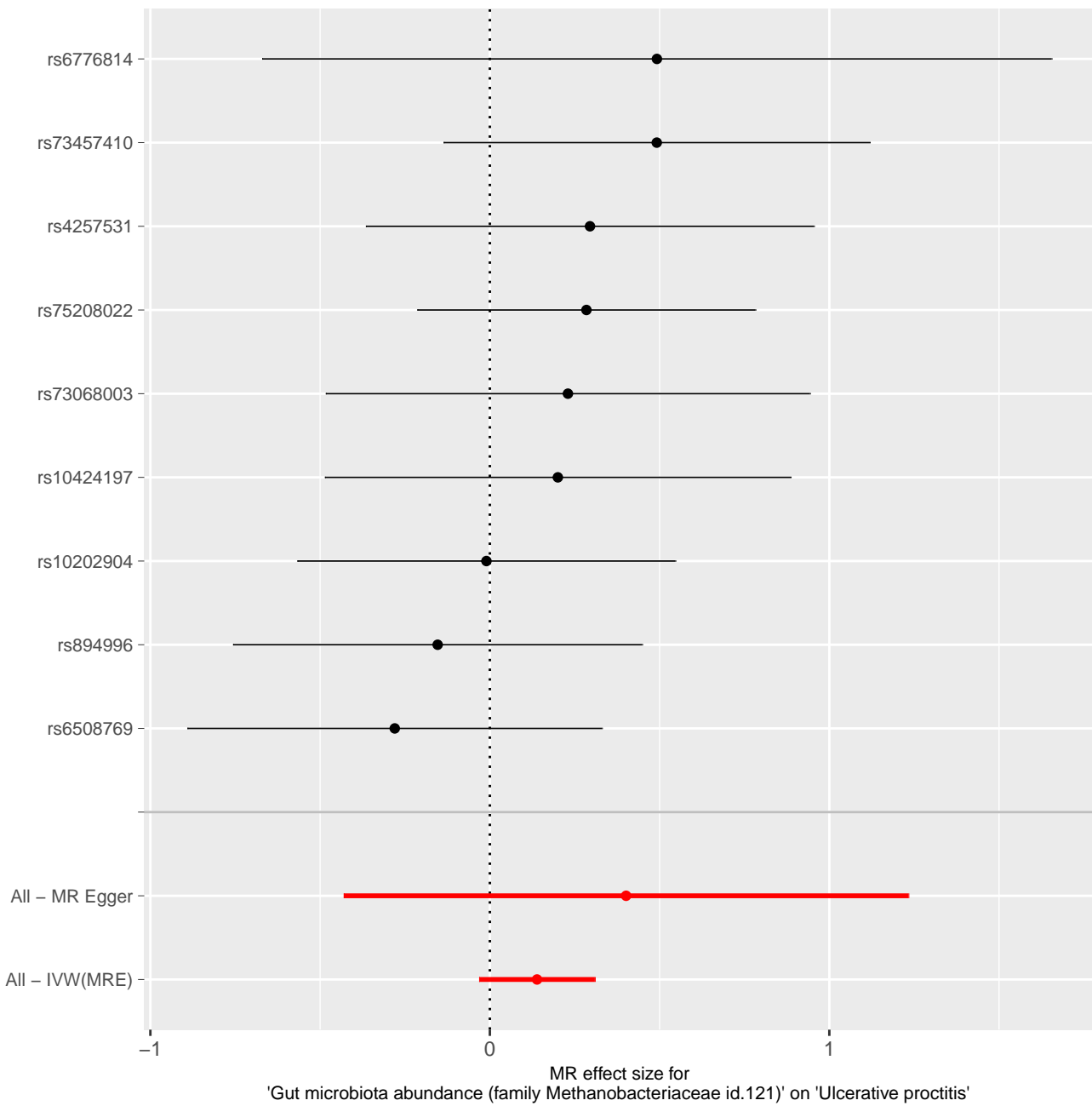


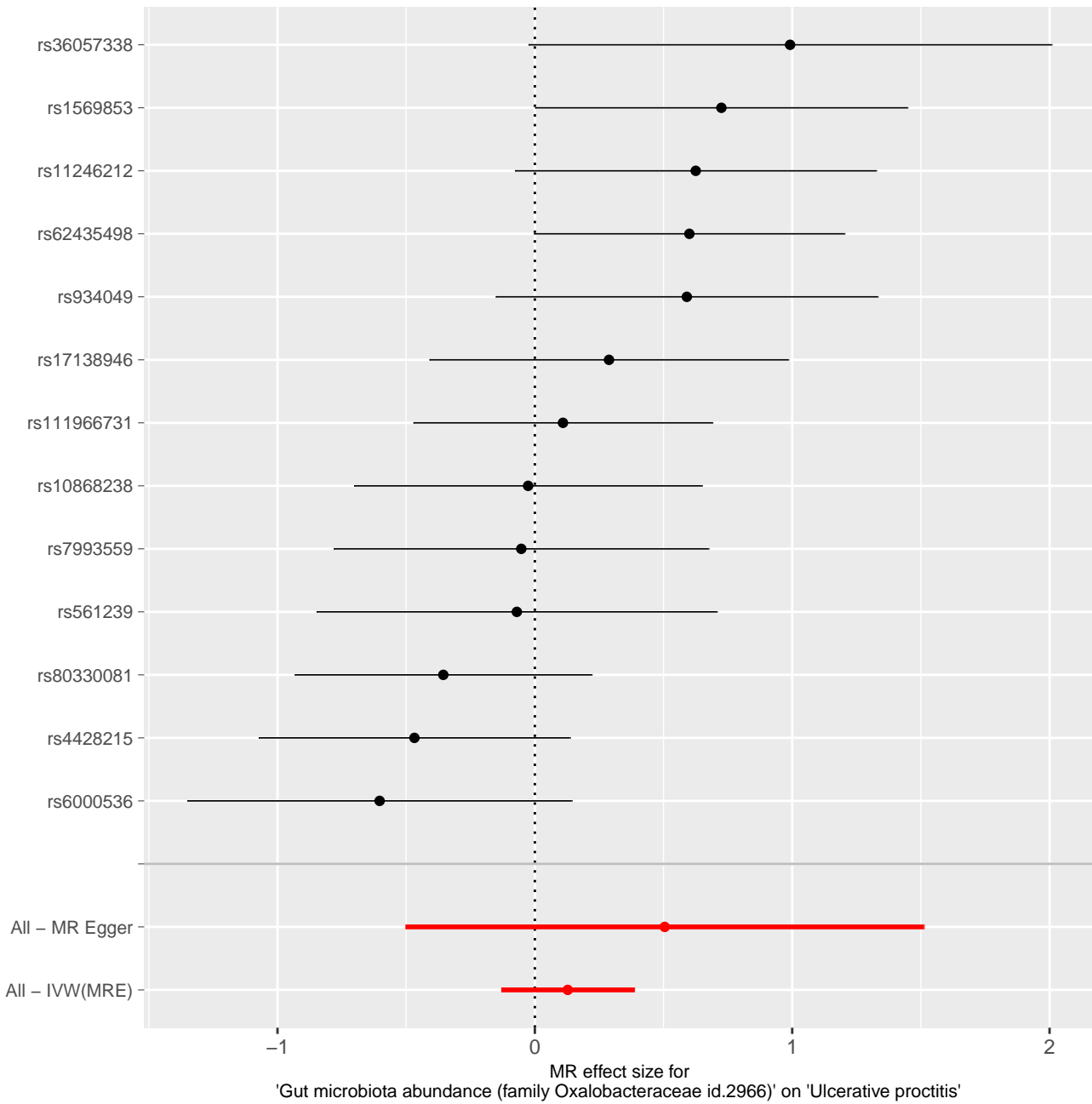


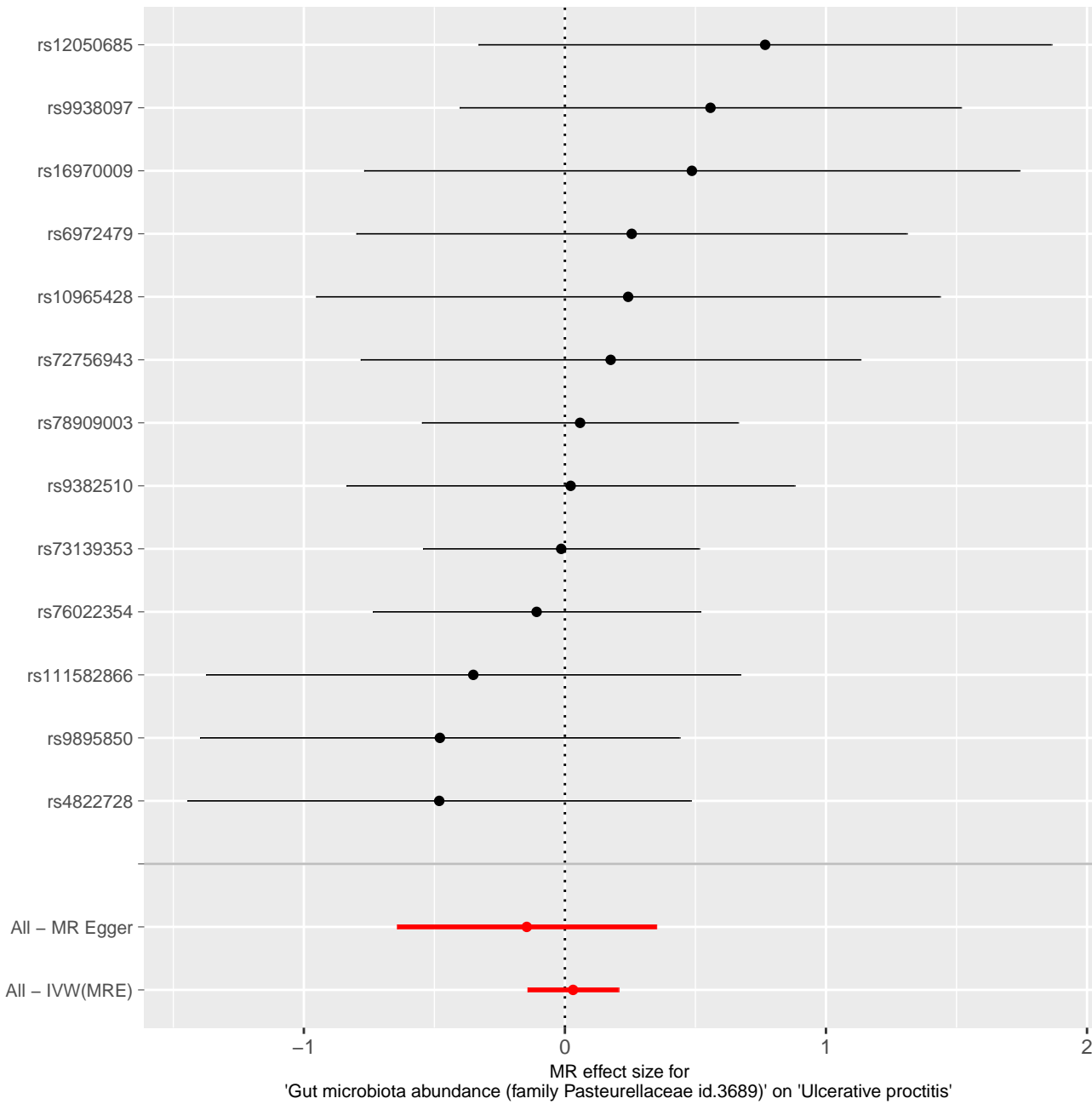


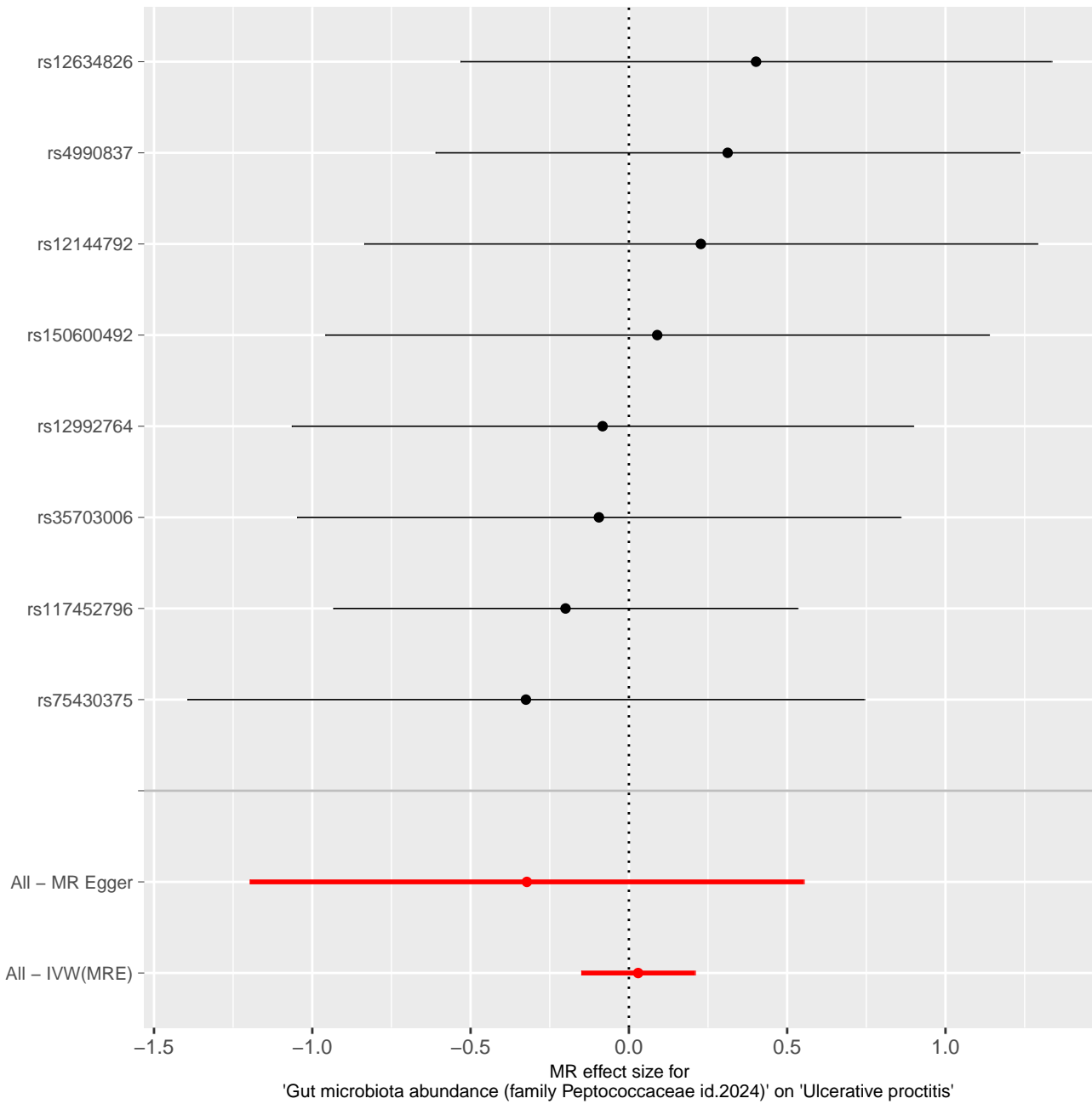


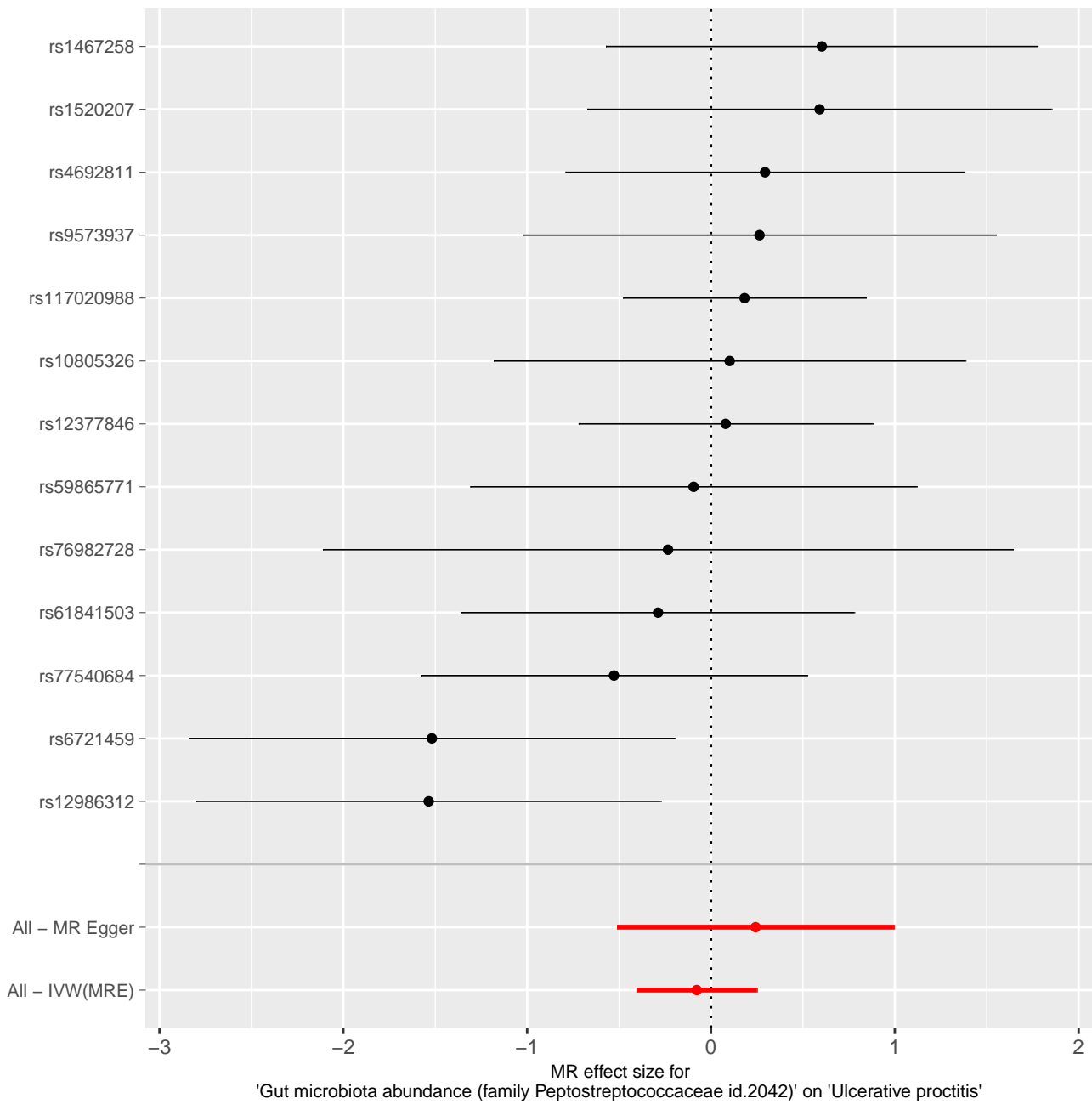


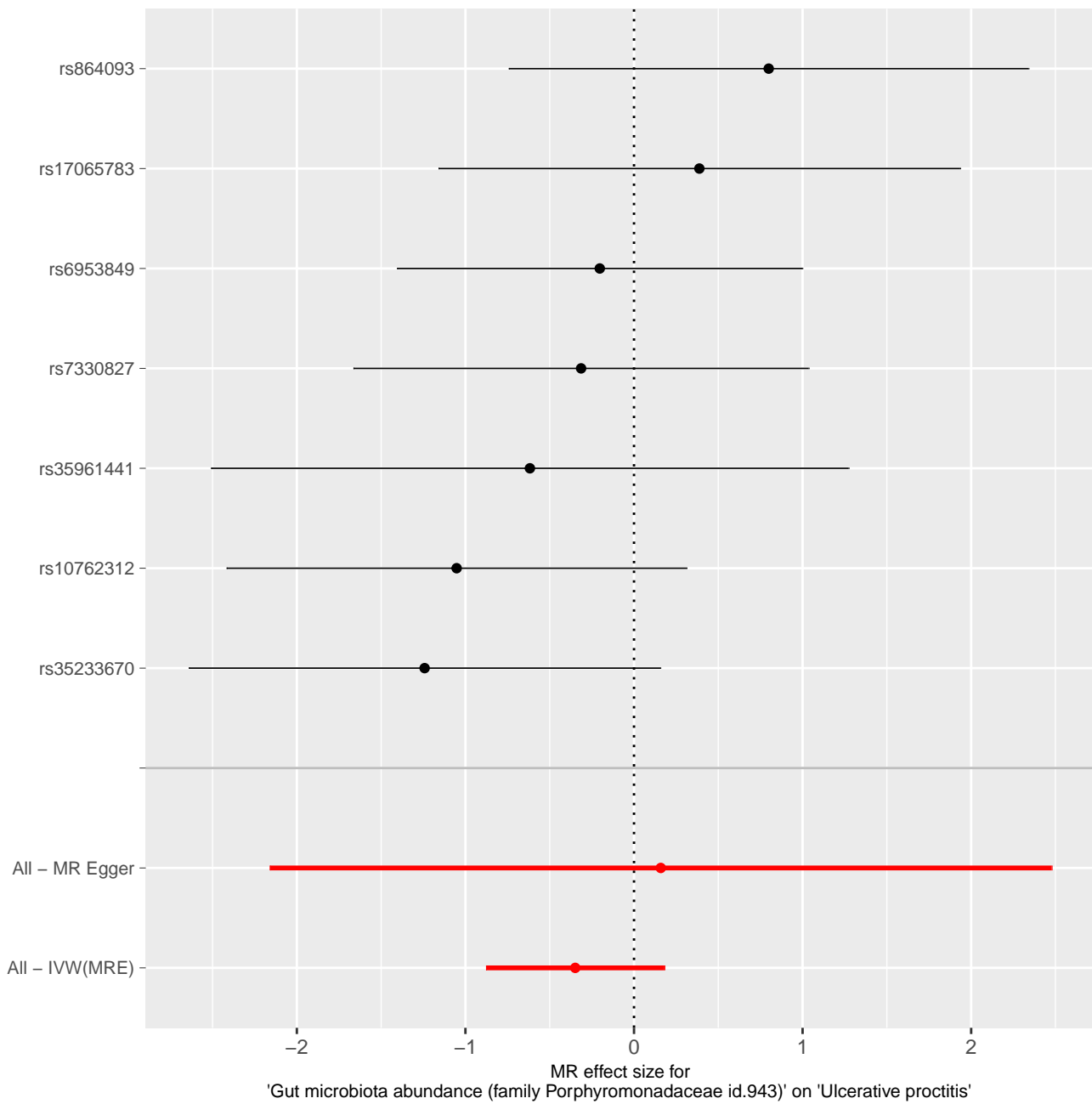




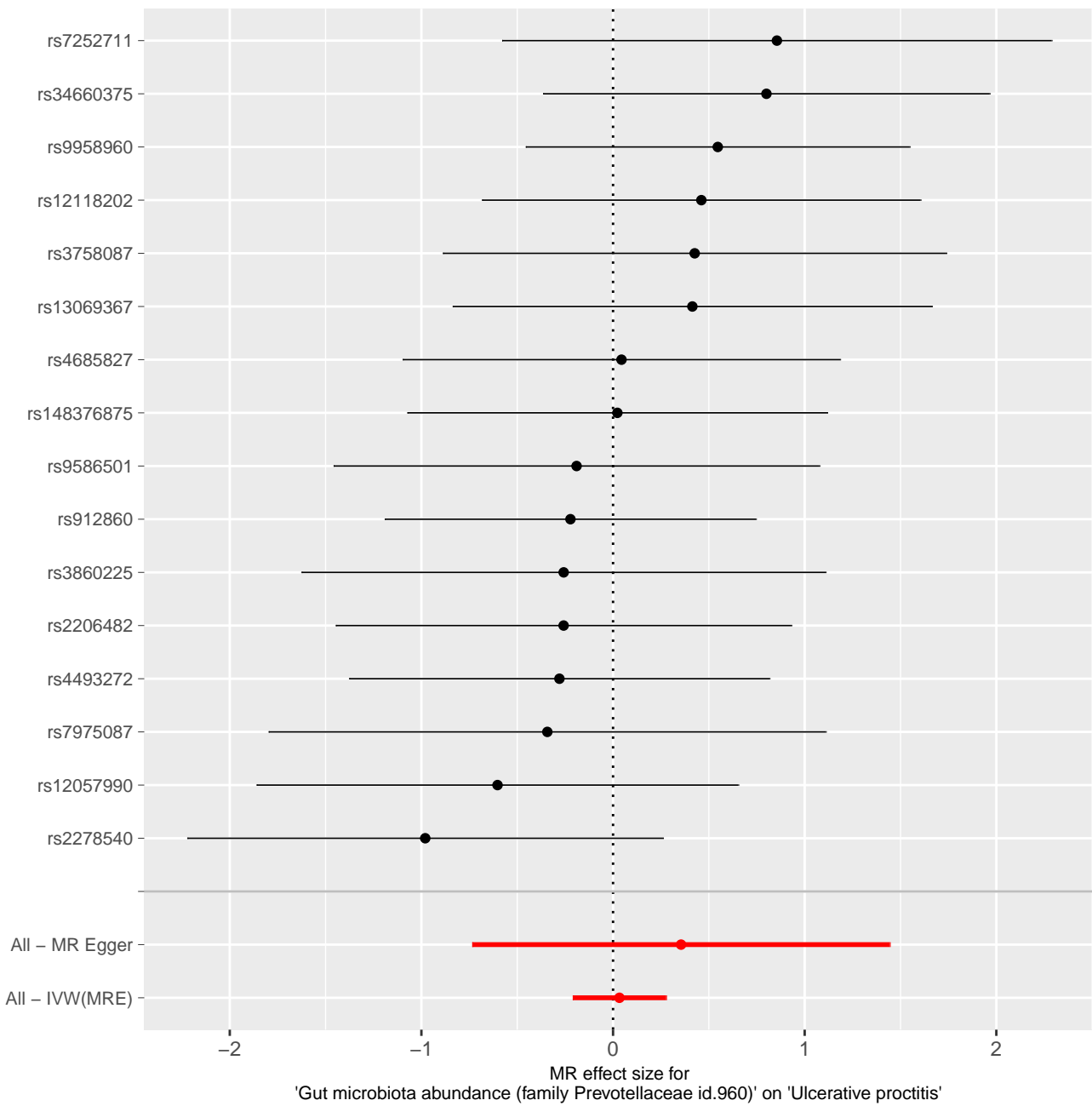


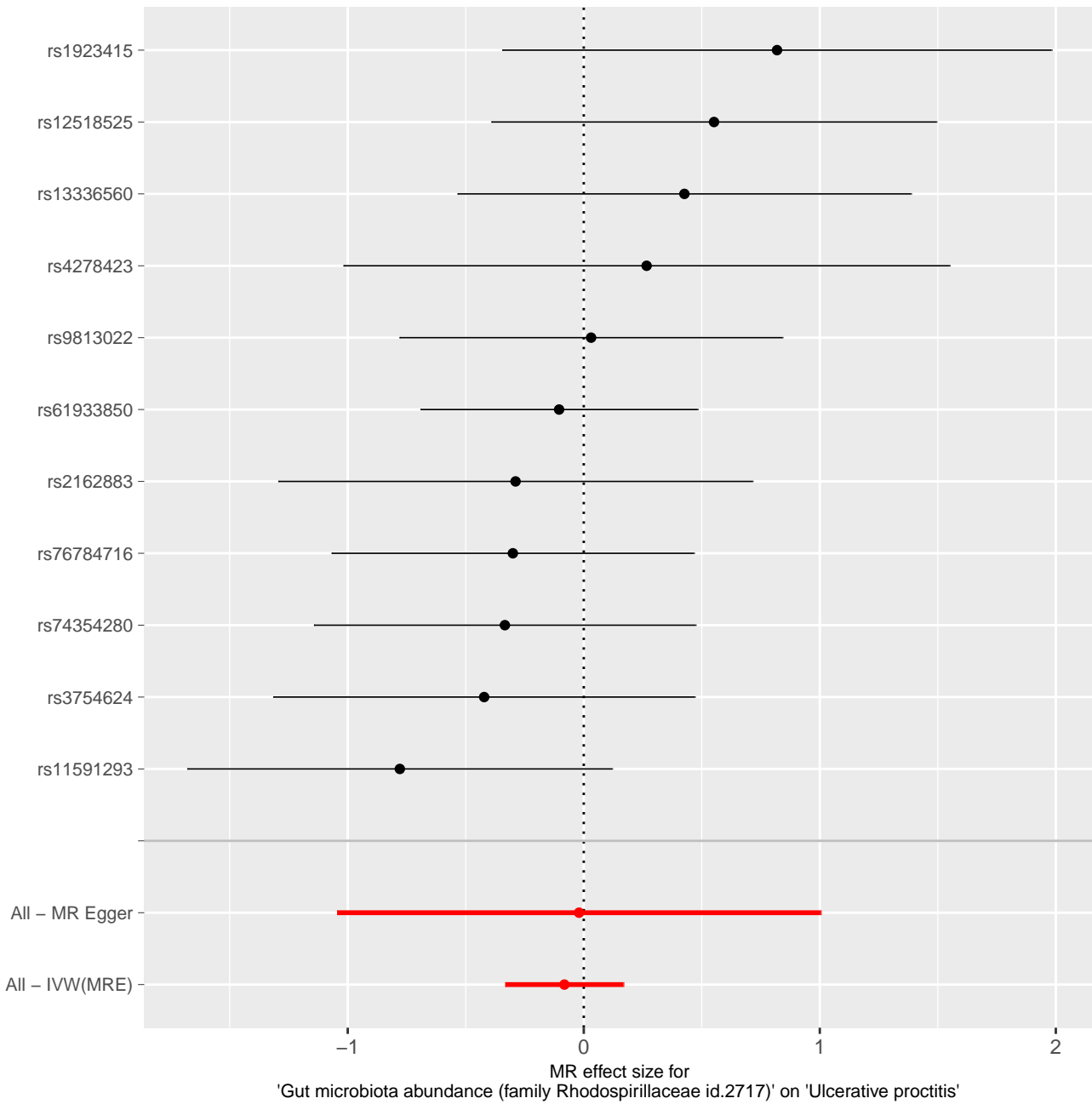


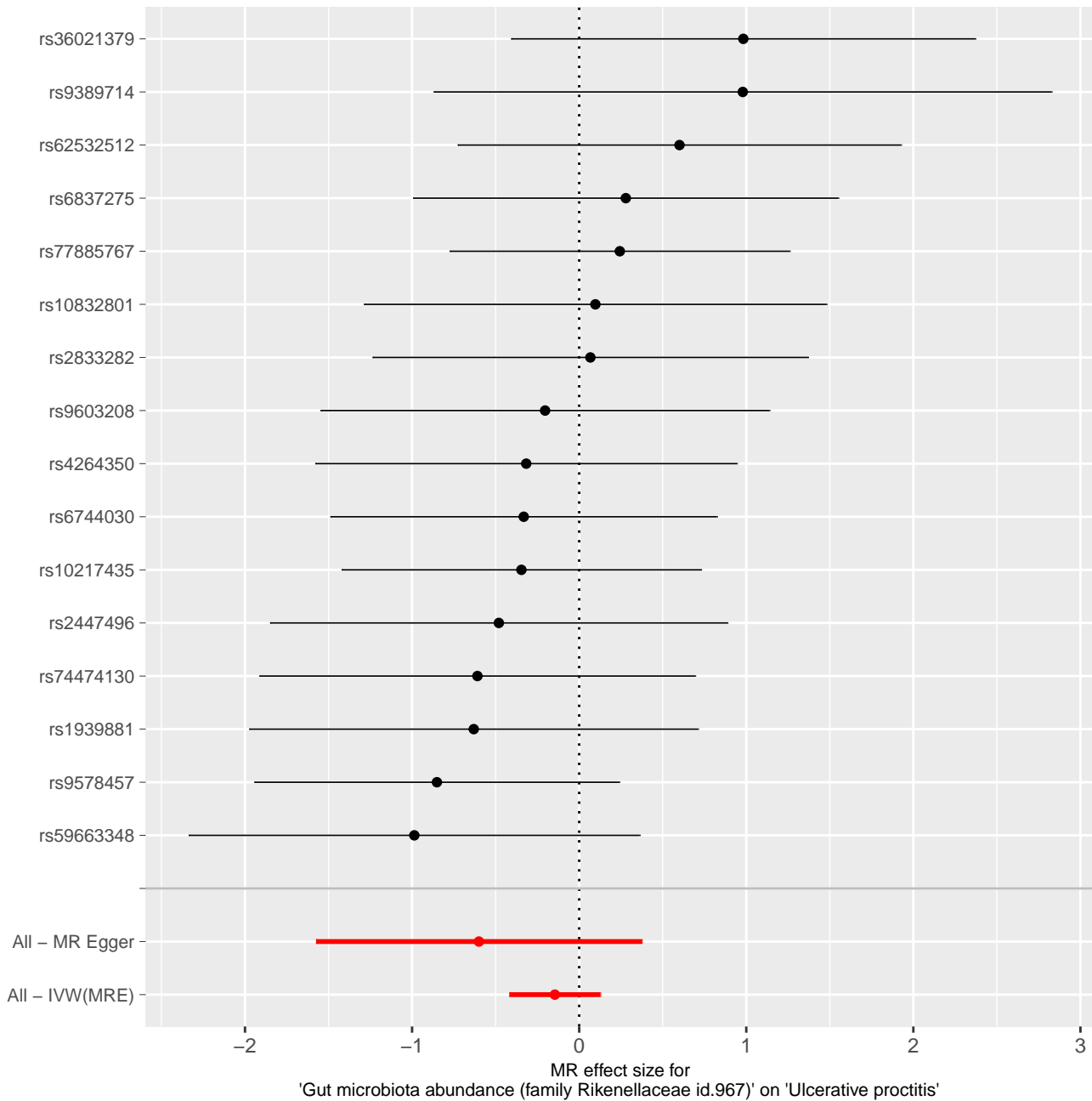


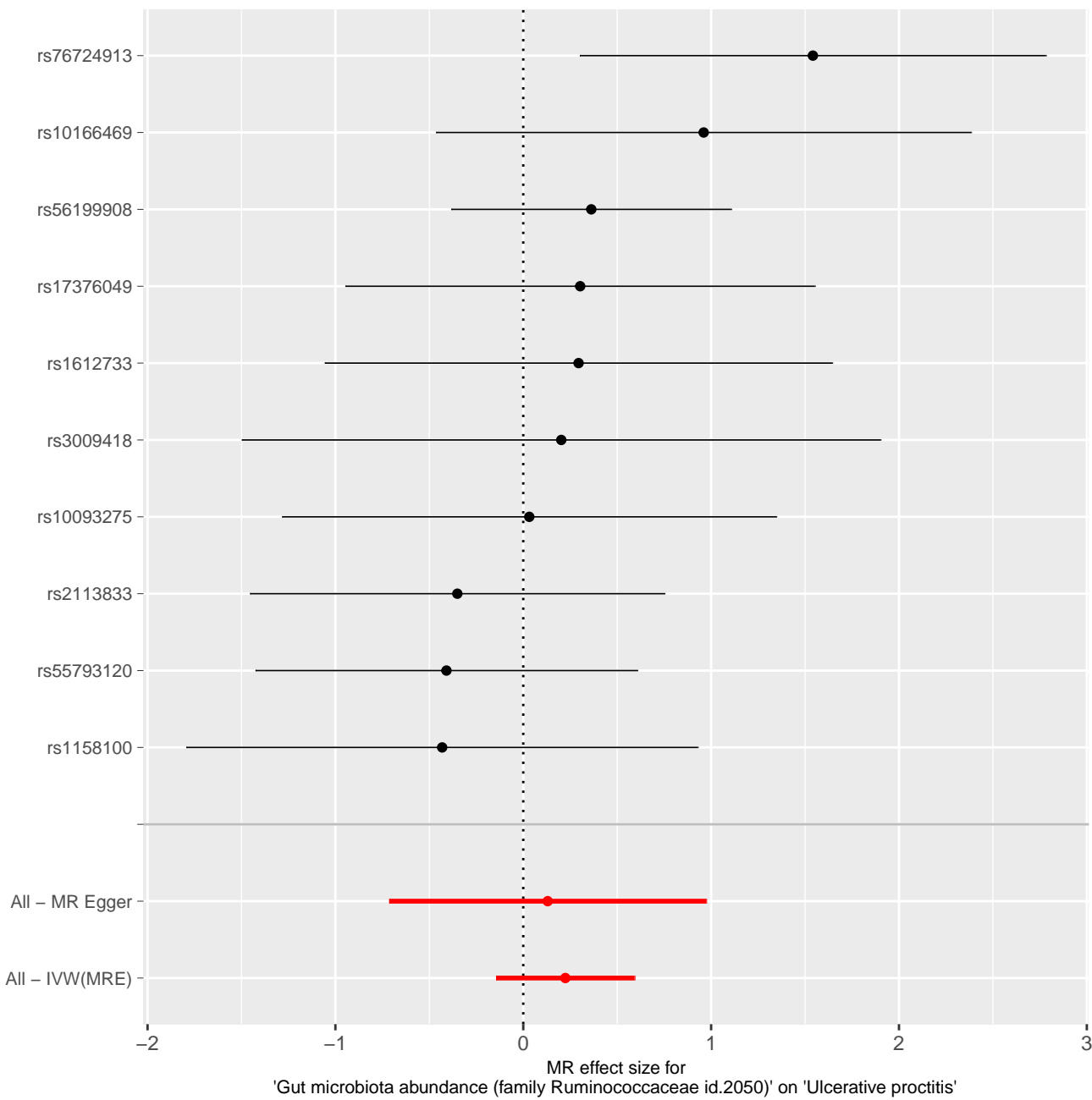


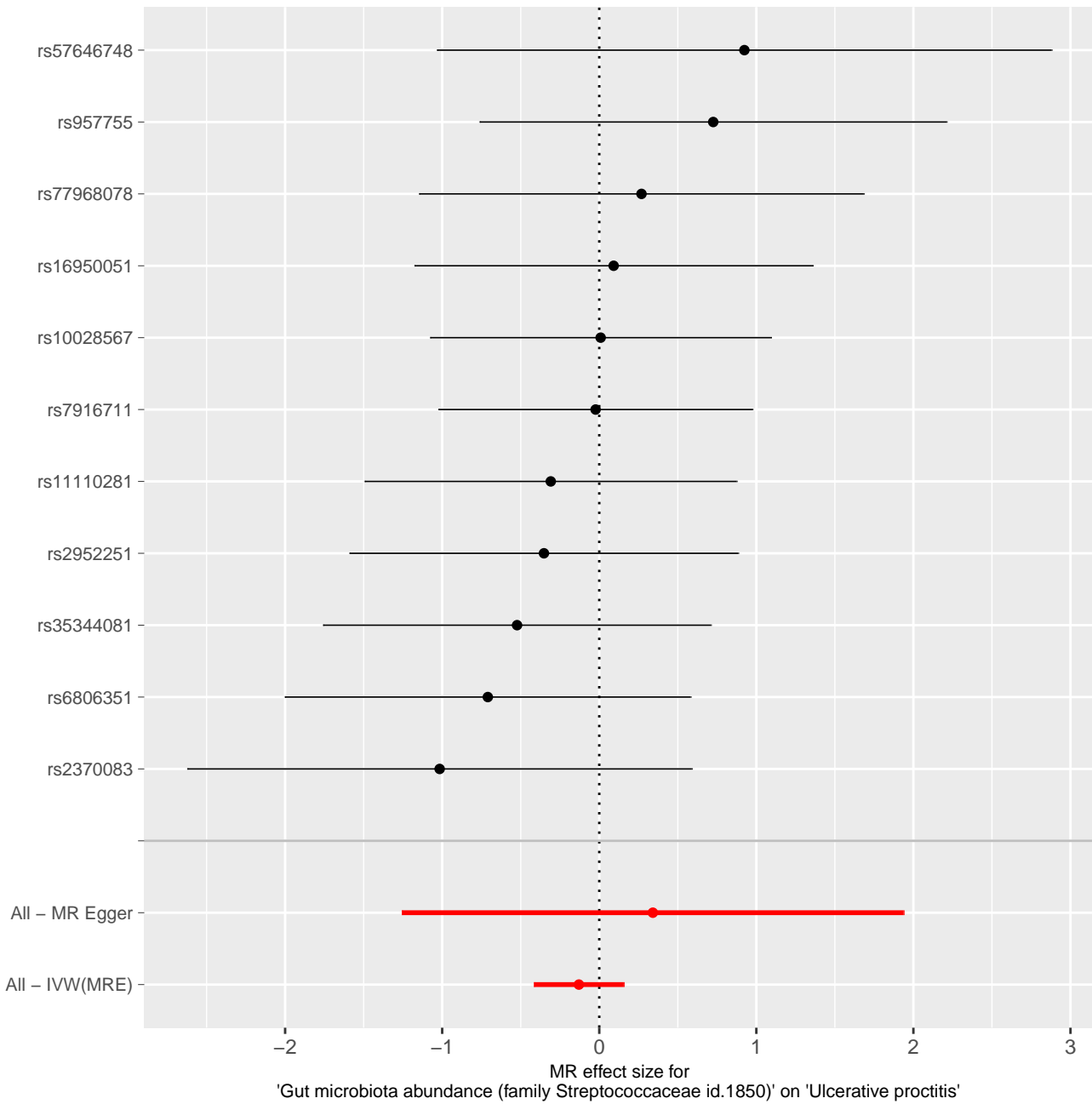
Batch 1375 : Gut microbiota abundance (family Prevotellaceae id.960) on Ulcerative proctitis

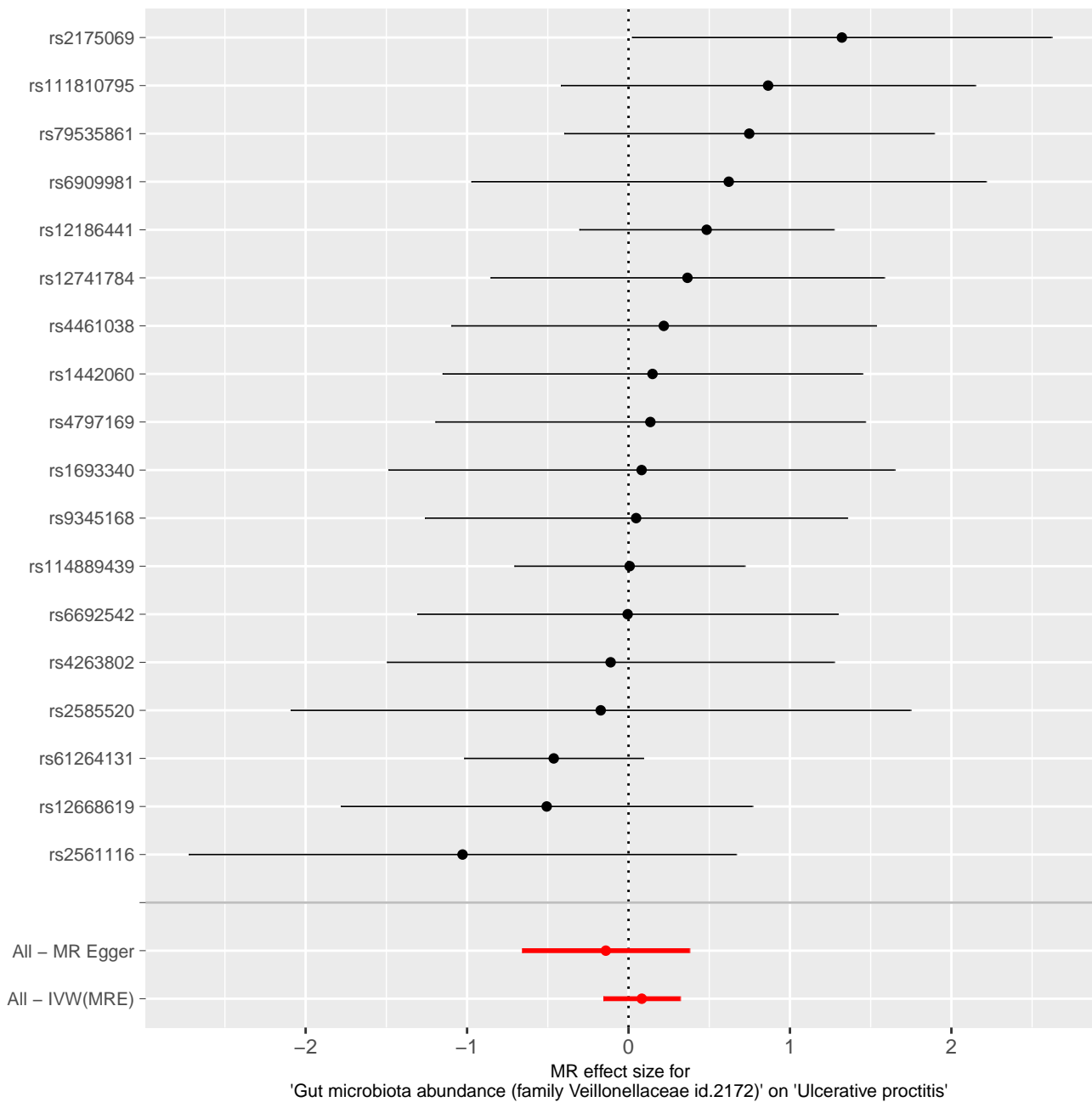


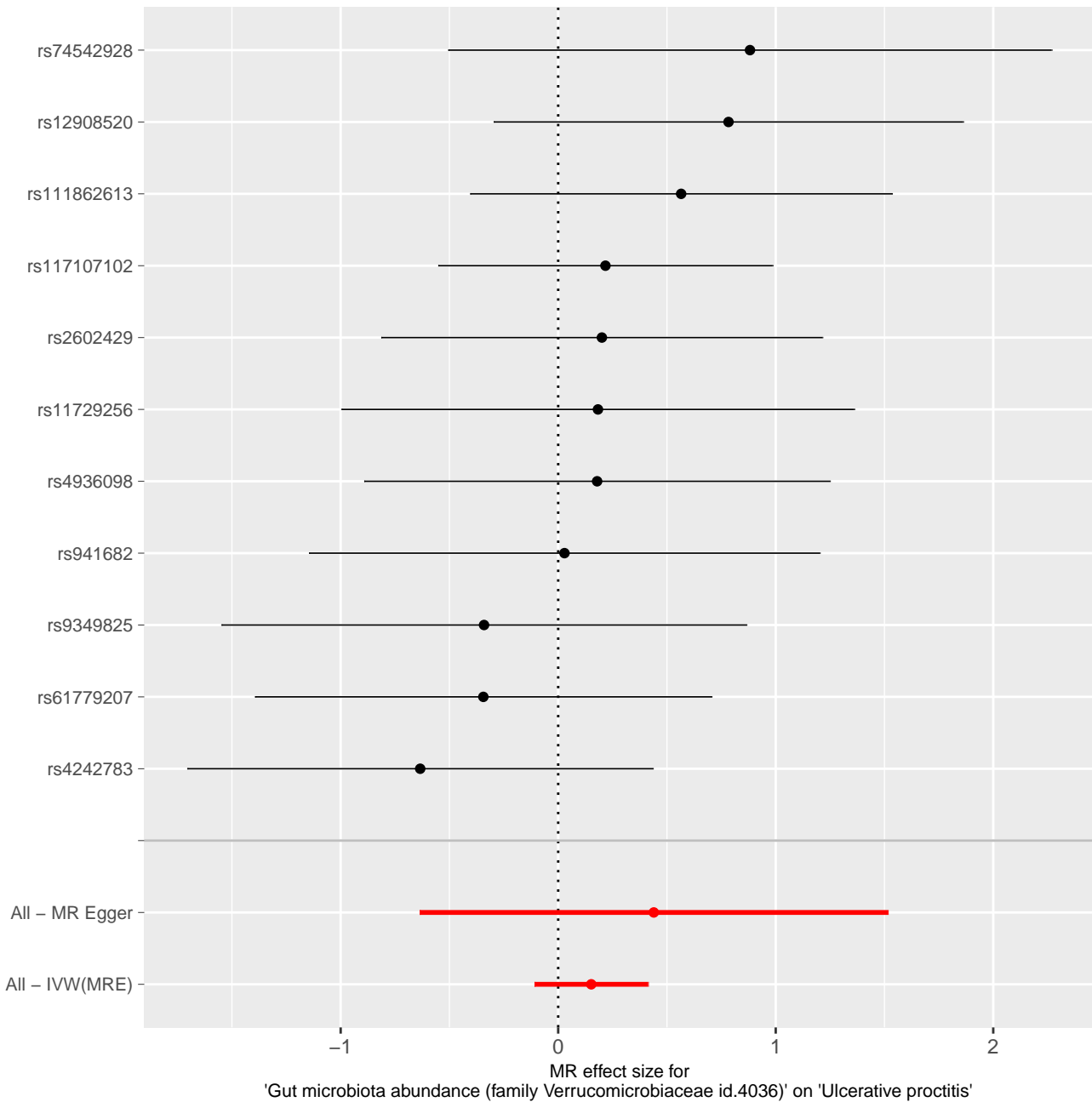




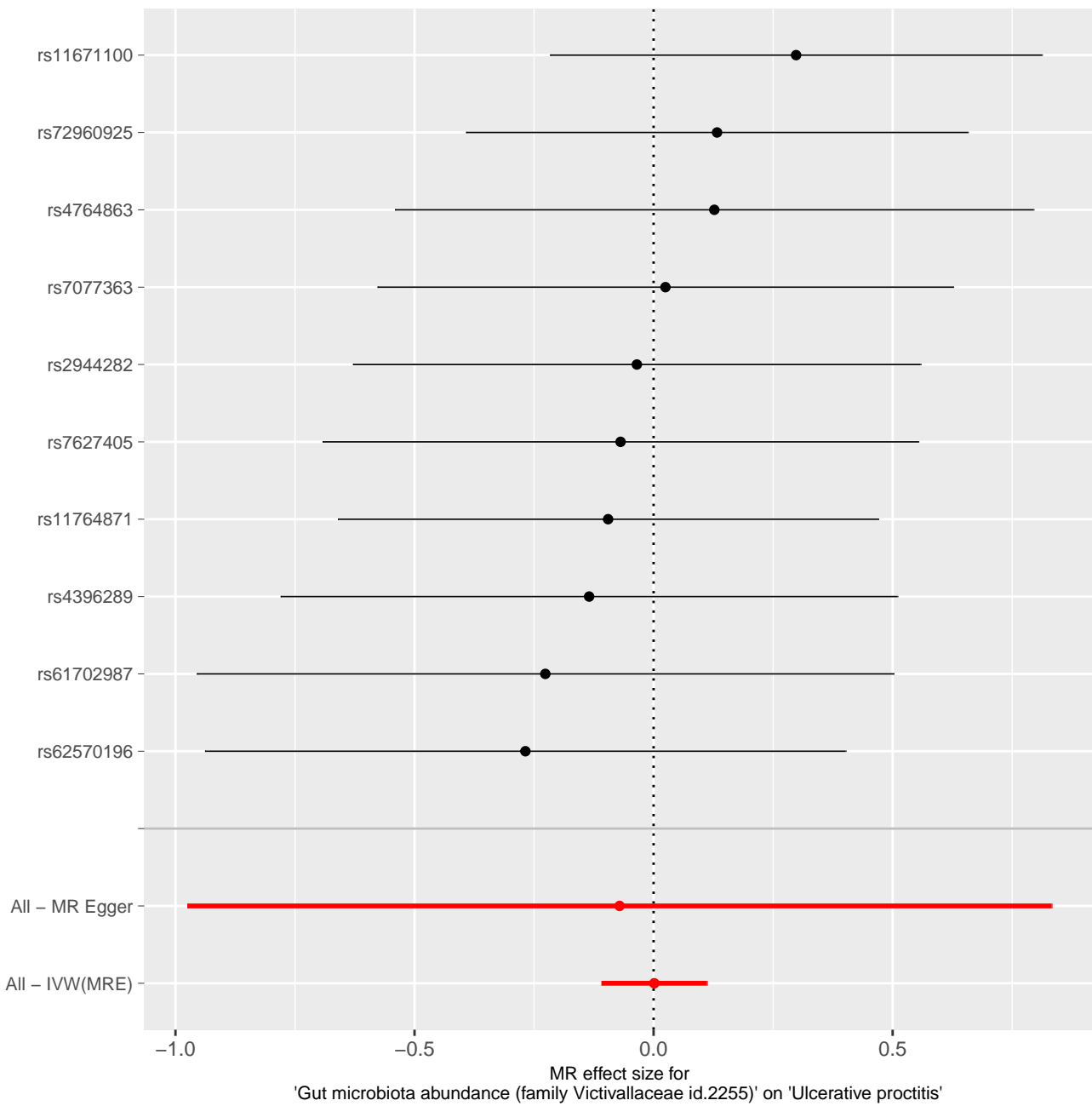


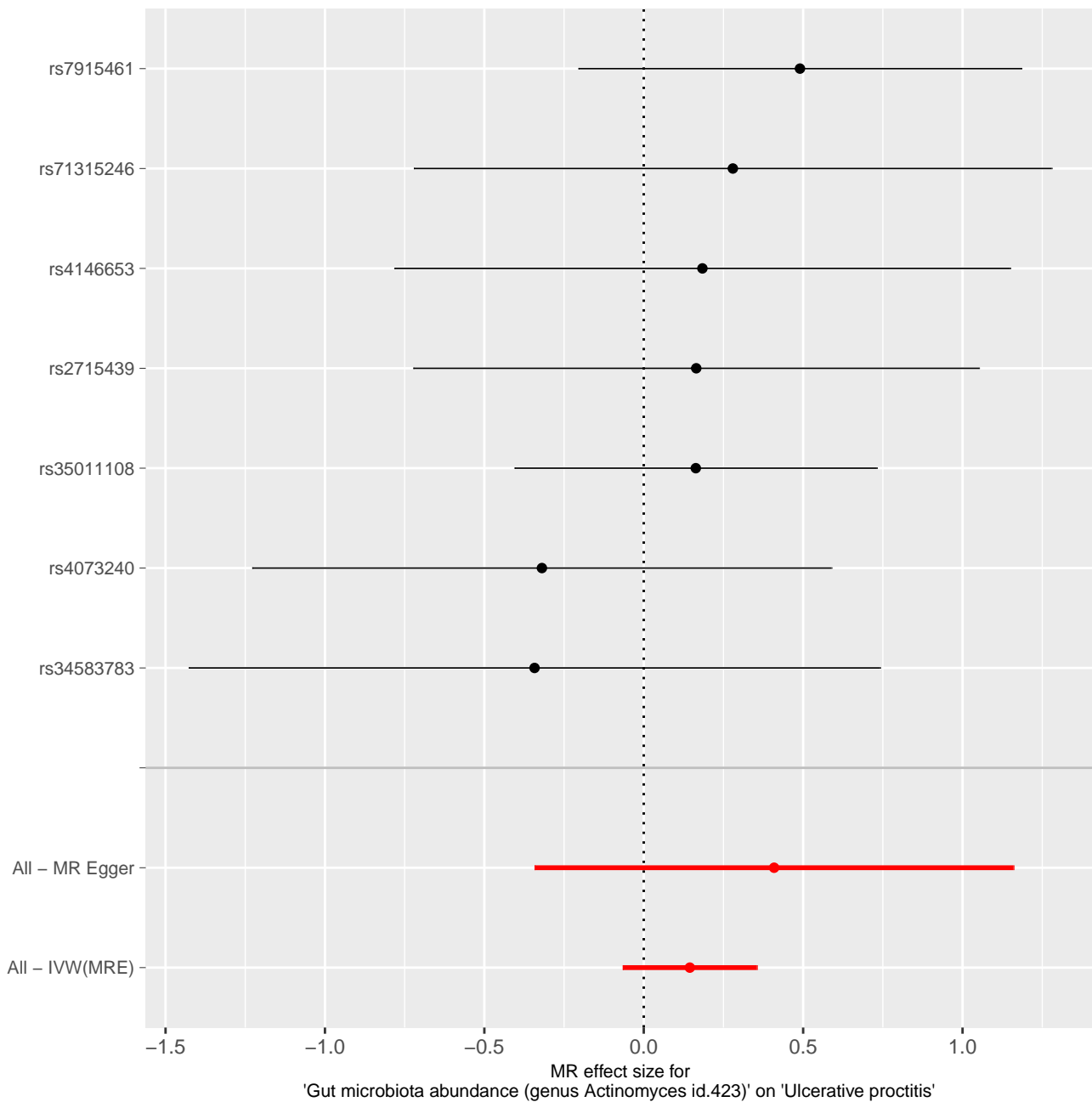


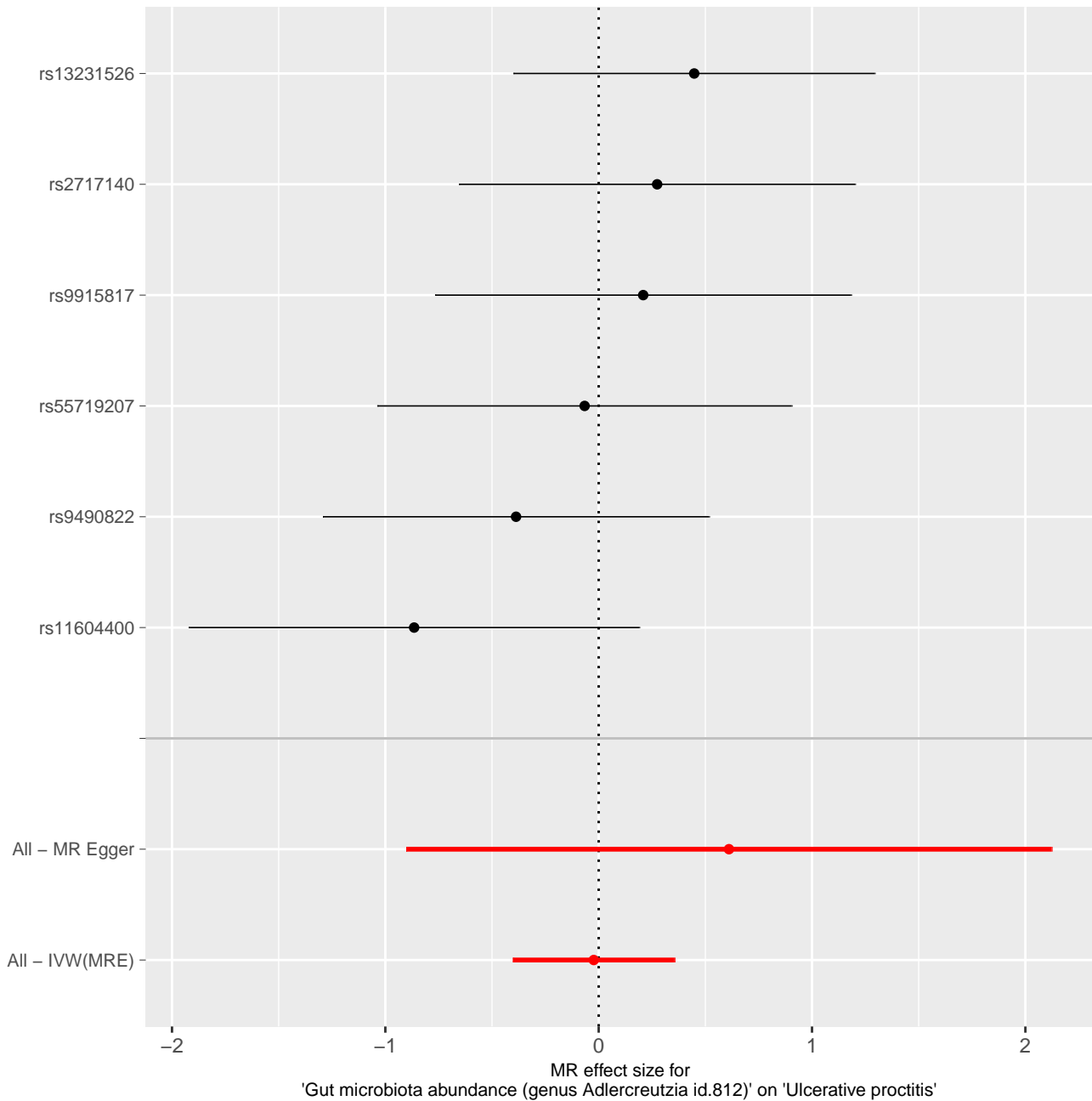




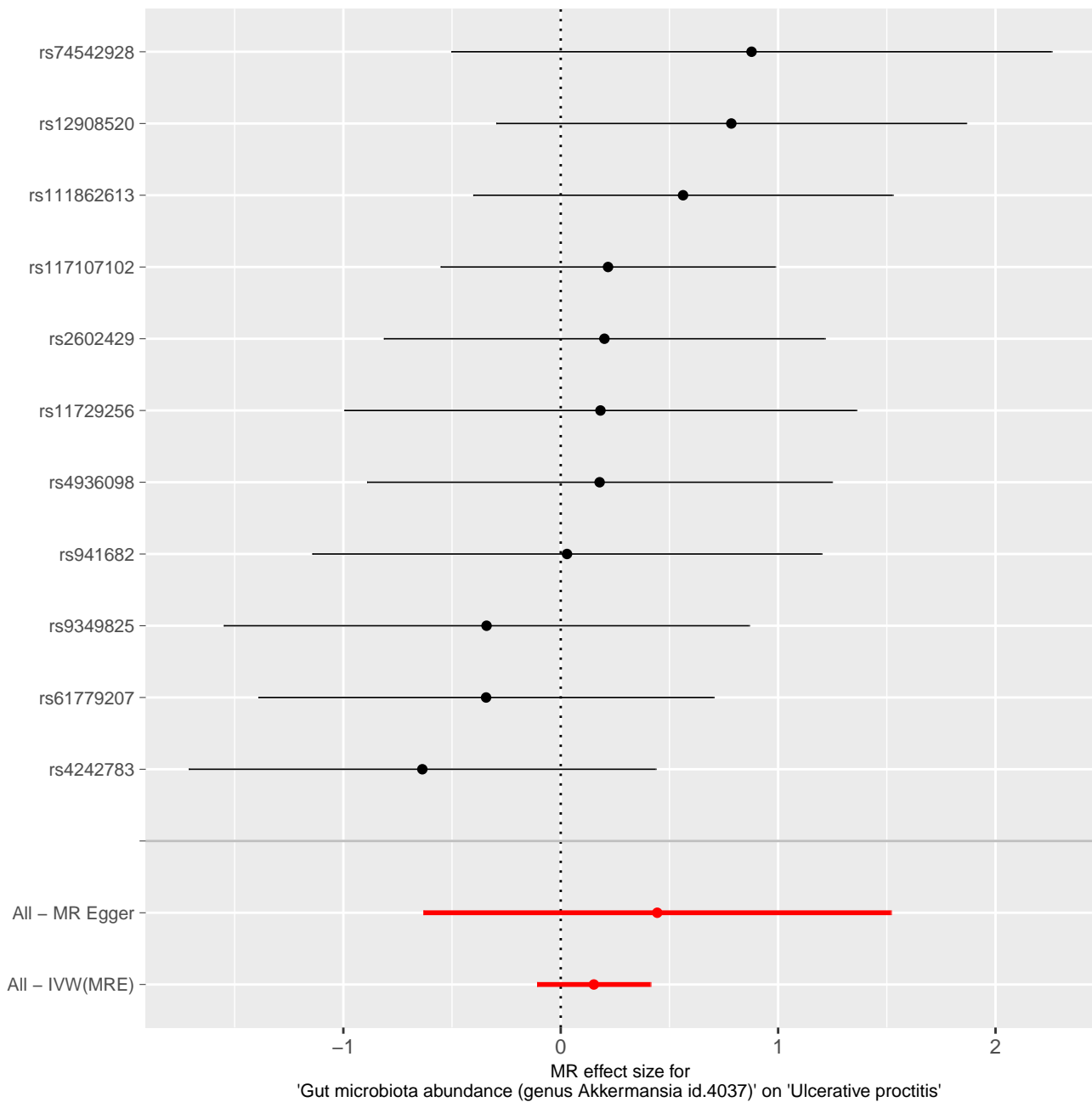
Batch 1382 : Gut microbiota abundance (family Victivallaceae id.2255) on Ulcerative proctitis

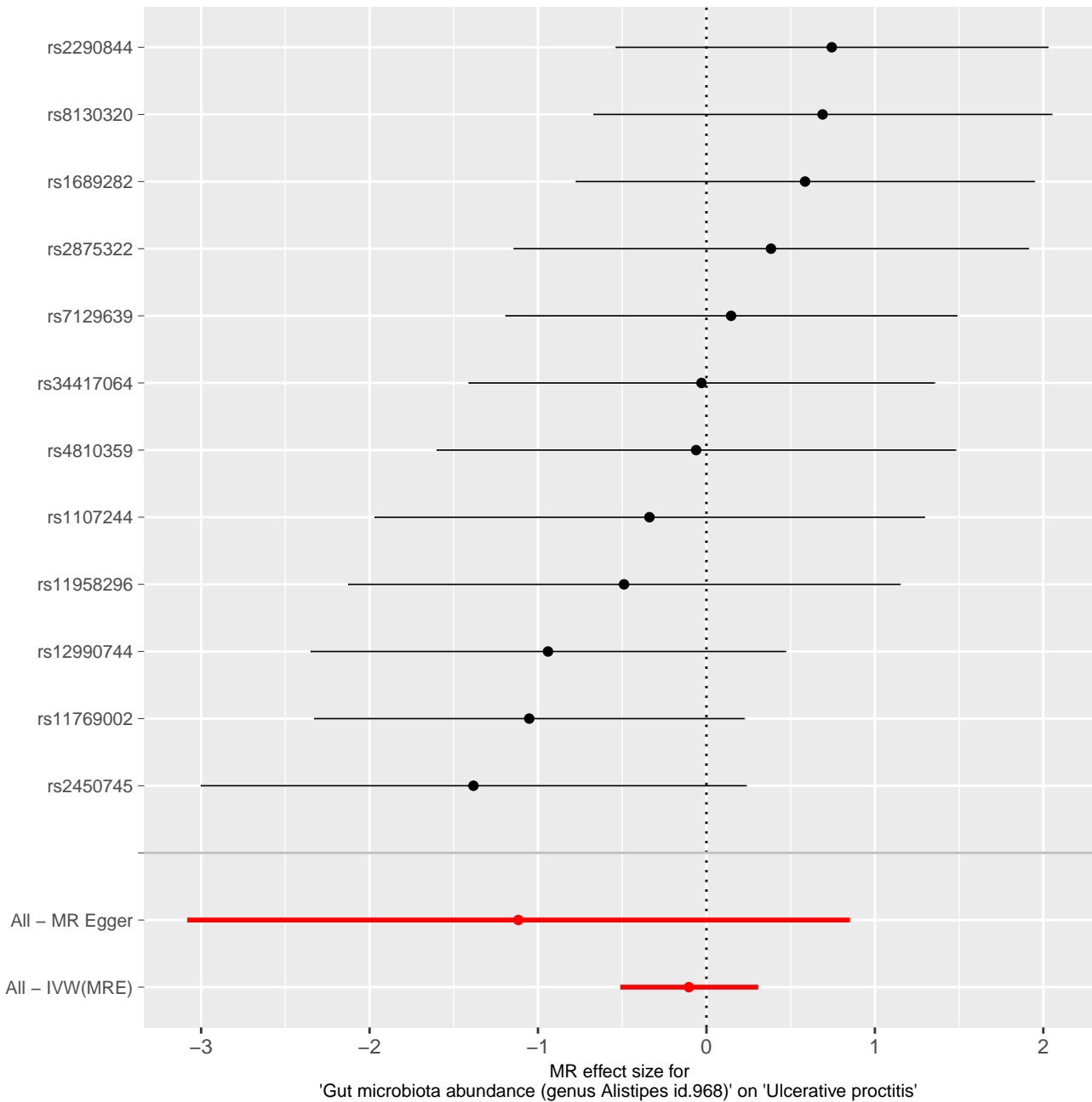


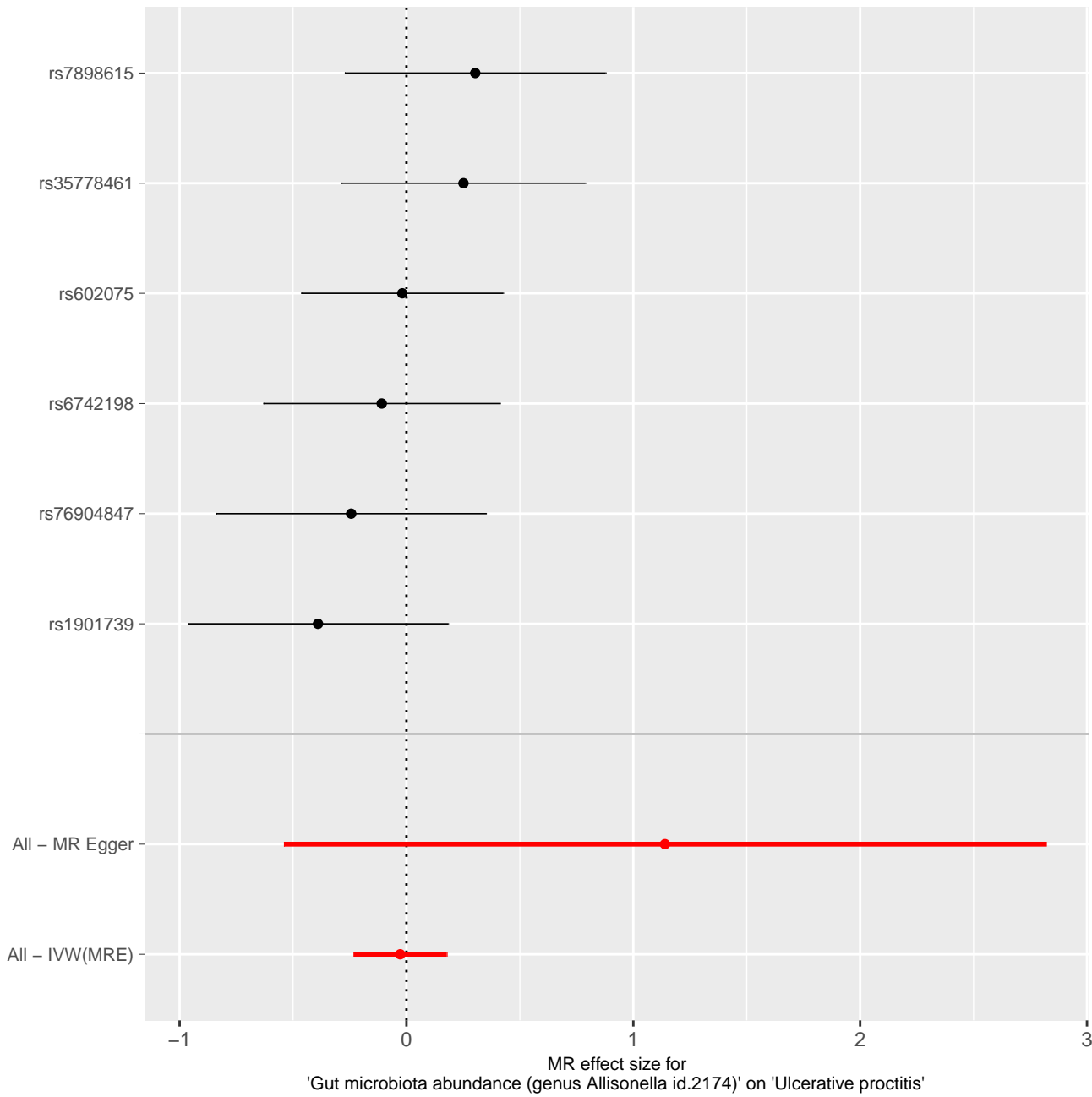


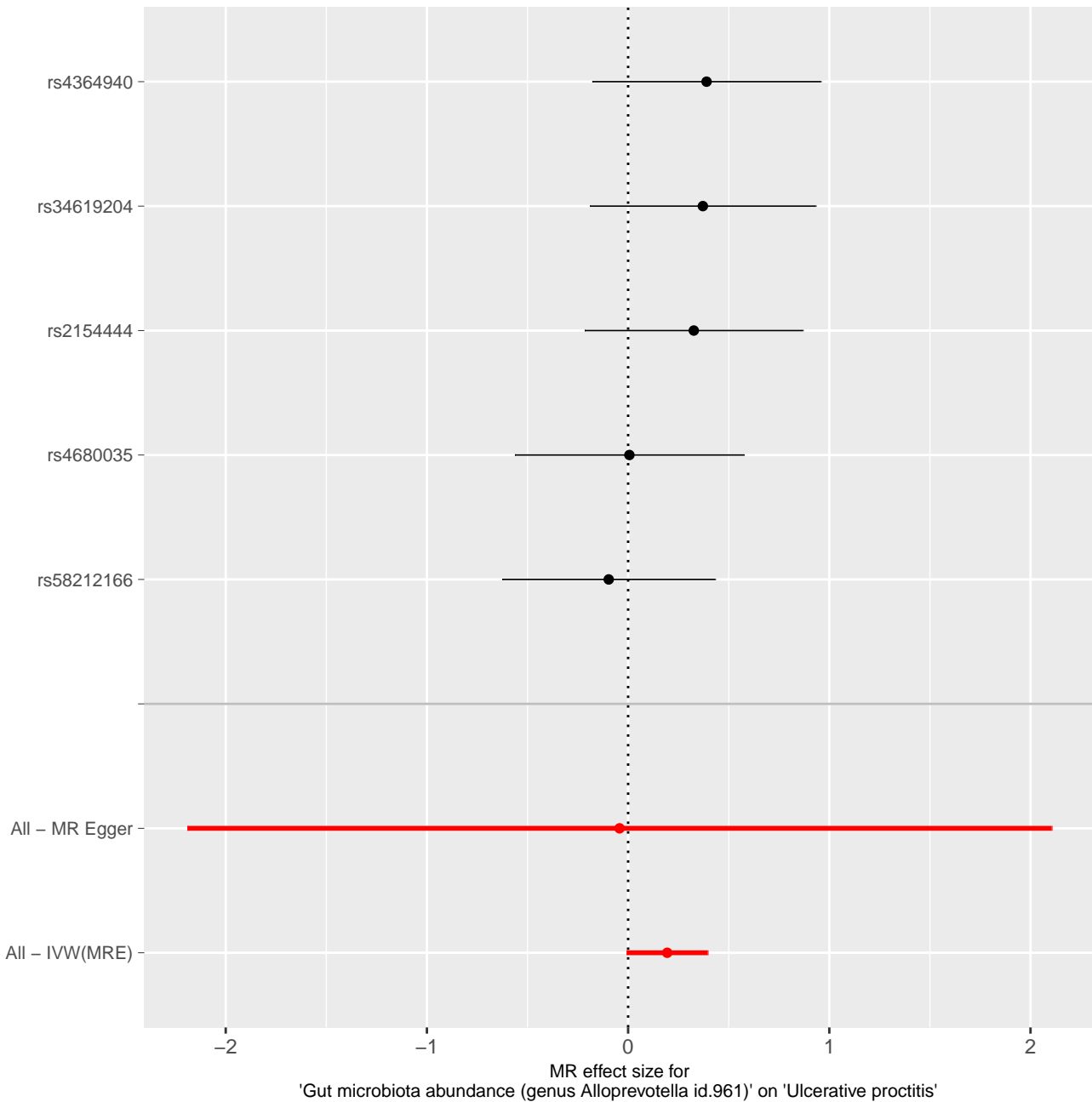


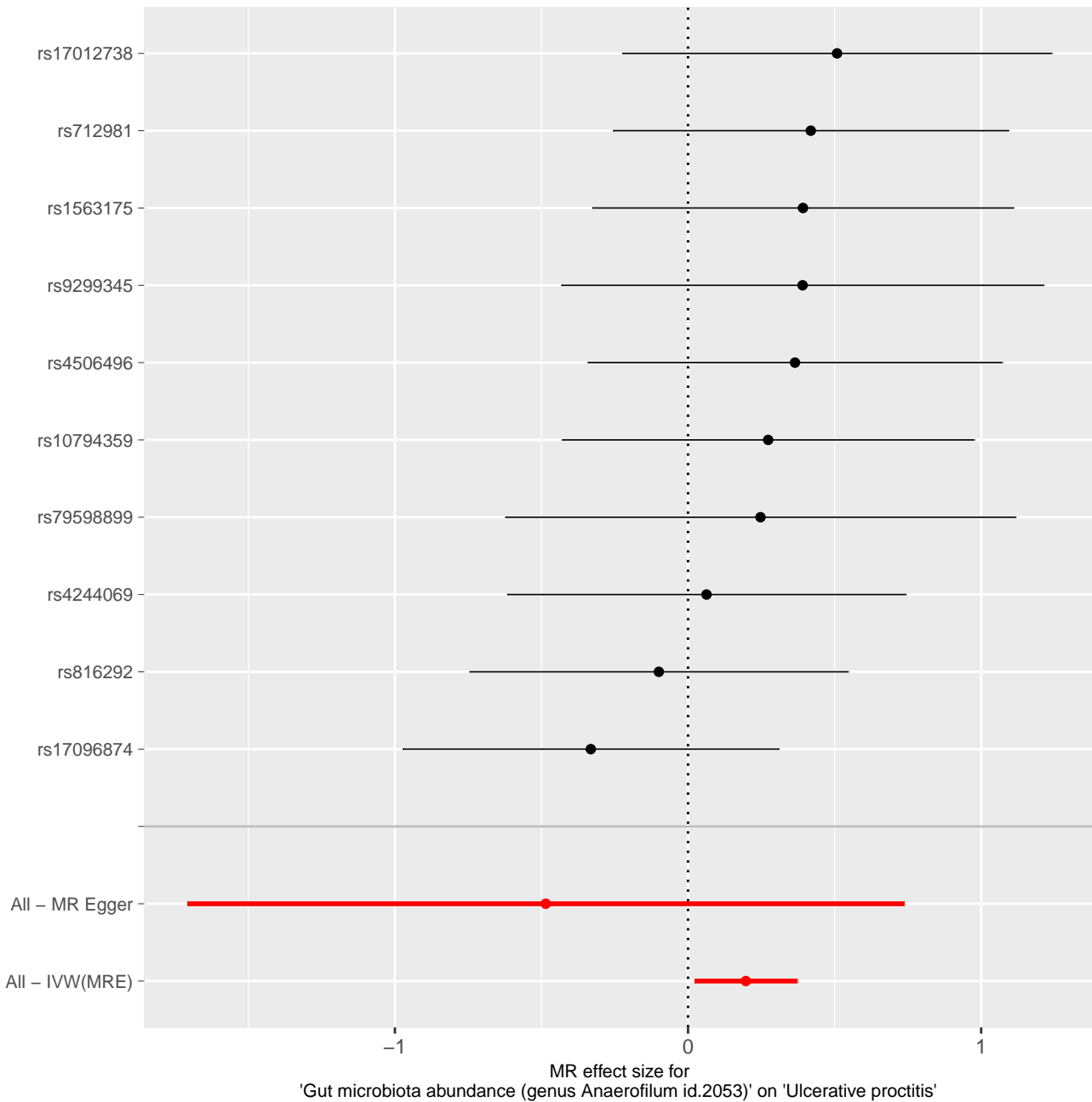
Batch 1385 : Gut microbiota abundance (genus Akkermansia id.4037) on Ulcerative proctitis



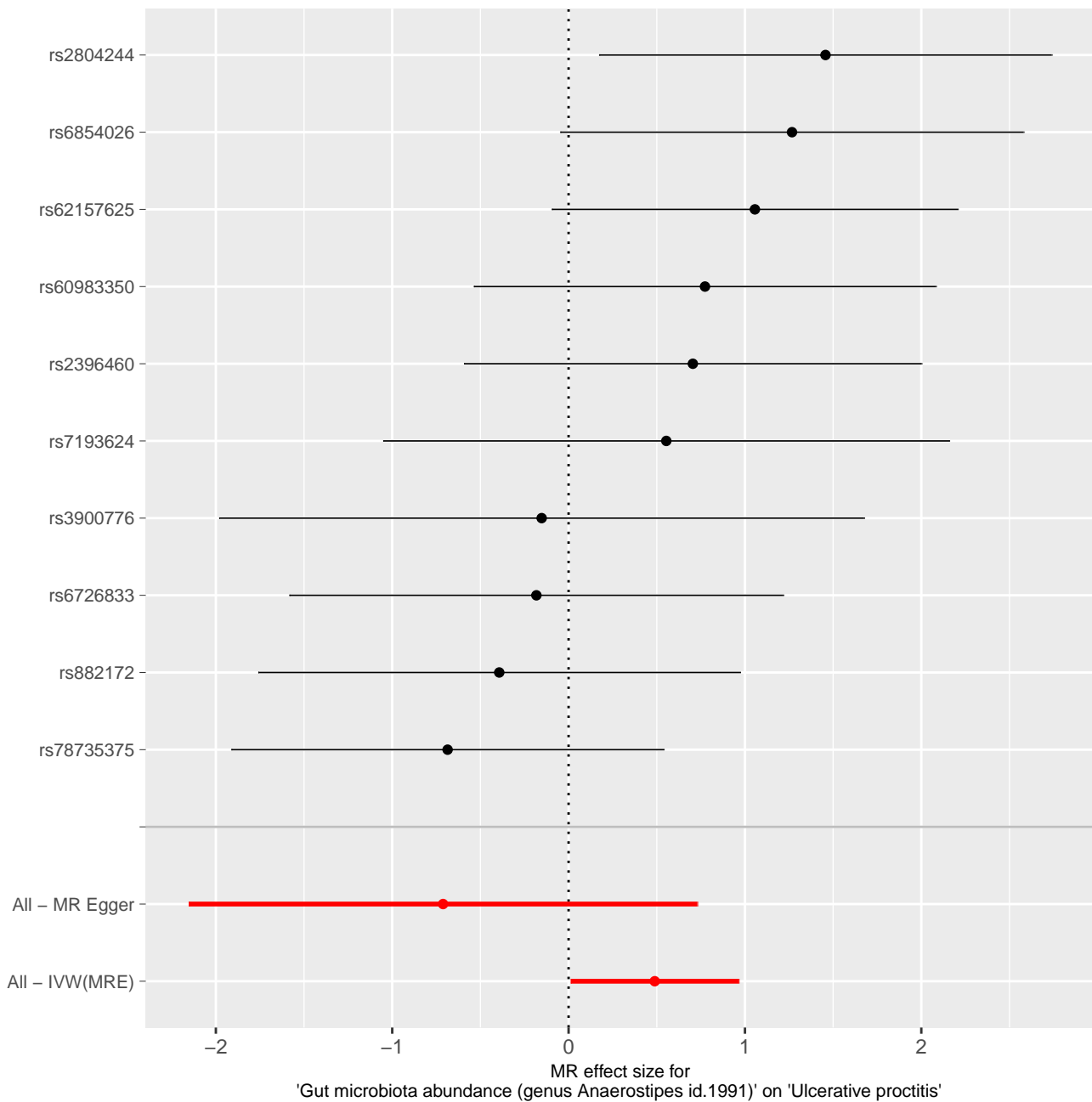


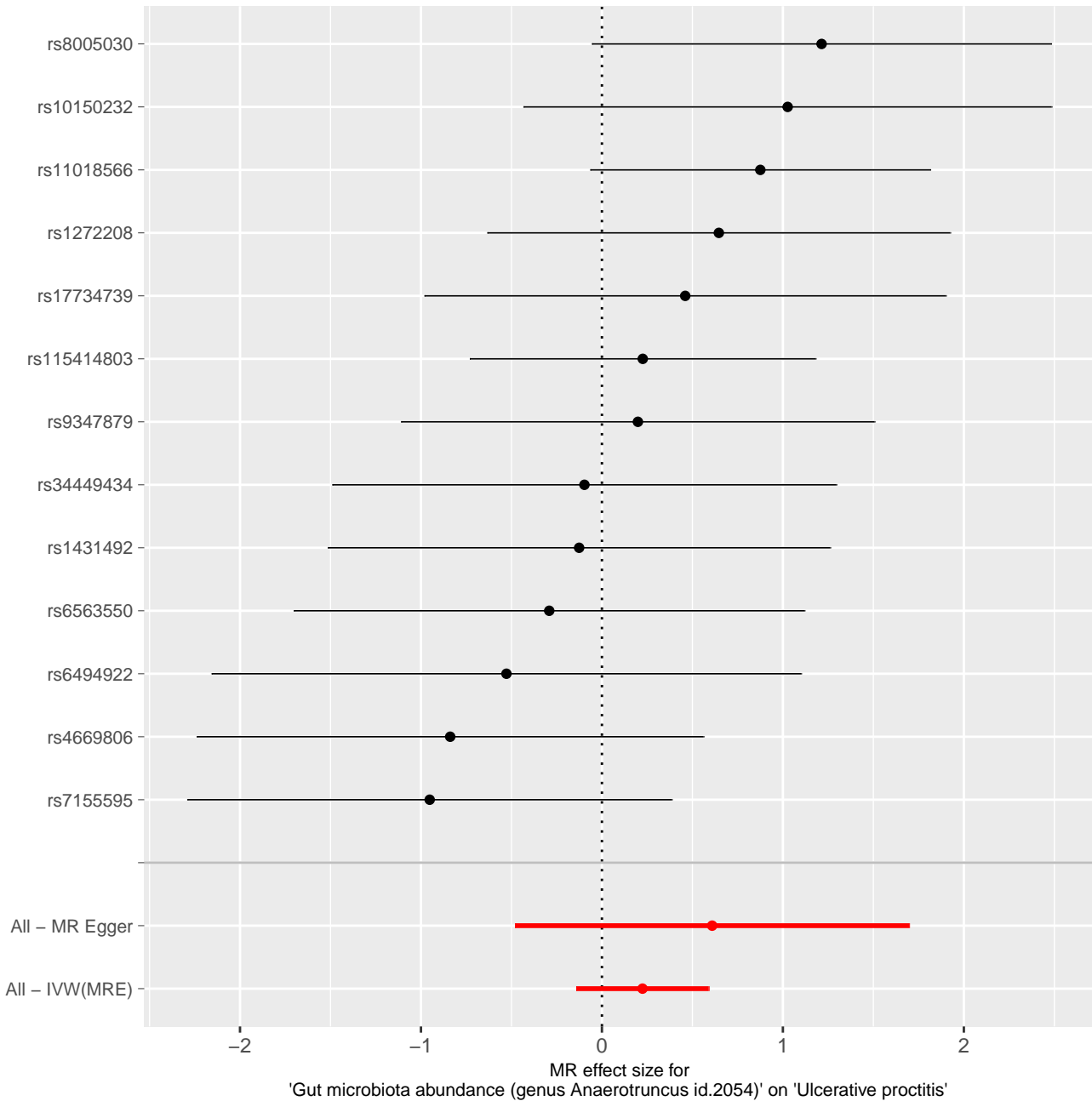


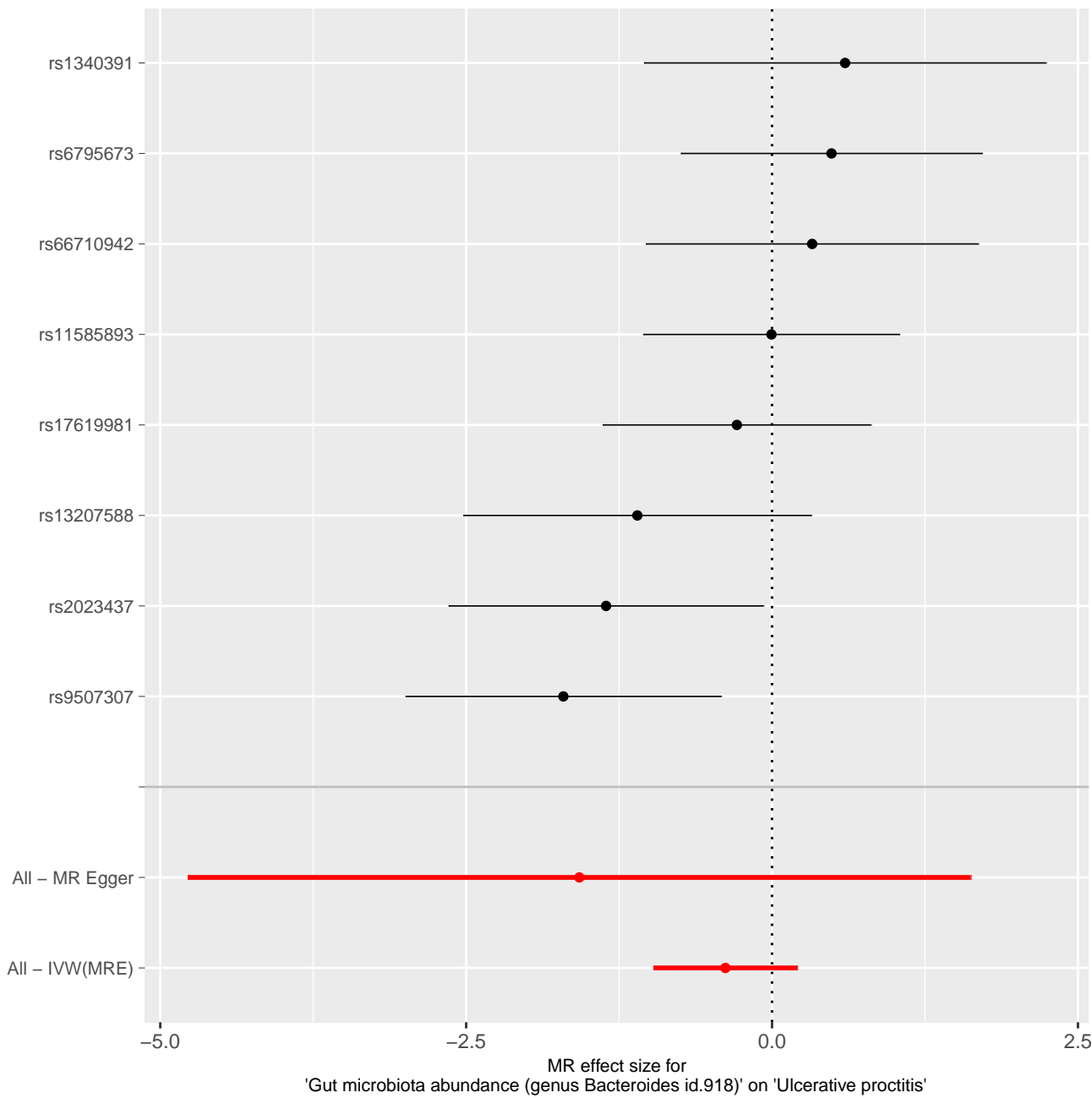


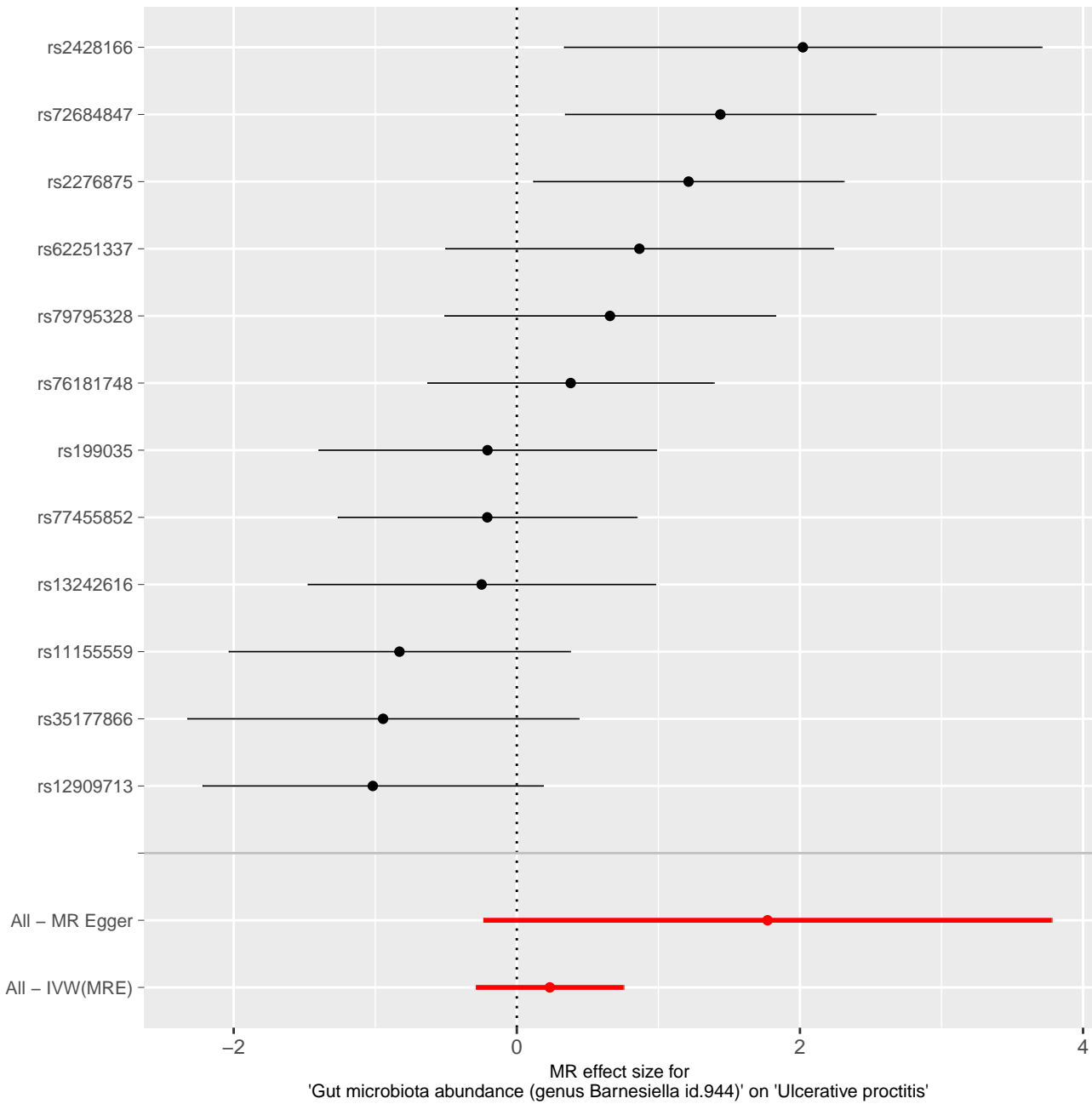


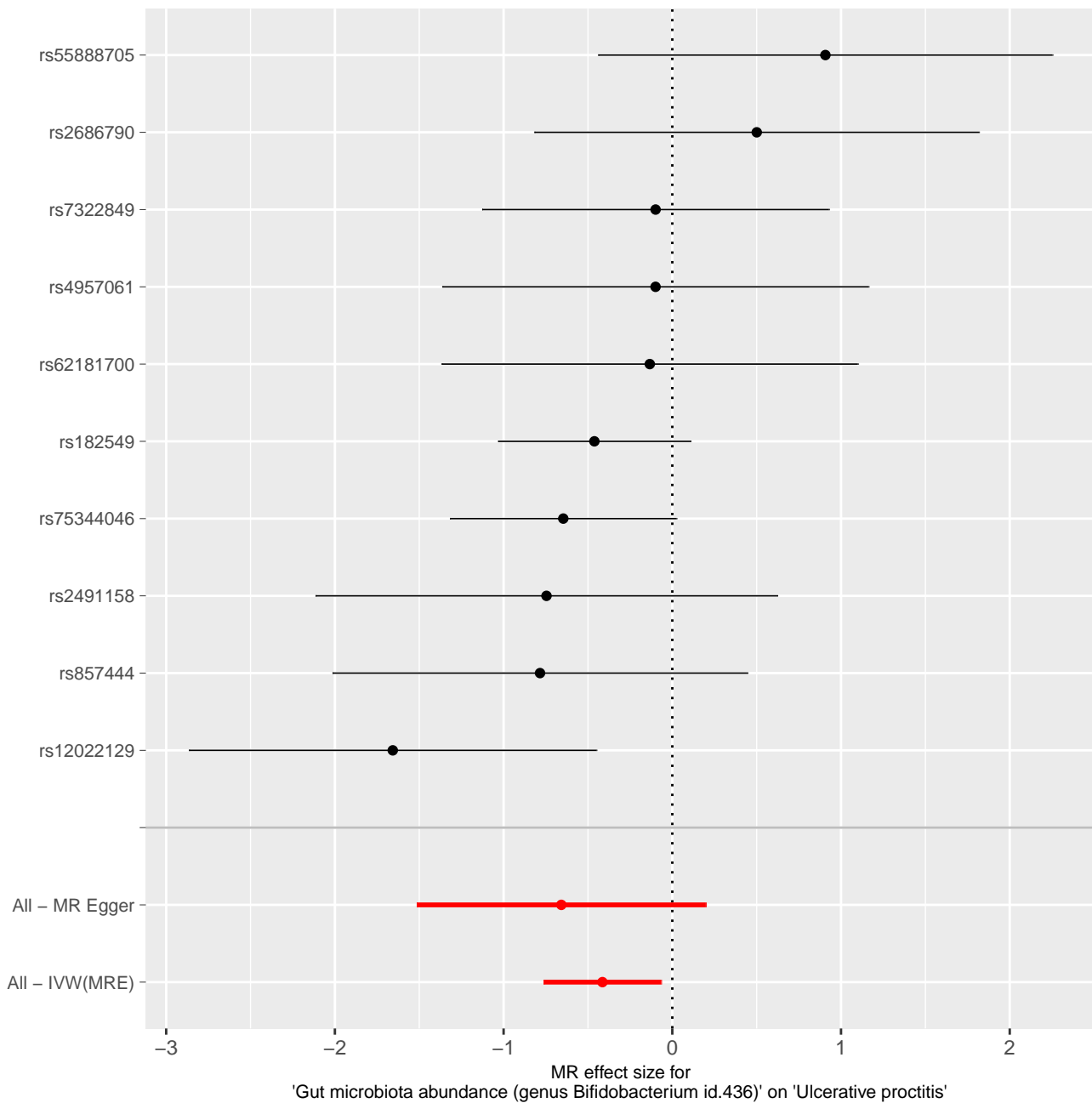
Batch 1390 : Gut microbiota abundance (genus Anaerostipes id.1991) on Ulcerative proctitis



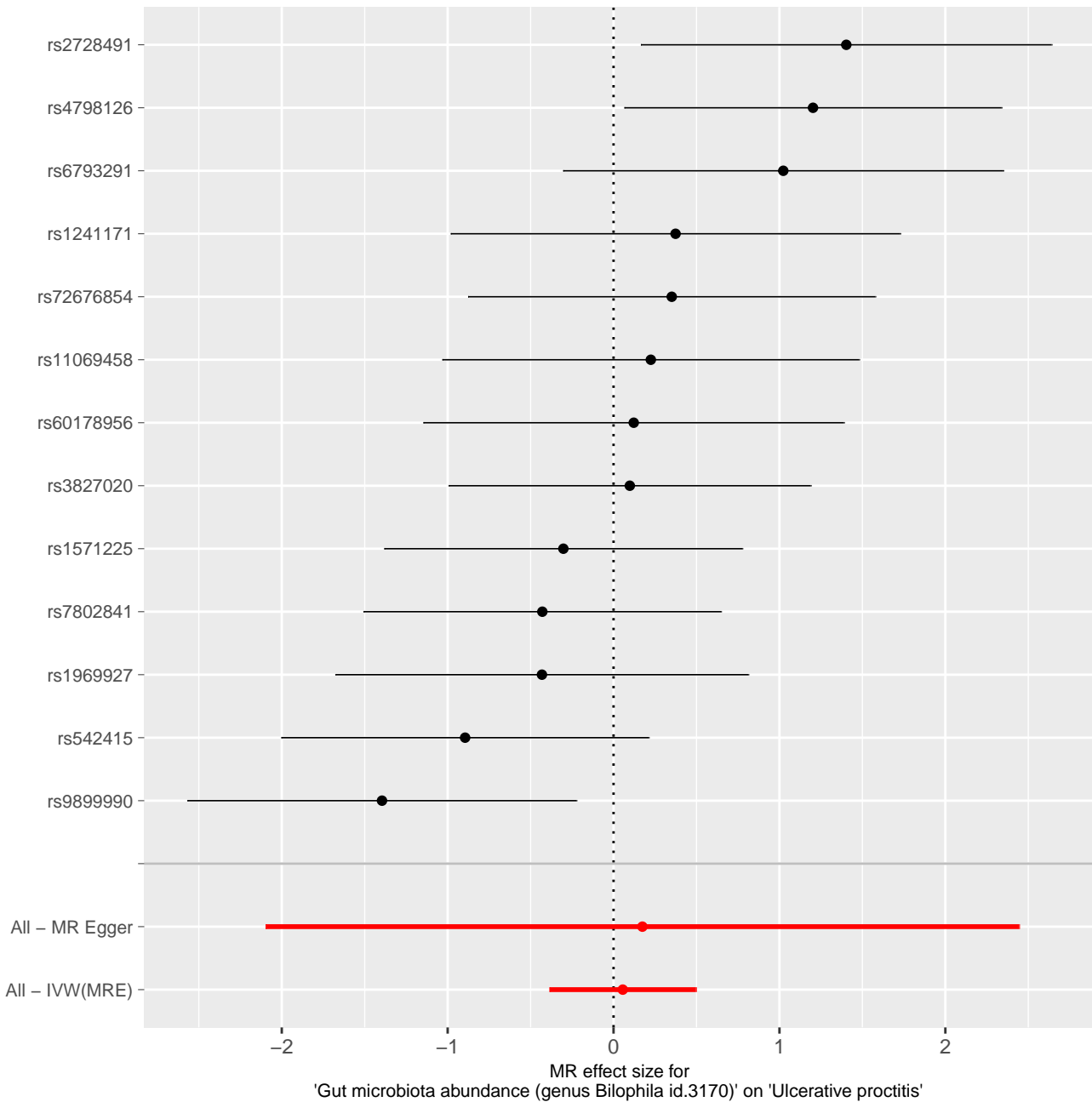


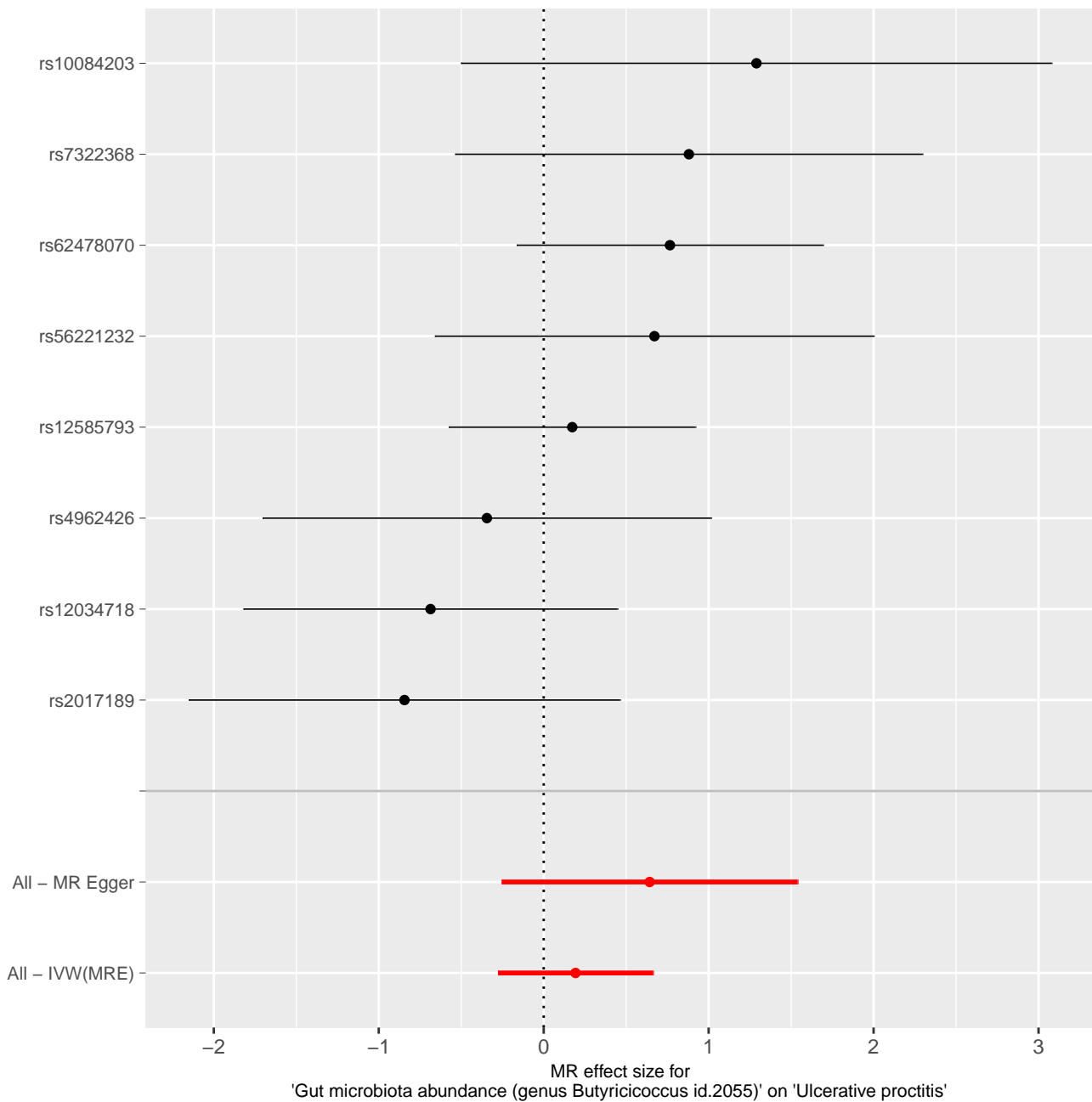




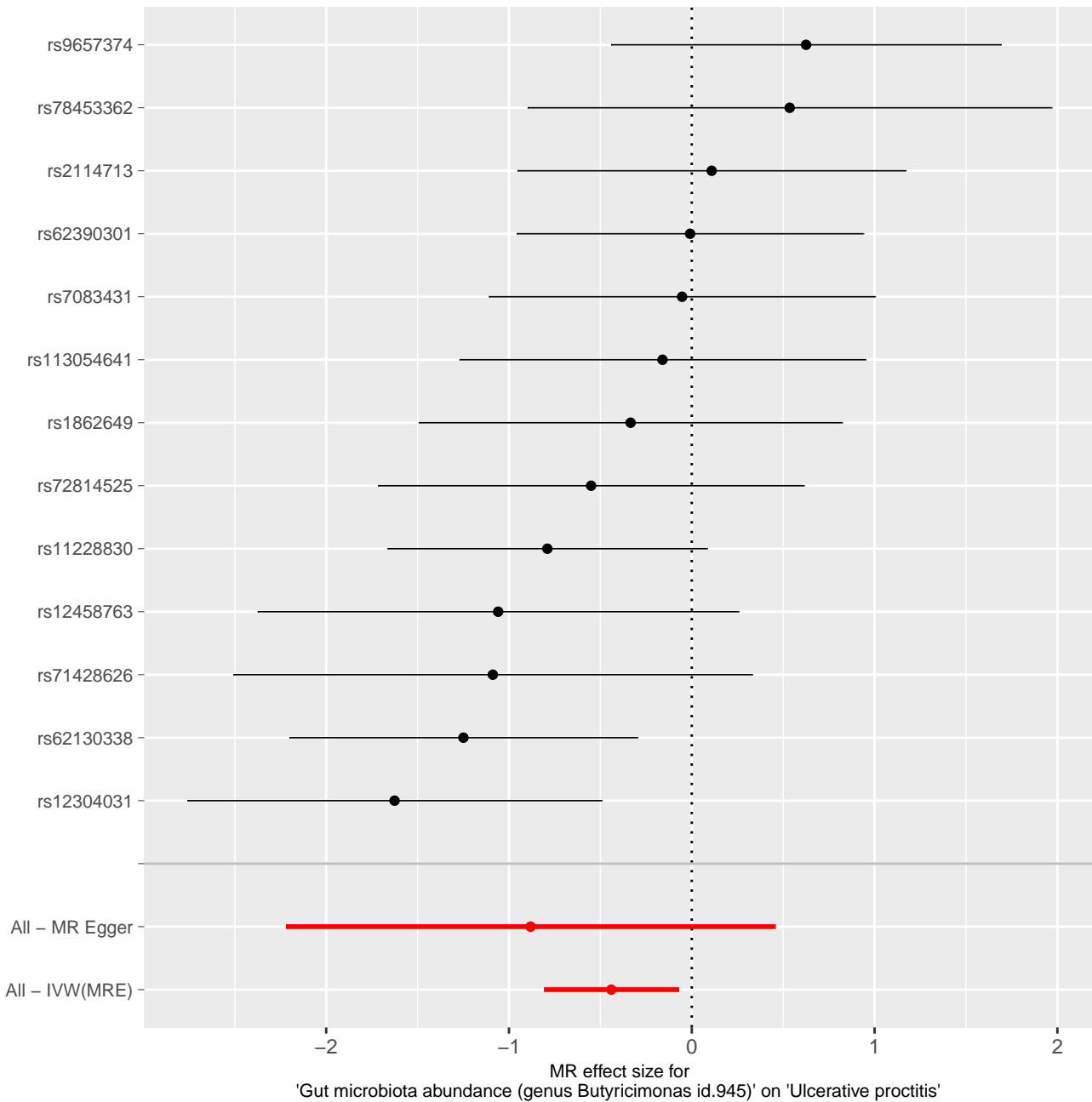


Batch 1395 : Gut microbiota abundance (genus Bilophila id.3170) on Ulcerative proctitis

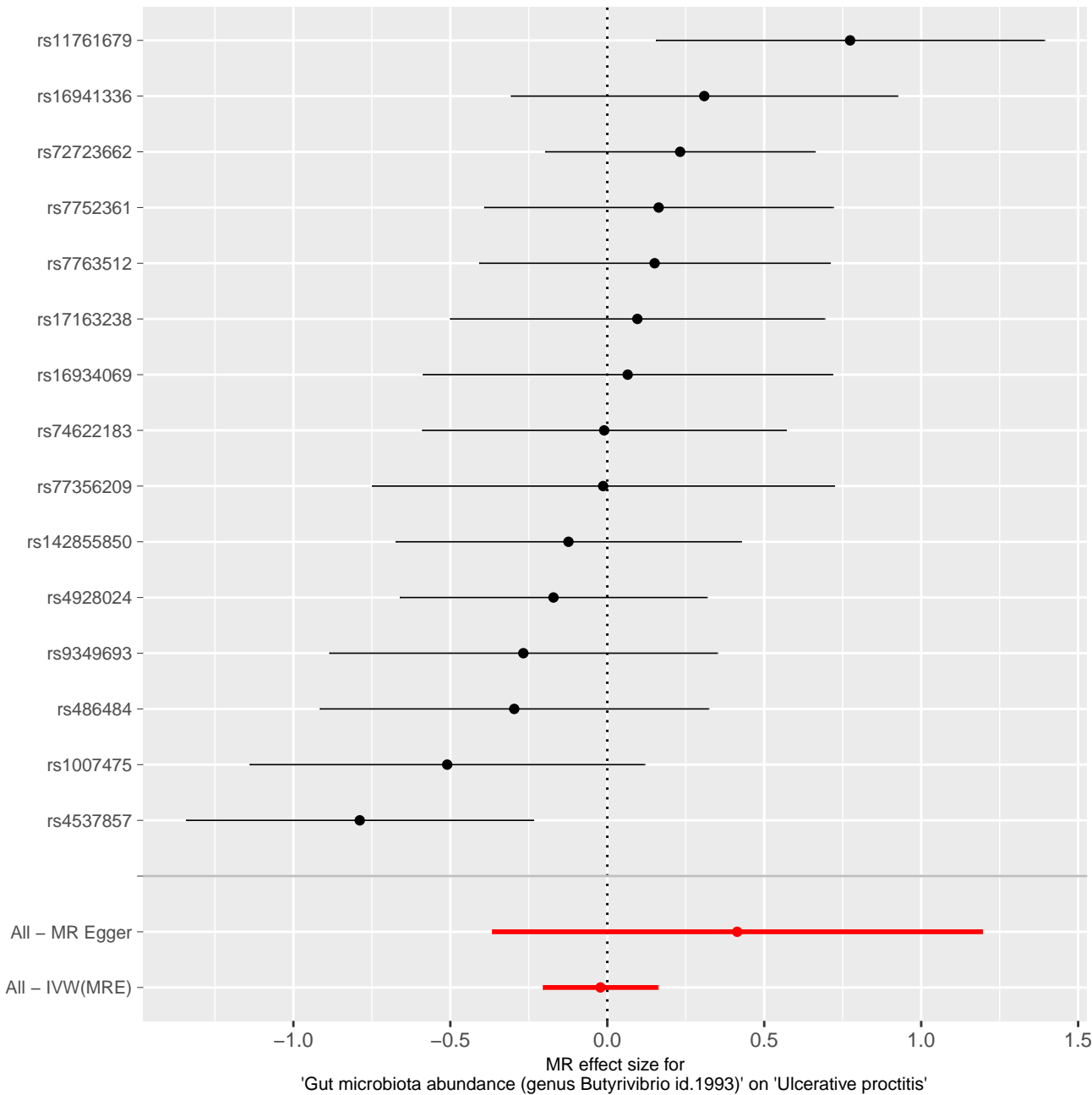


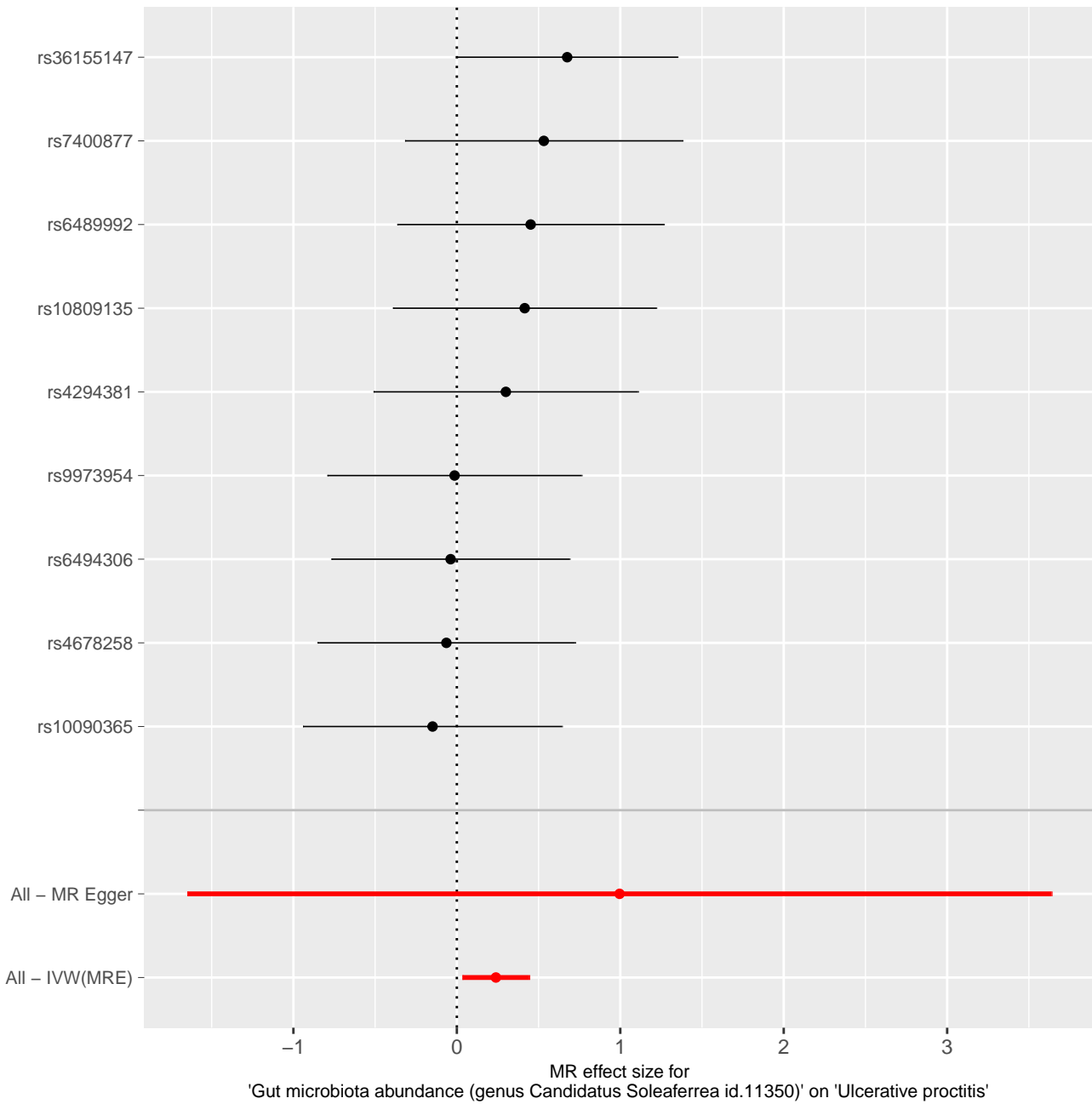


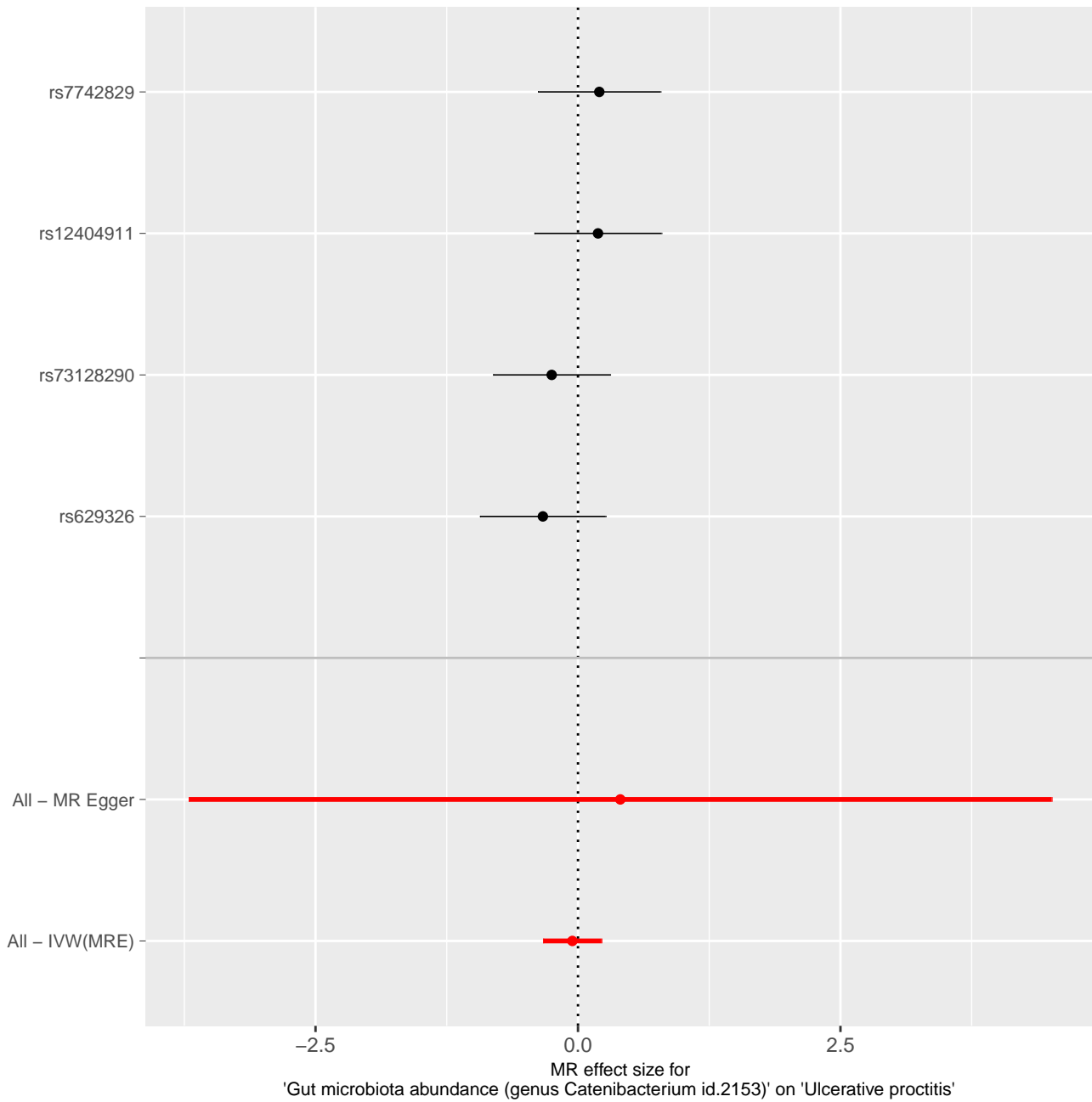
Batch 1398 : Gut microbiota abundance (genus Butyricimonas id.945) on Ulcerative proctitis

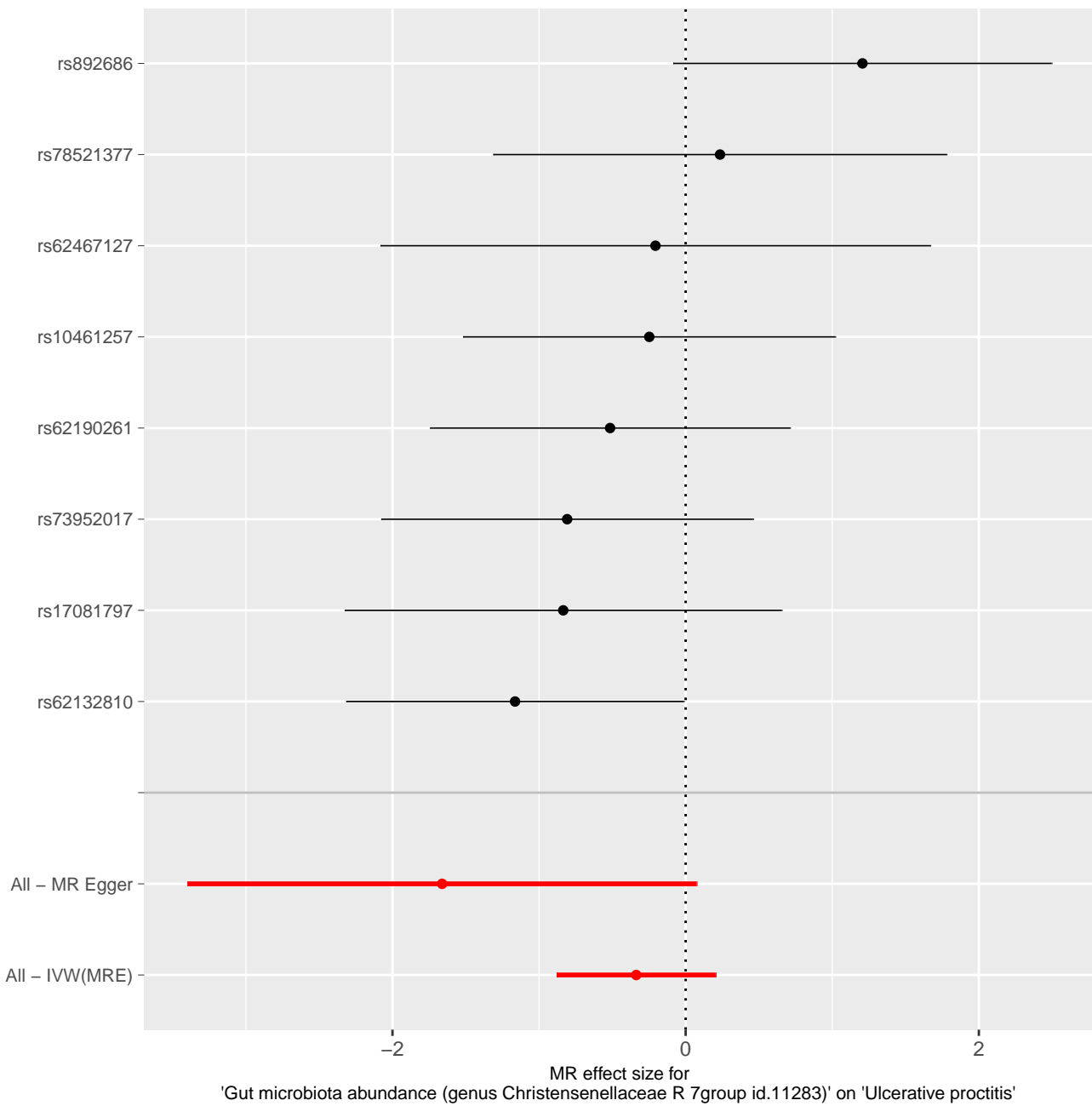


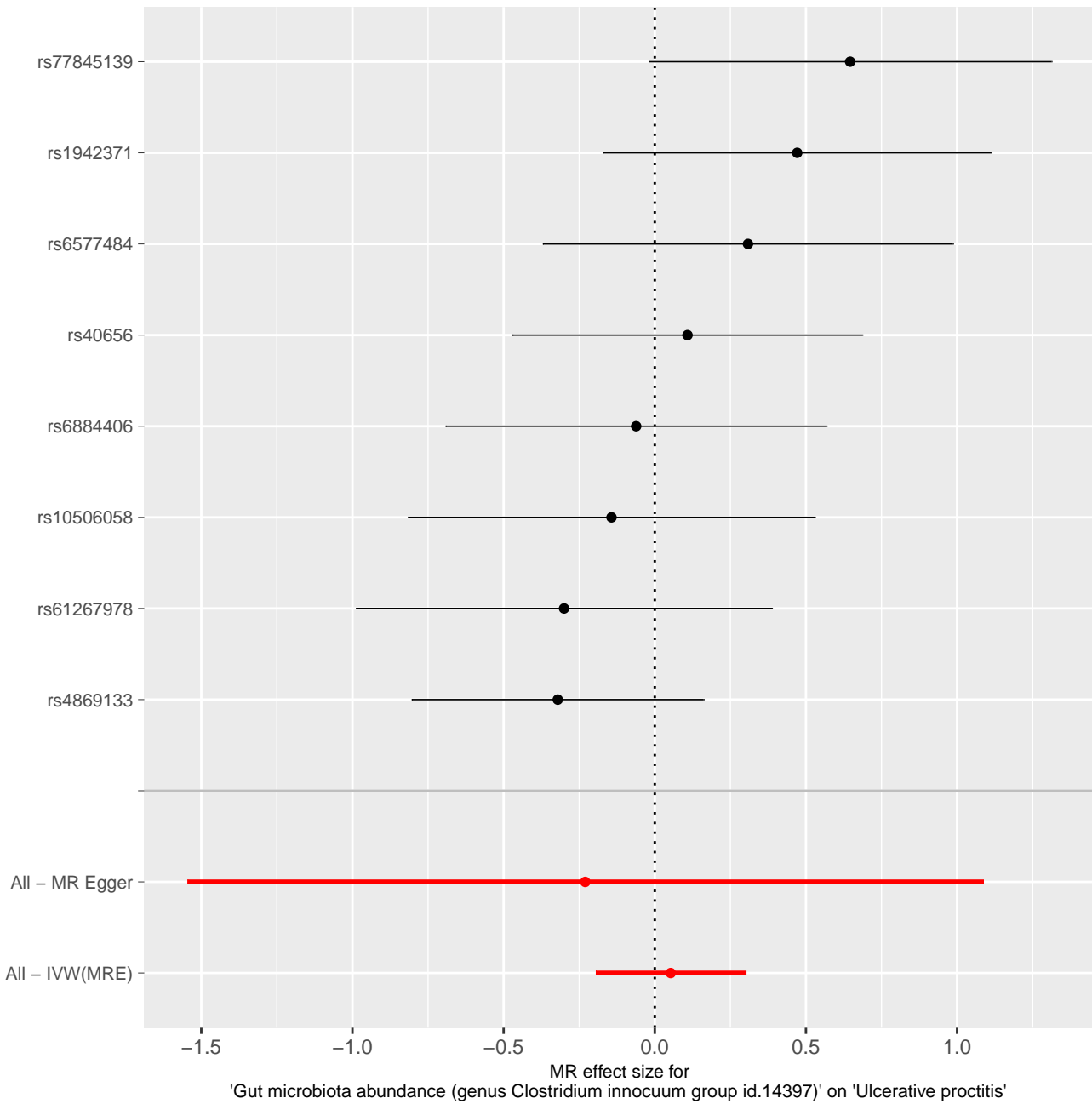
Batch 1399 : Gut microbiota abundance (genus *Butyrivibrio* id.1993) on Ulcerative proctitis

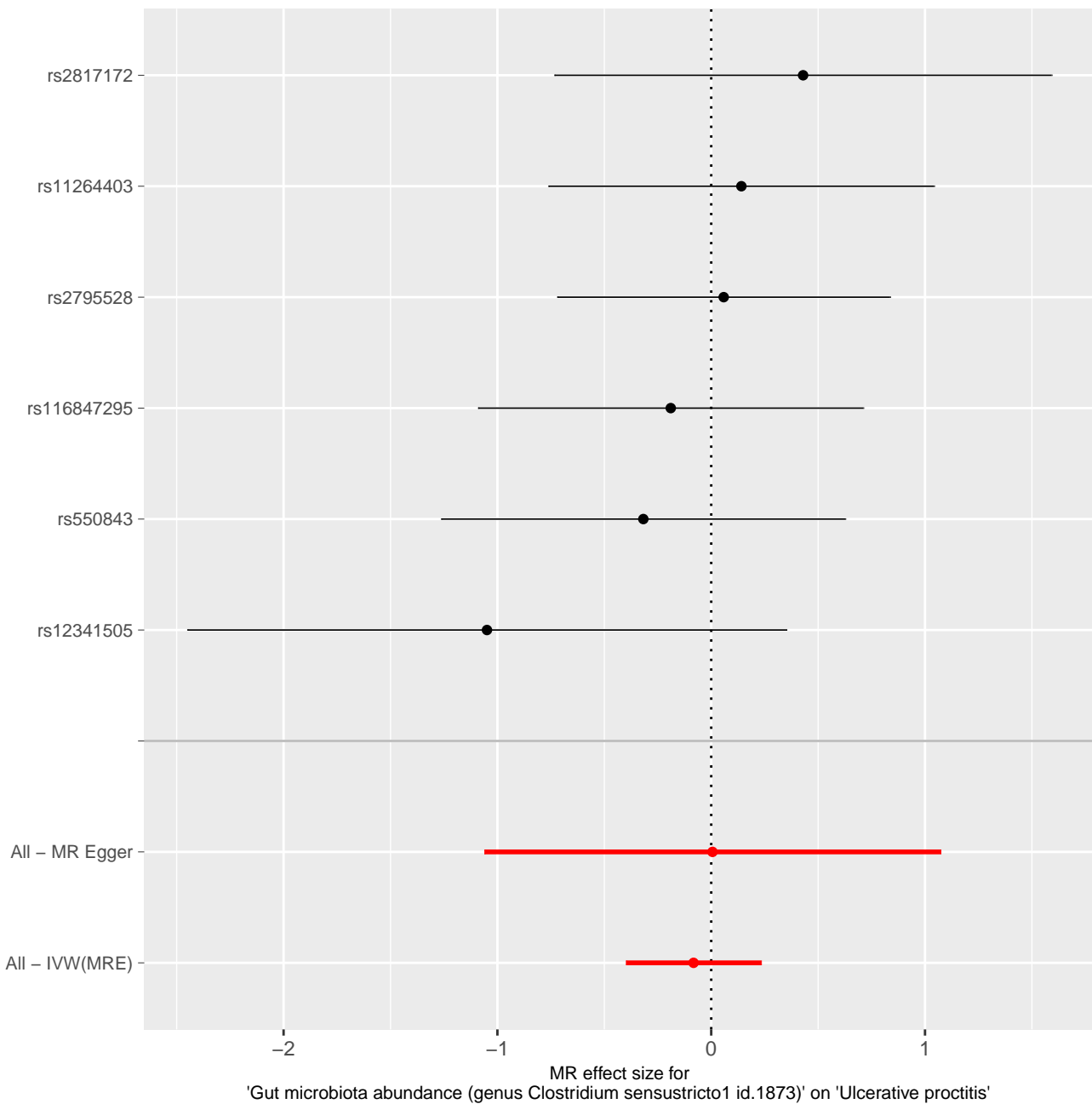


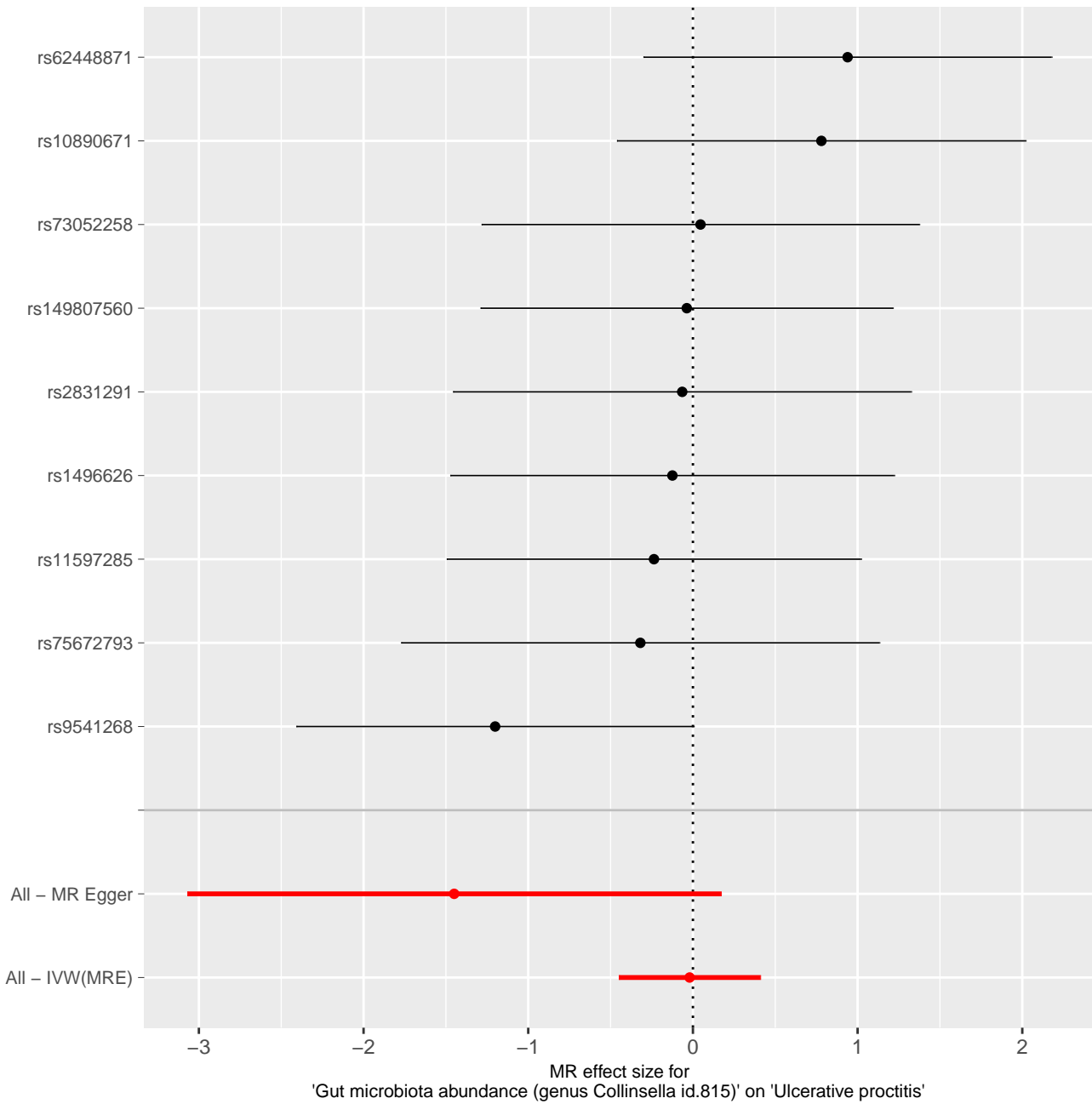


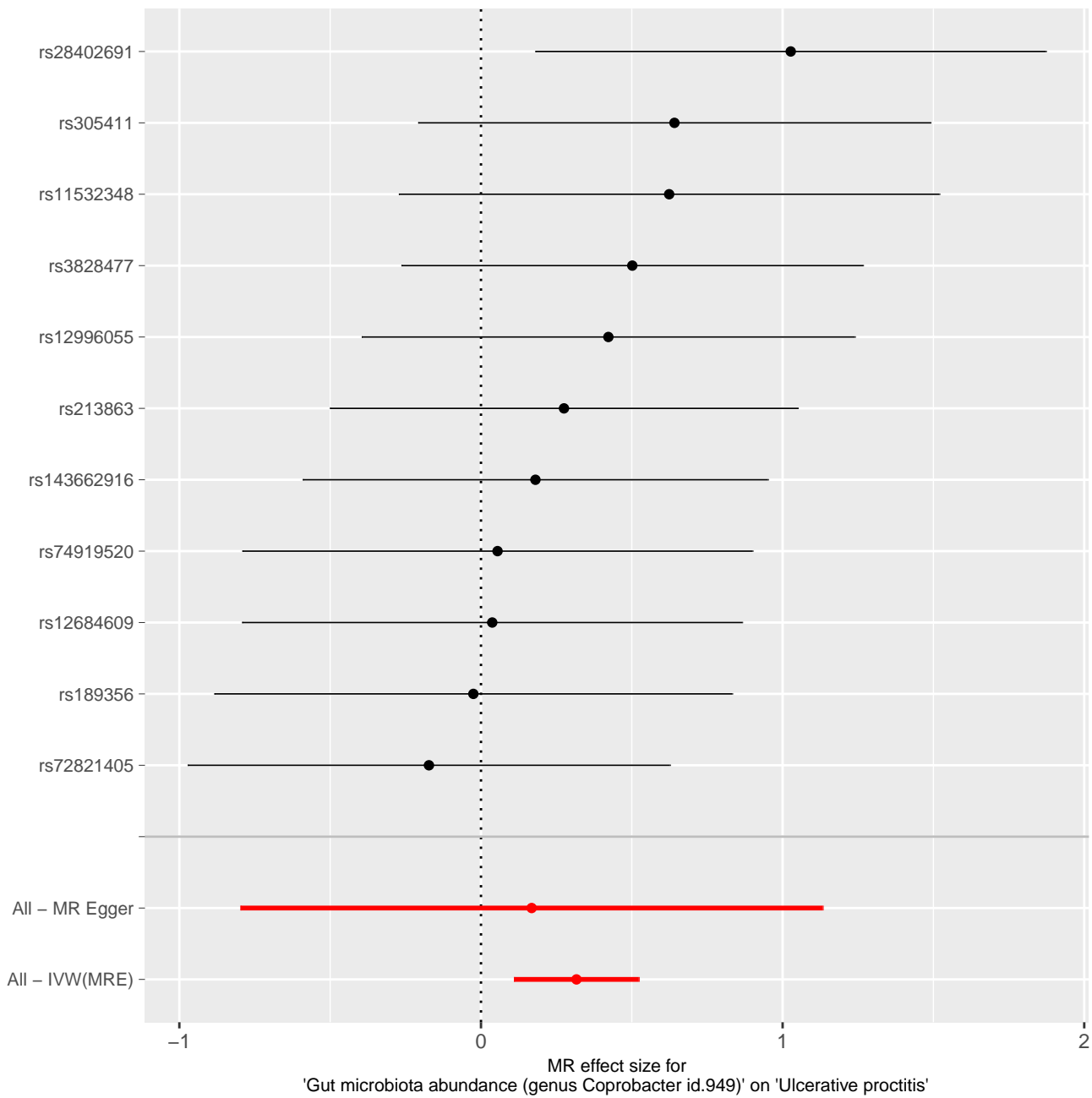


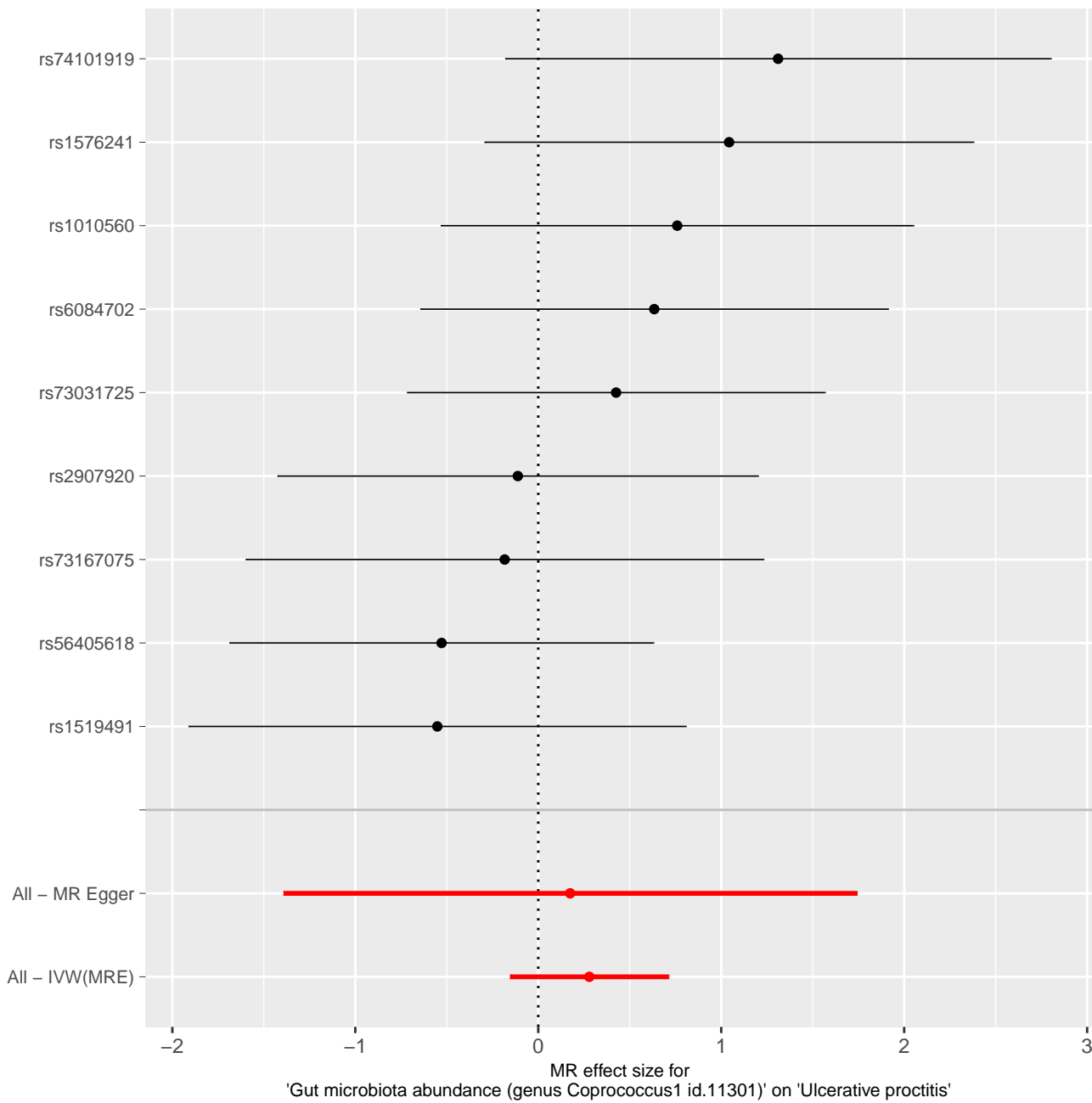


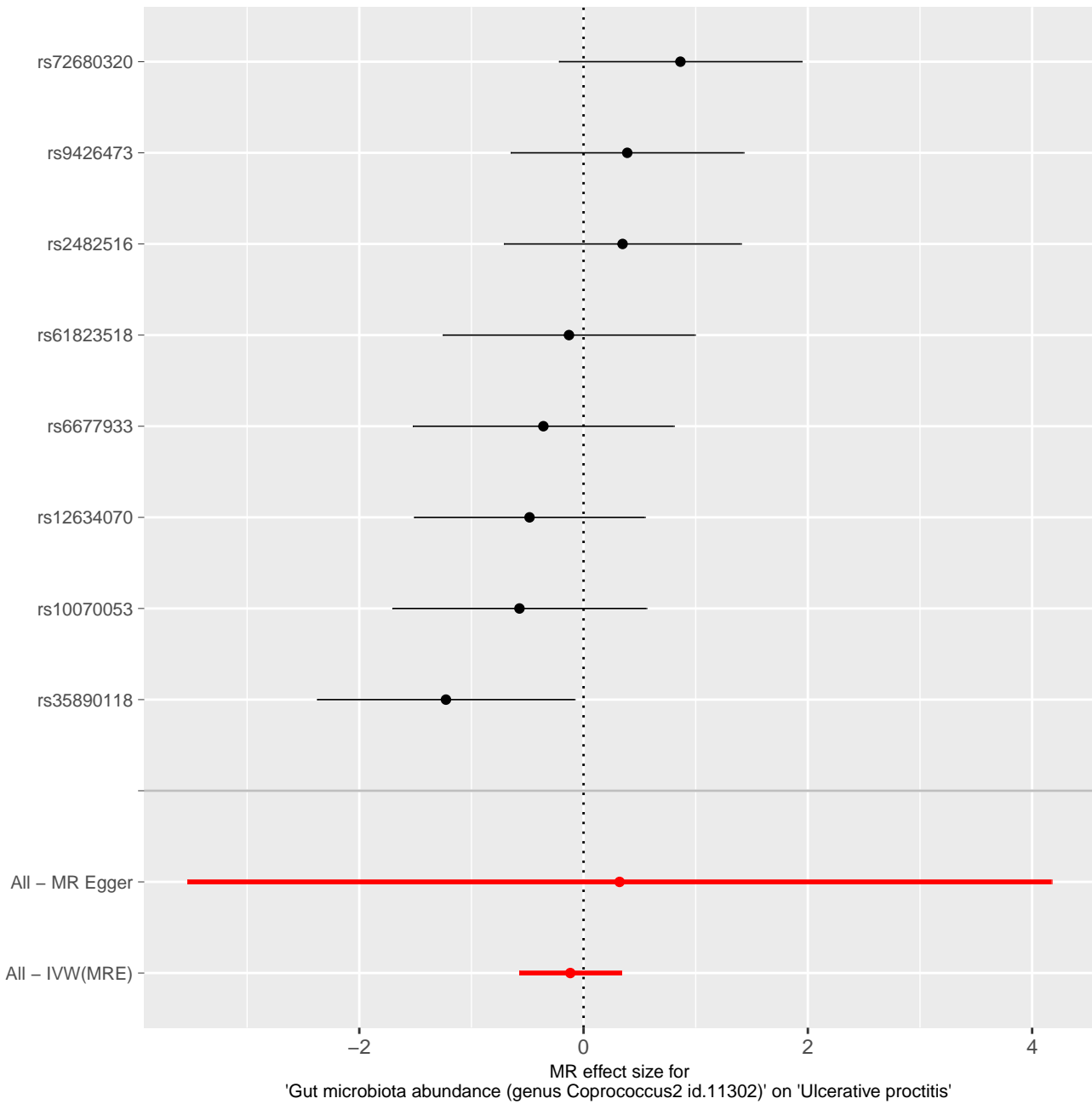


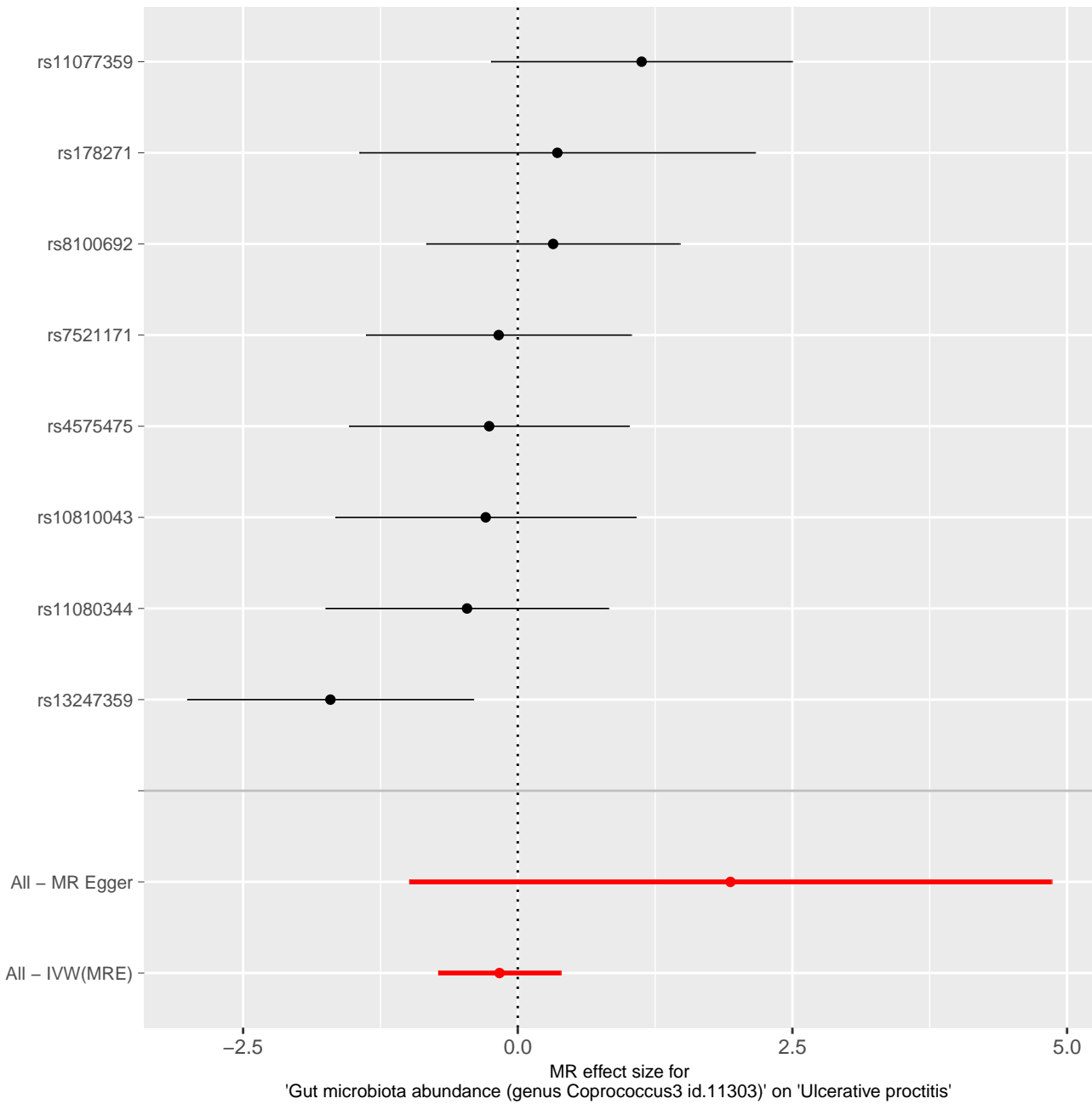


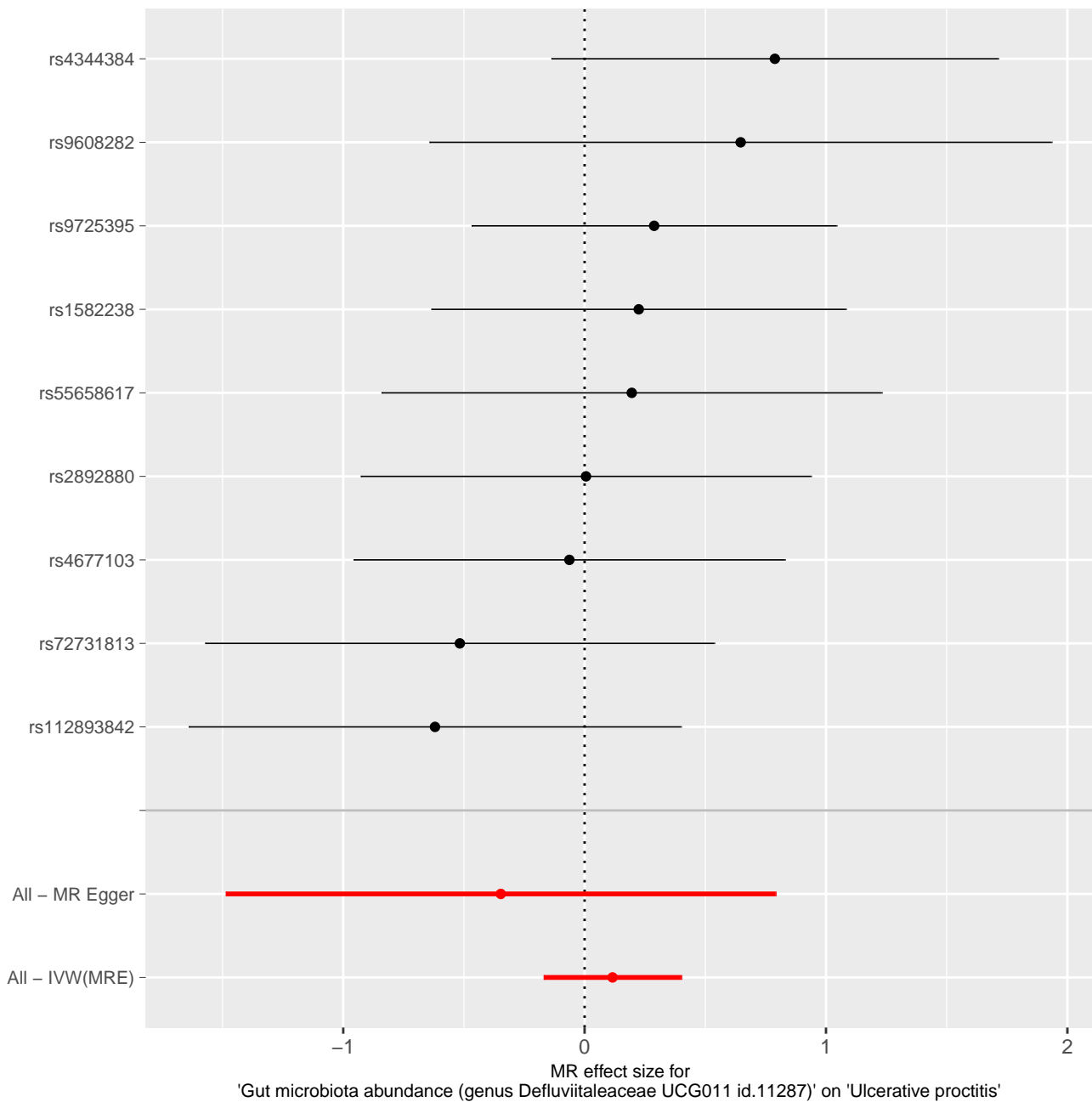


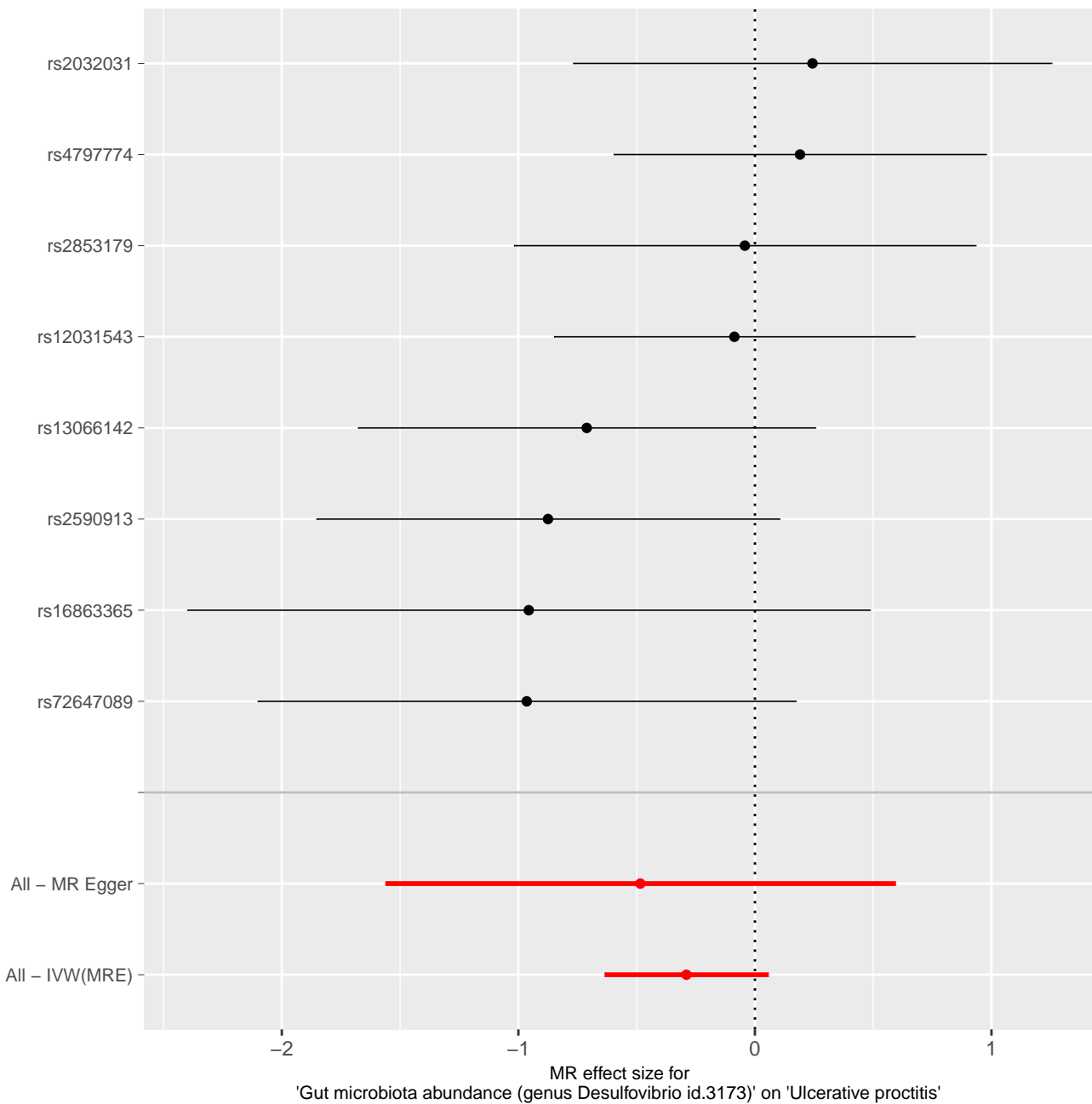




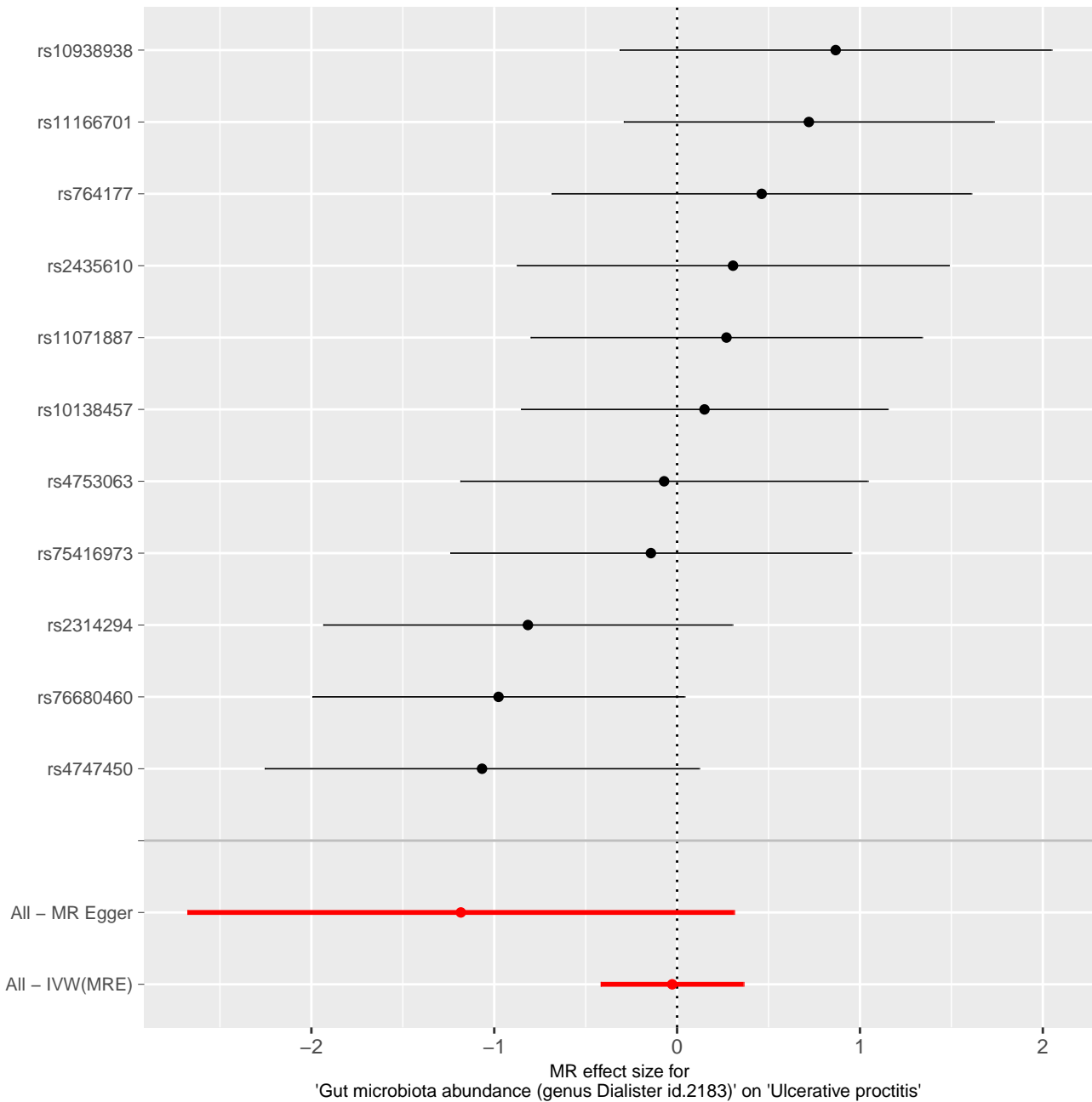


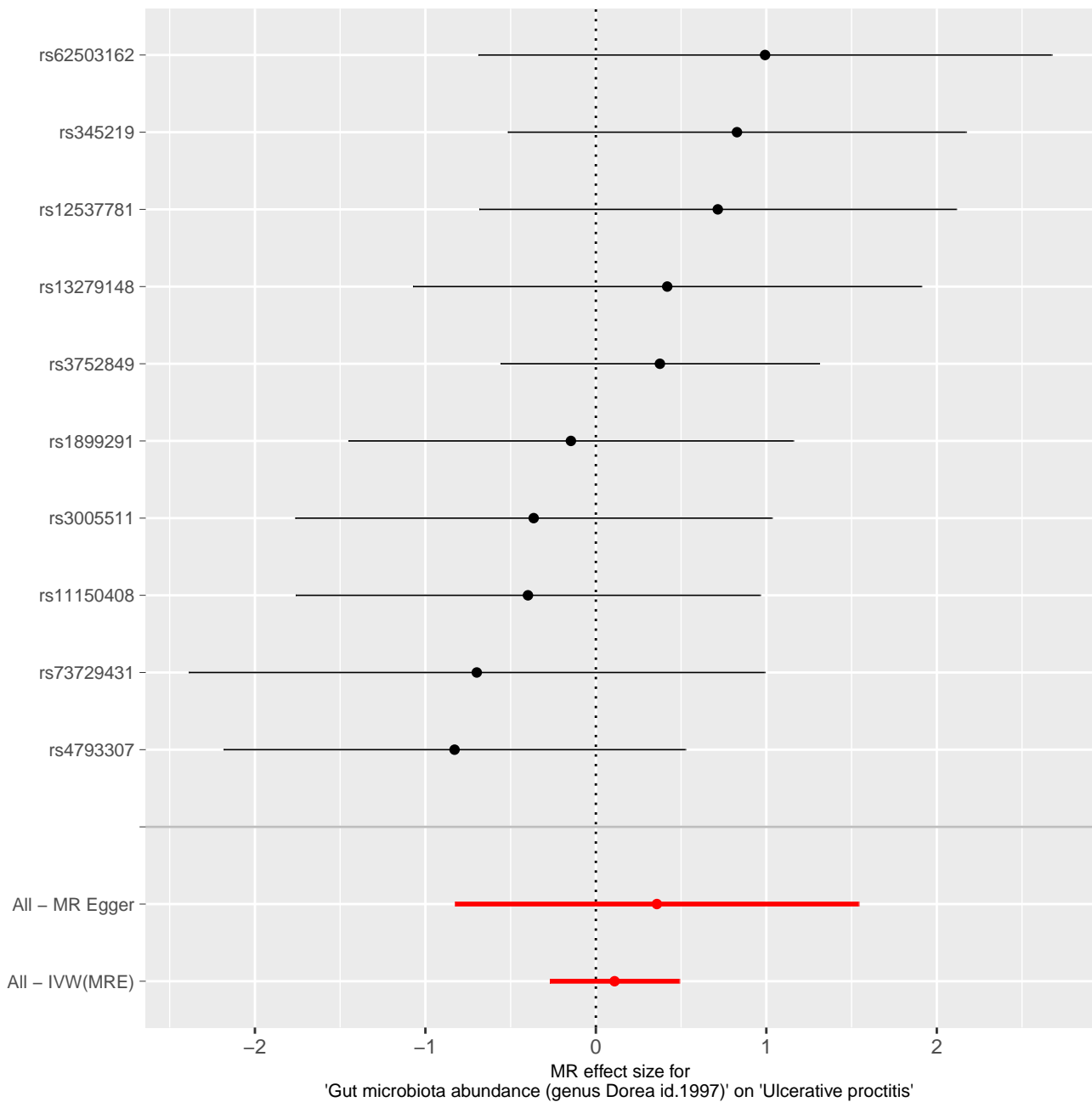


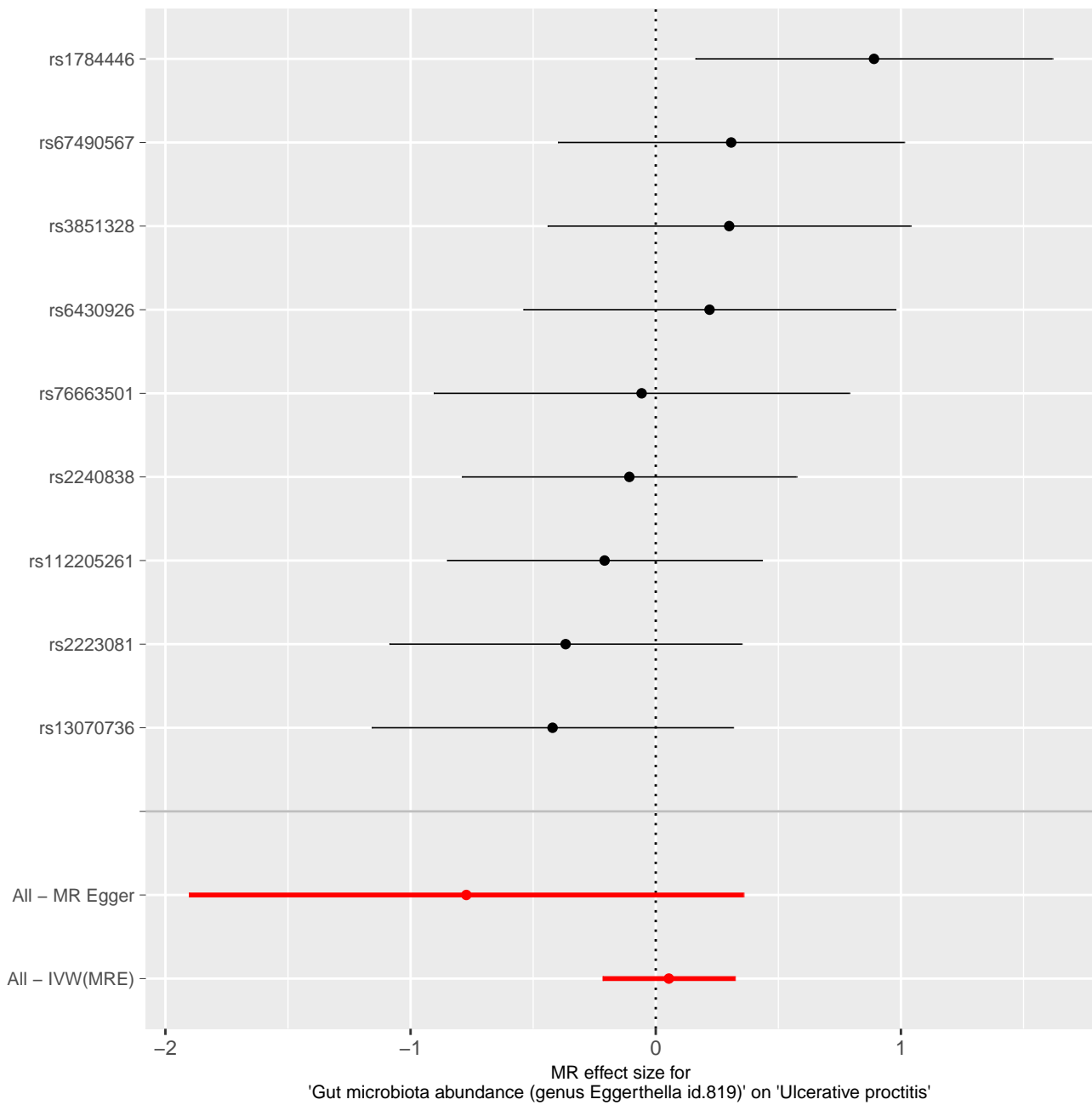




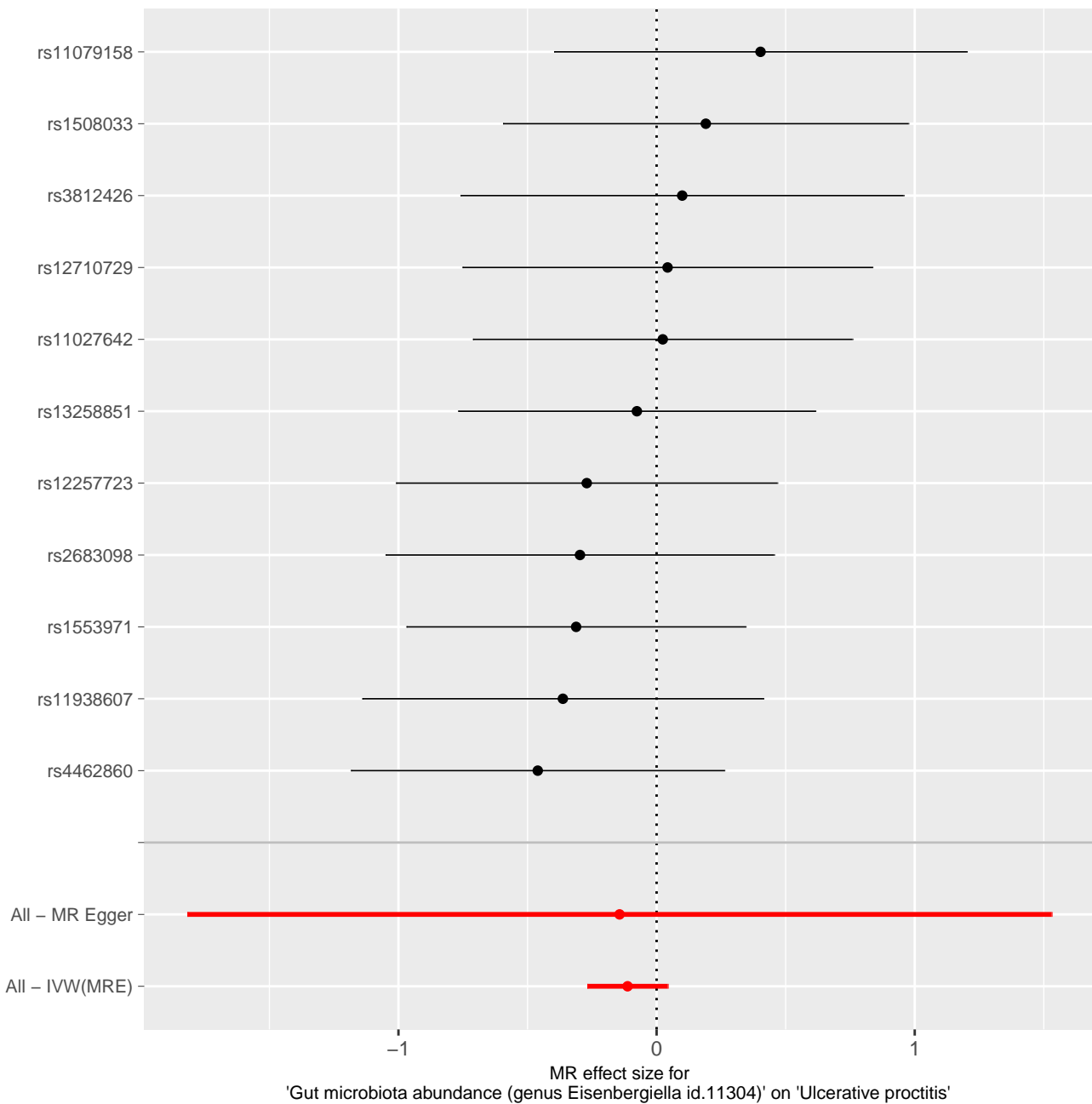
Batch 1412 : Gut microbiota abundance (genus Dialister id.2183) on Ulcerative proctitis

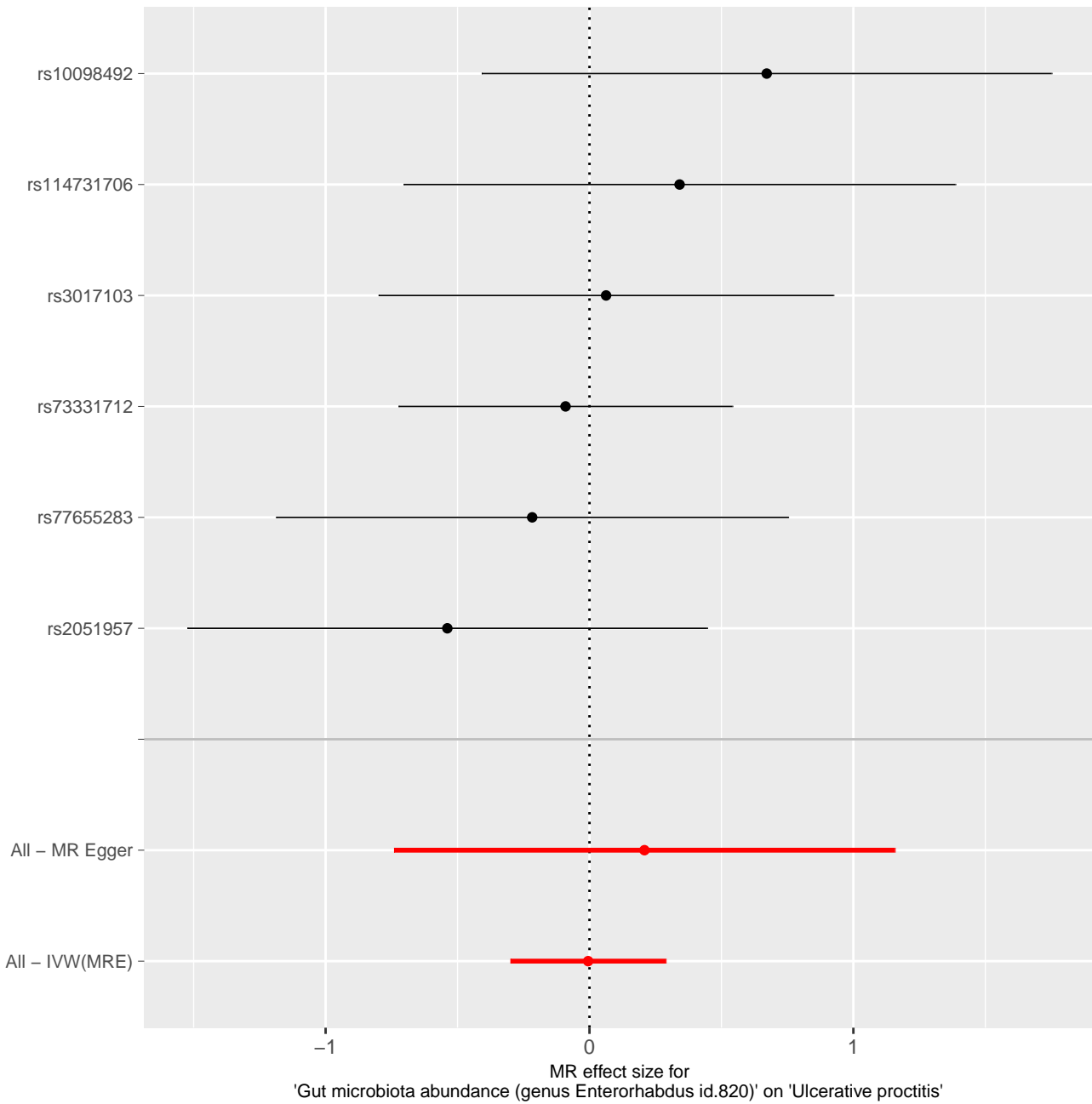


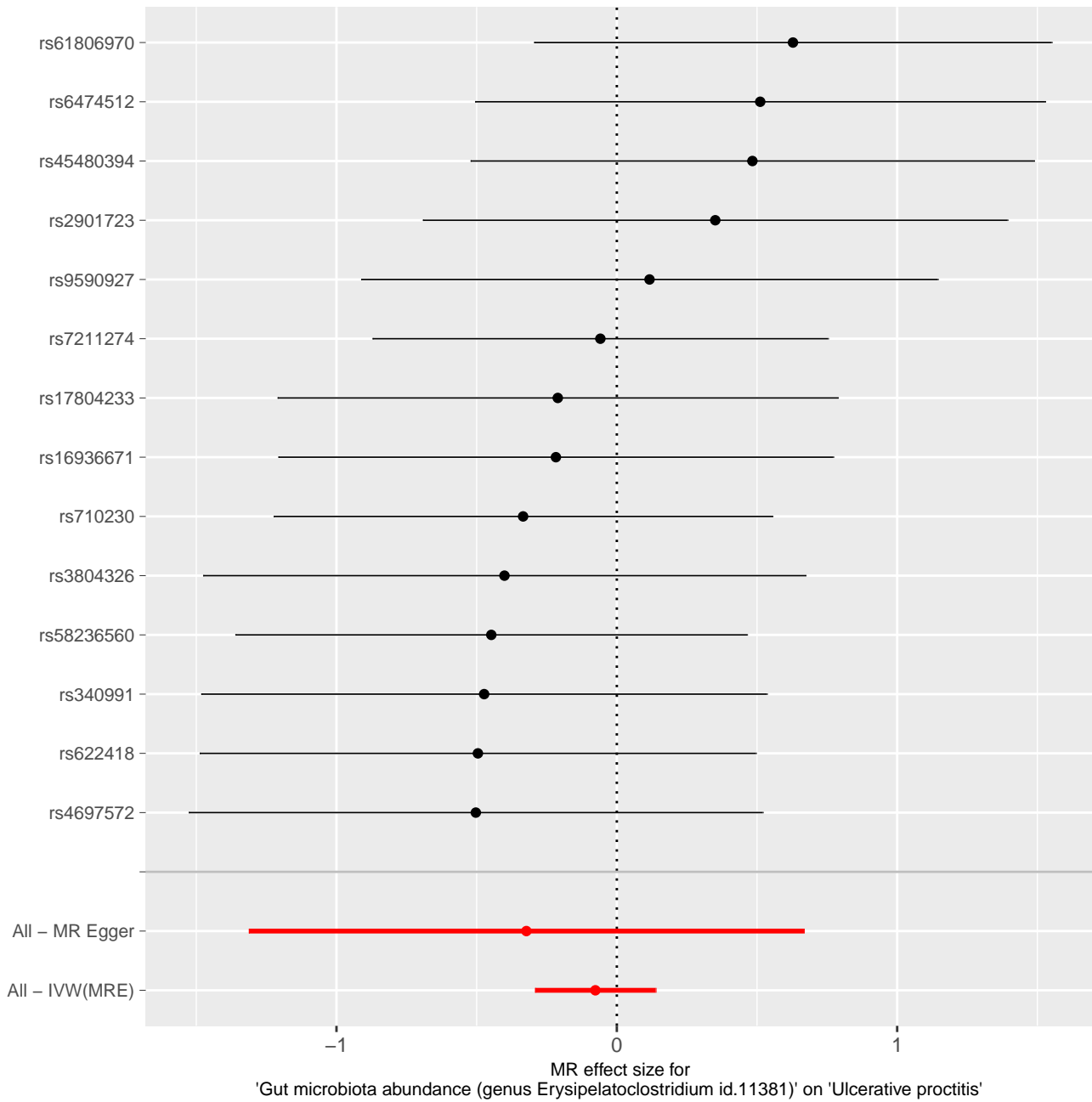


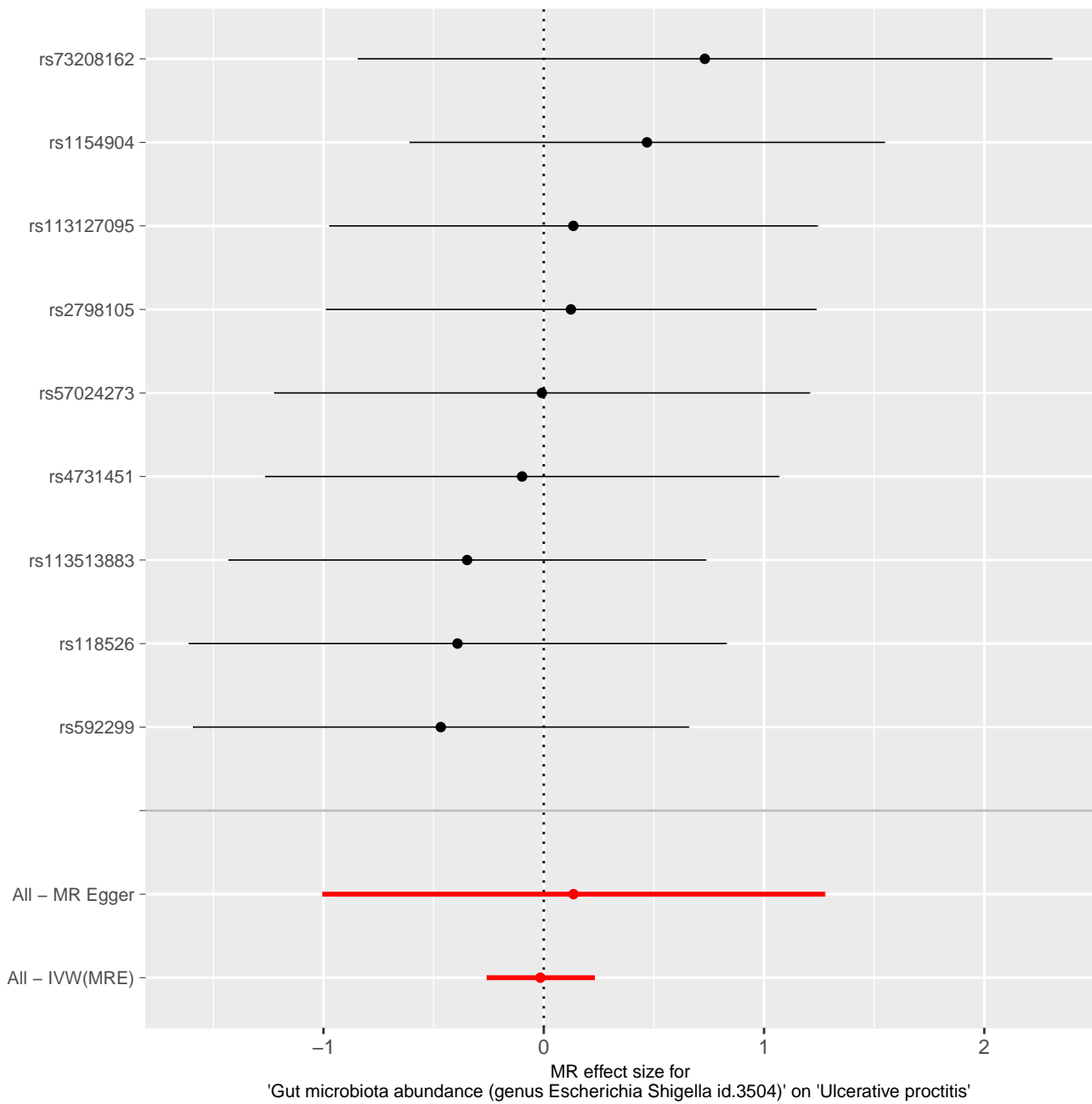


Batch 1415 : Gut microbiota abundance (genus Eisenbergiella id.11304) on Ulcerative proctitis

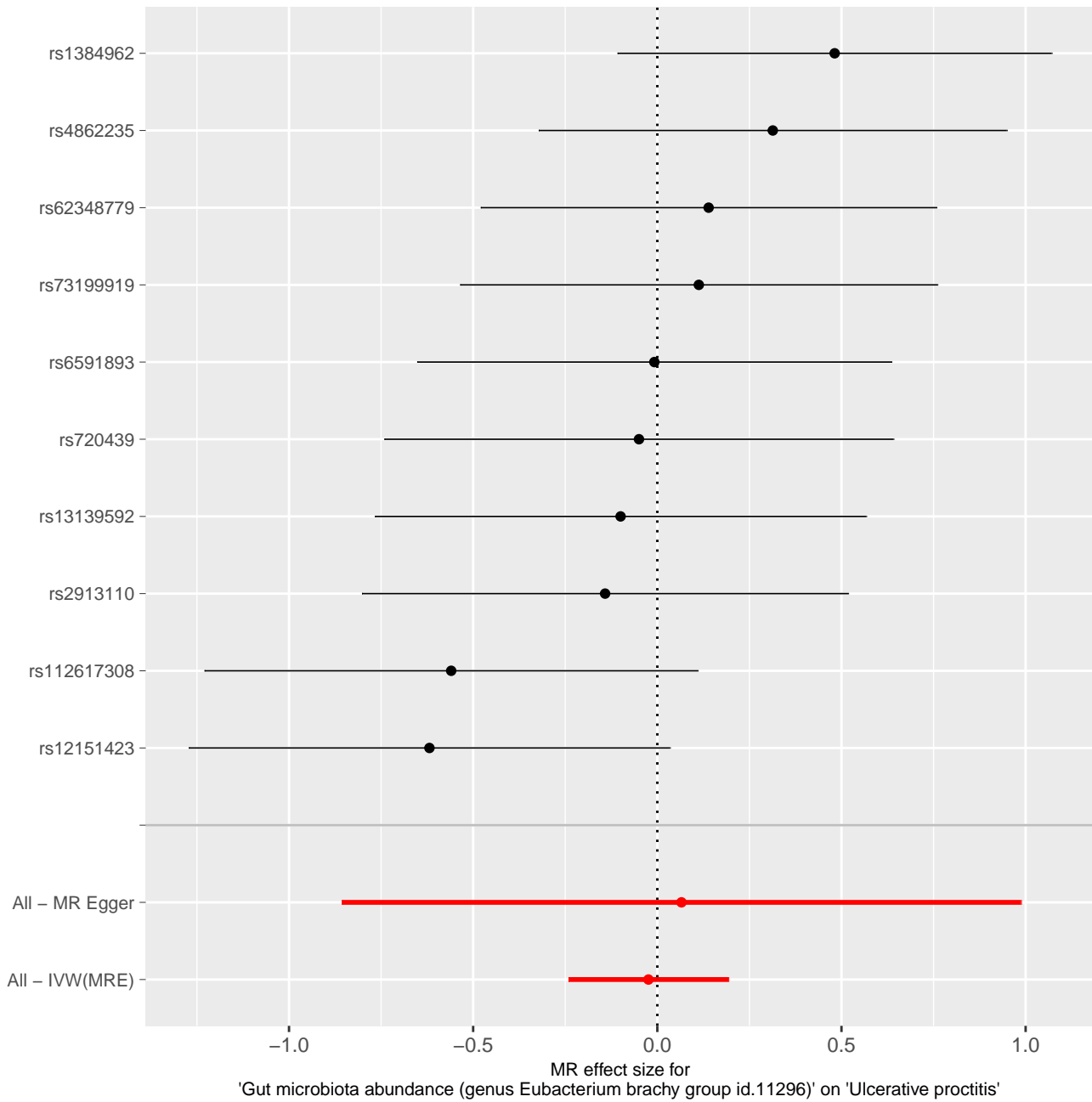


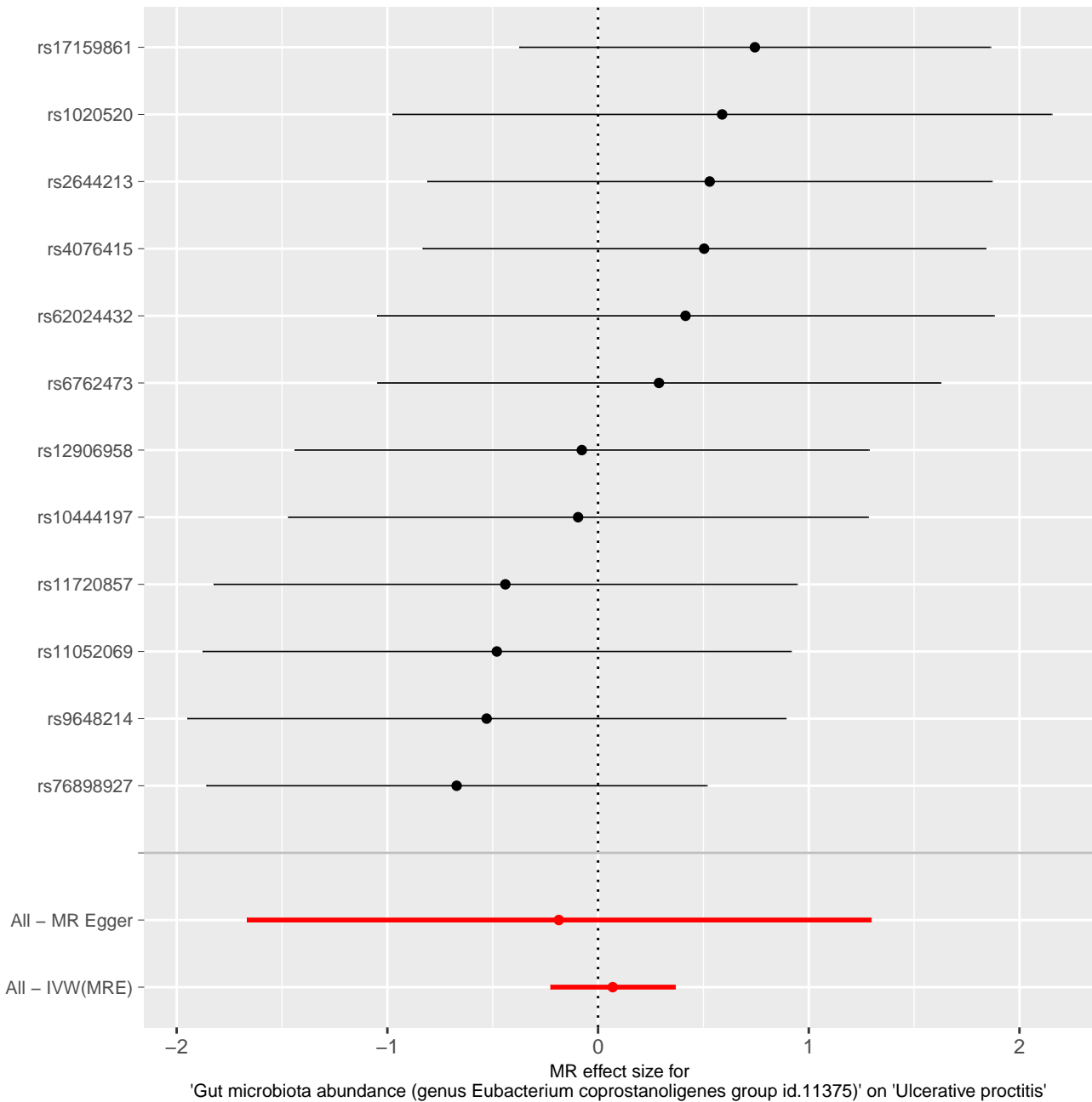


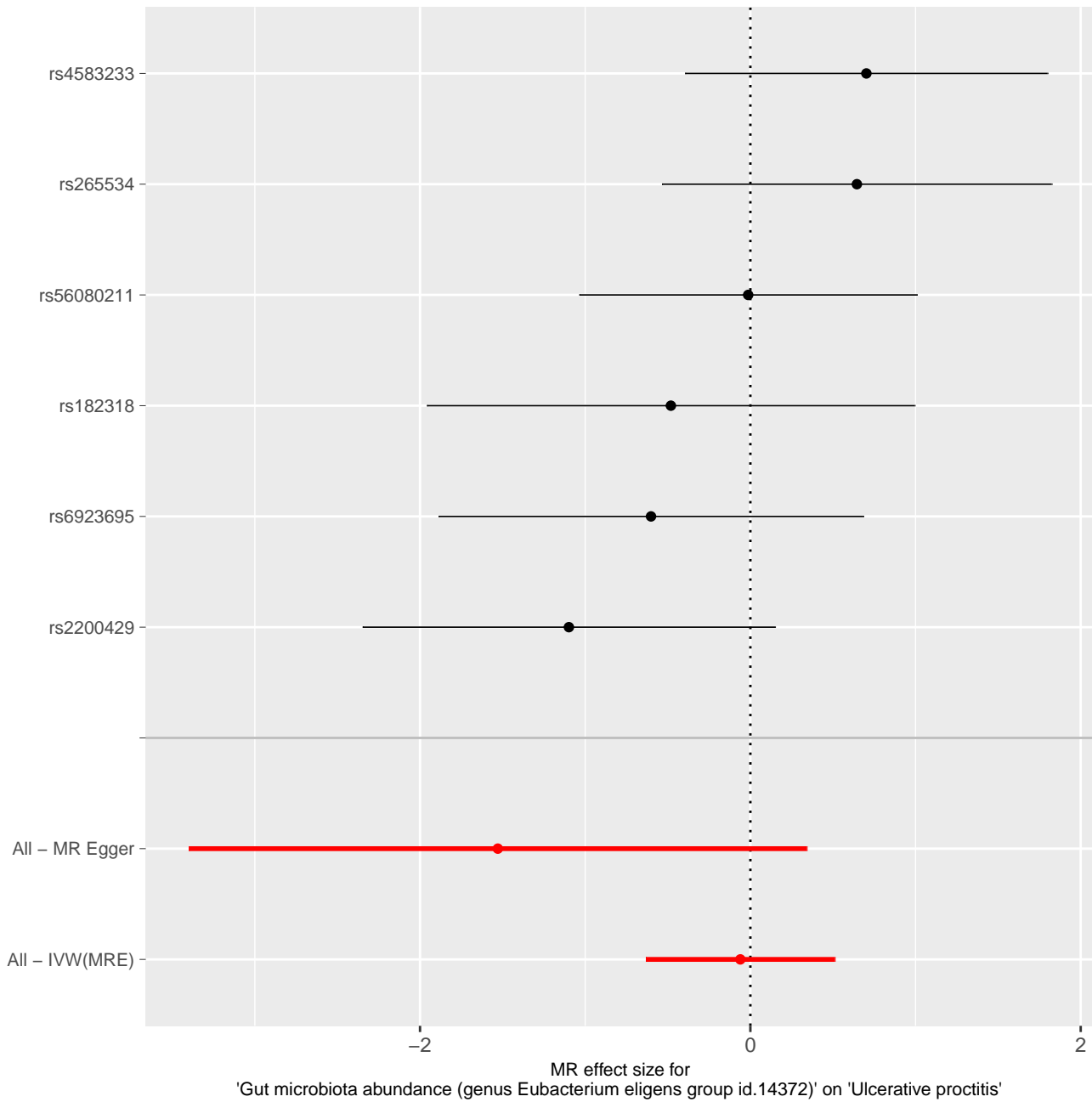


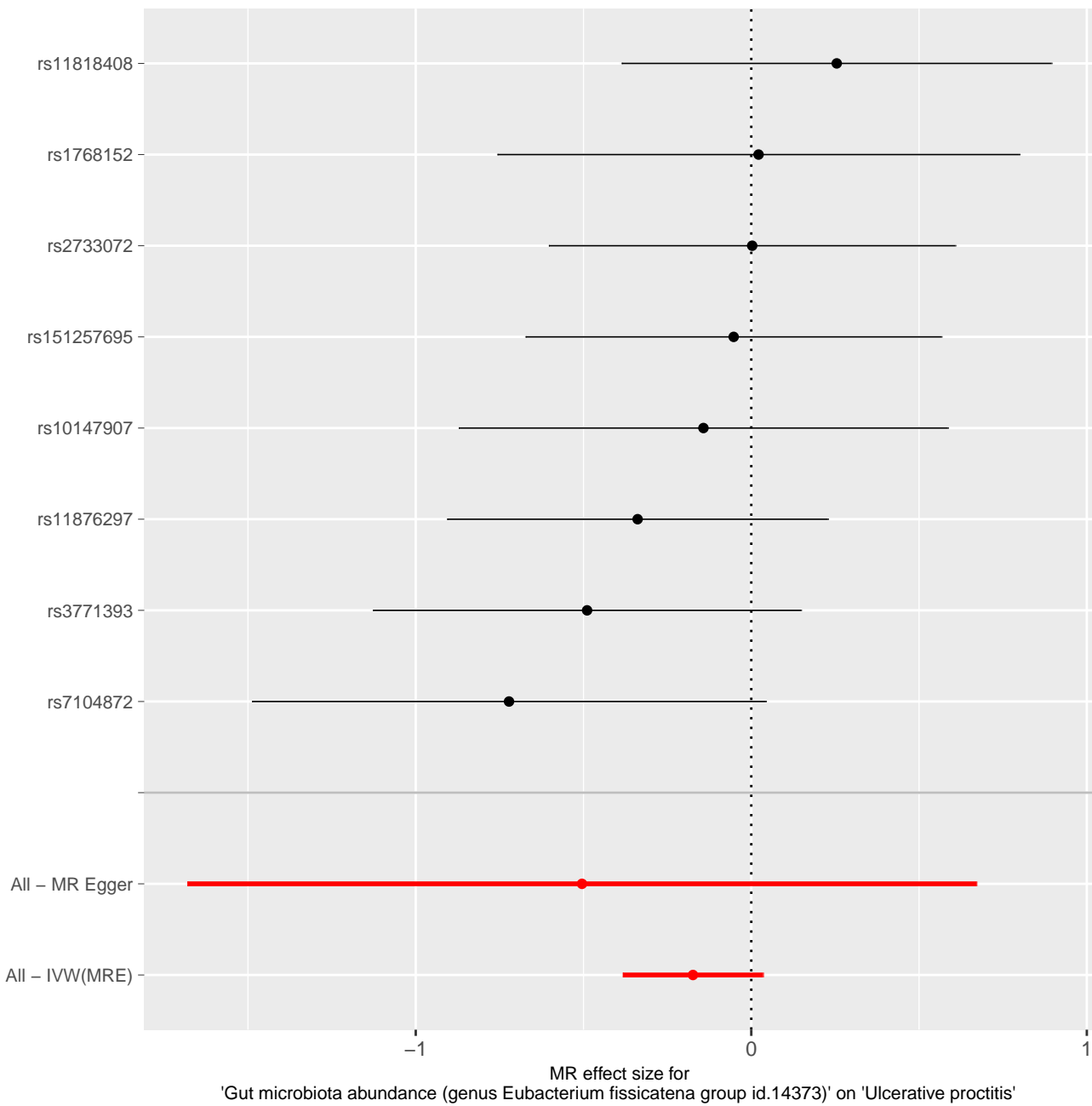


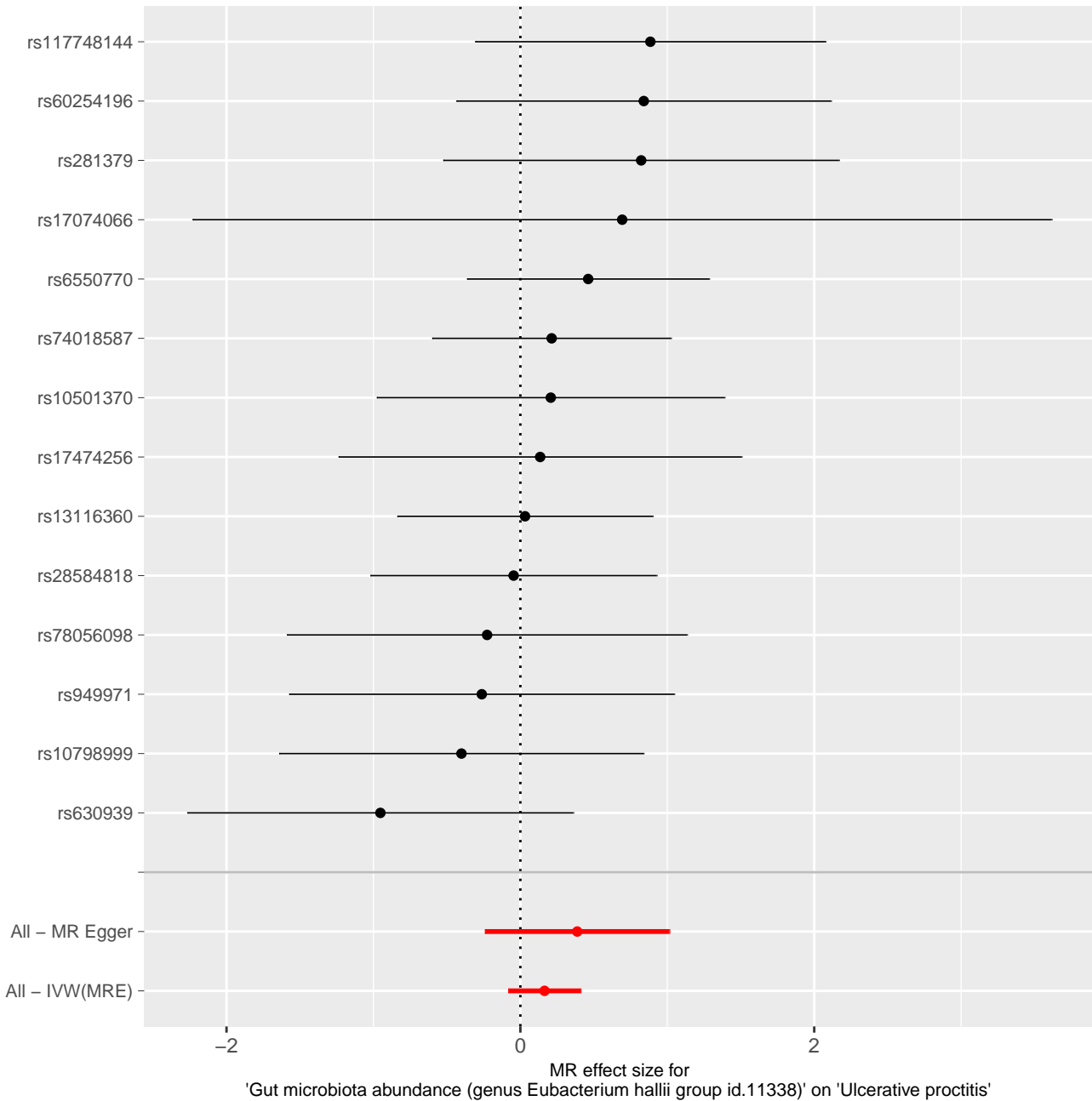
Batch 1420 : Gut microbiota abundance (genus Eubacterium brachy group id.11296) on Ulcerative proctitis

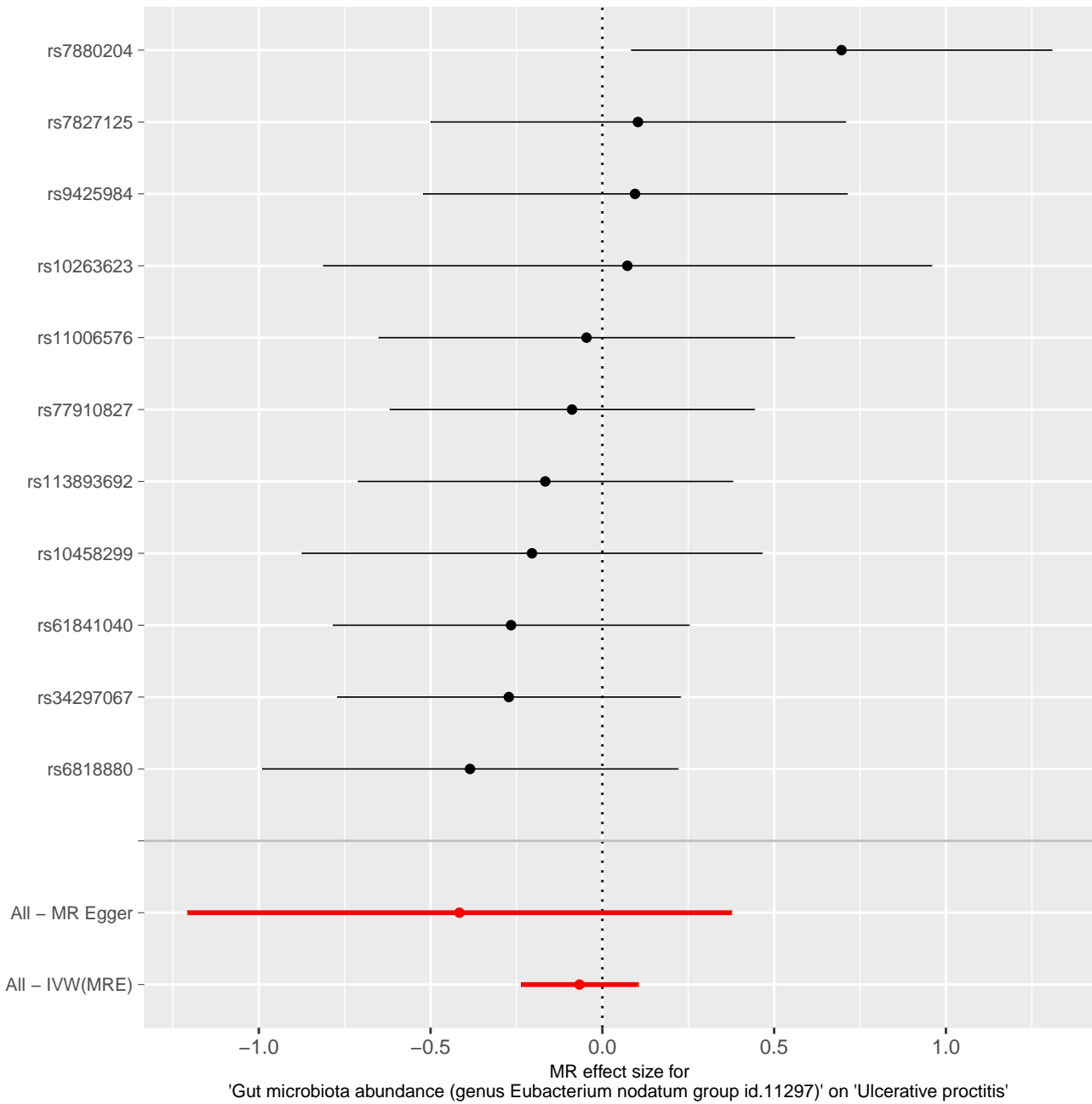


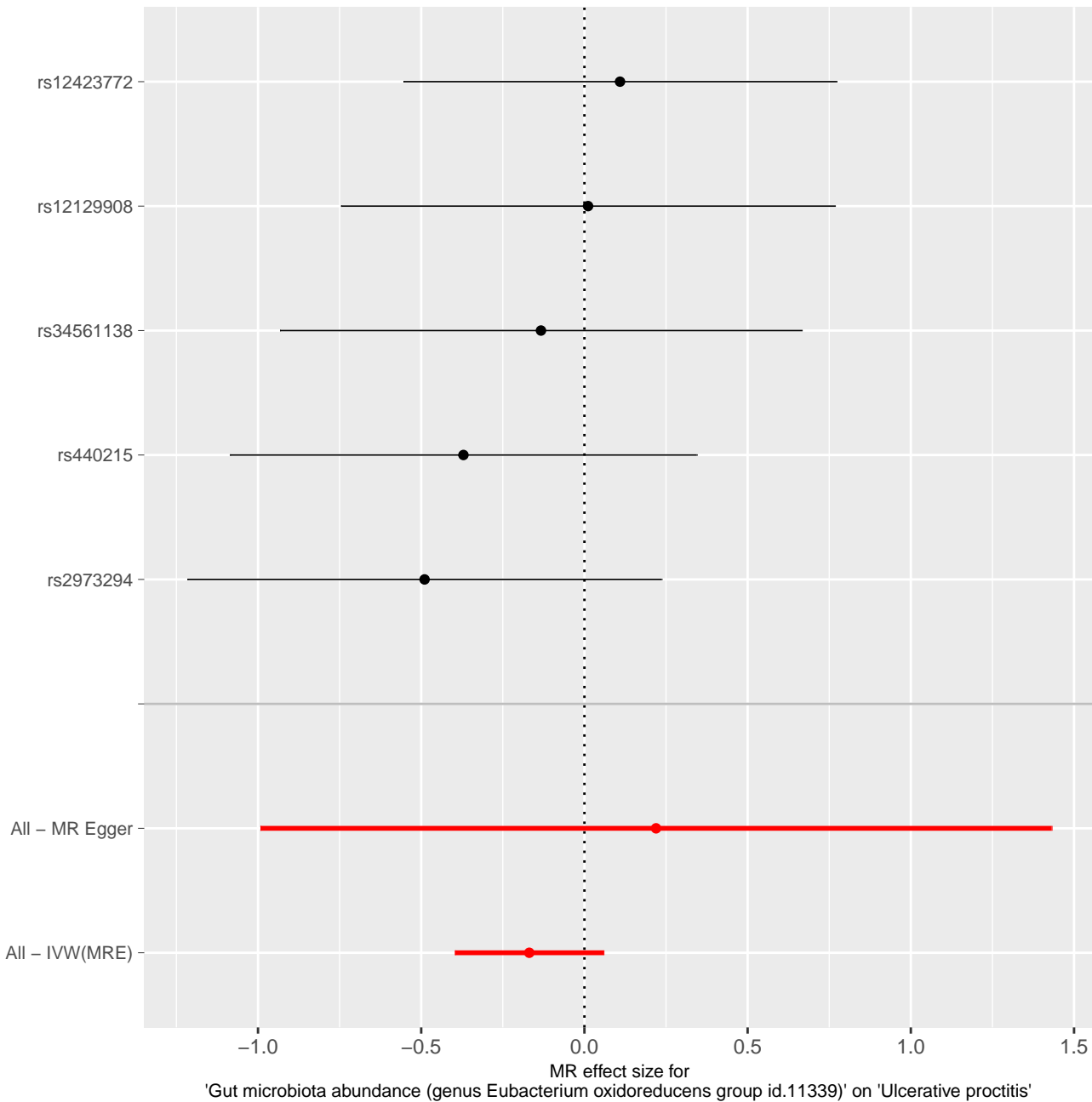


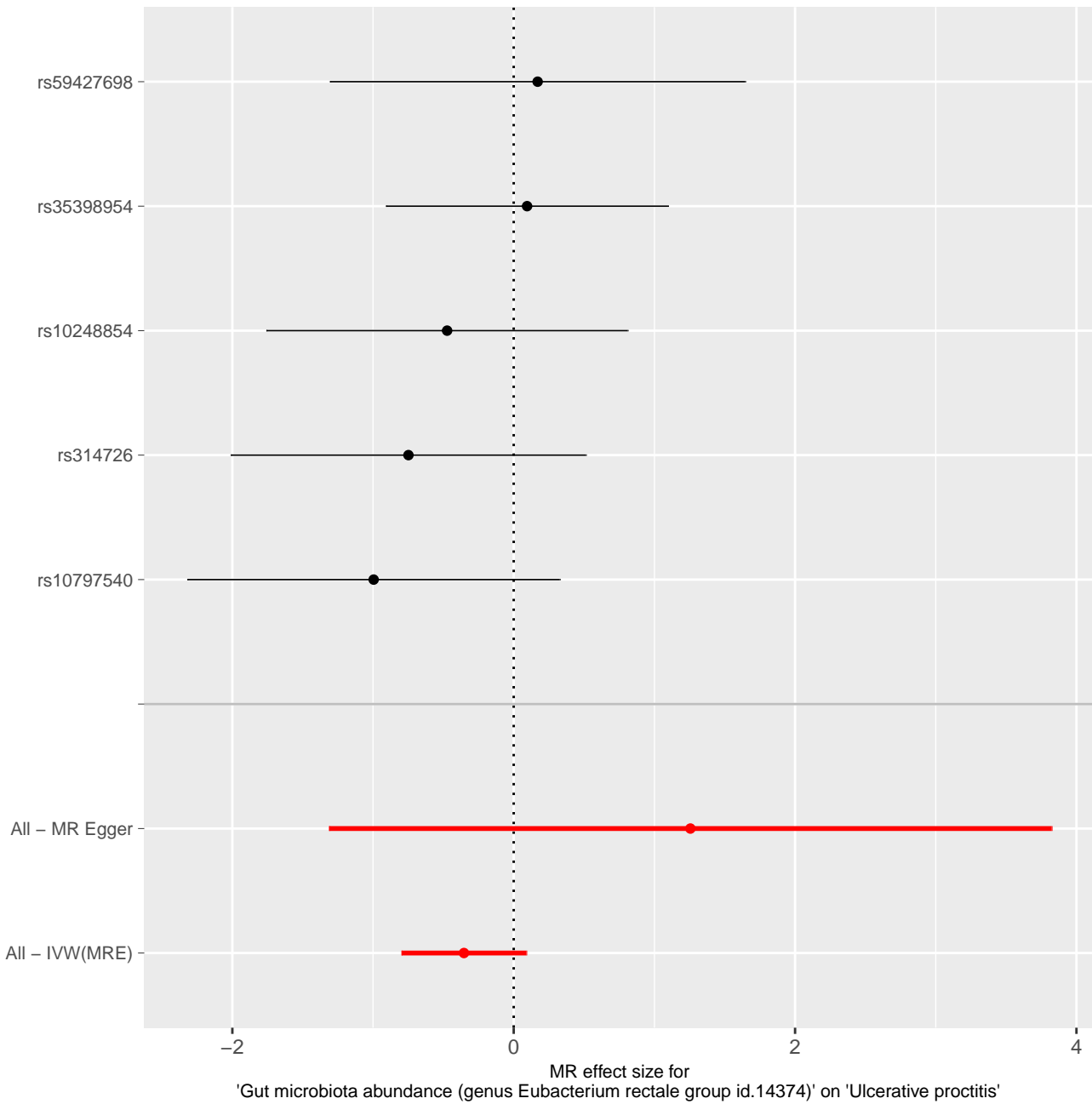


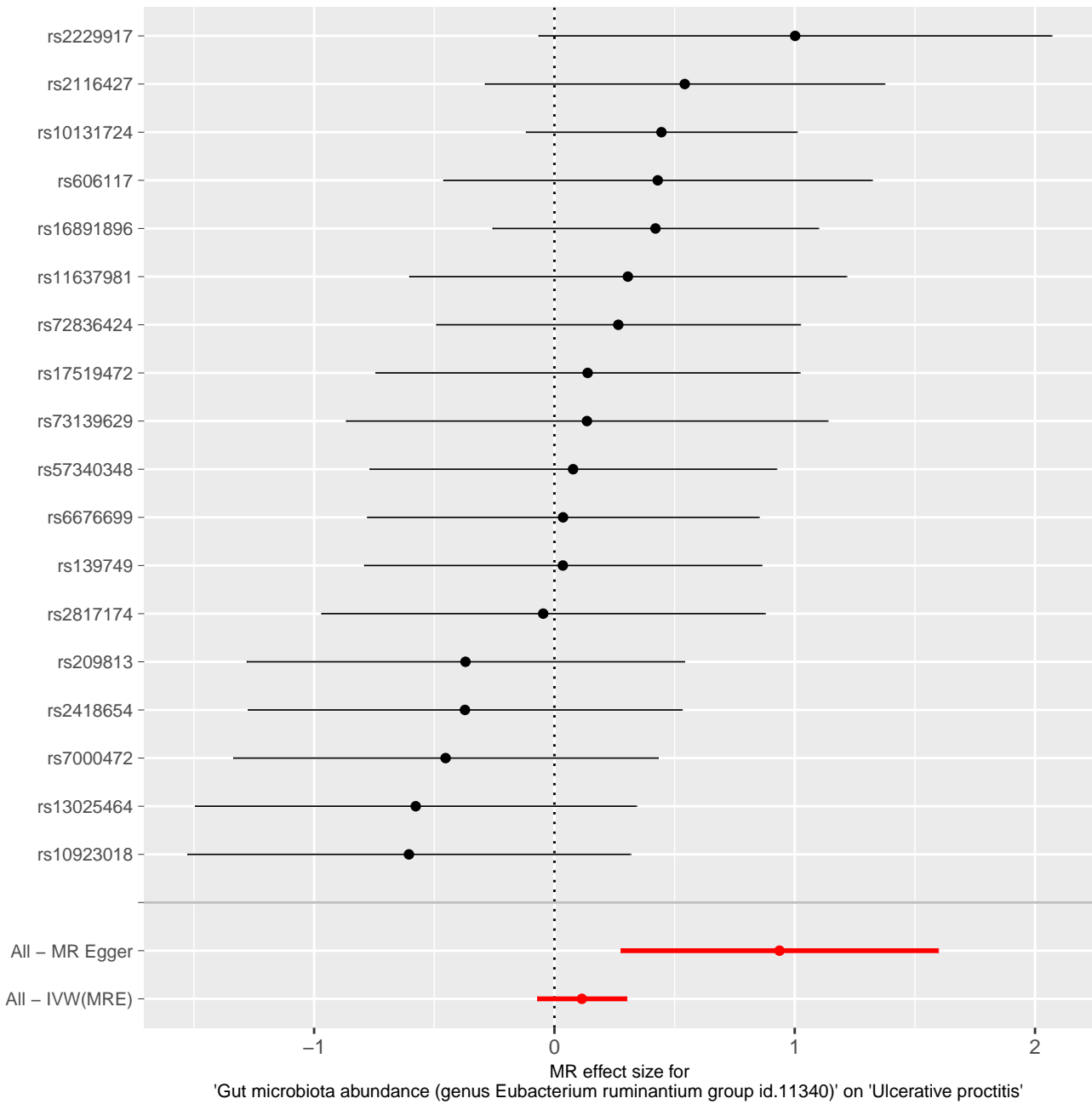


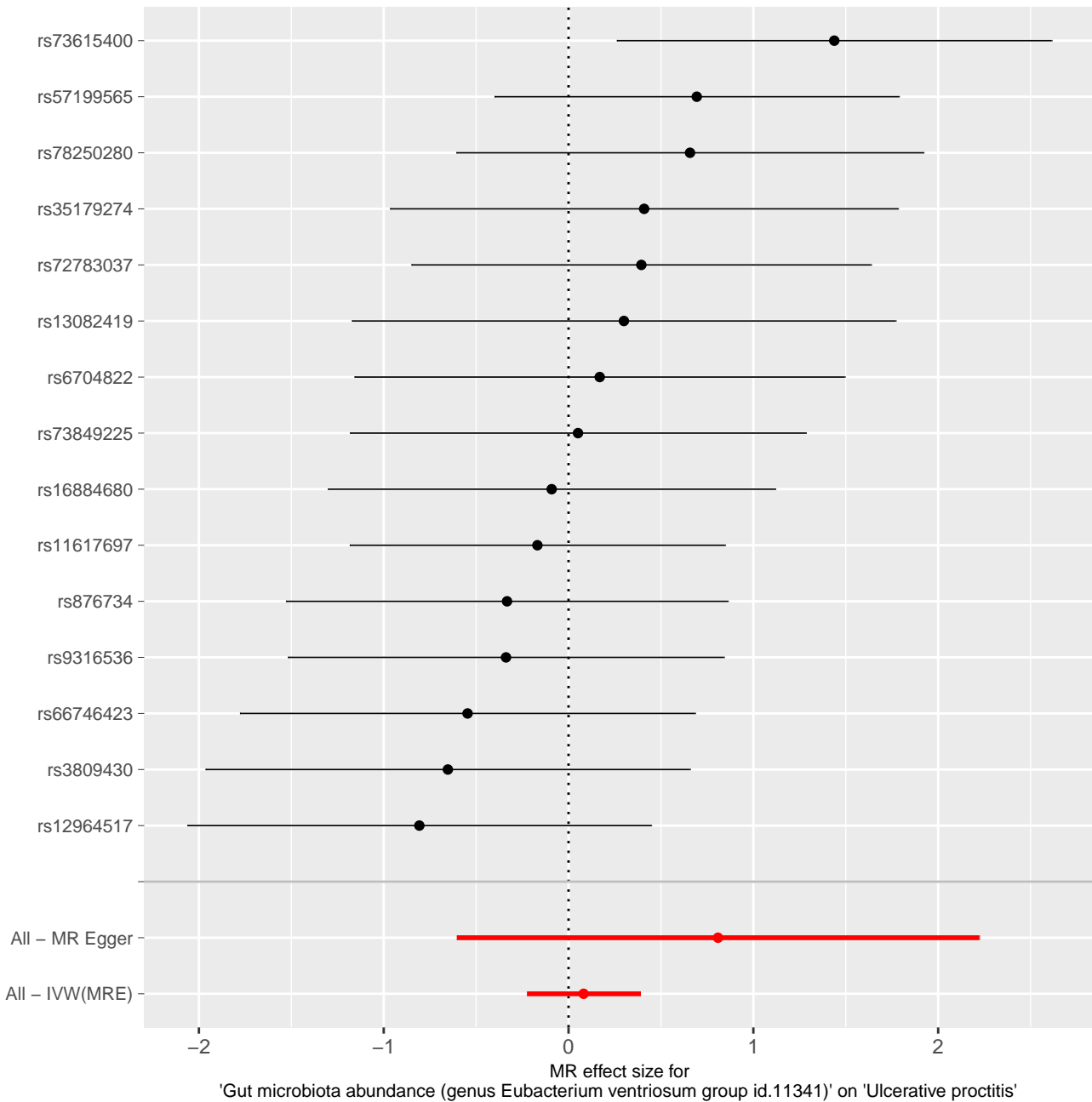


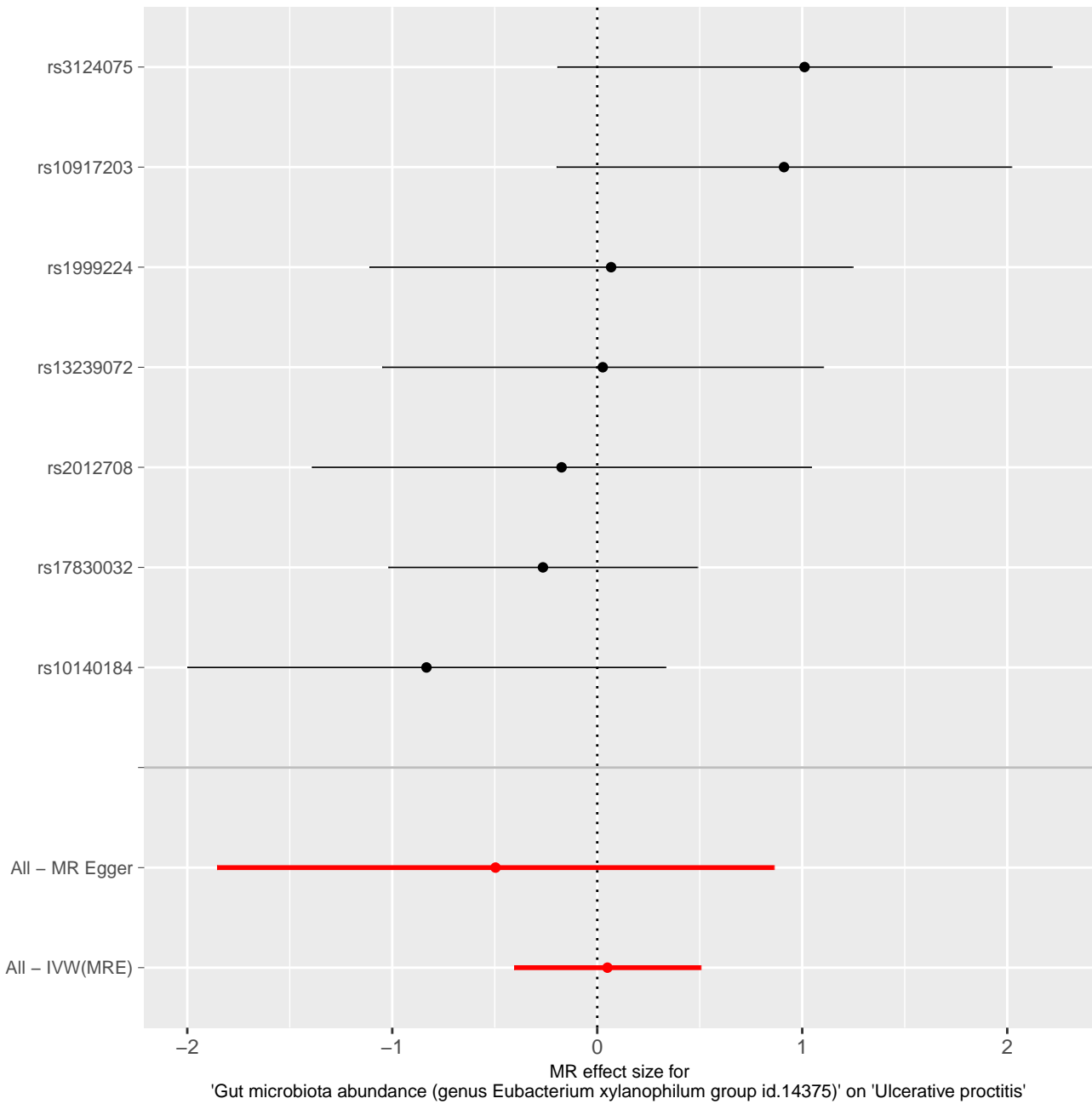


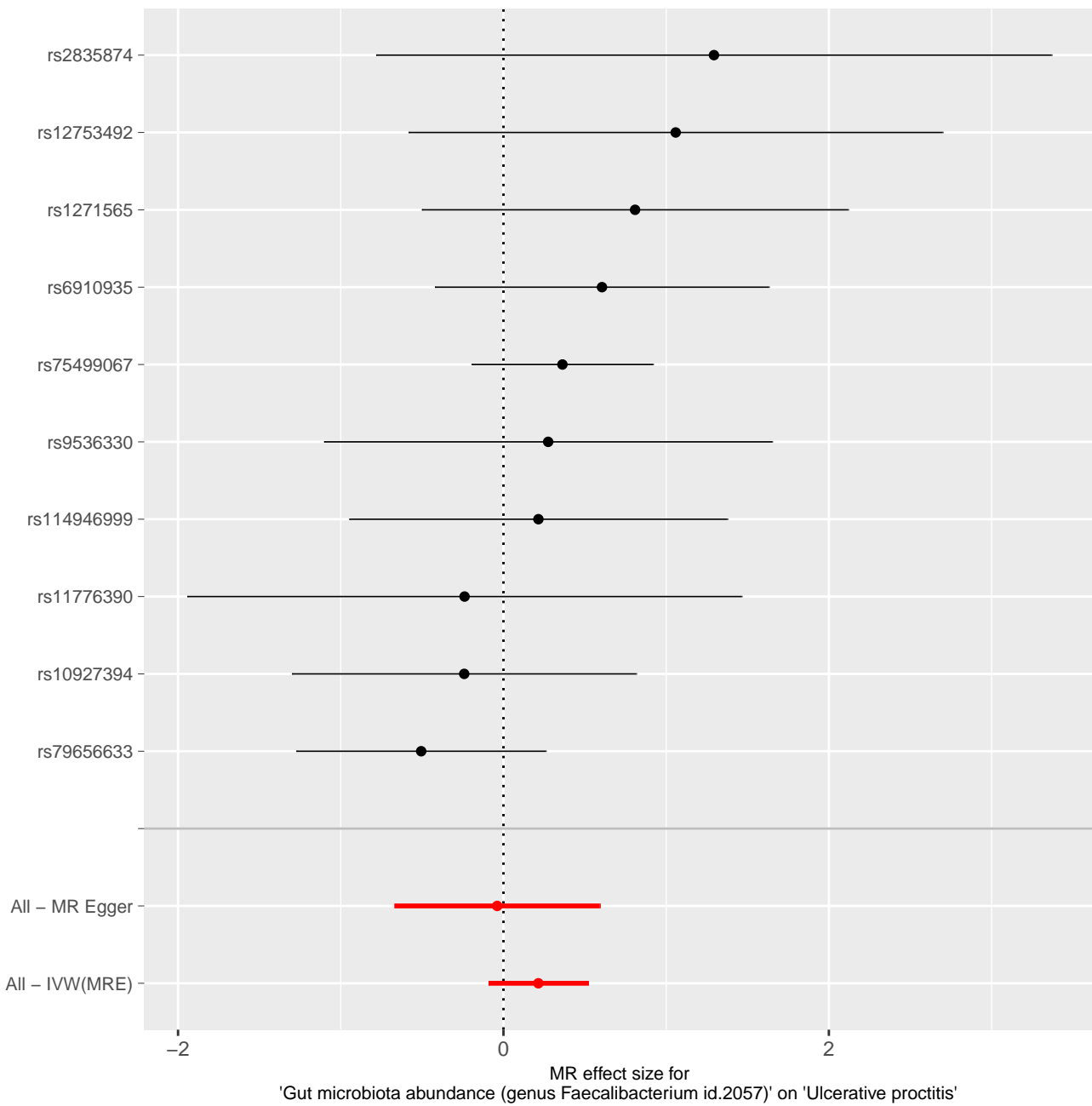


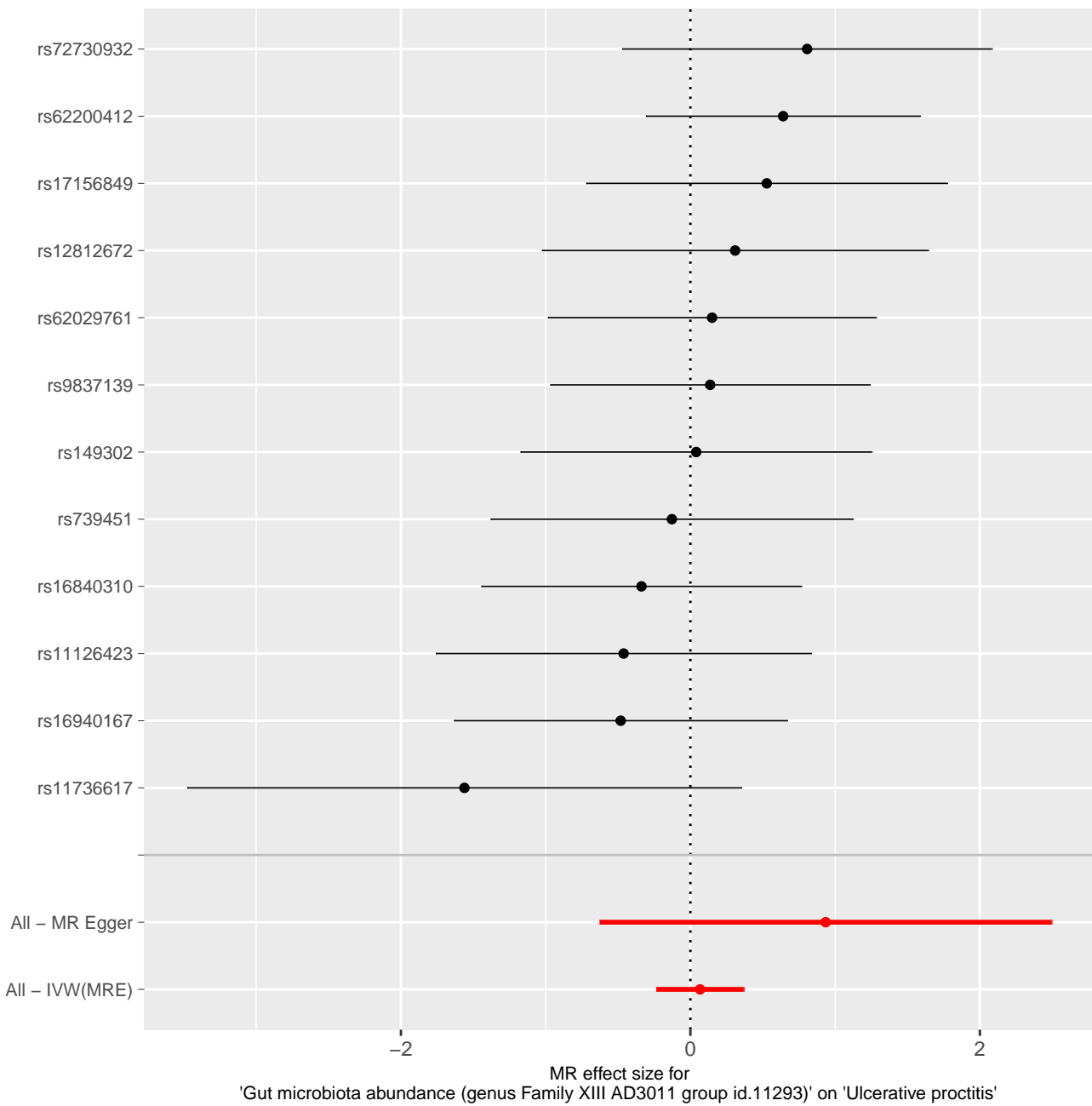


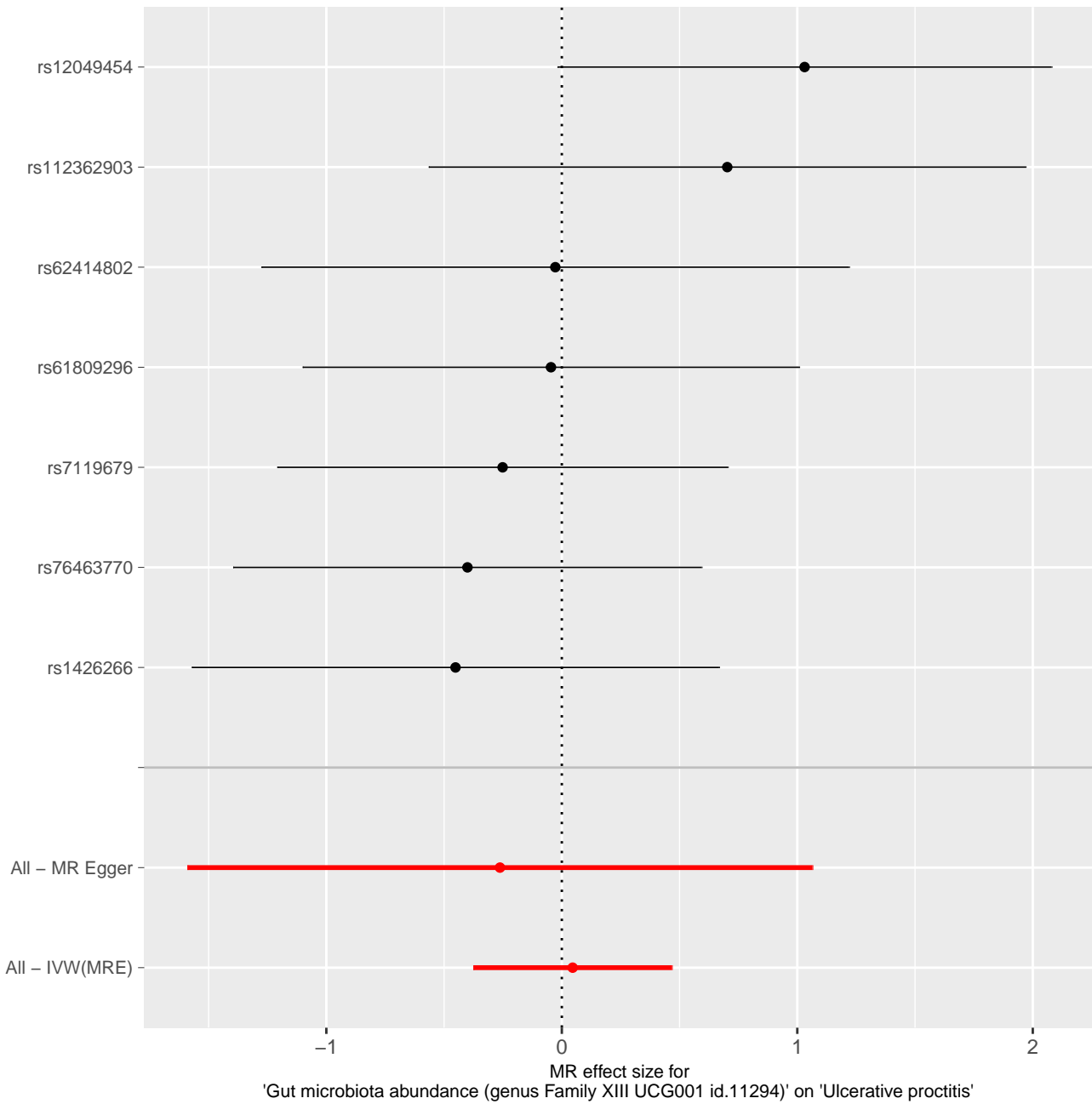


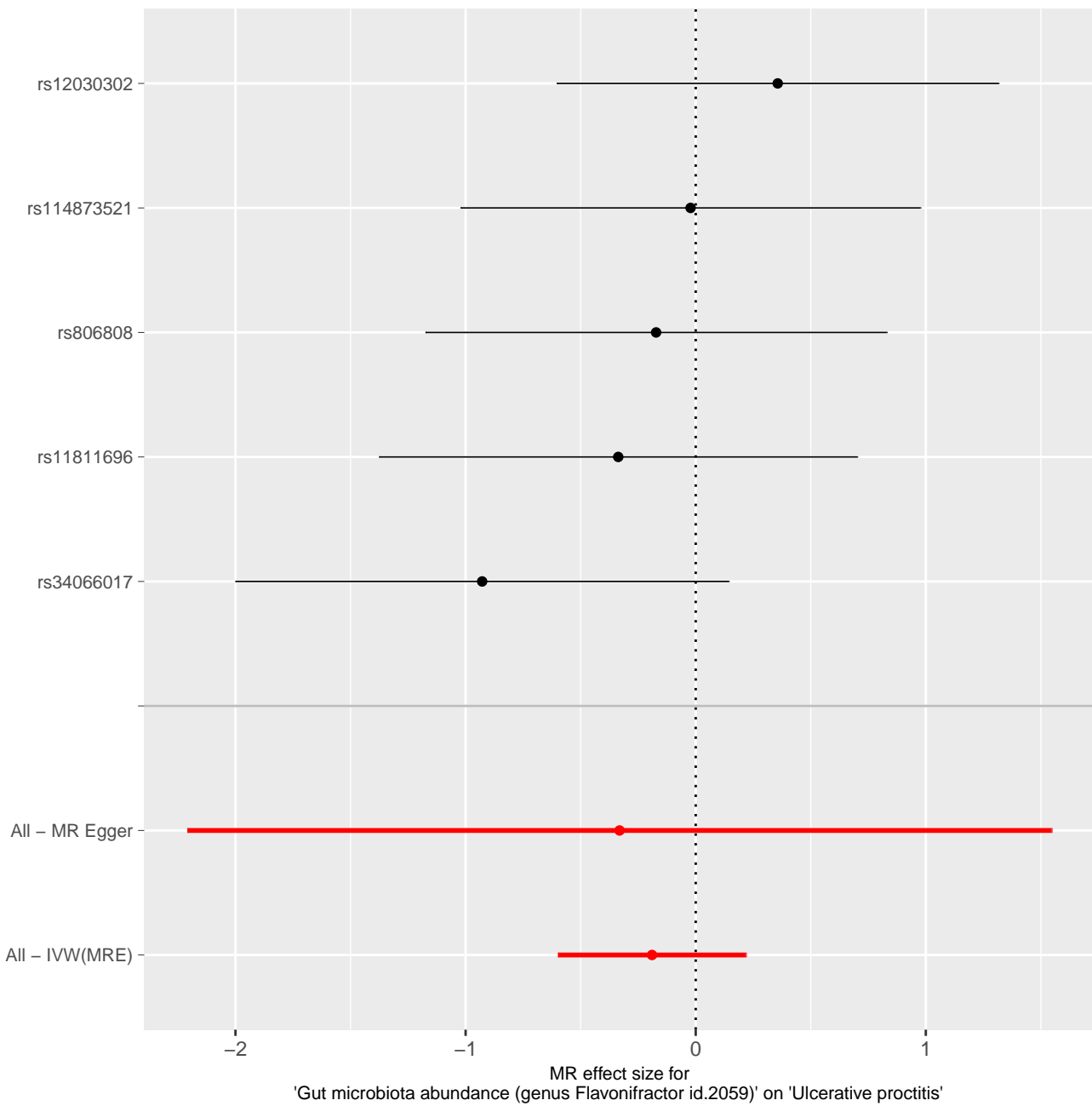




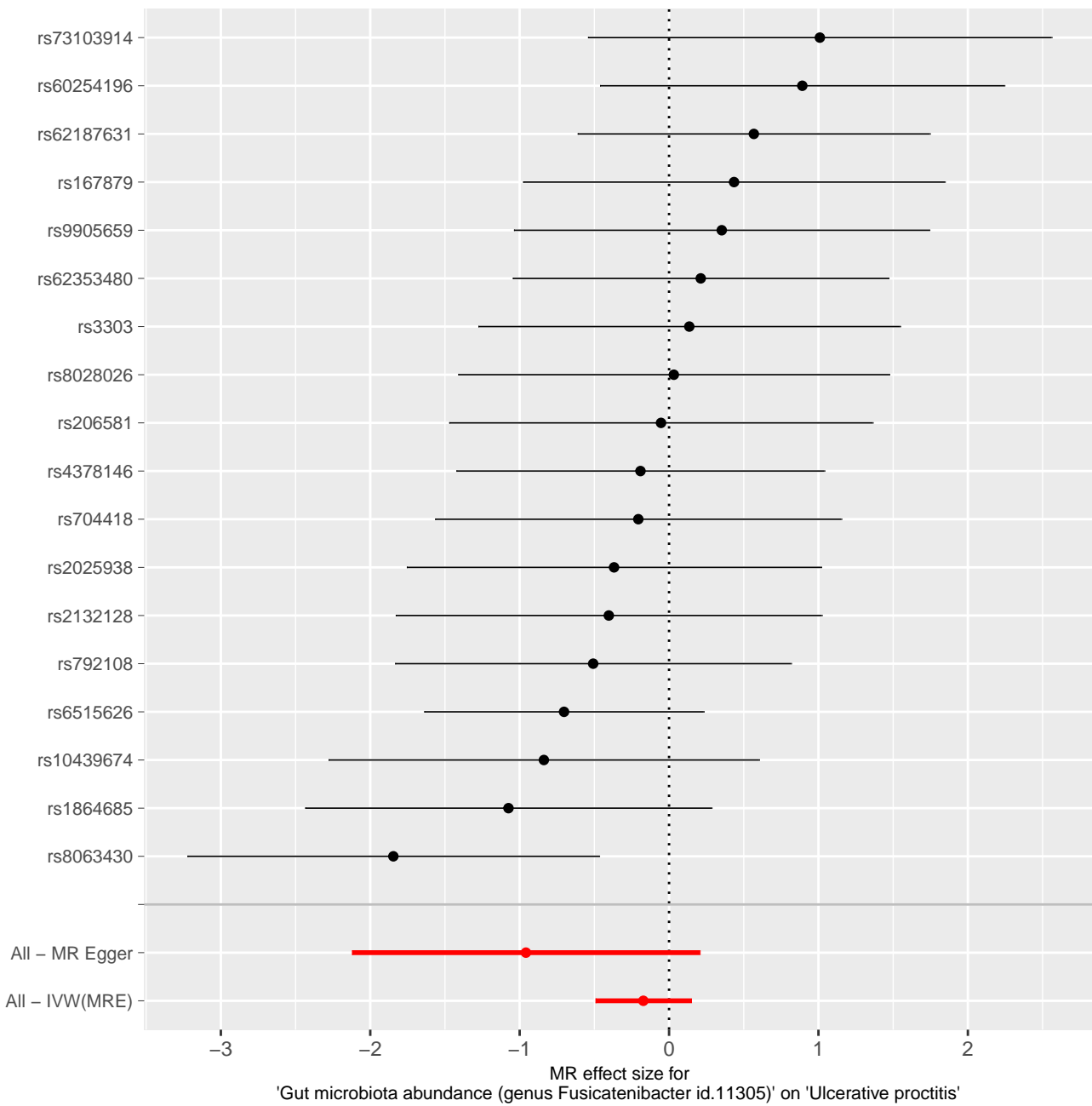


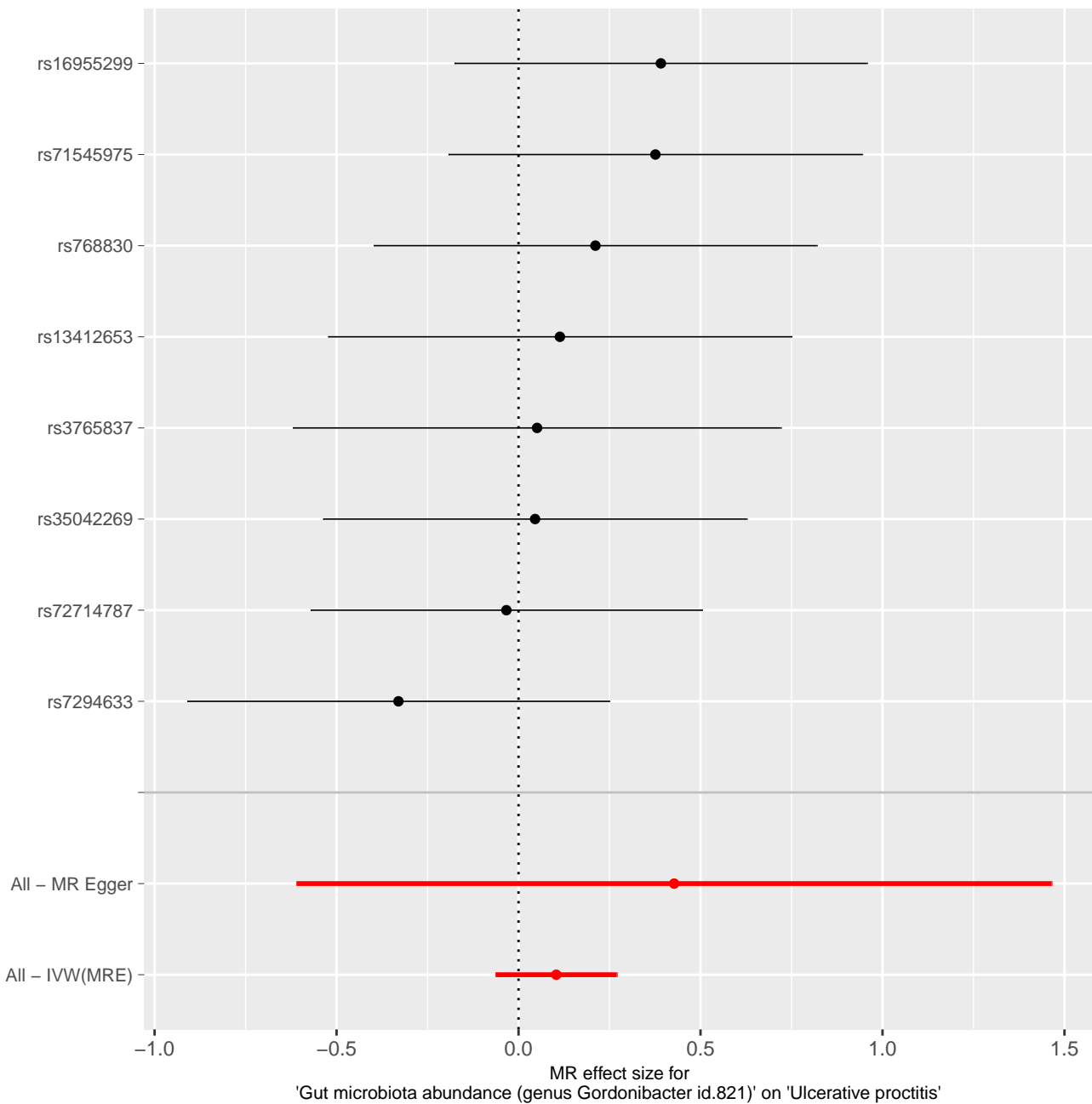


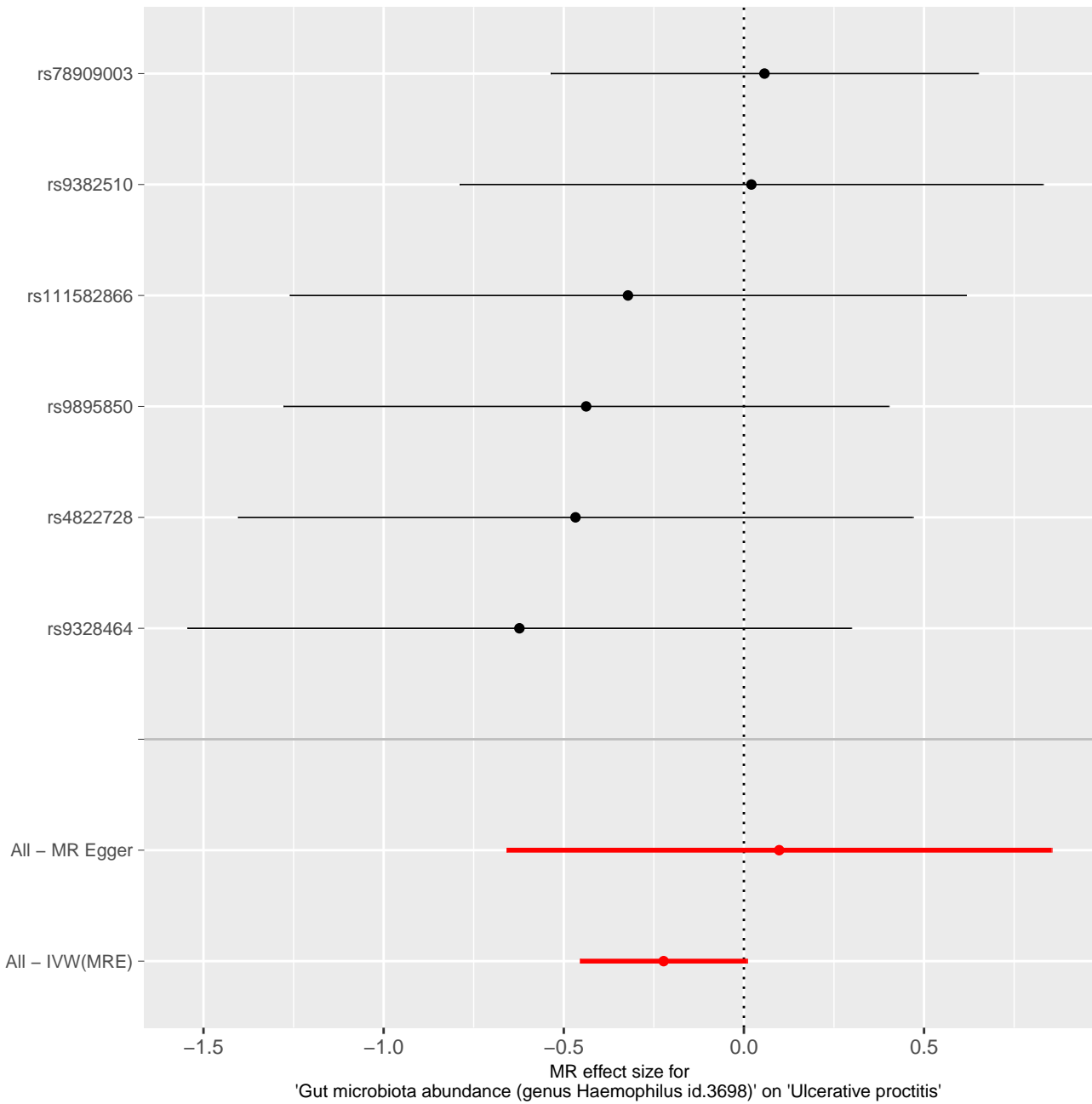


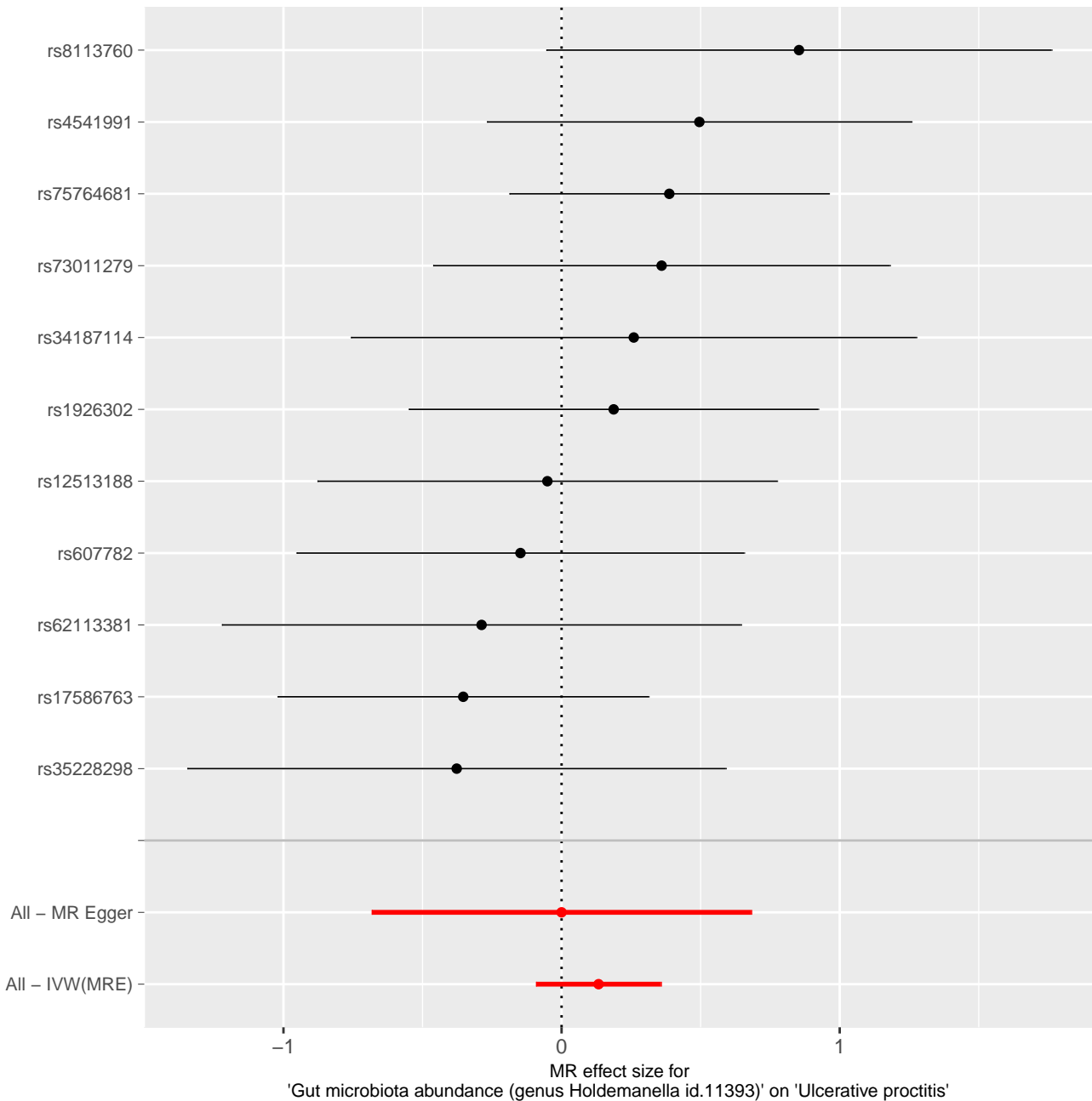


Batch 1435 : Gut microbiota abundance (genus Fusicatenibacter id.11305) on Ulcerative proctitis

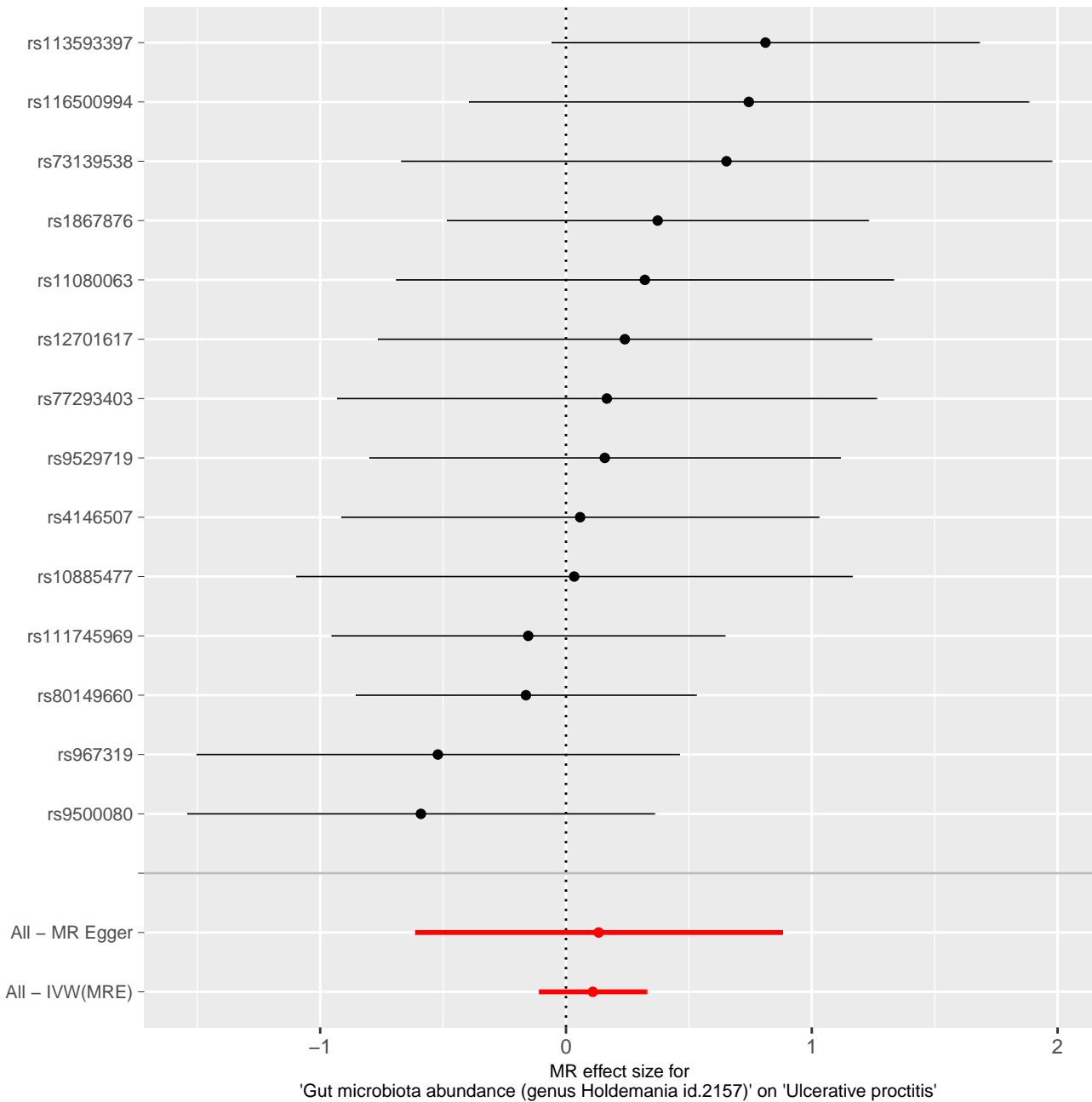


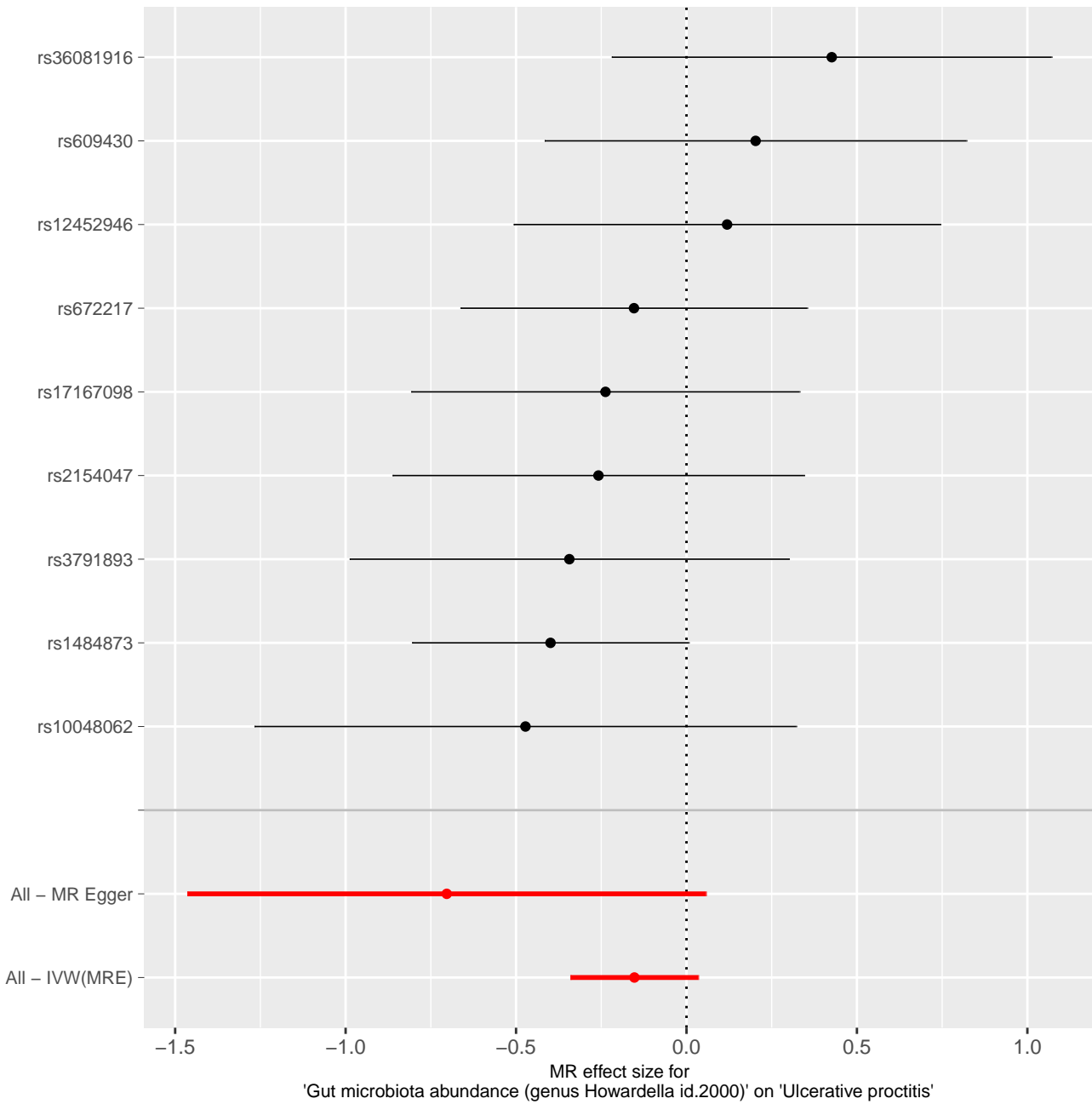


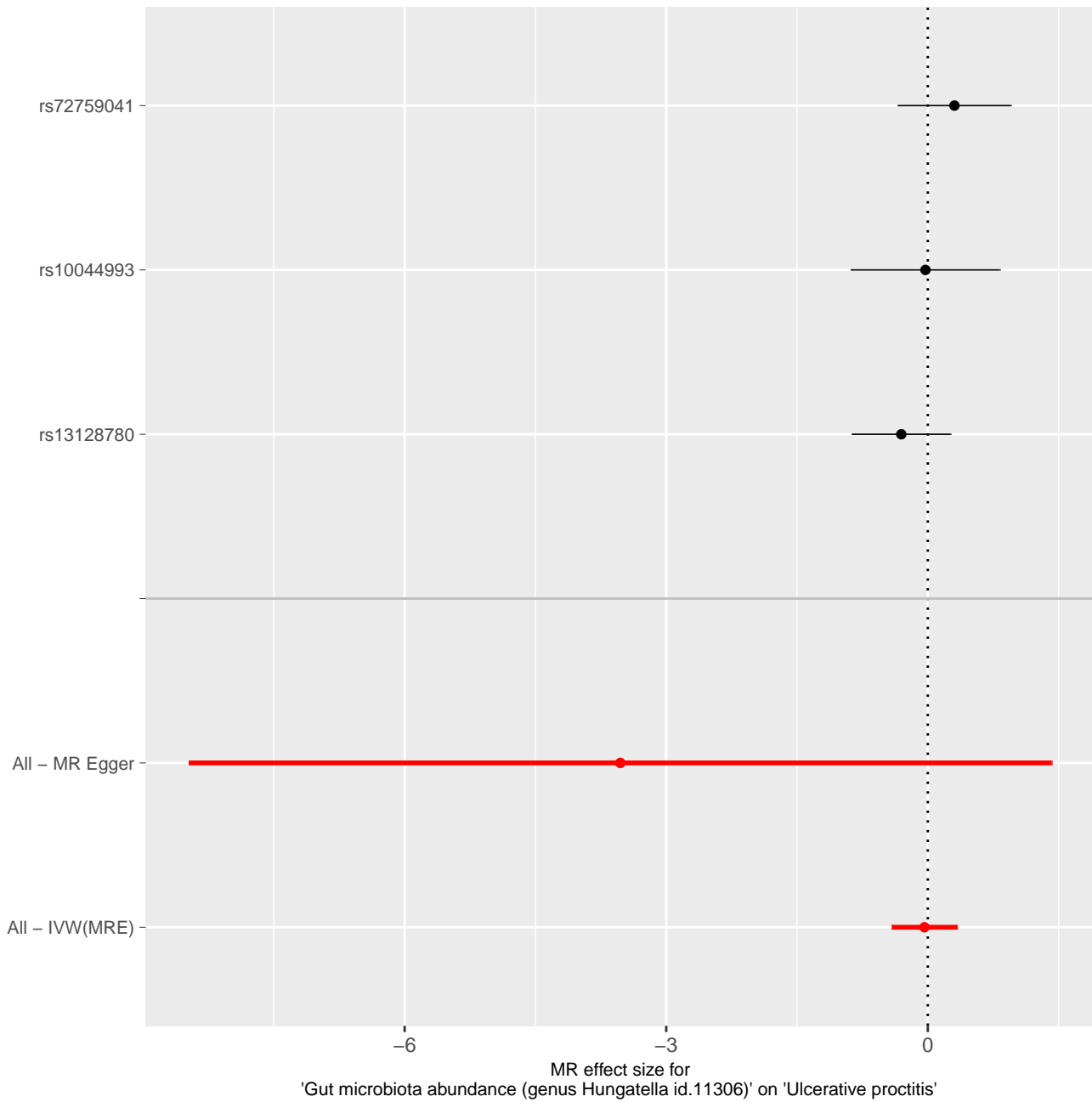


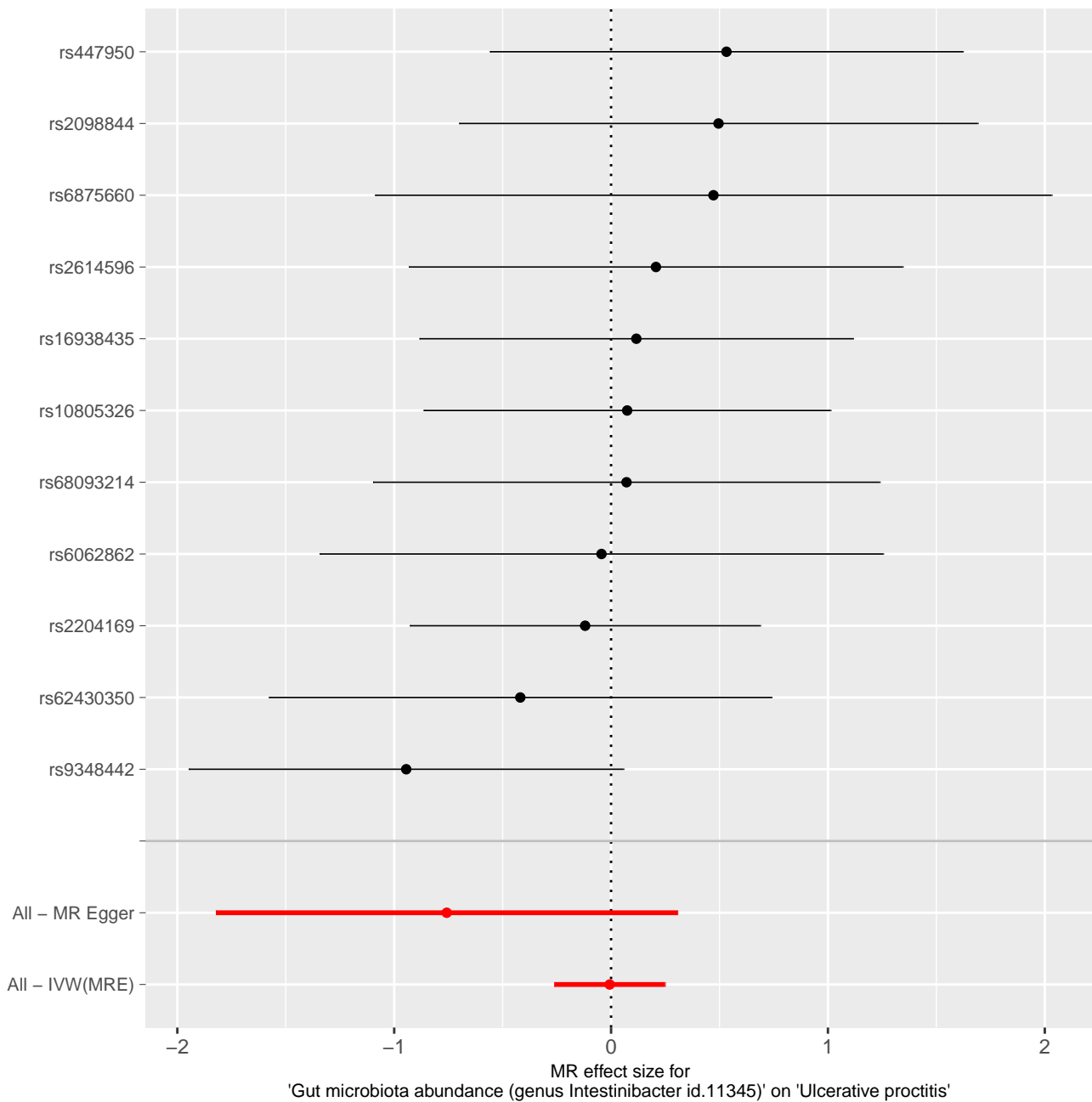


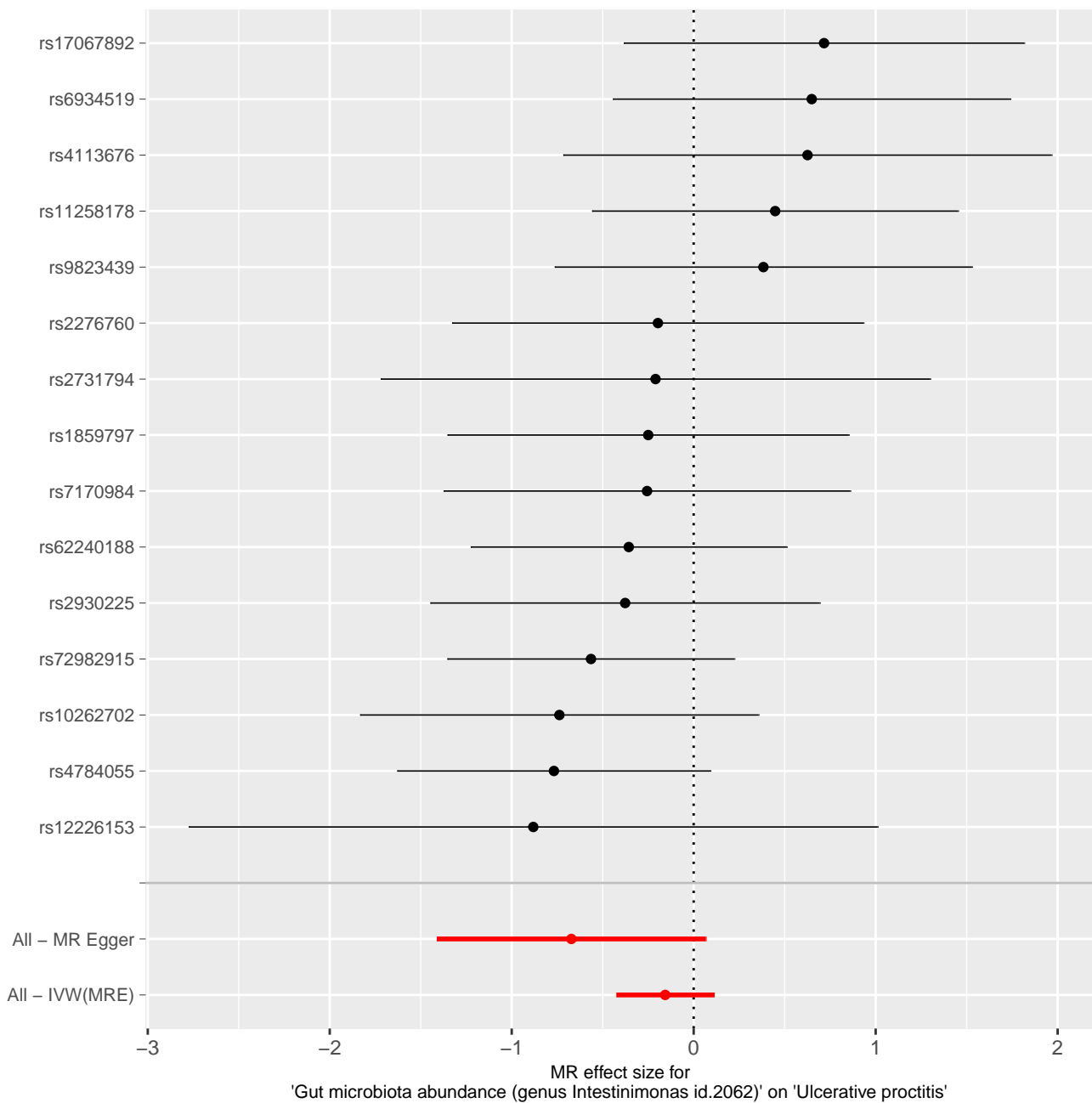
Batch 1439 : Gut microbiota abundance (genus Holdemania id.2157) on Ulcerative proctitis

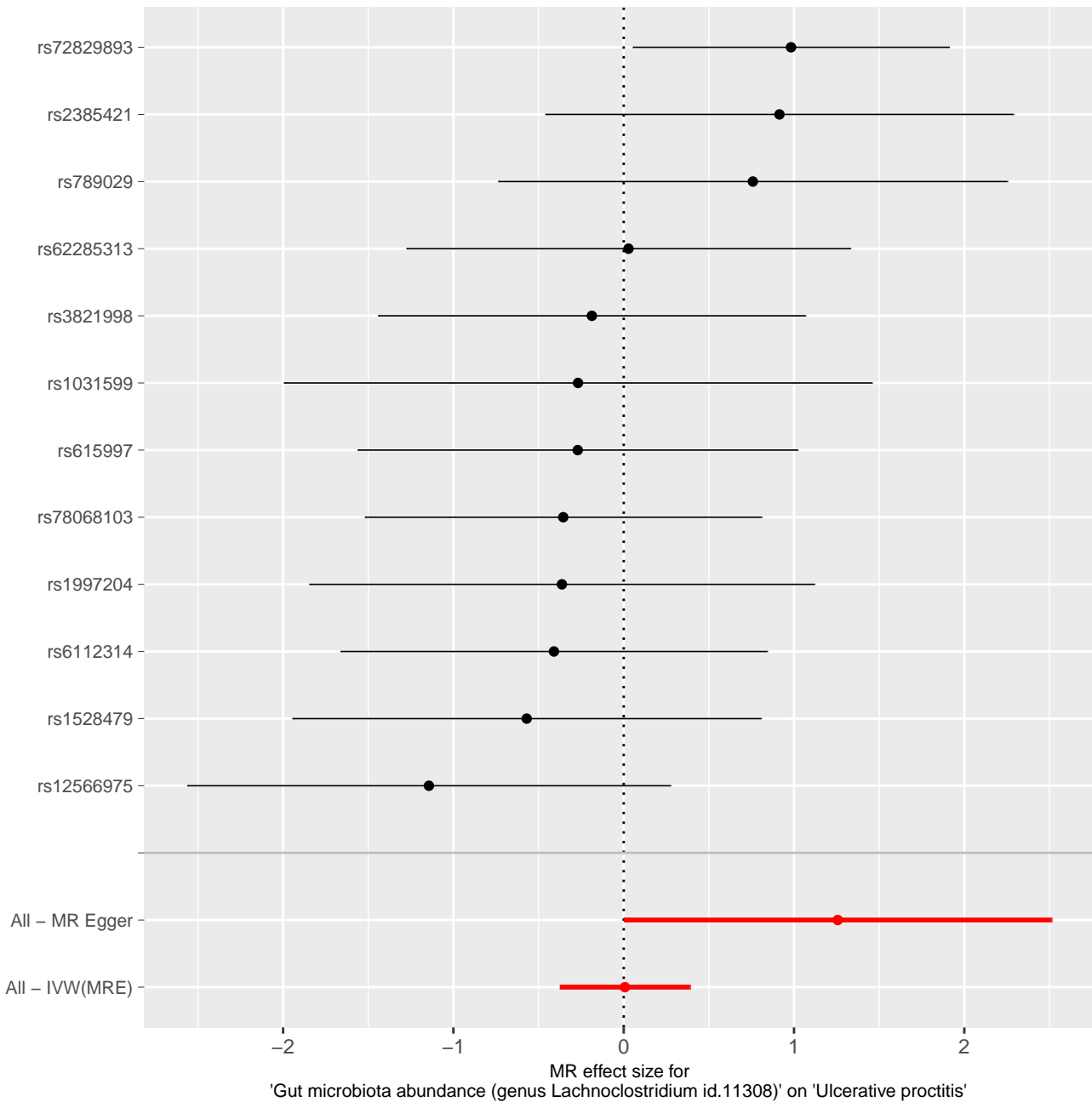


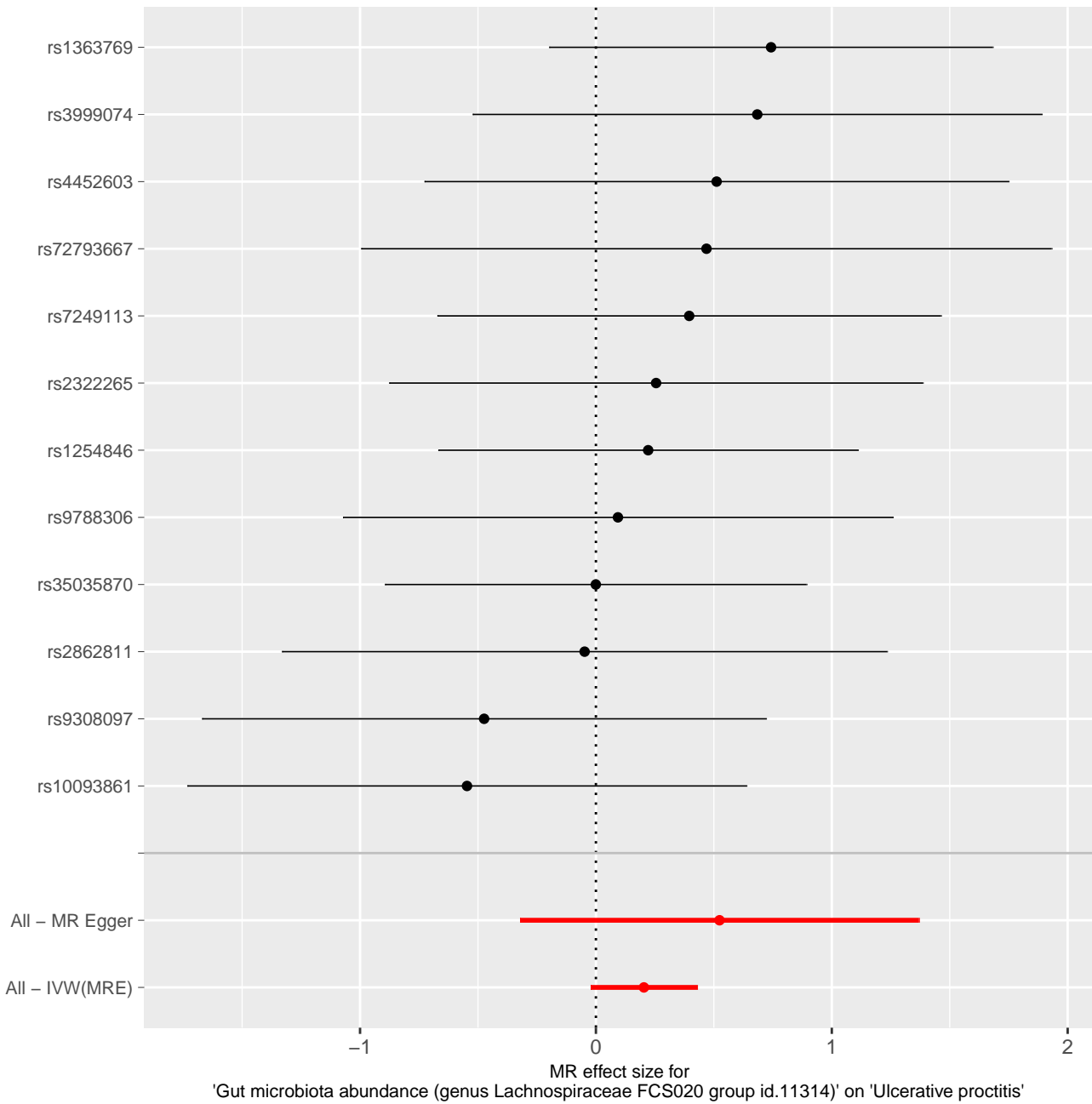


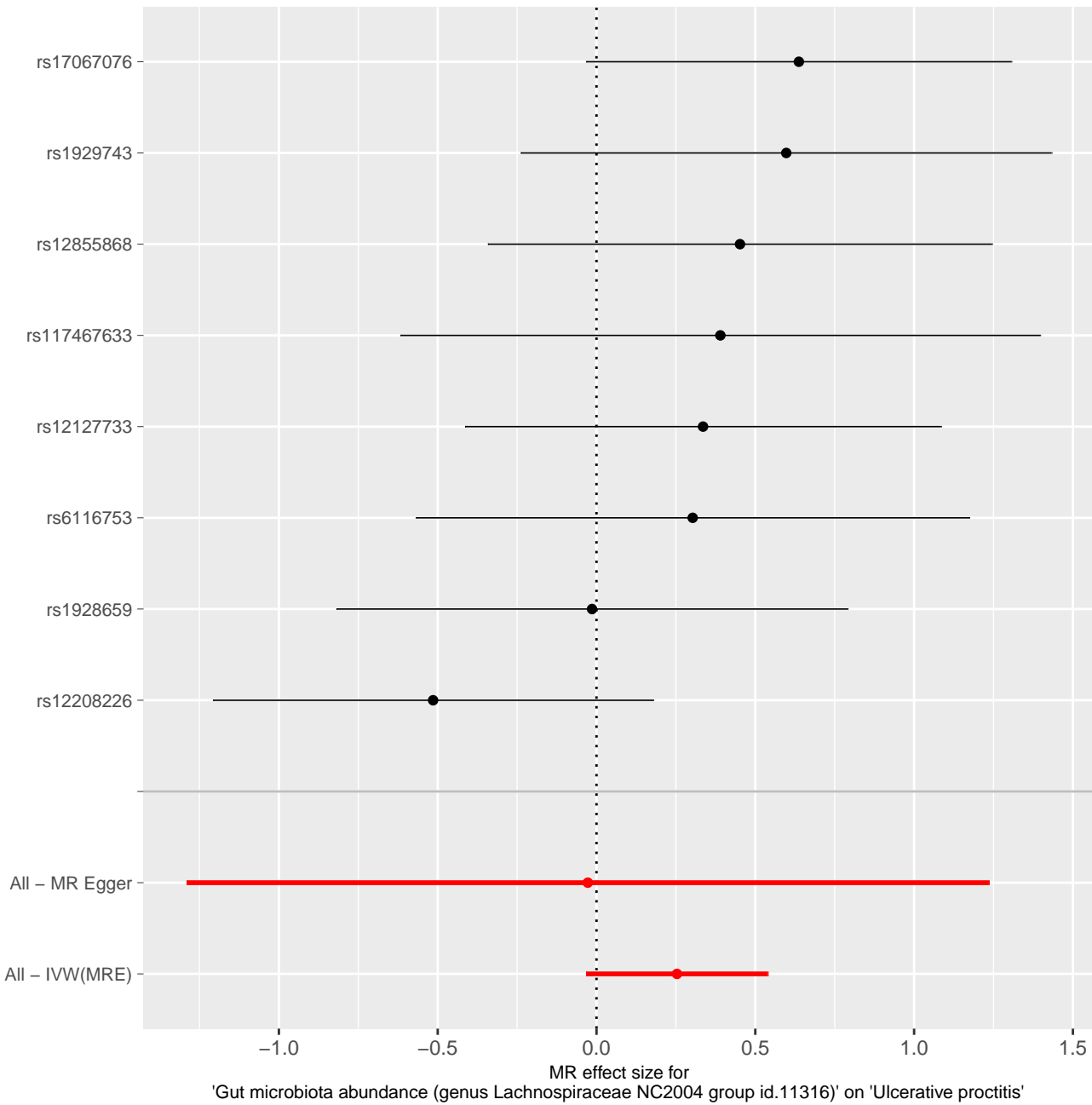


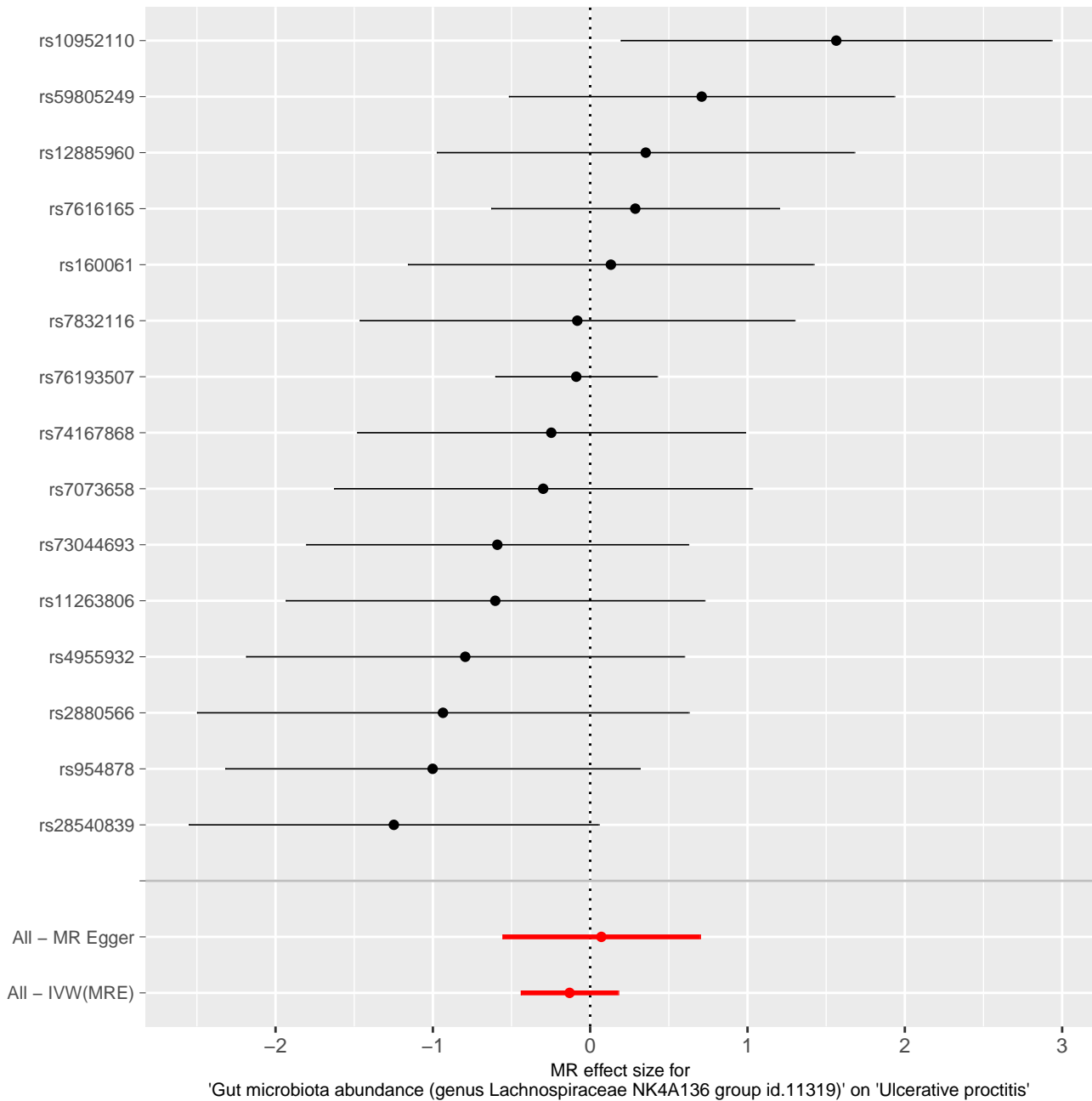


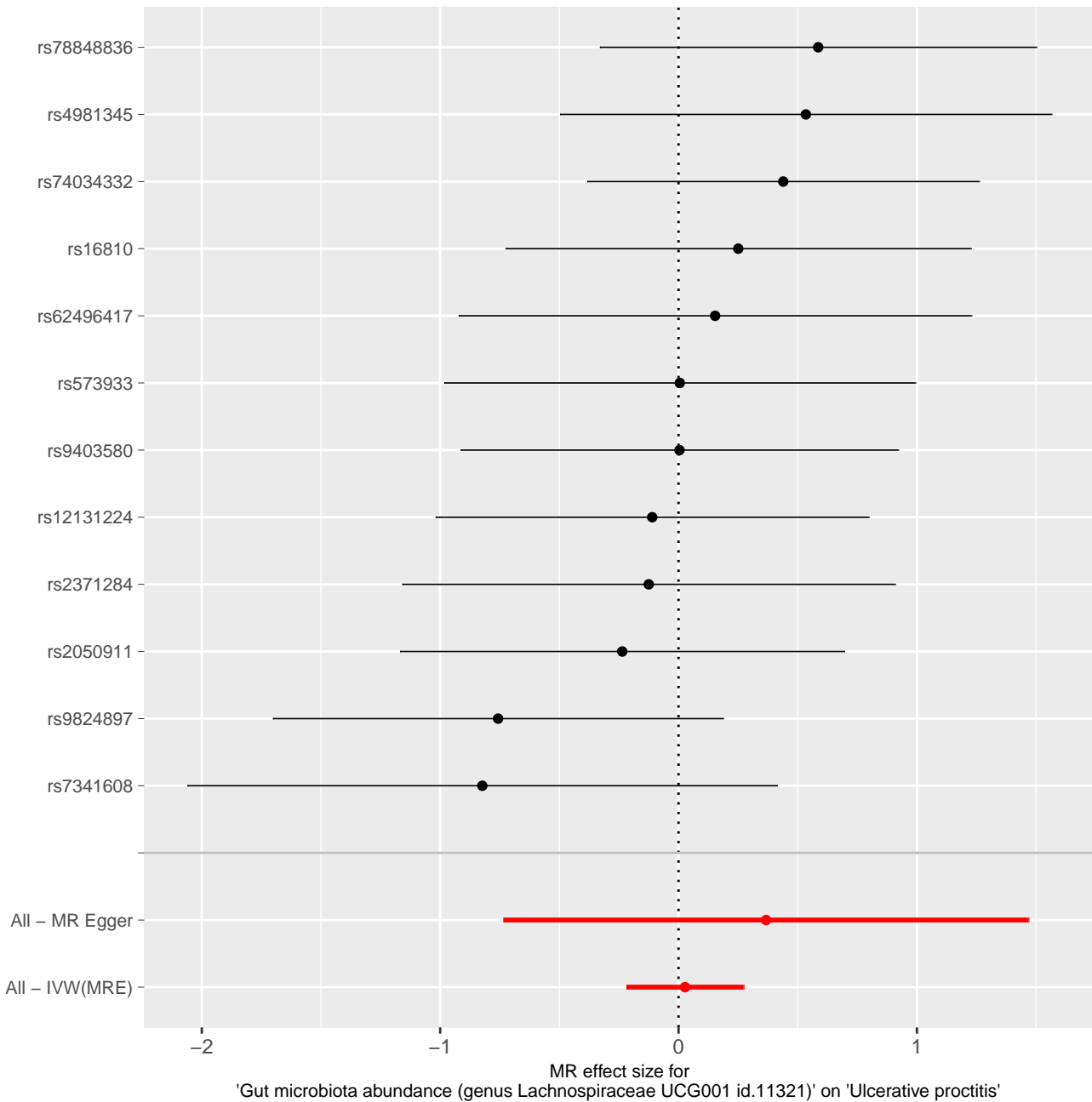


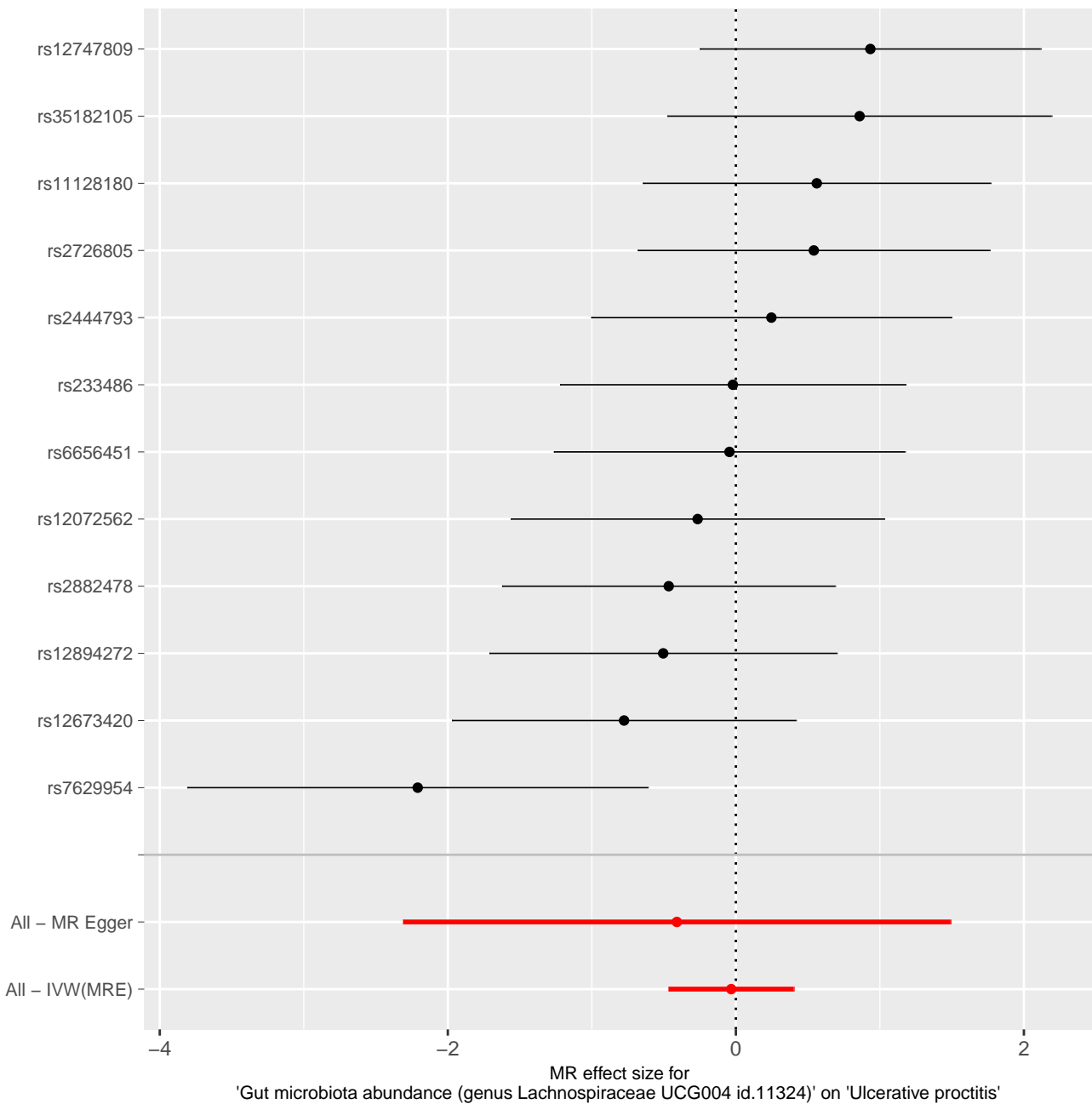


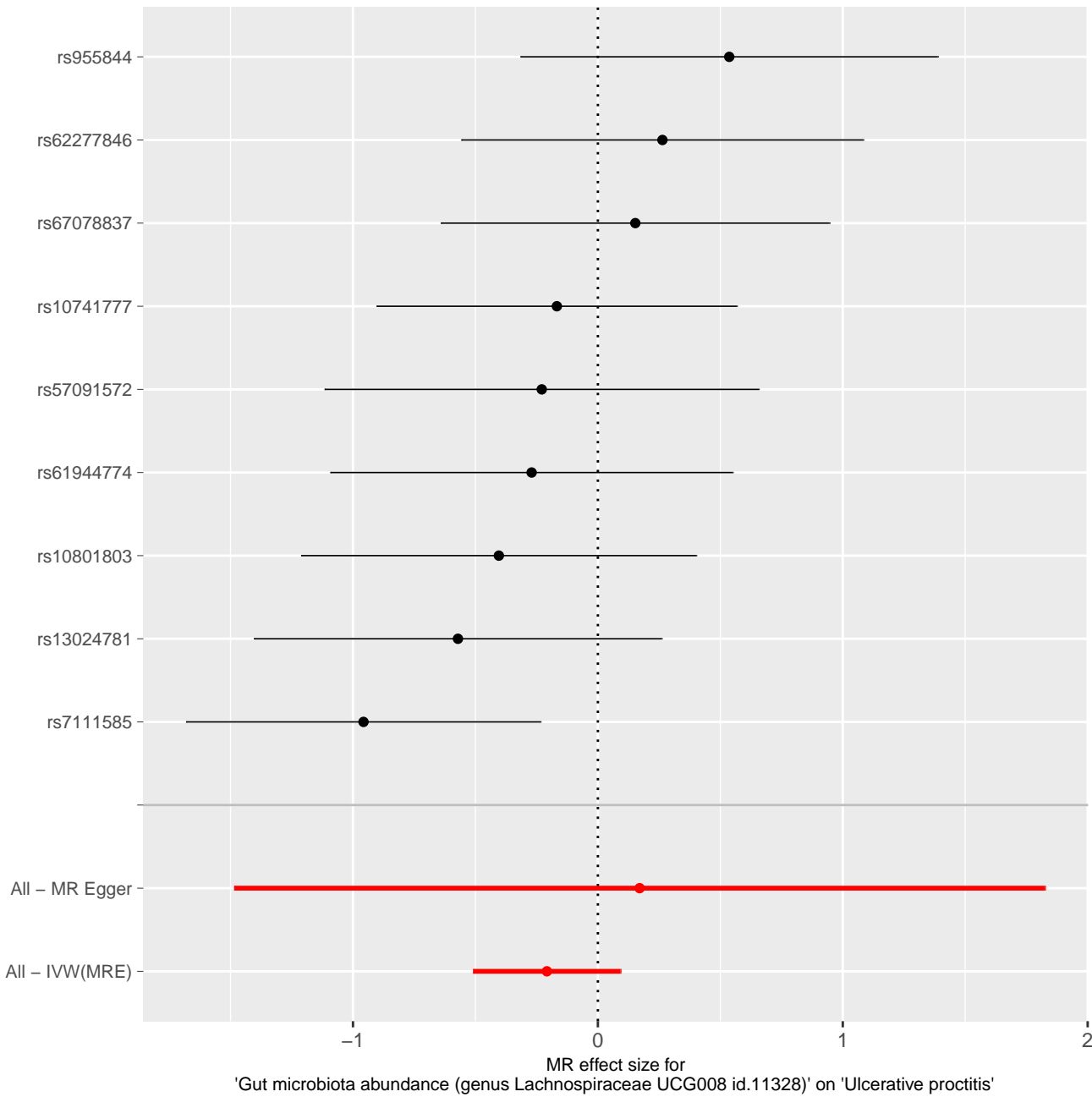


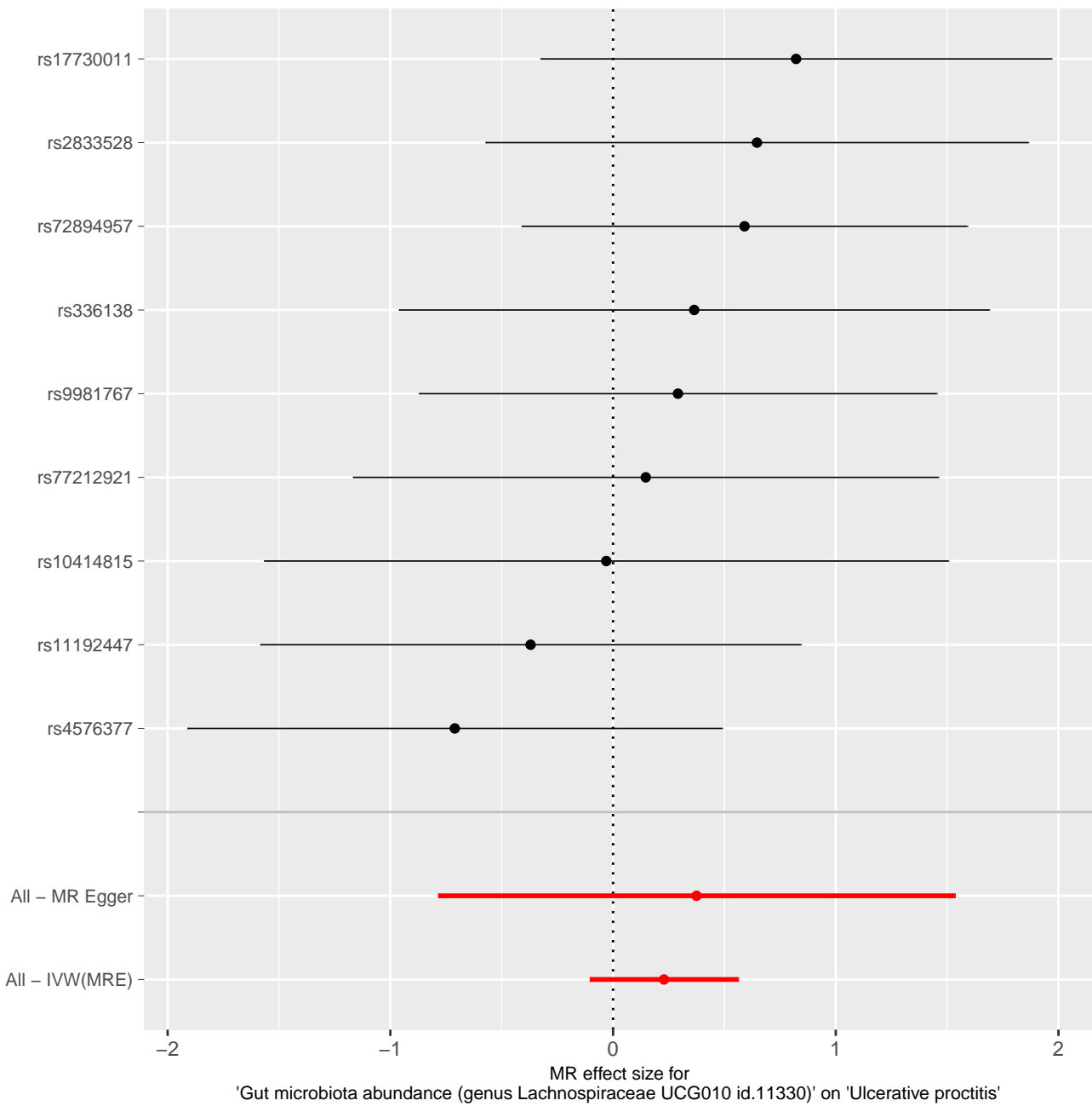


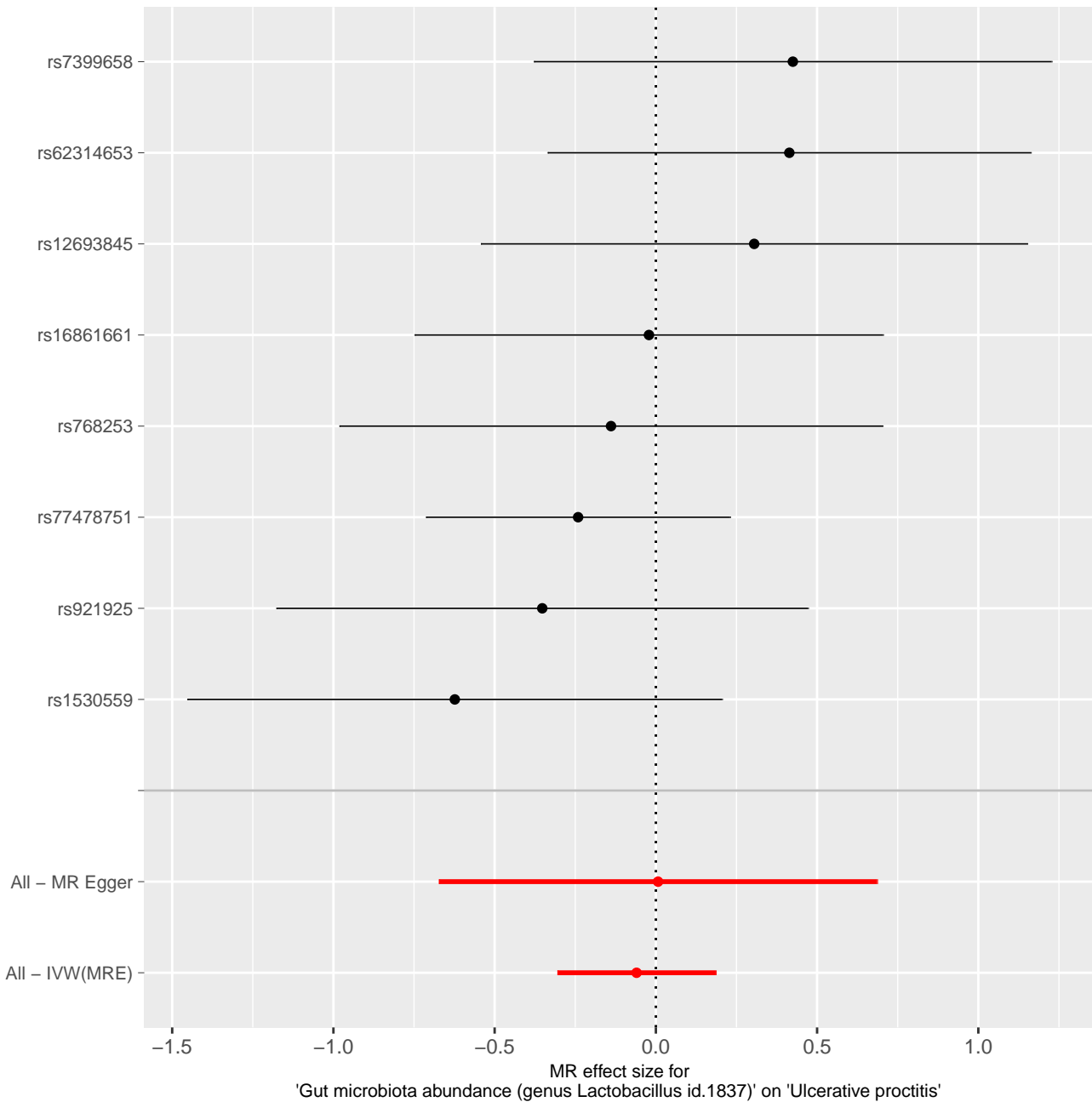


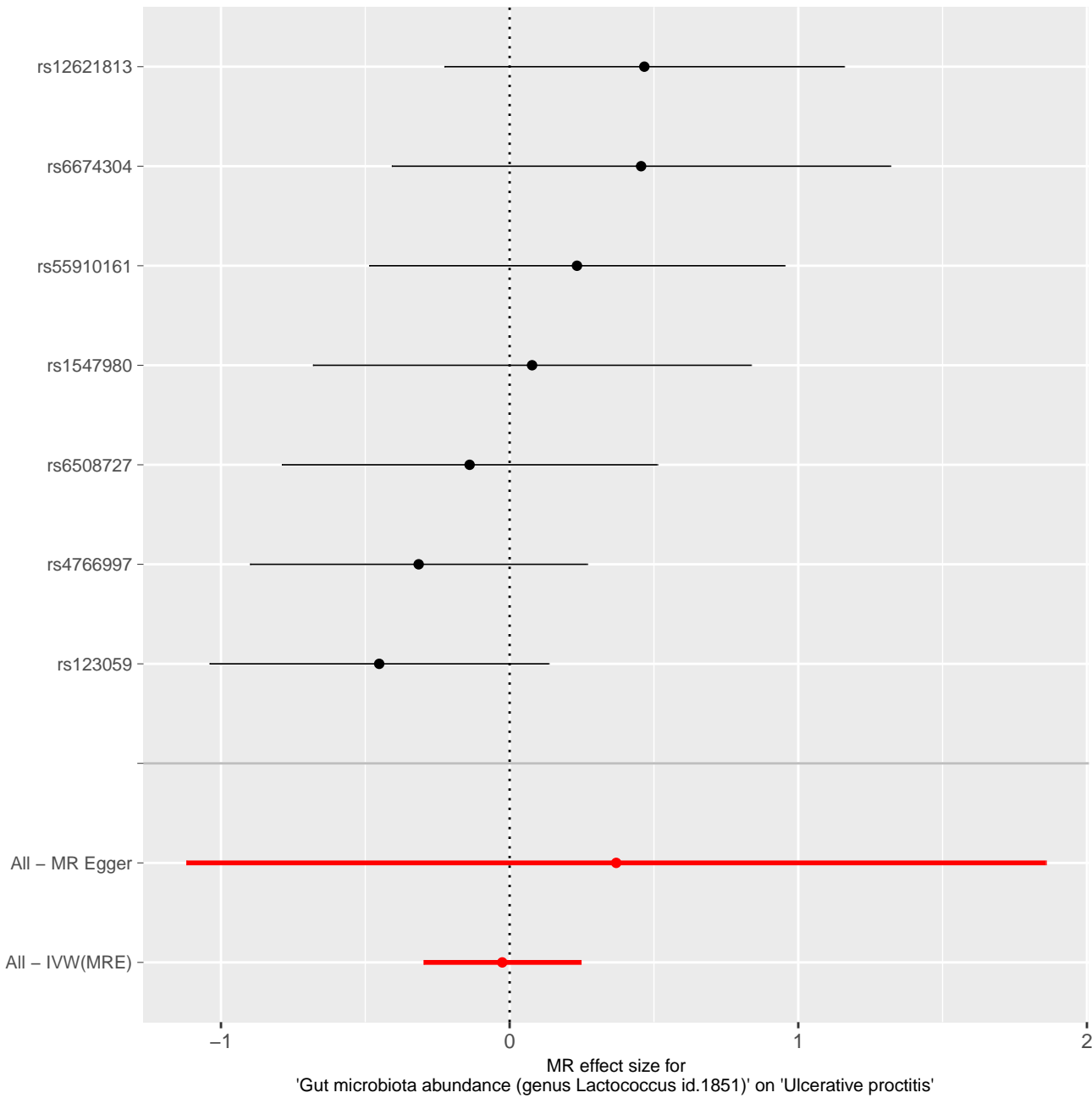


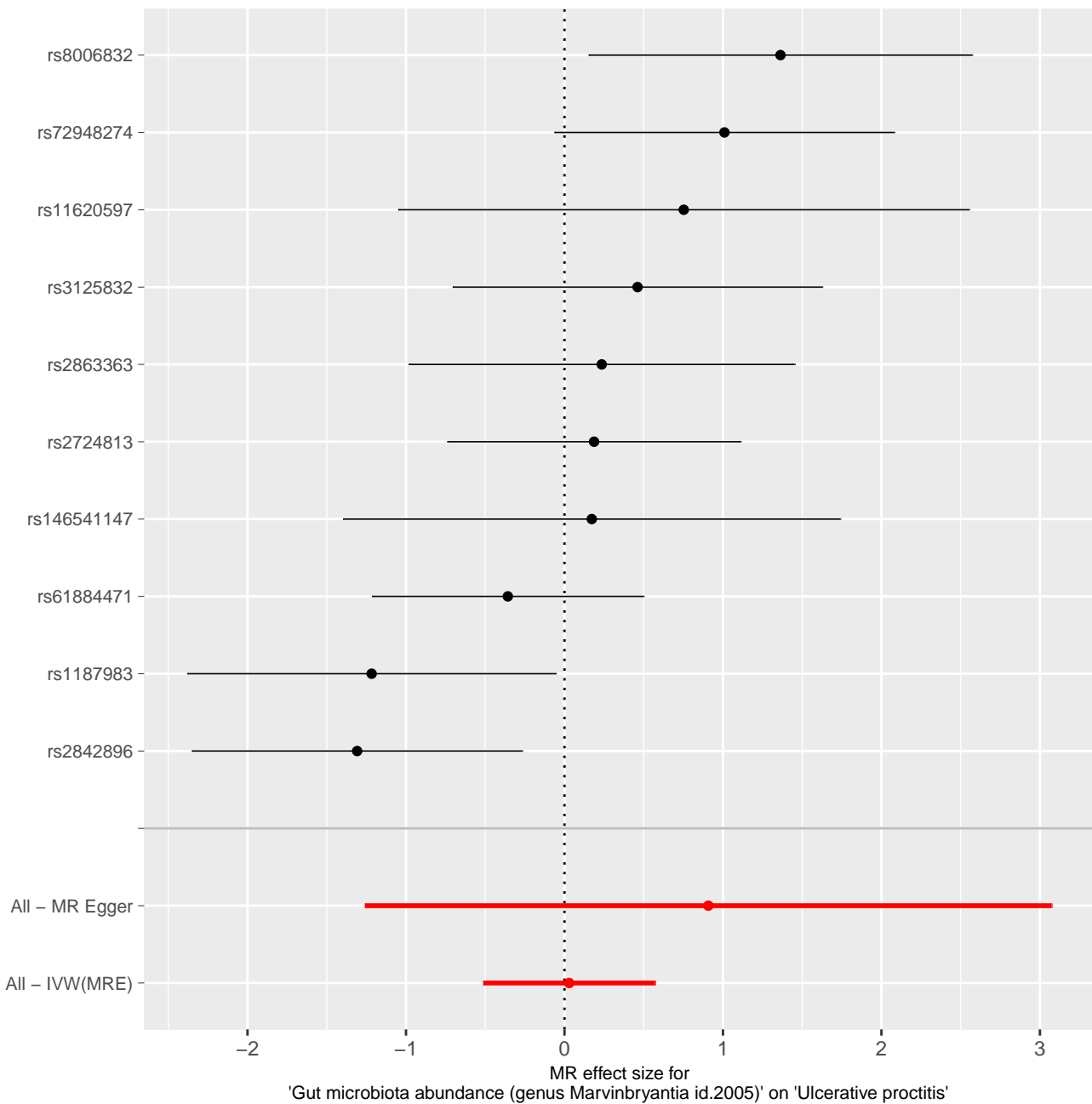


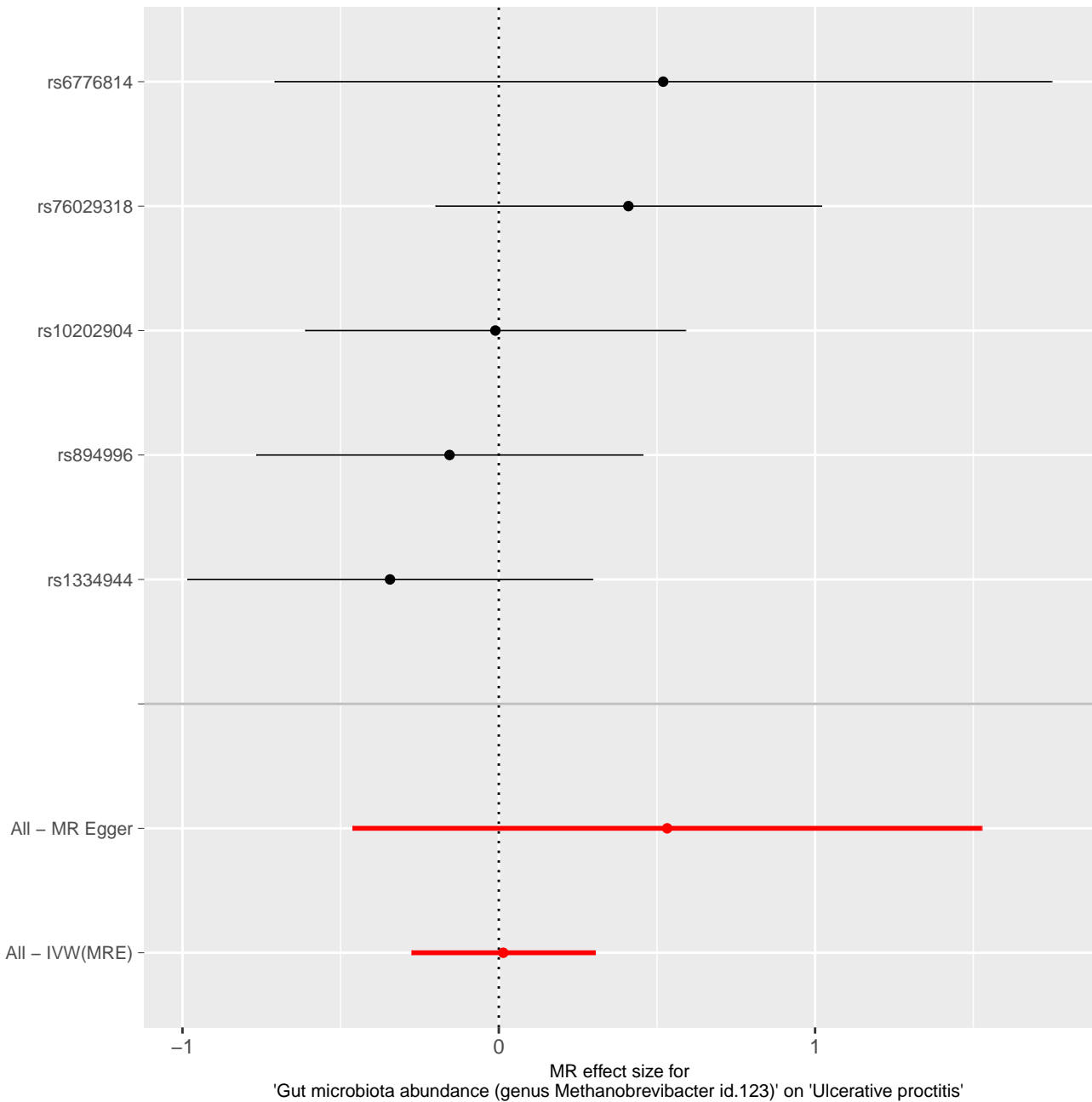


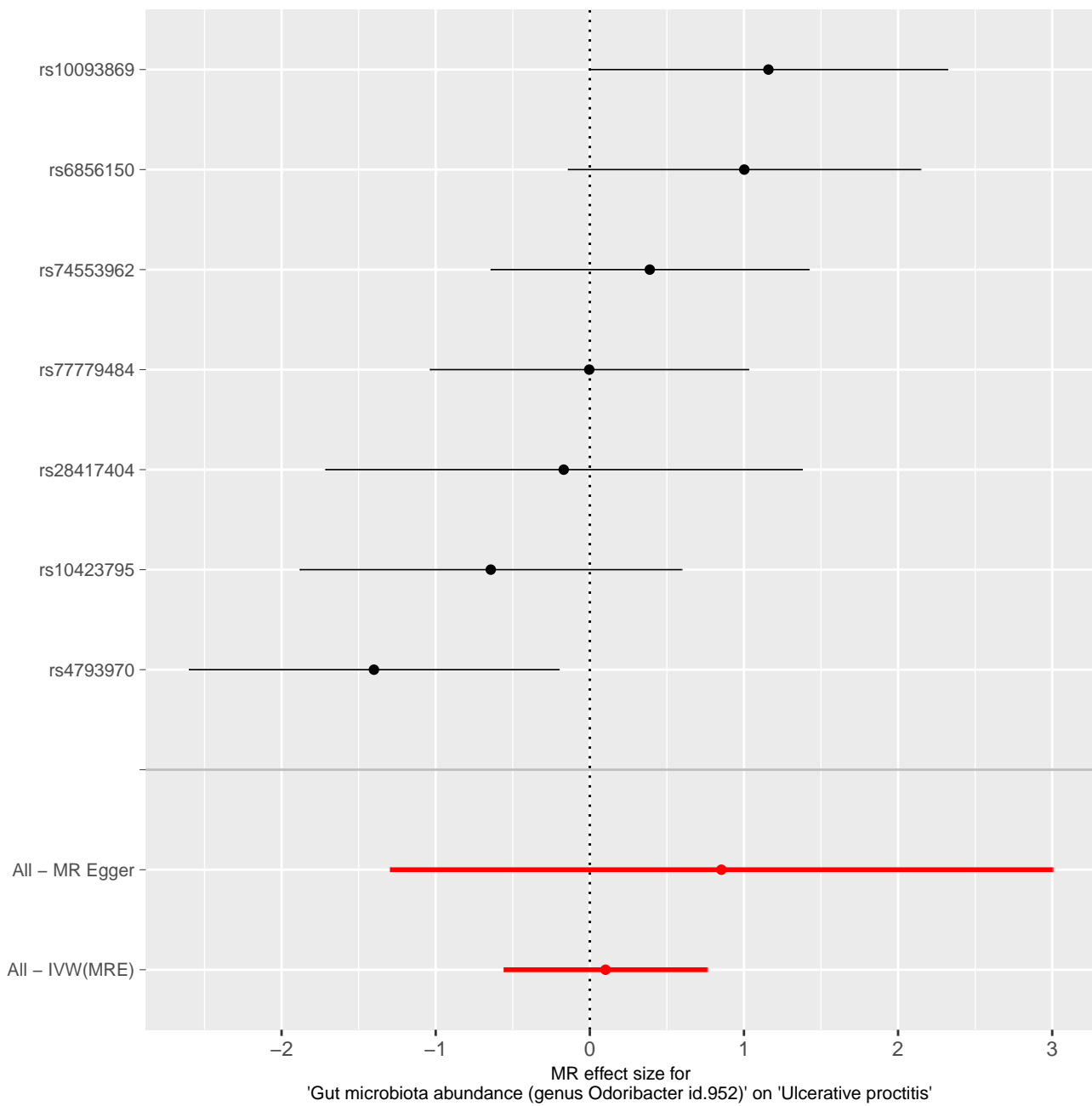


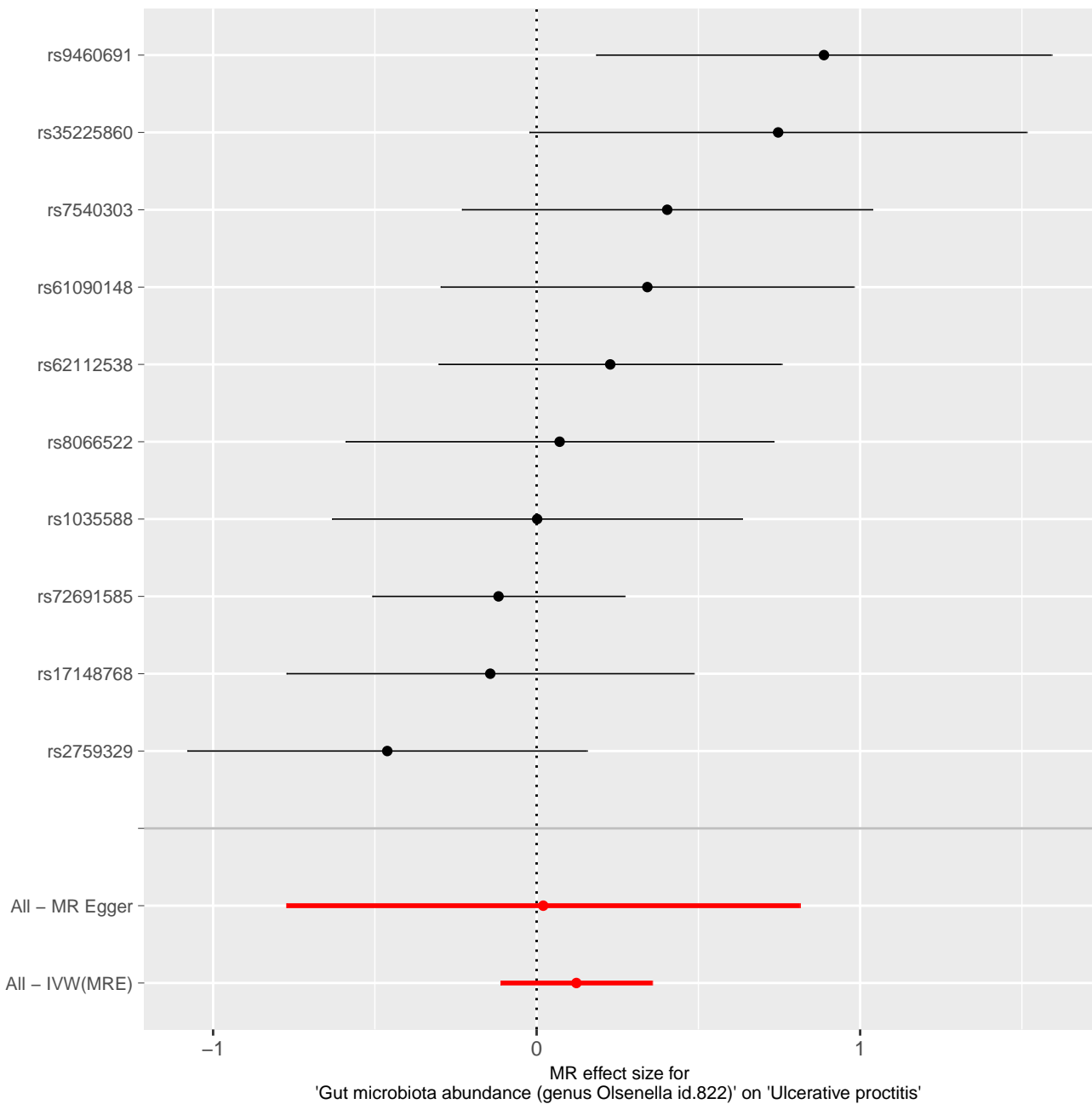




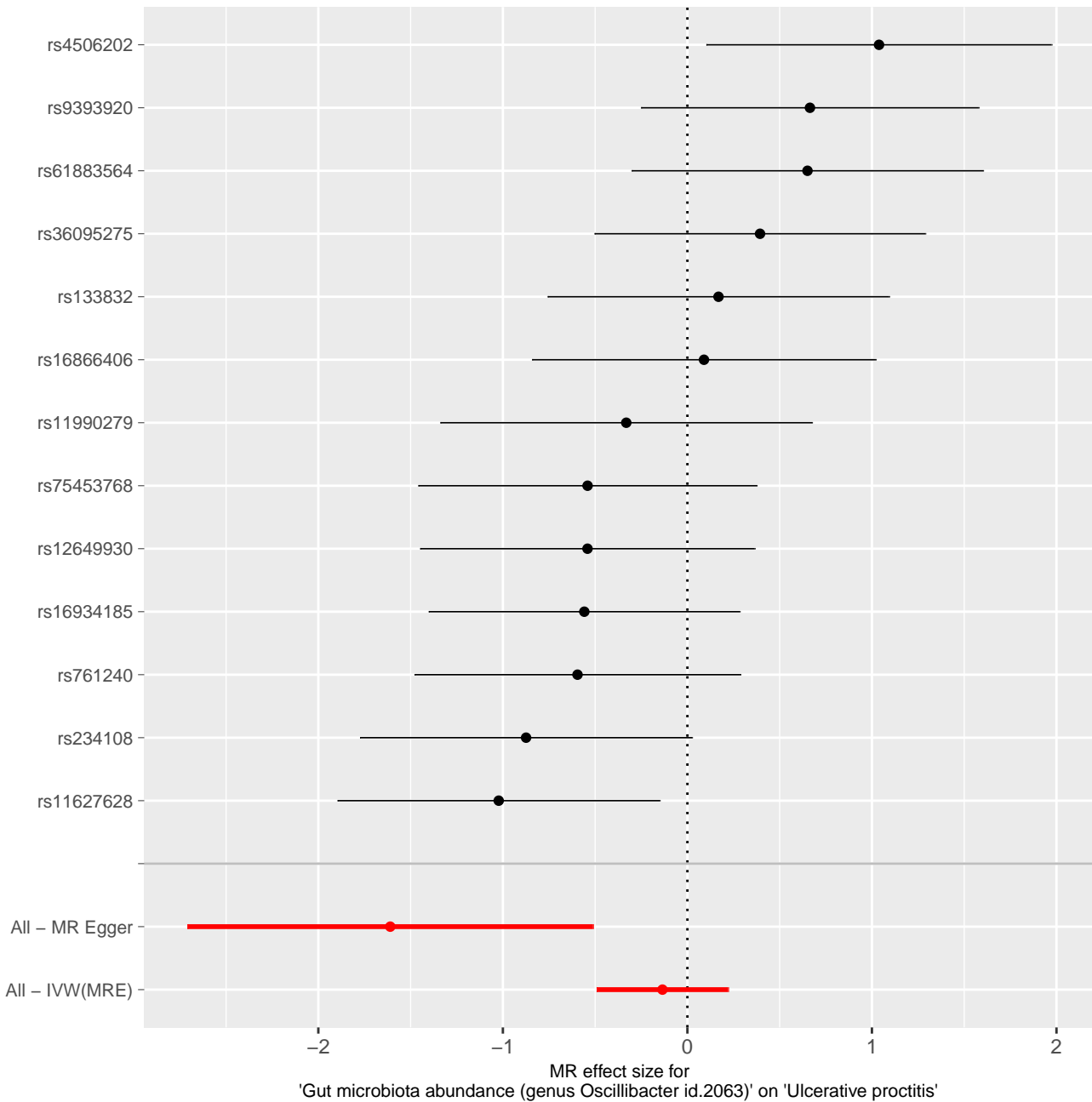


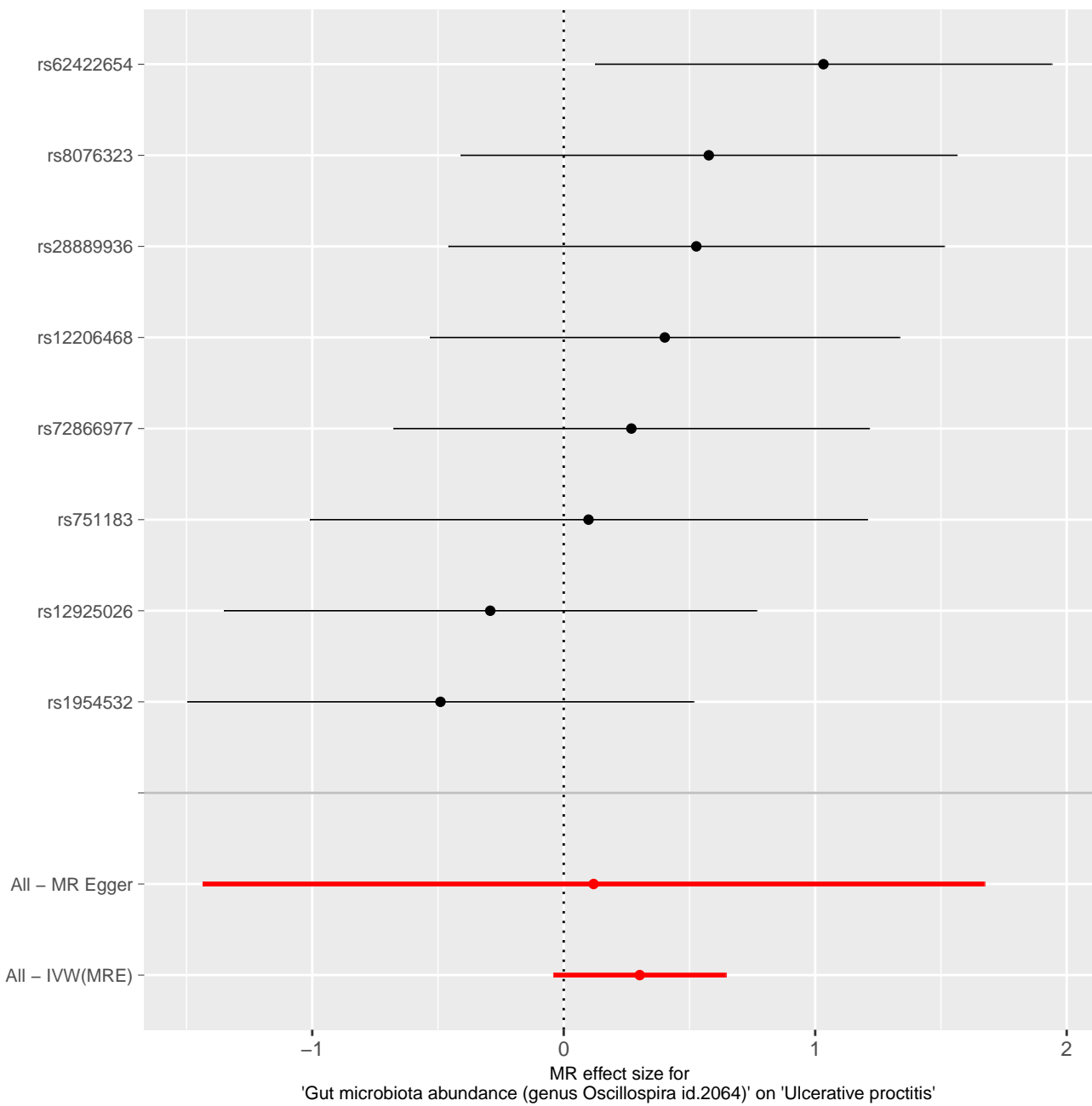


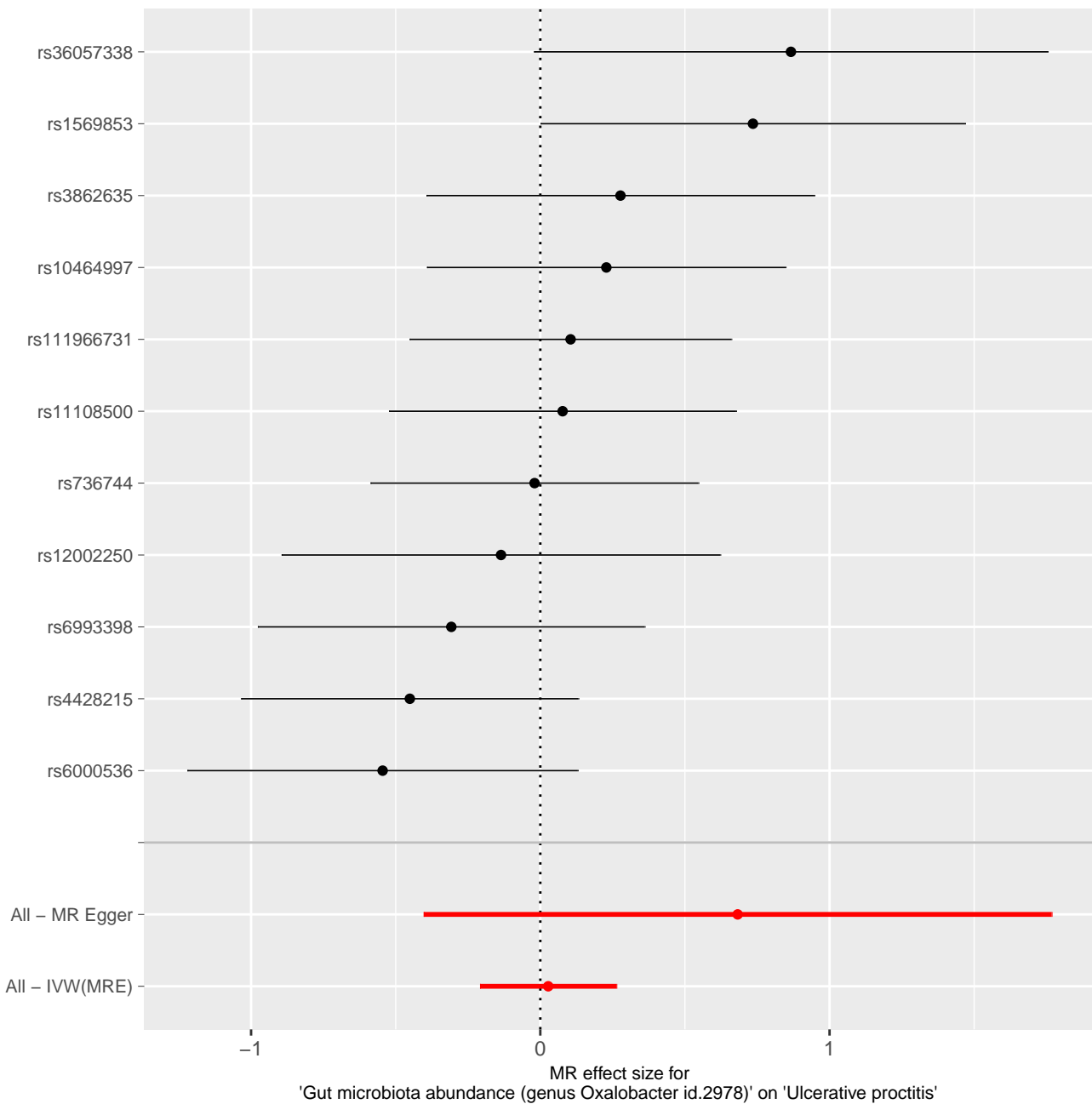


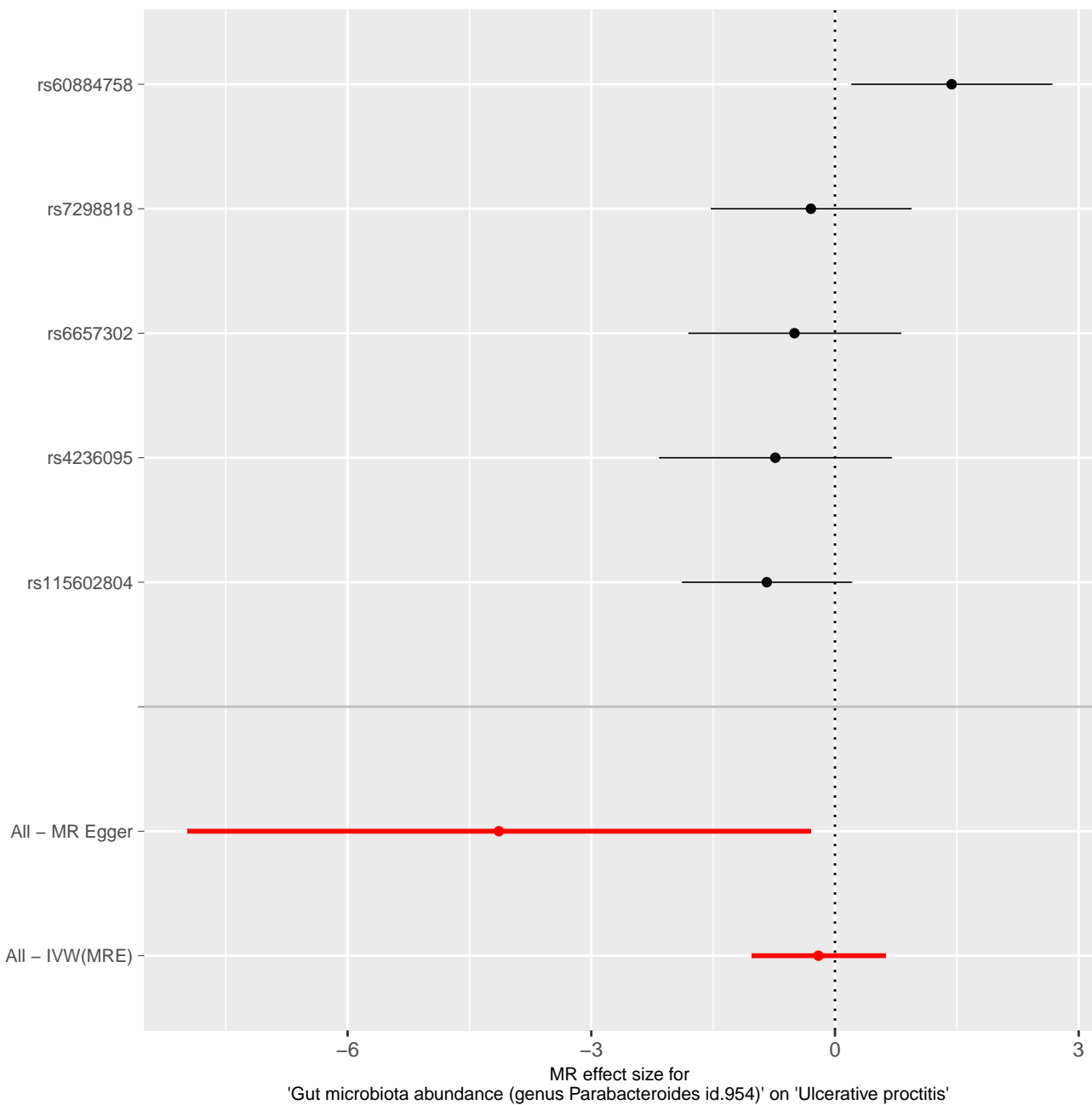


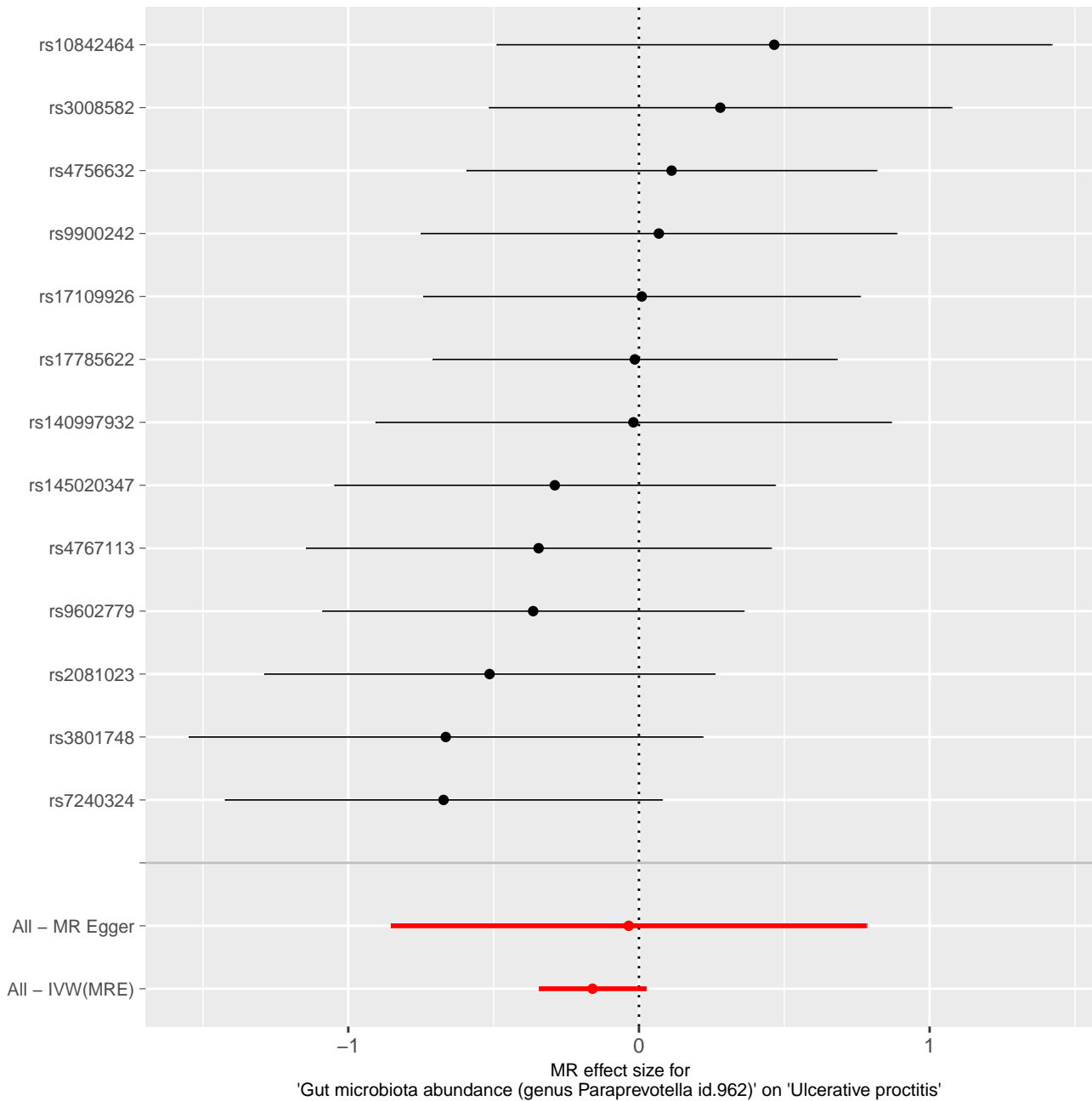
Batch 1460 : Gut microbiota abundance (genus Oscillibacter id.2063) on Ulcerative proctitis



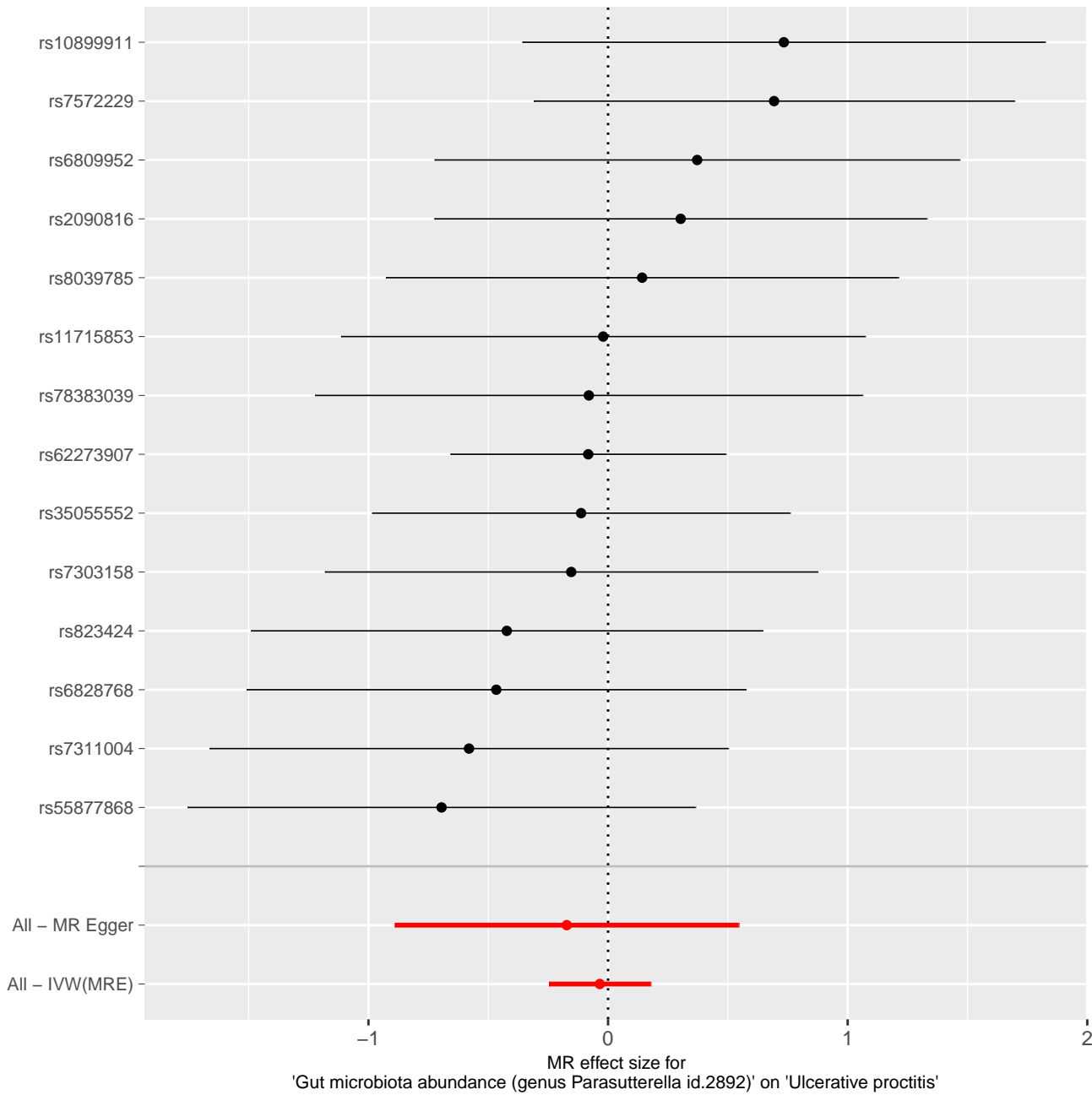


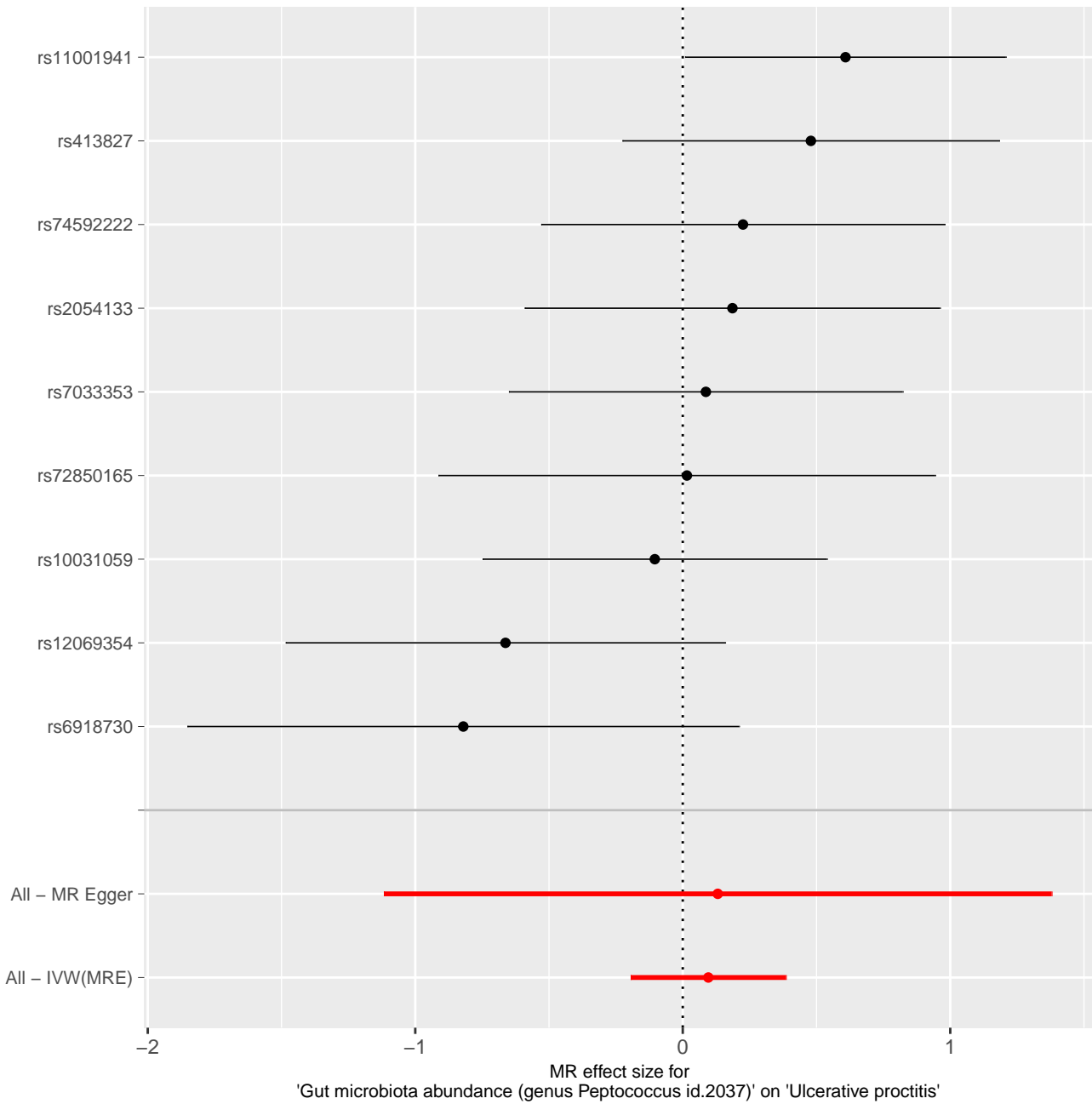


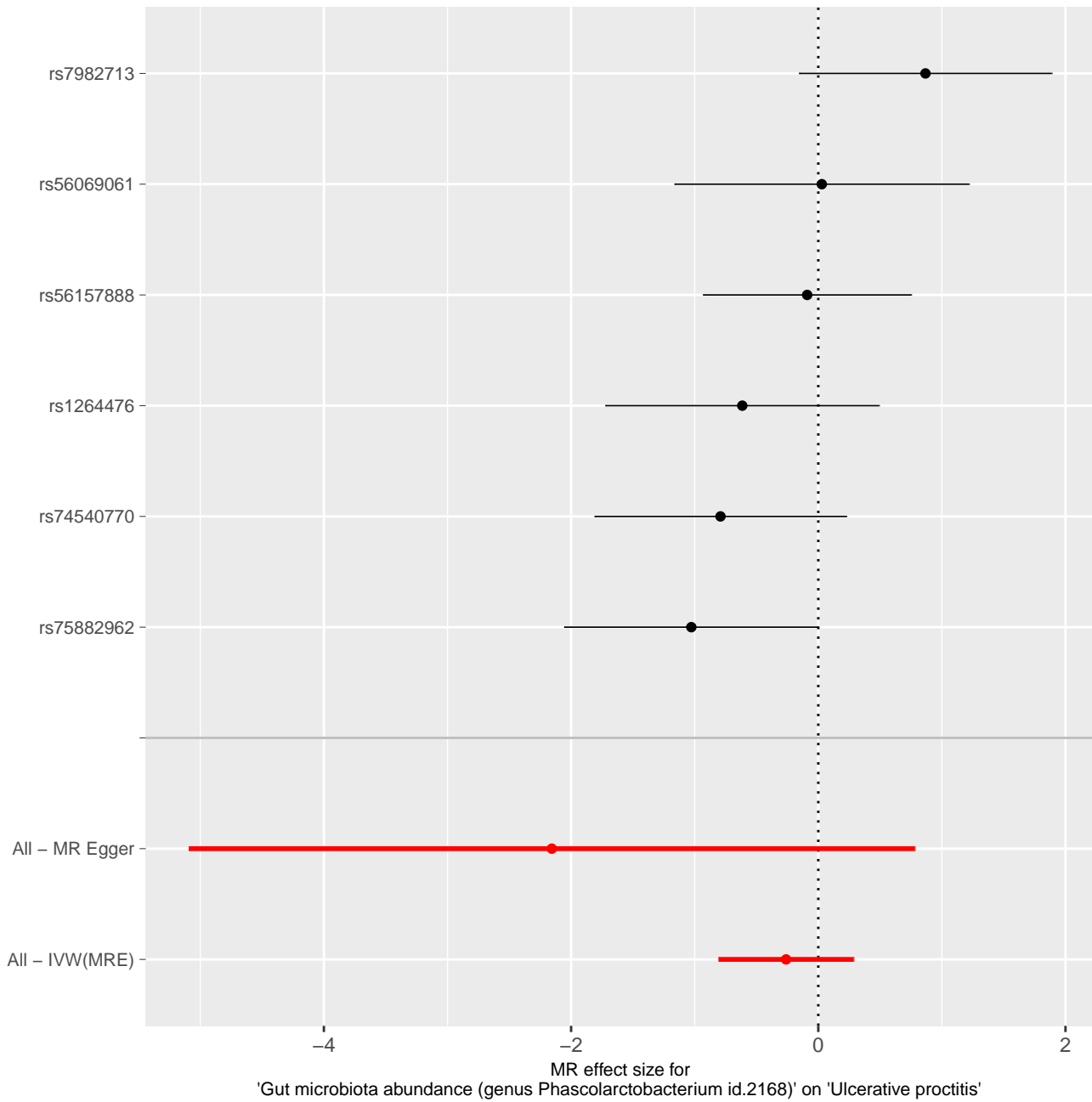


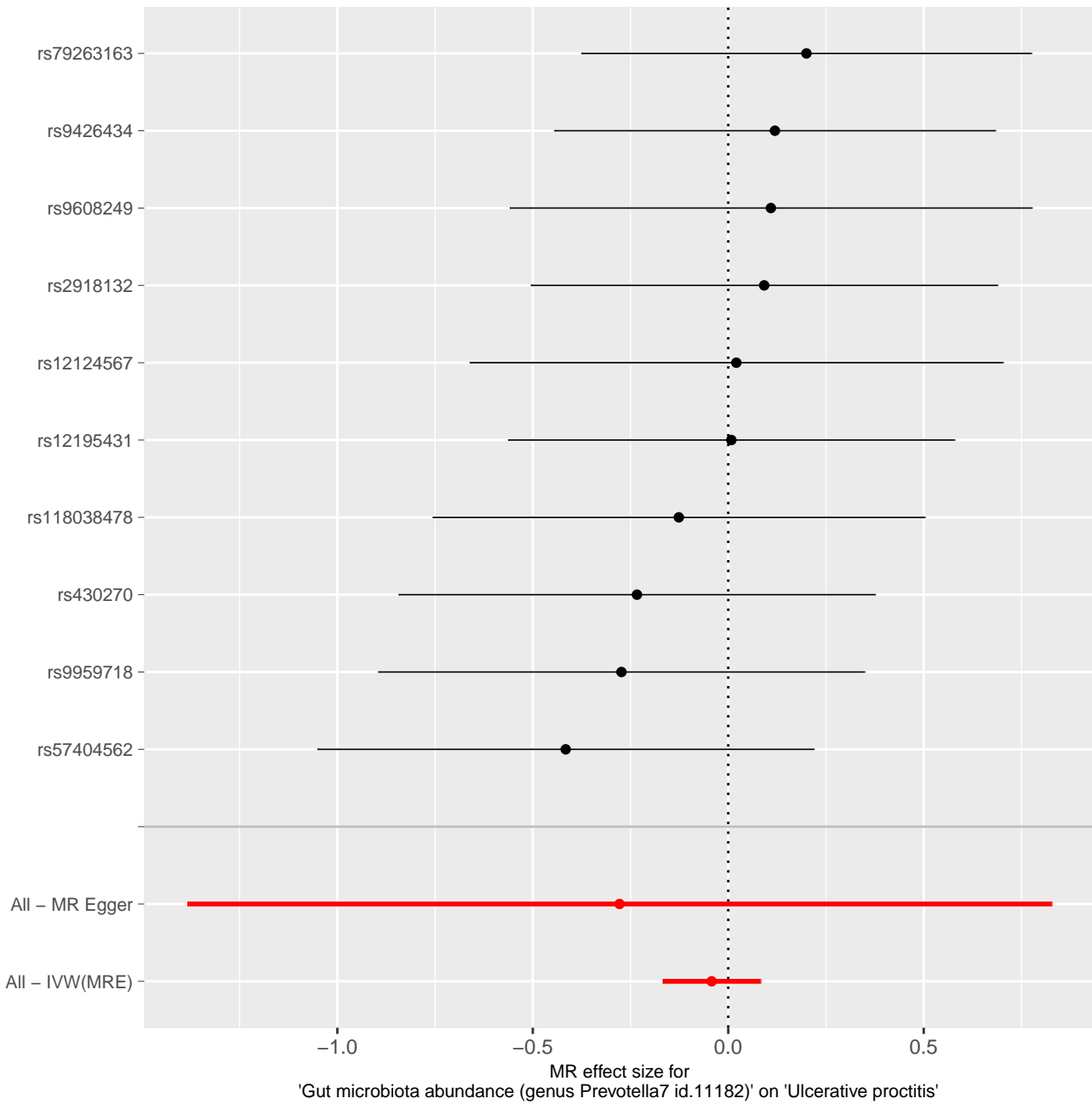


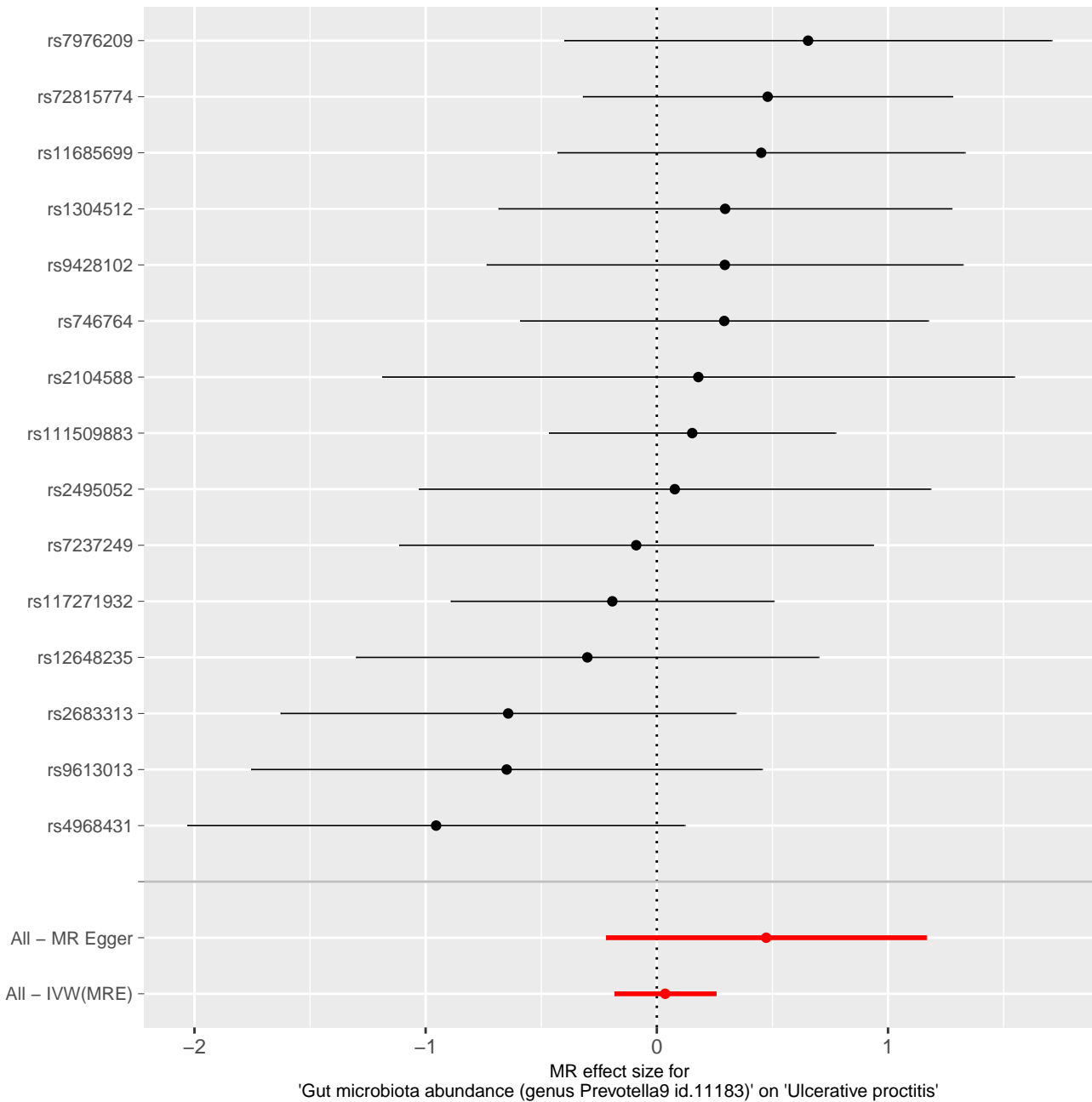
Batch 1465 : Gut microbiota abundance (genus Parasutterella id.2892) on Ulcerative proctitis

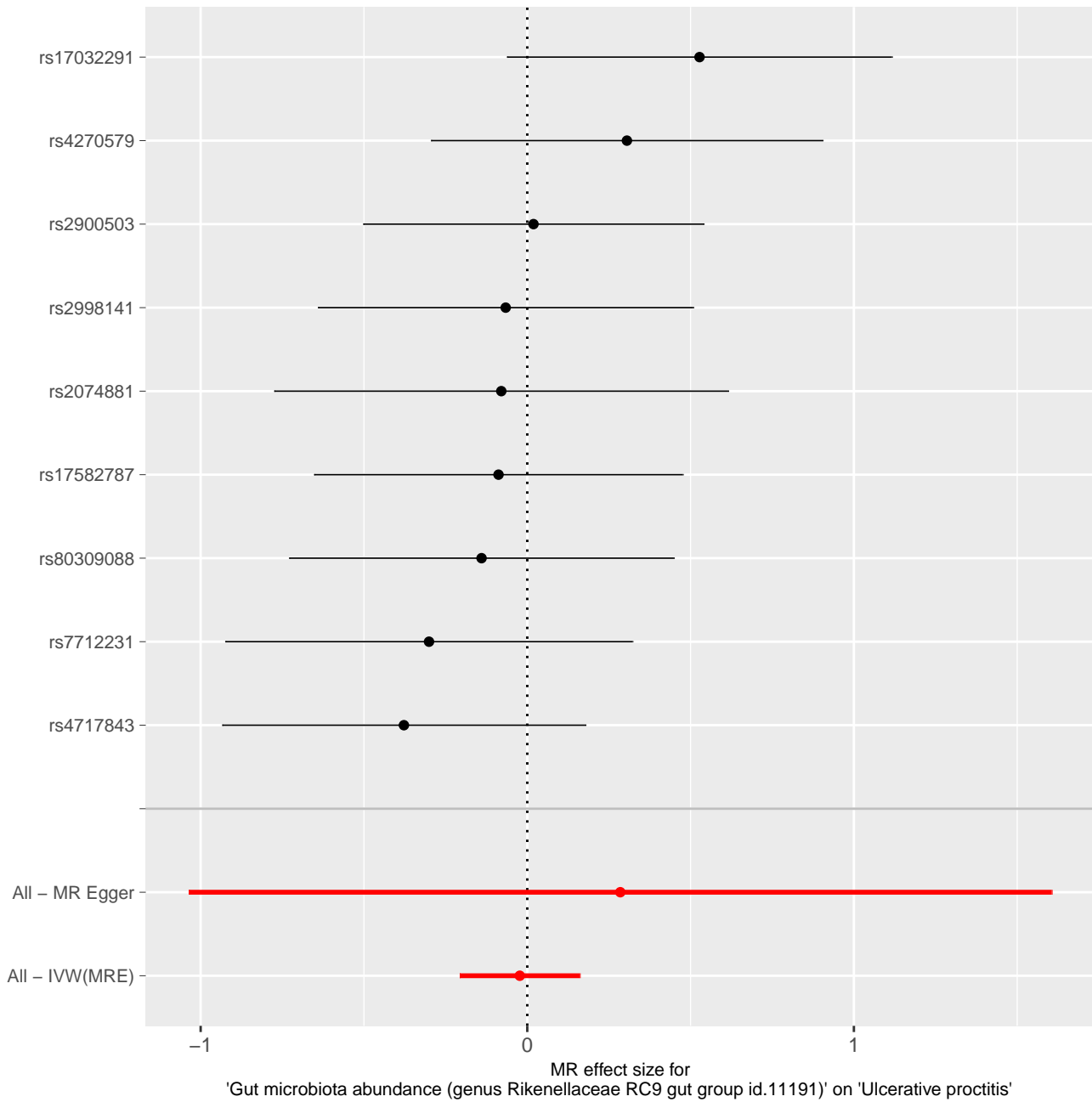


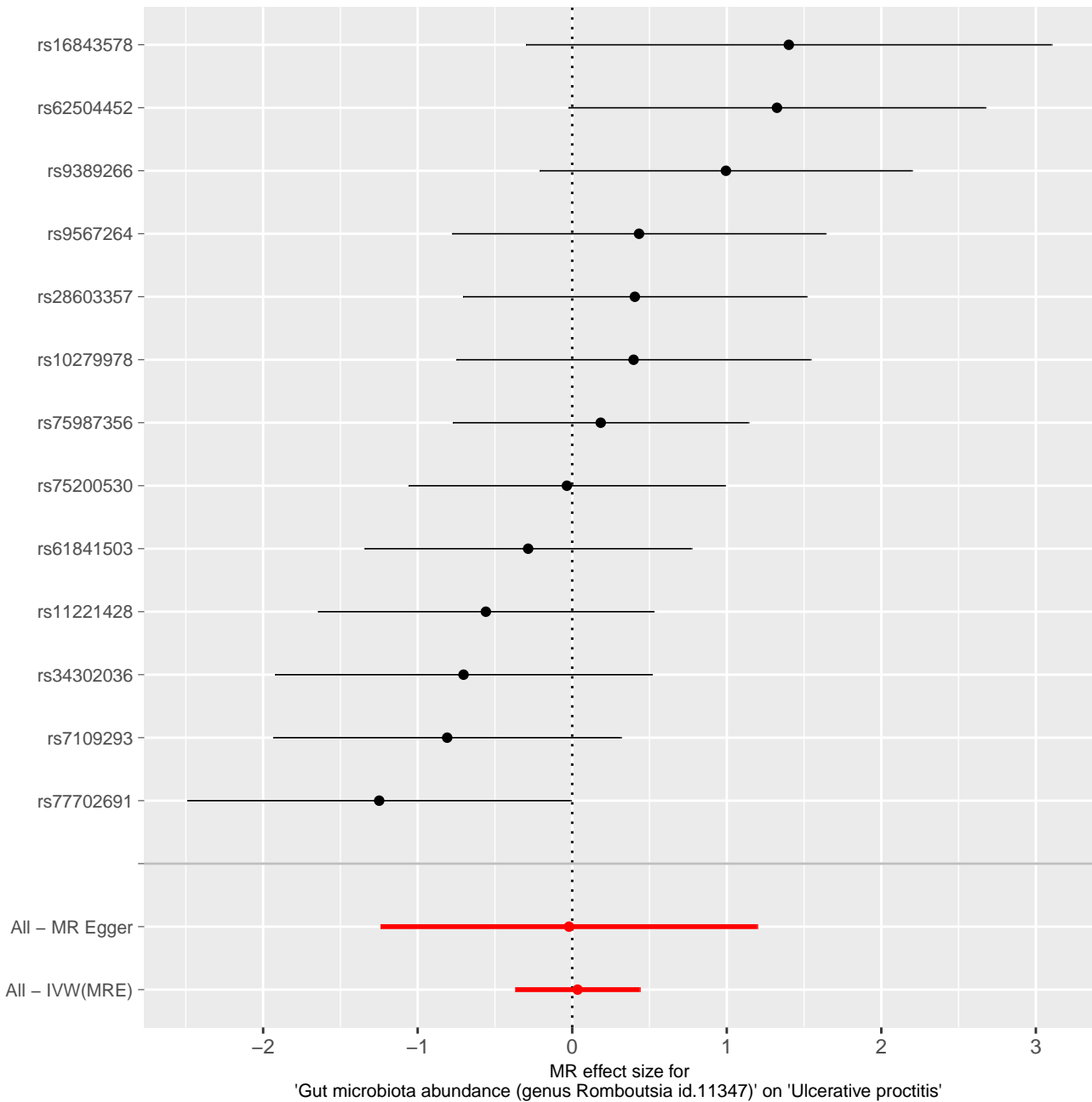


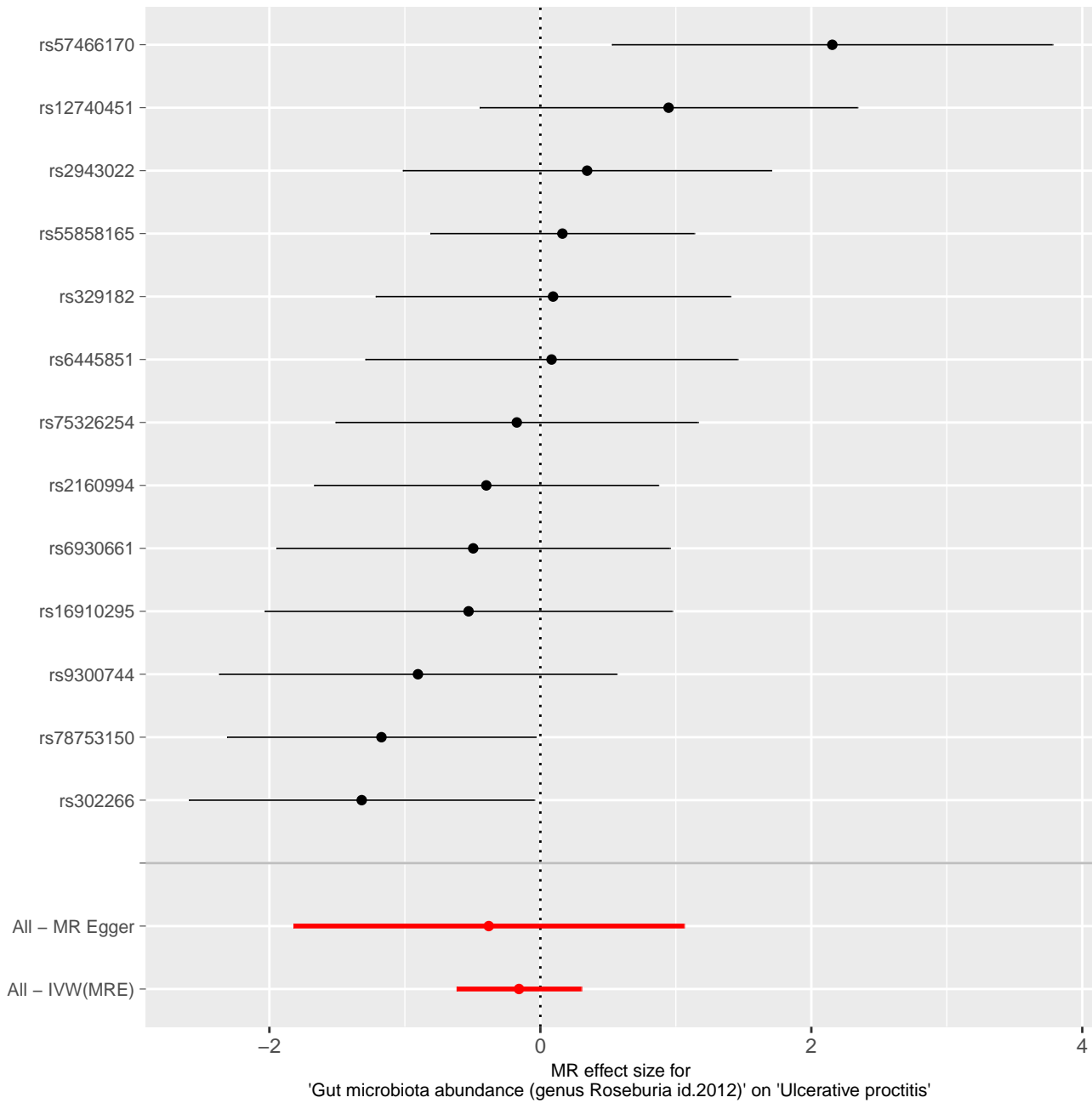




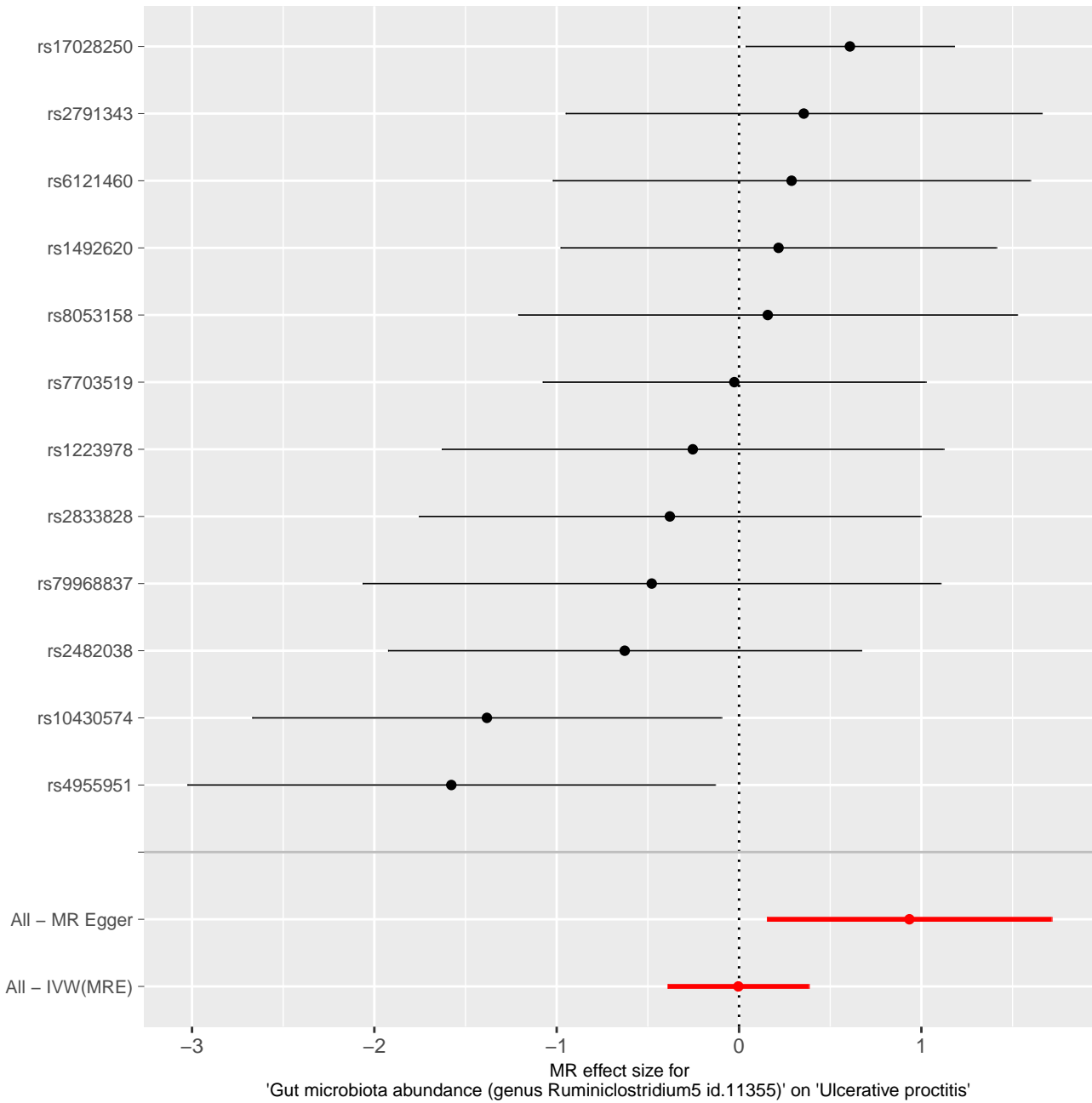


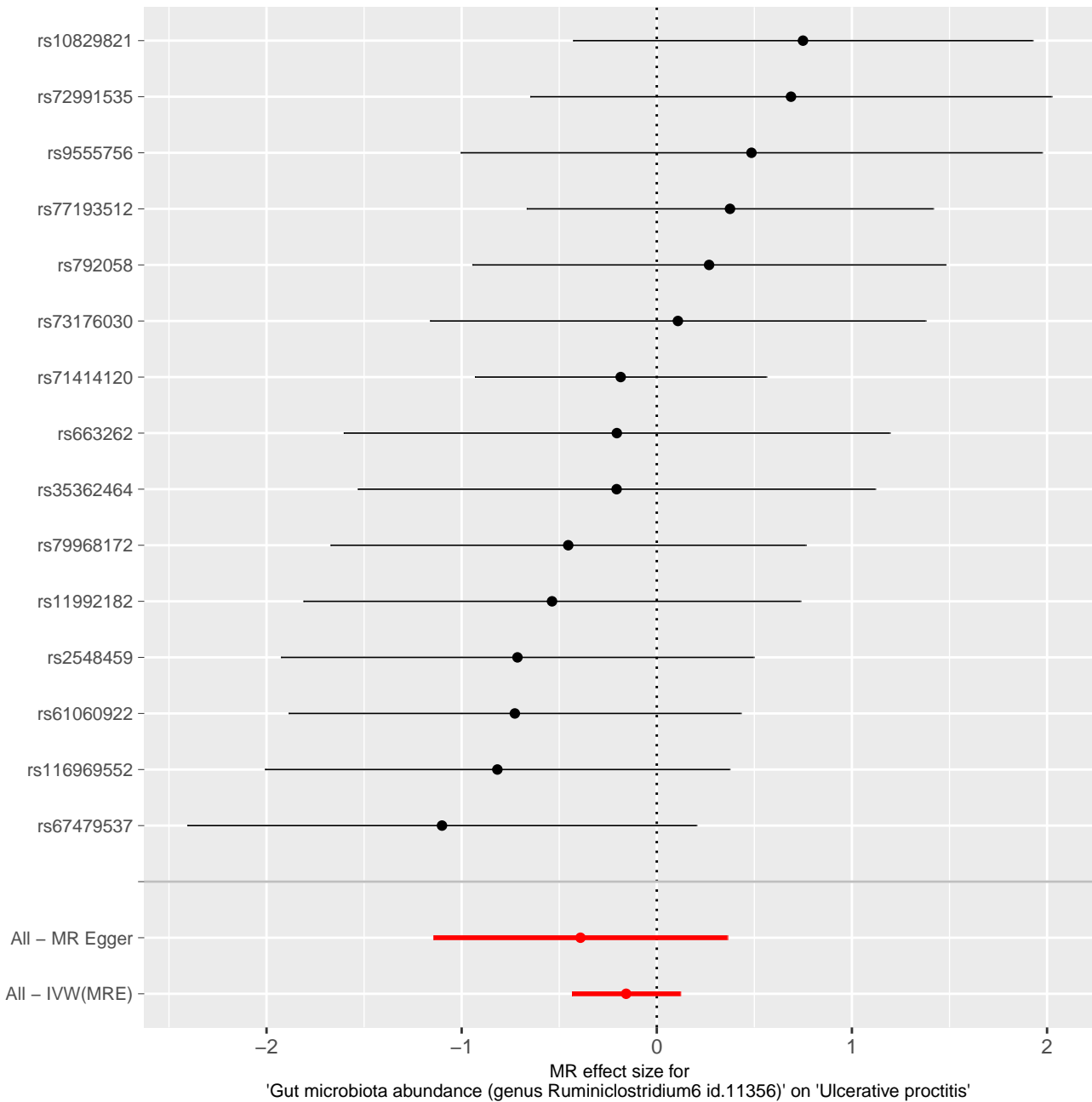


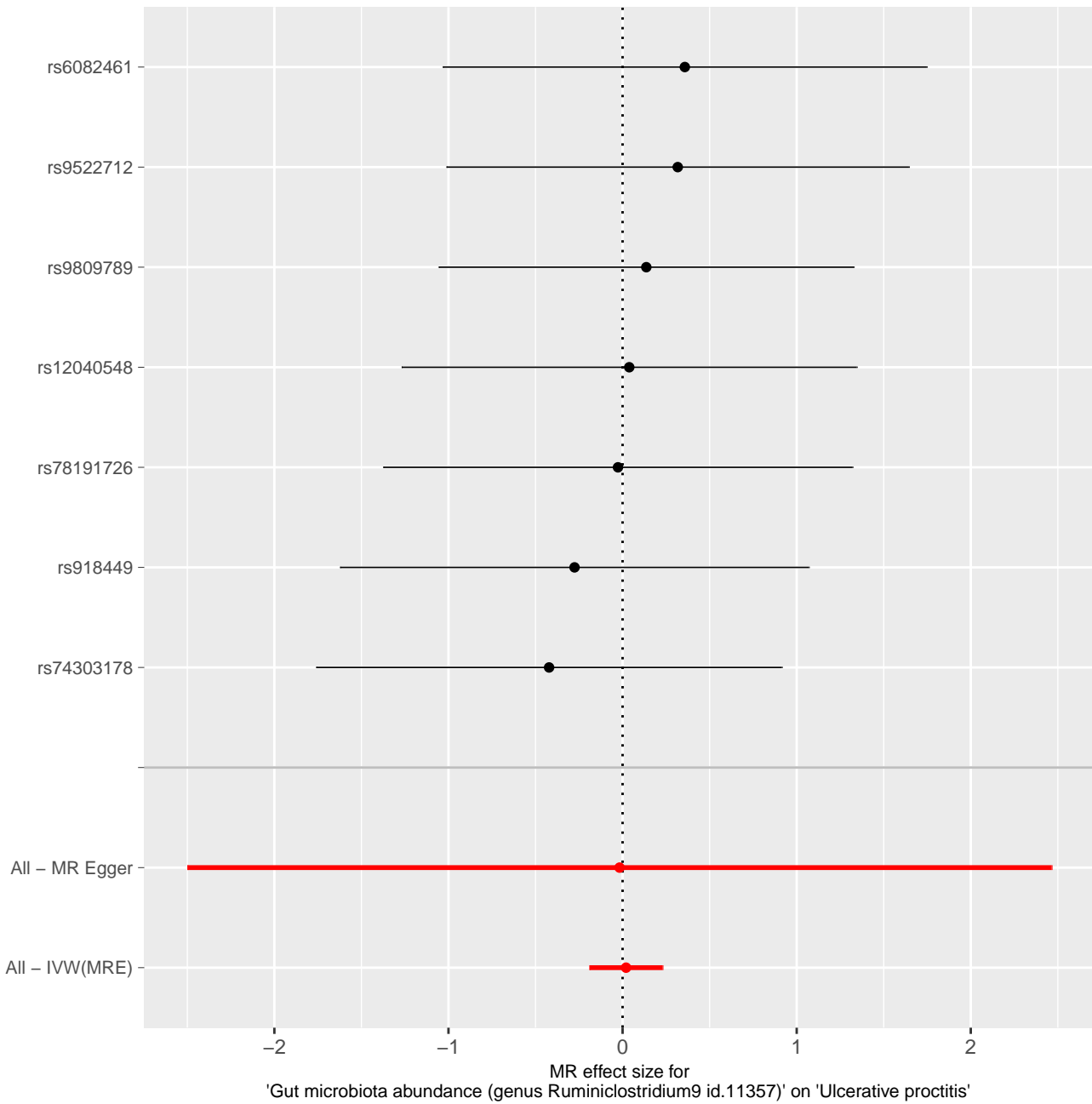


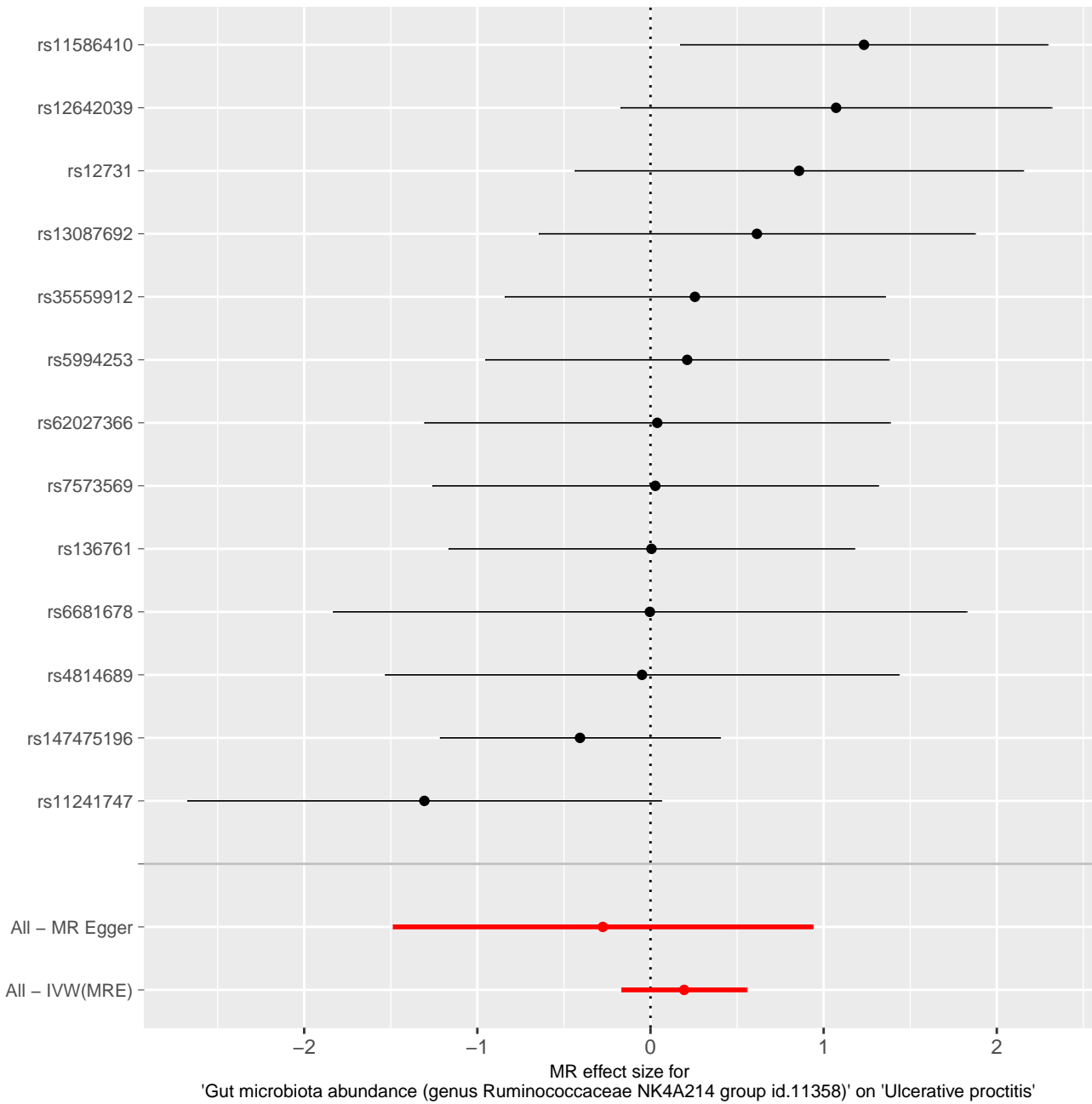


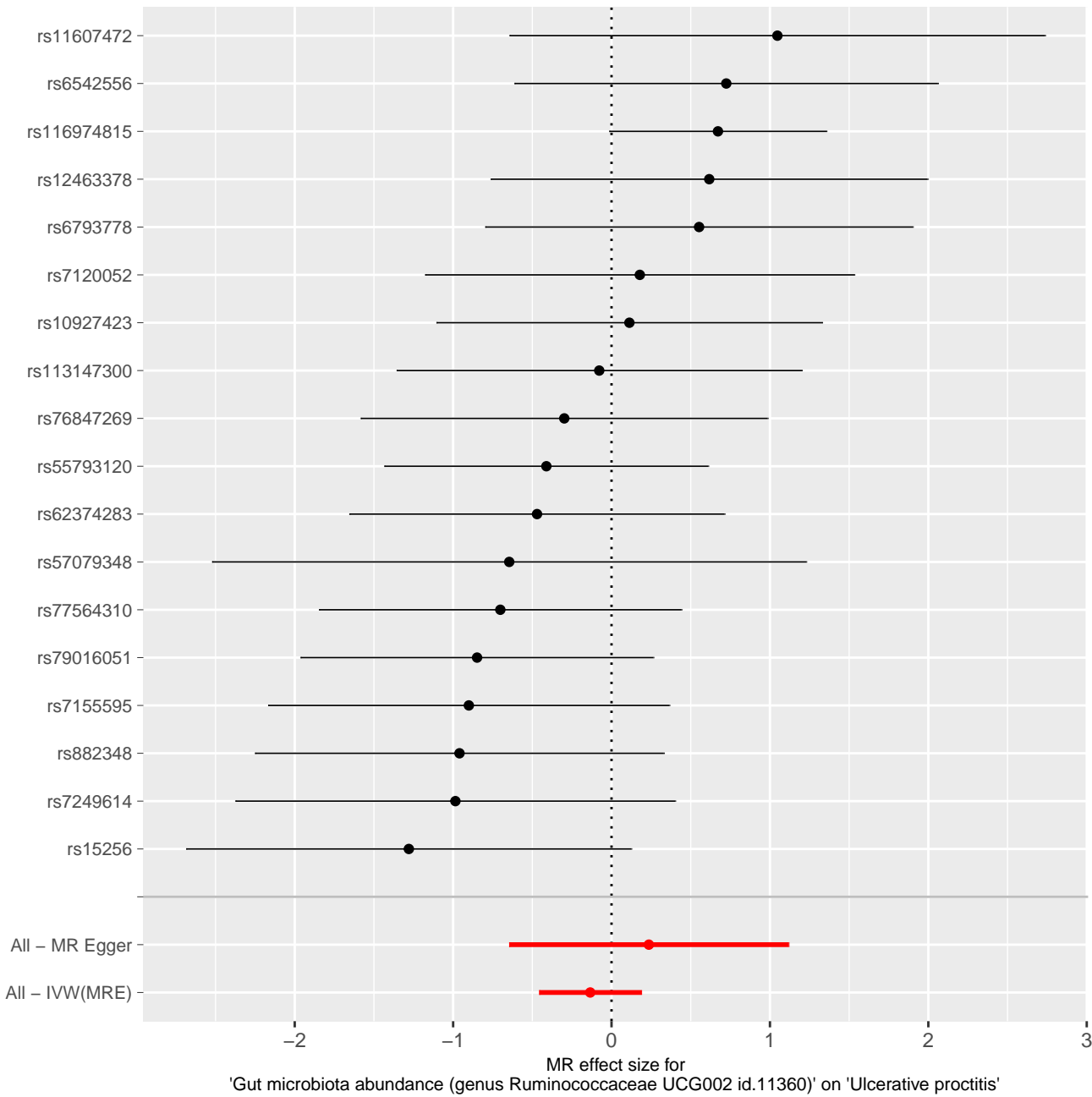
Batch 1473 : Gut microbiota abundance (genus Ruminiclostridium5 id.11355) on Ulcerative proctitis

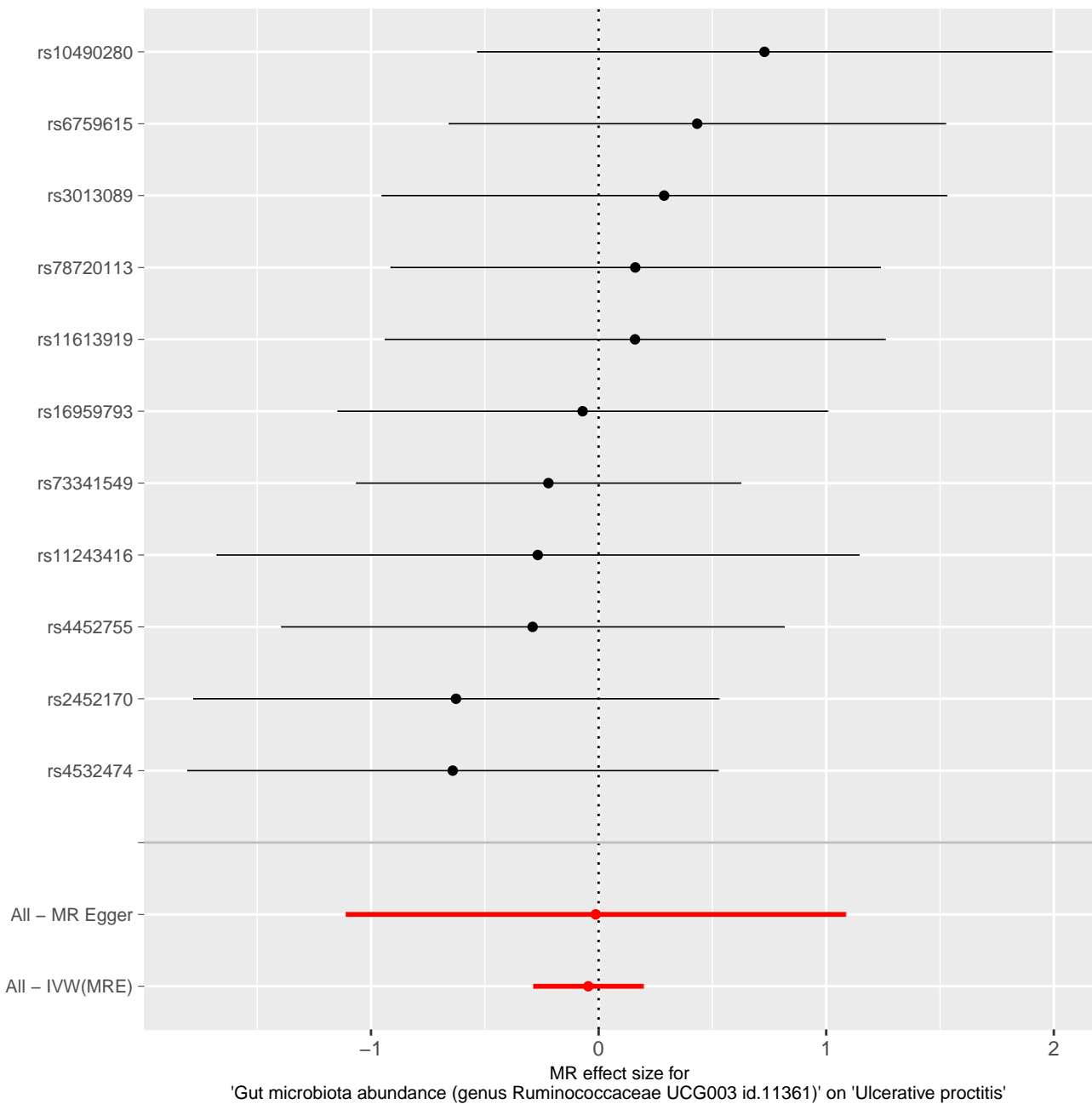


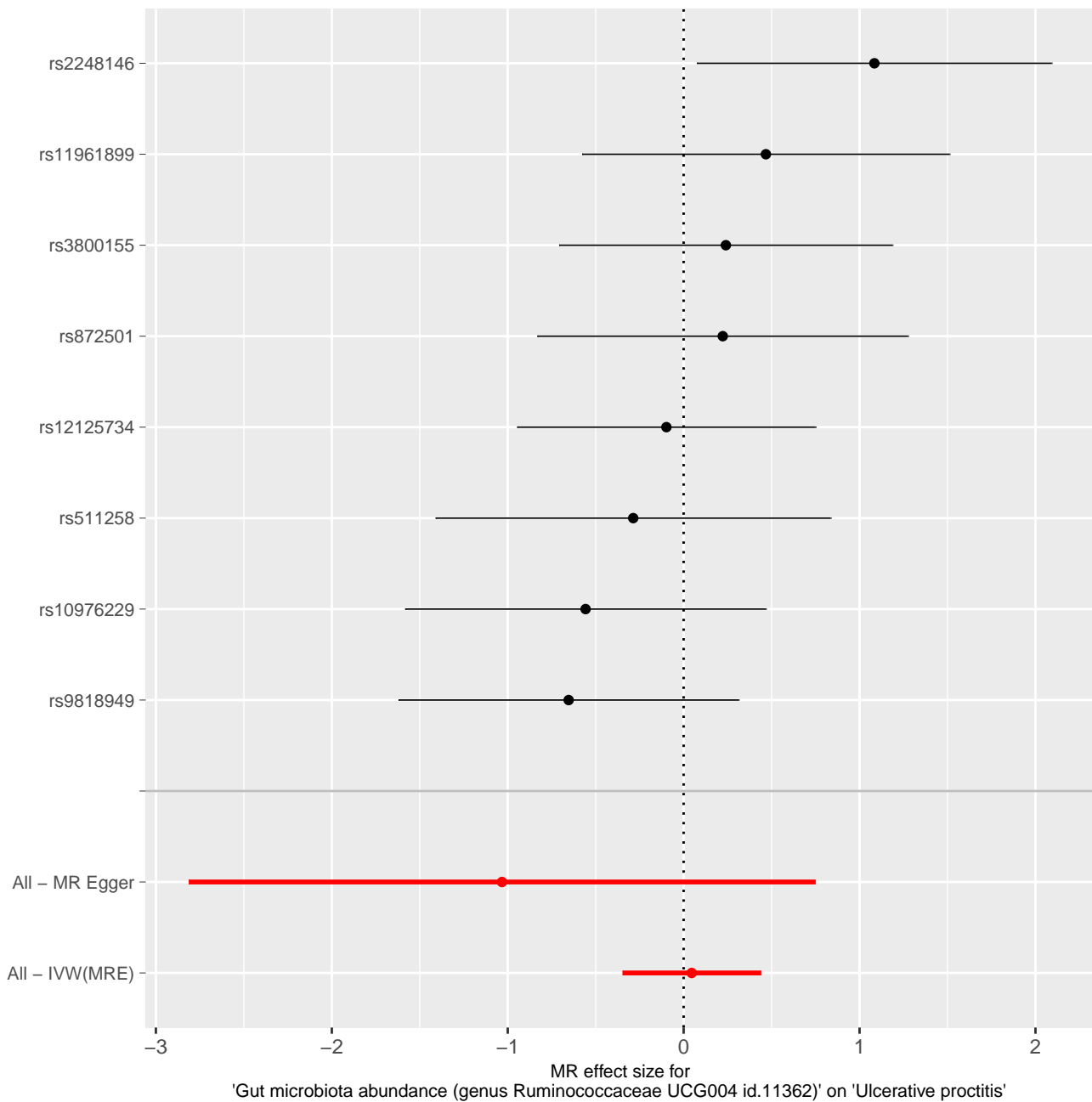


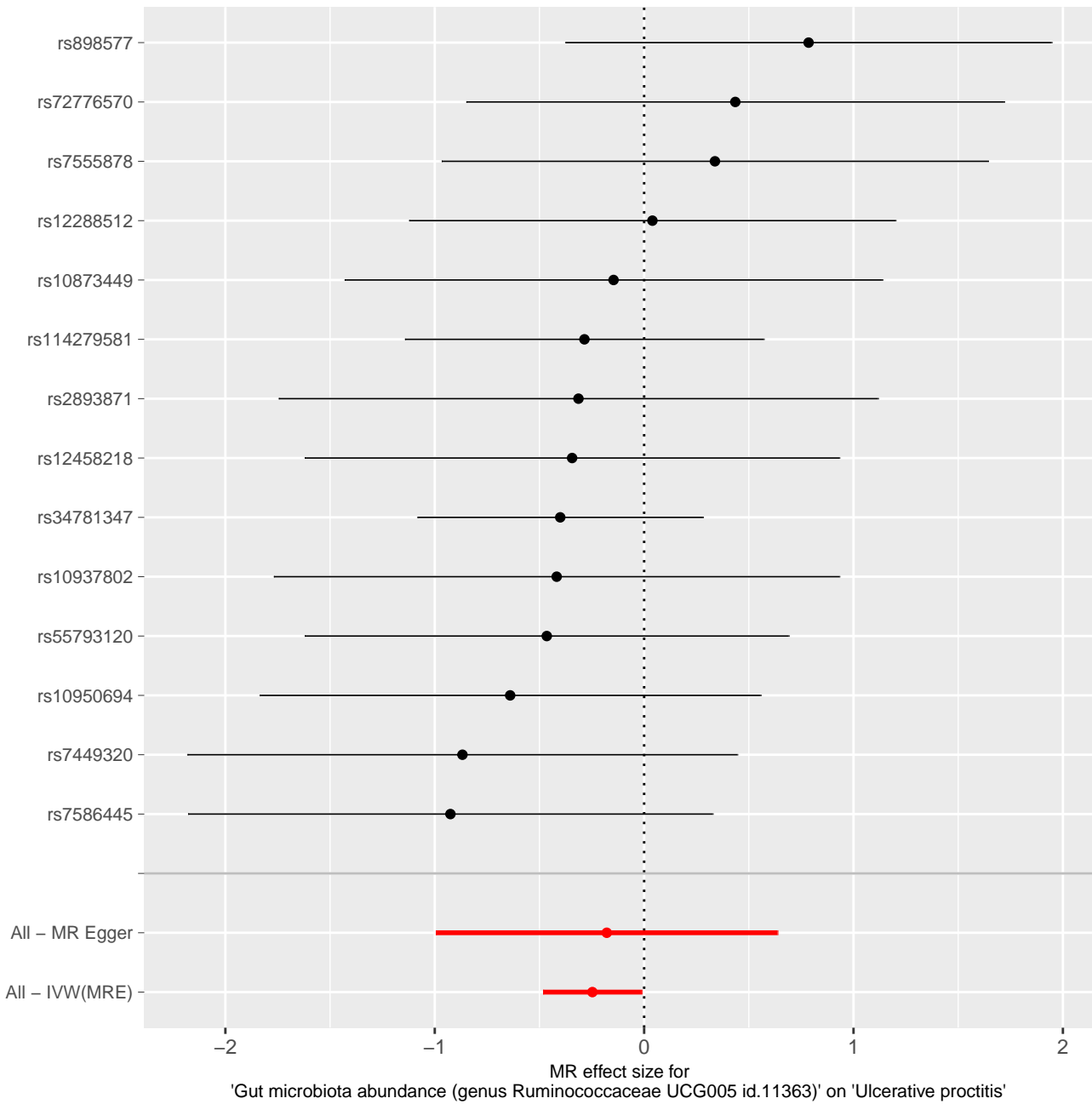


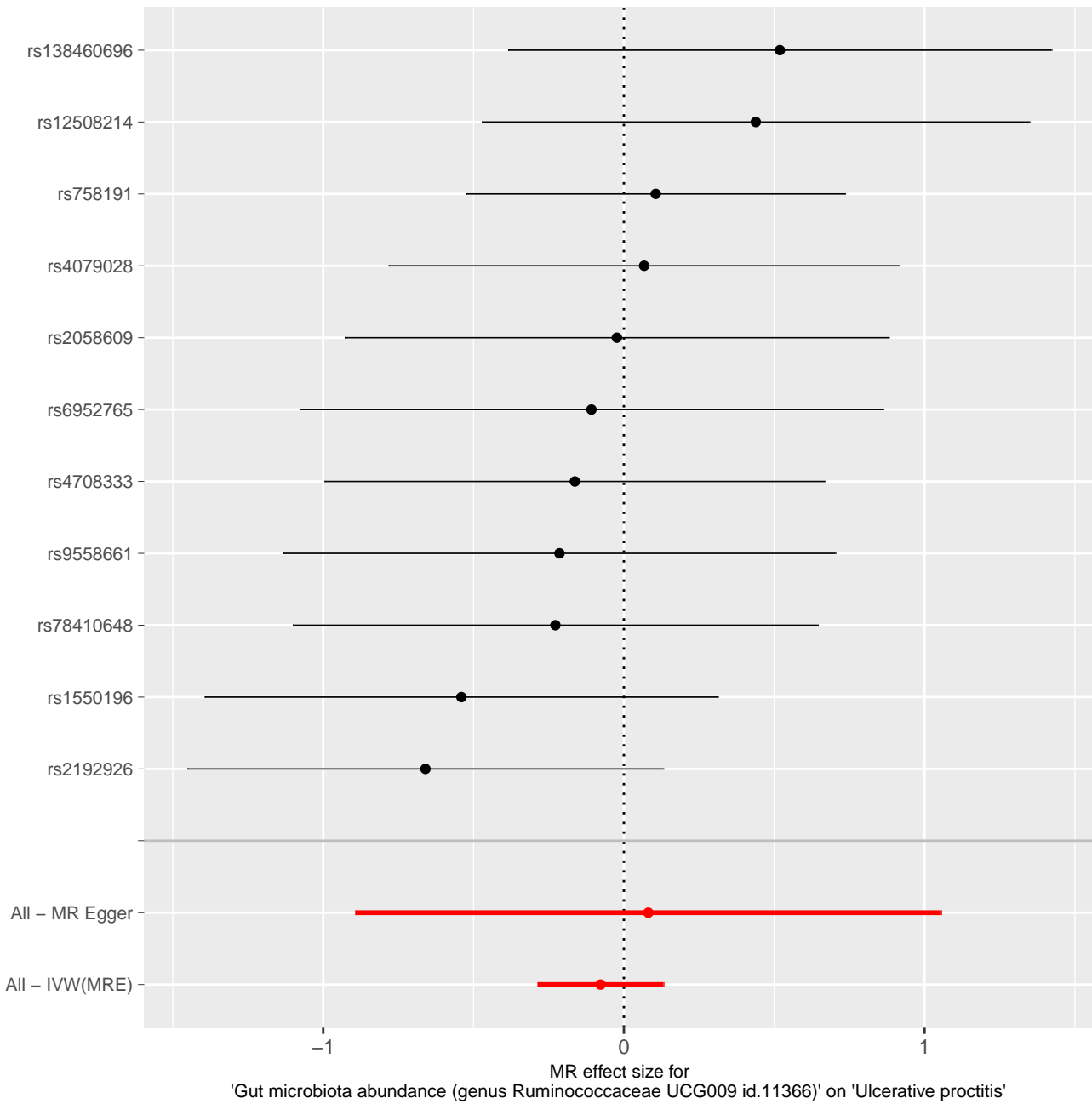


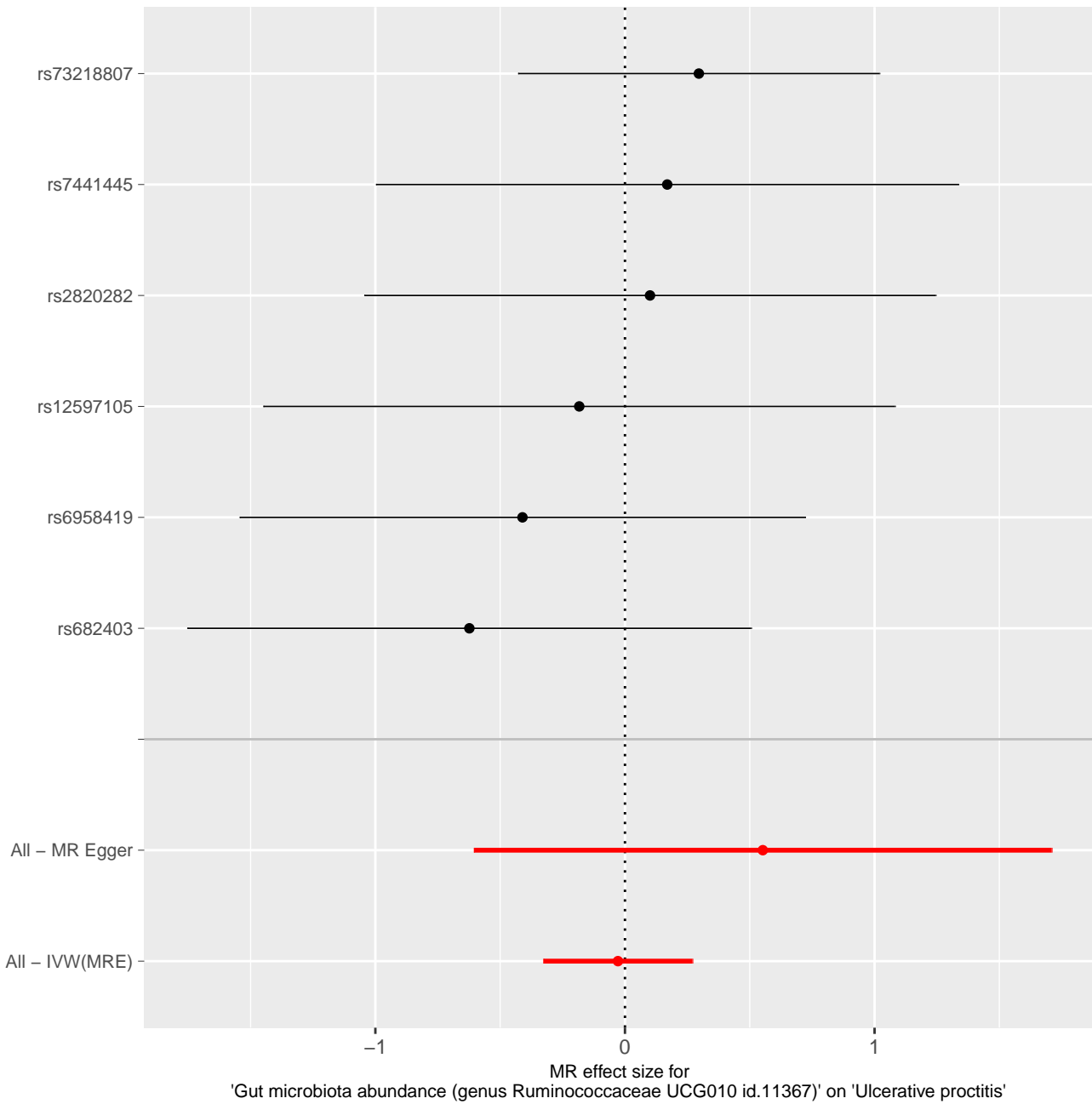


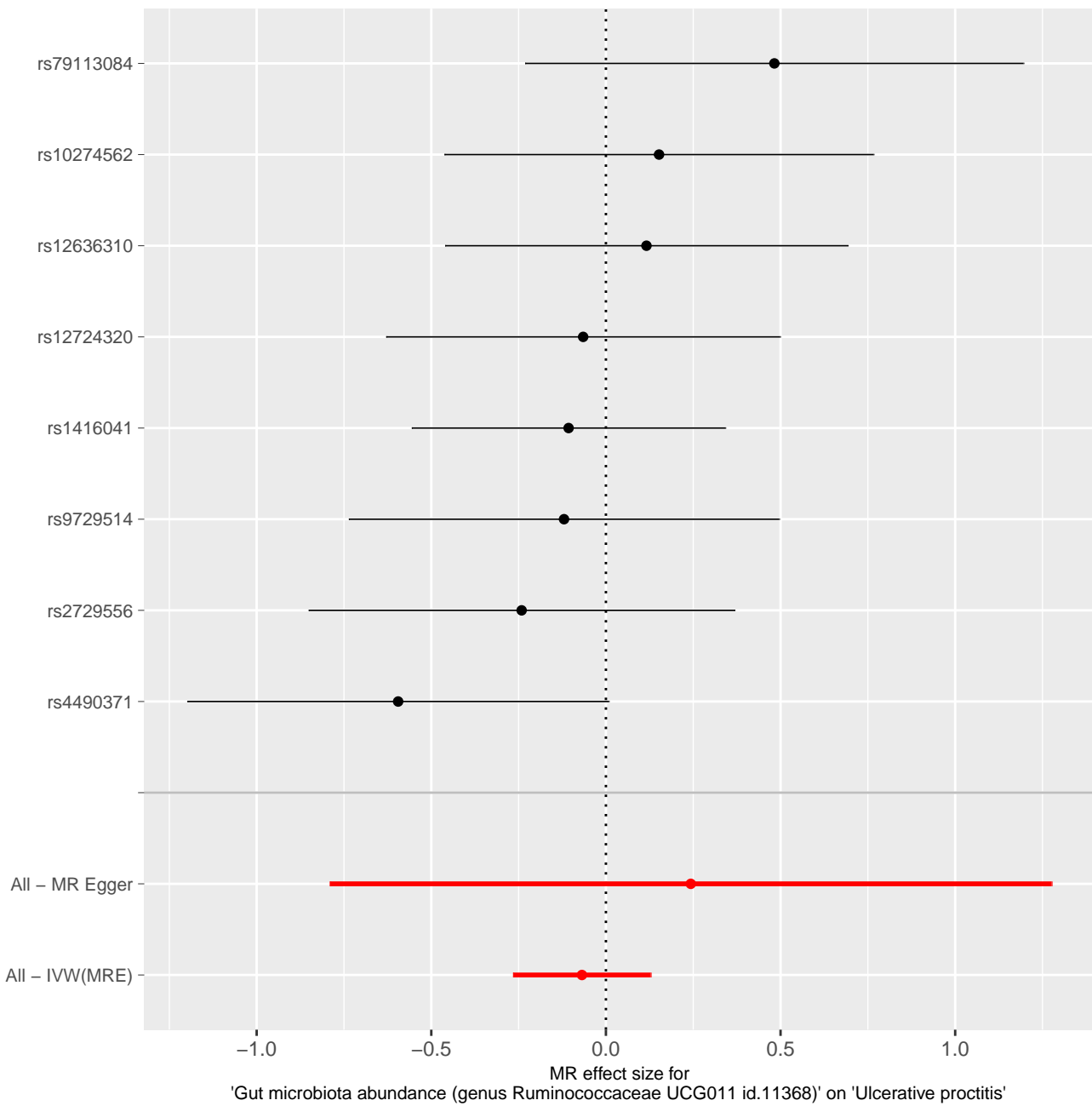


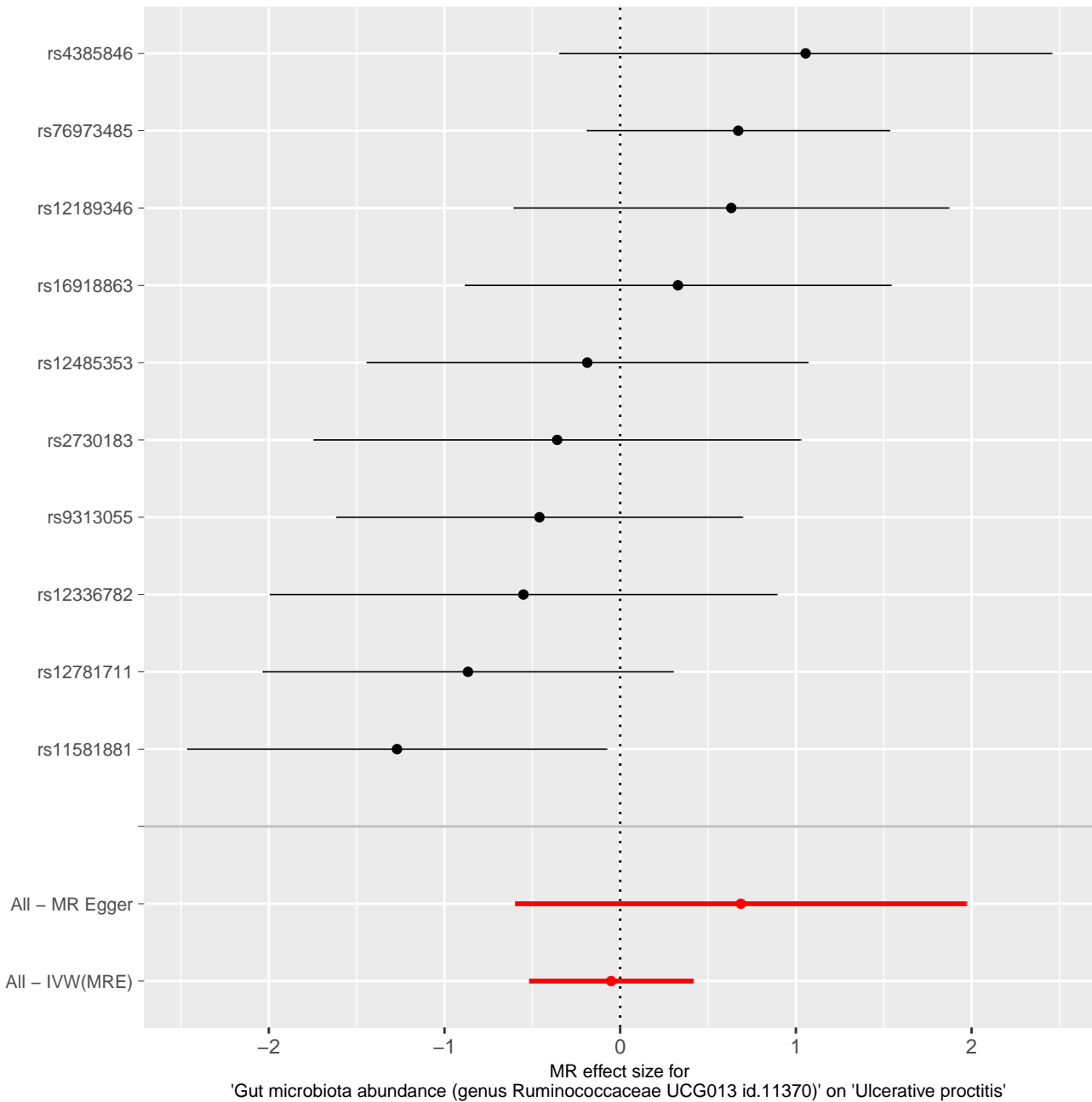


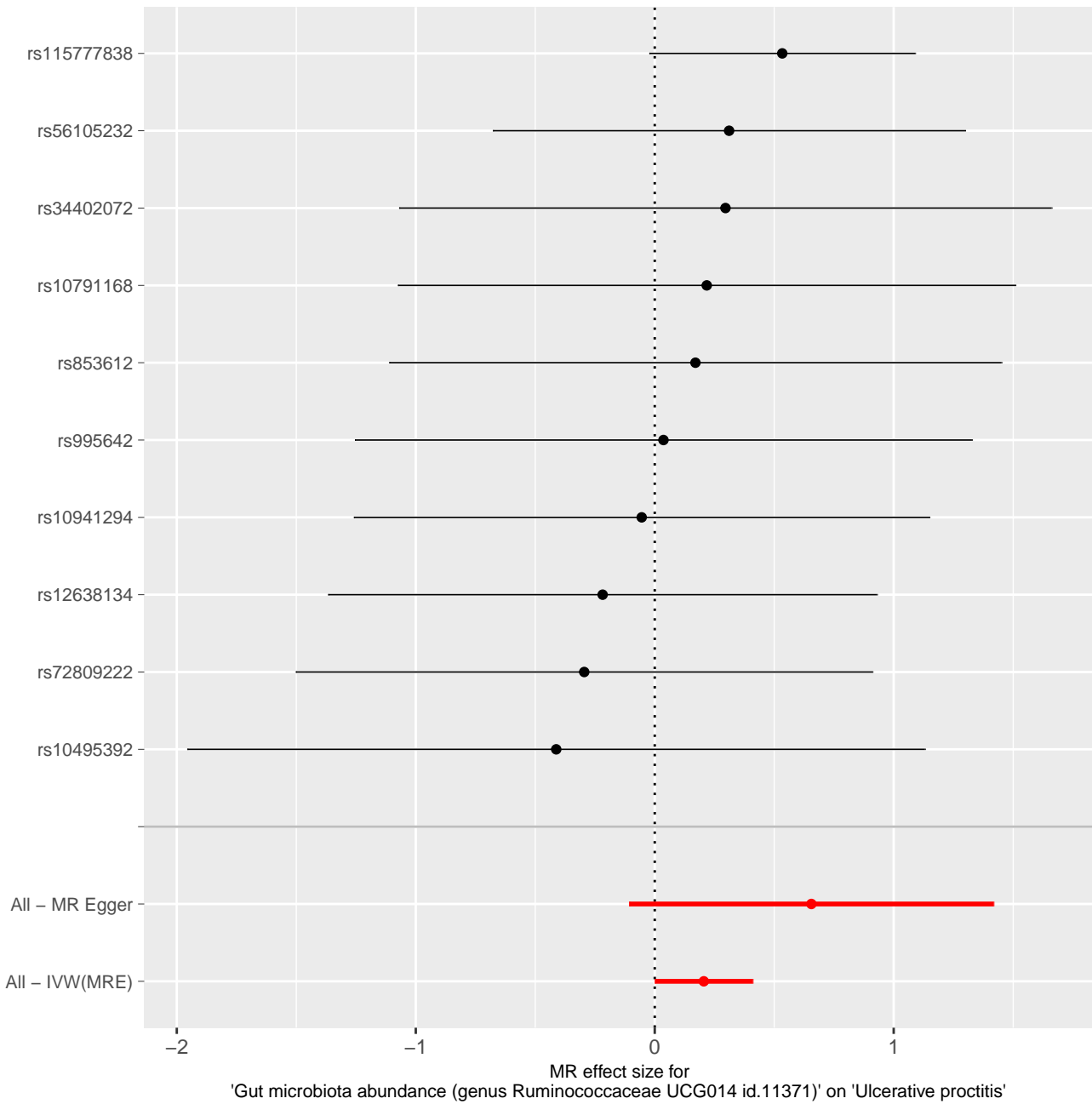


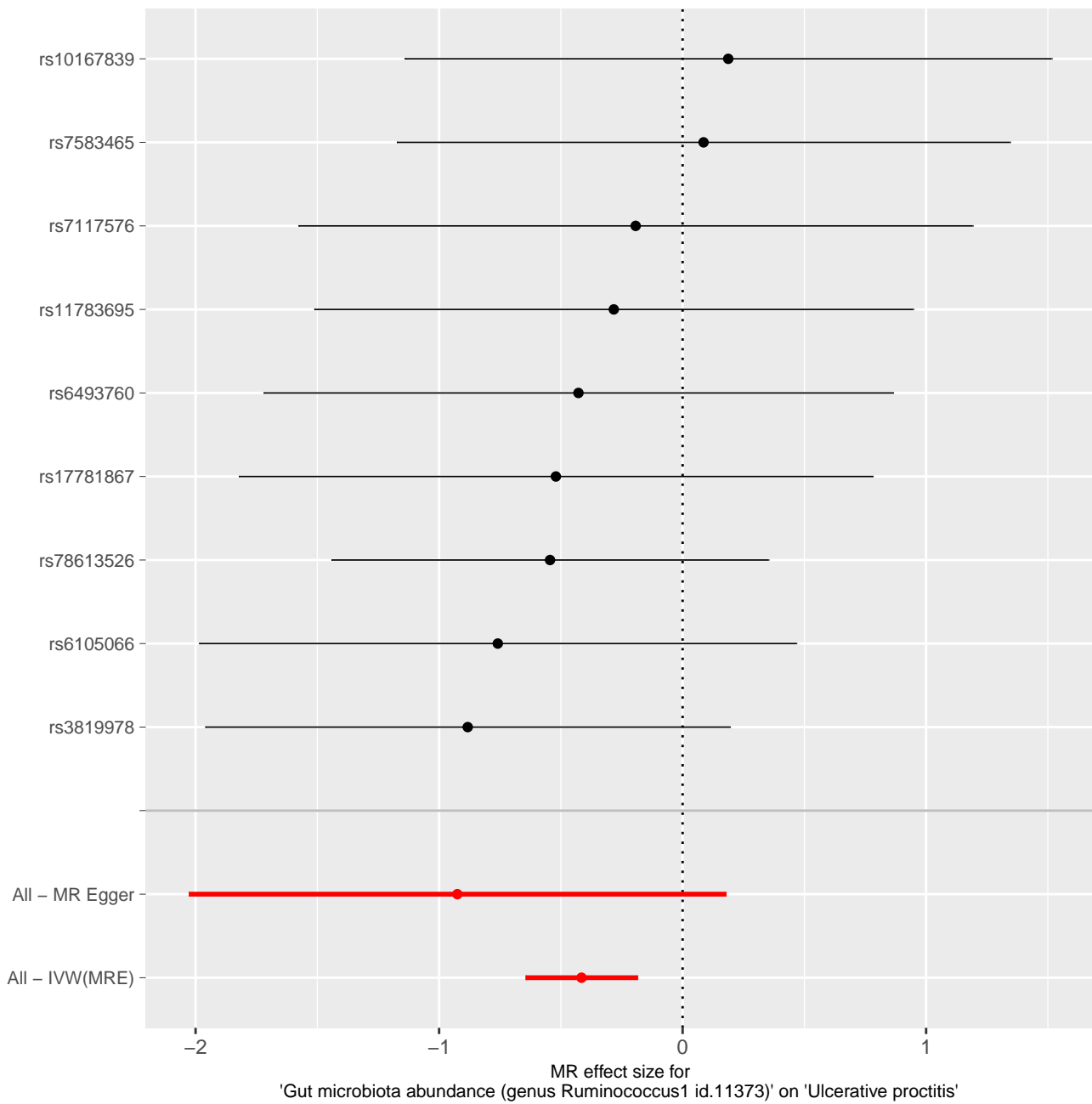


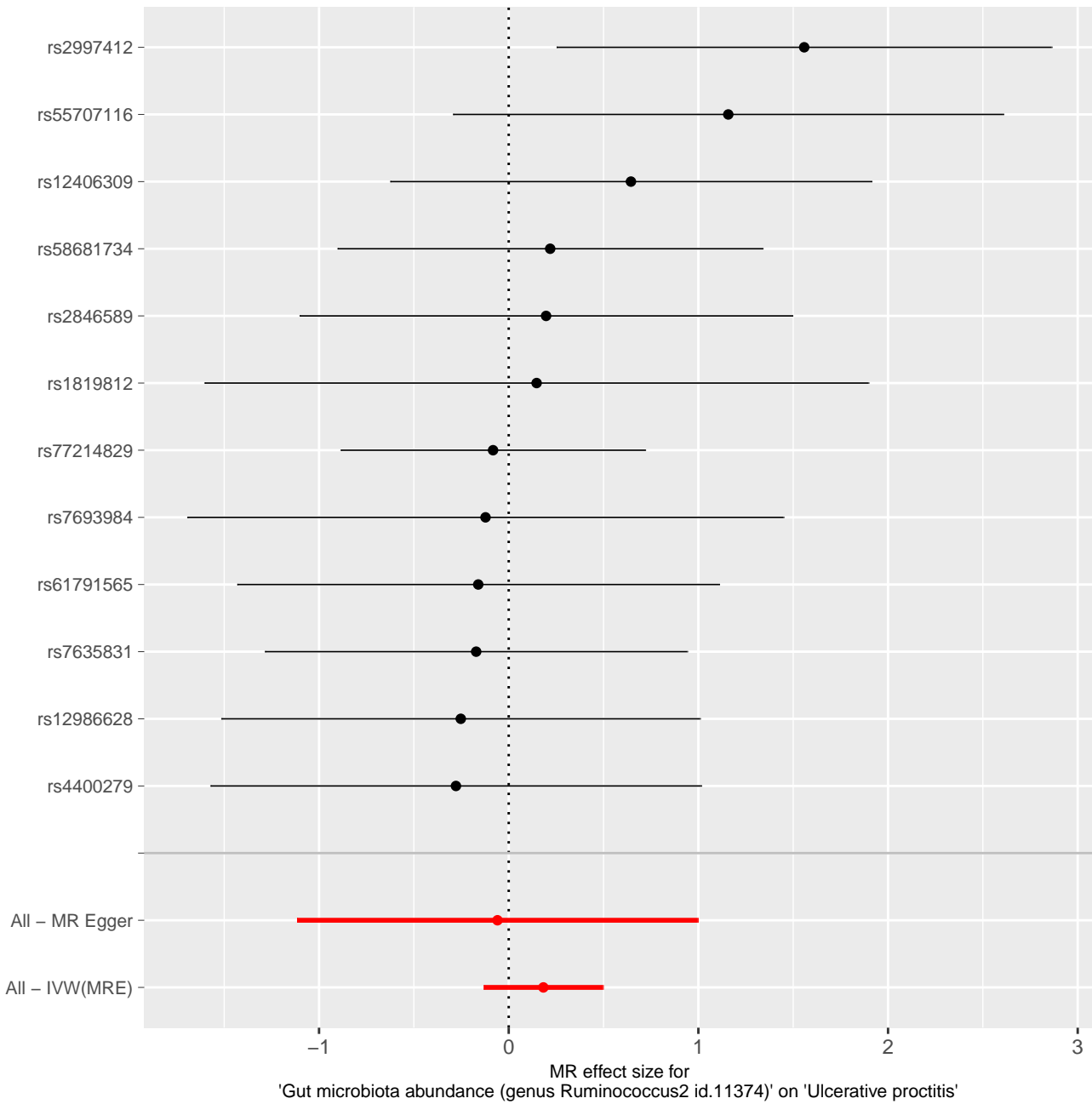


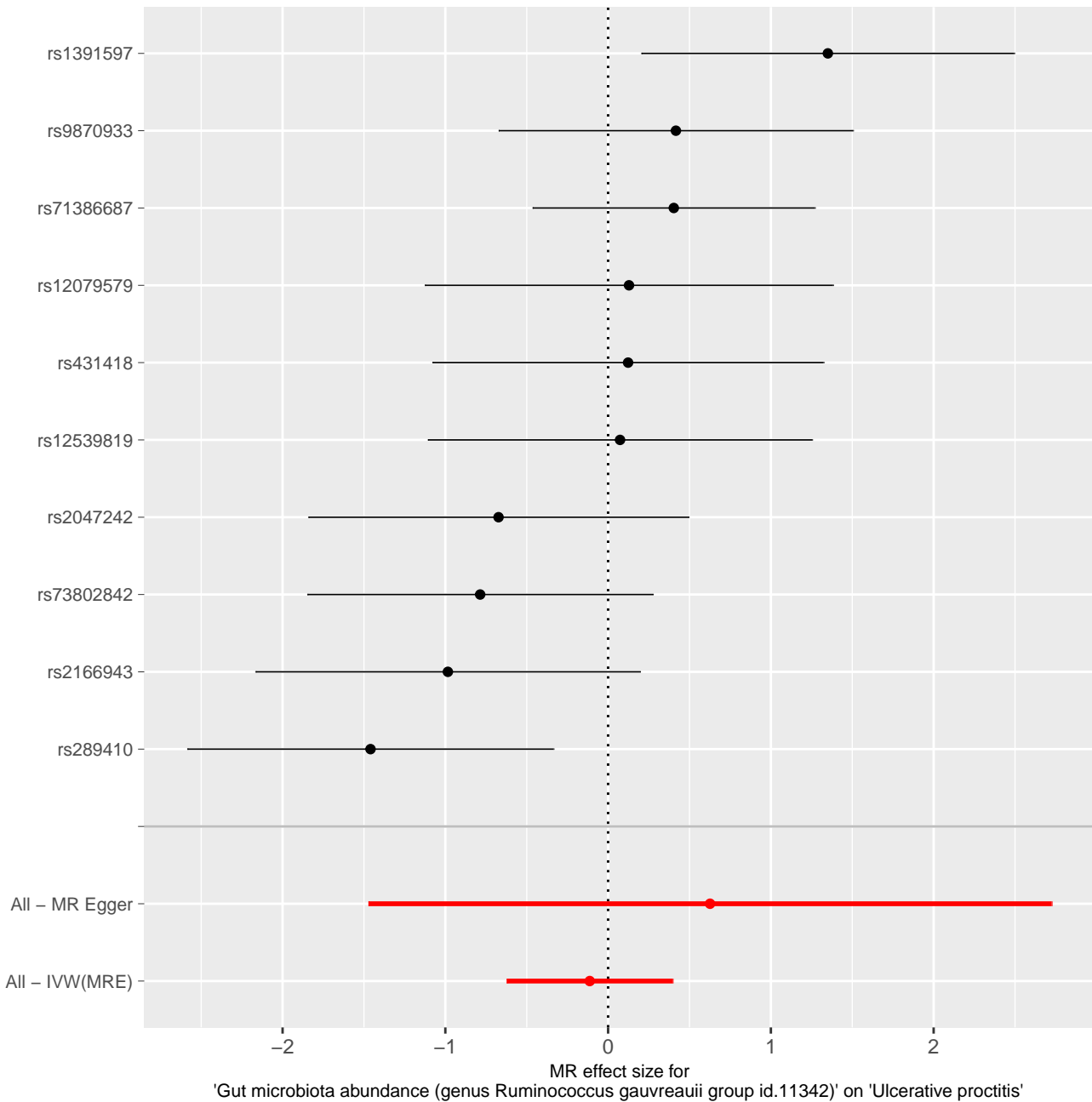


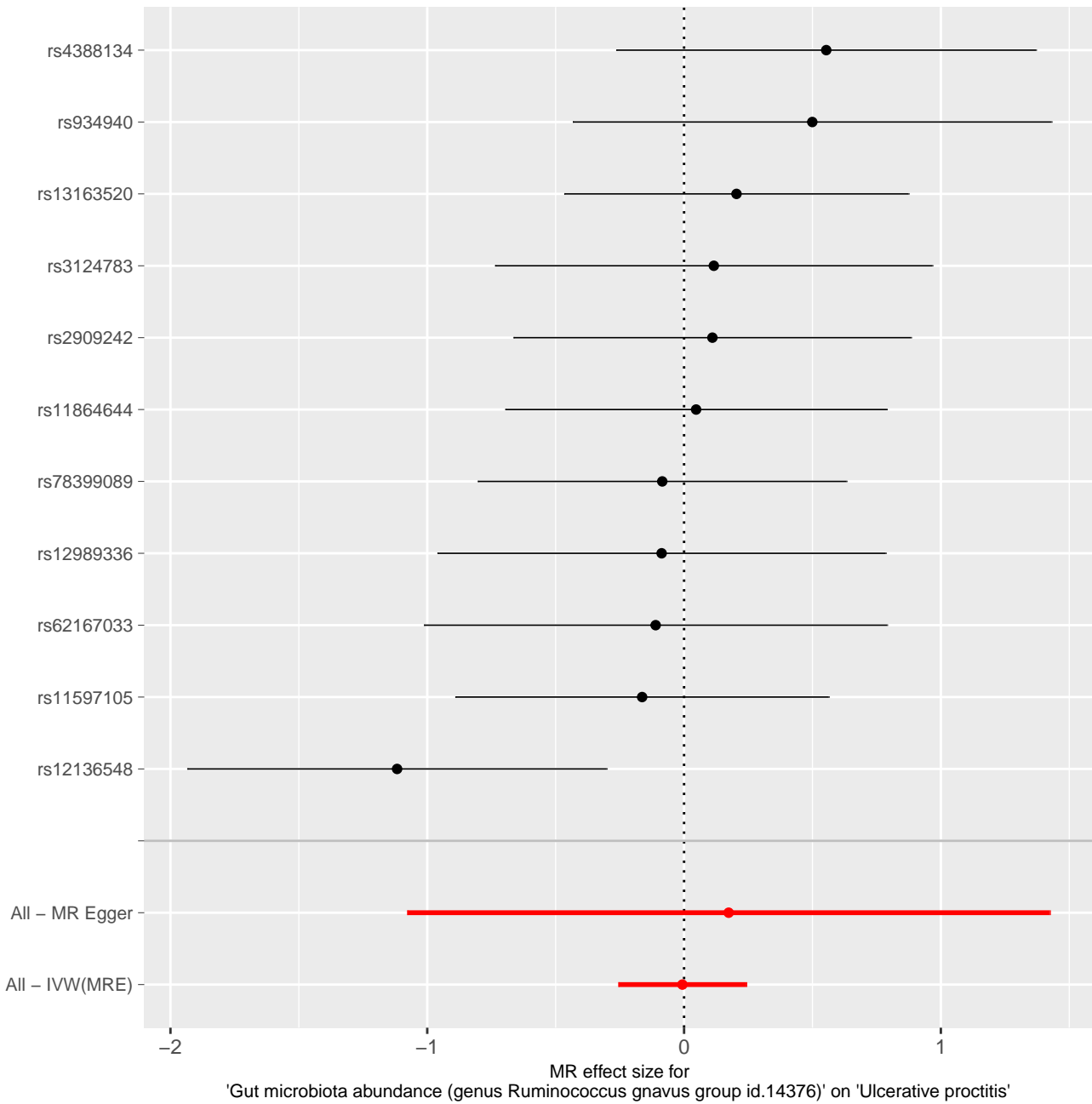


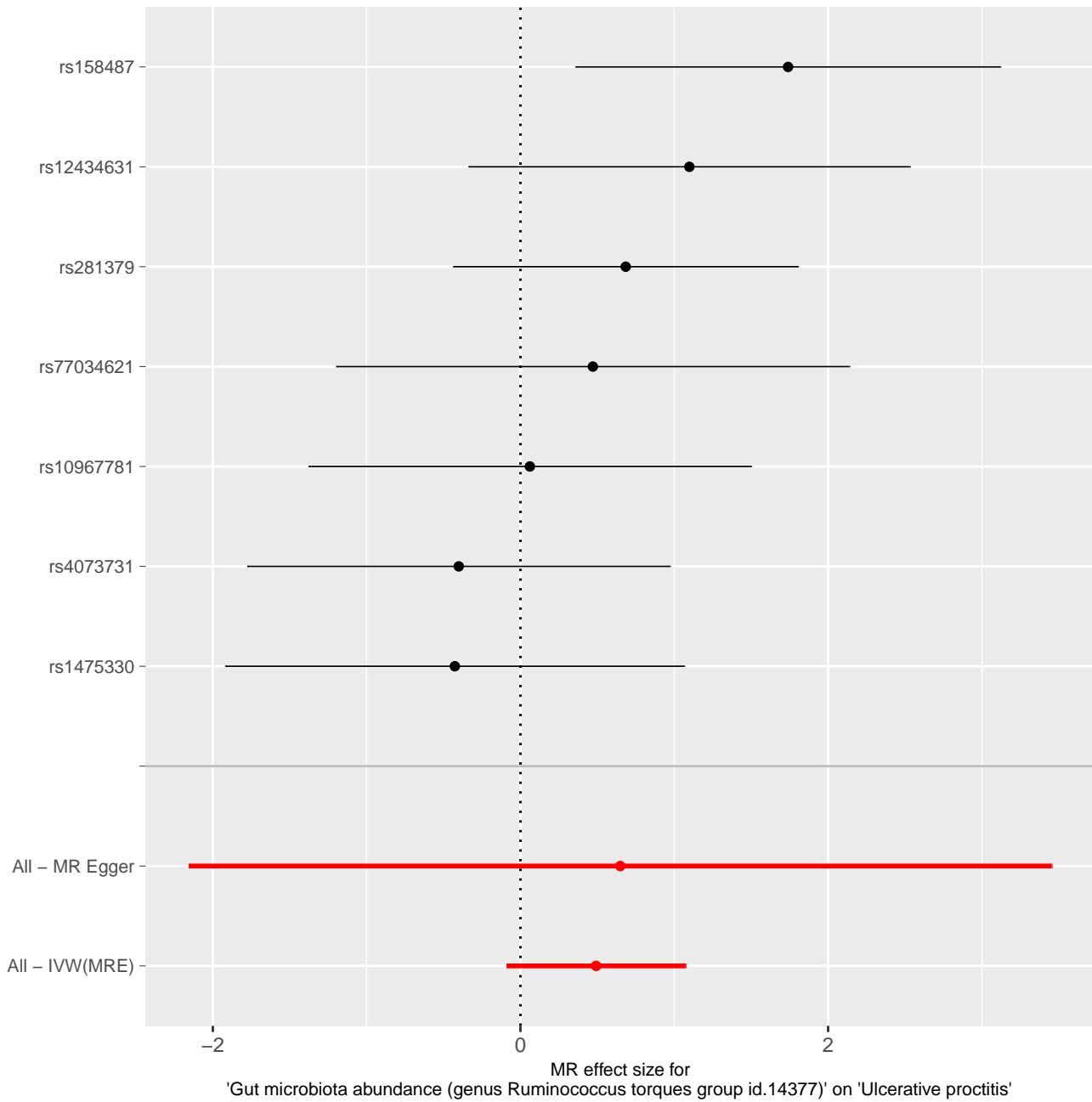


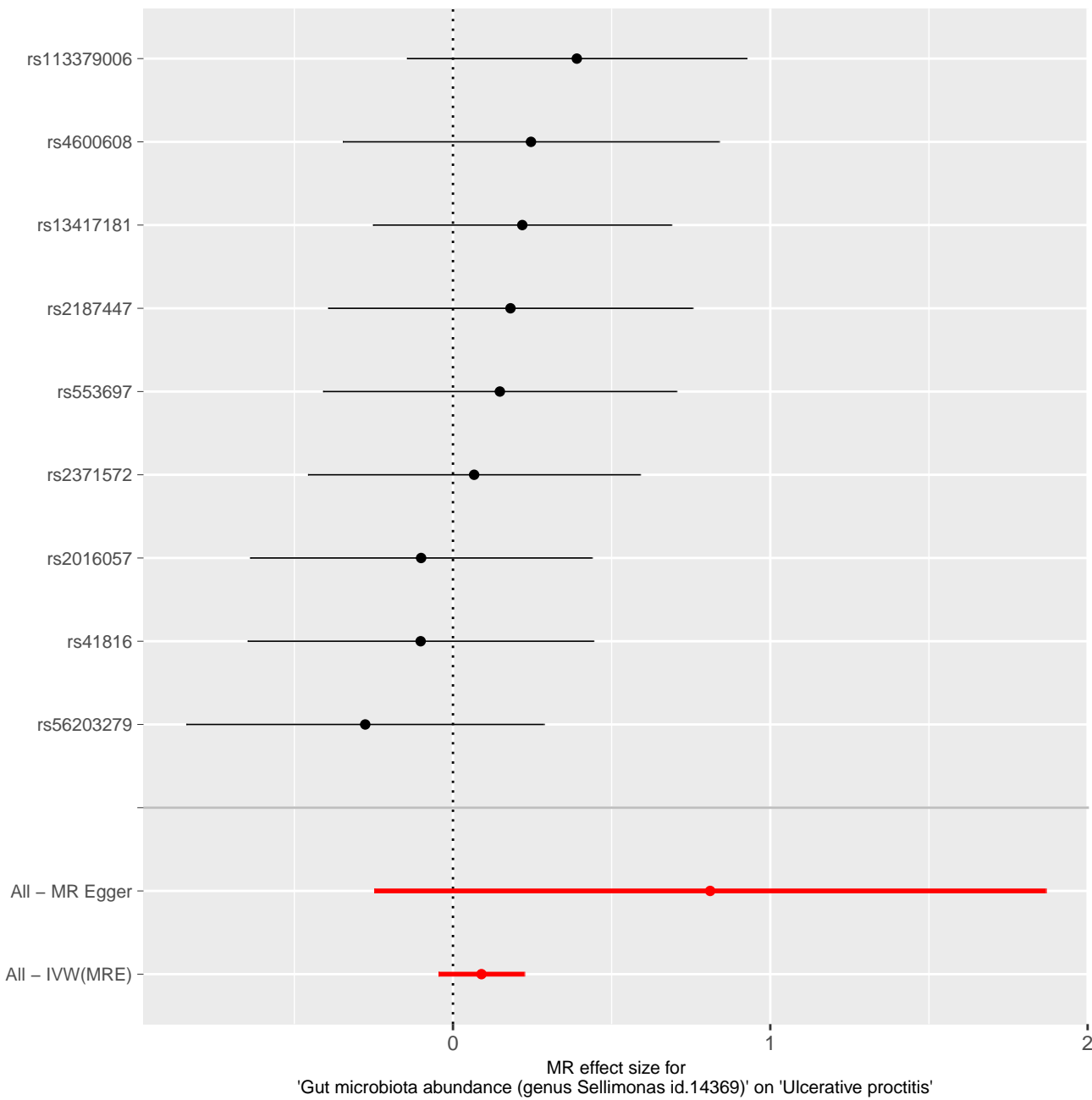


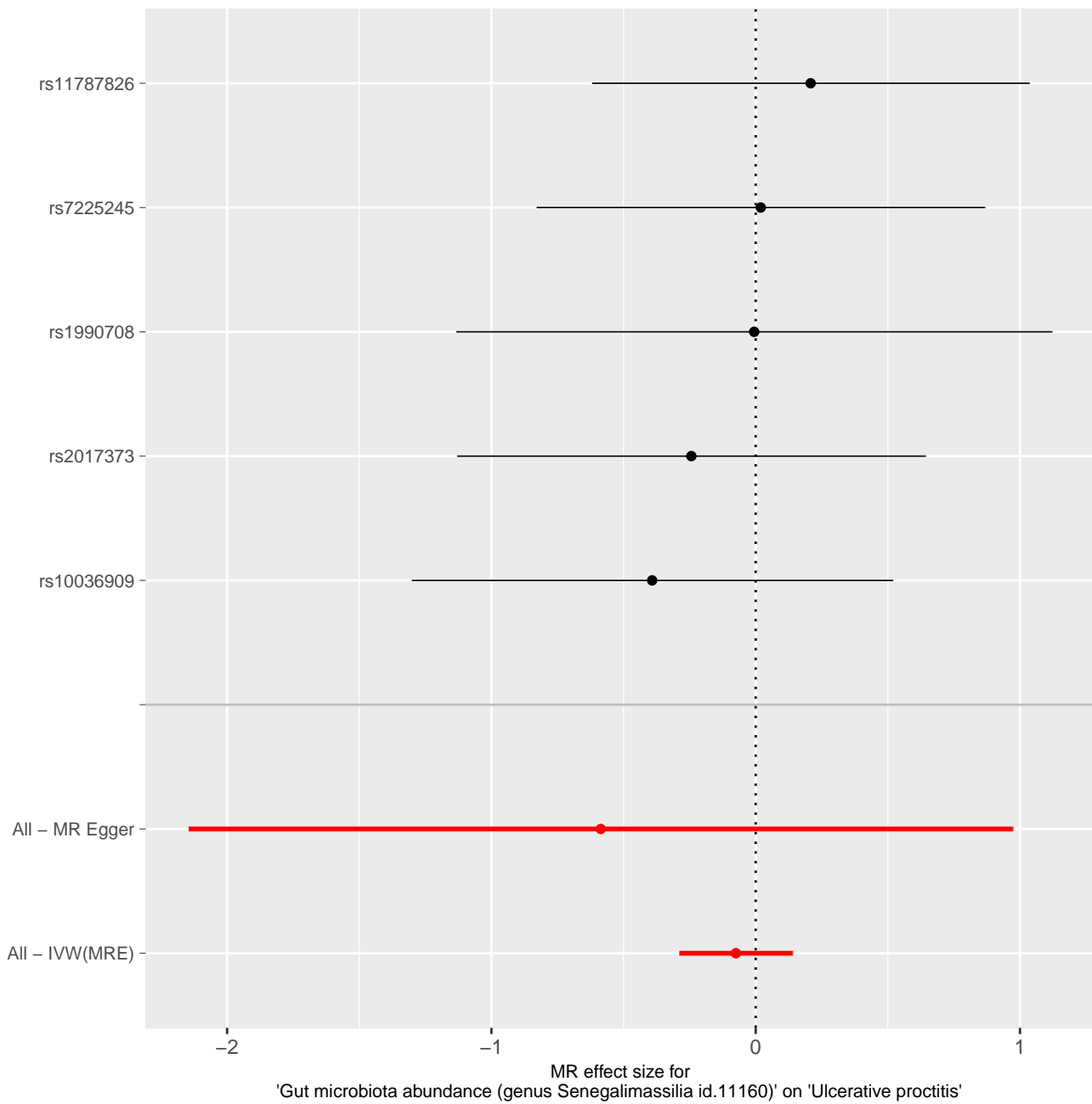


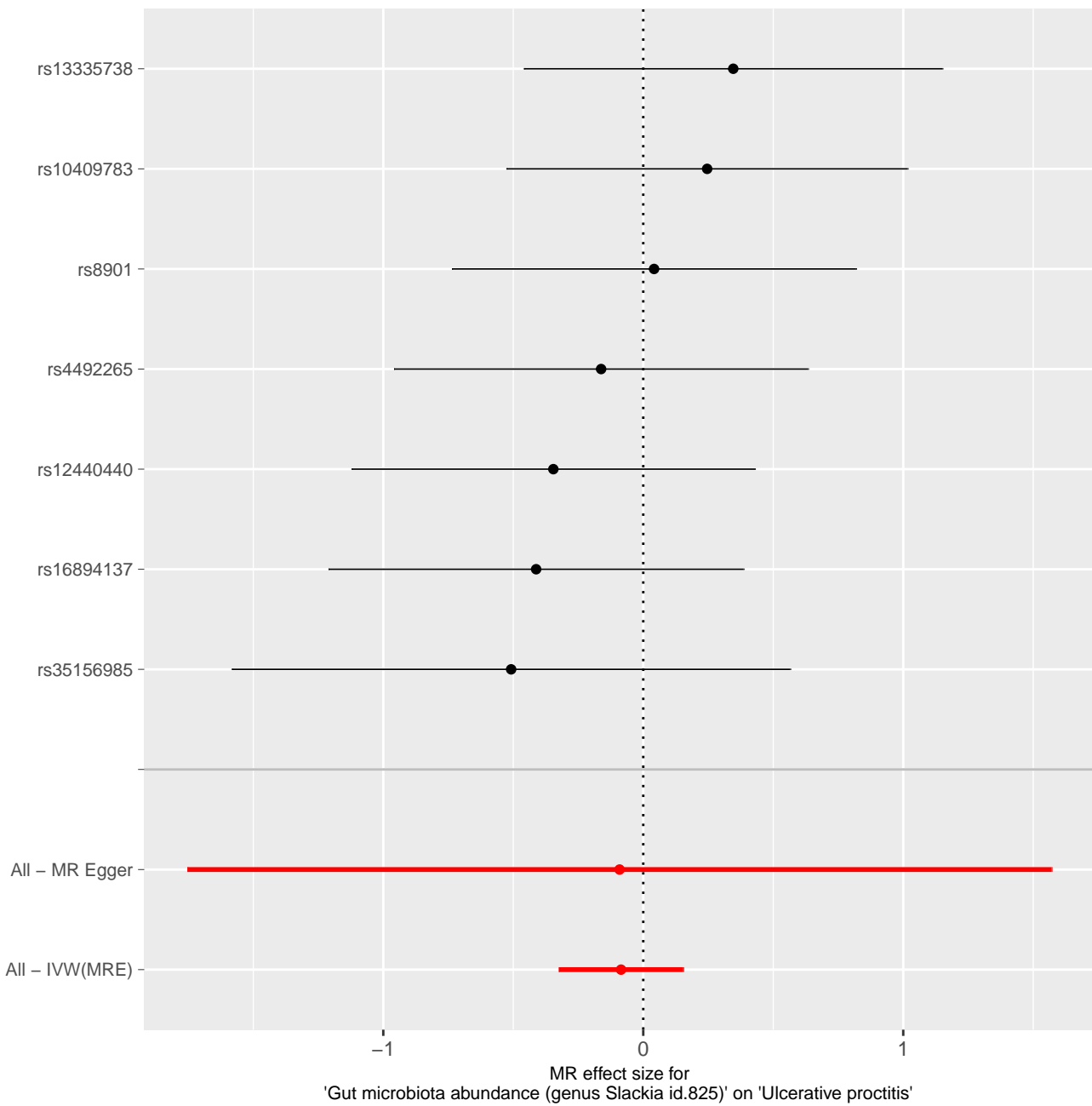


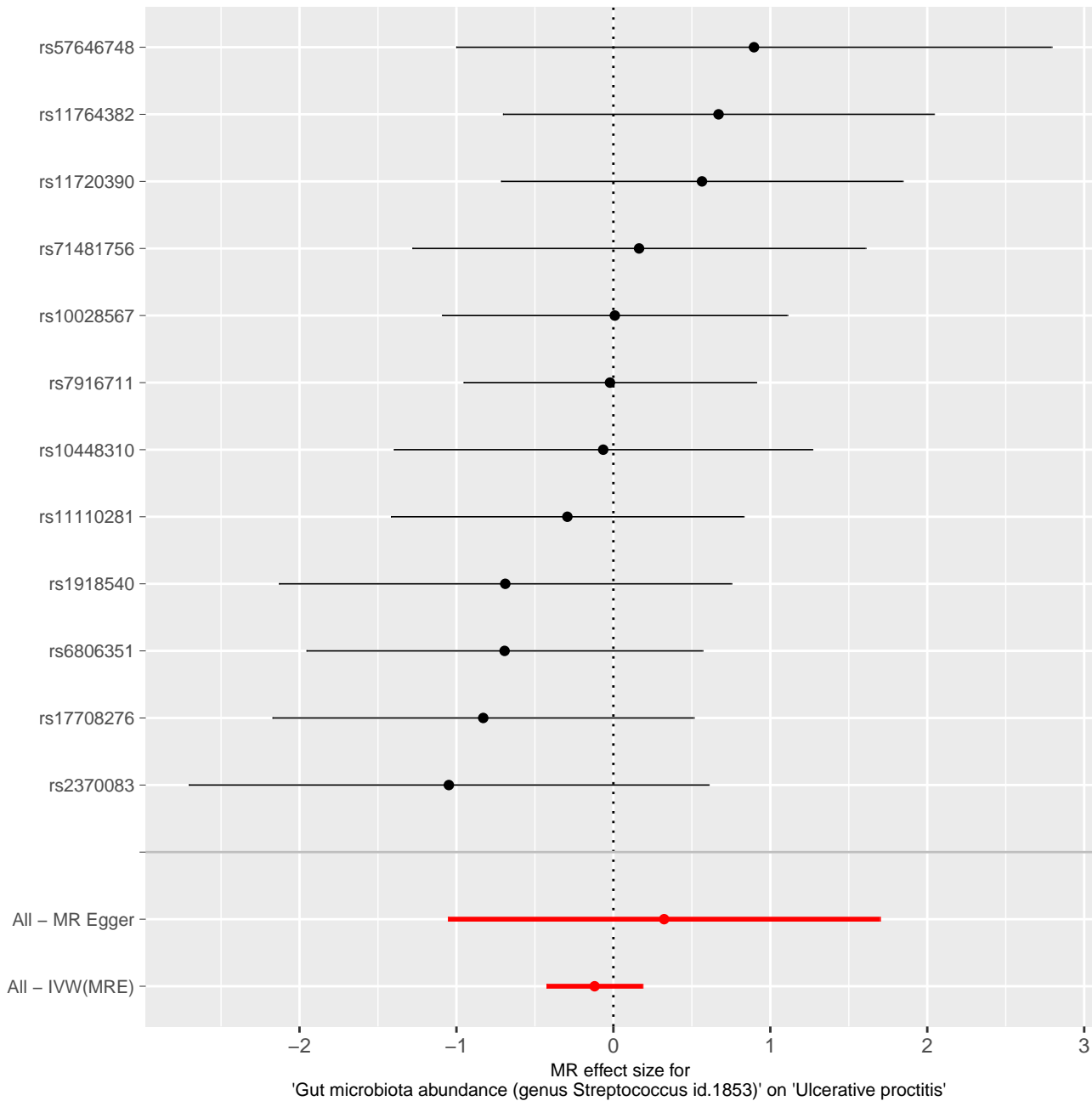


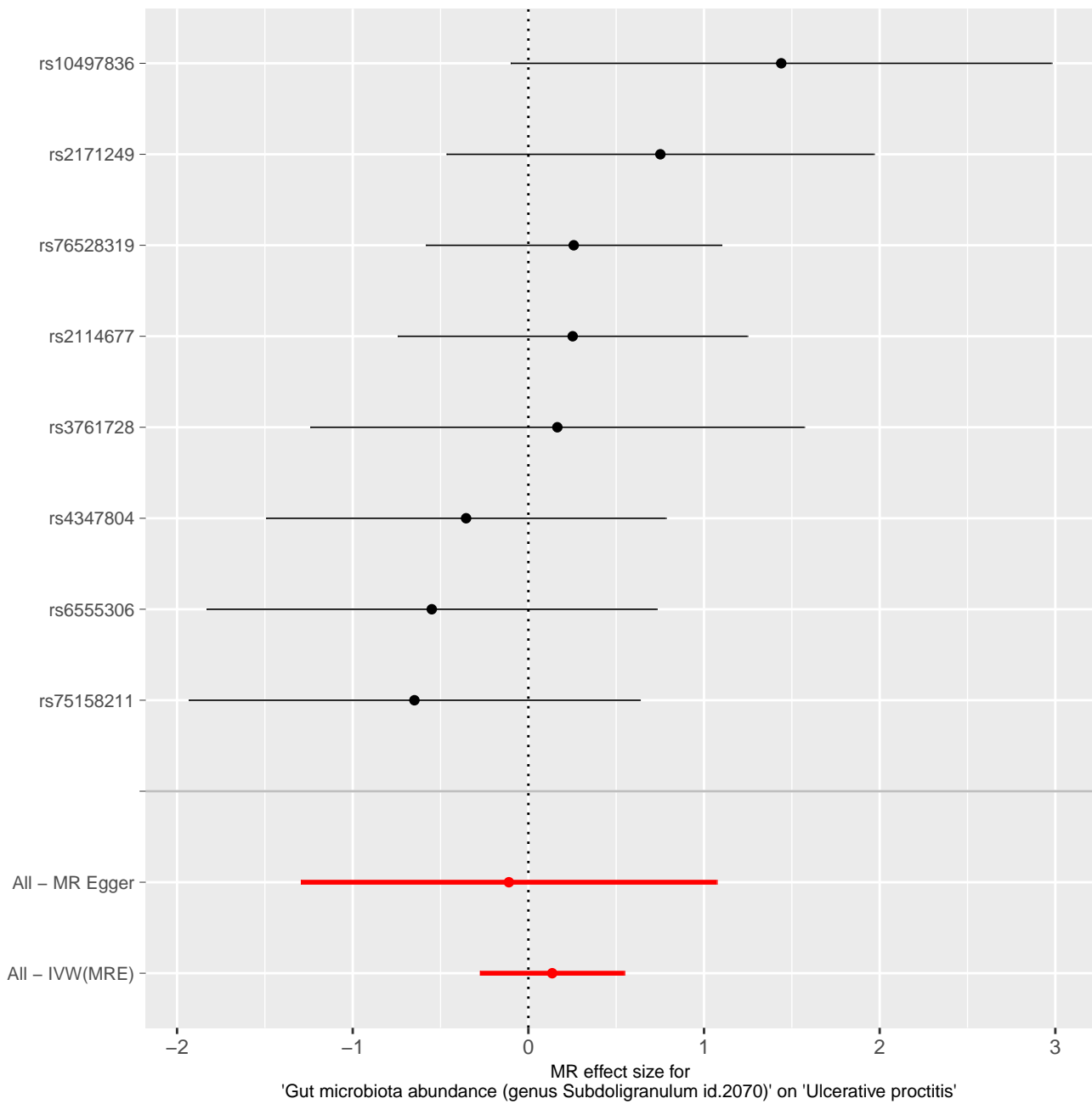


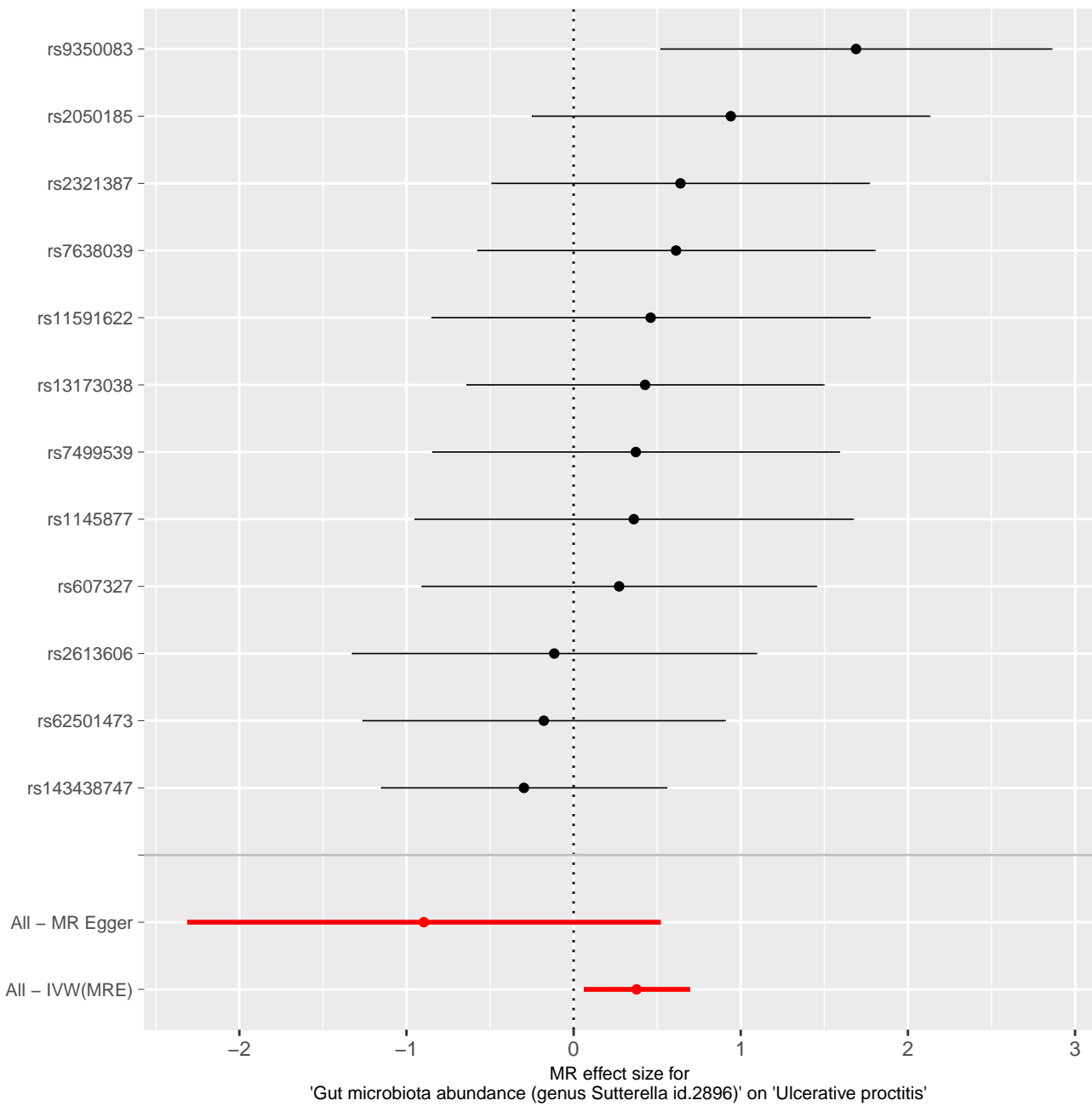


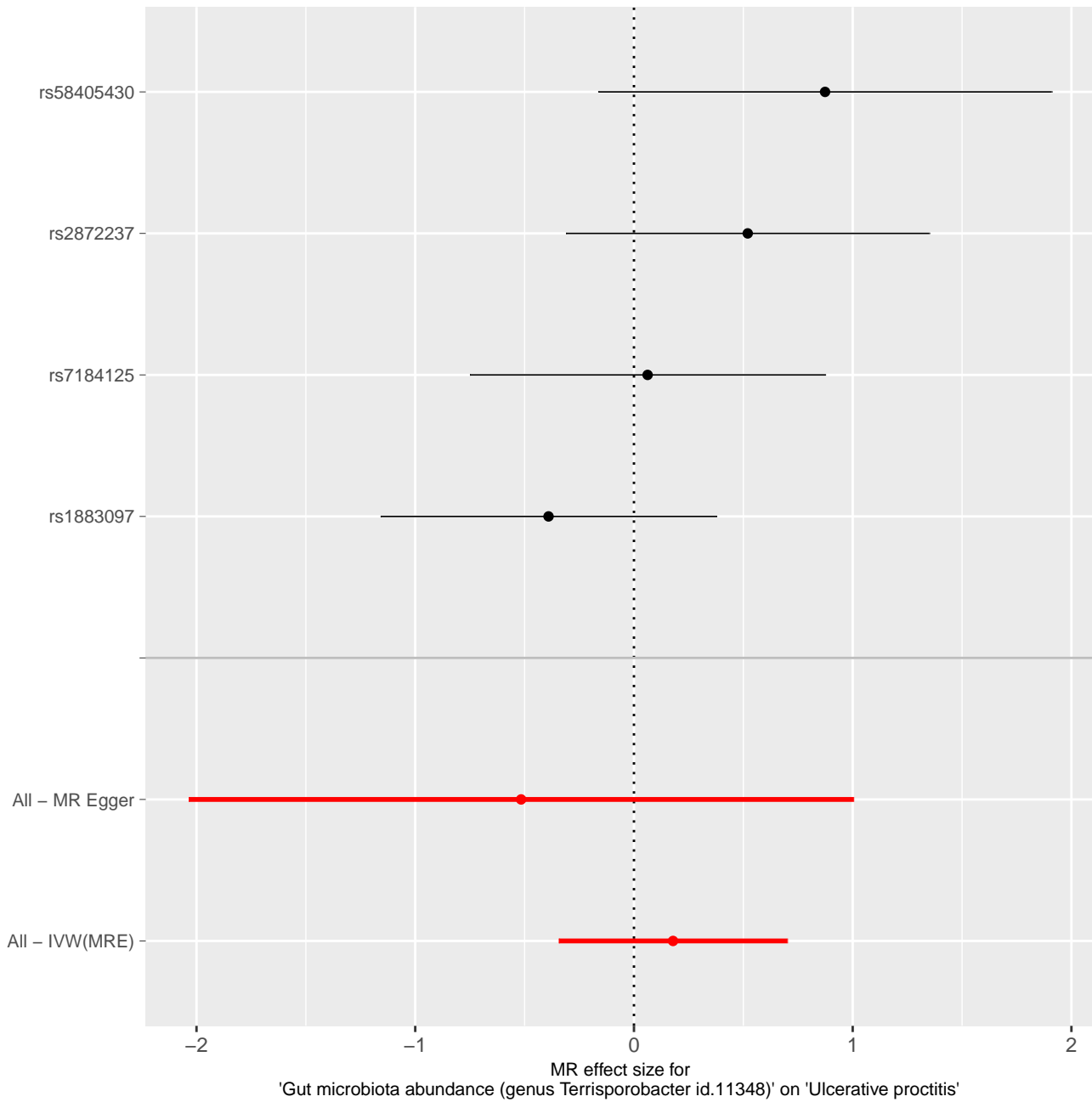


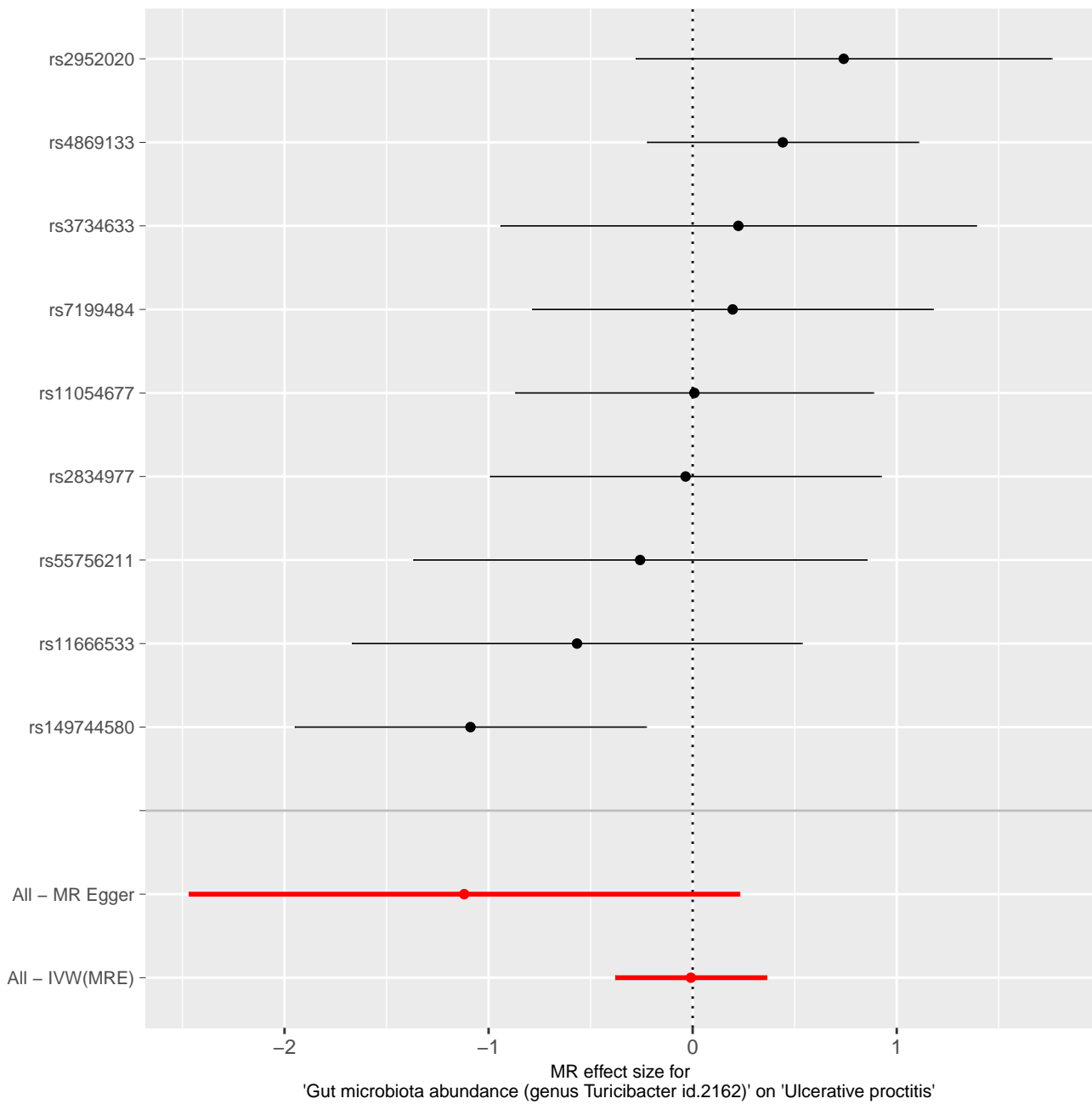


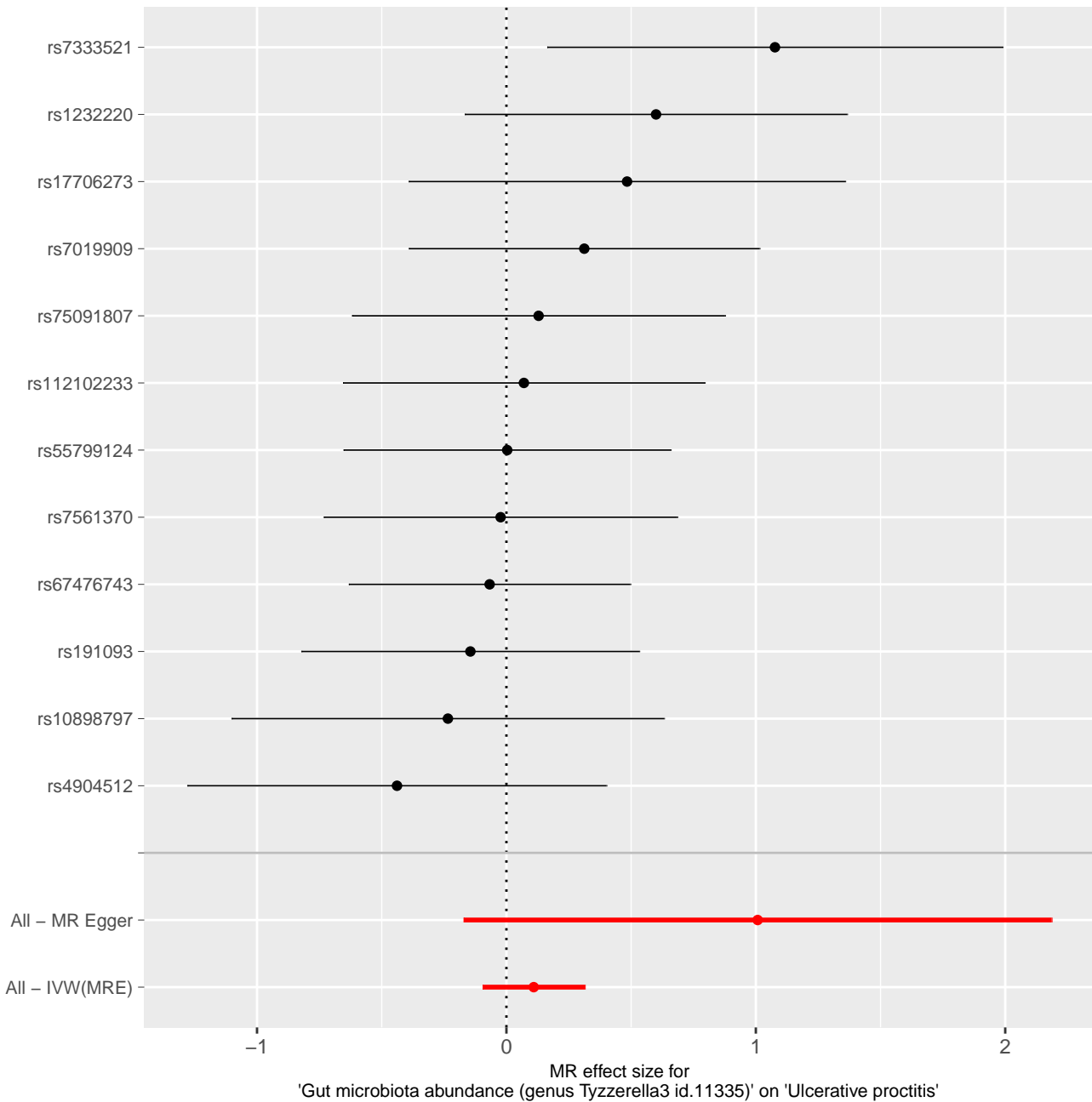


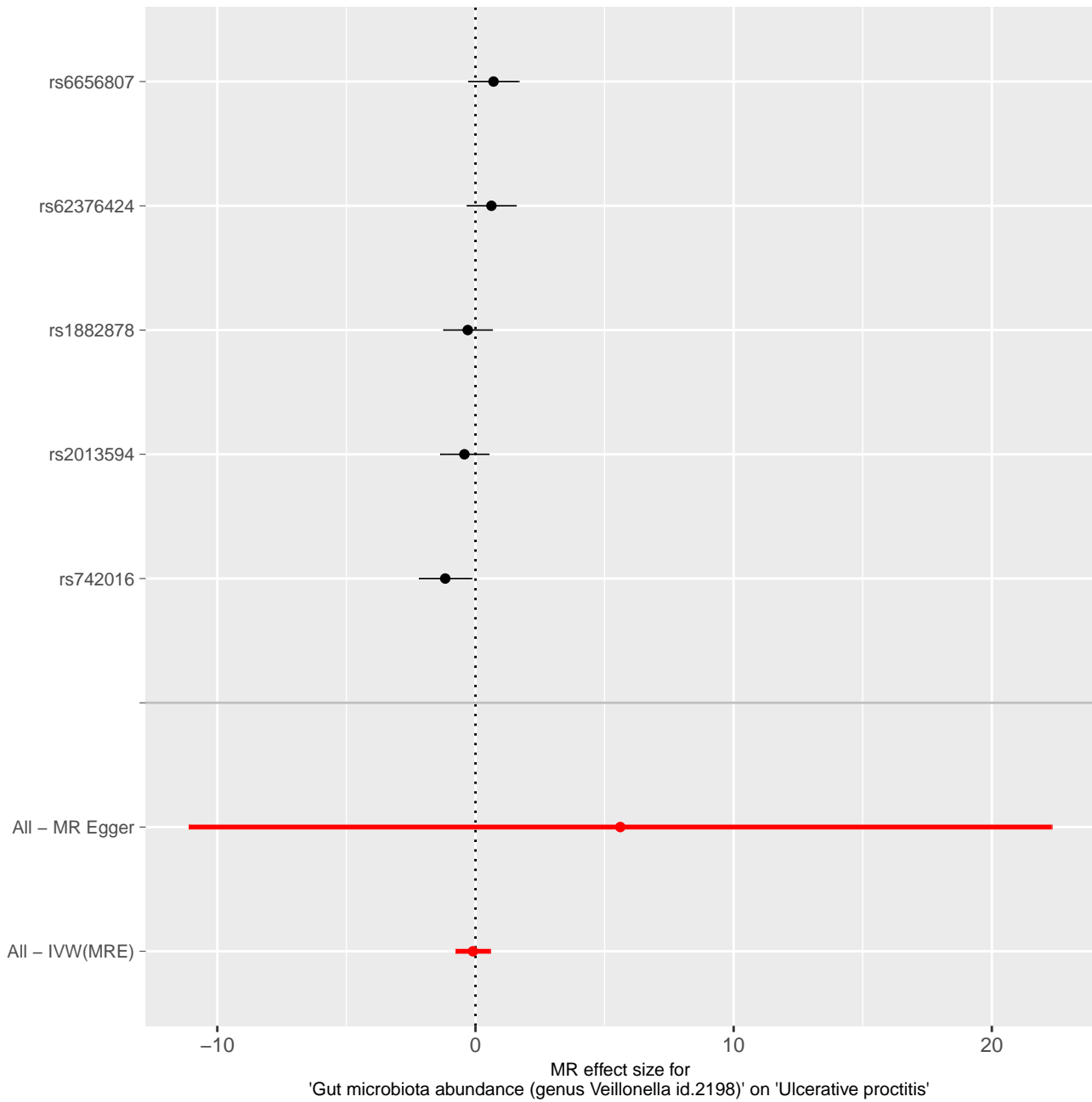


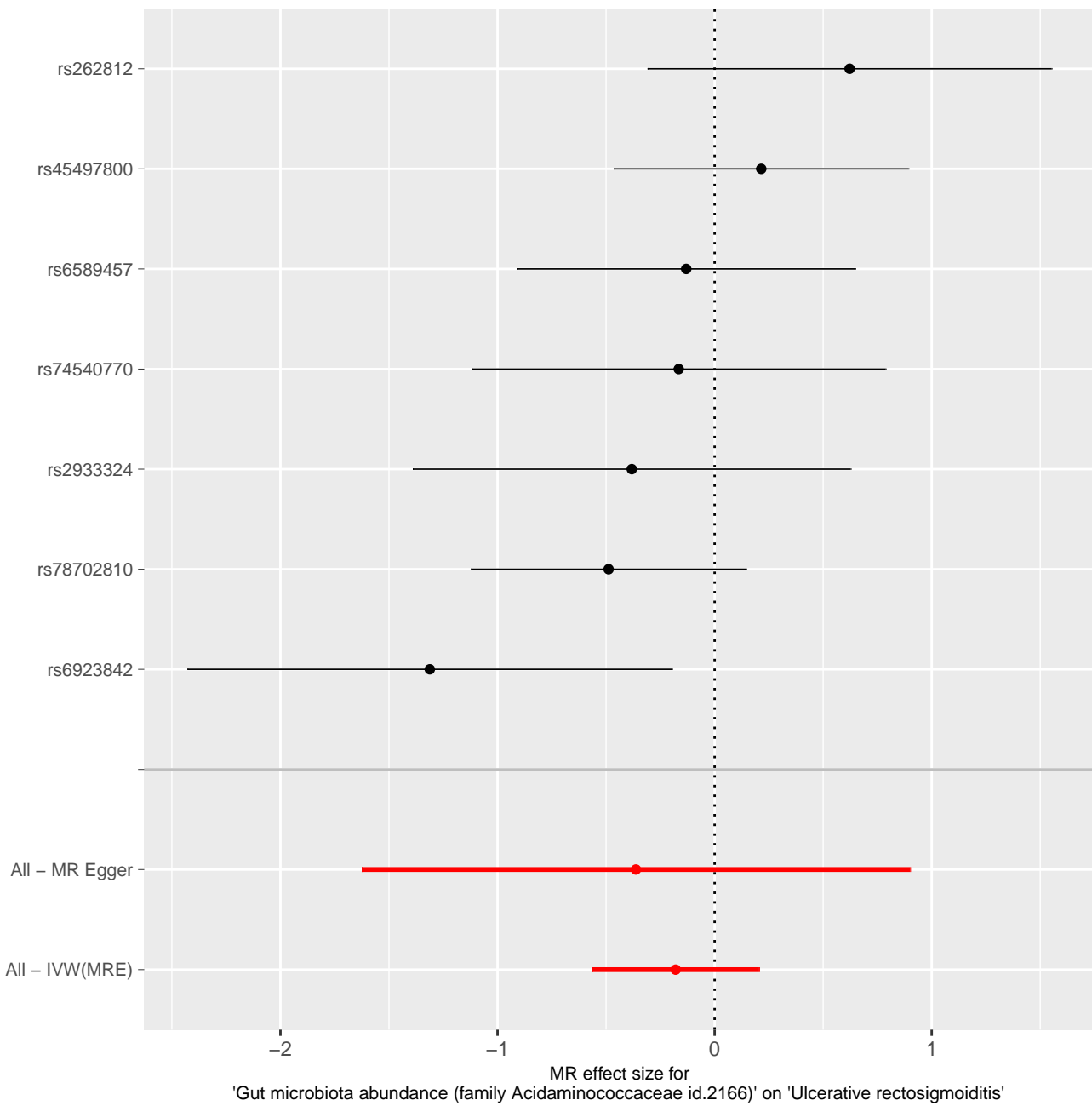


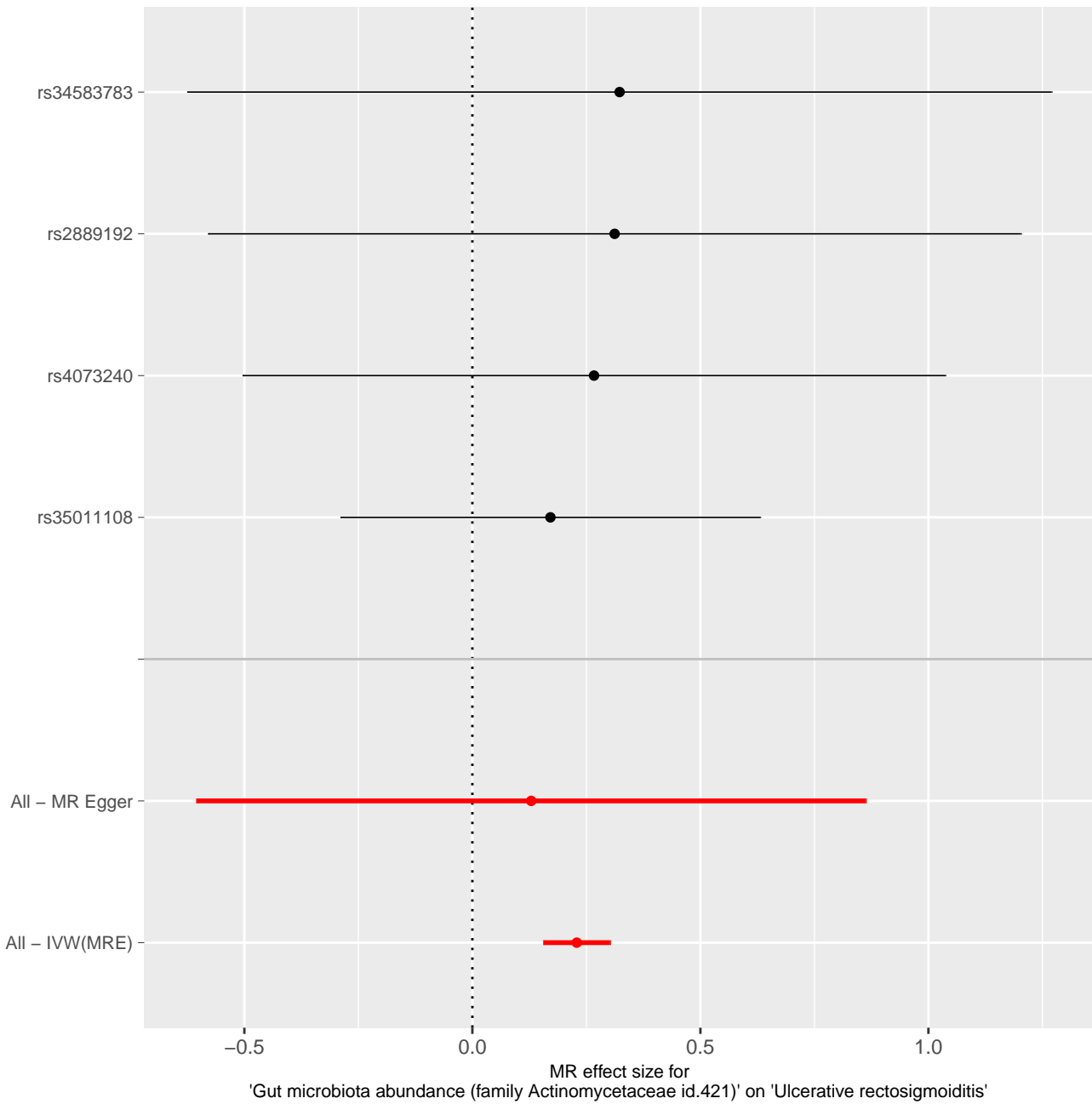


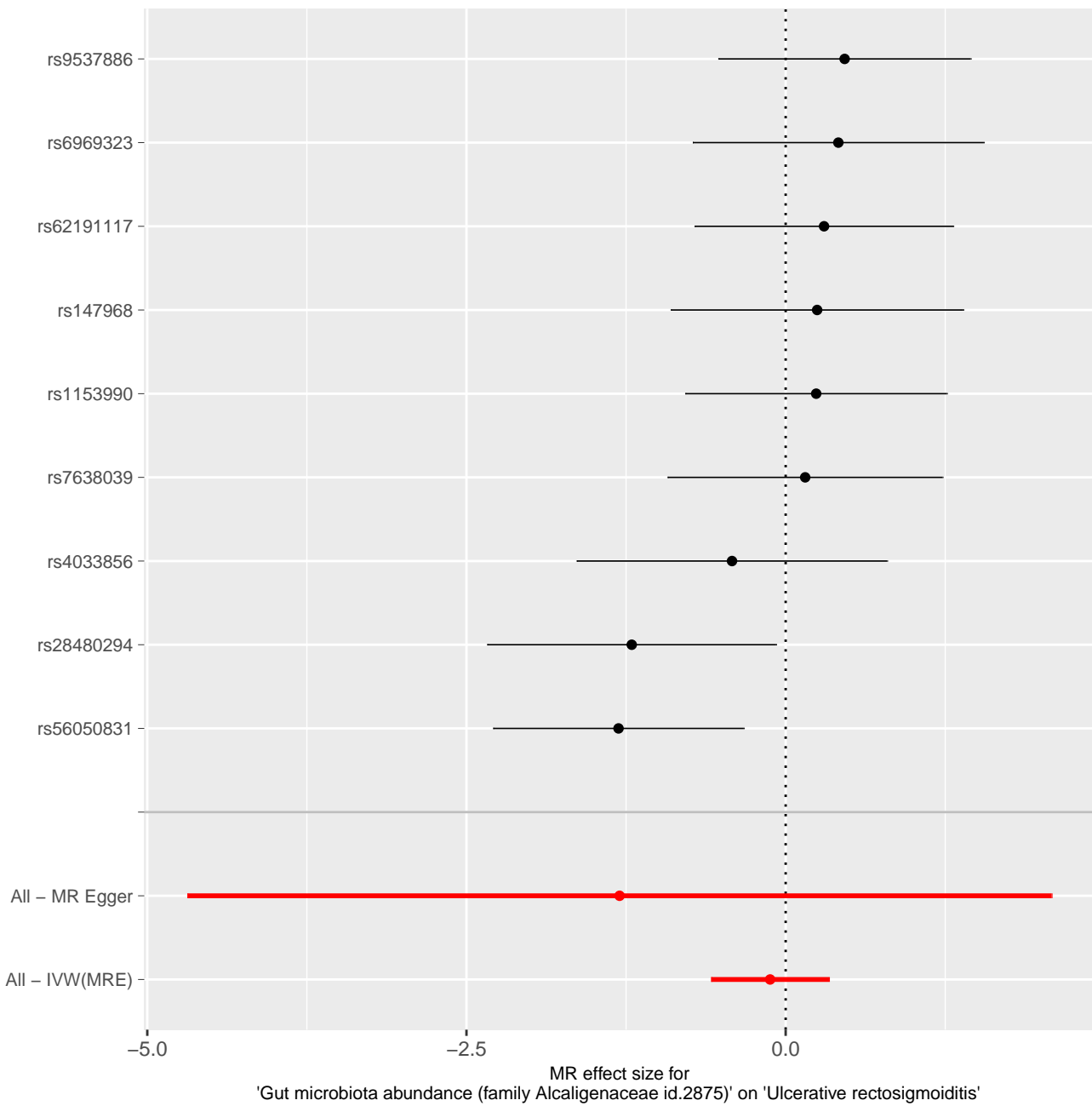


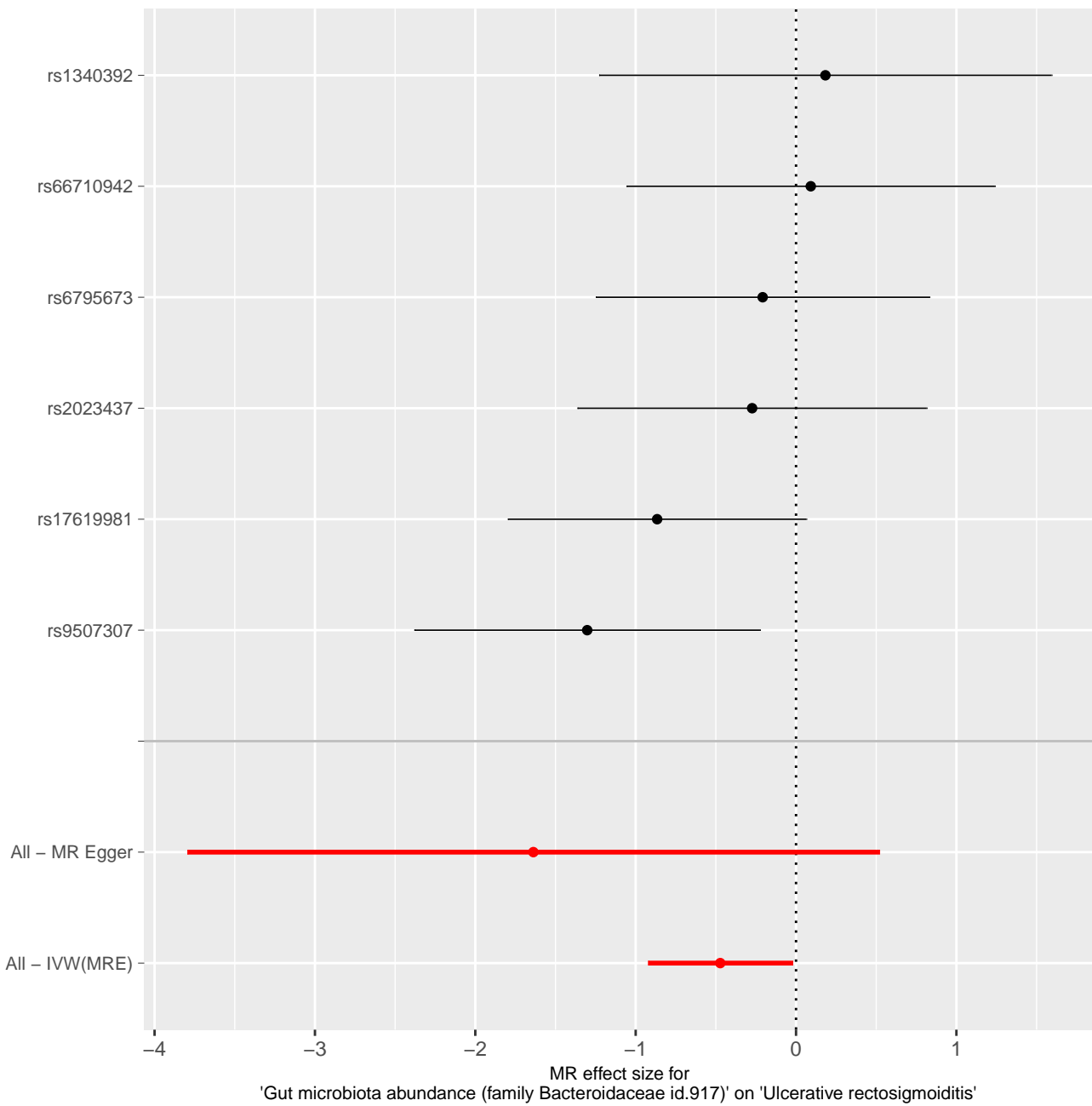


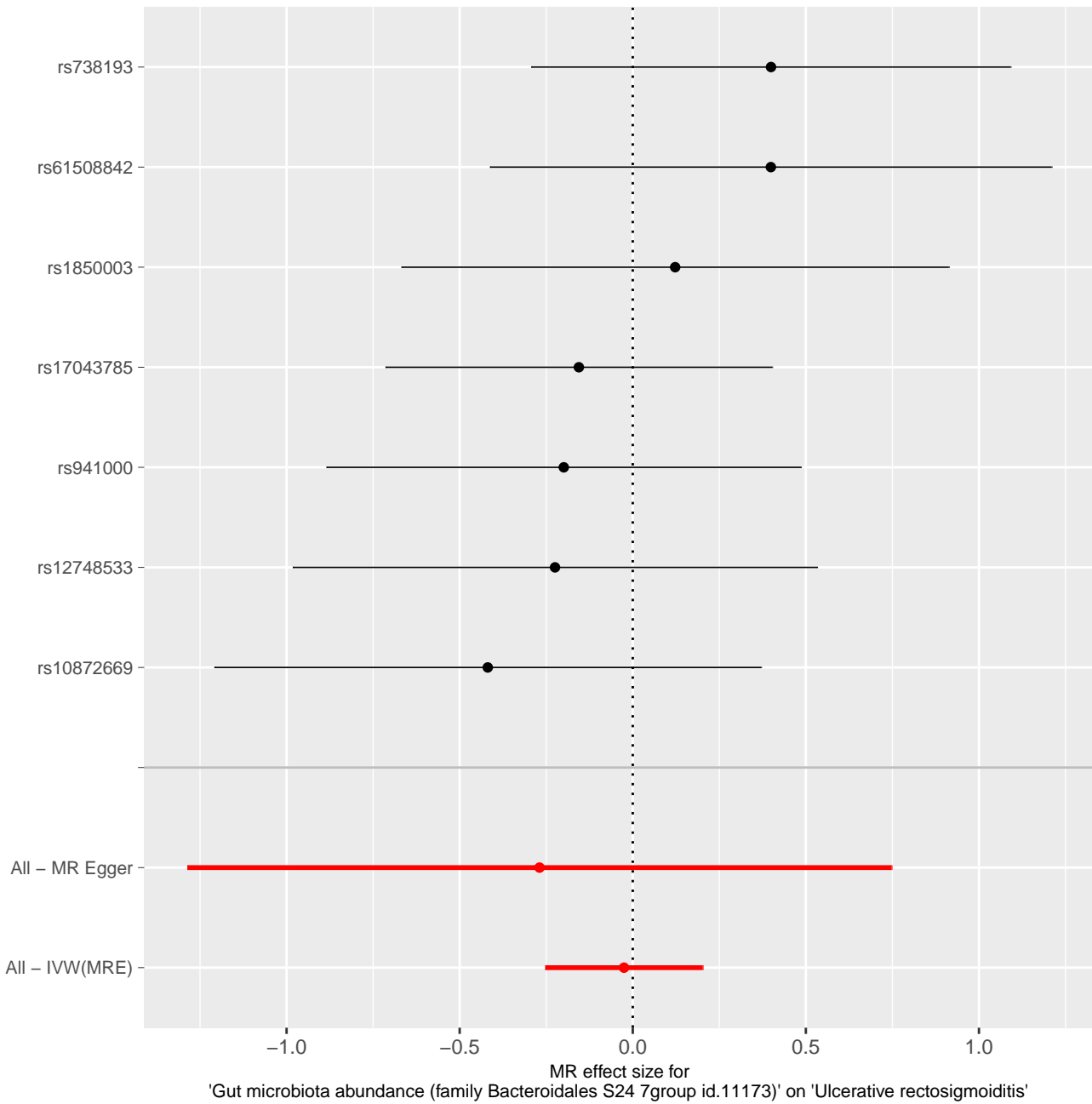


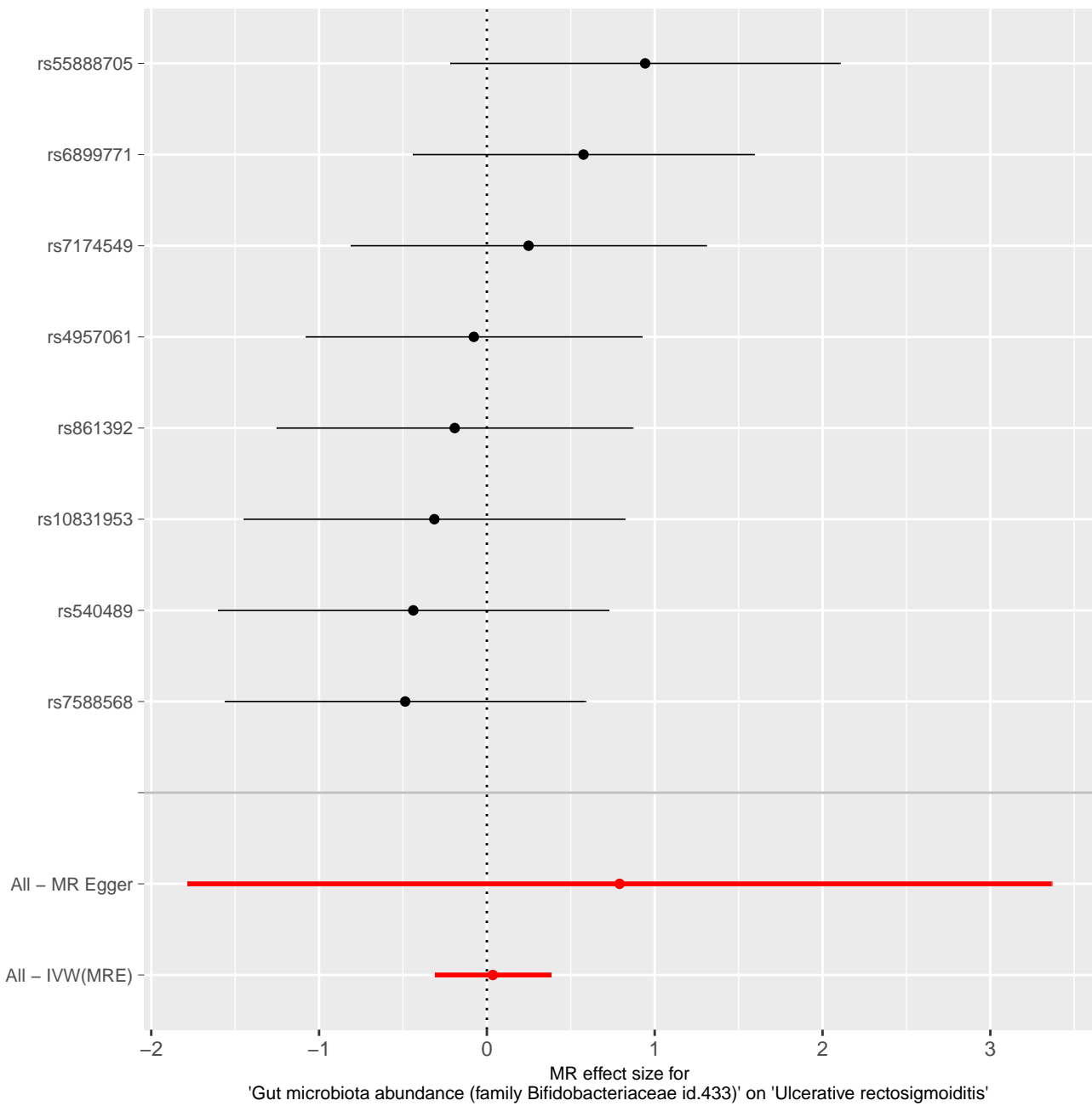


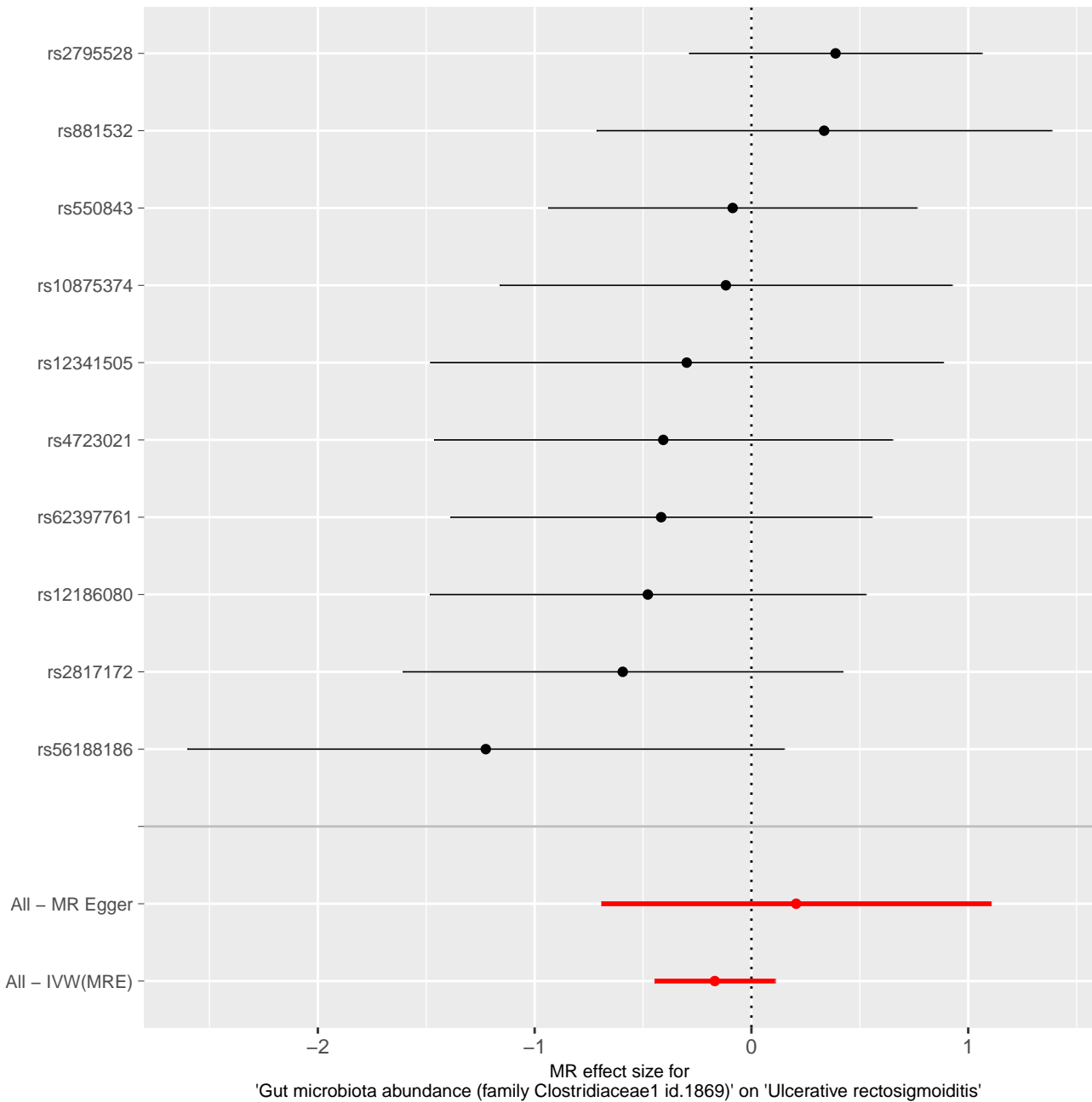


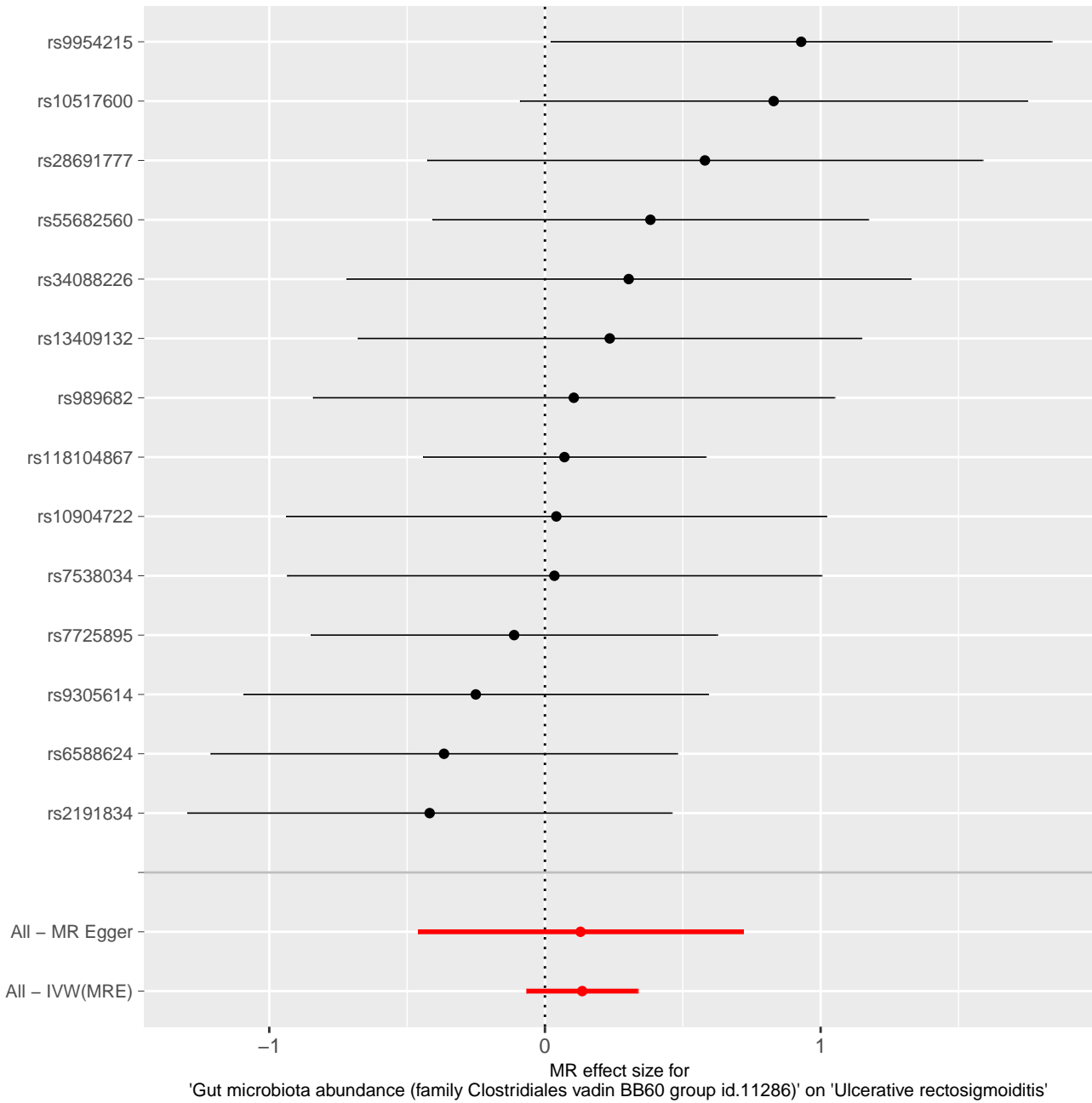


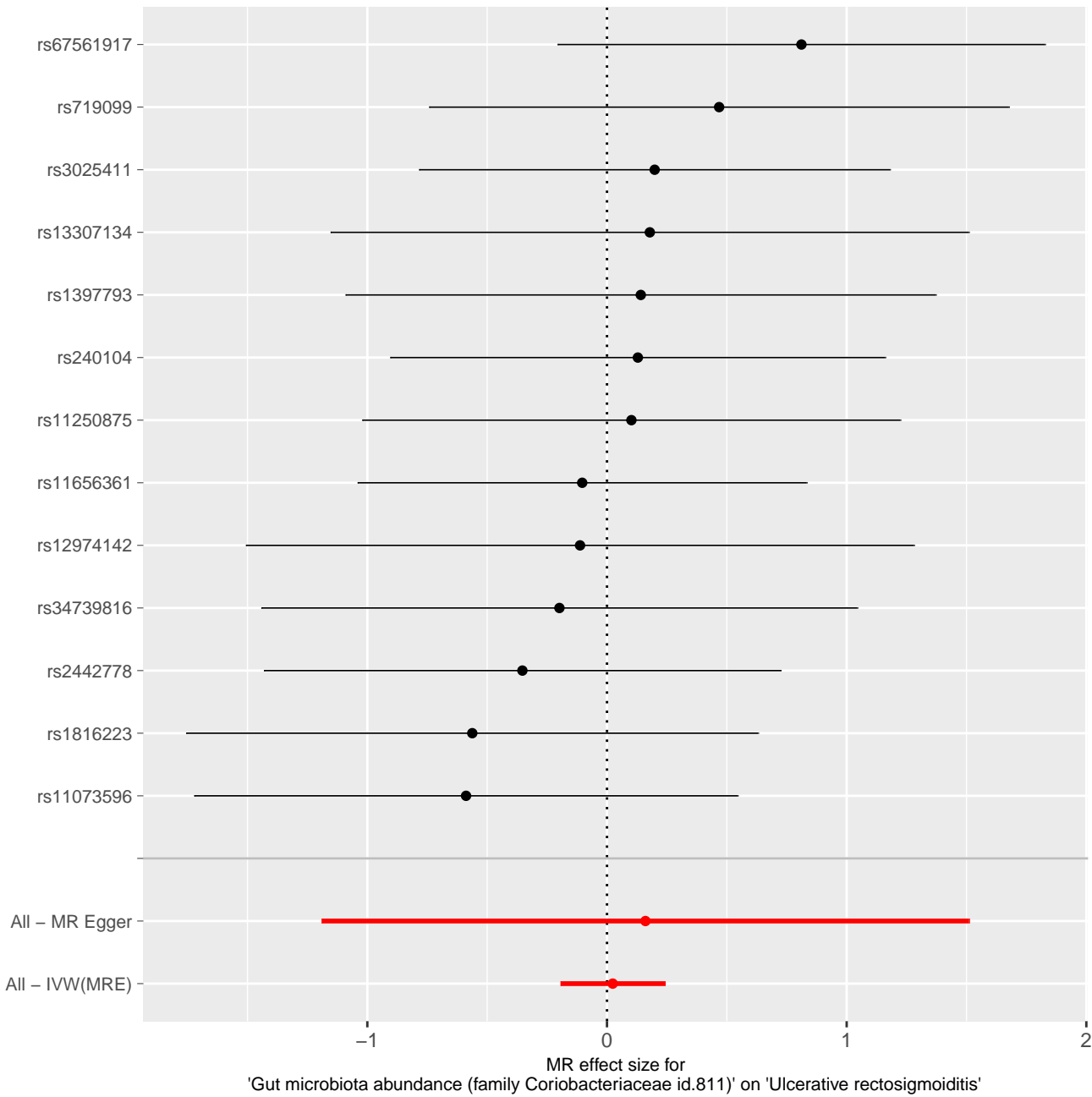


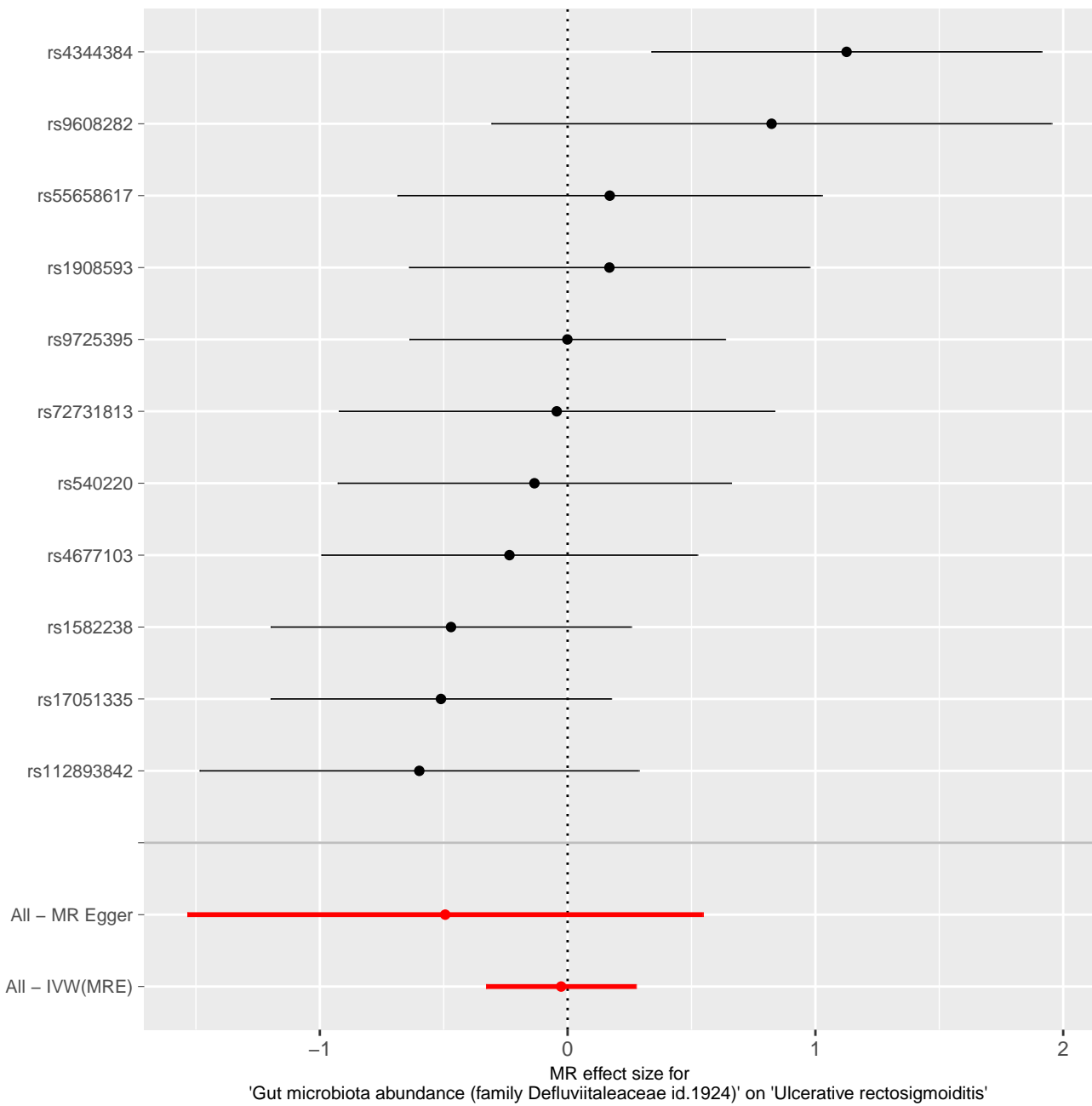


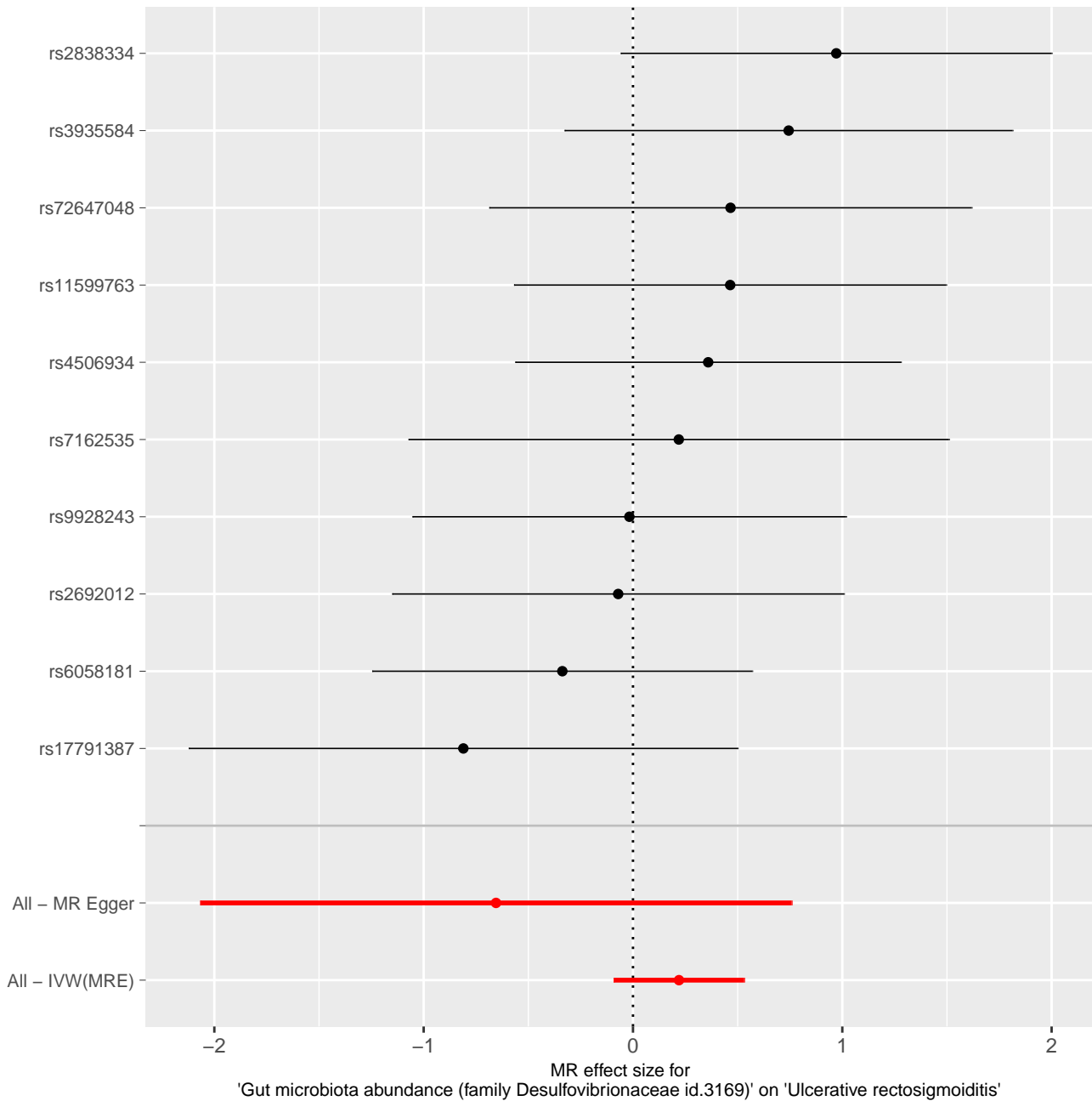


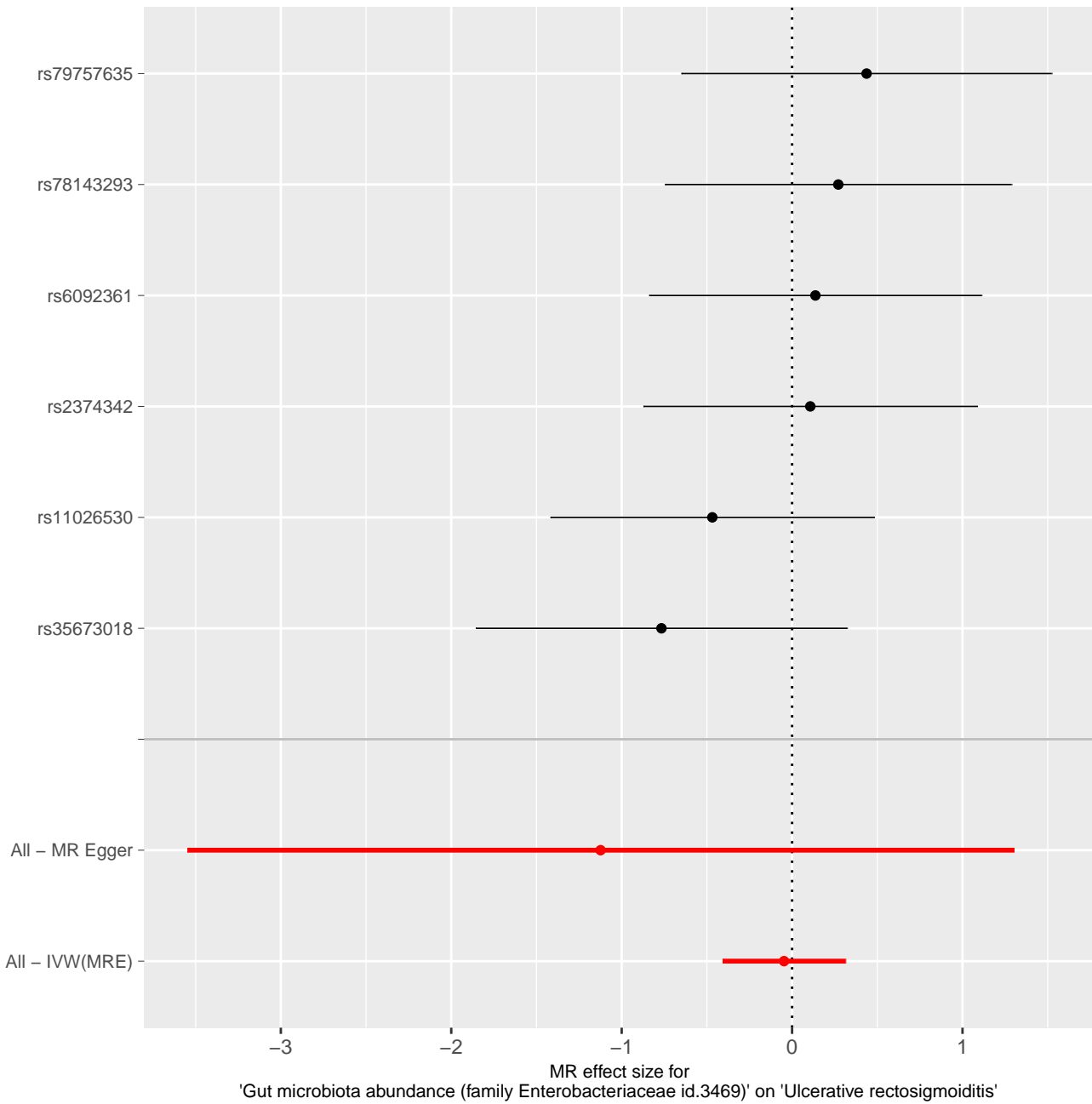


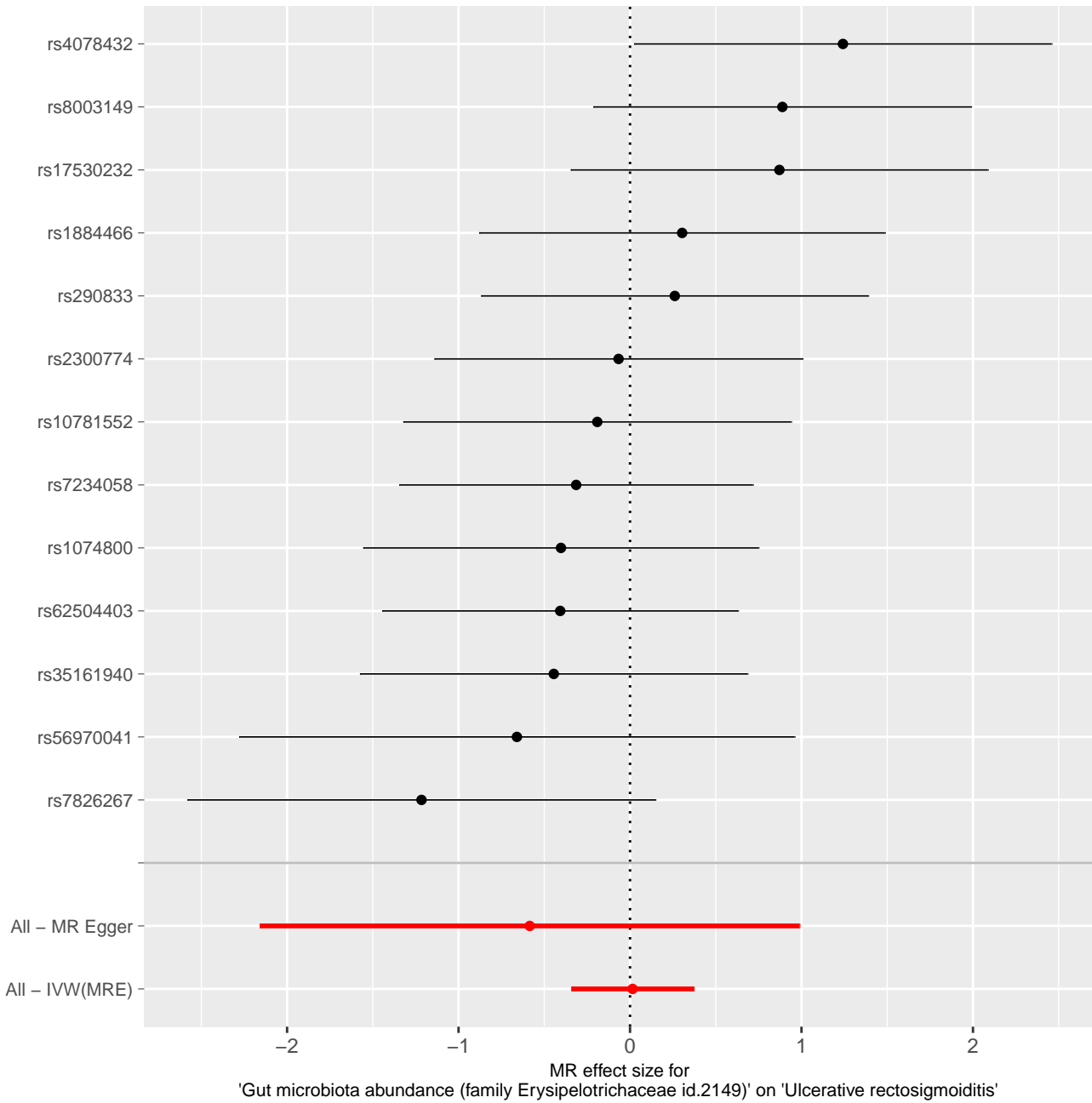


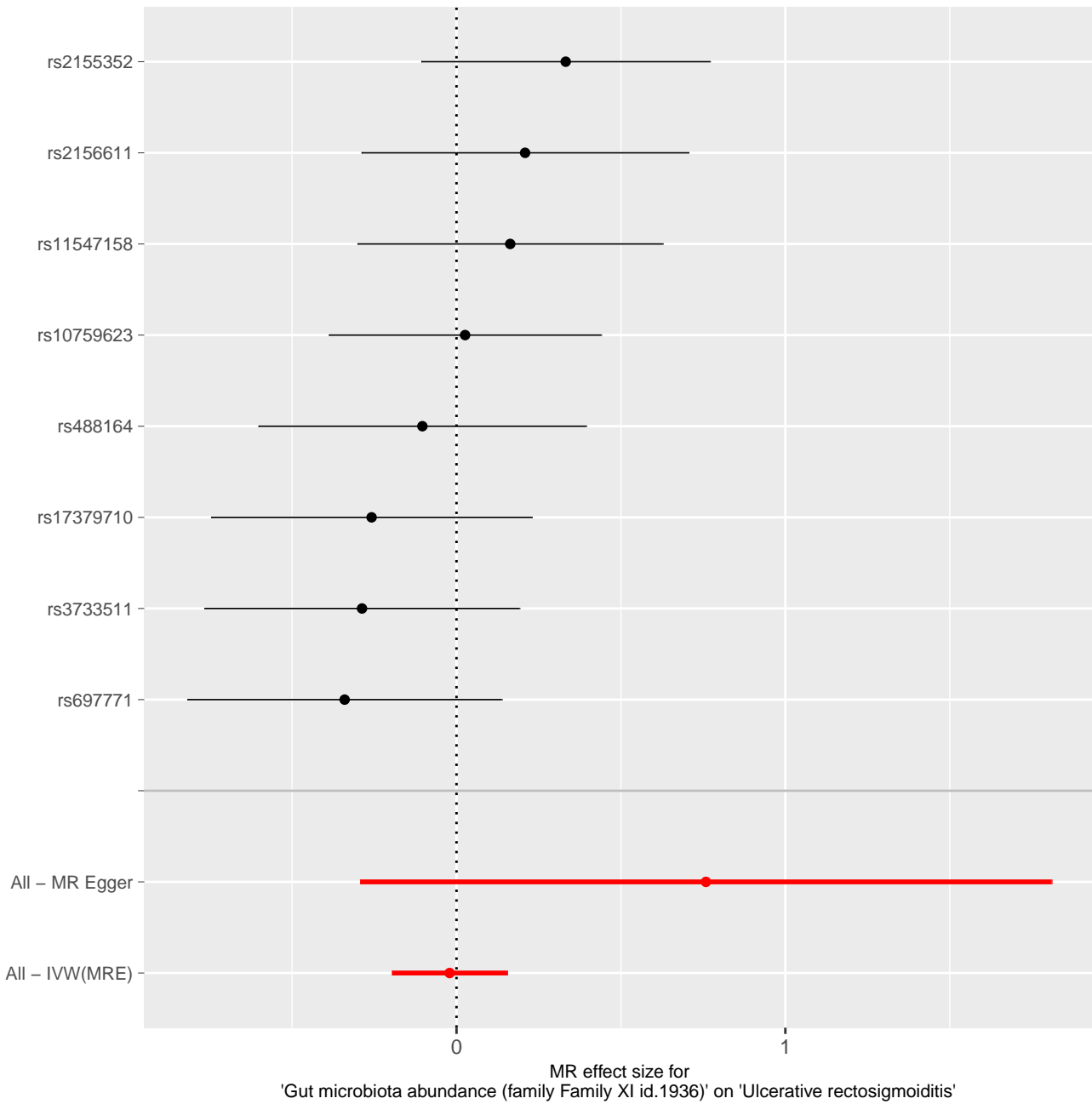


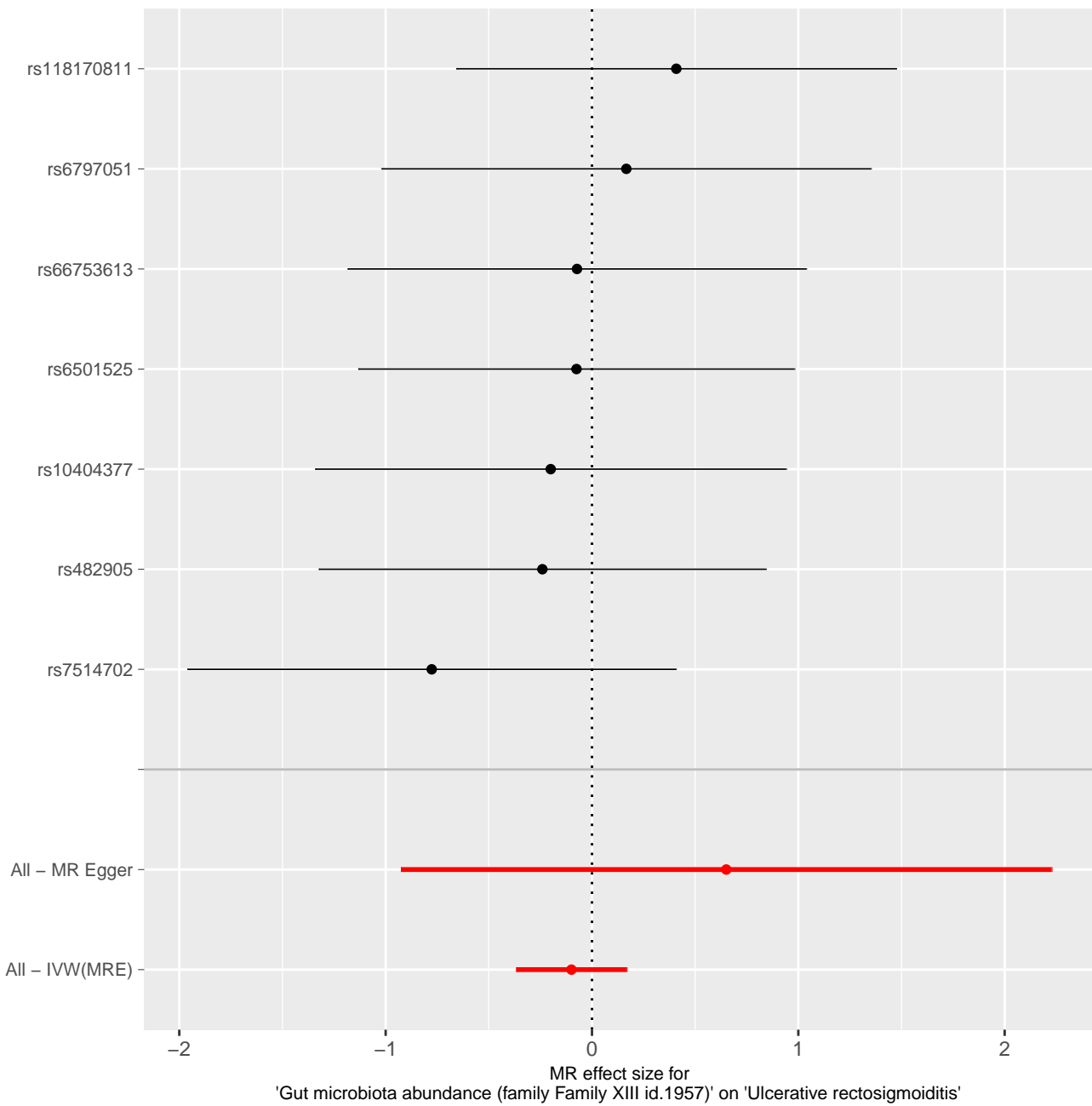


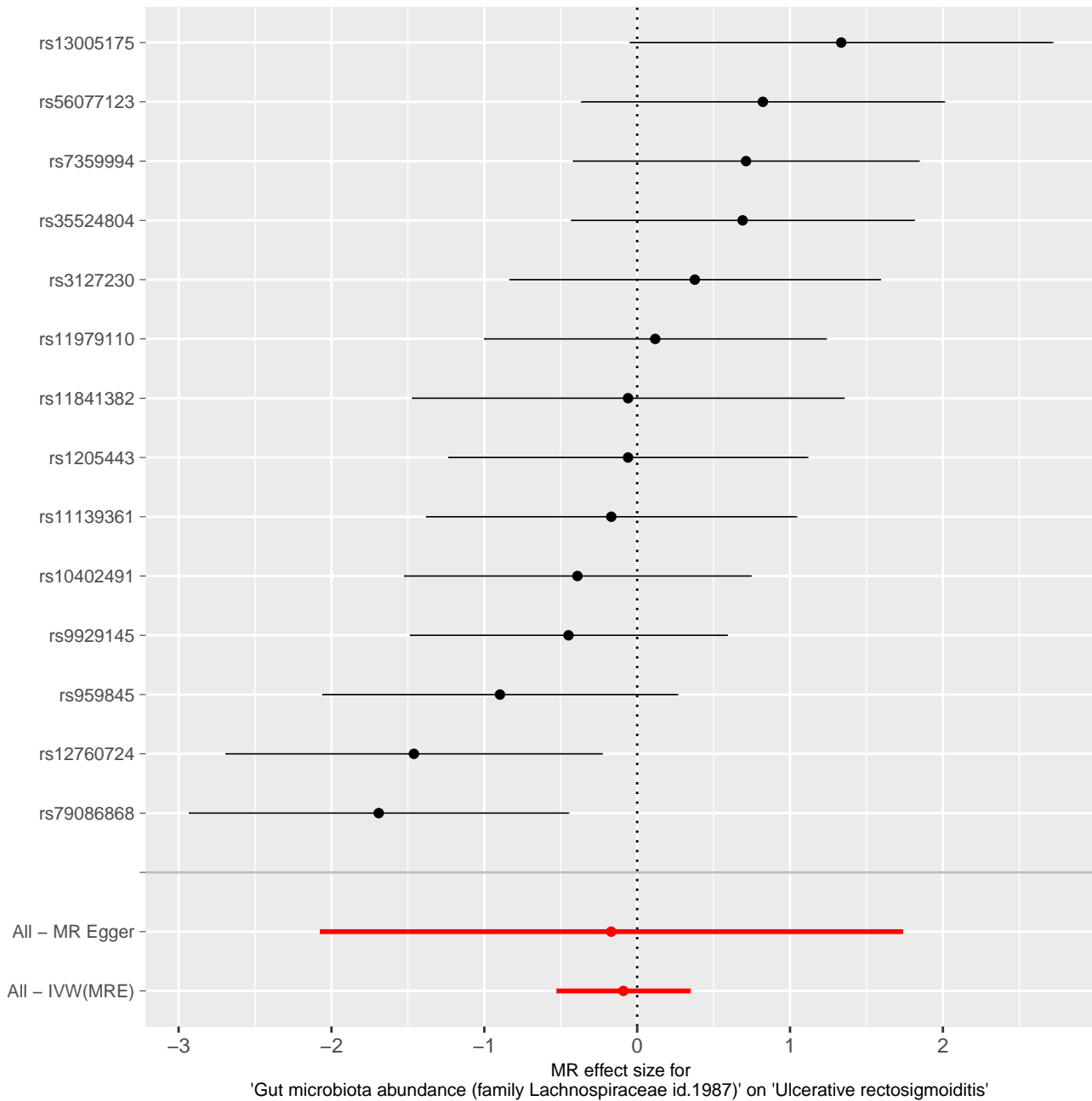


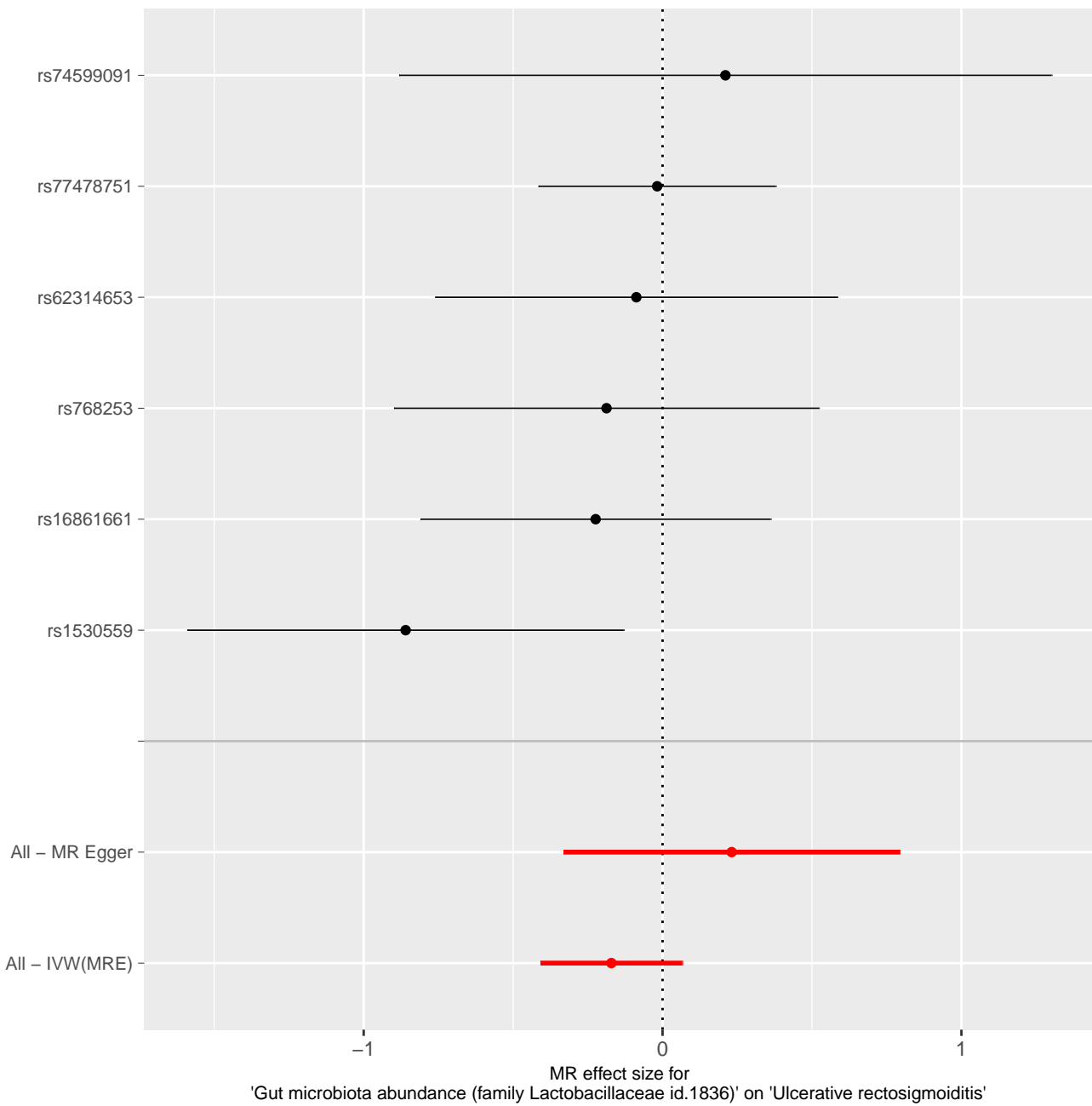


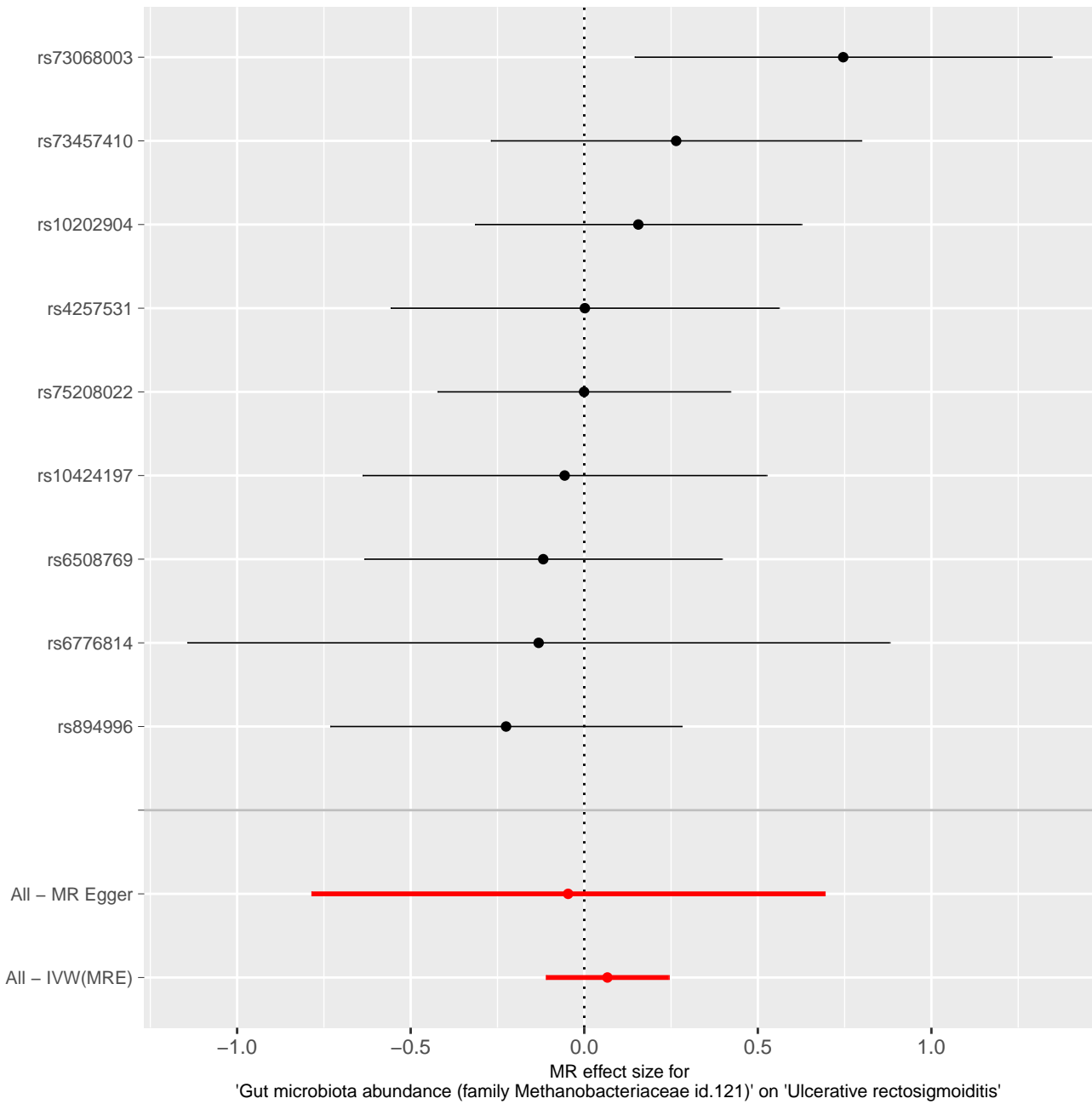


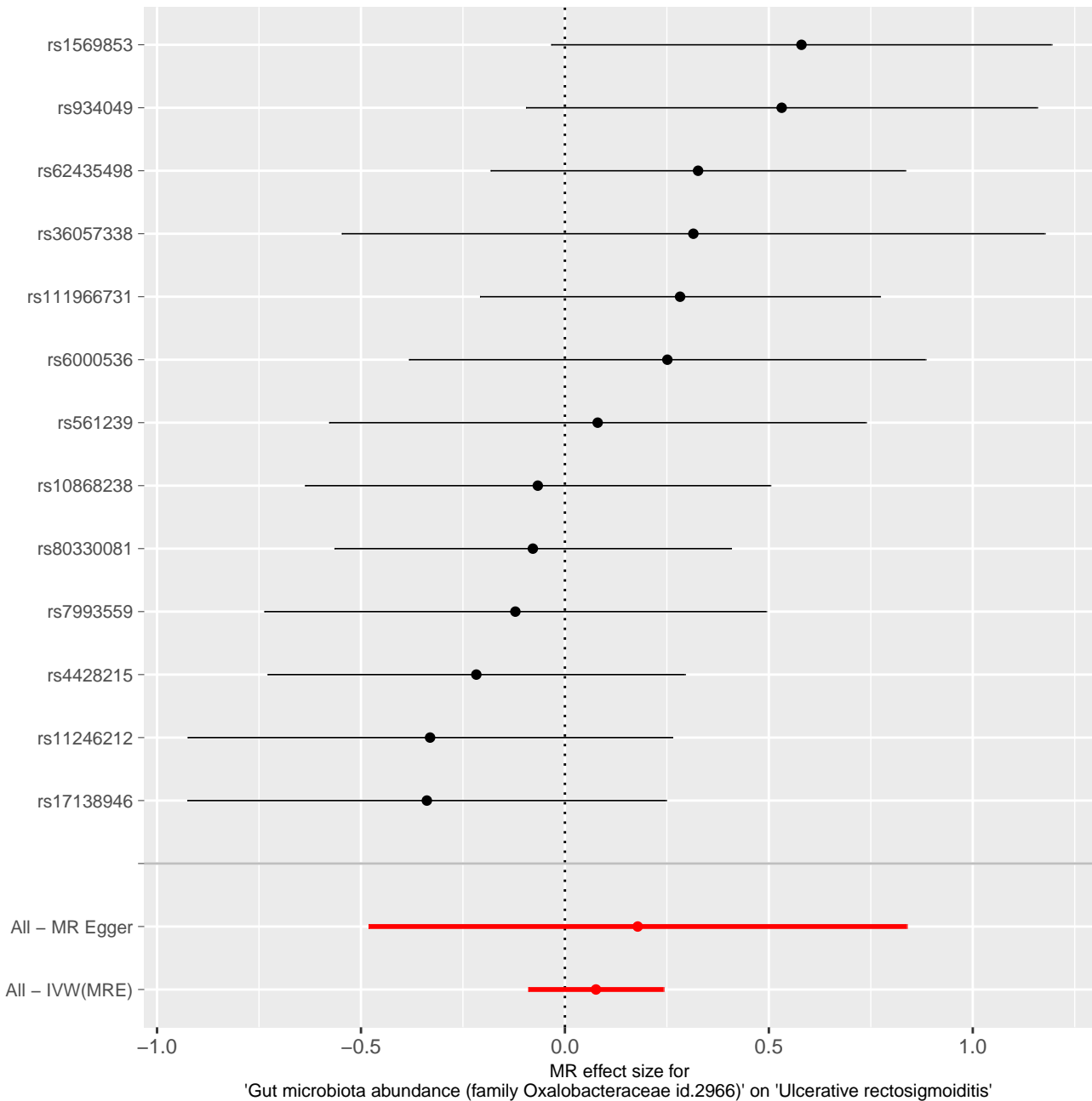


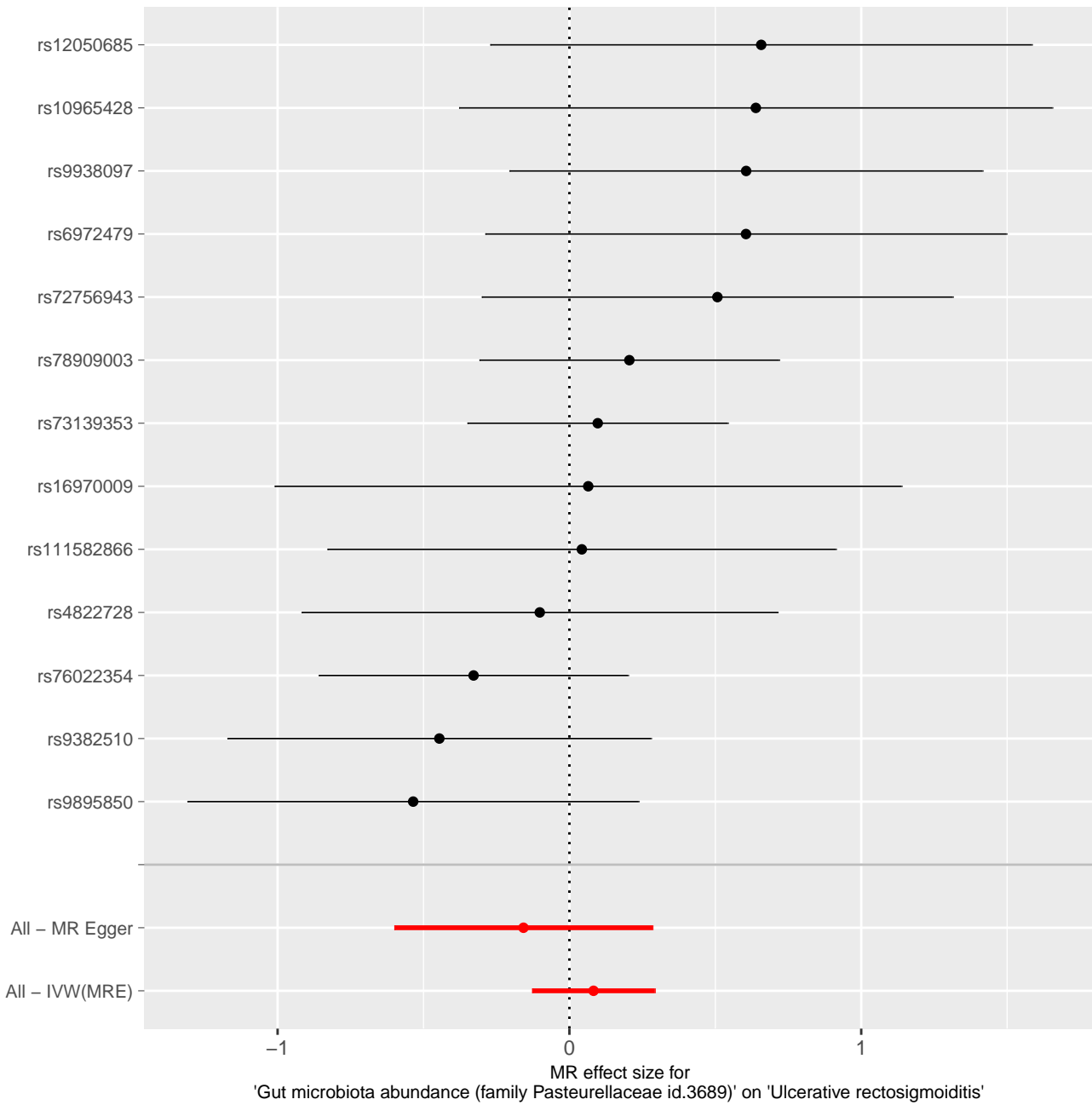


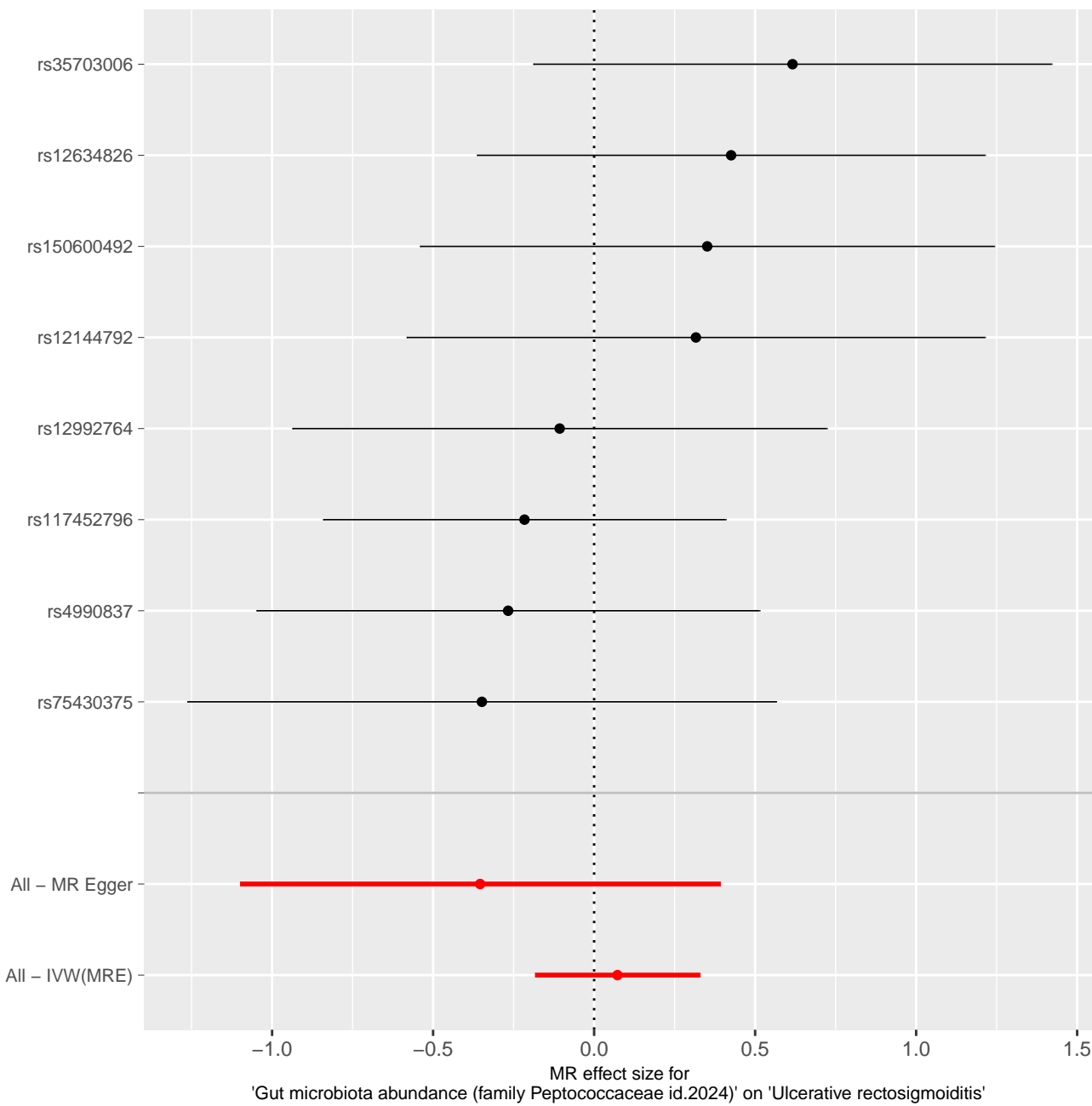


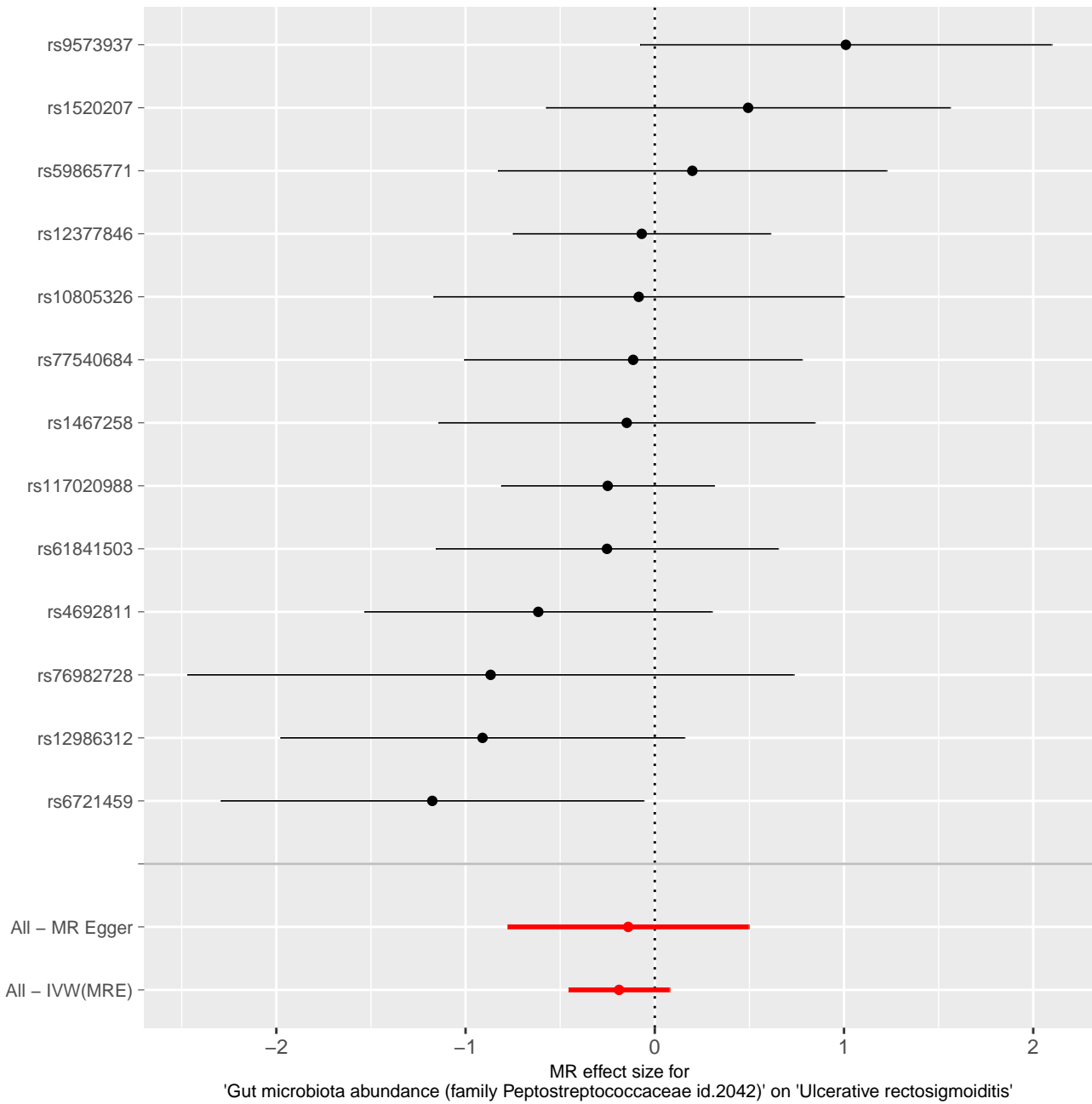


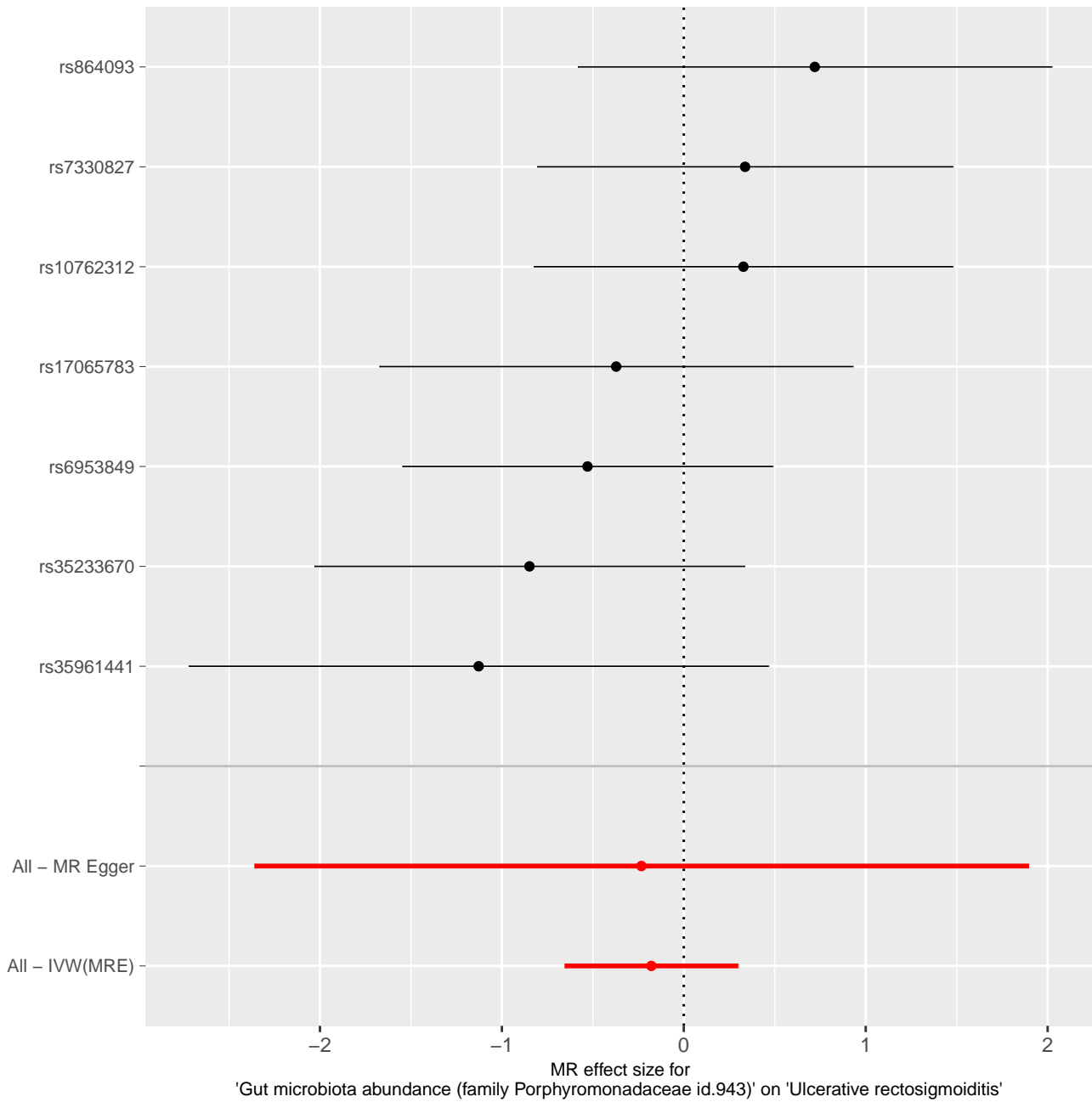


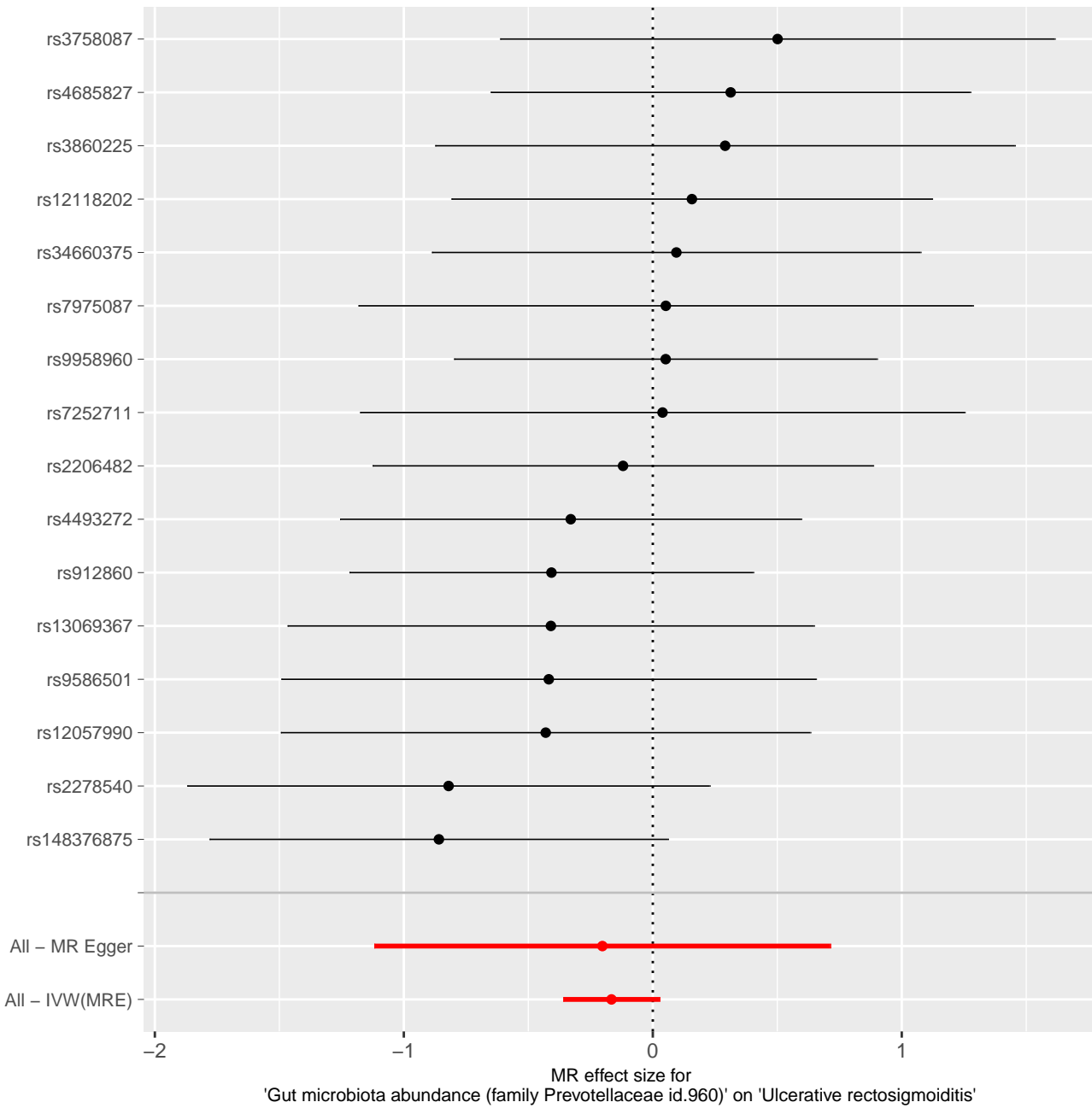


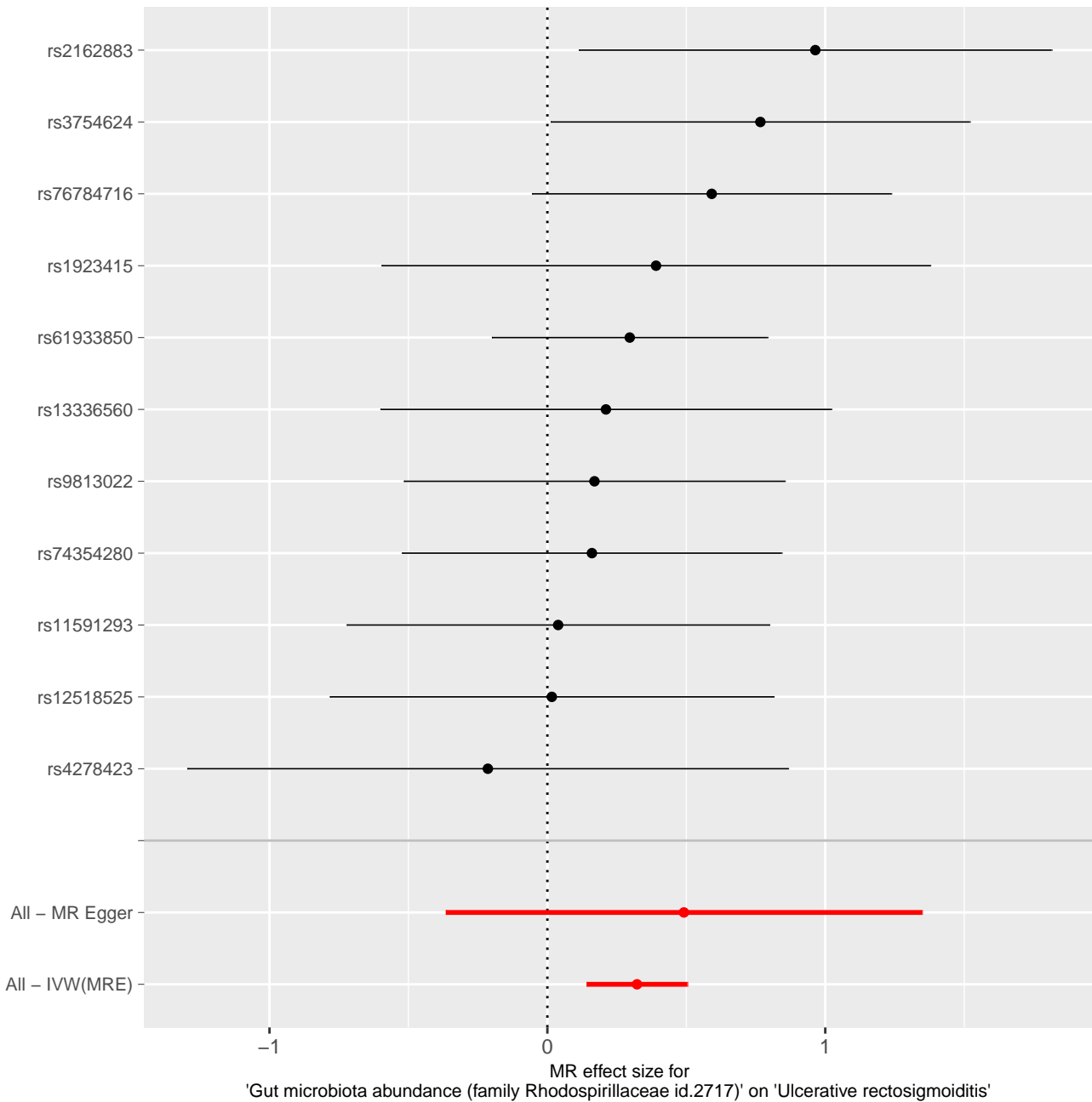


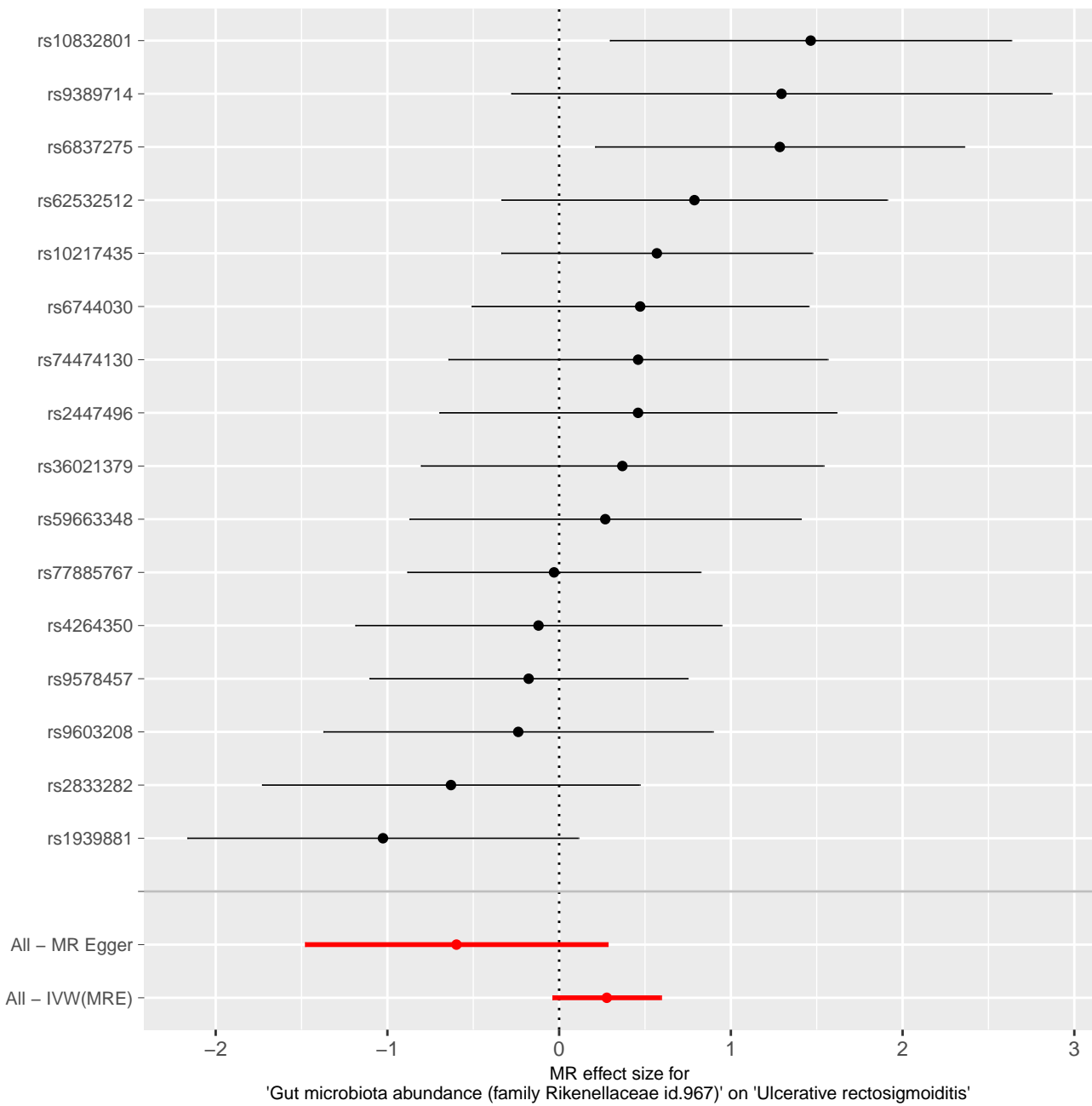


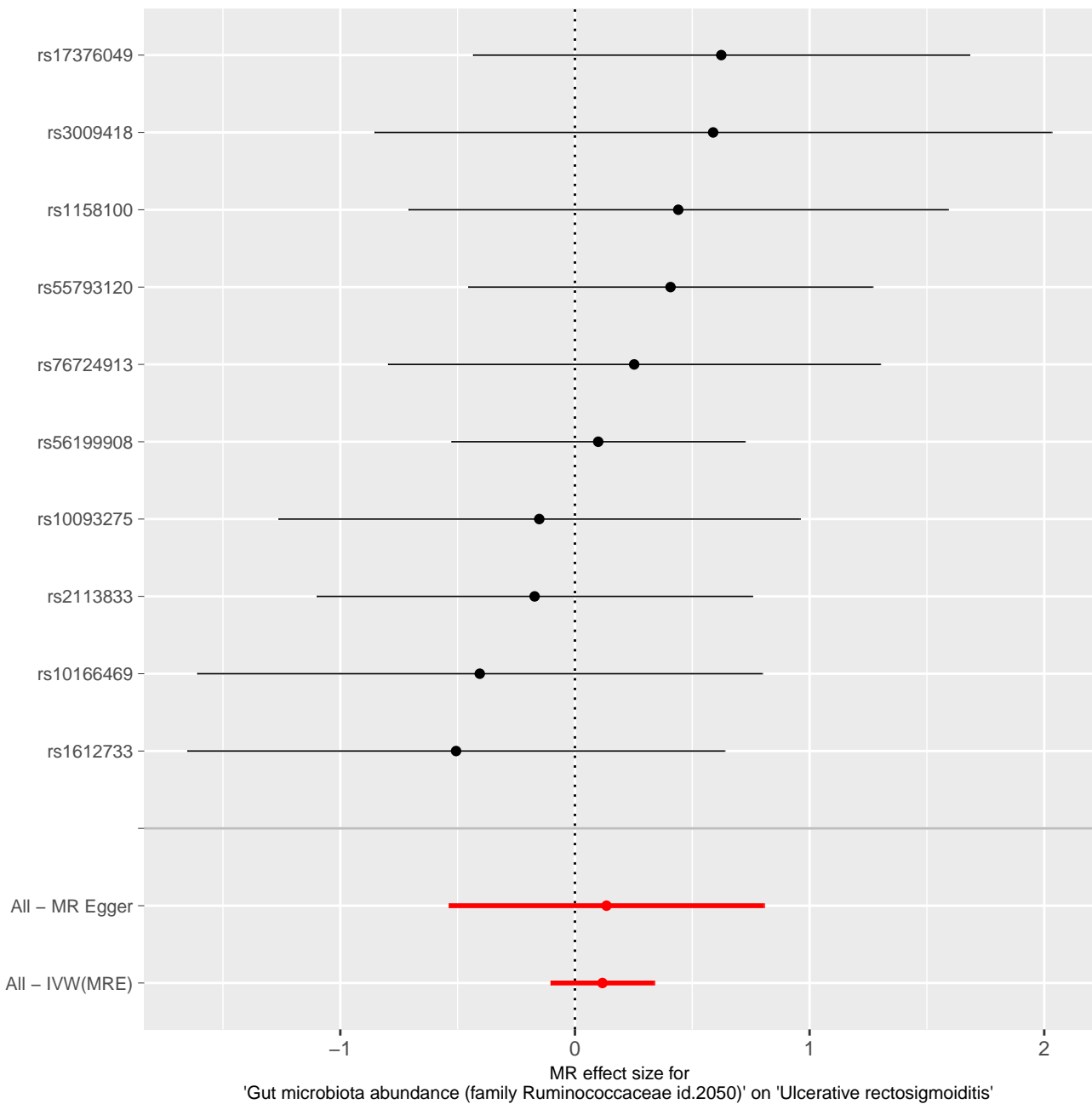


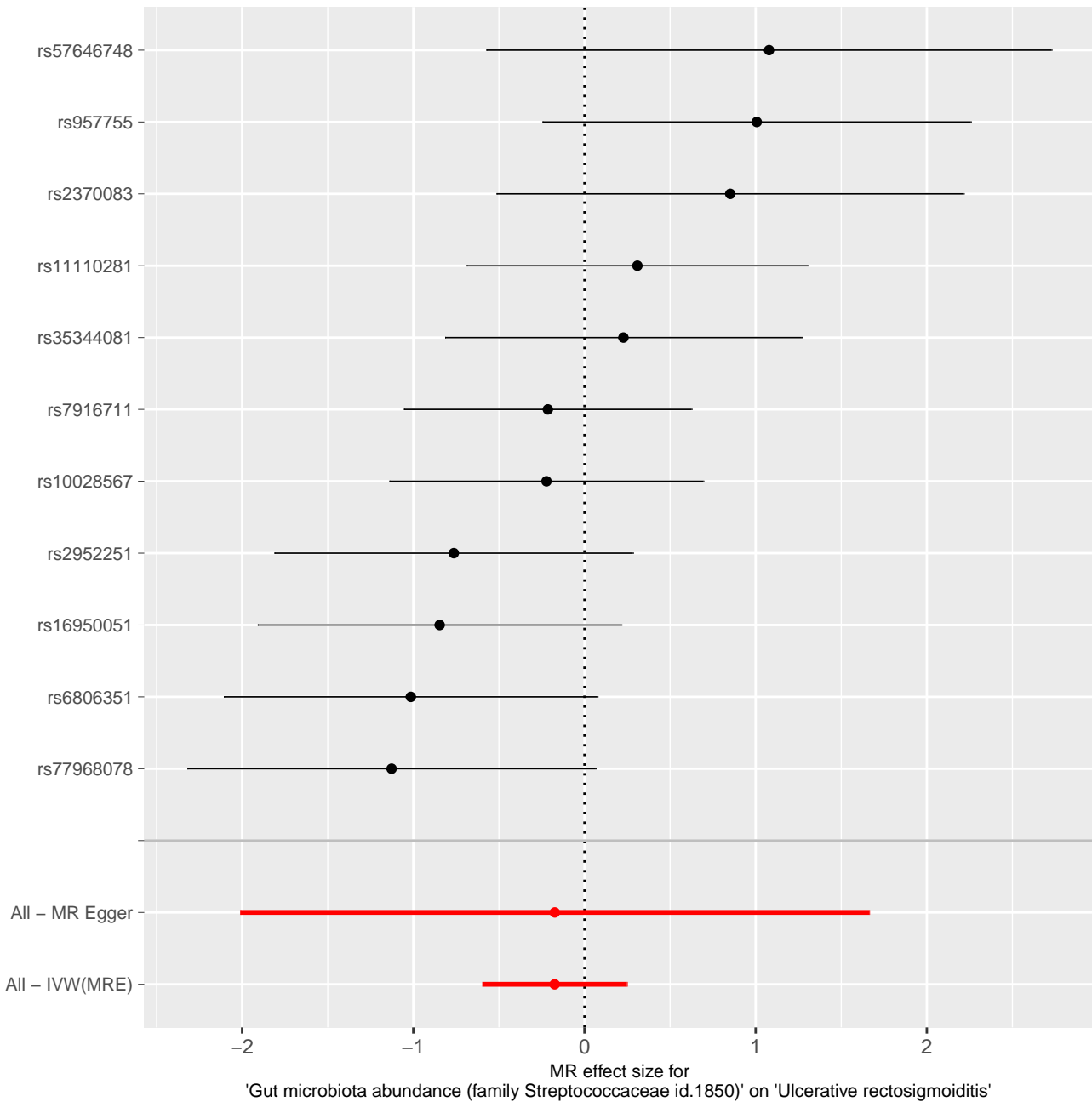


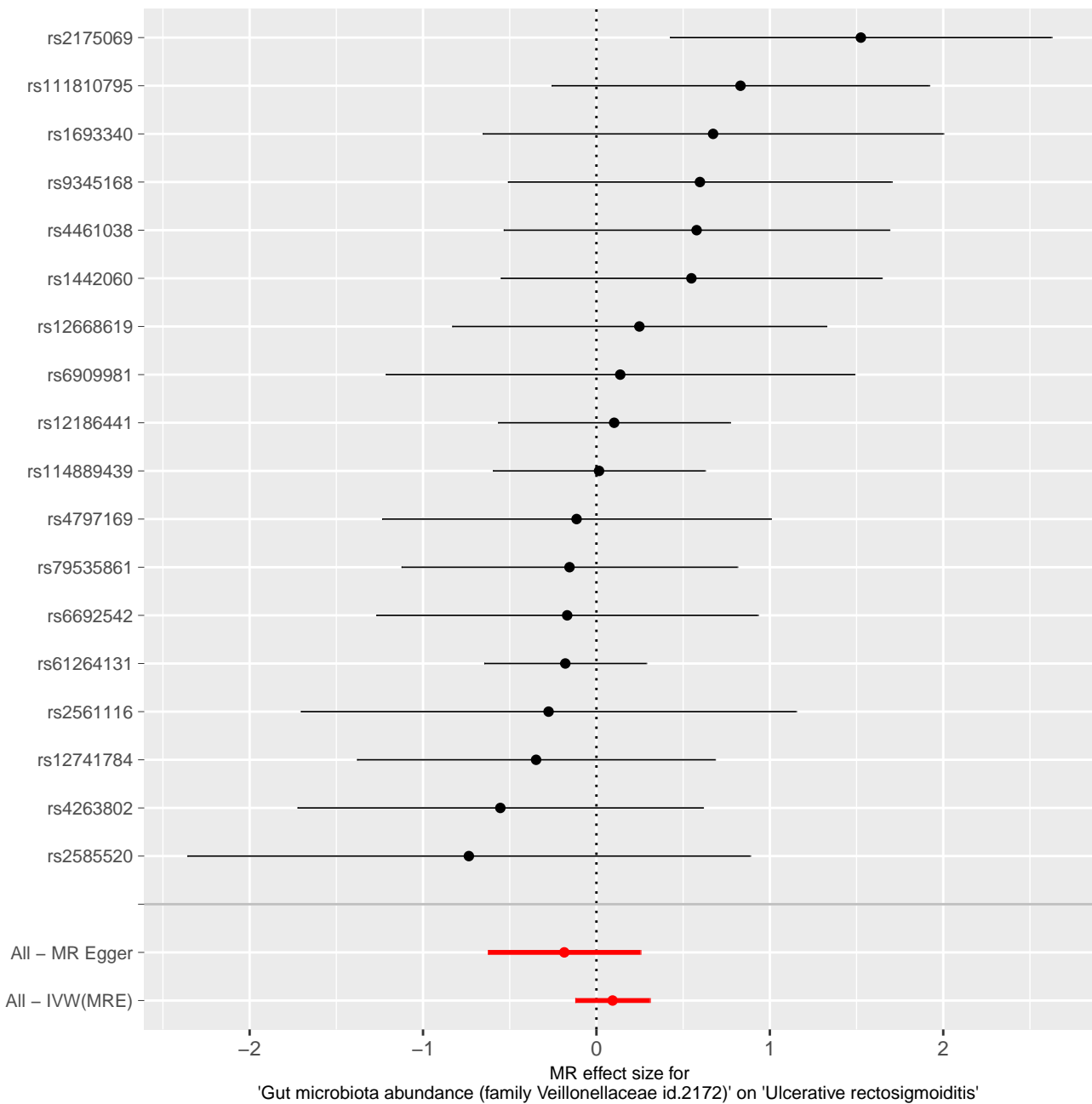


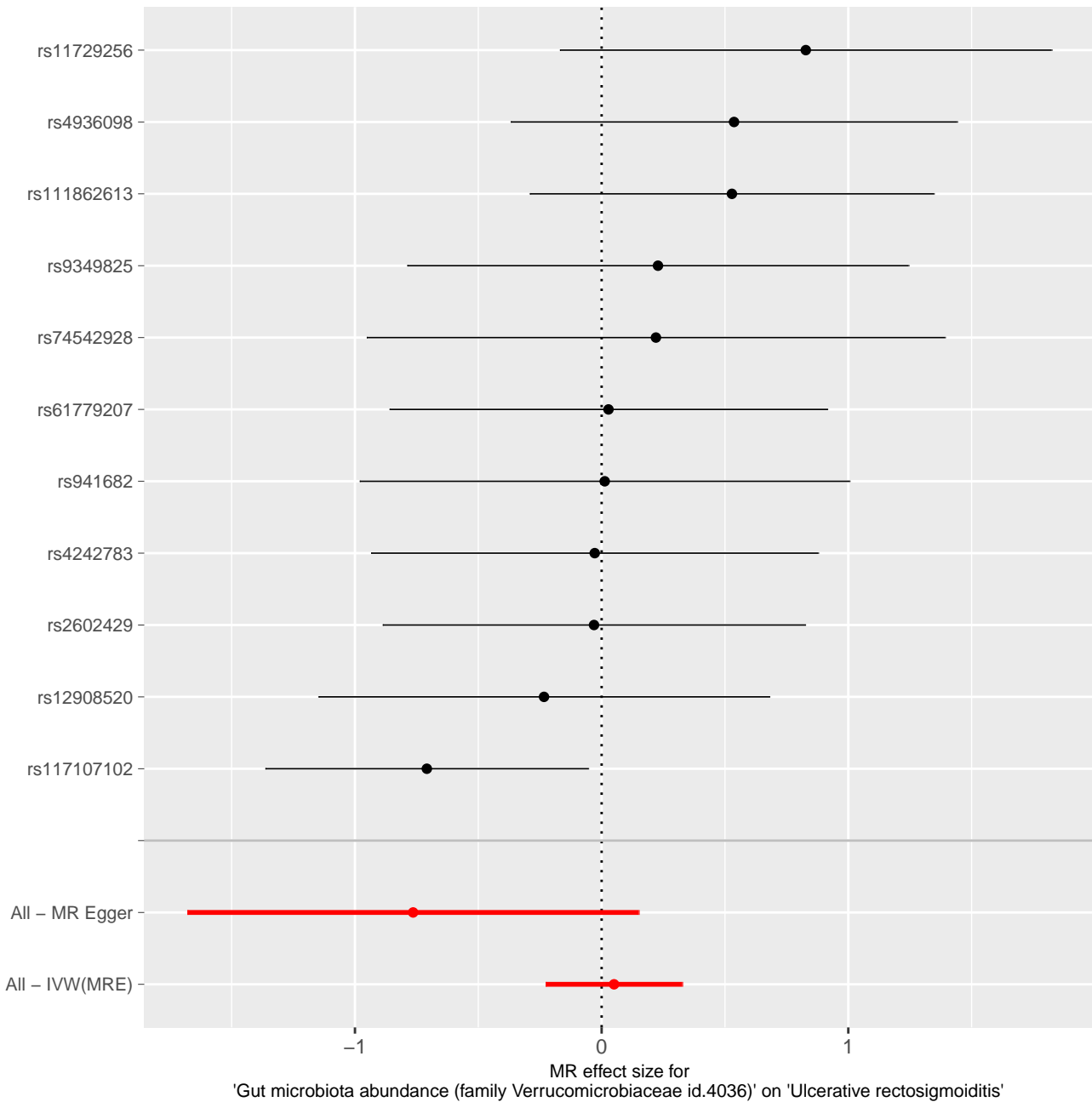


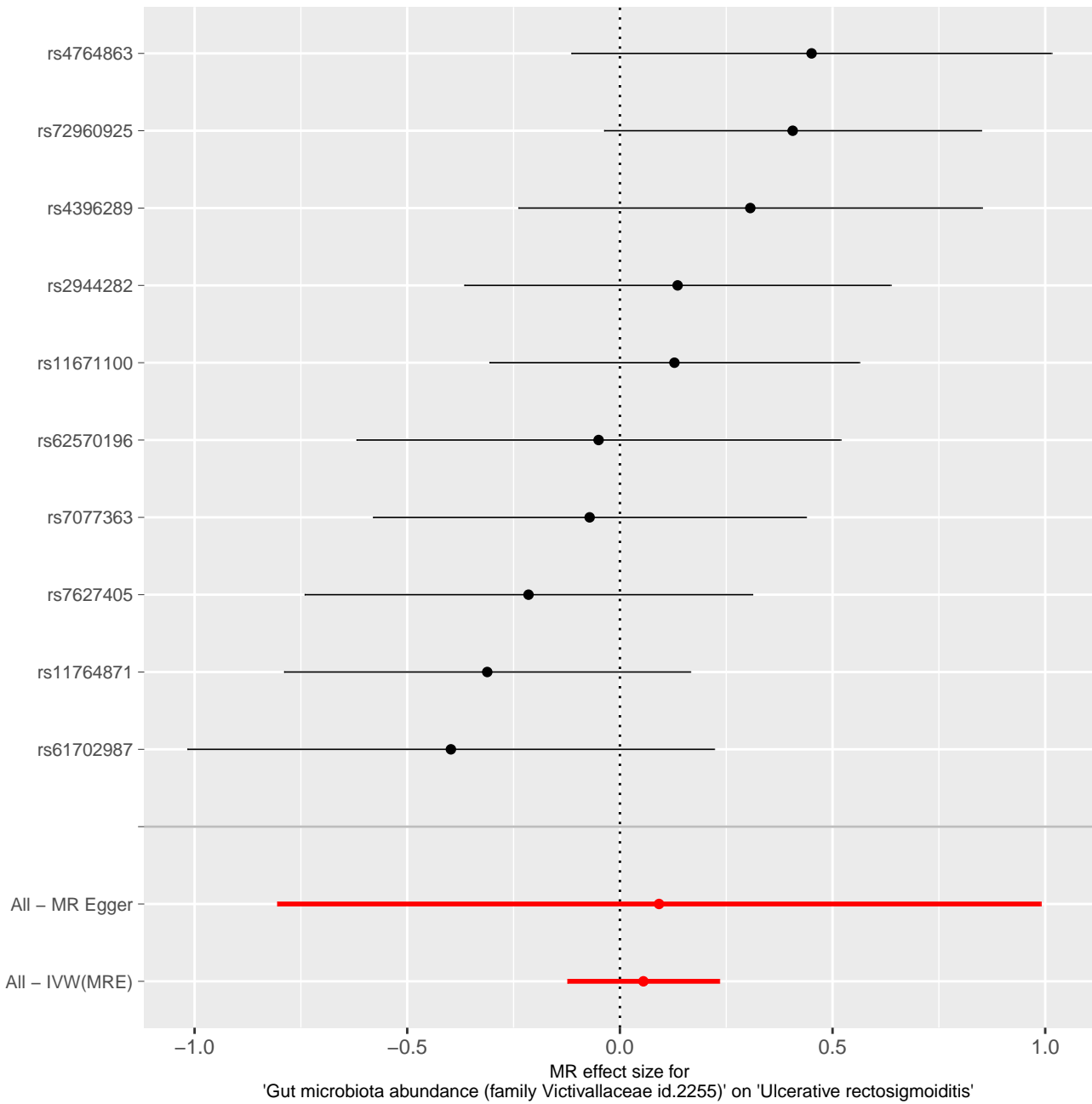


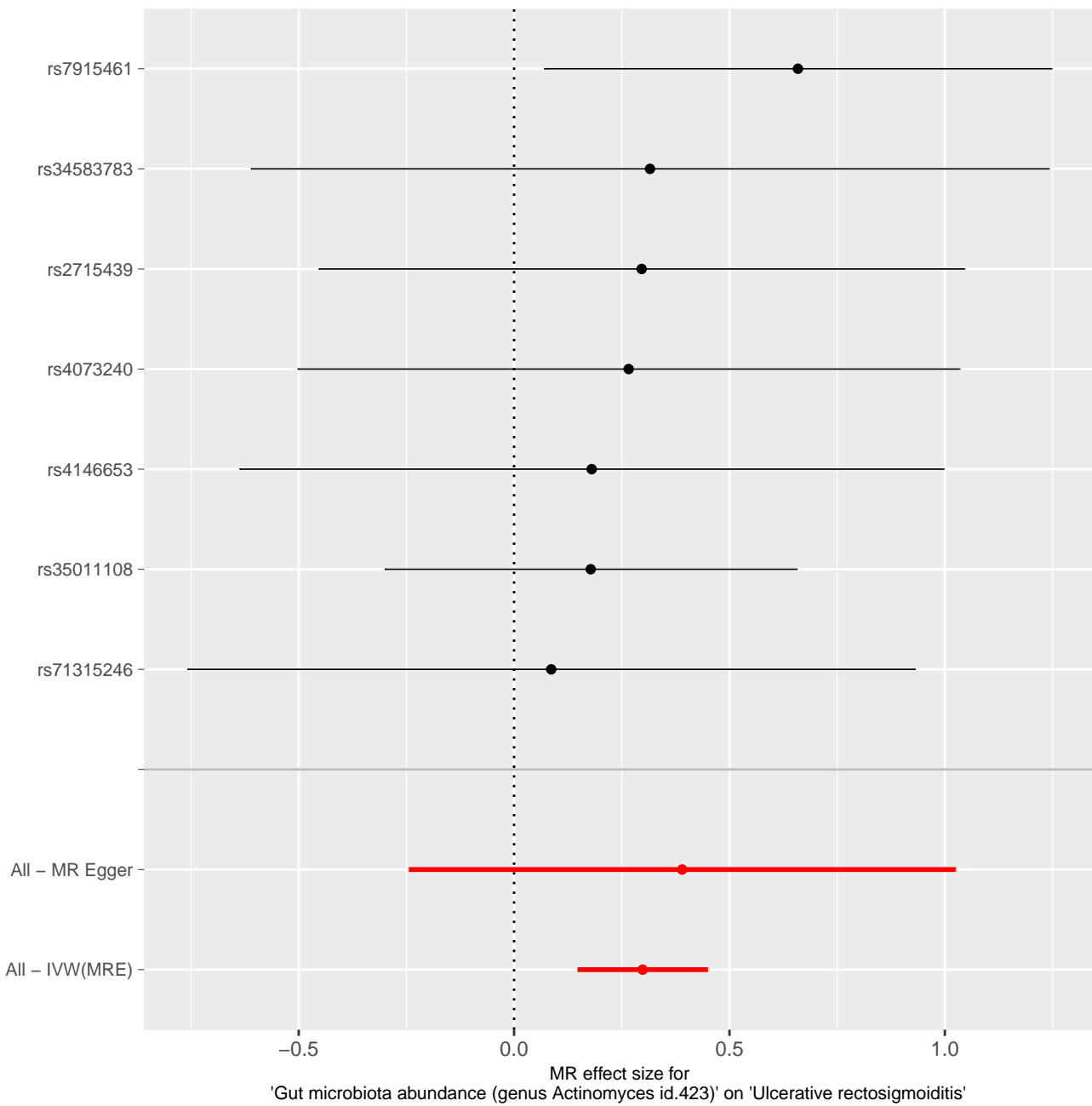


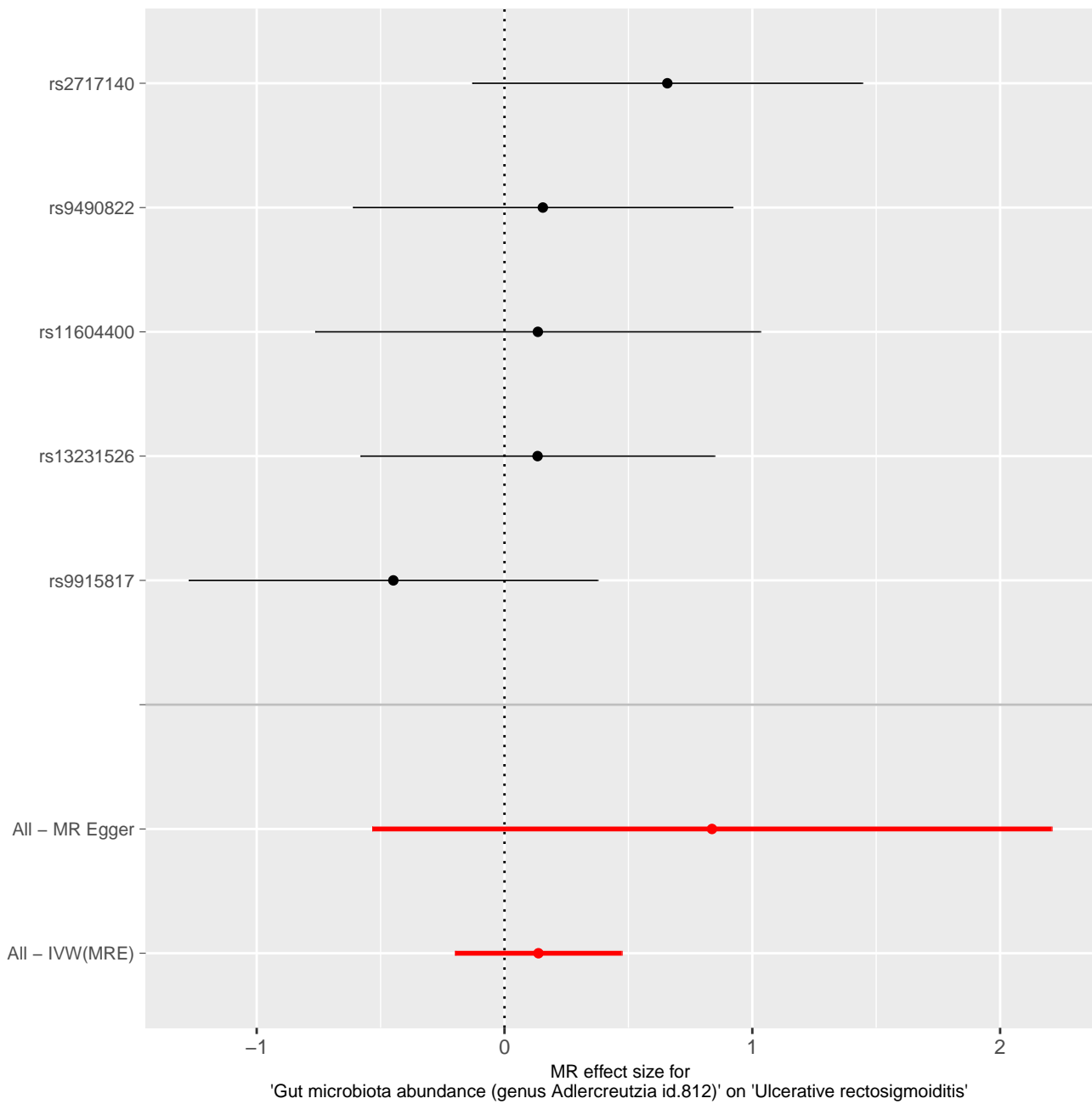




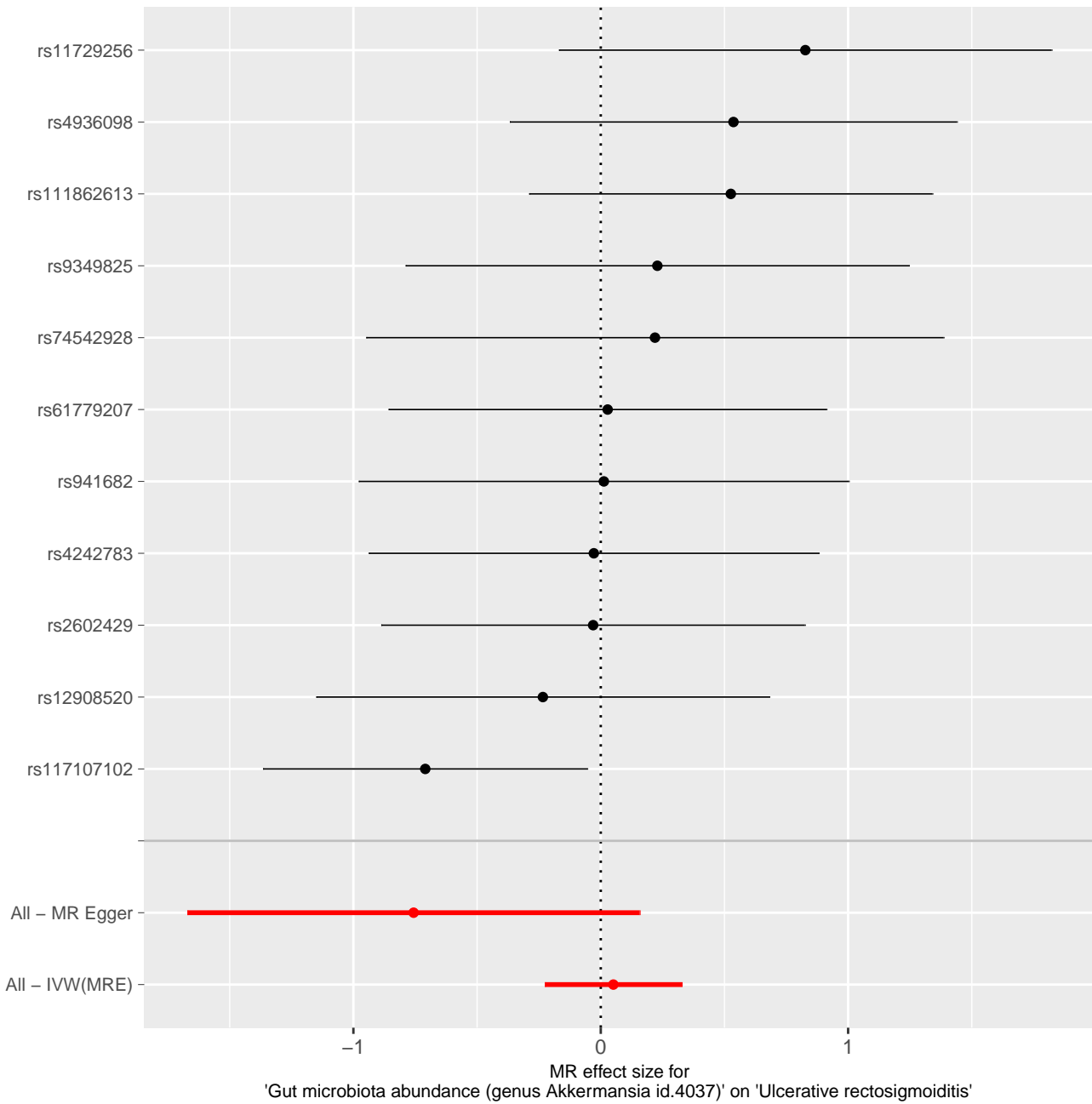


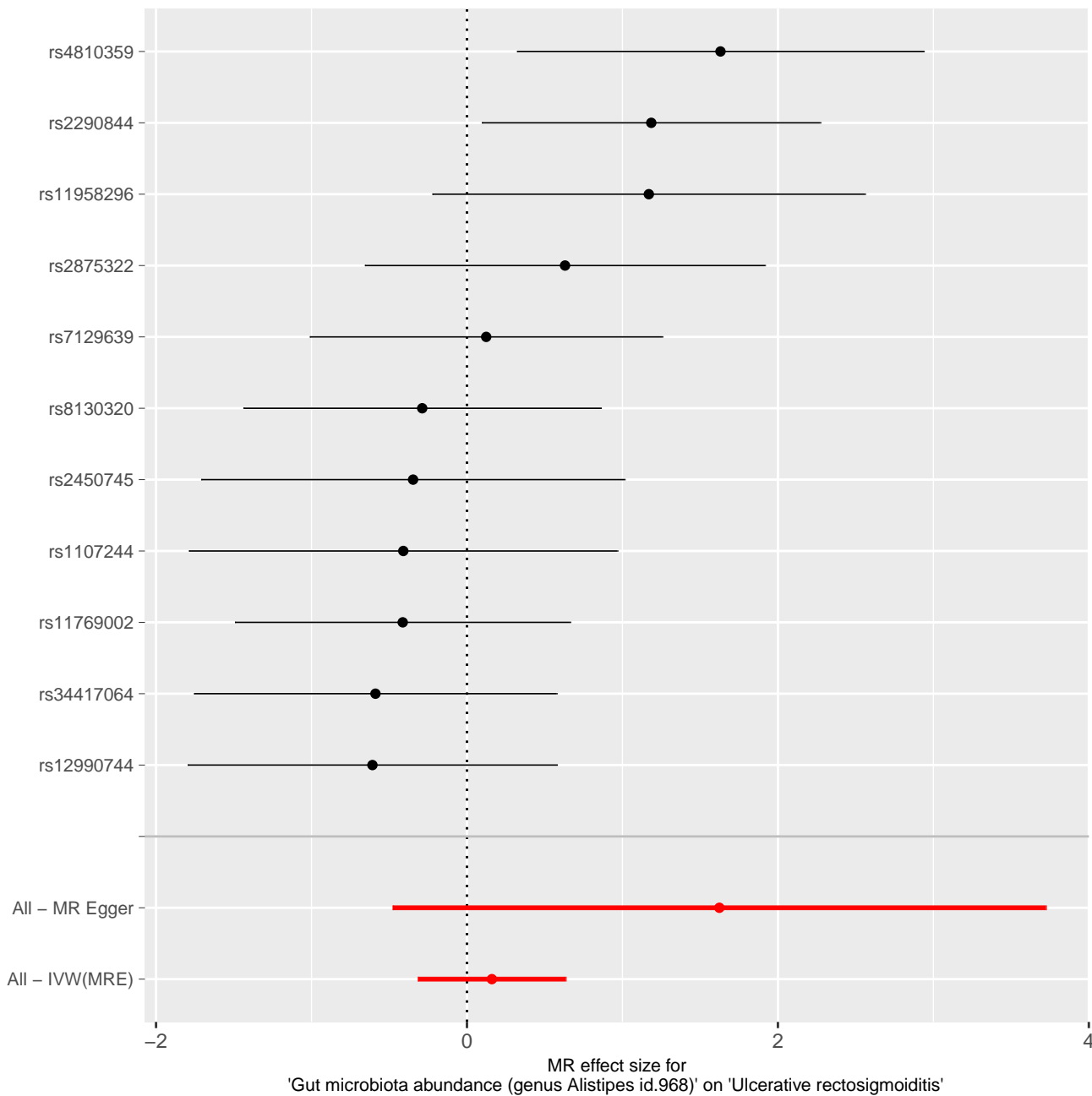


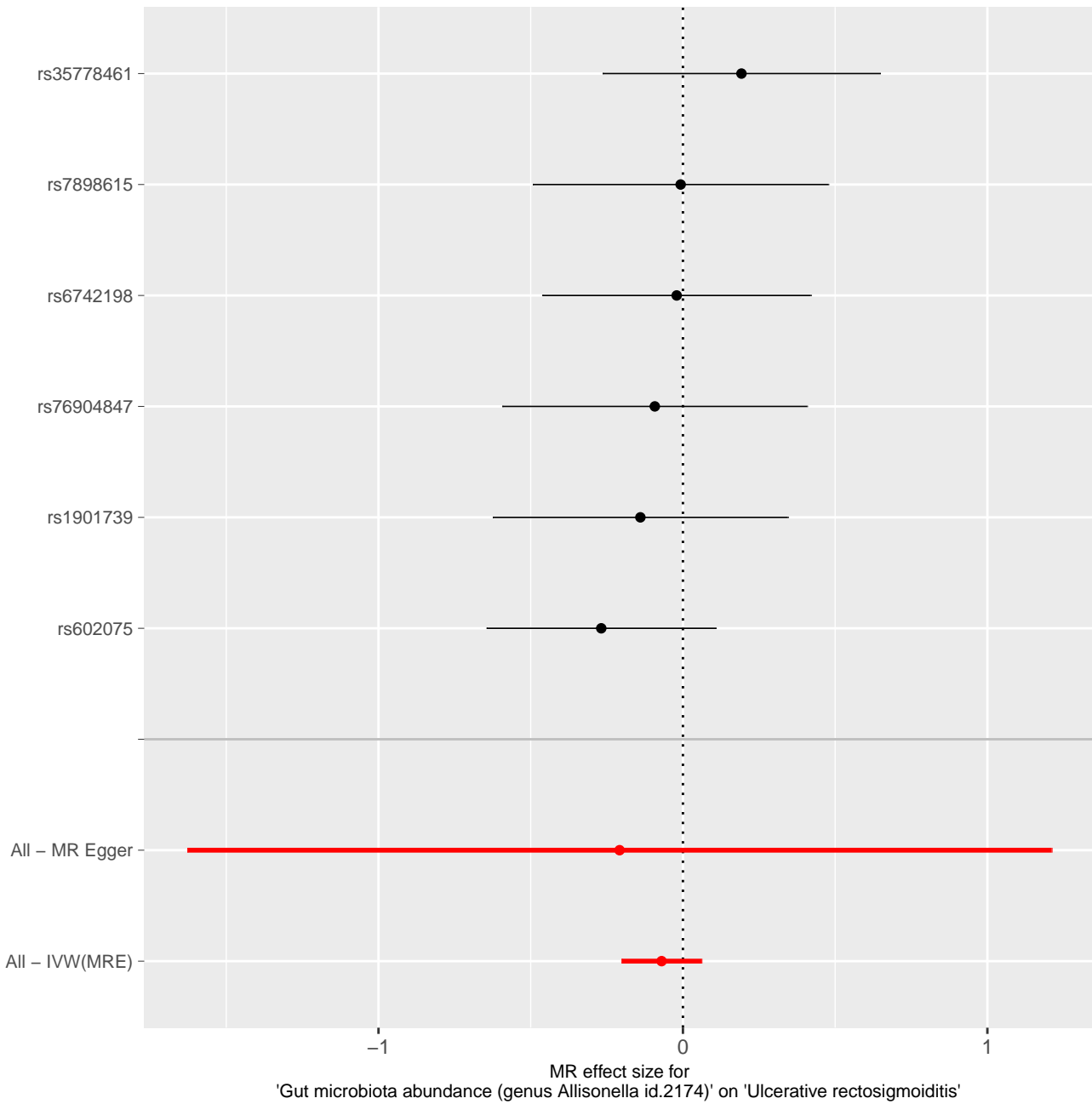


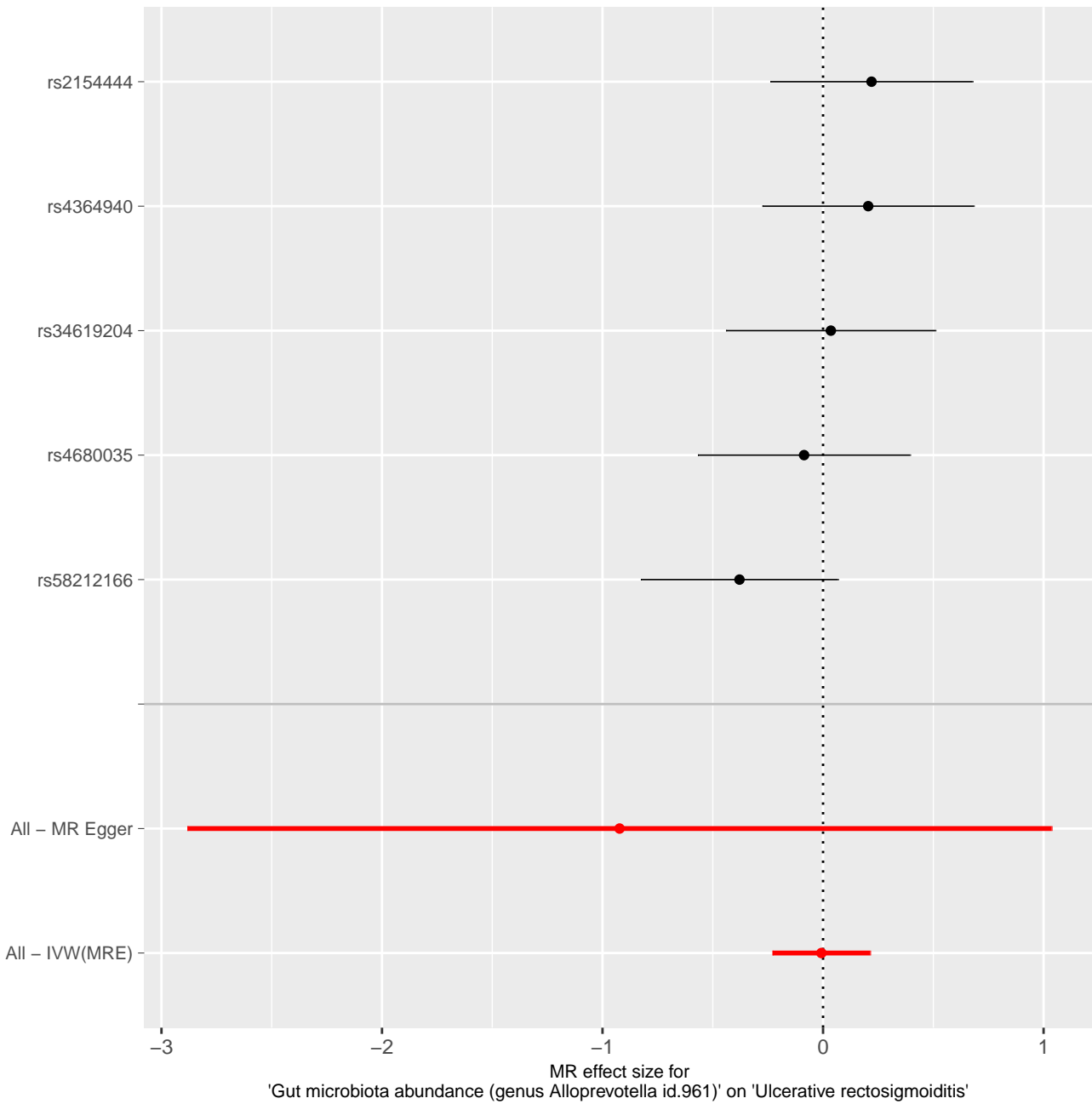


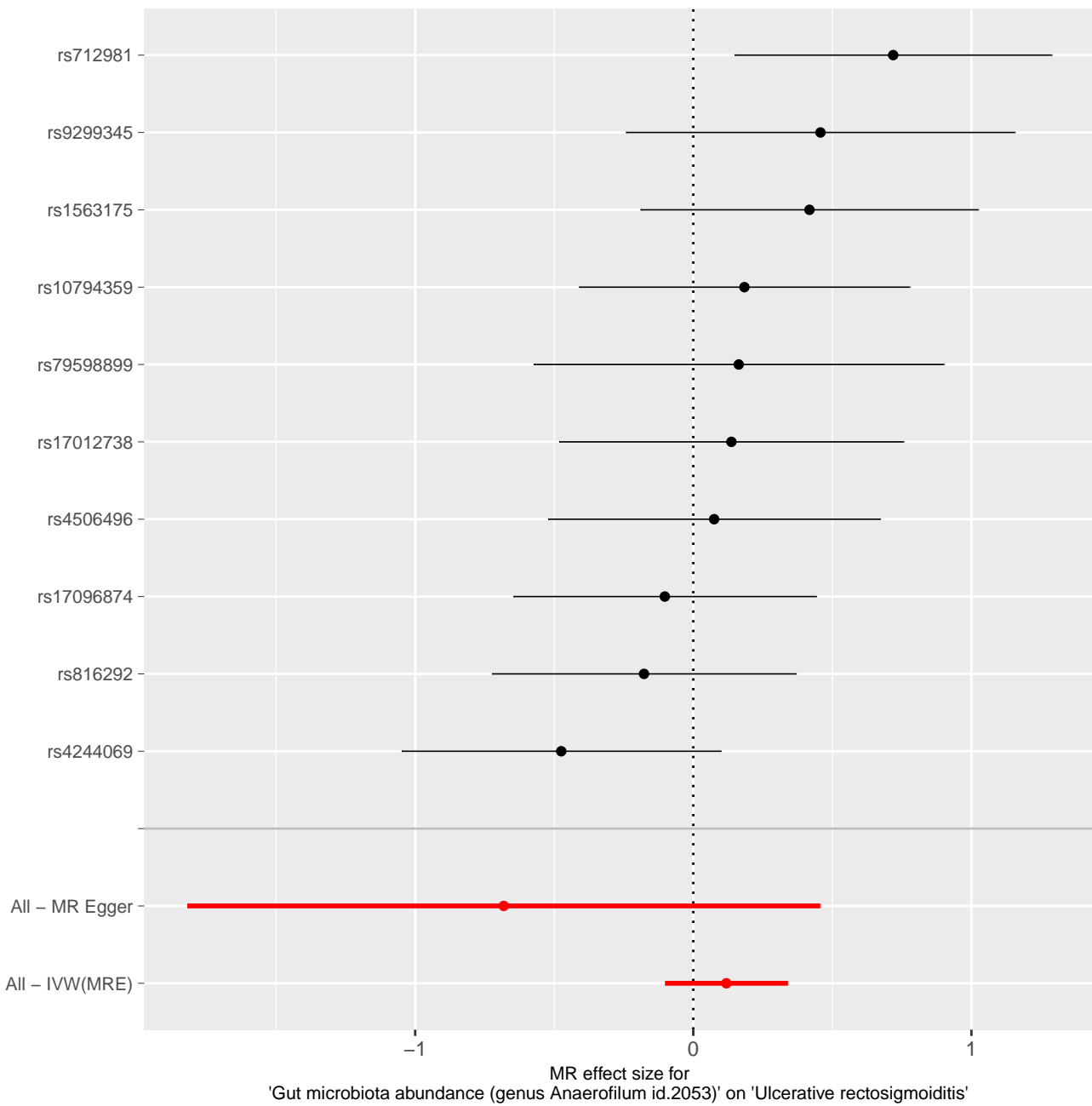
Batch 1535 : Gut microbiota abundance (genus Akkermansia id.4037) on Ulcerative rectosigmoiditis

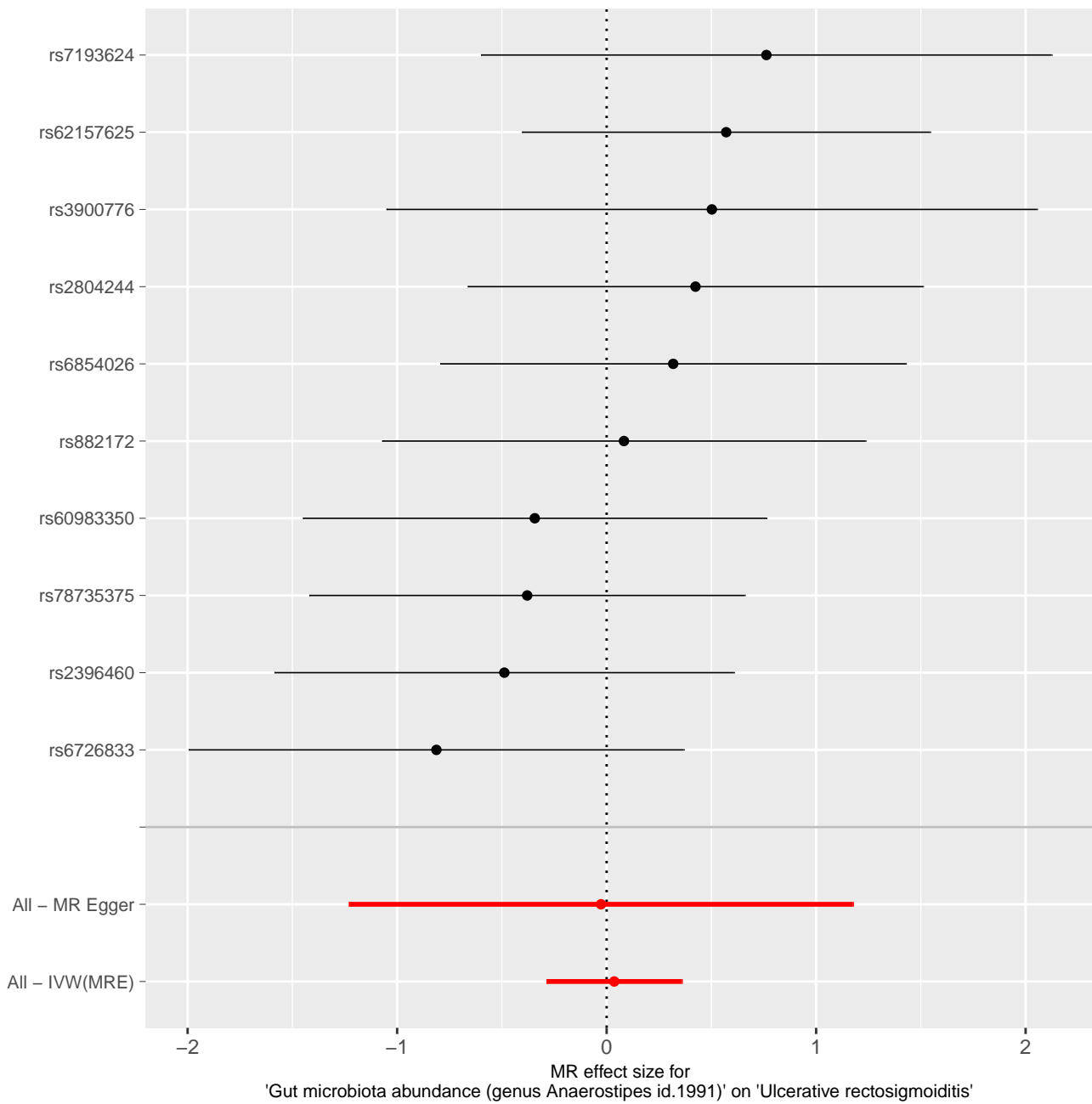


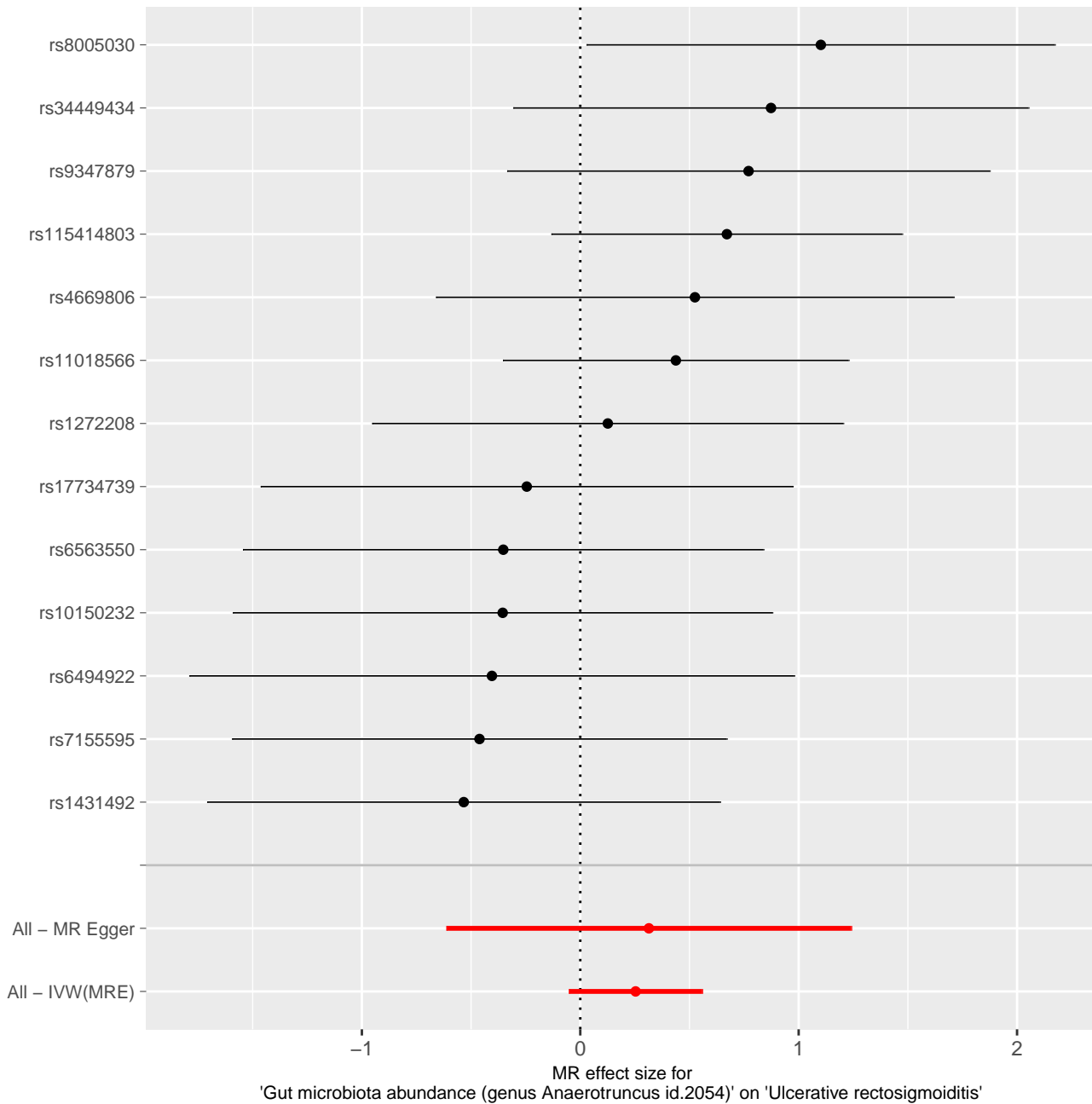


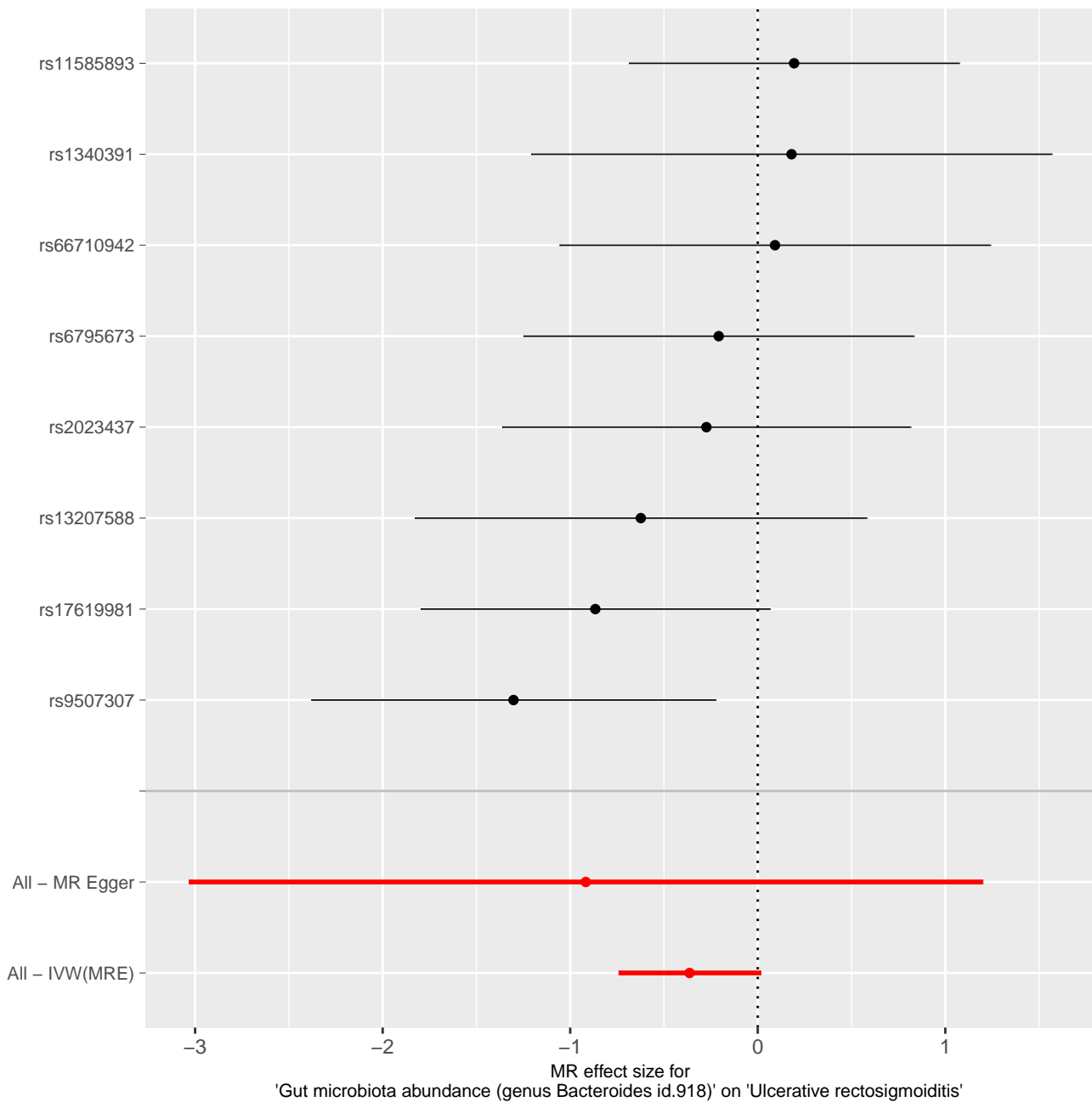


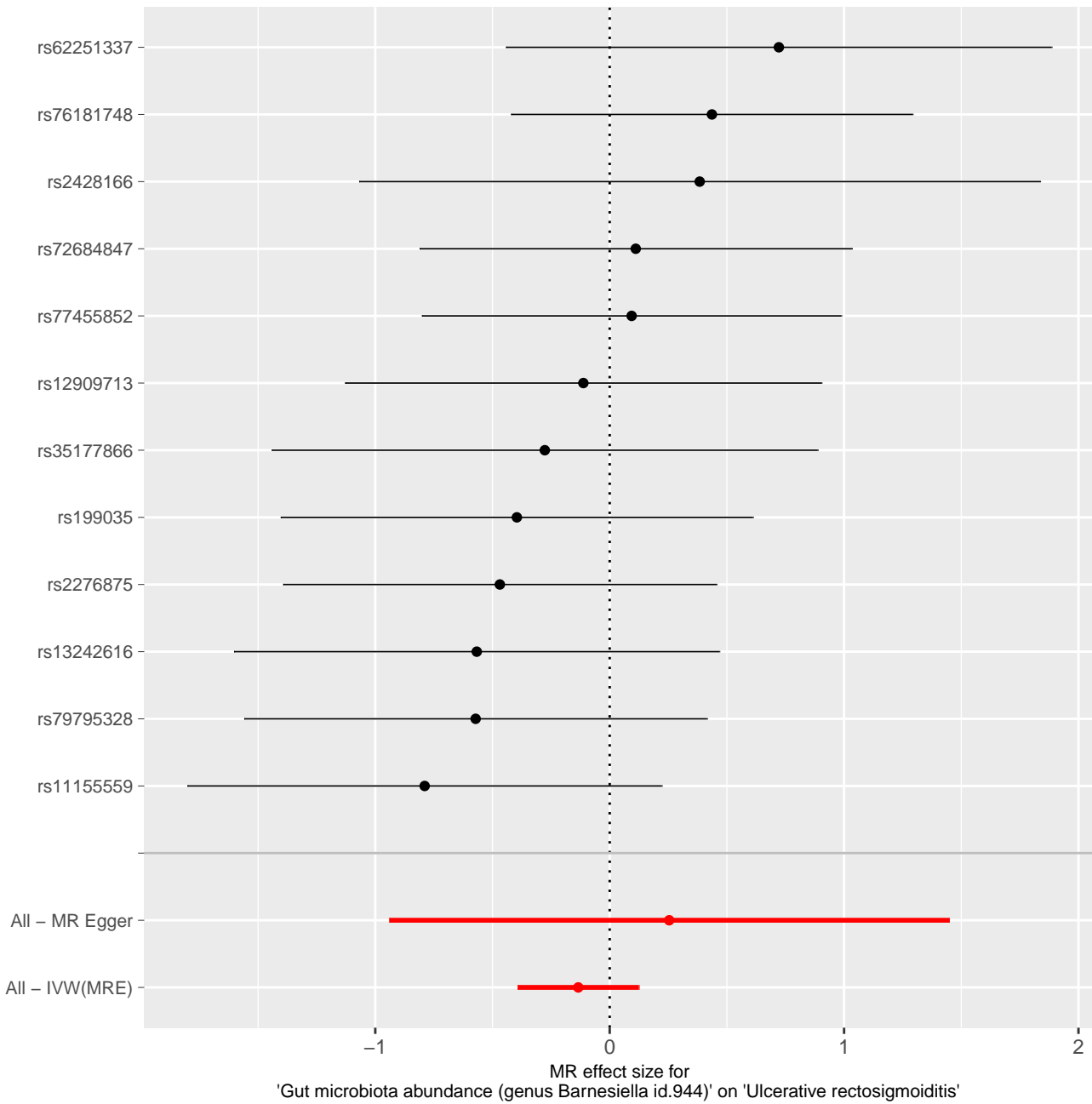


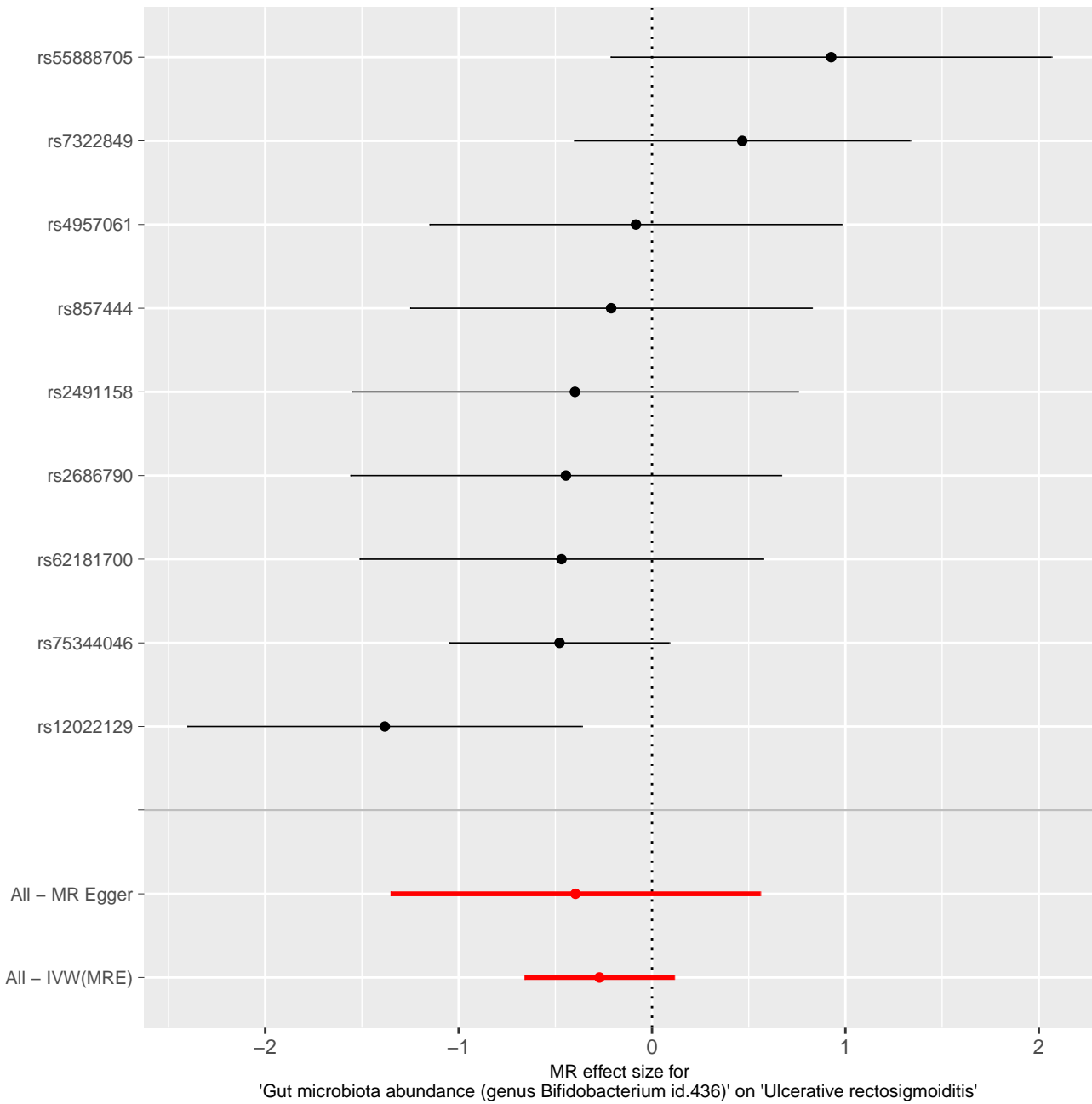


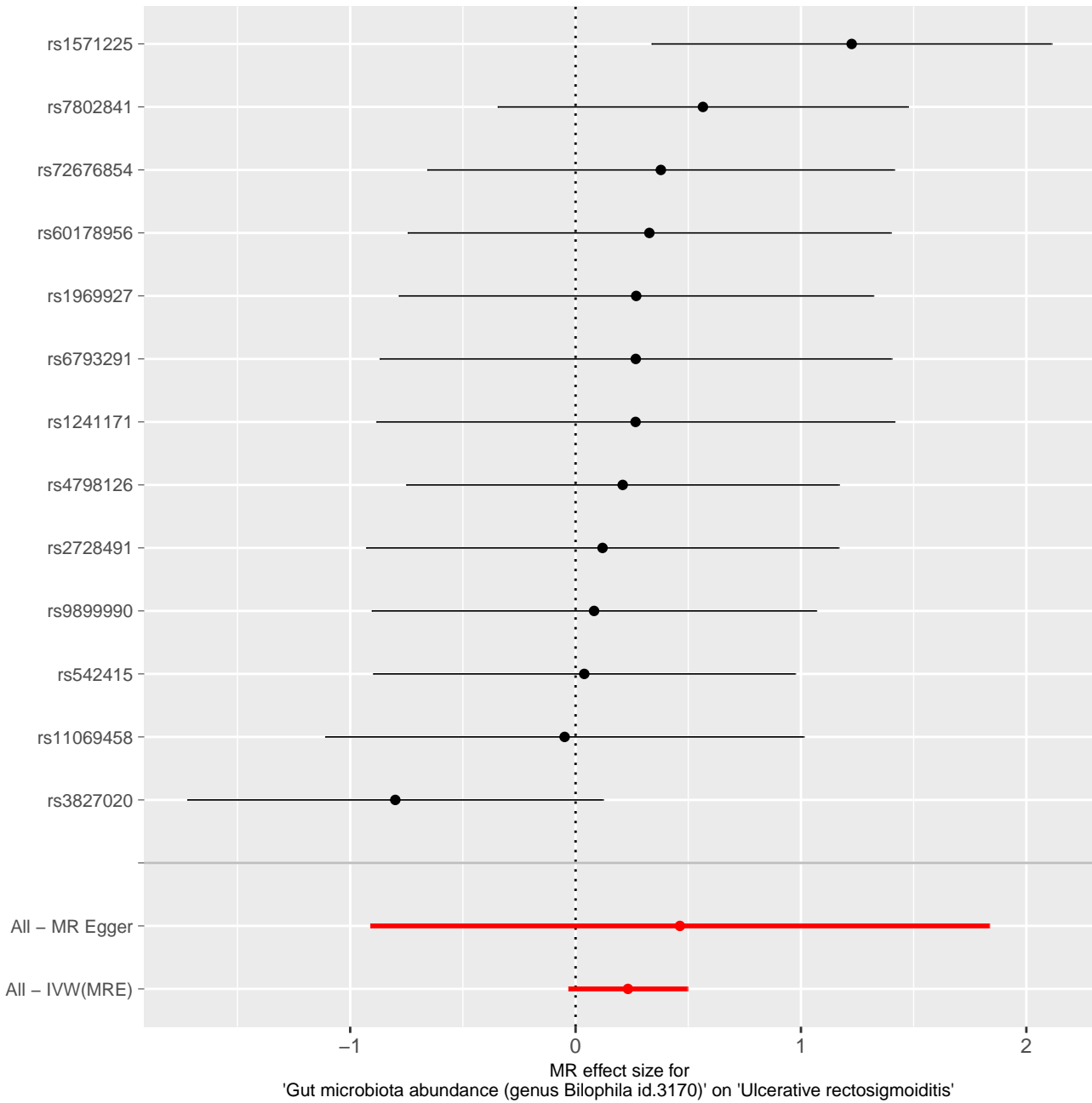


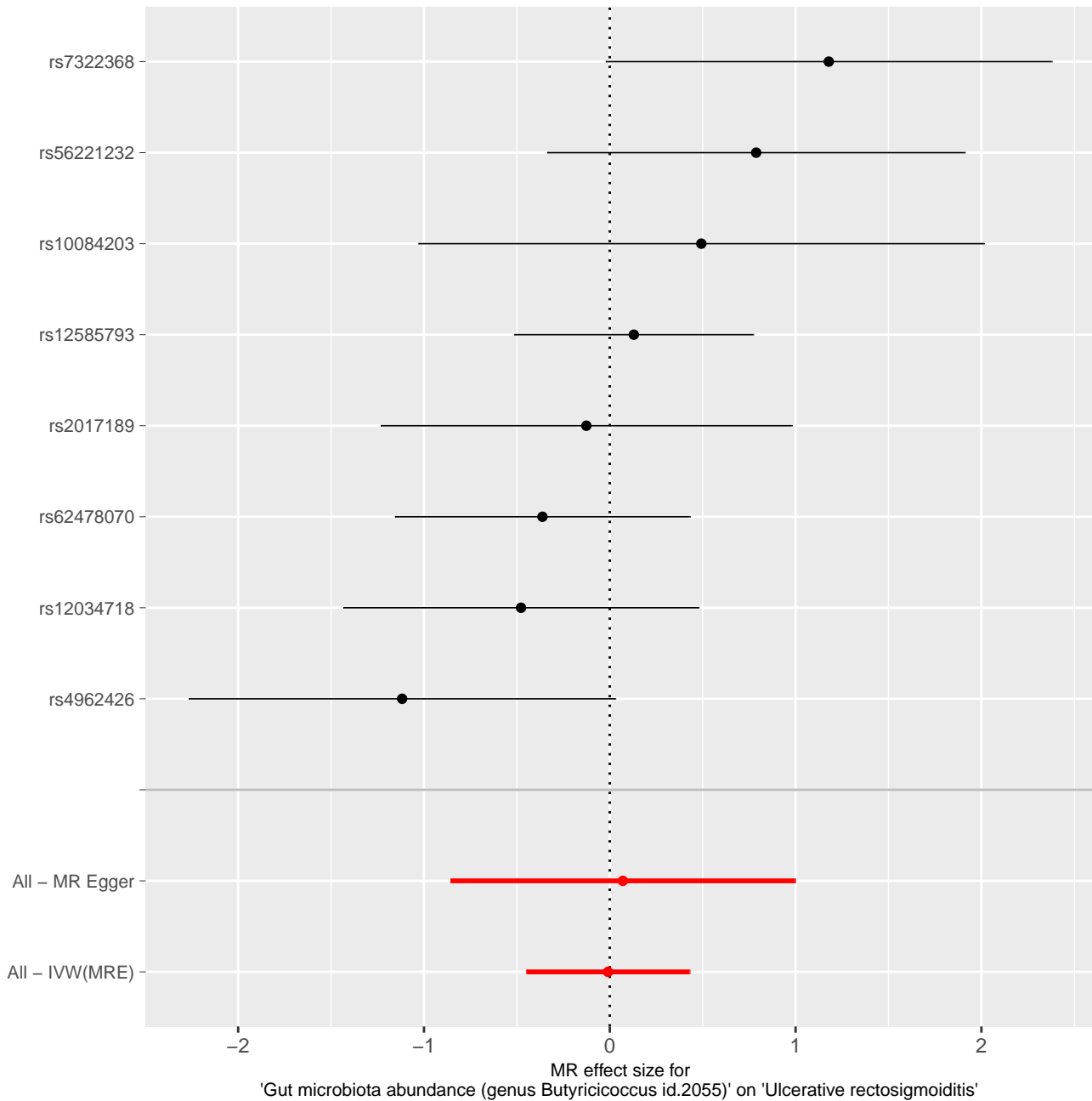


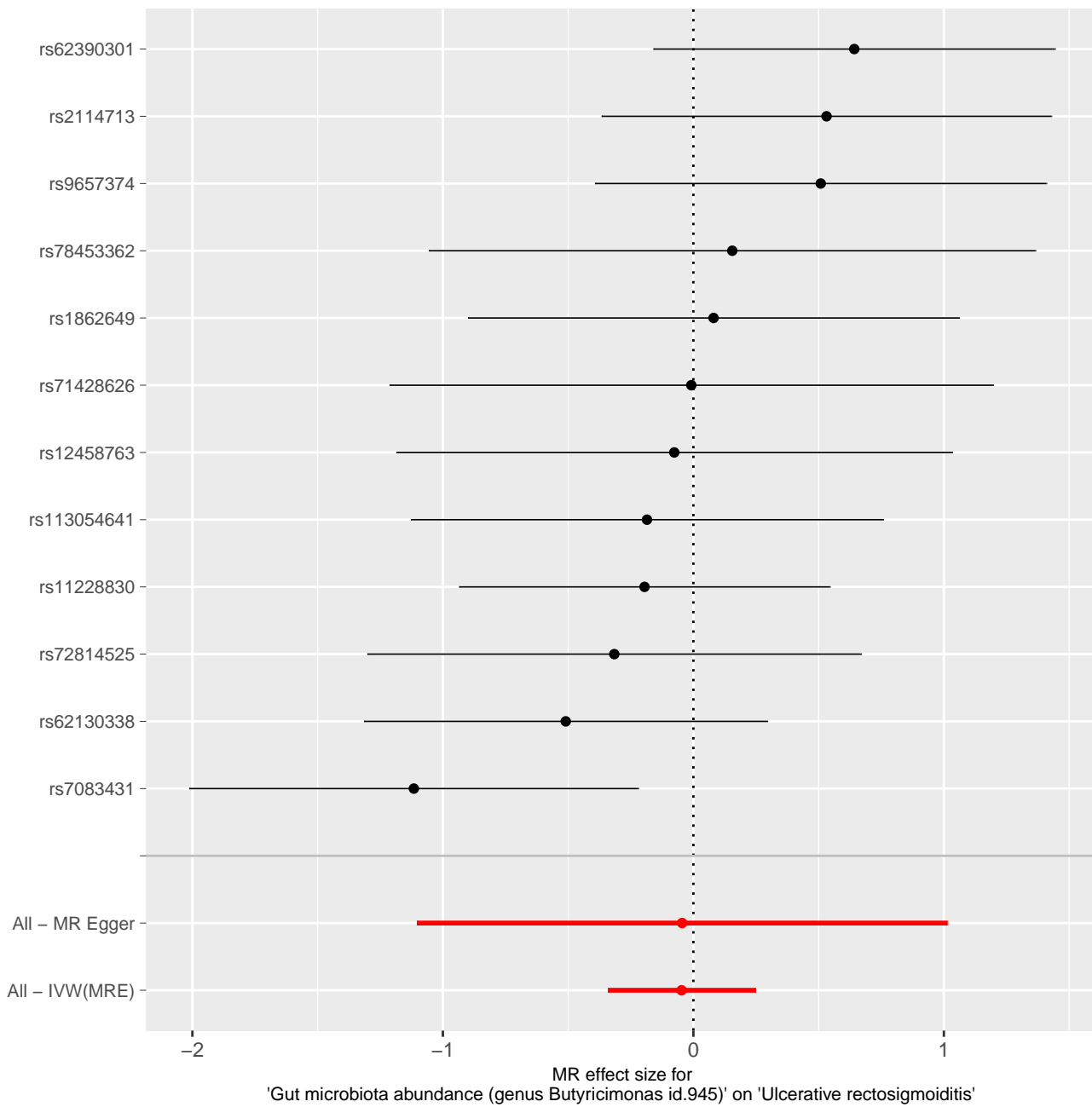


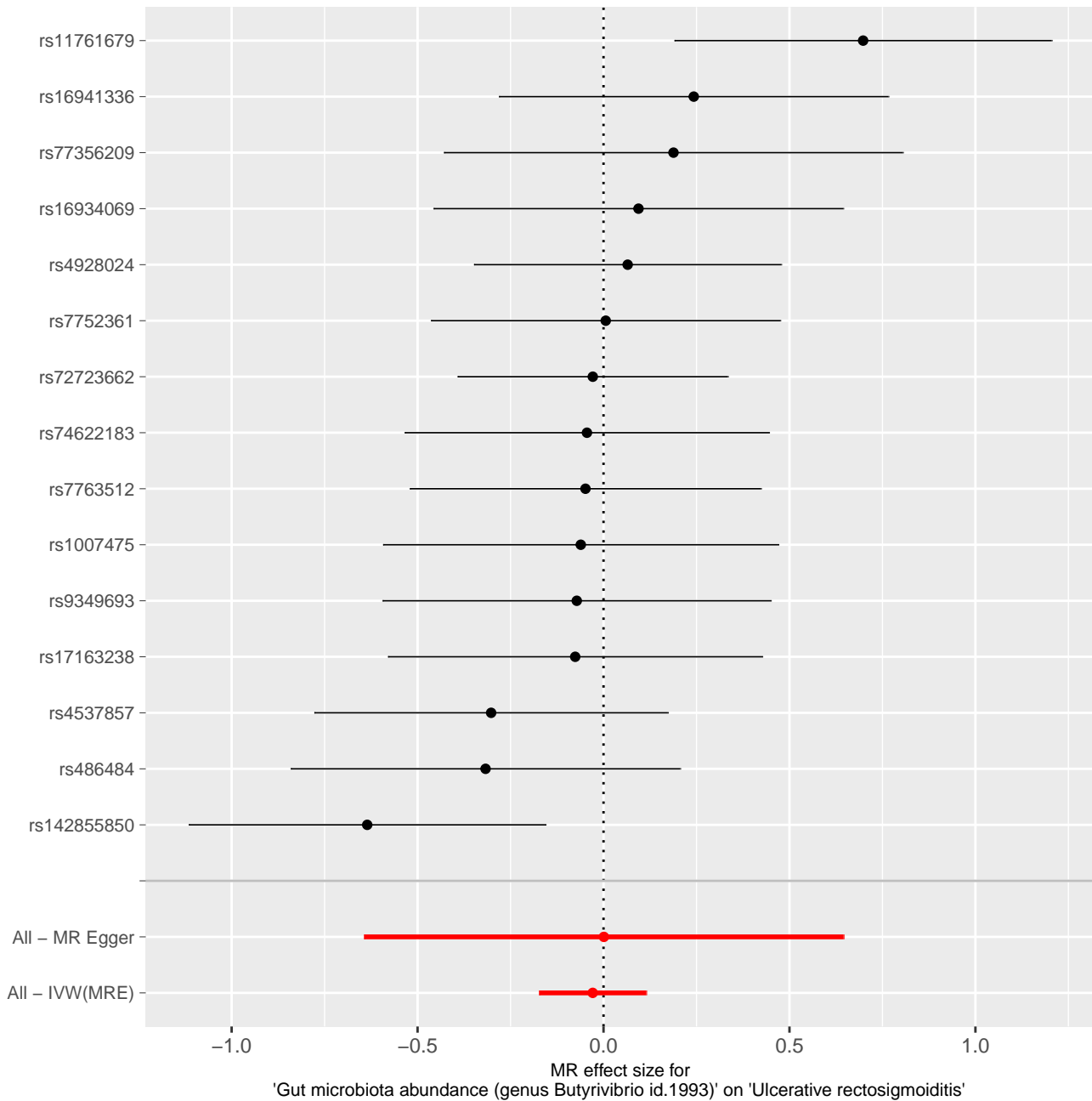


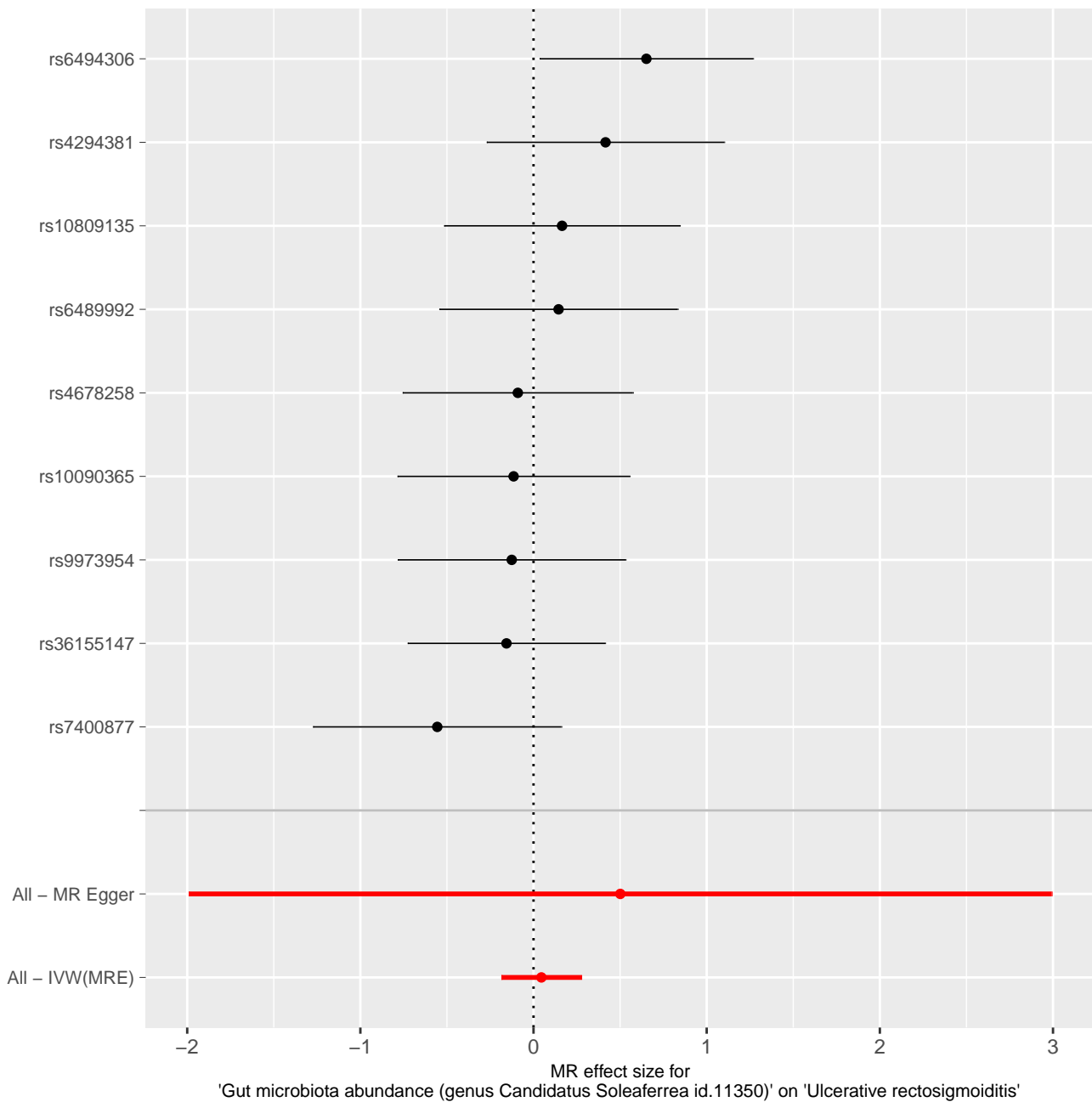


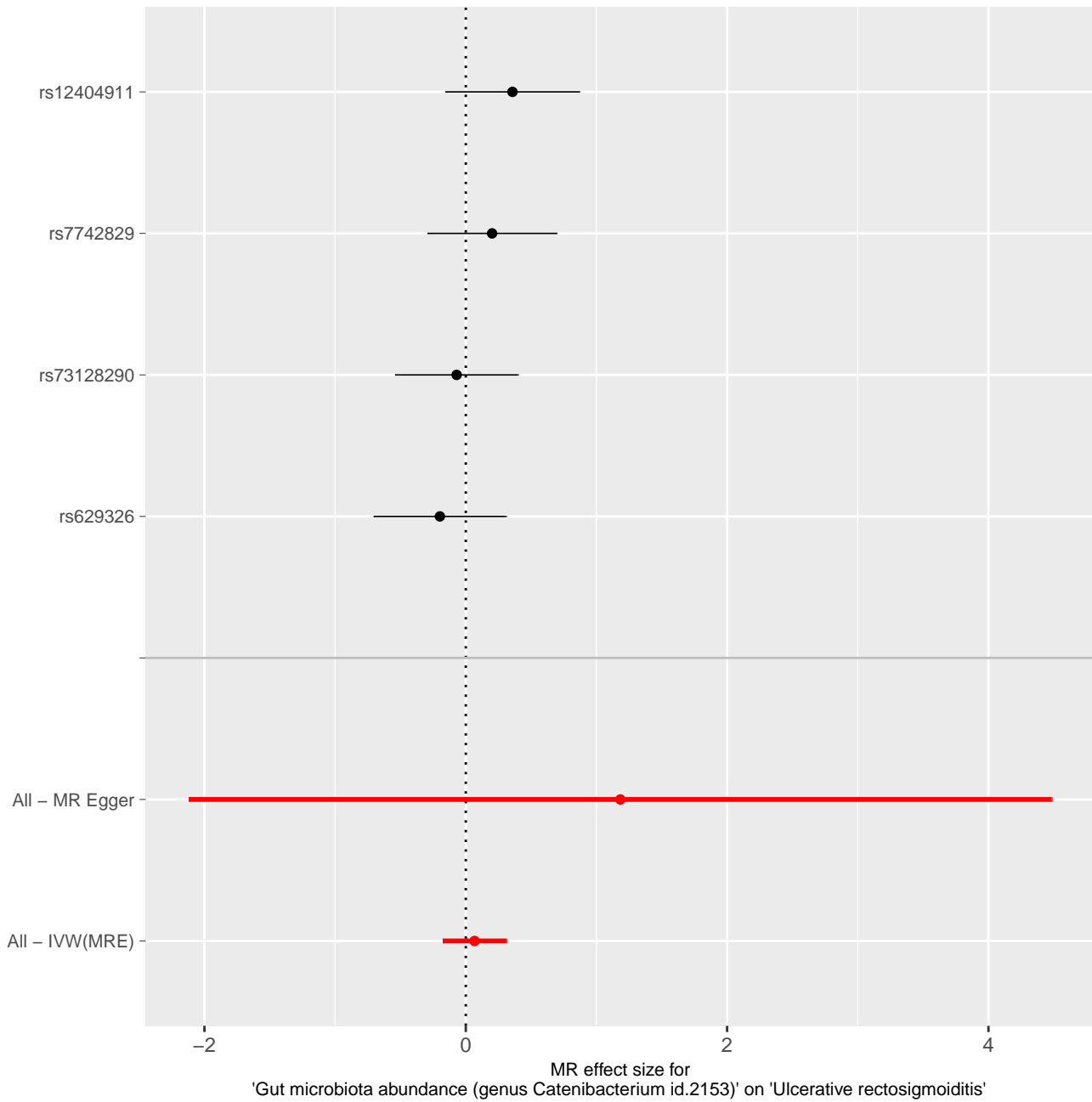


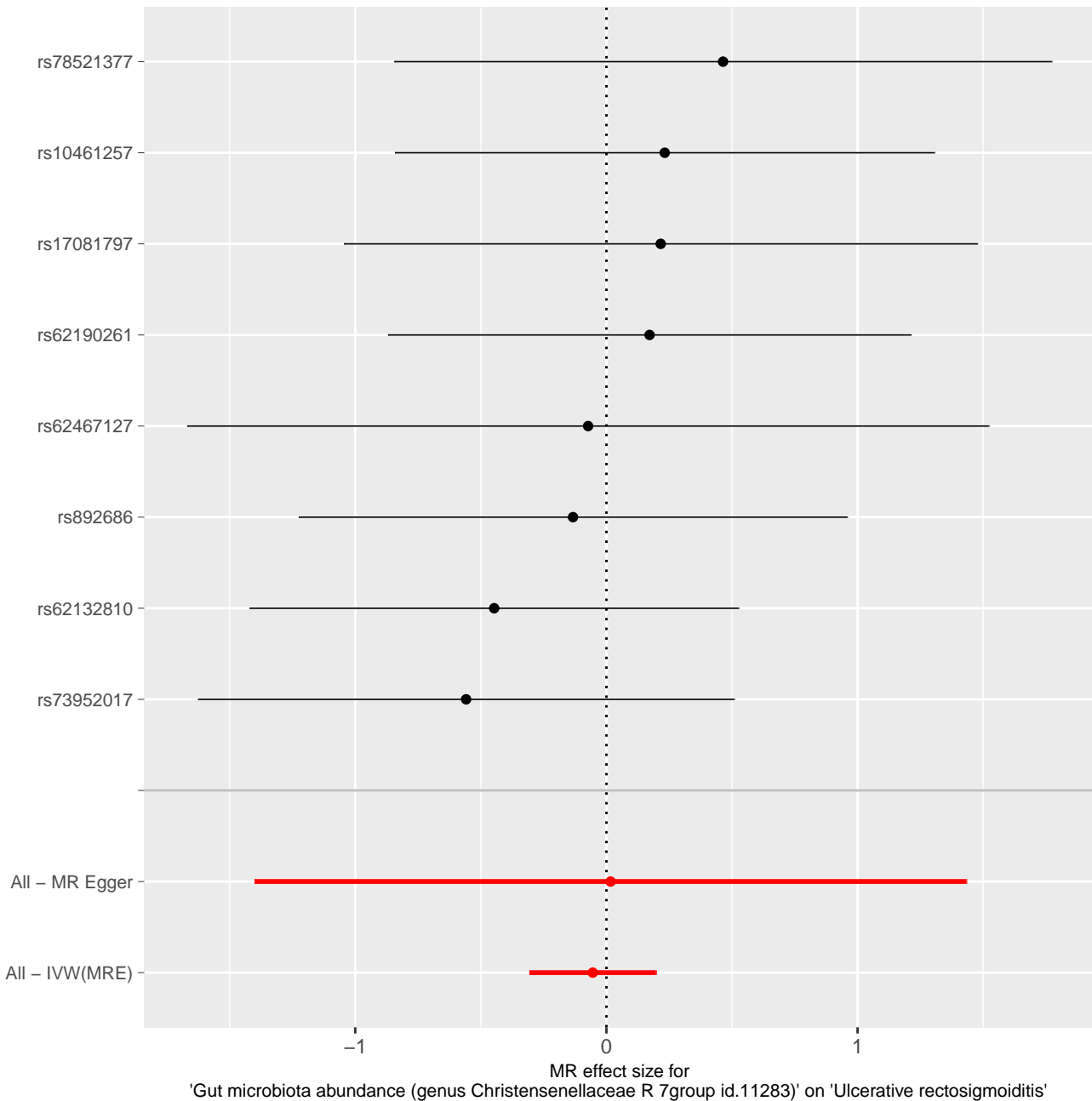




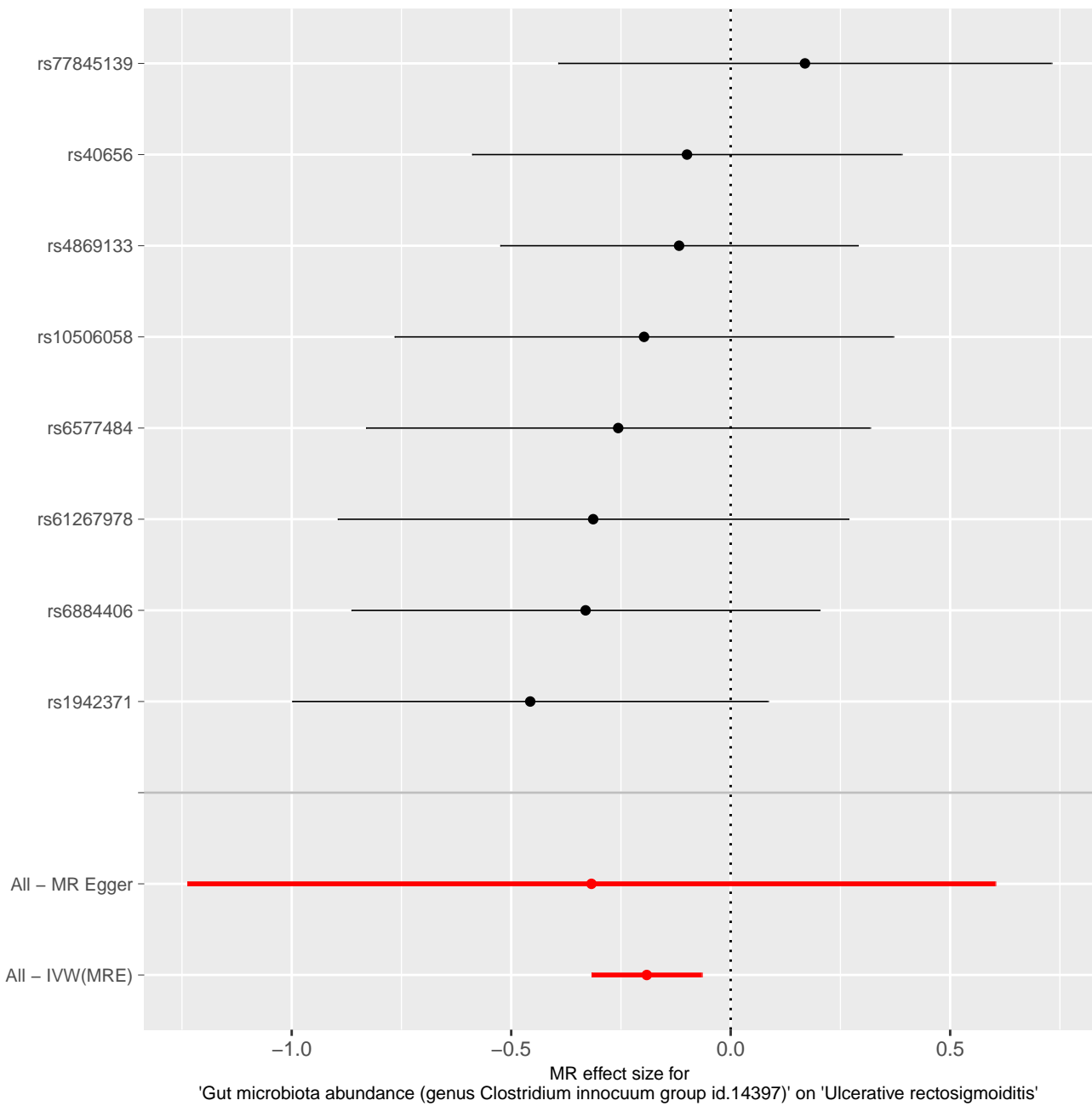


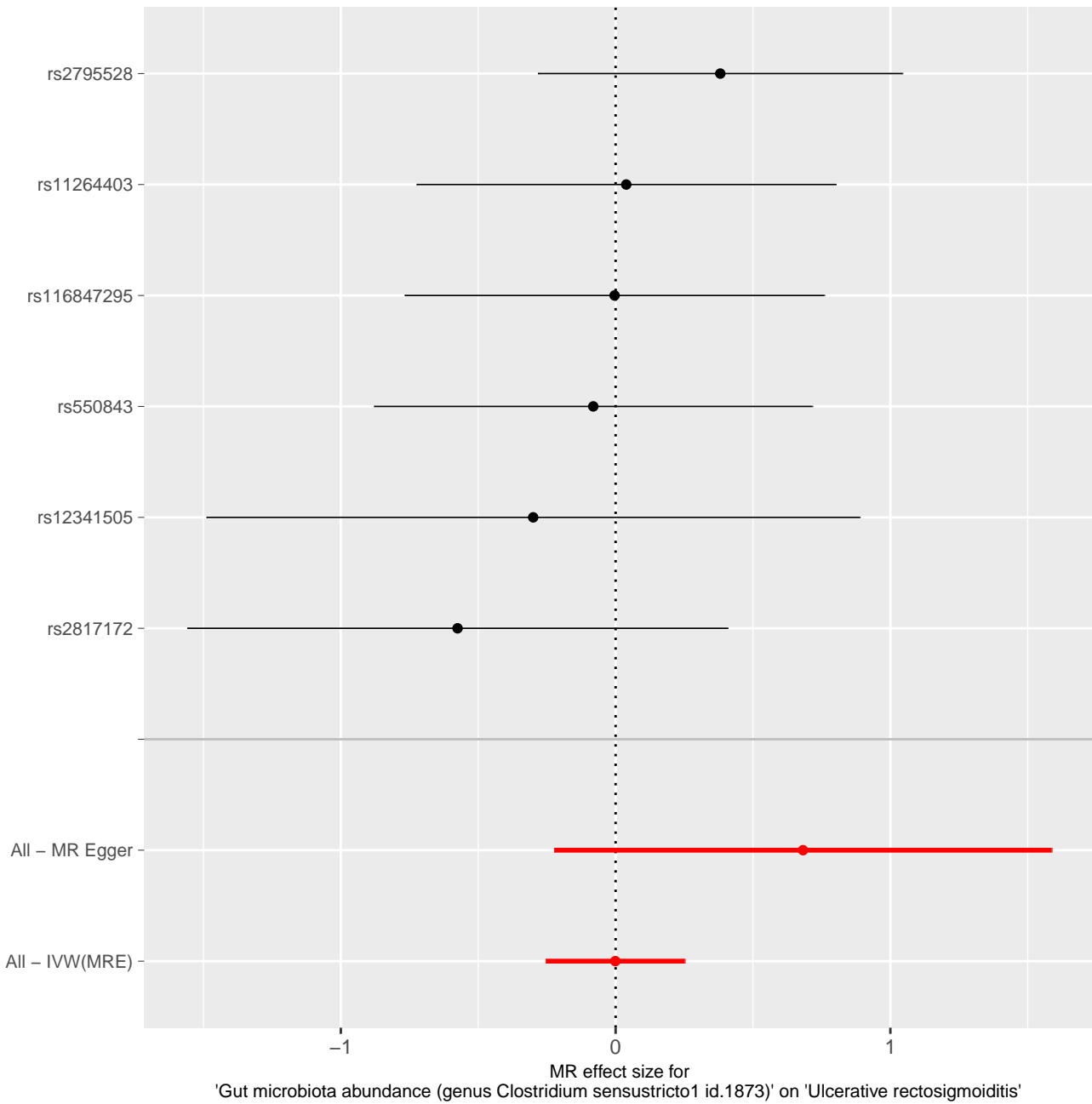


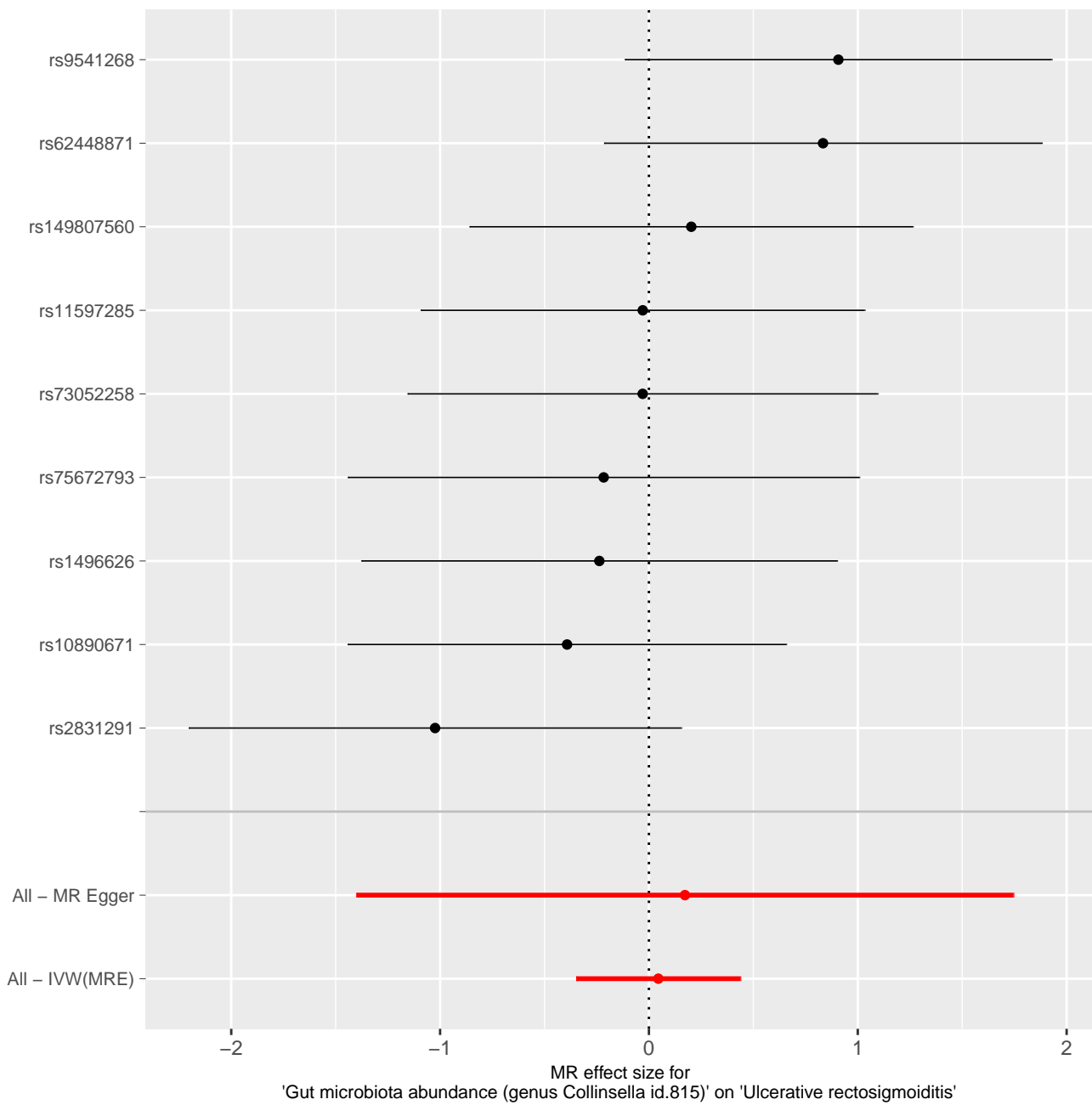


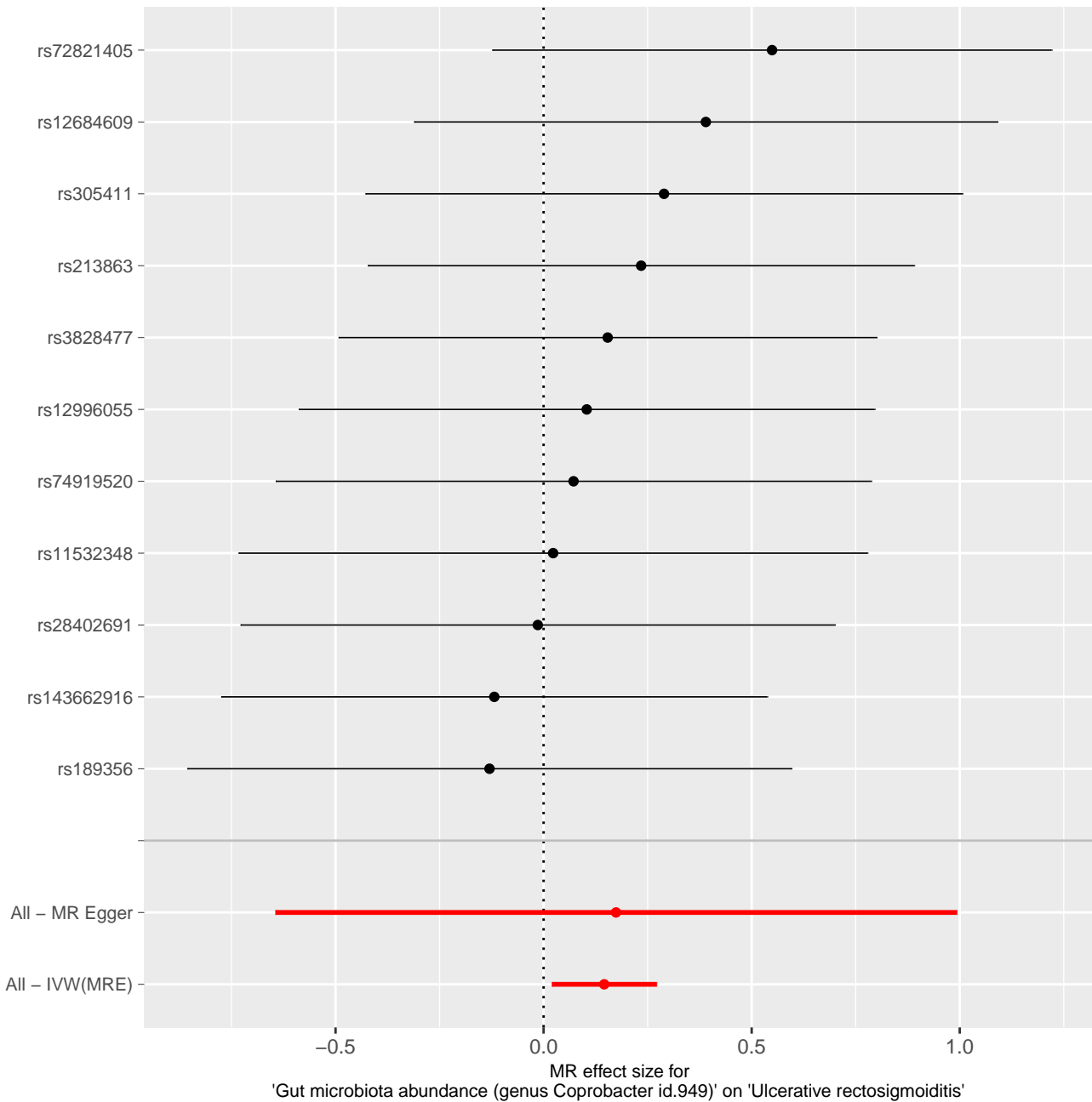


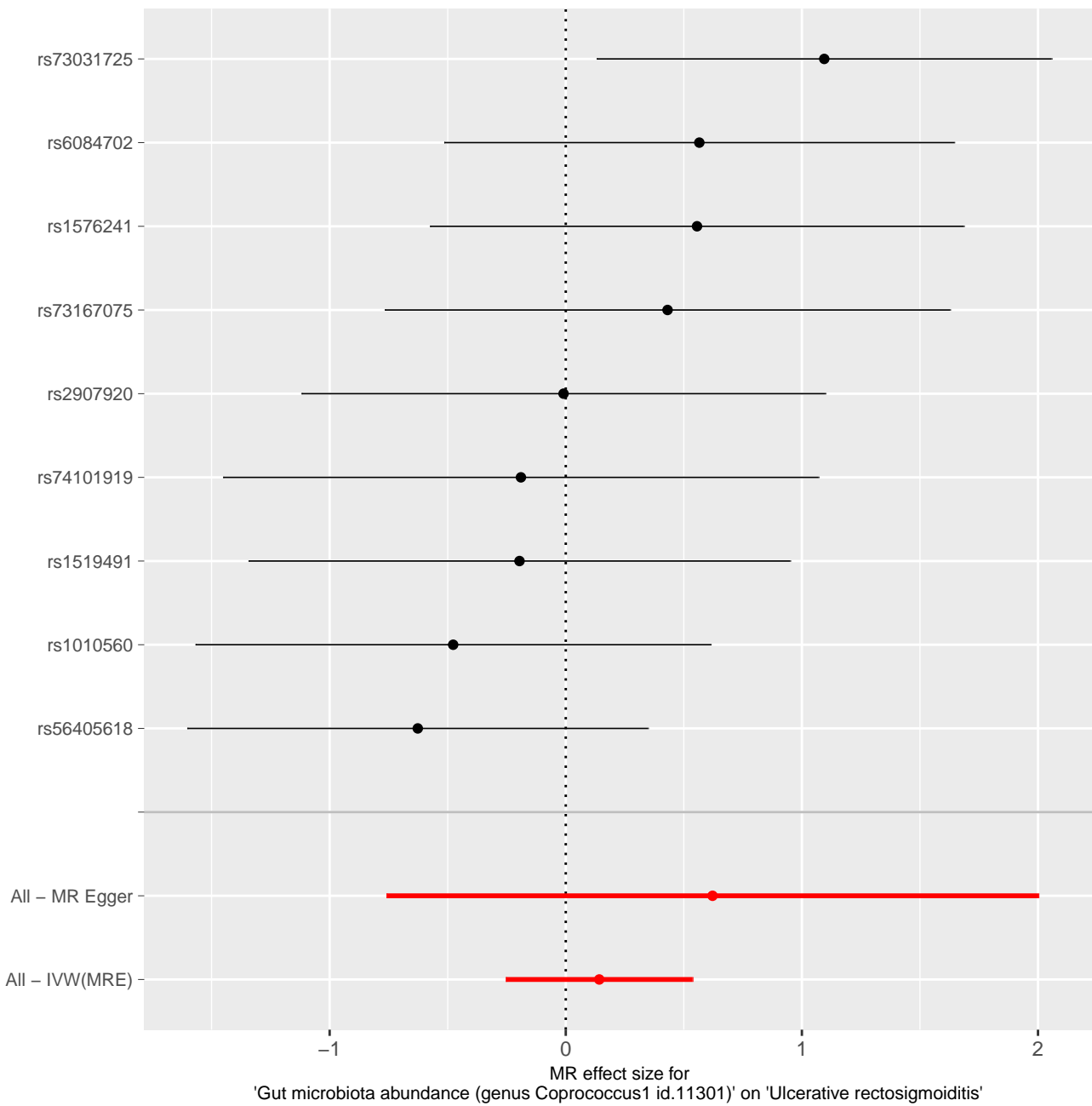
'Gut microbiota abundance (genus Christensenellaceae R 7group id.11283)' on 'Ulcerative rectosigmoiditis'

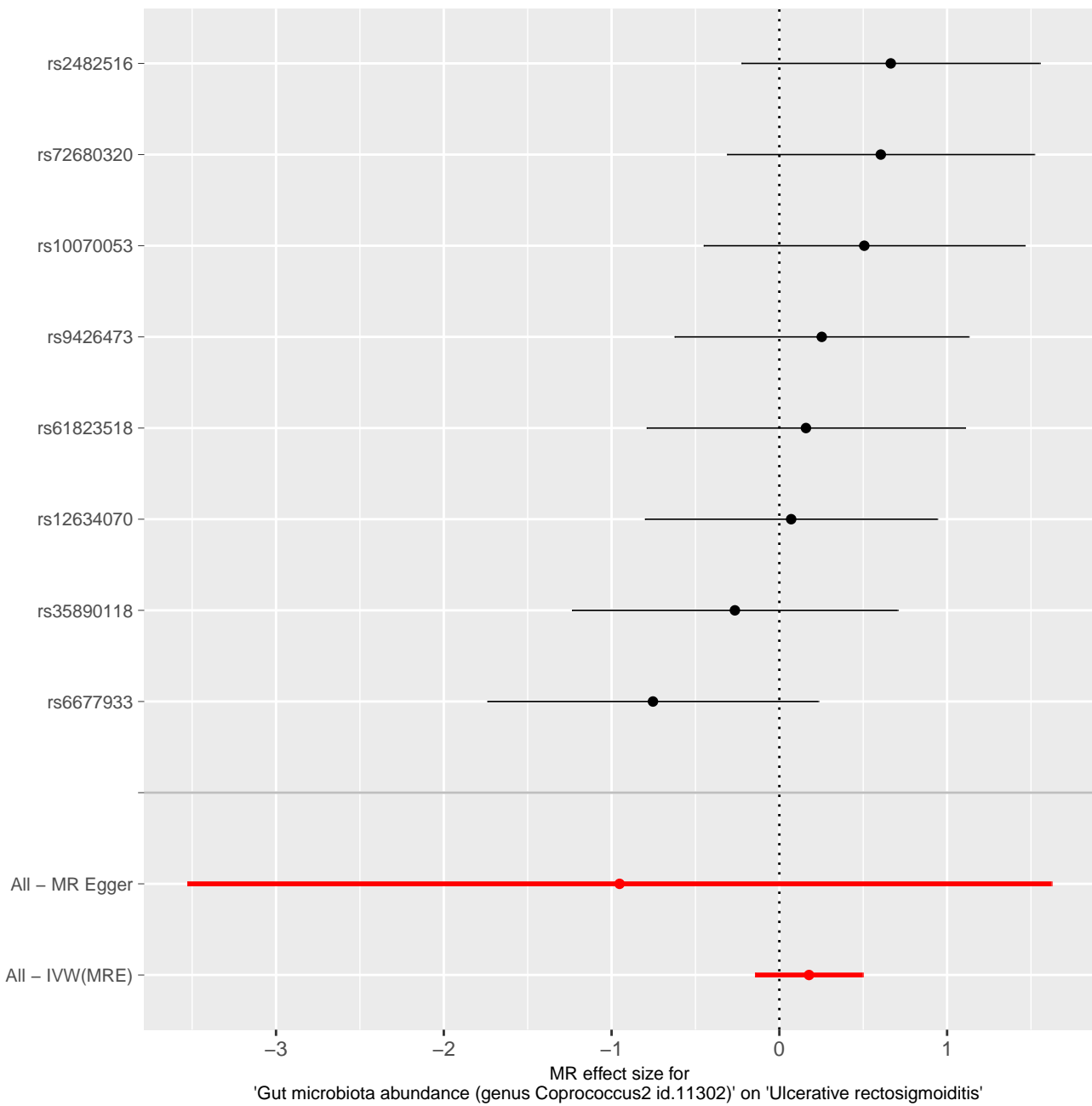


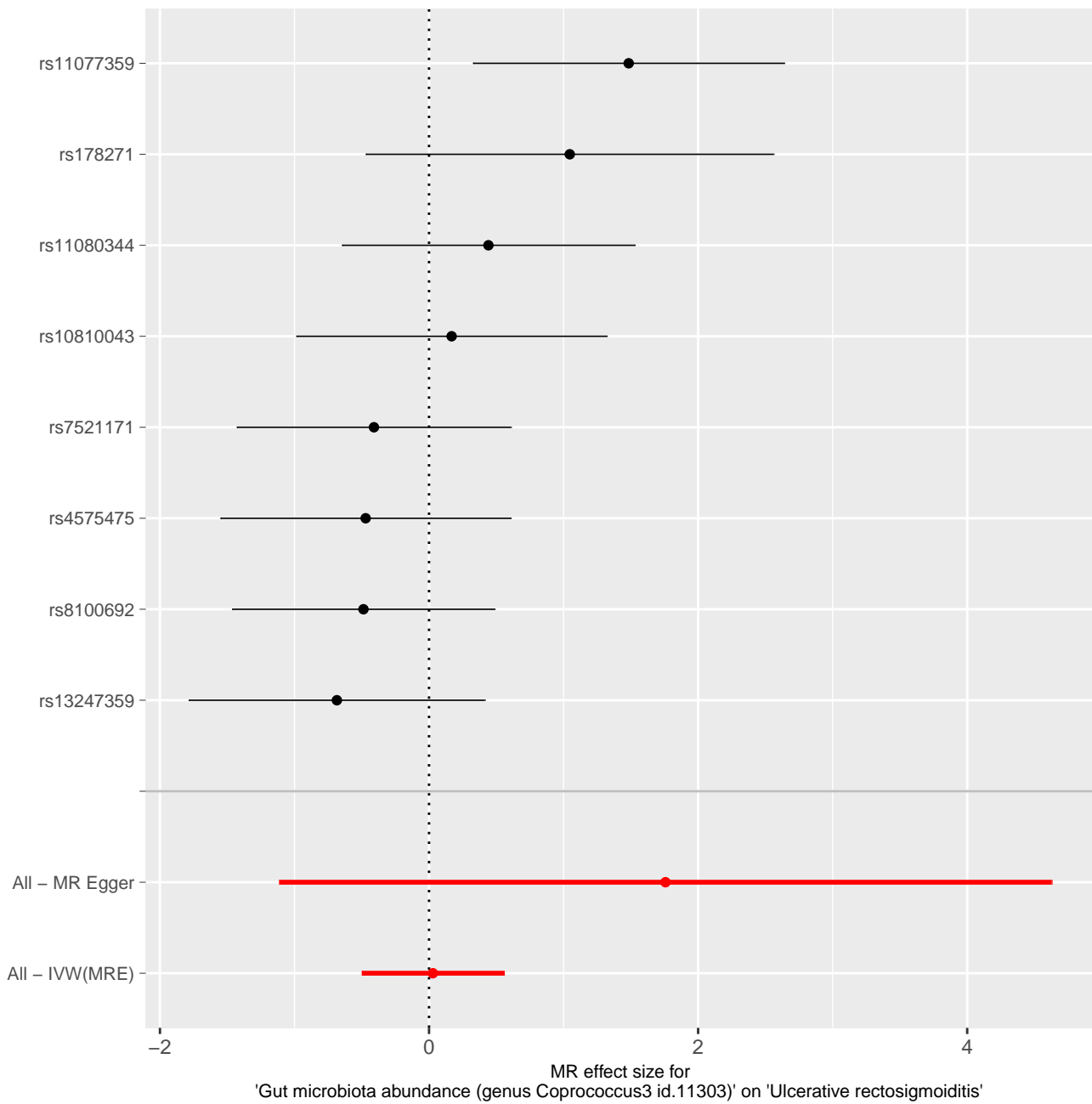


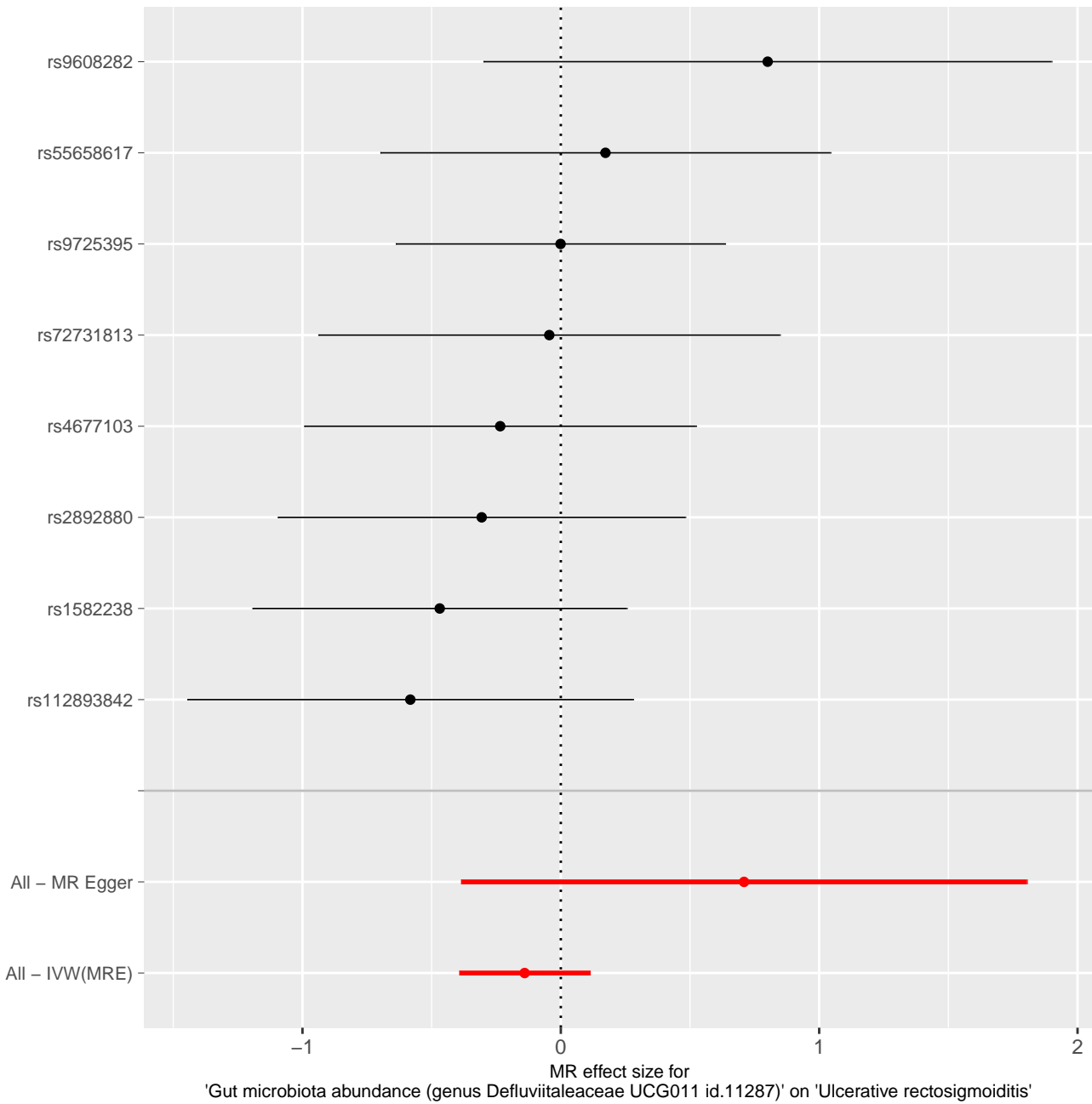


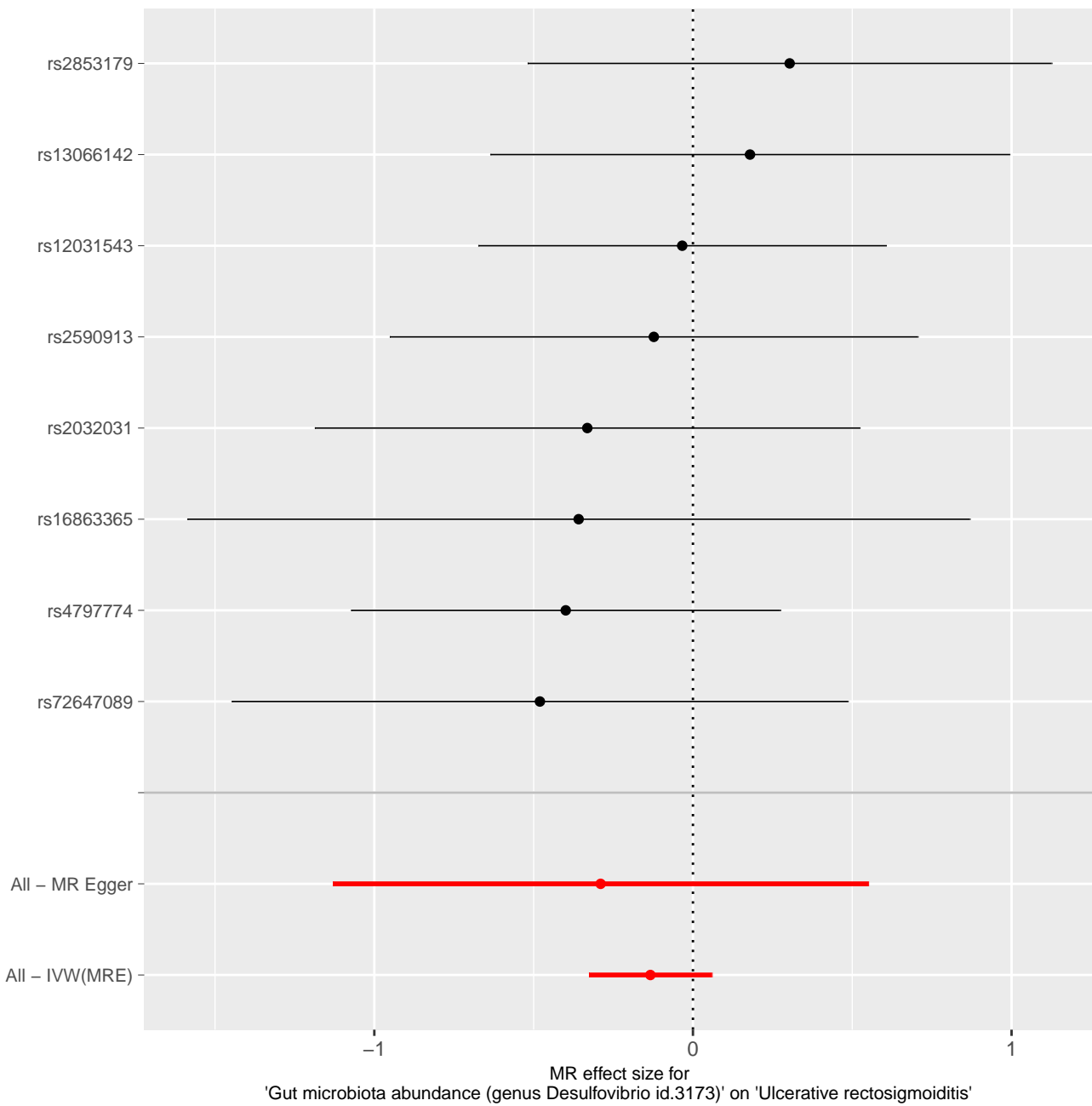




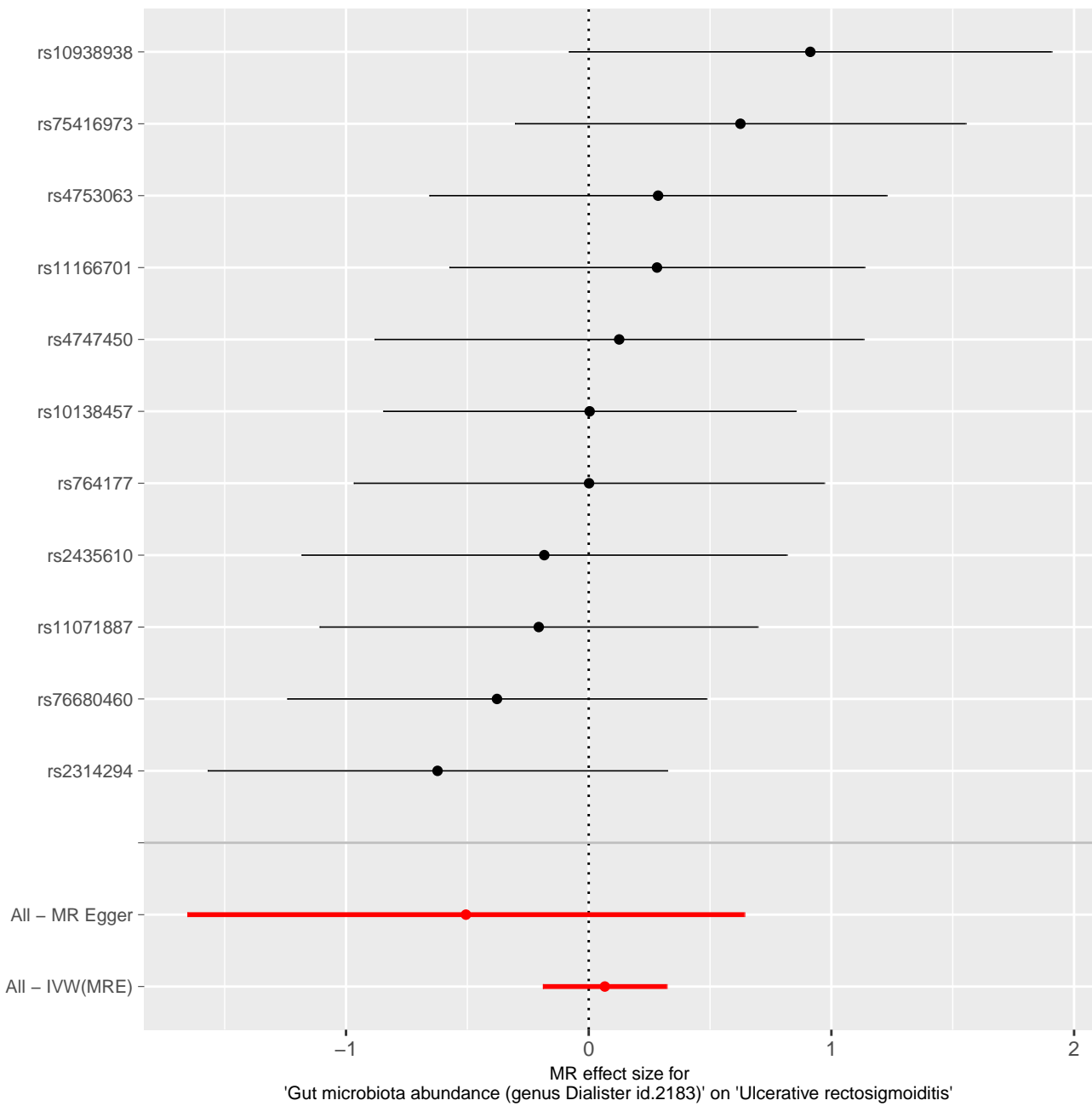




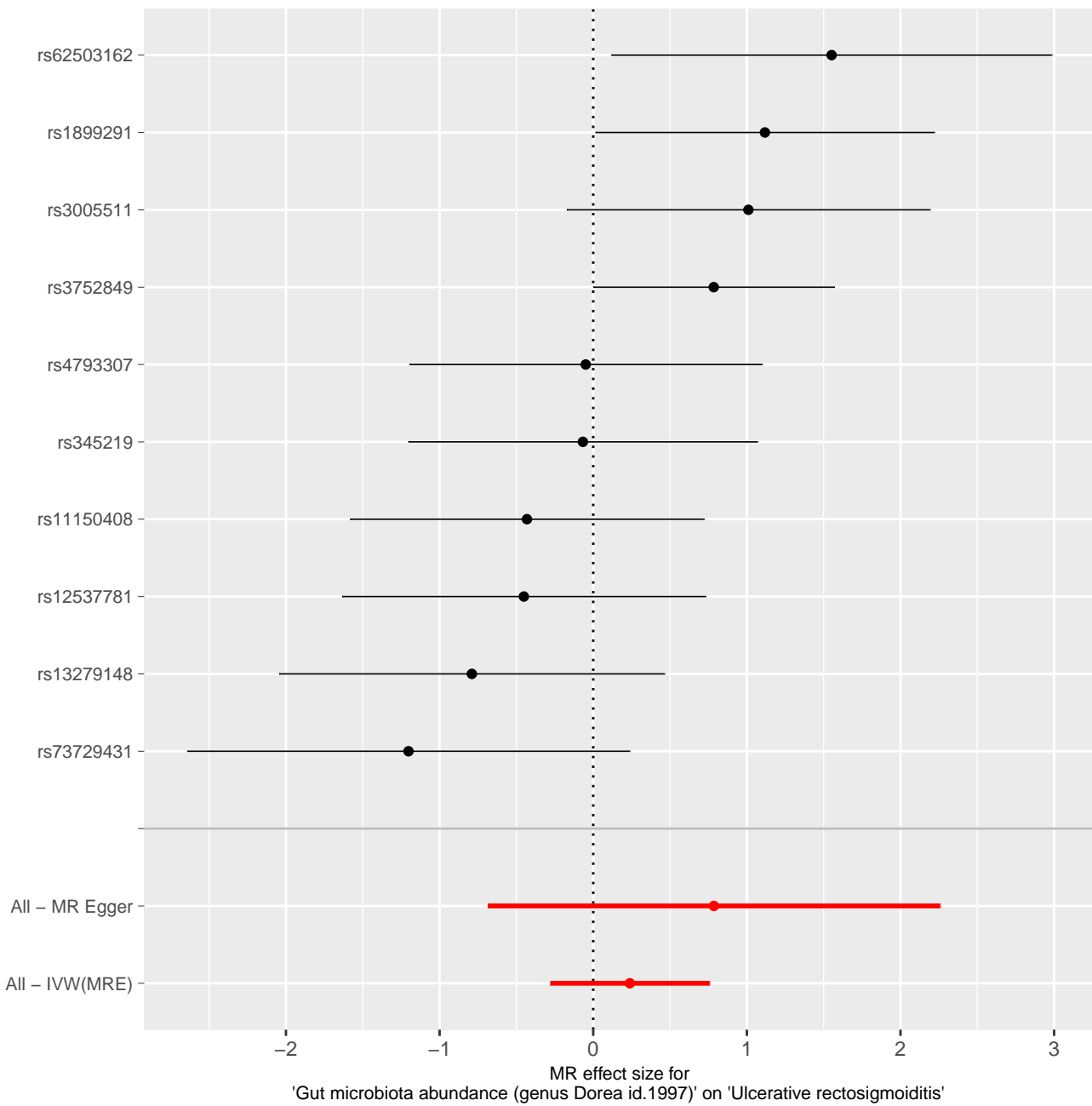


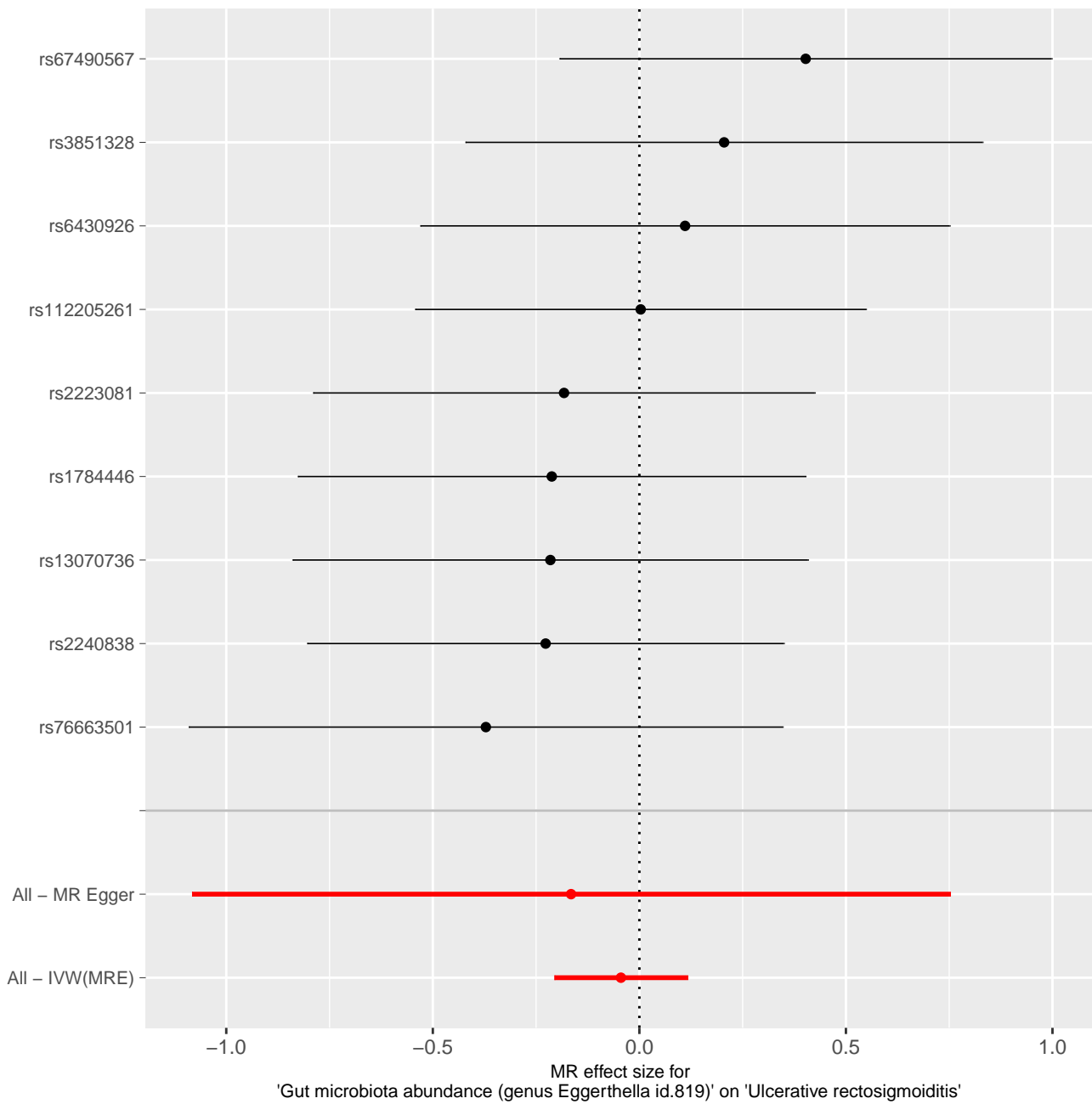


Batch 1562 : Gut microbiota abundance (genus Dialister id.2183) on Ulcerative rectosigmoiditis

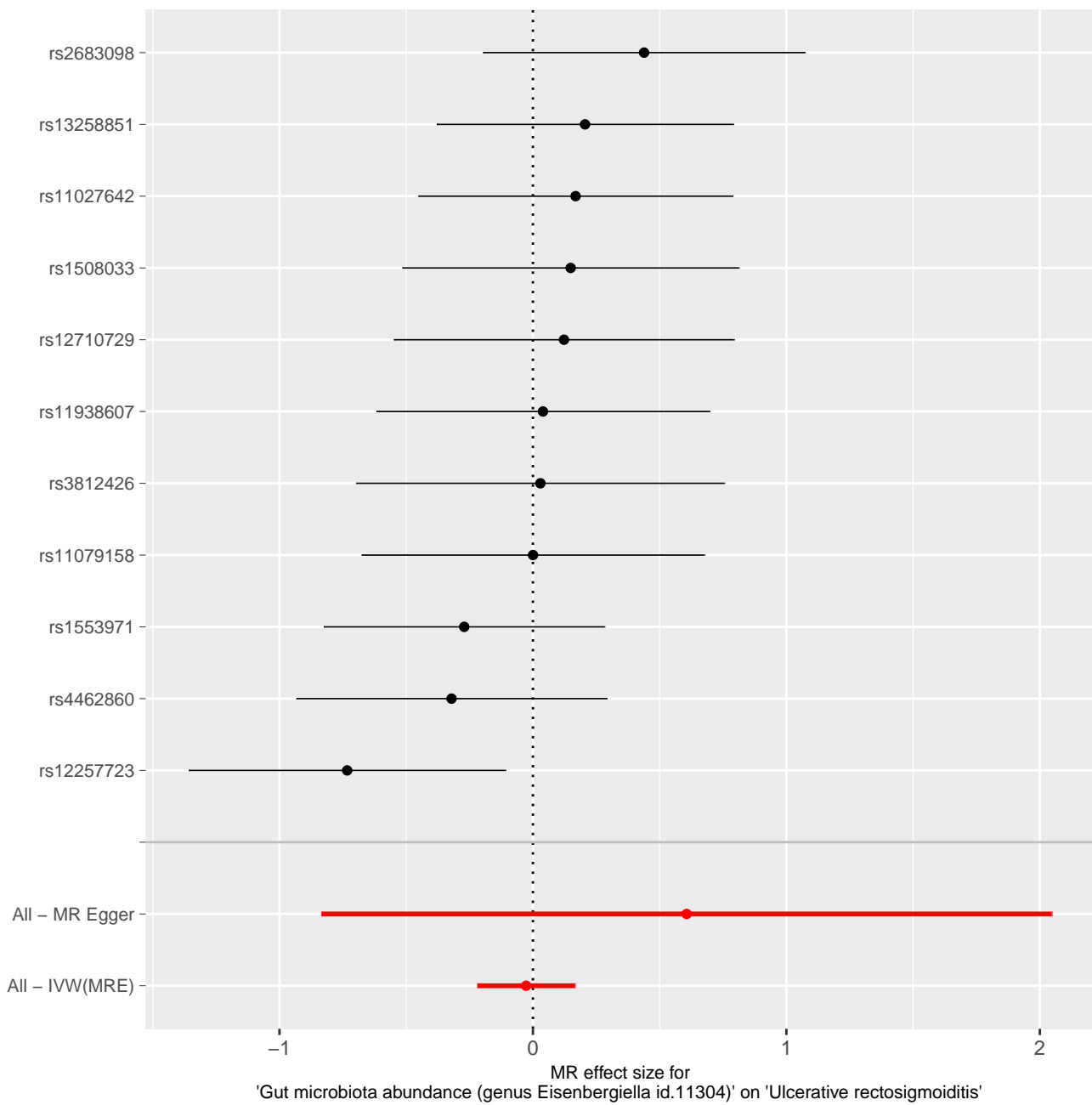


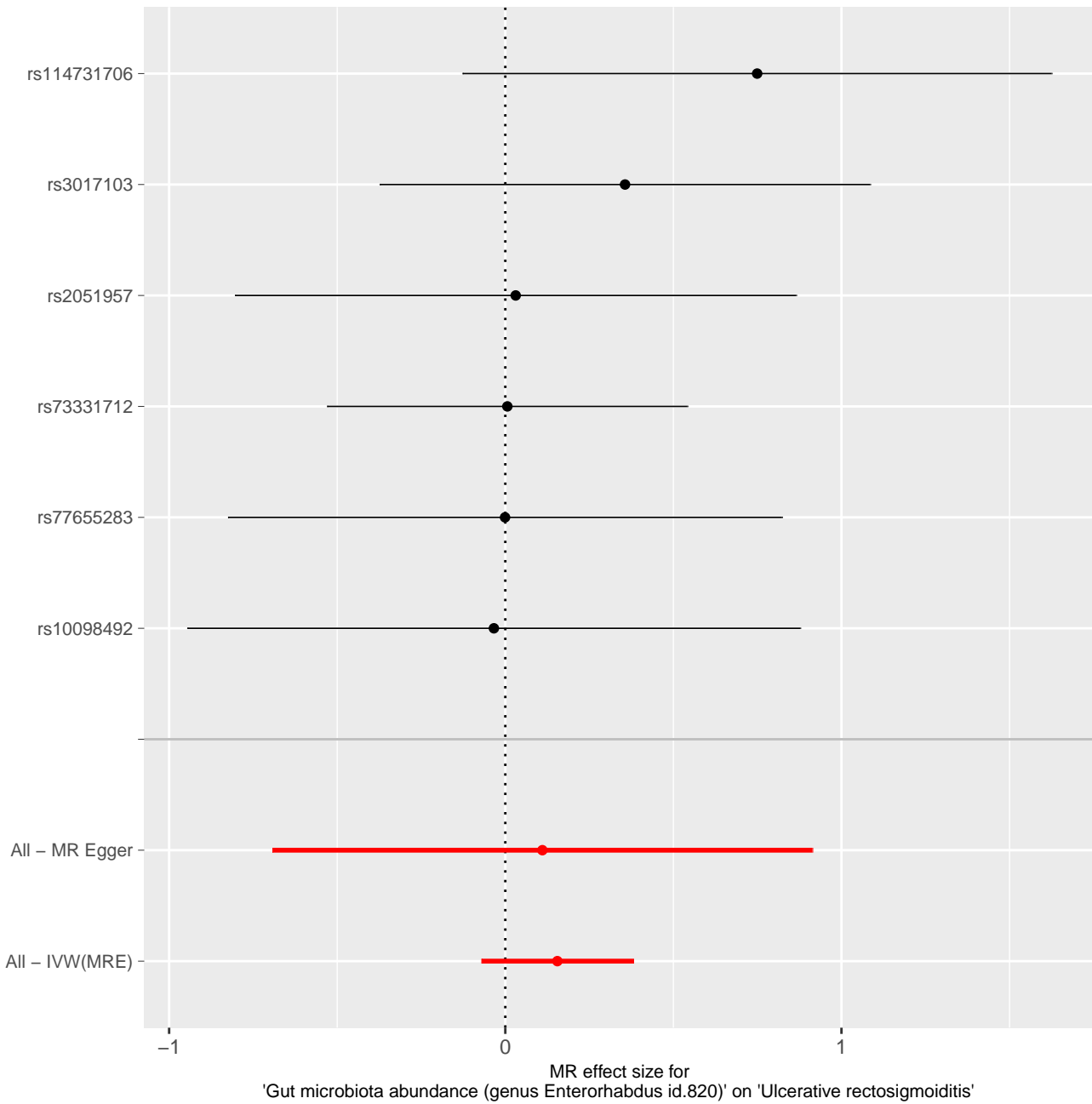
Batch 1563 : Gut microbiota abundance (genus Dorea id.1997) on Ulcerative rectosigmoiditis

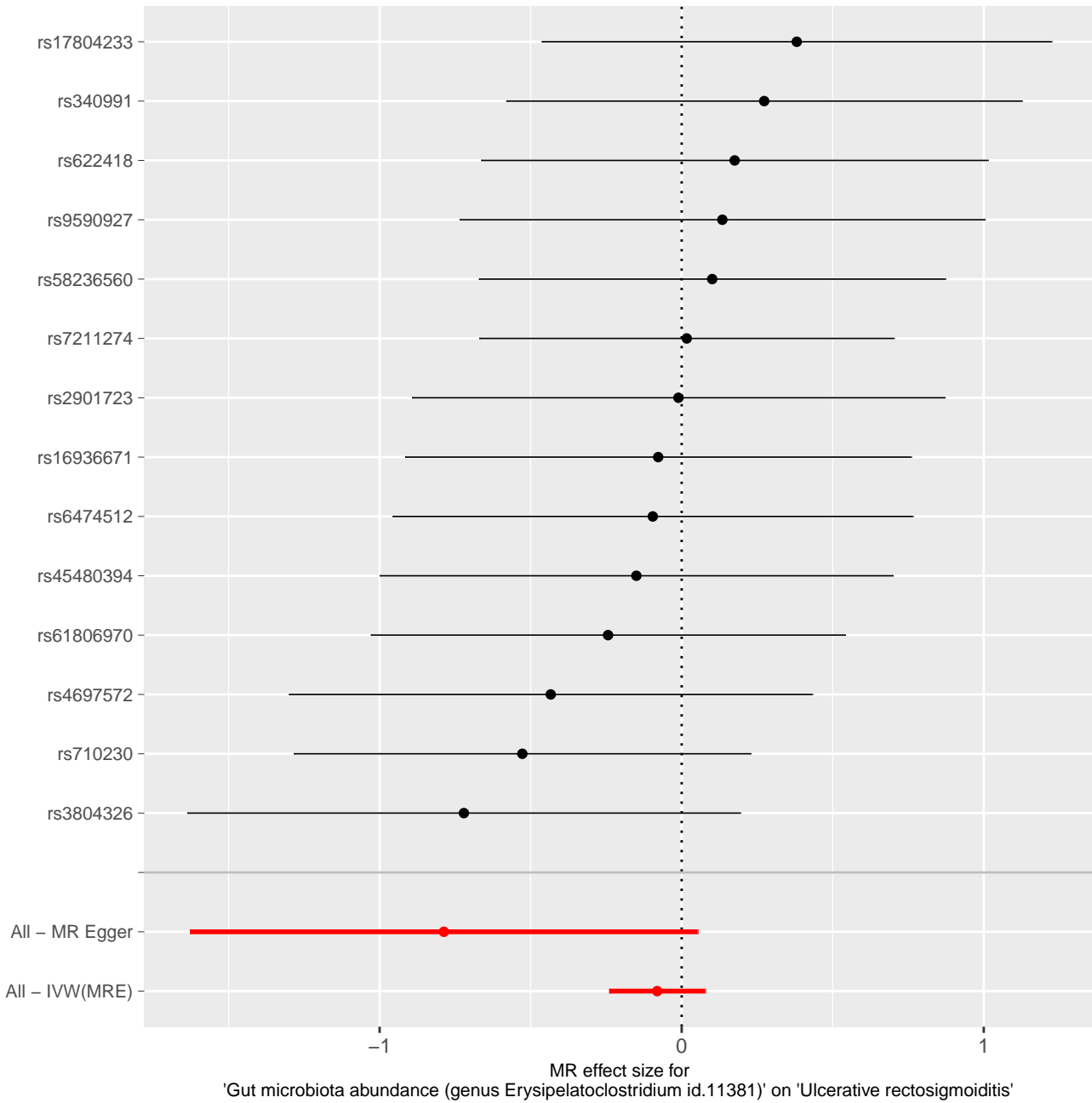


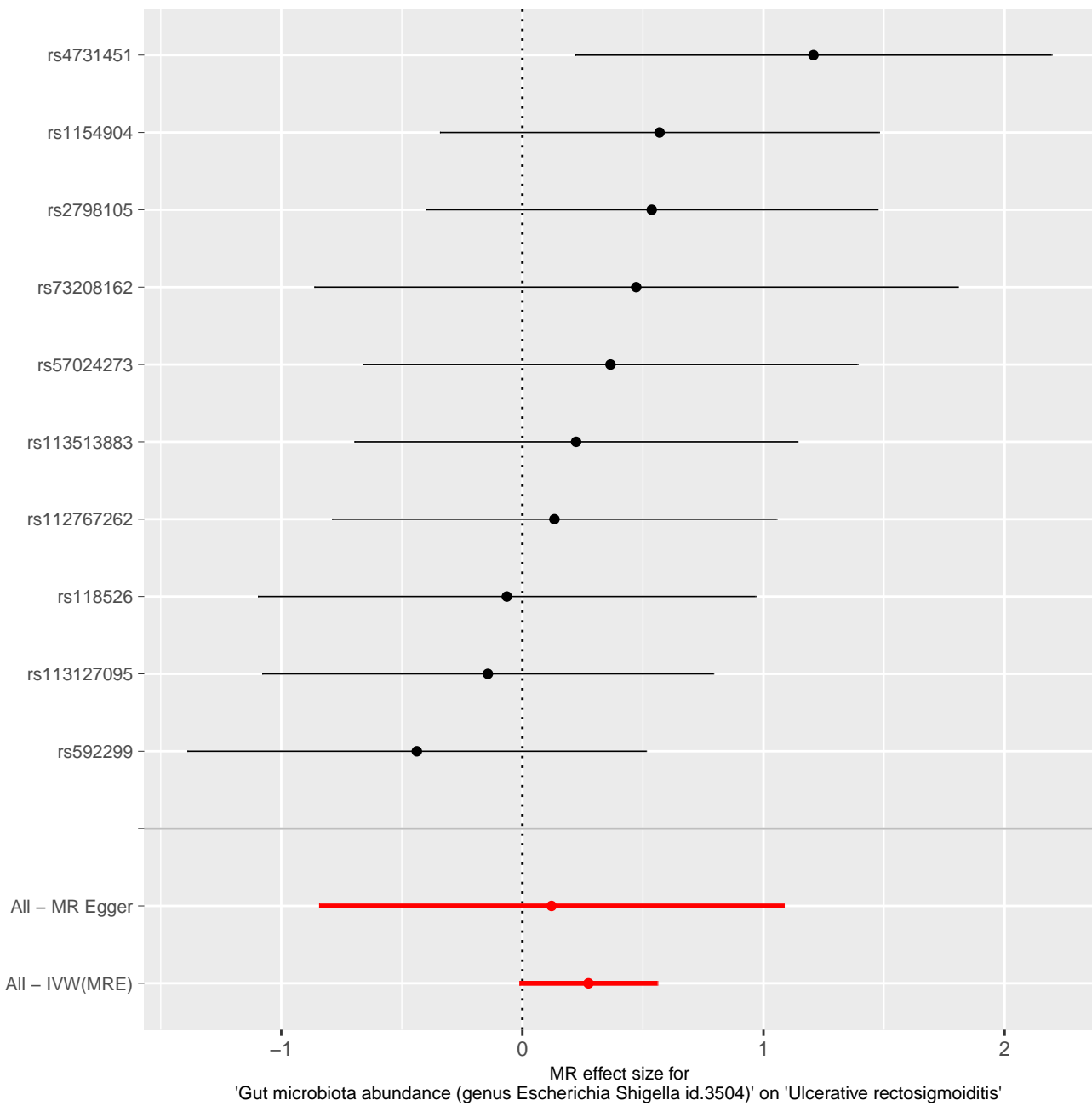


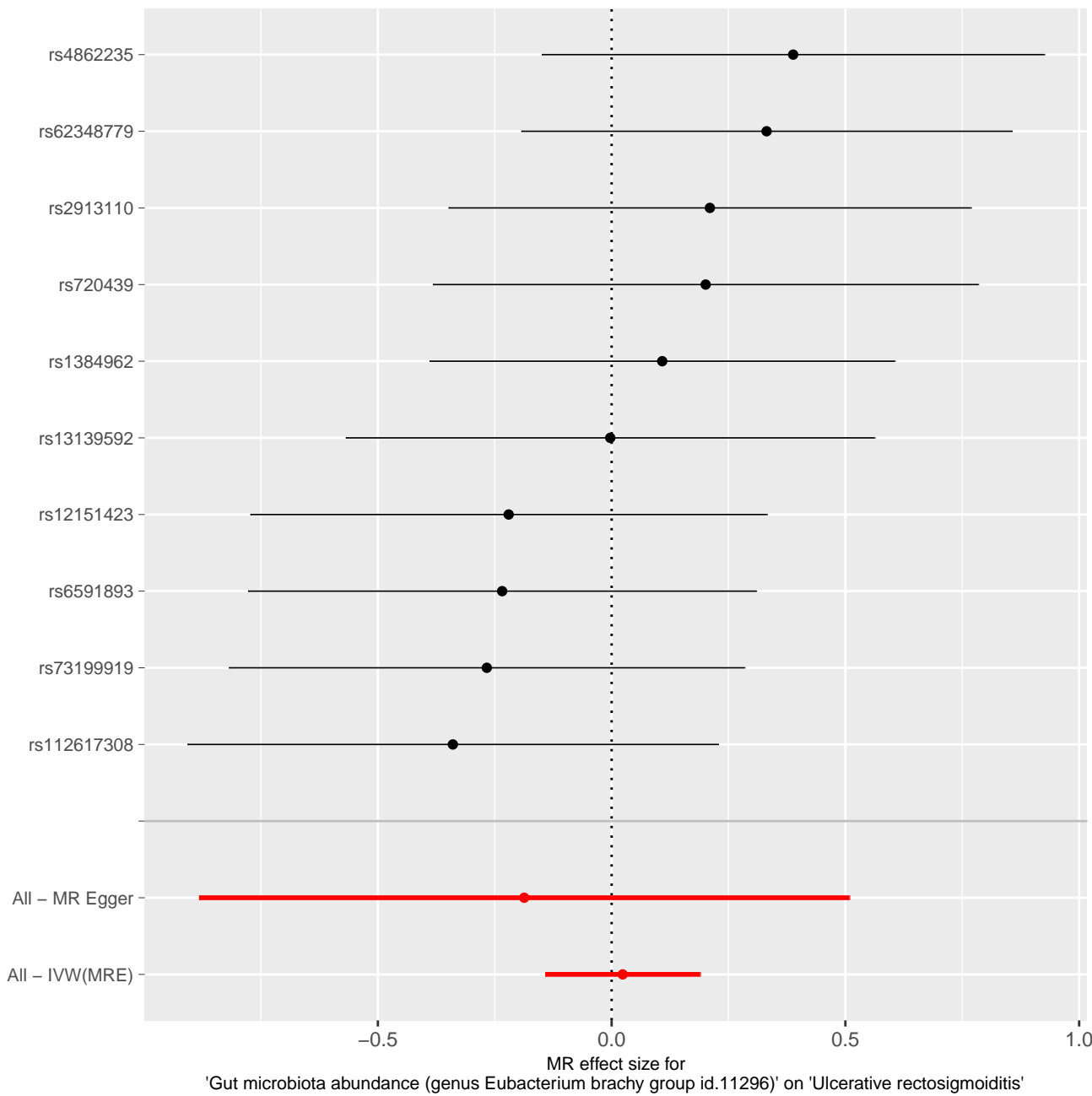
Batch 1565 : Gut microbiota abundance (genus Eisenbergiella id.11304) on Ulcerative rectosigmoiditis

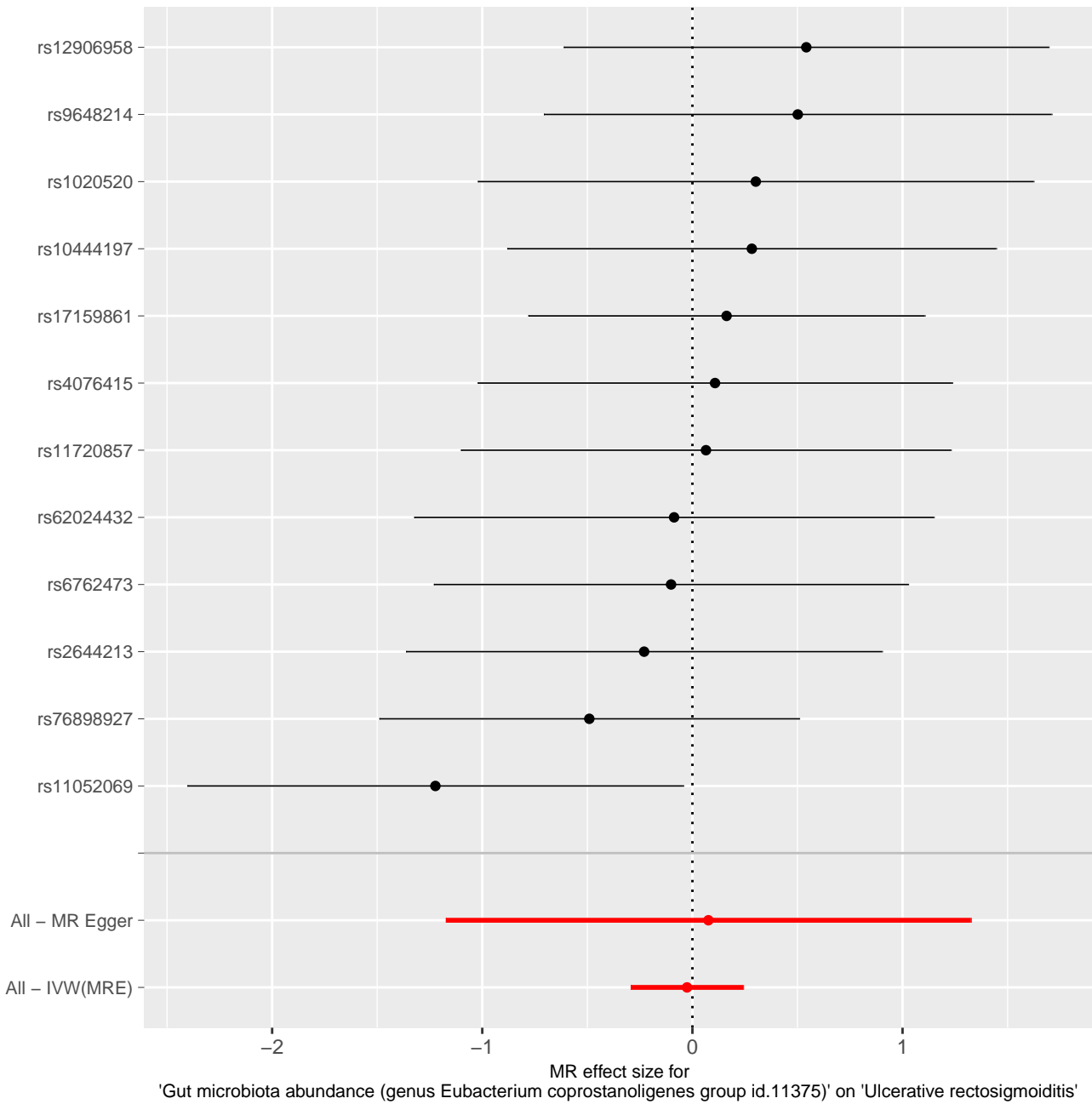


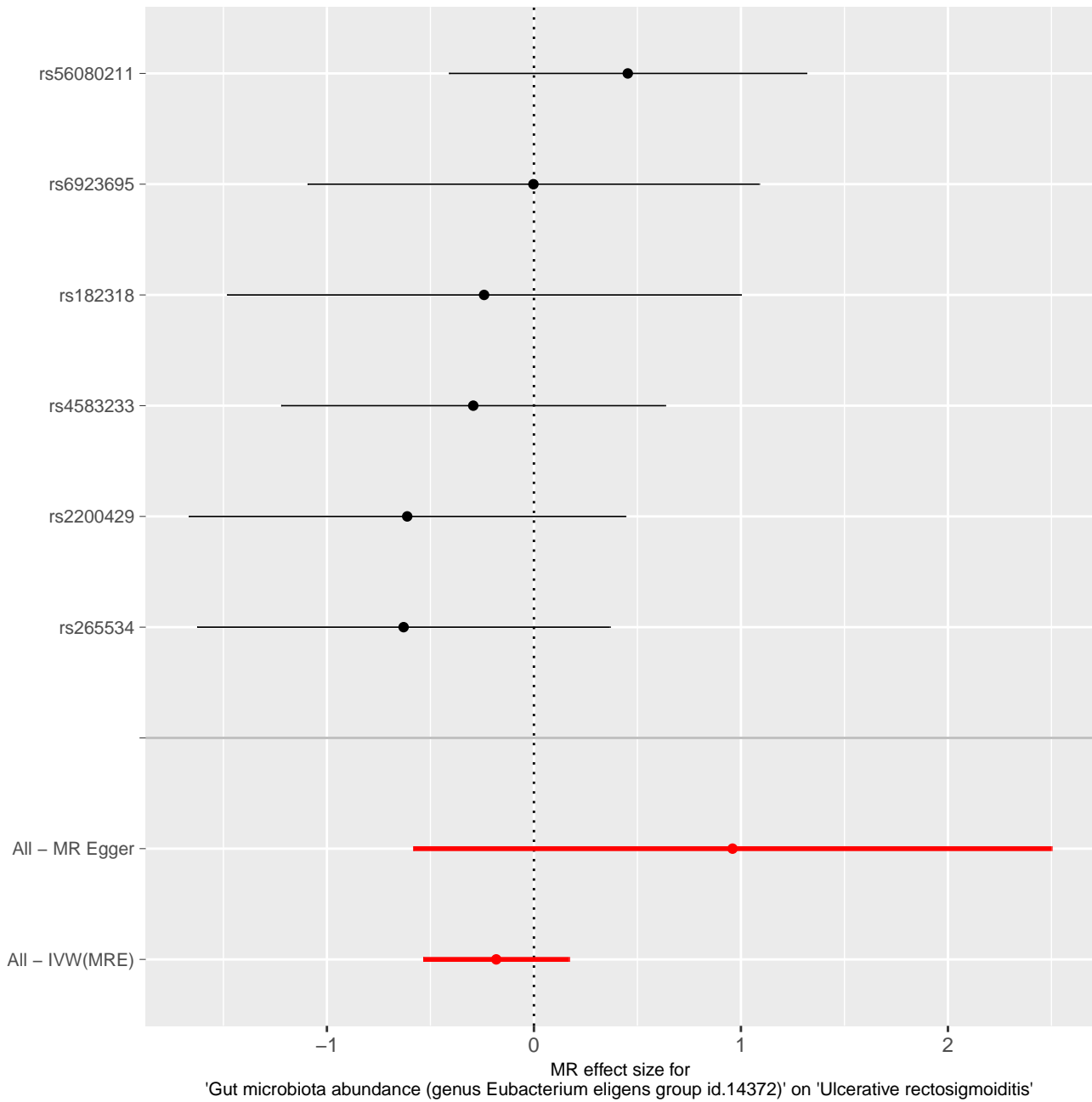


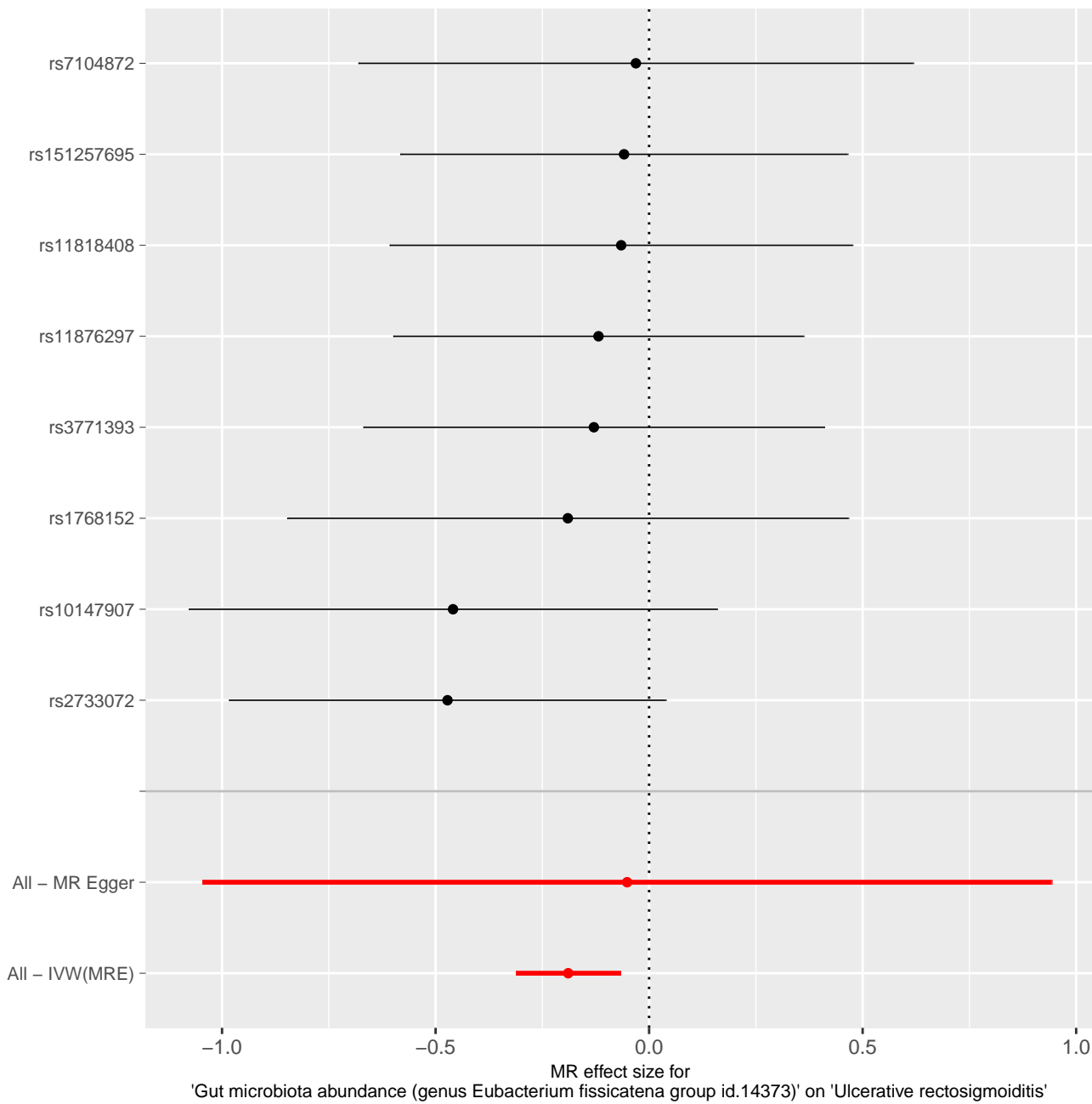


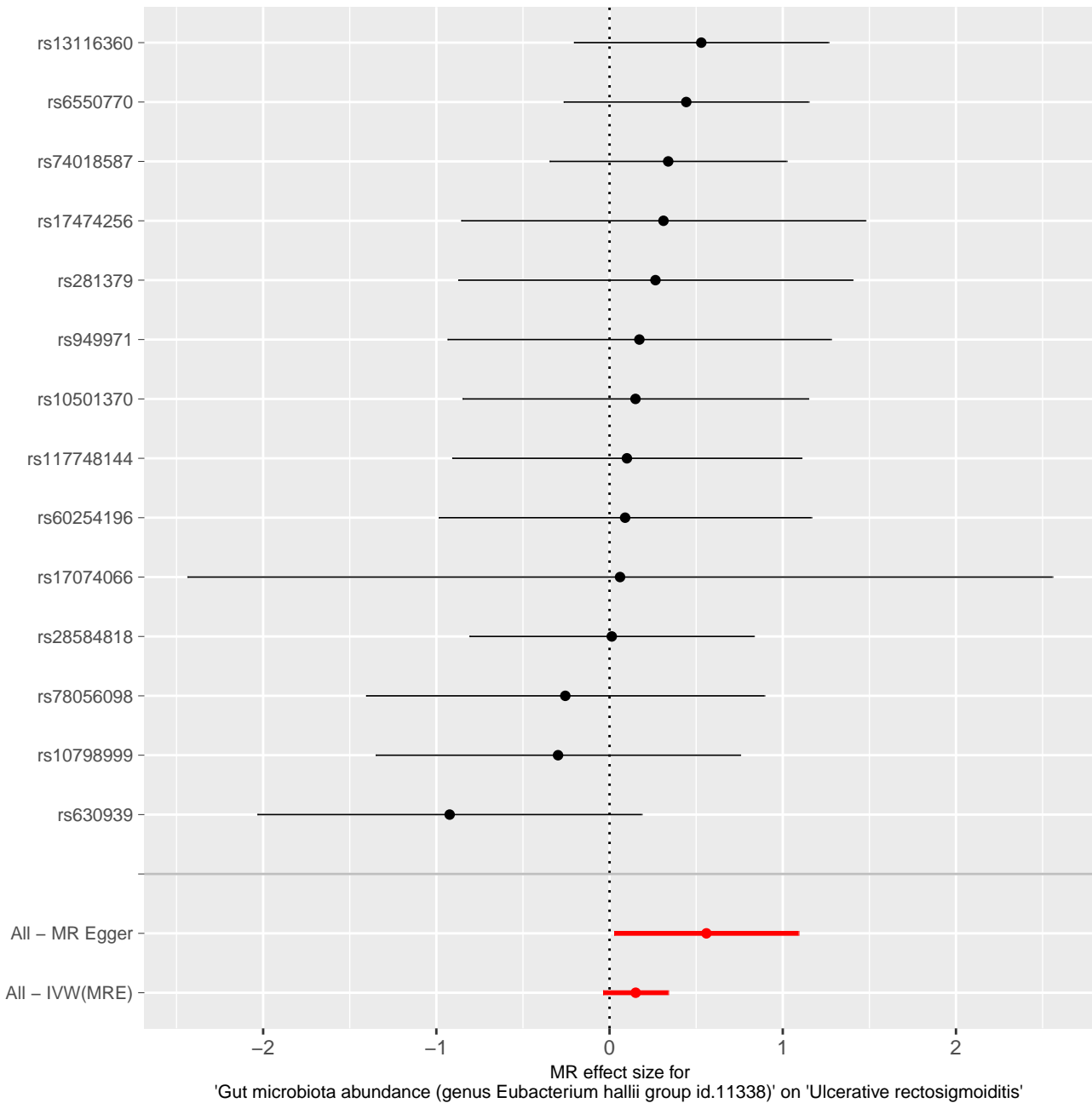


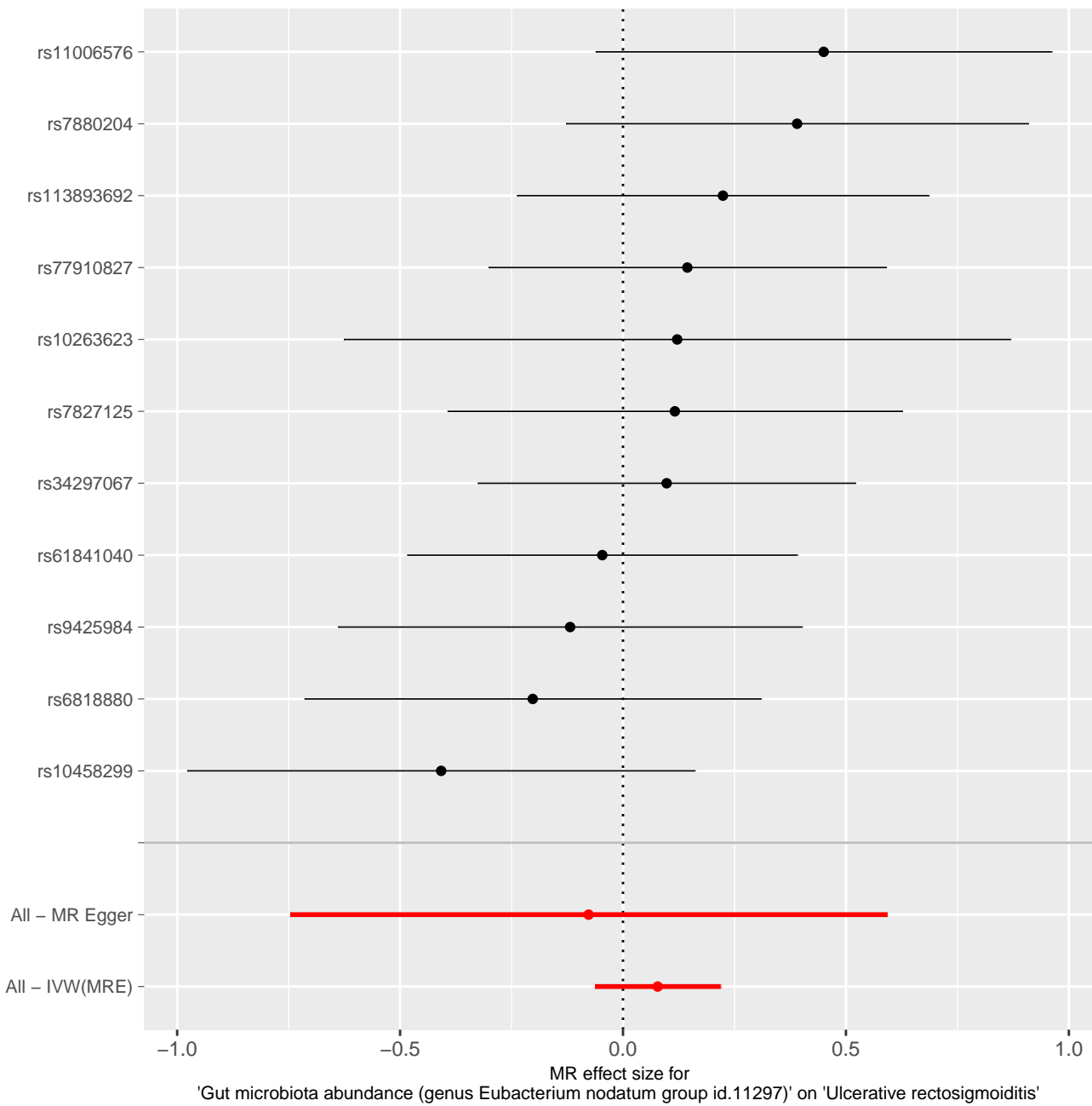


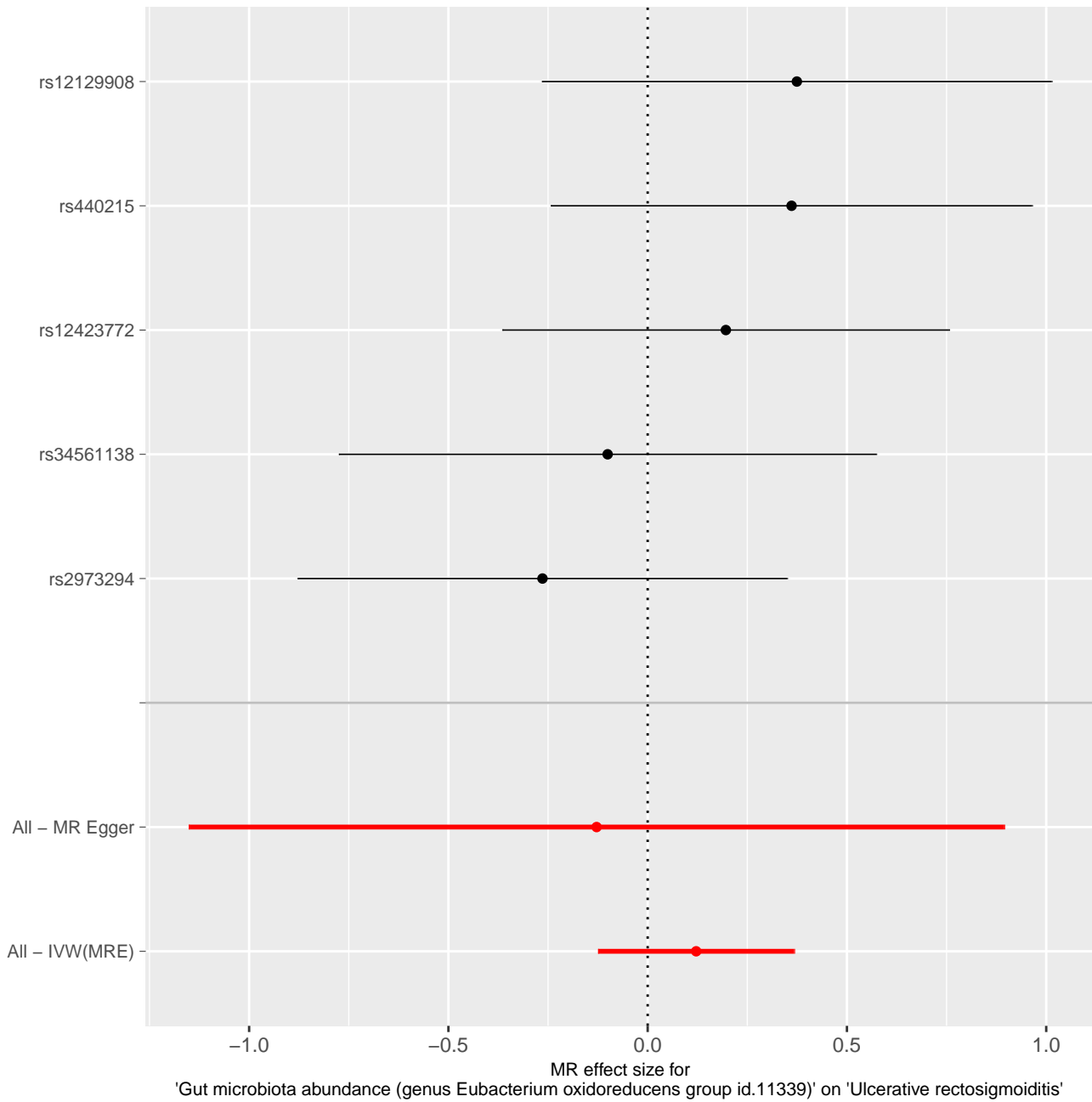


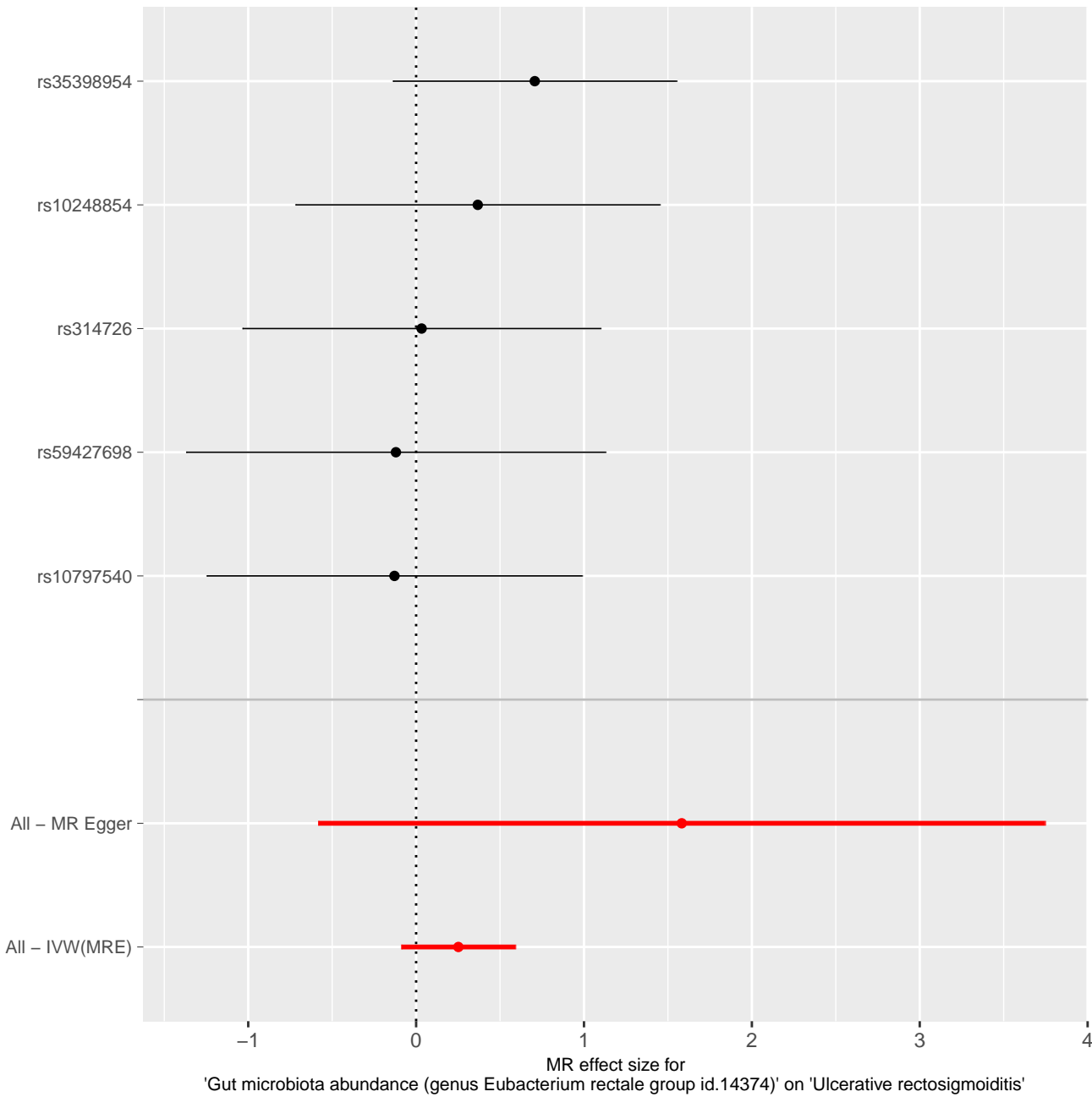


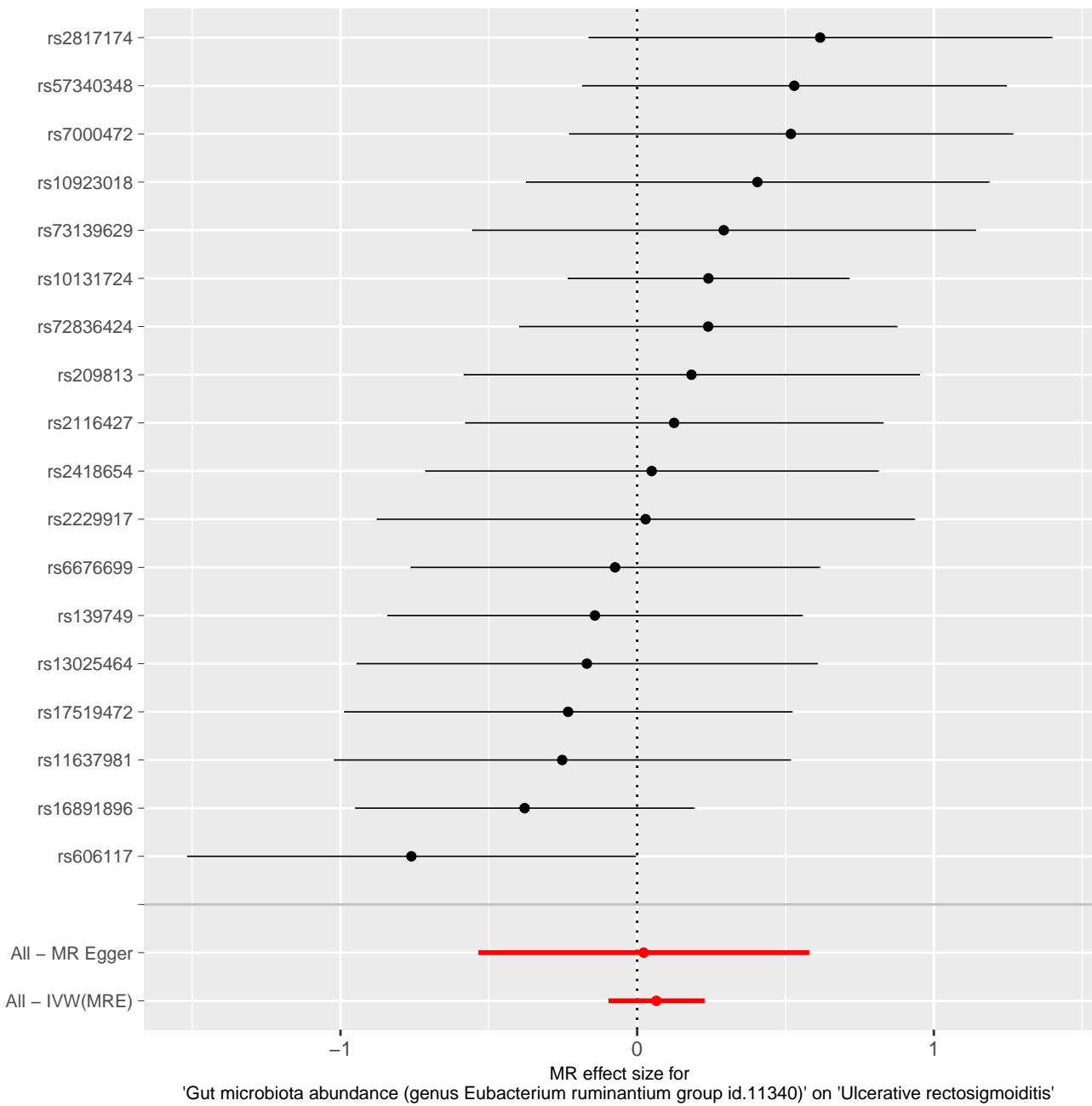


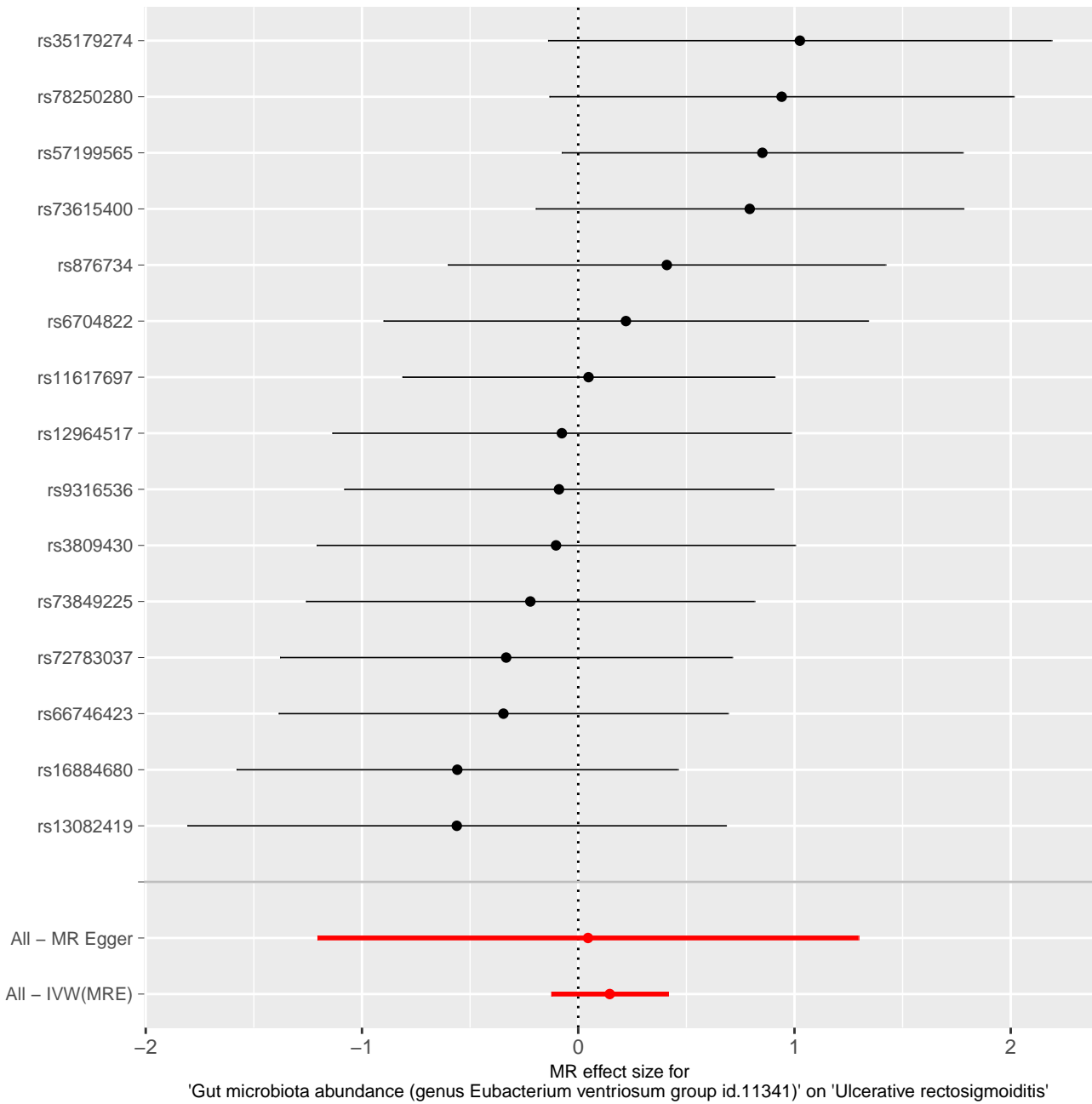


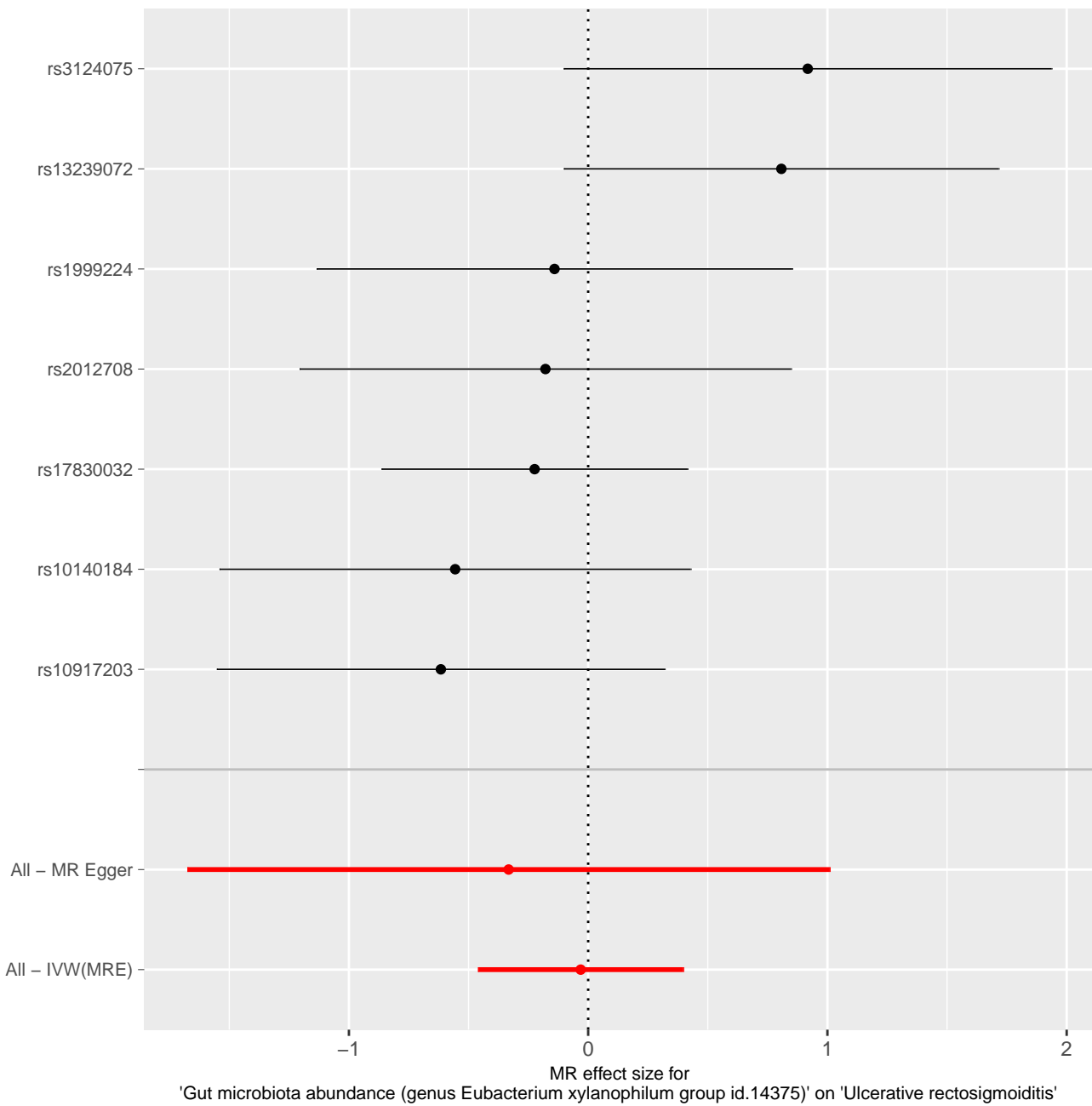


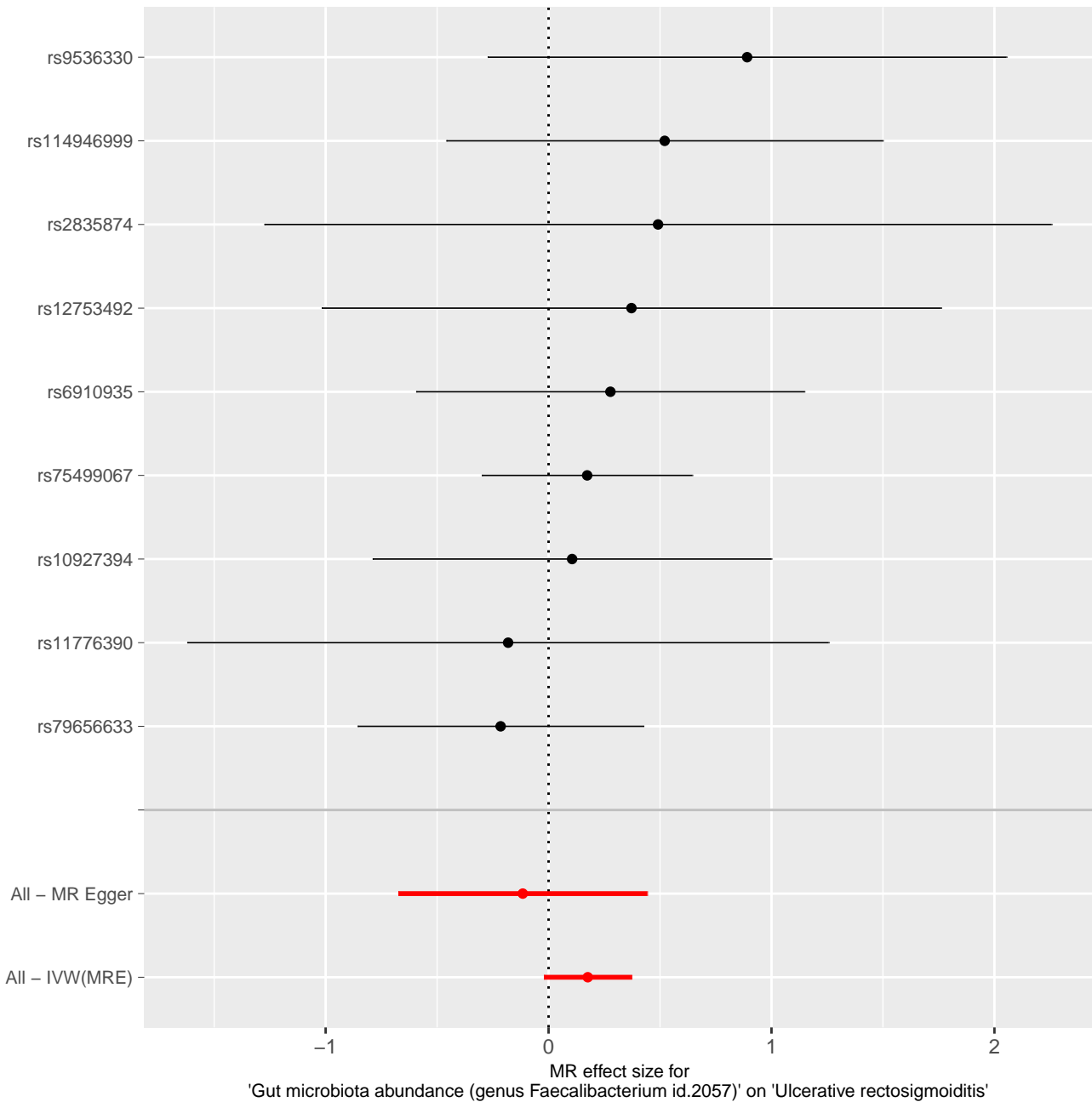


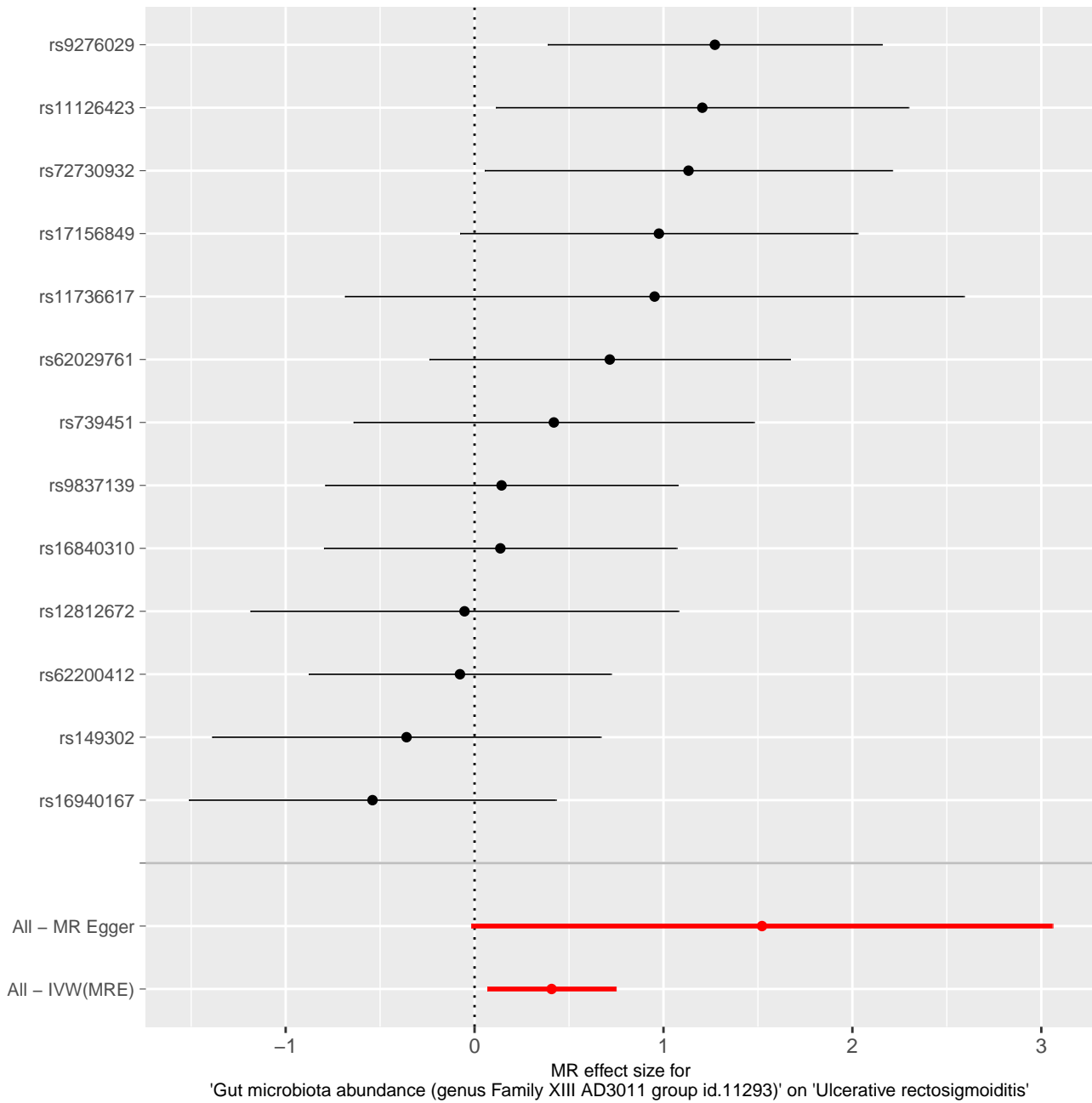


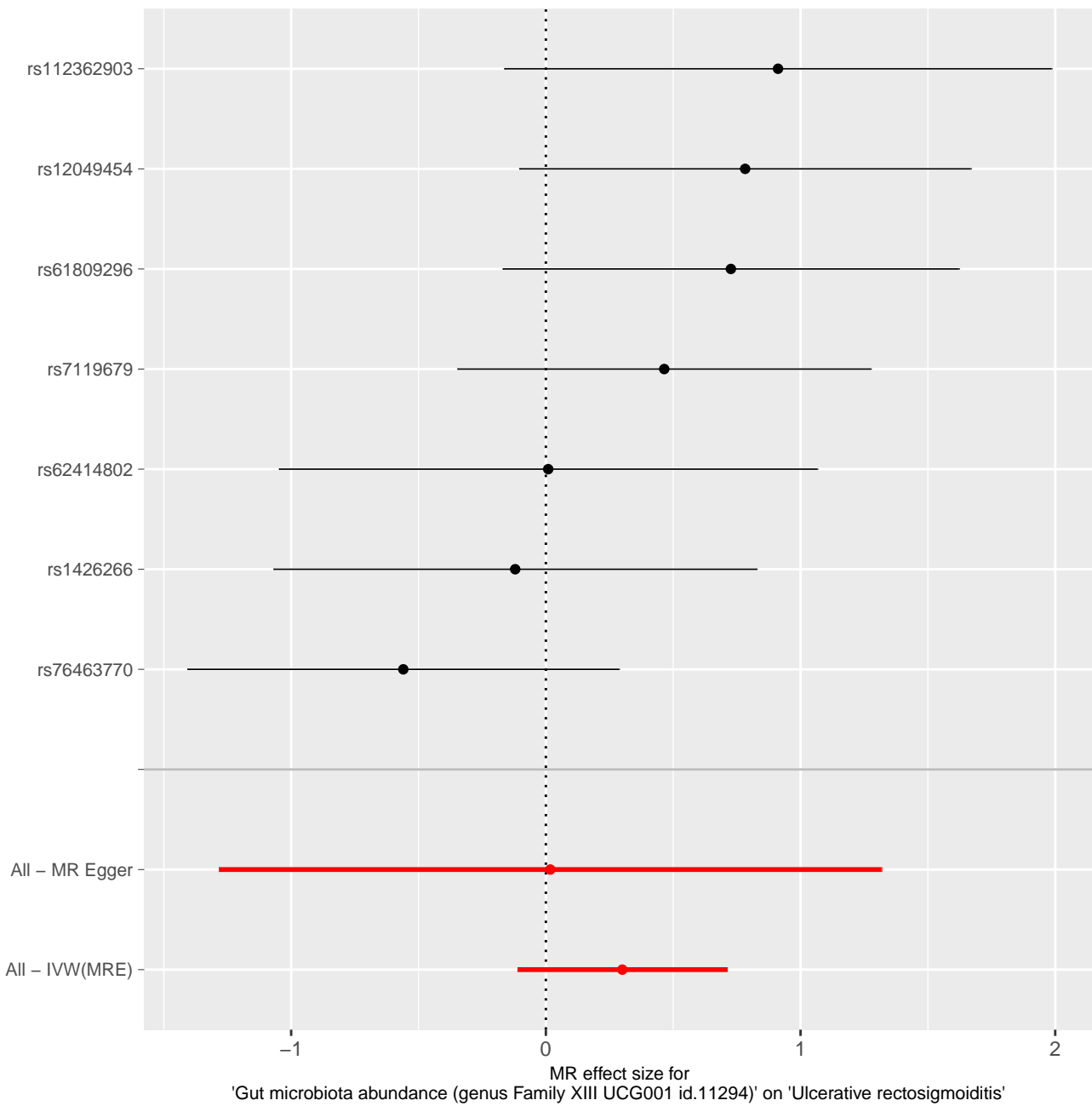


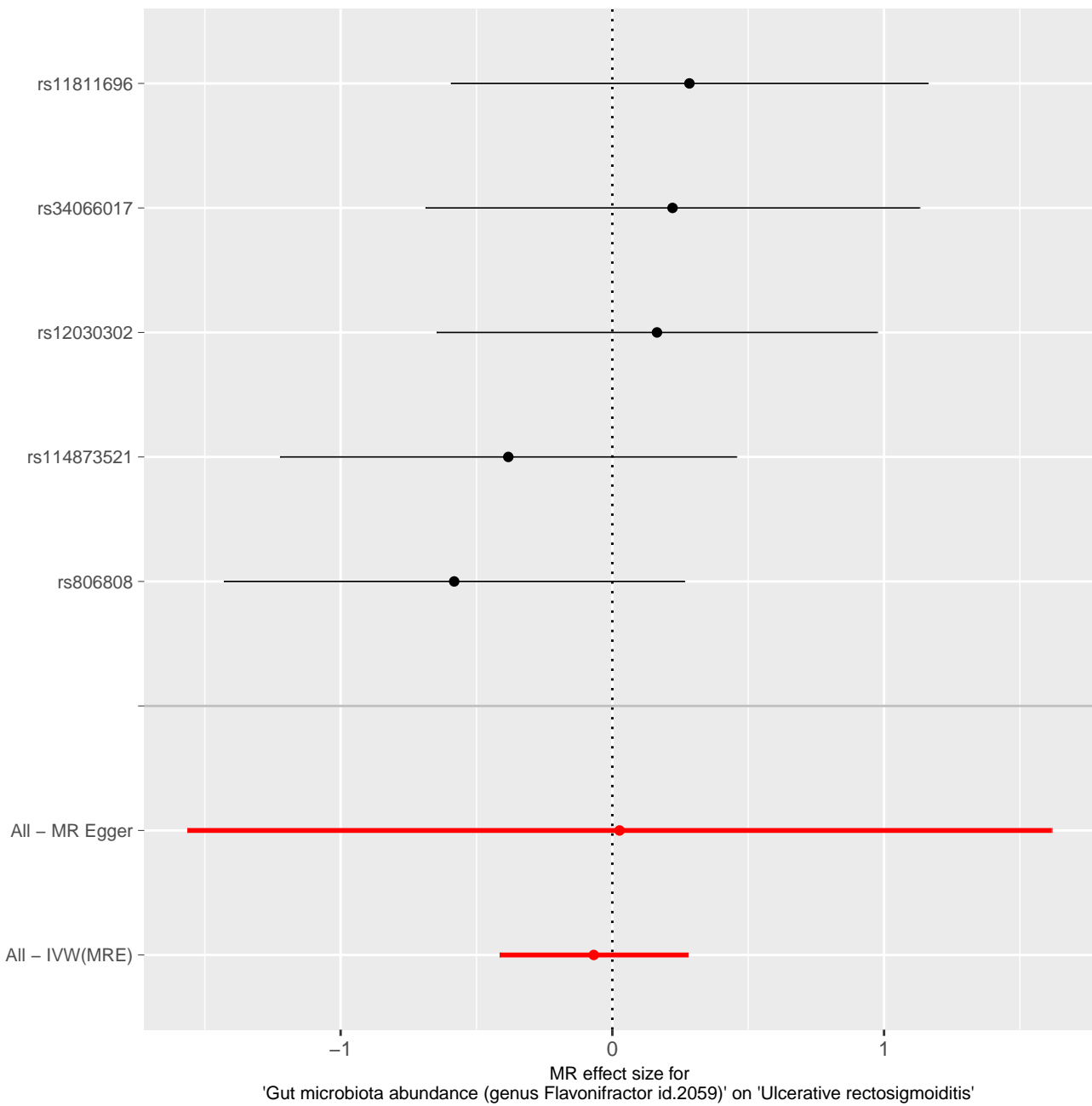




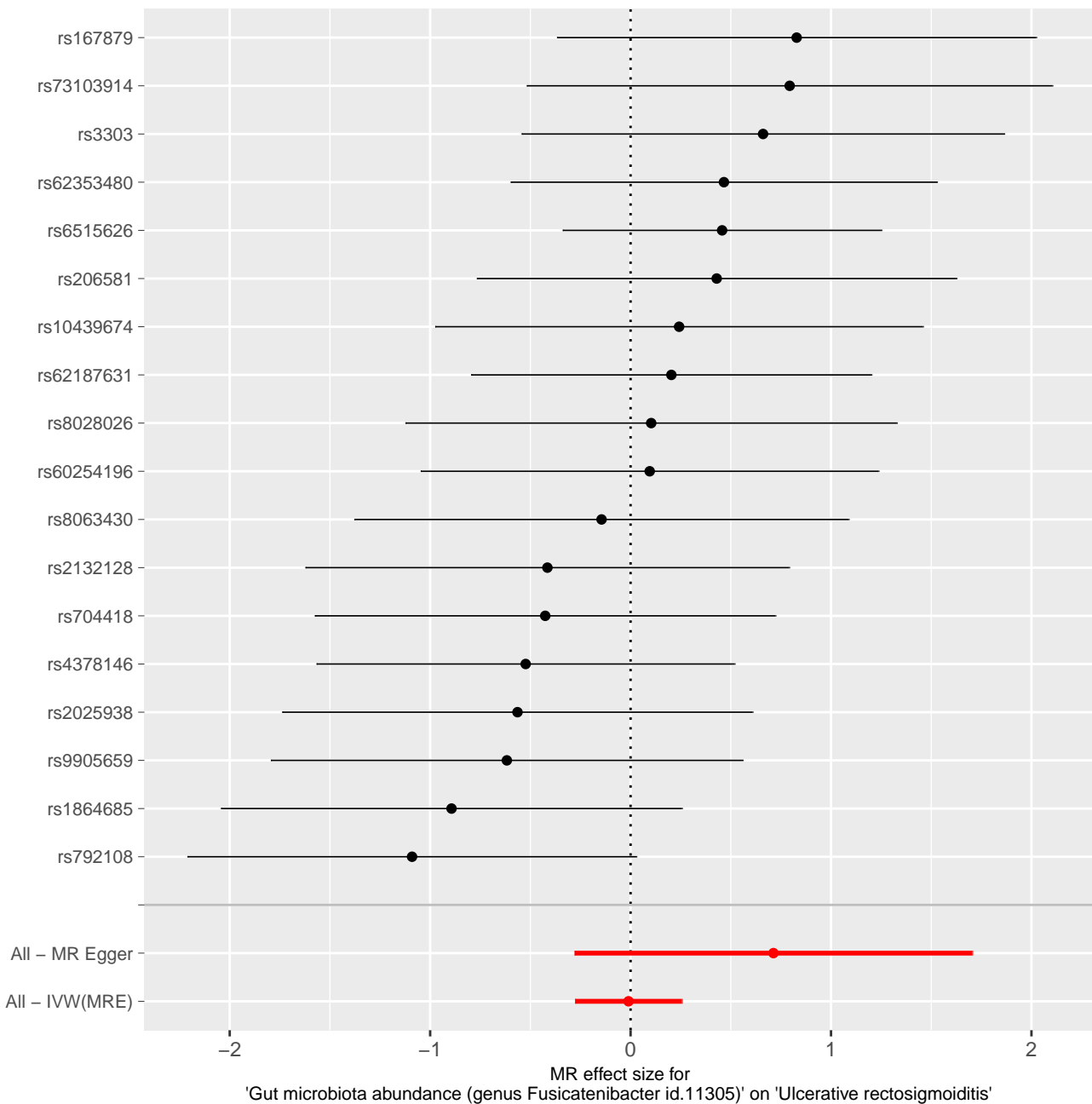


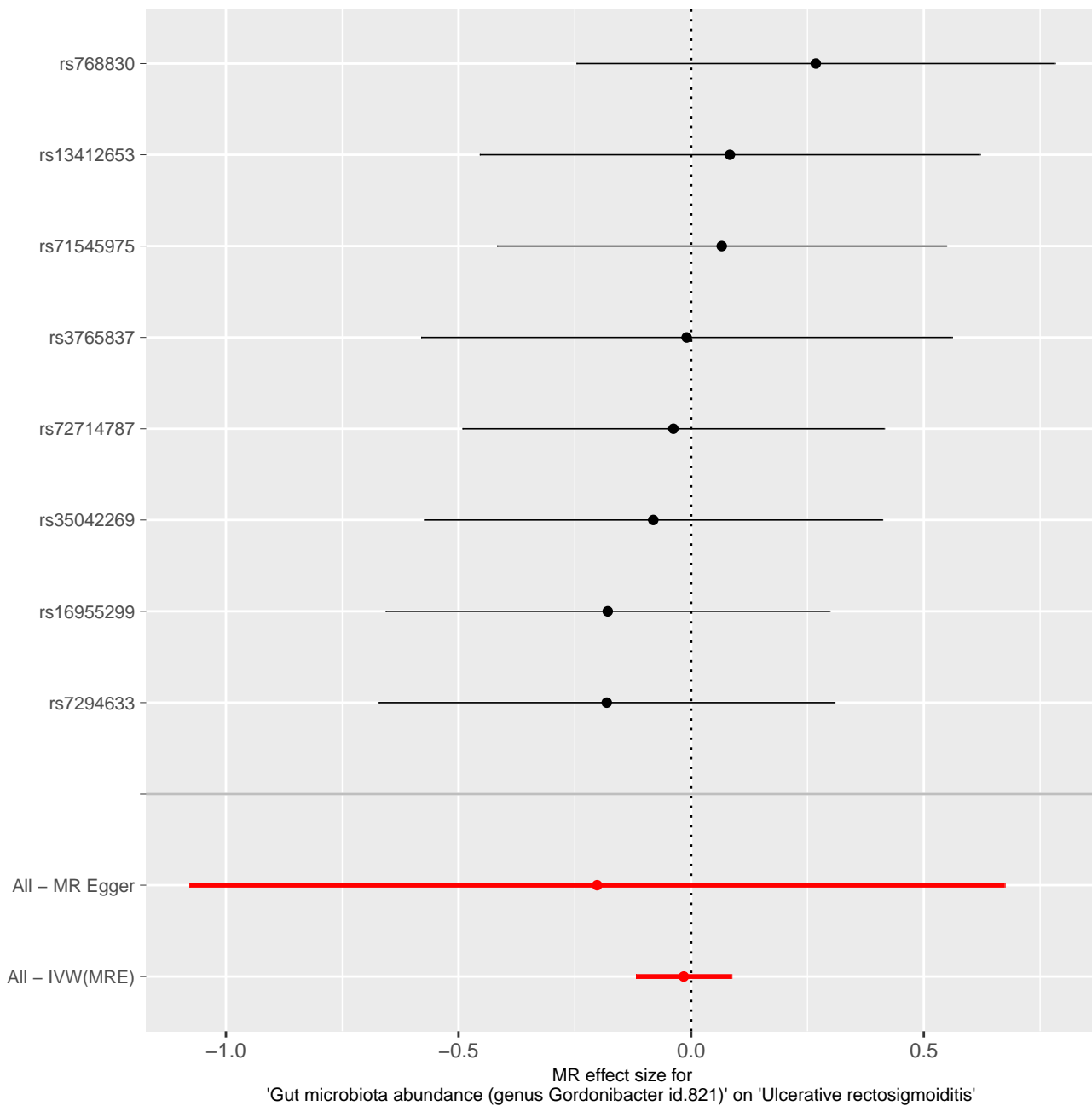


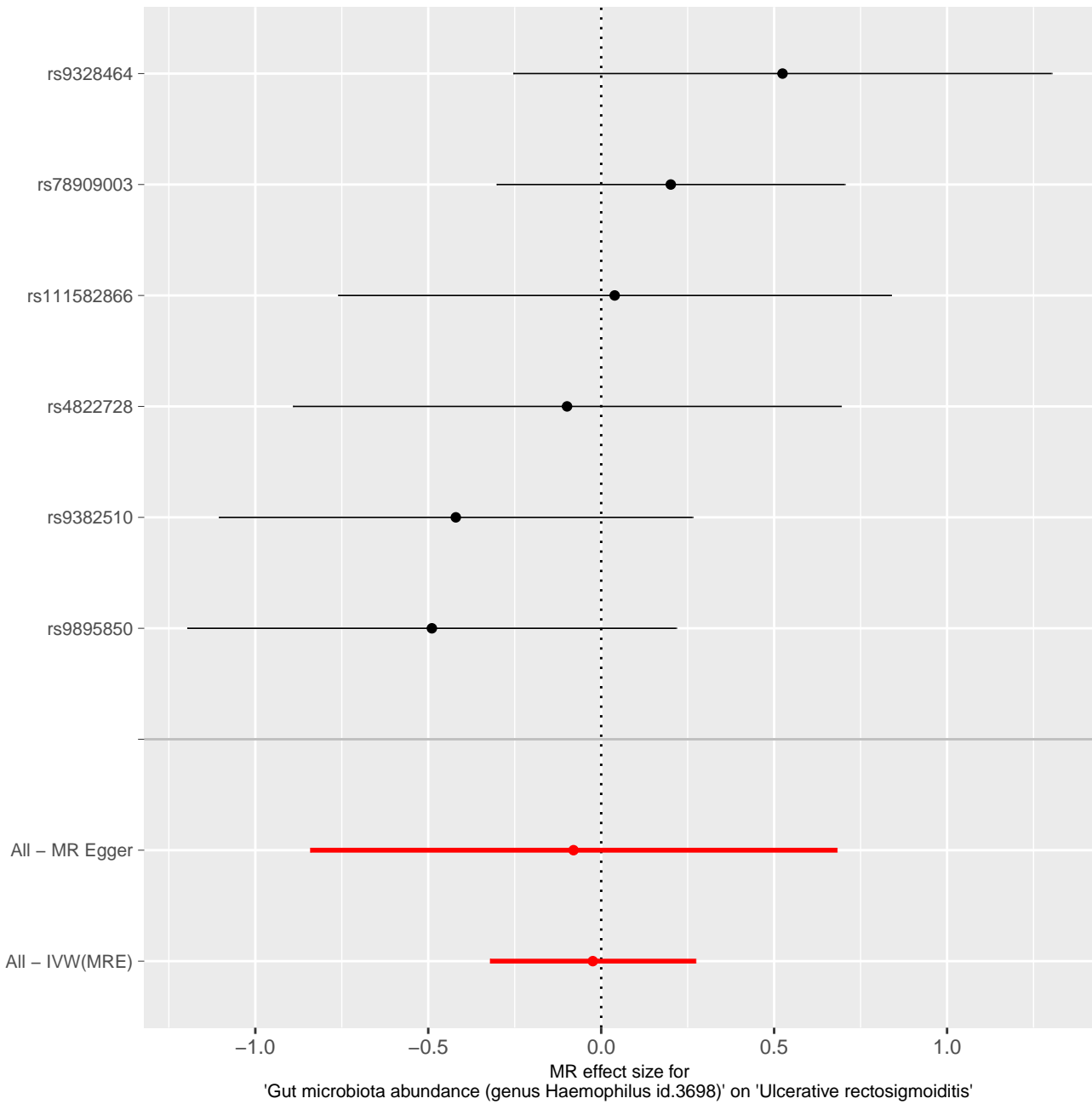


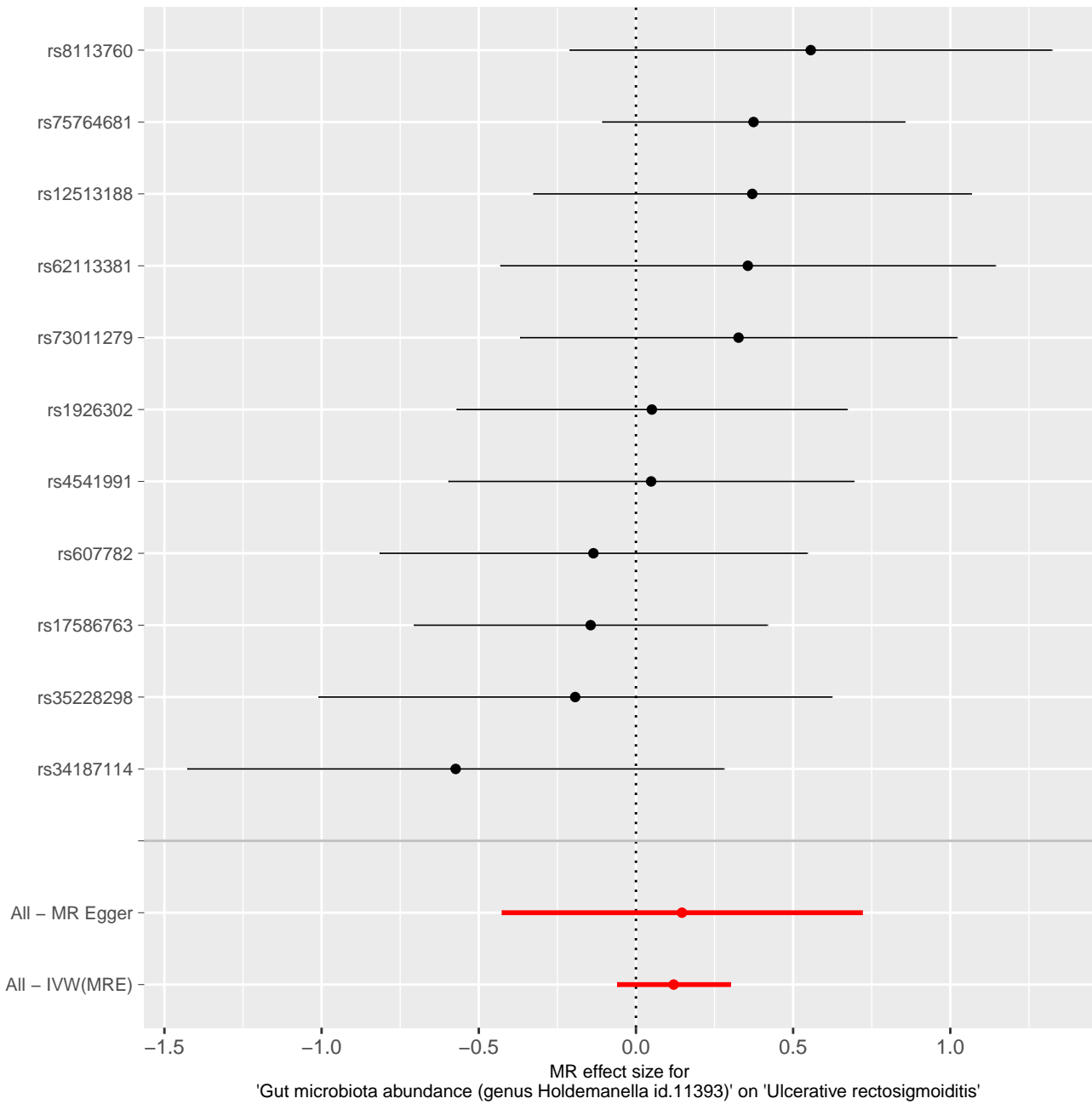


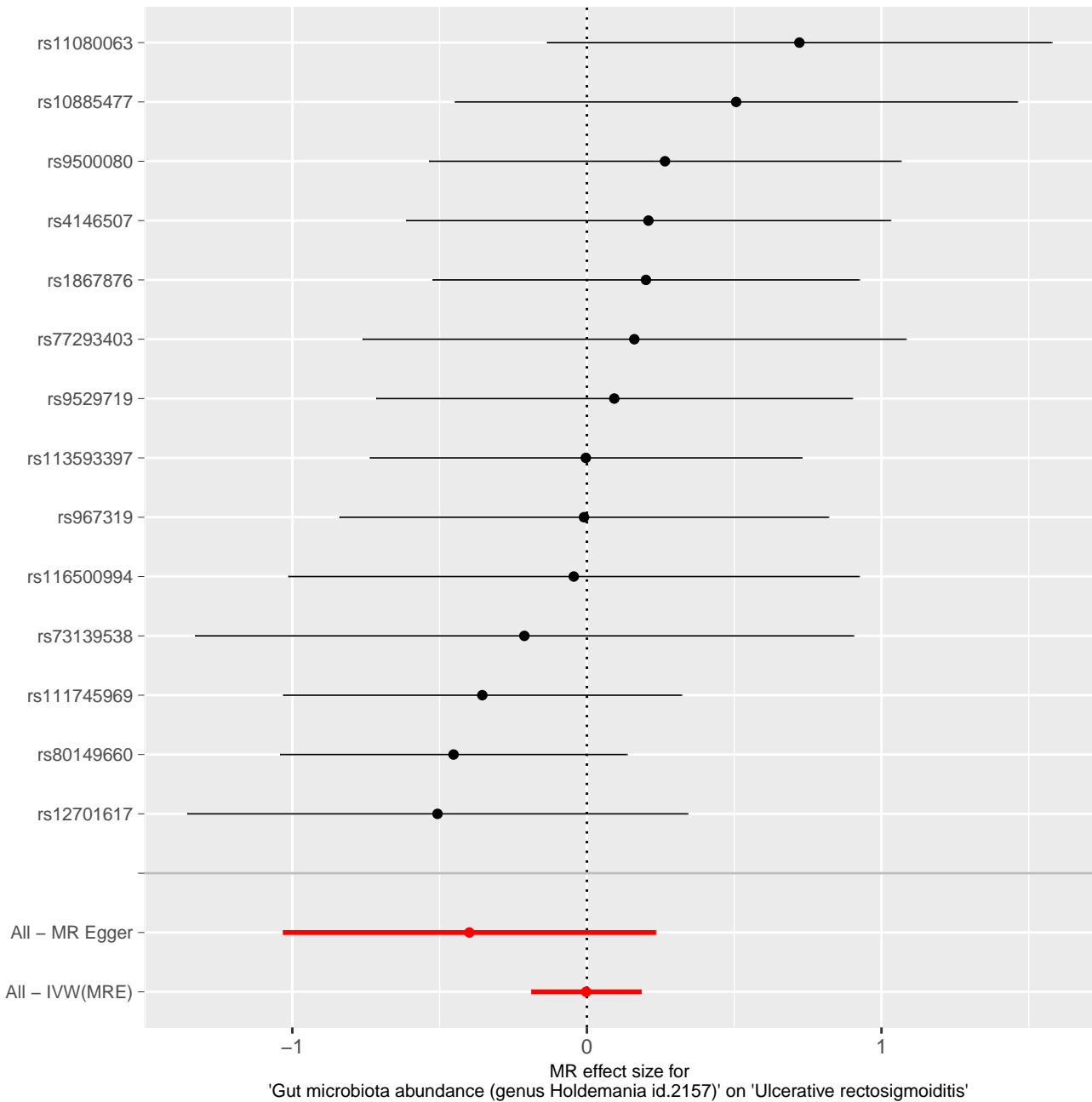
Batch 1585 : Gut microbiota abundance (genus Fusicatenibacter id.11305) on Ulcerative rectosigmoiditis

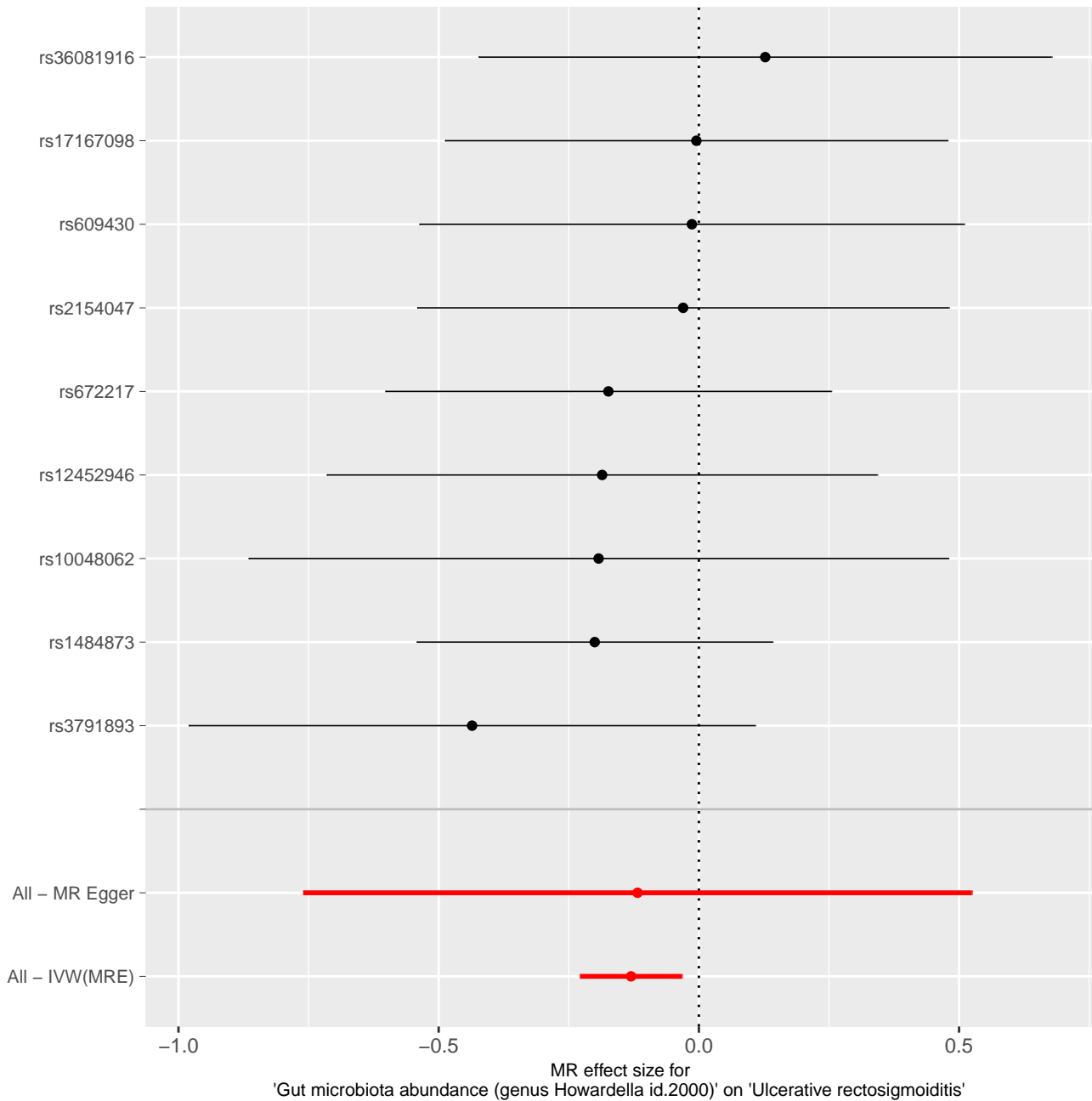


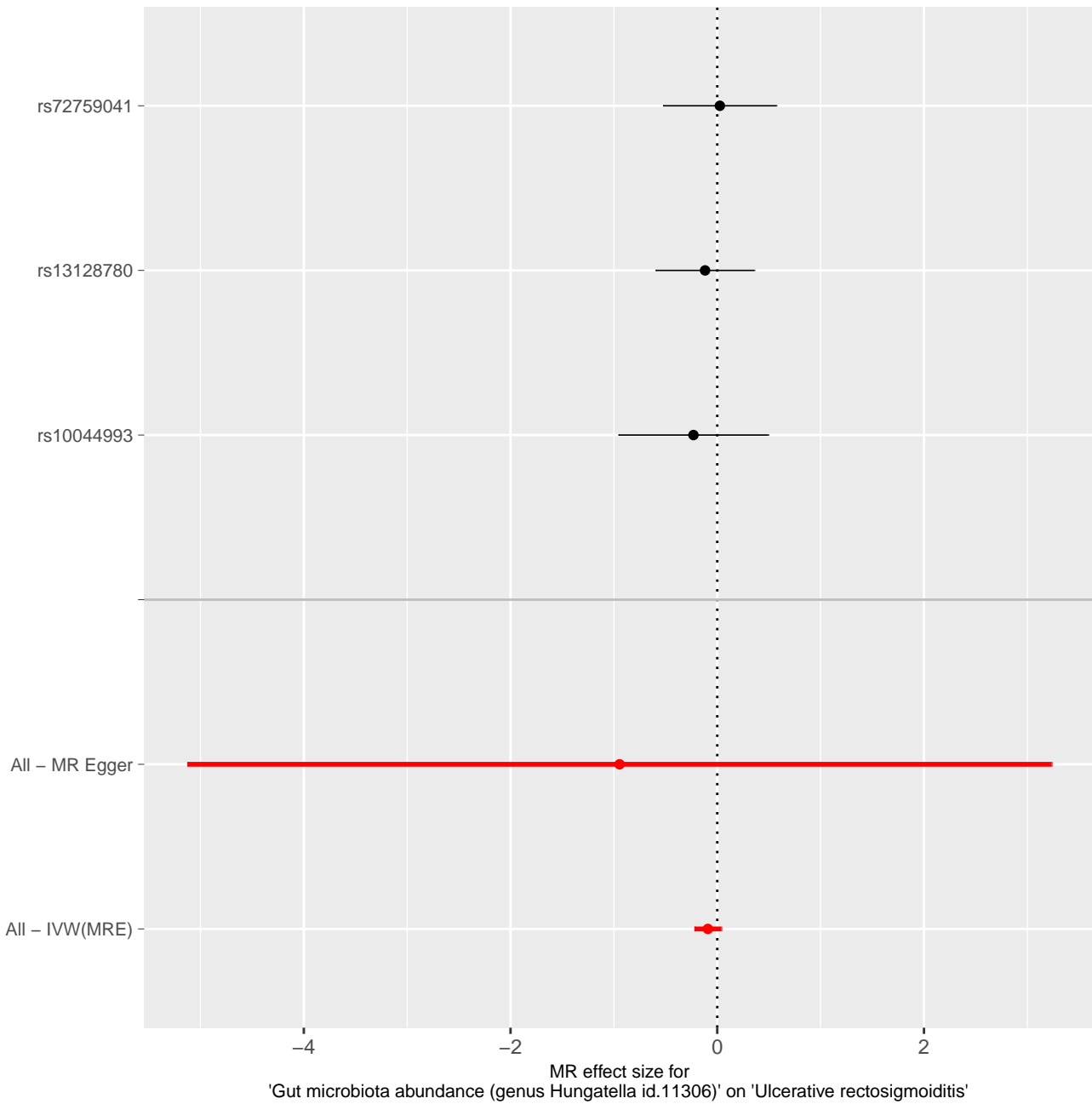




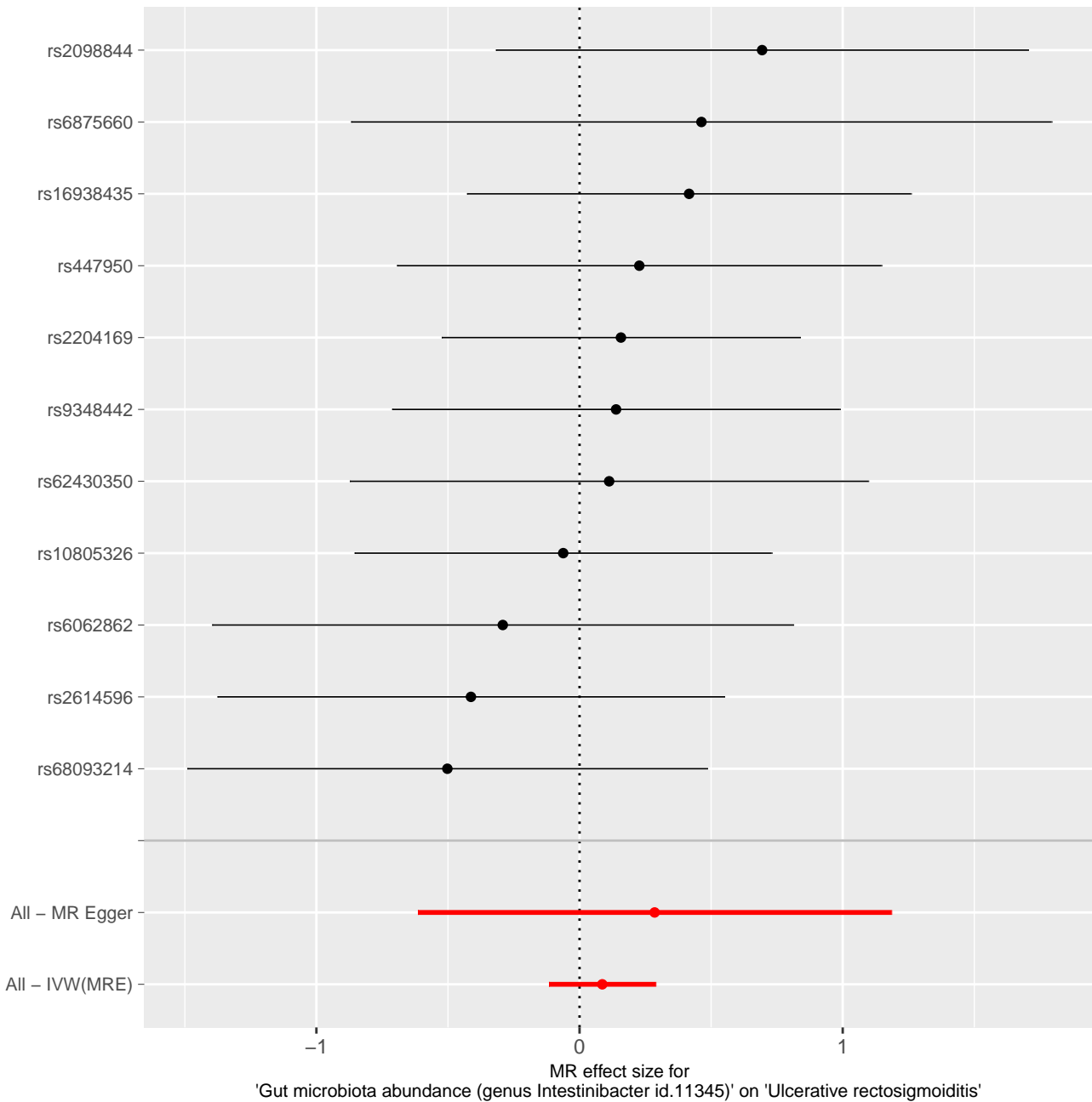


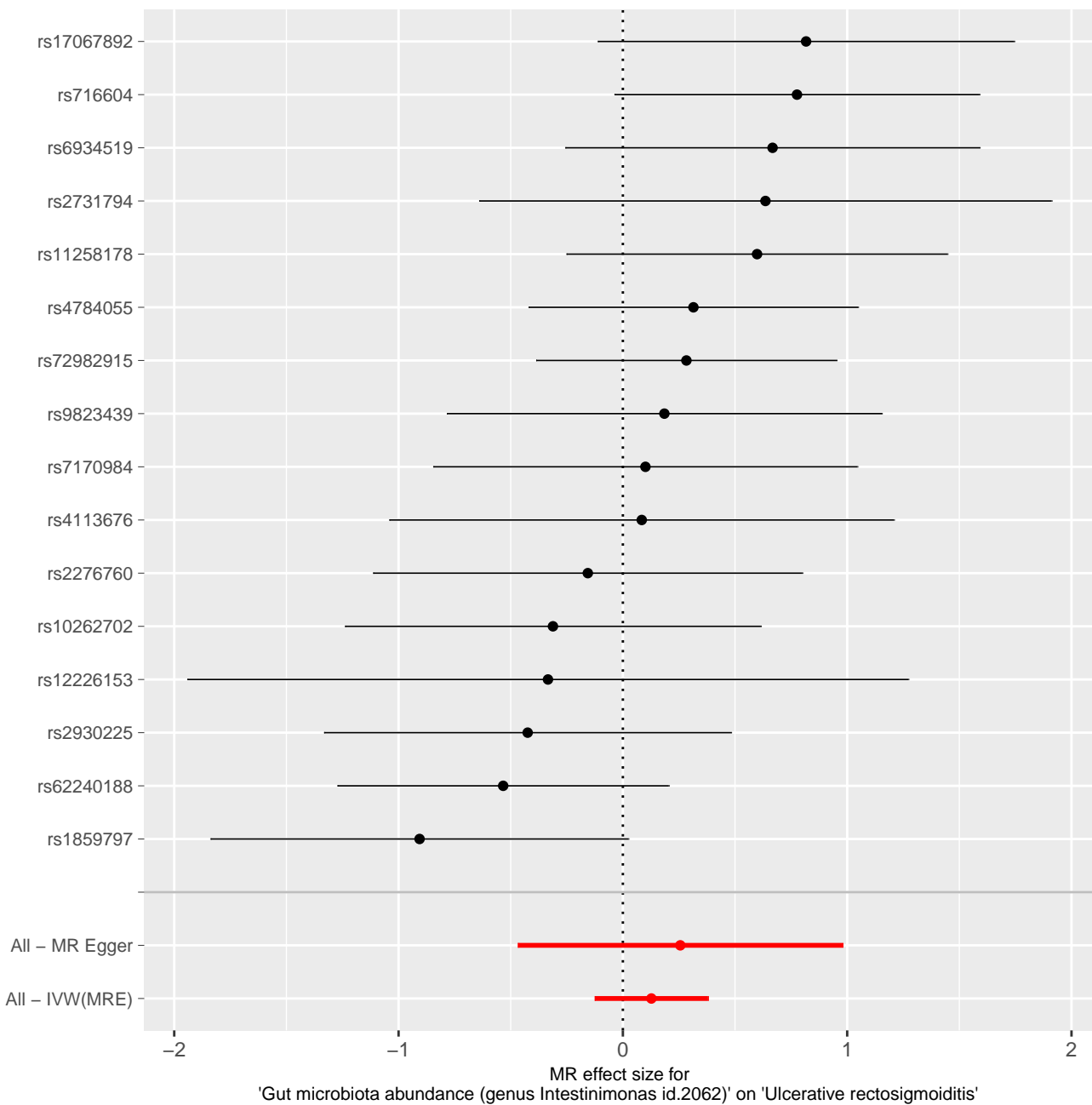


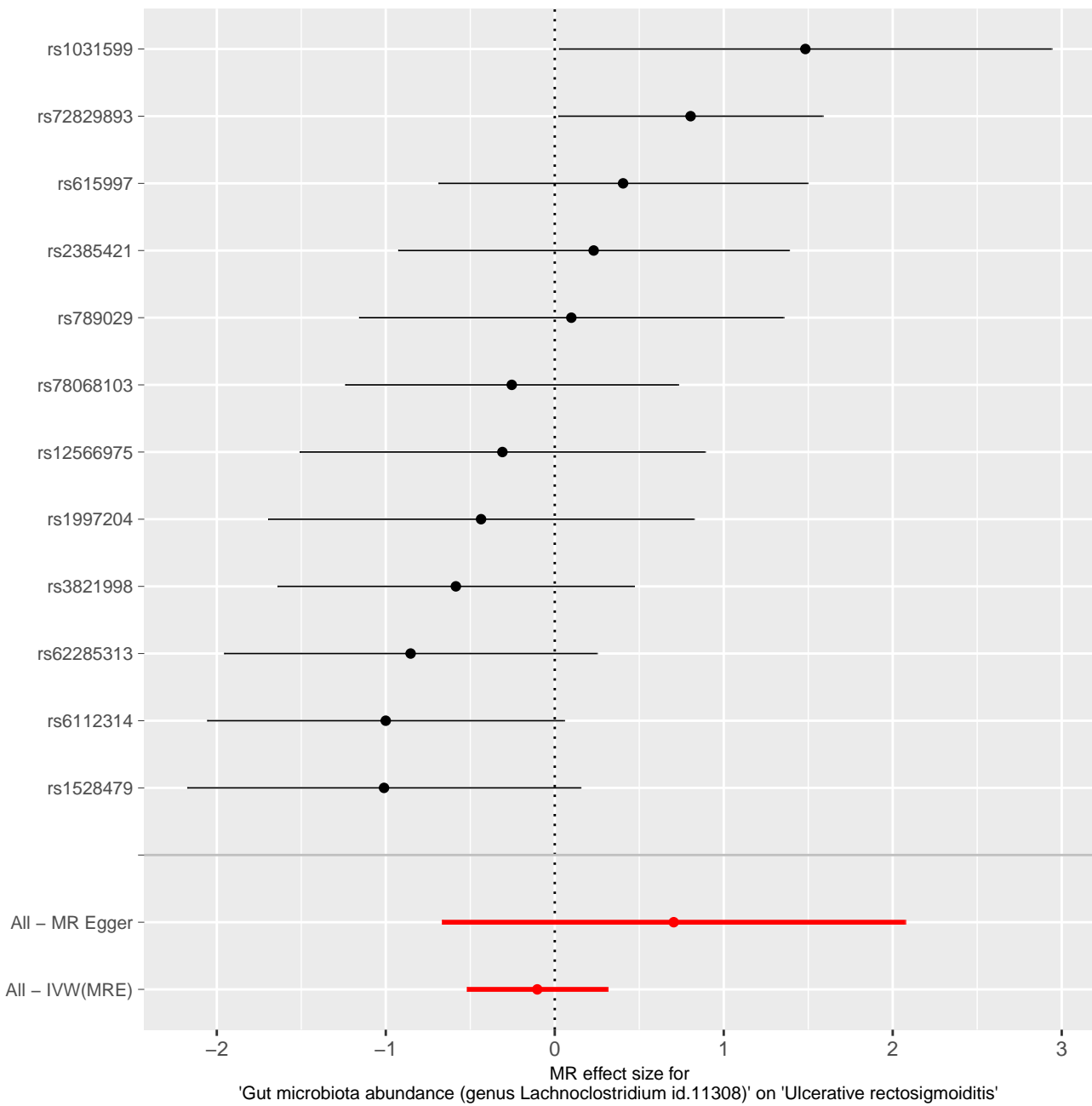


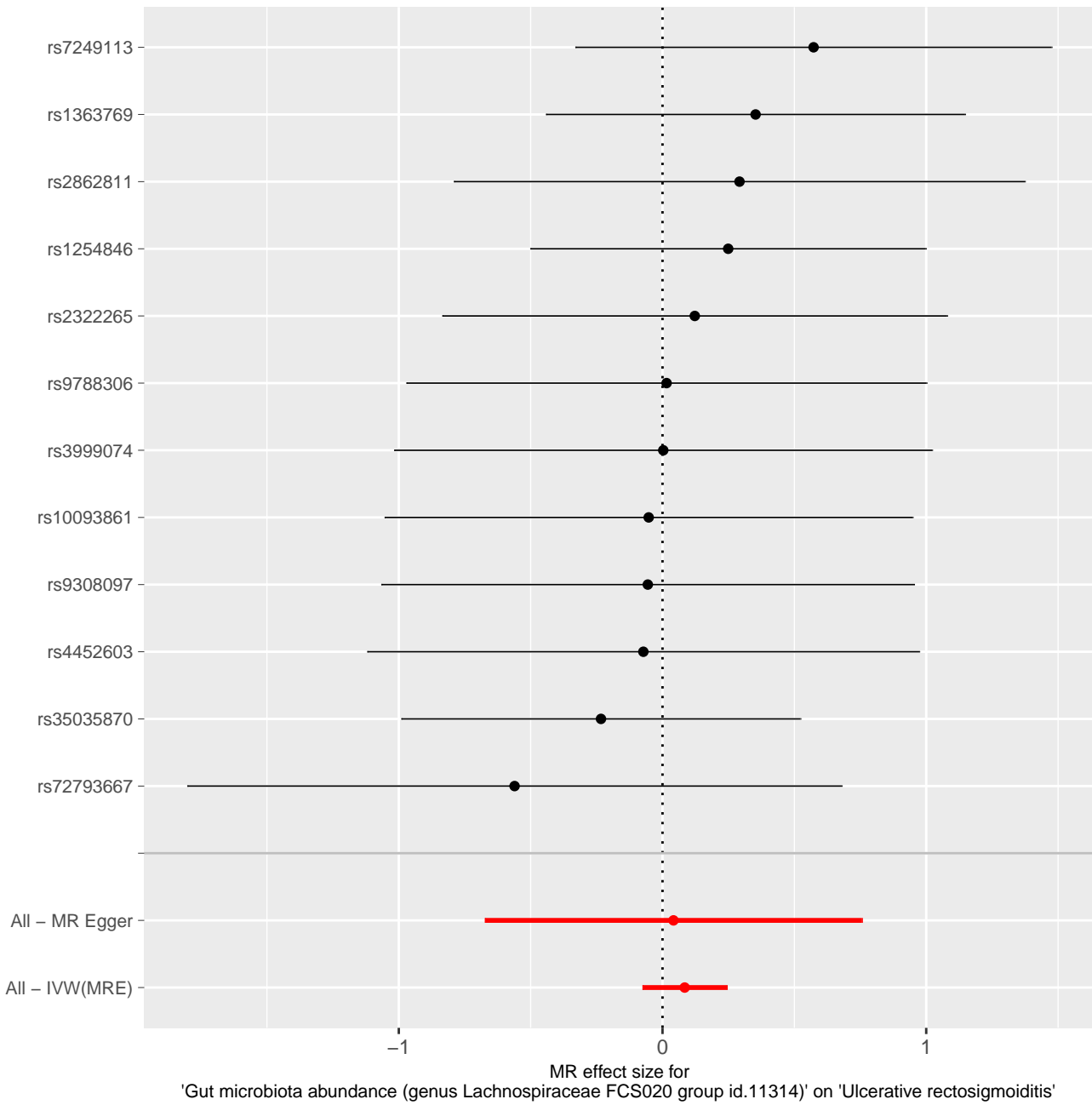


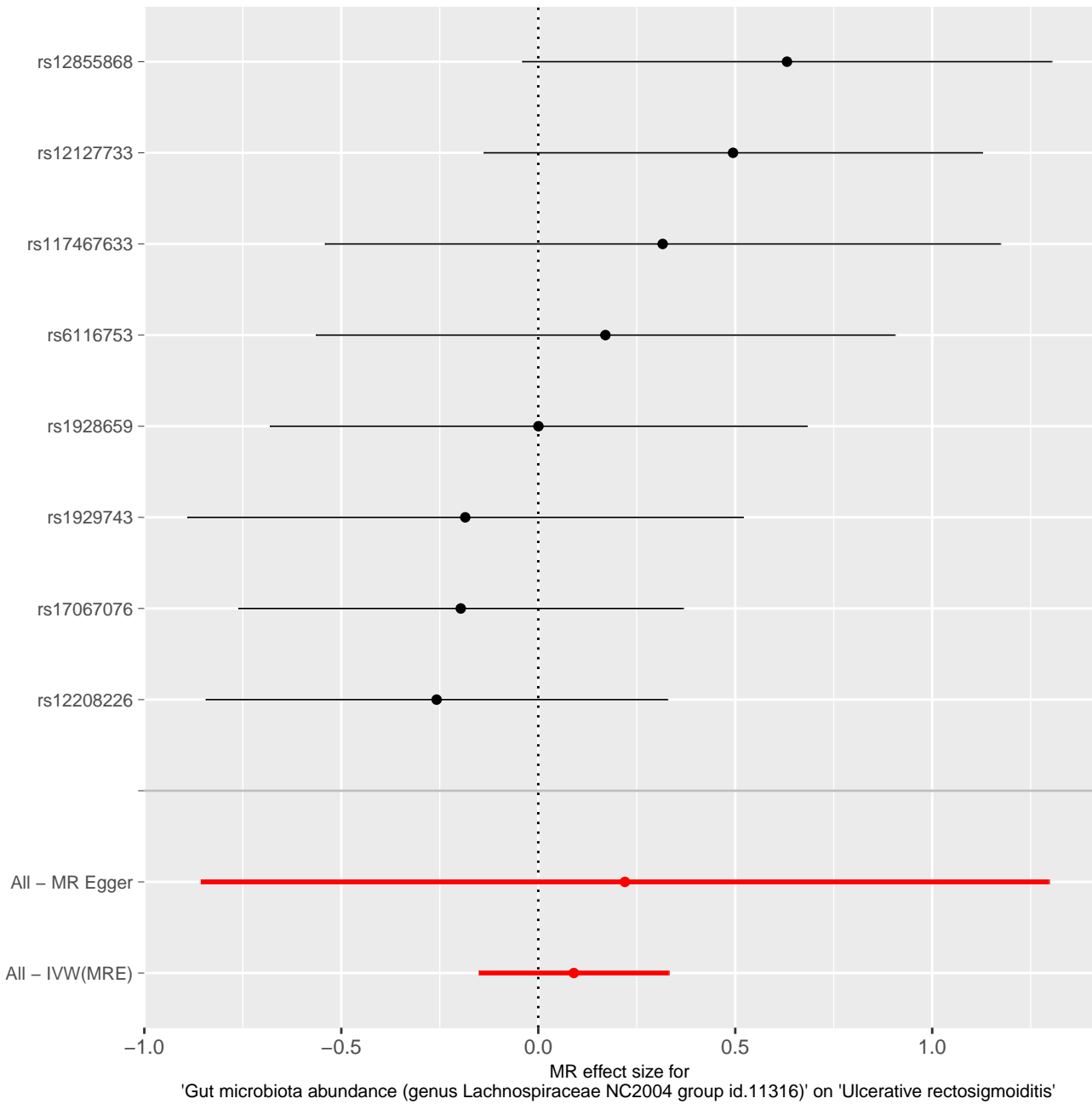
Batch 1592 : Gut microbiota abundance (genus Intestinibacter id.11345) on Ulcerative rectosigmoiditis

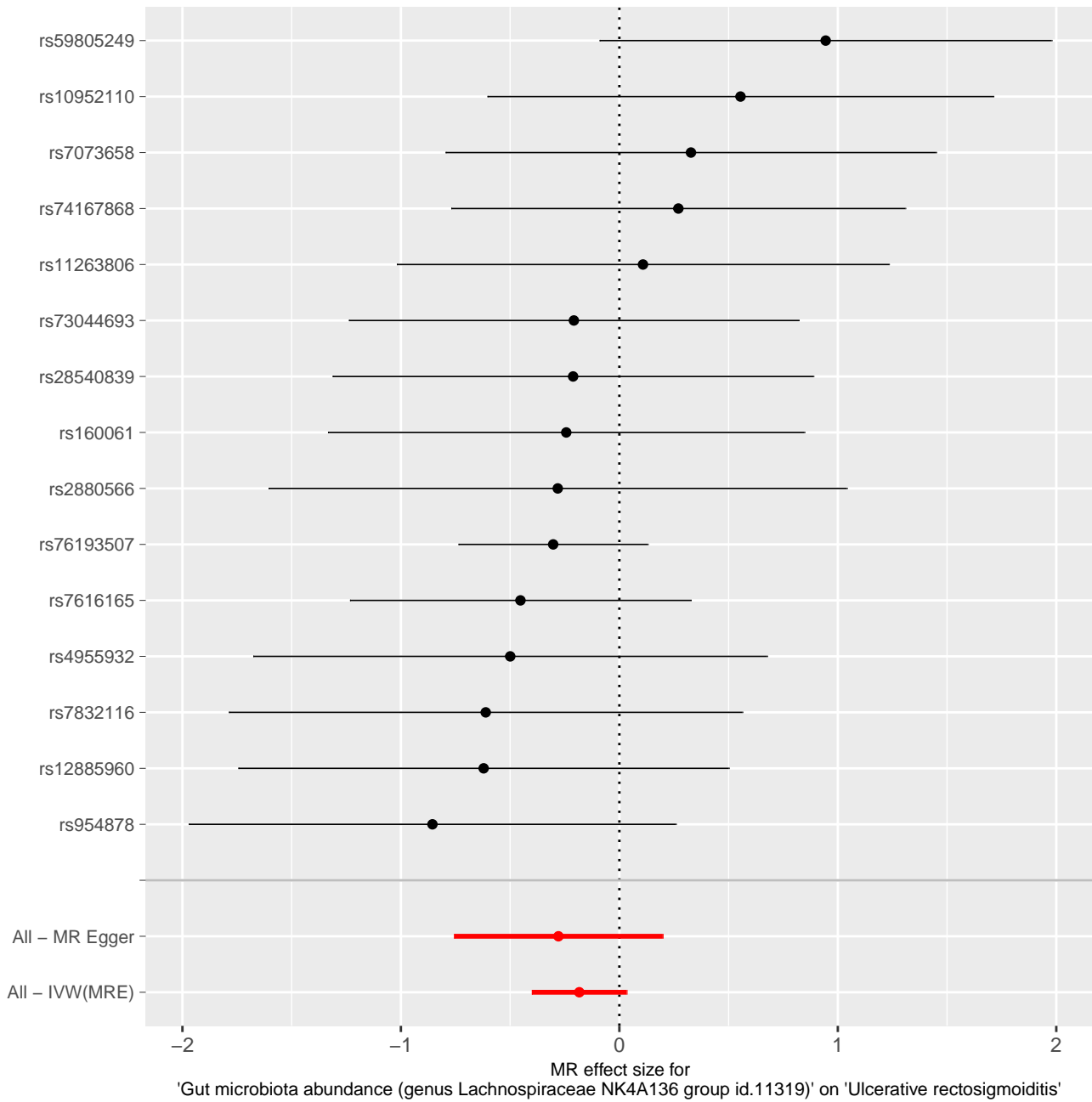


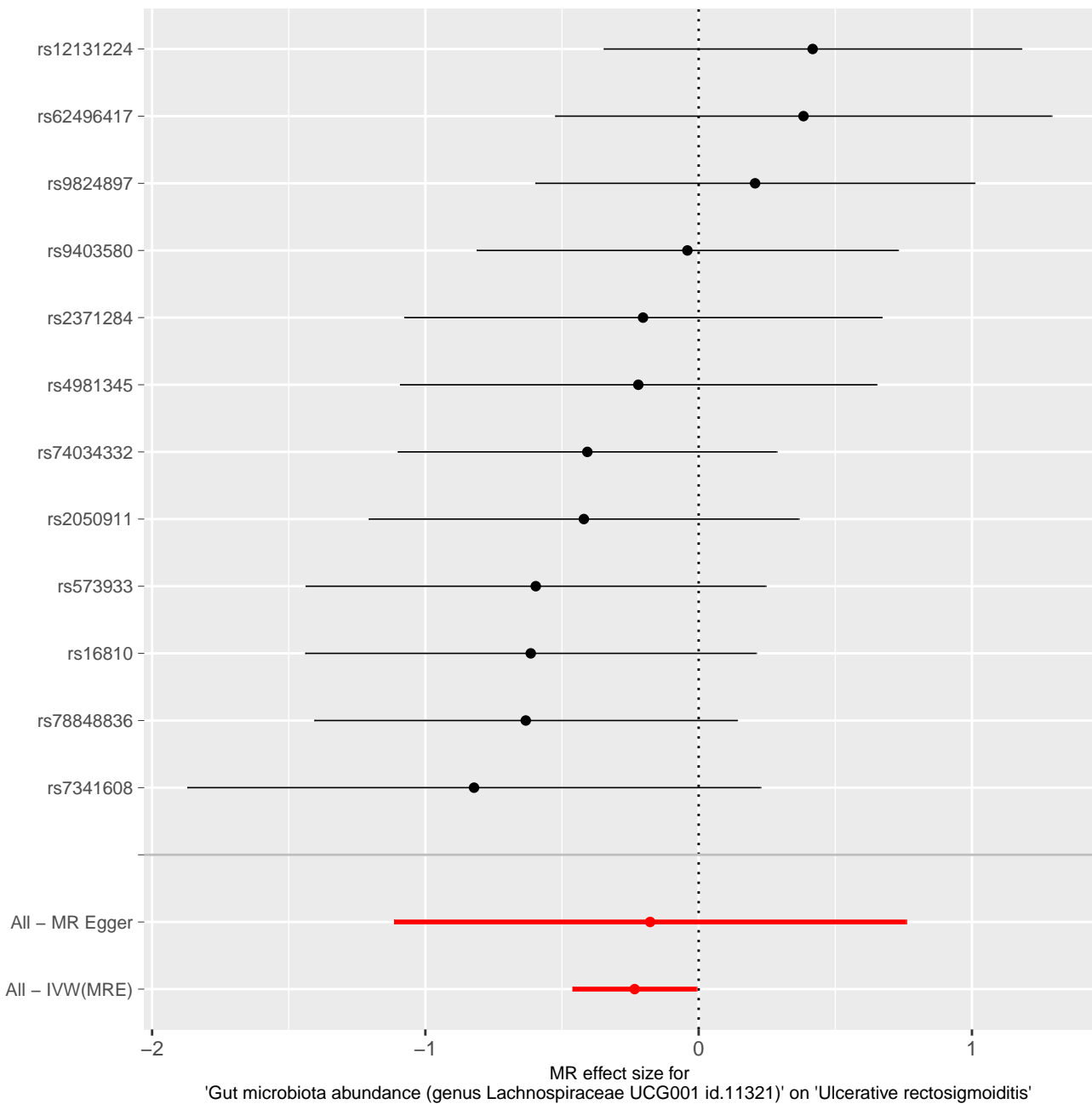


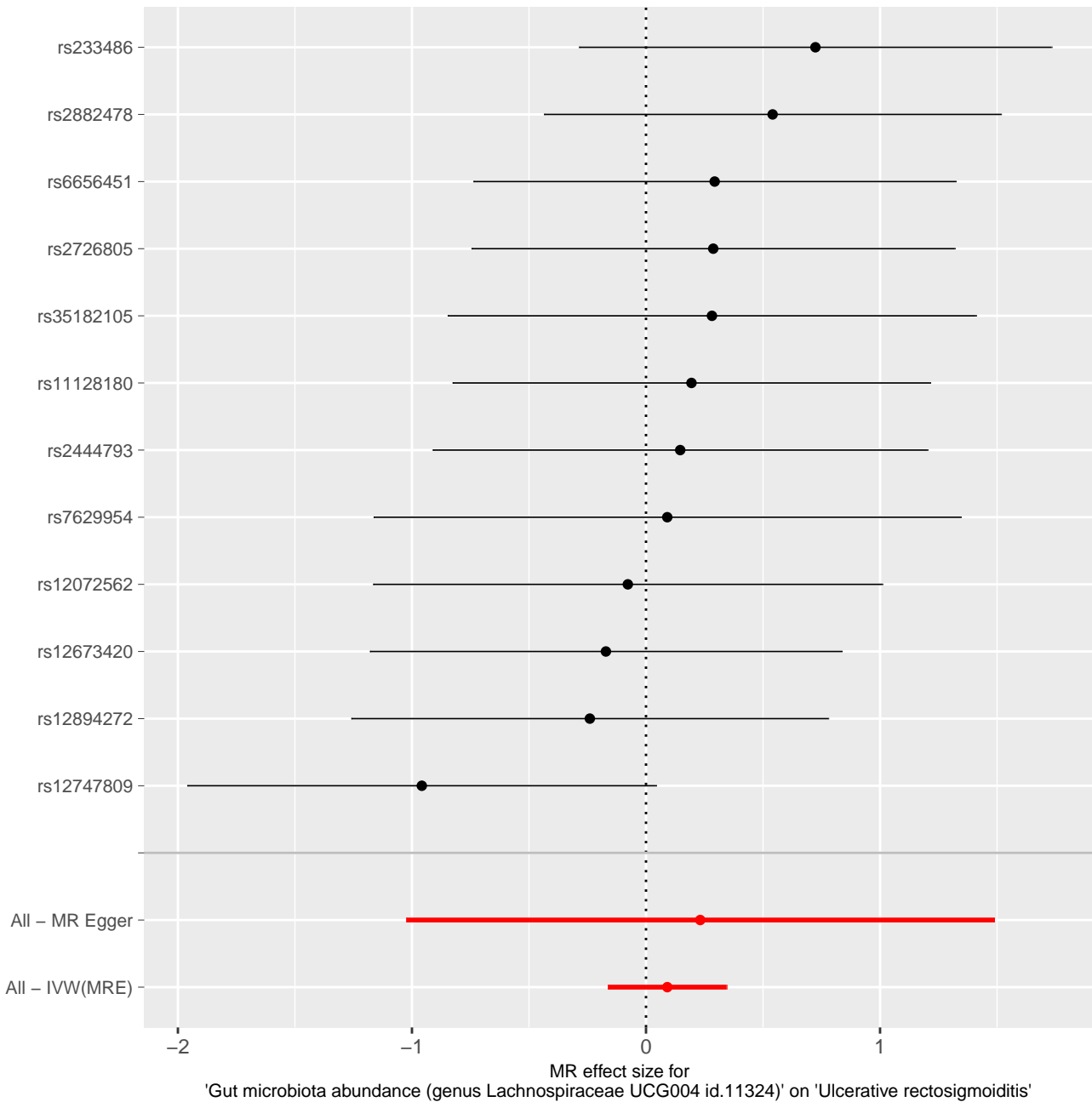


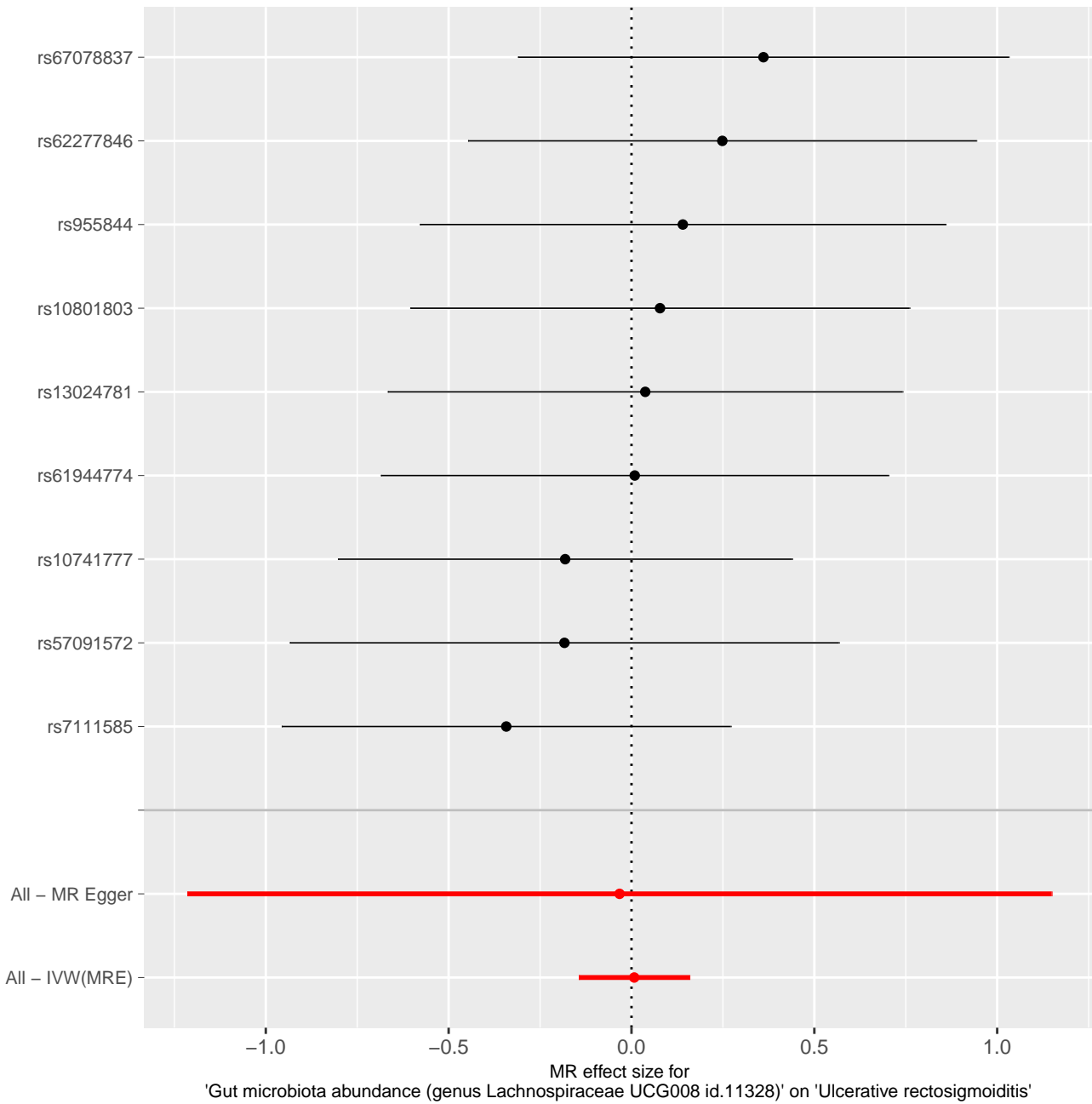


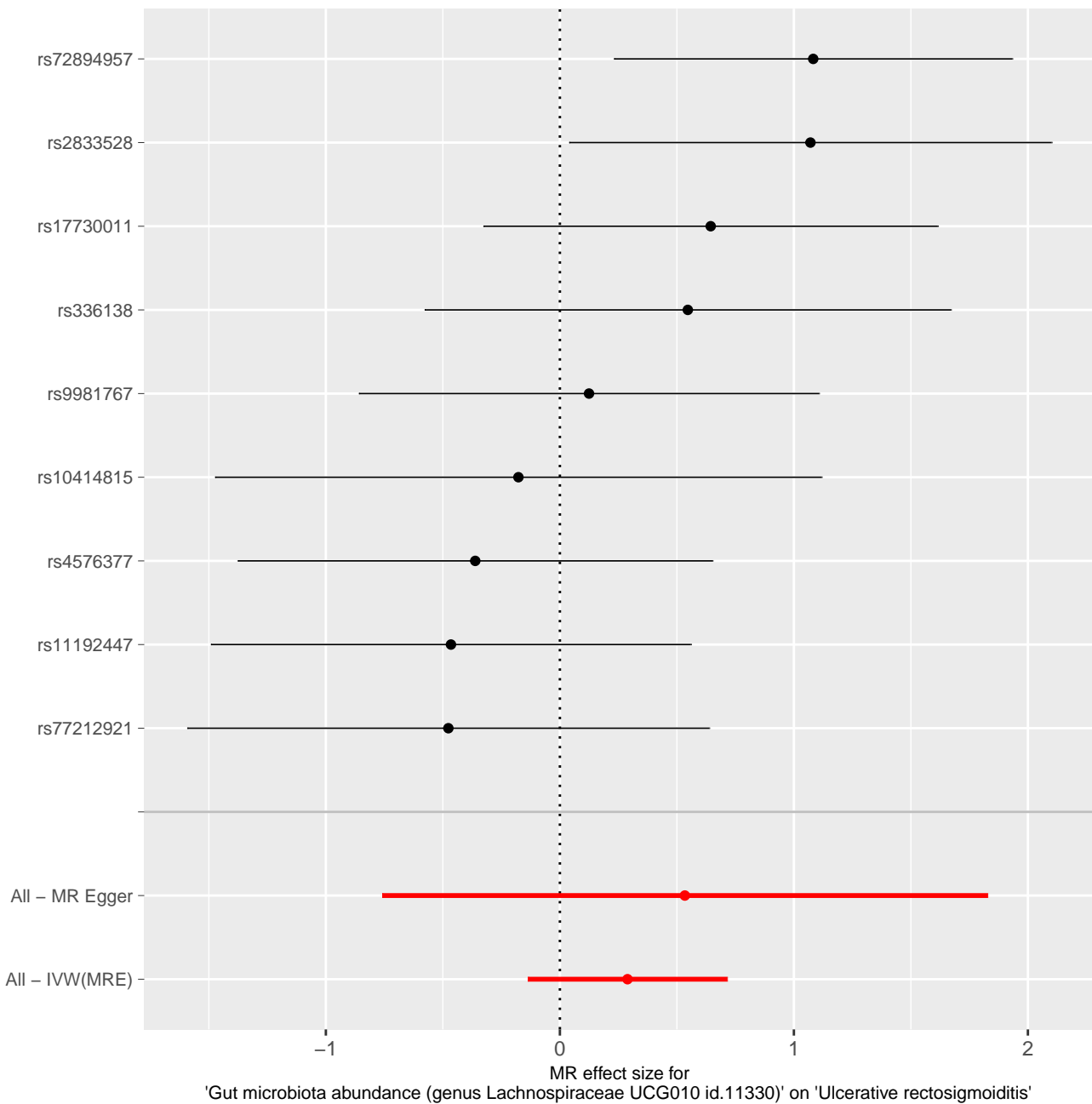


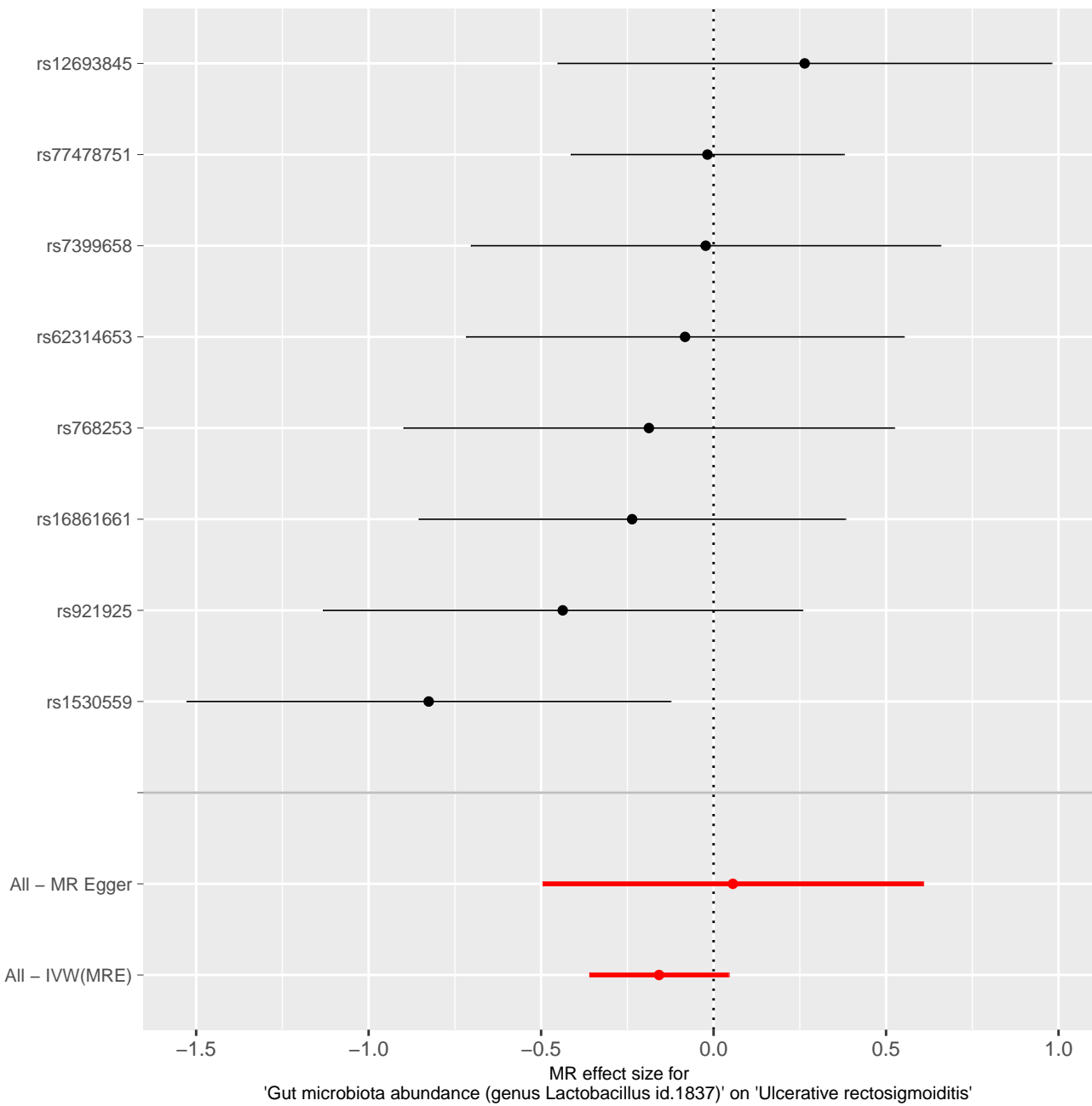


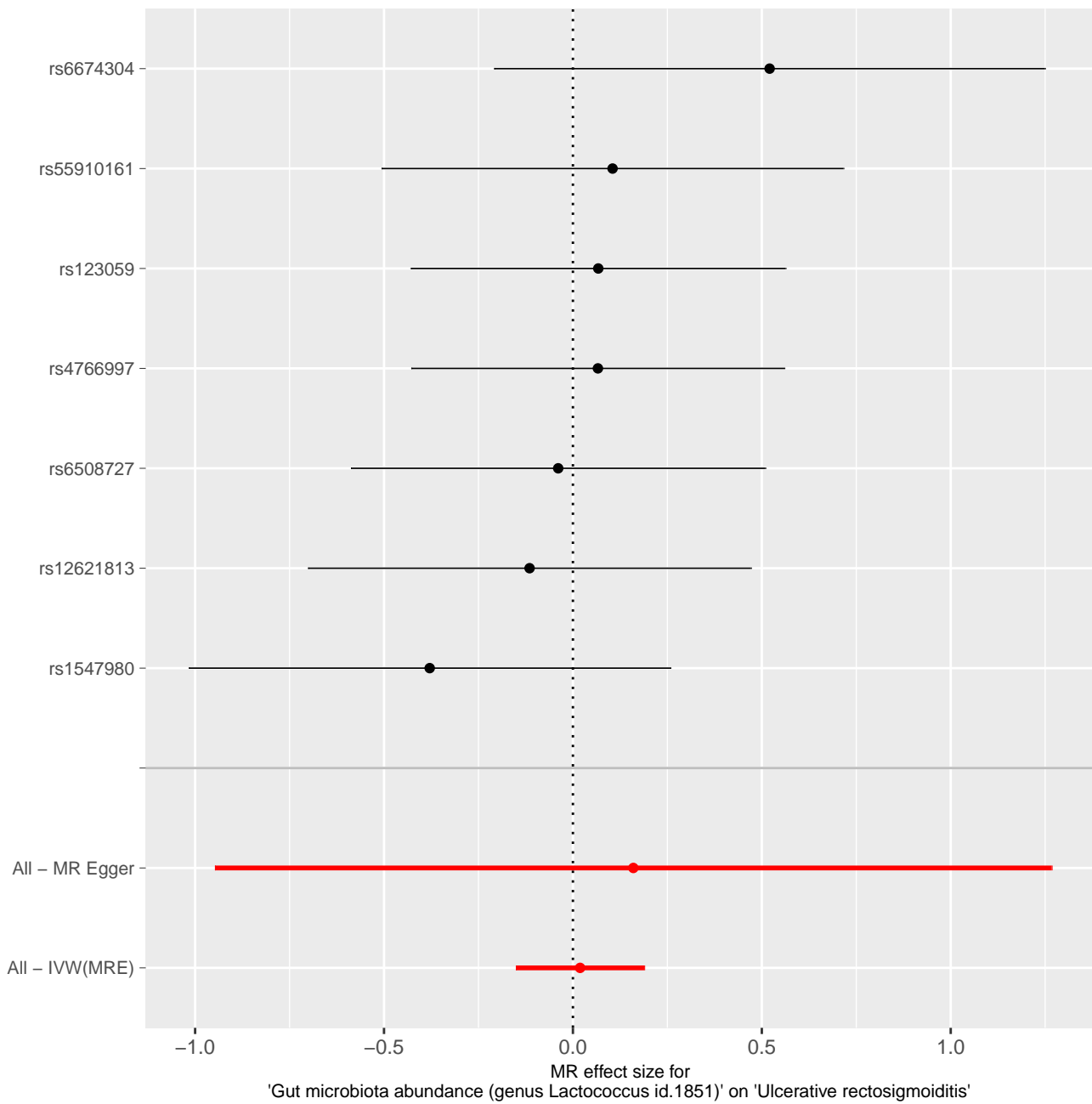


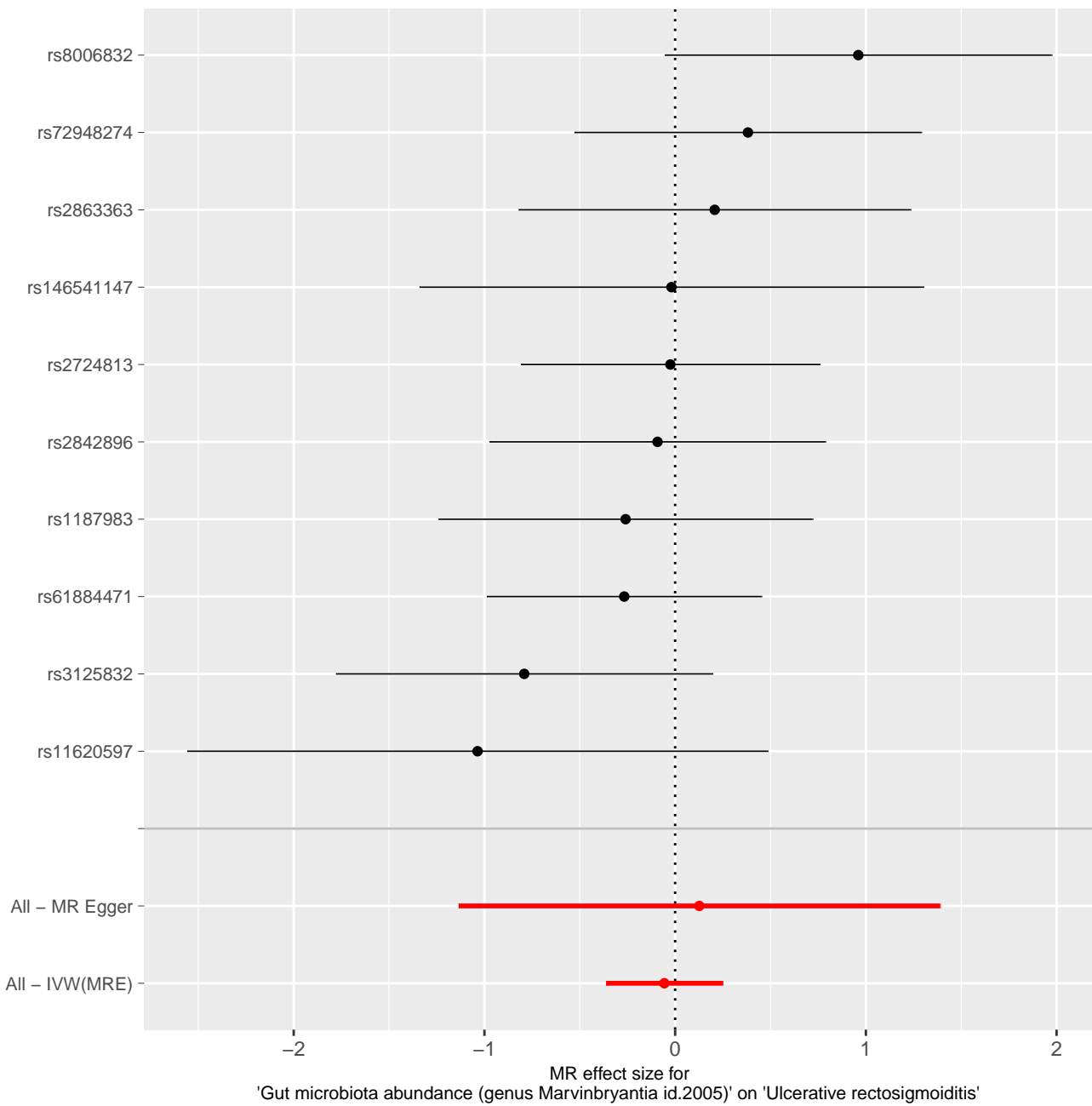


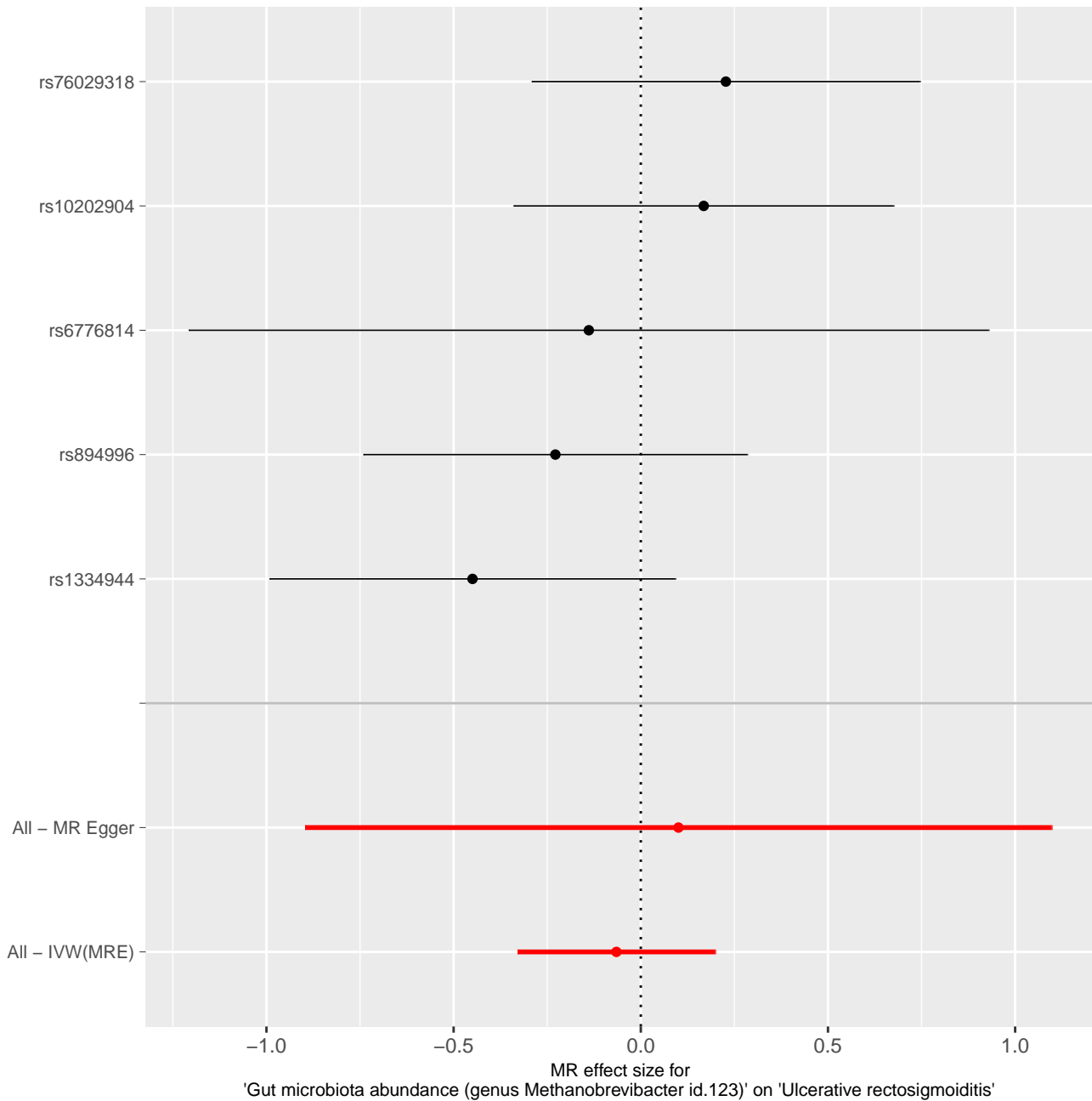


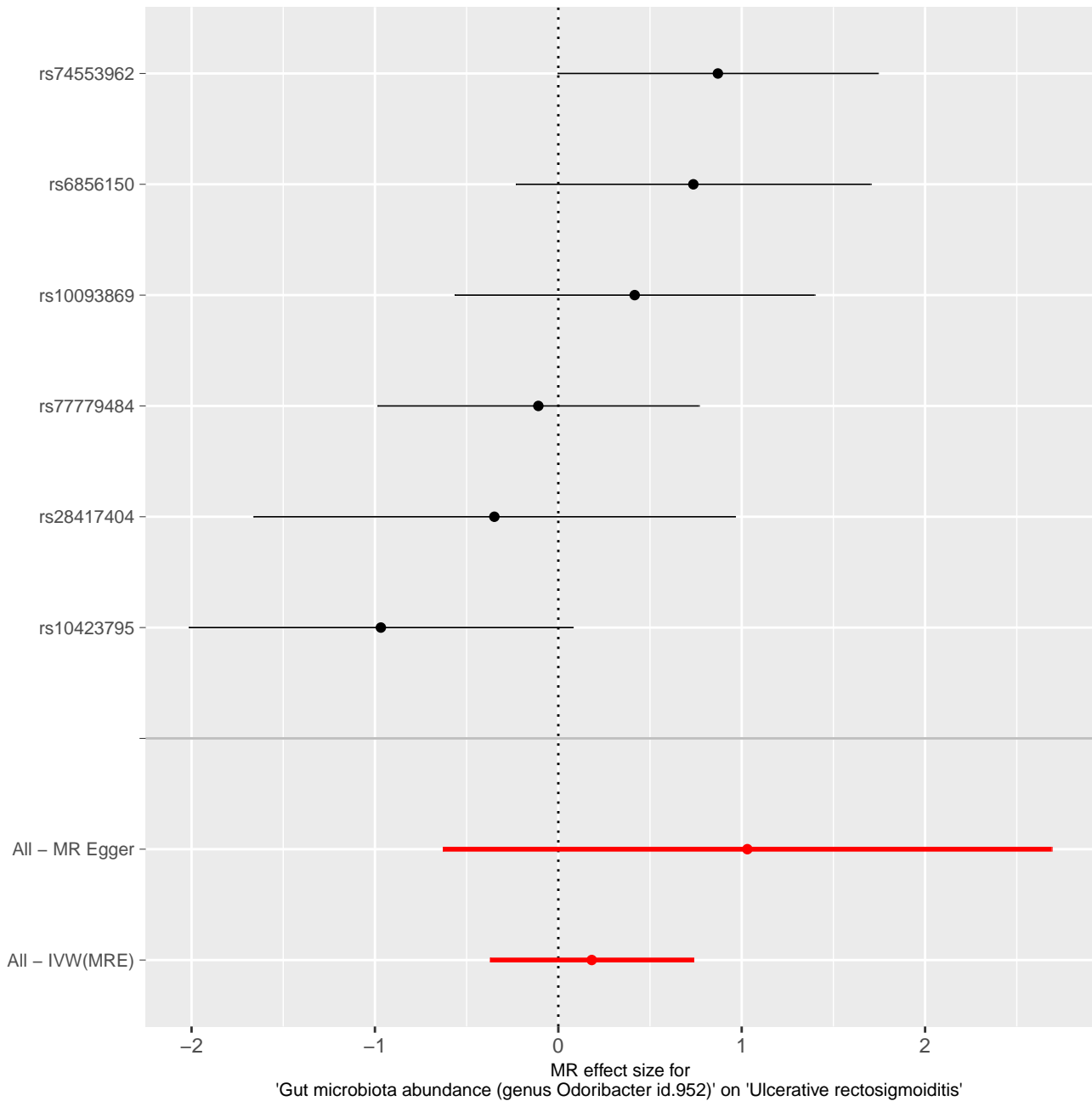


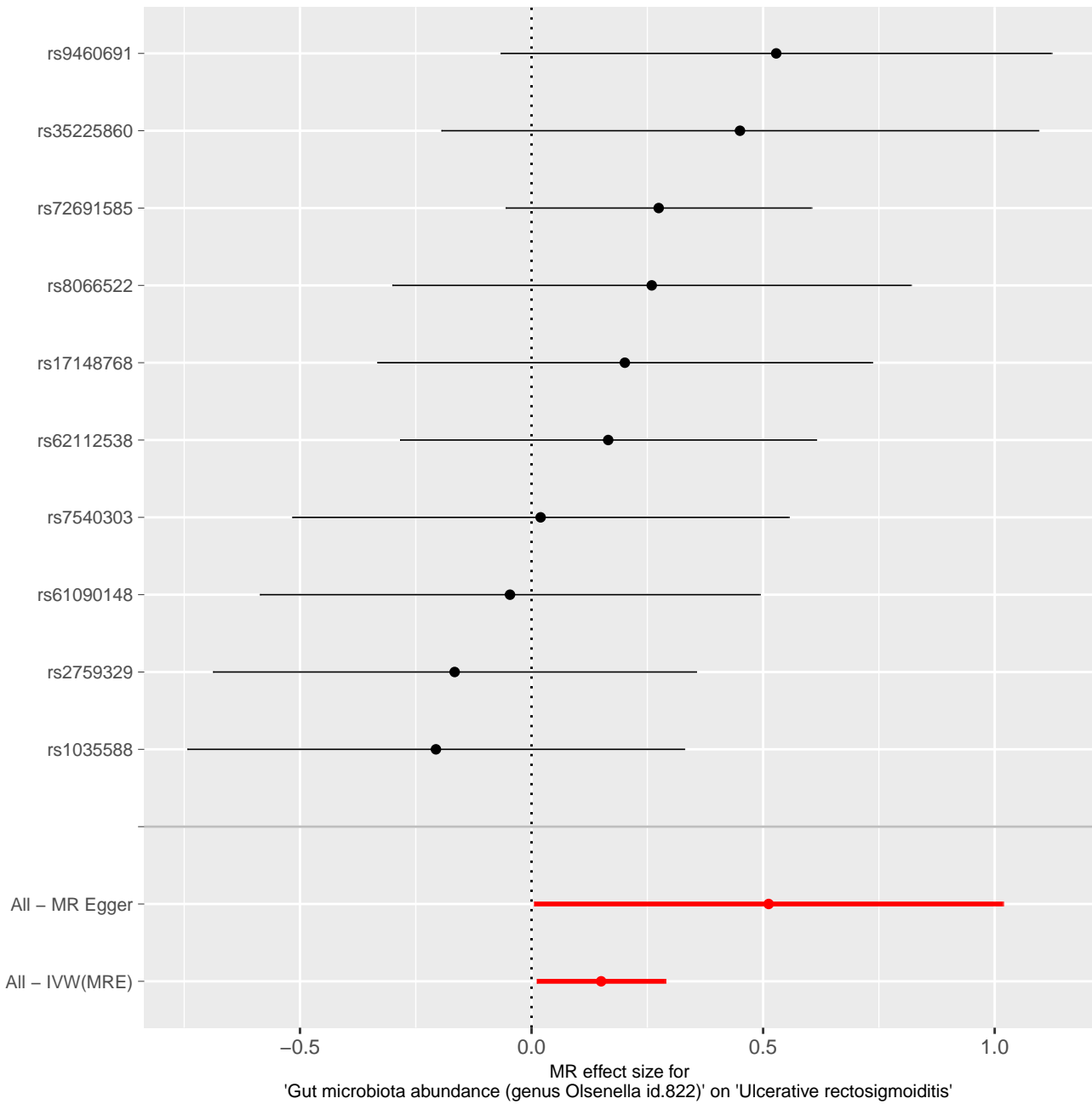




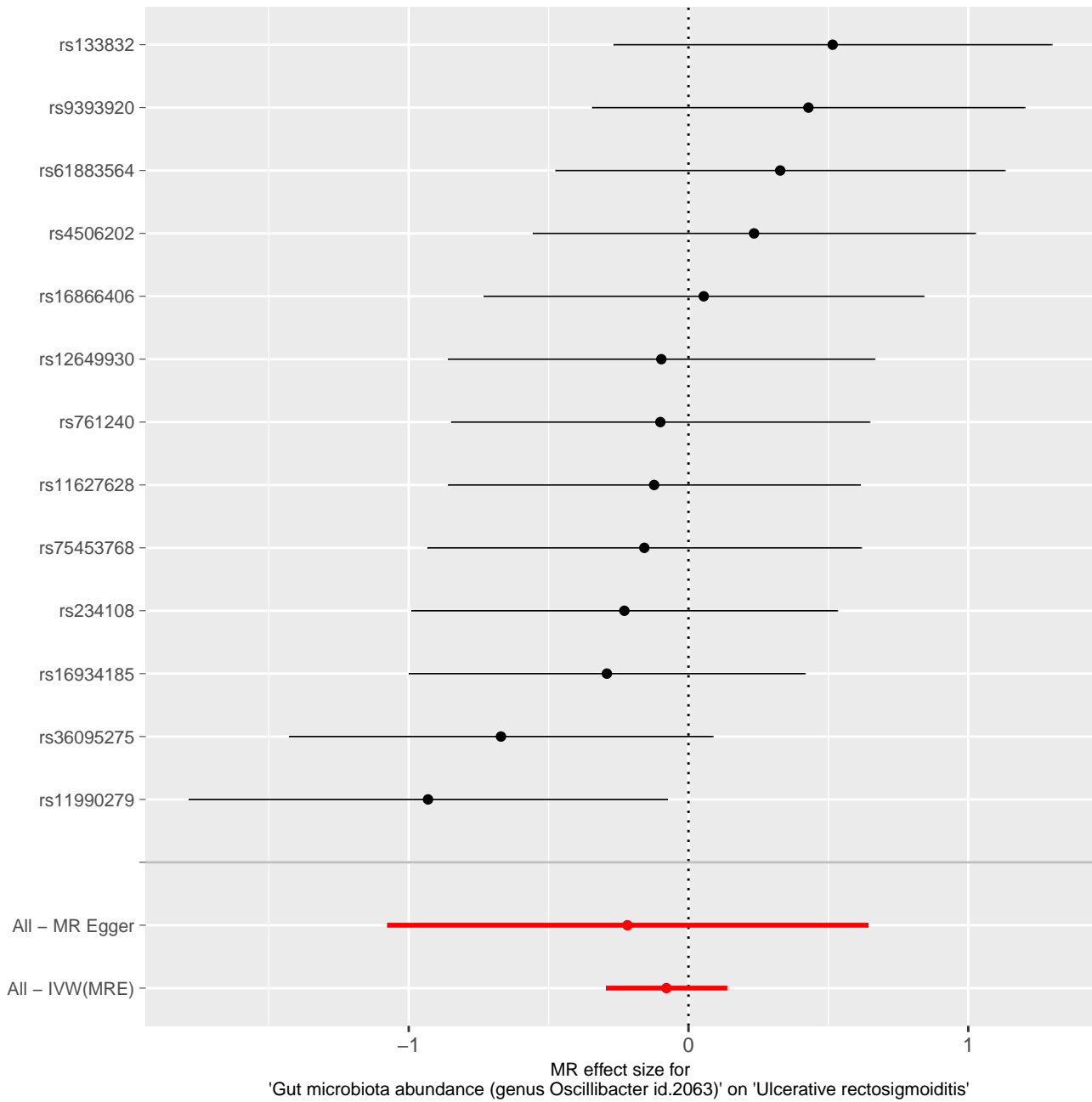


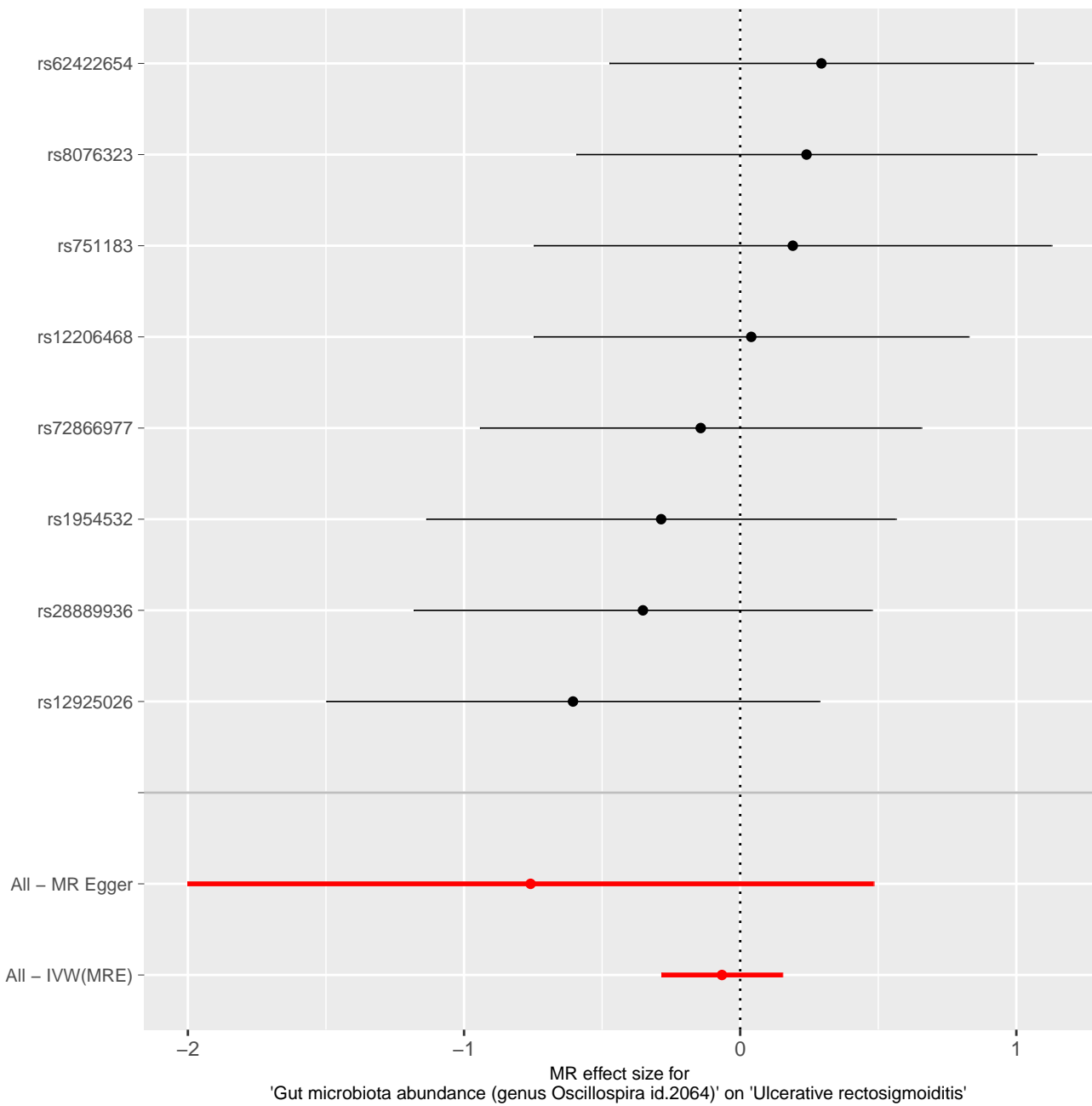


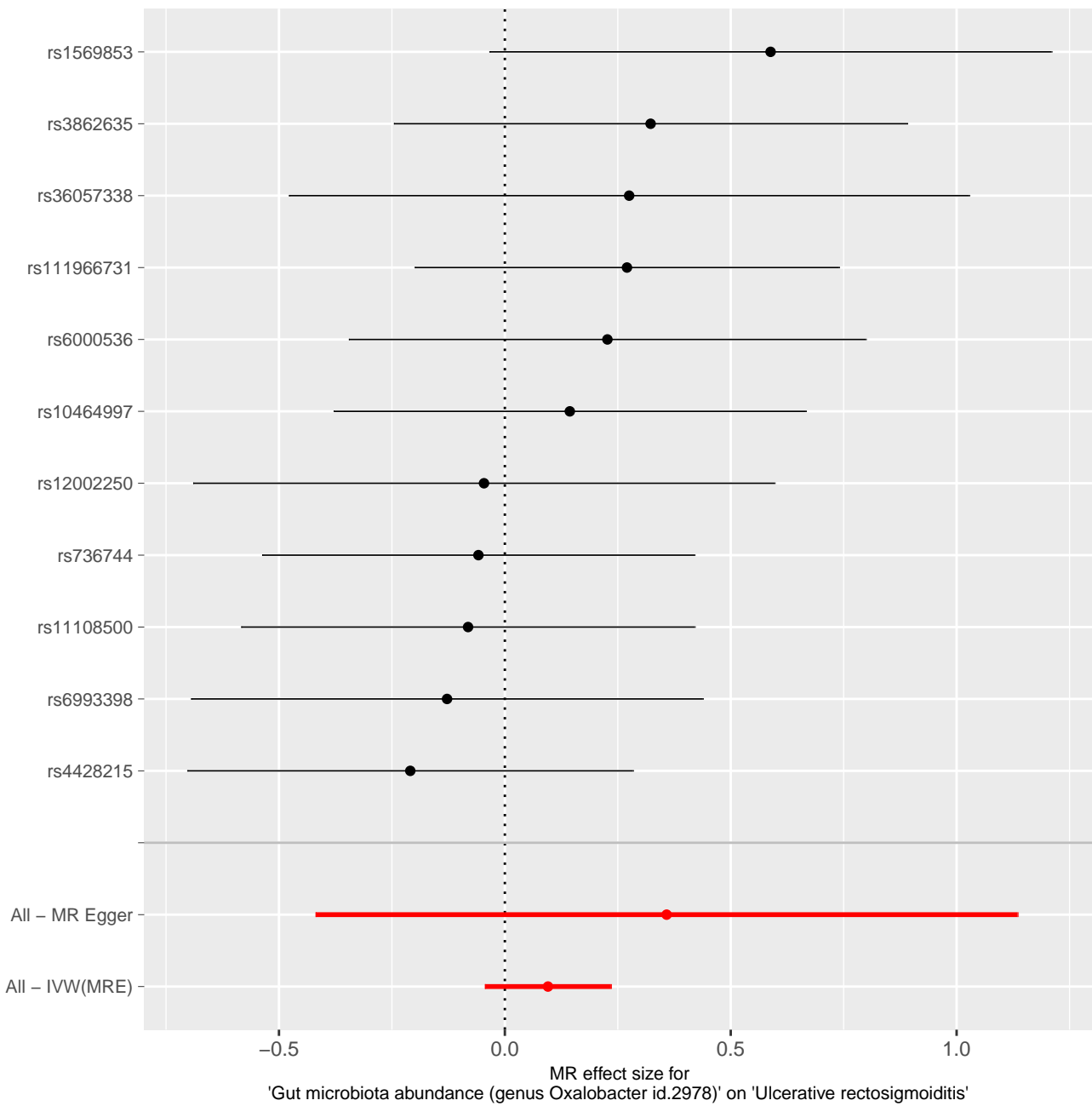


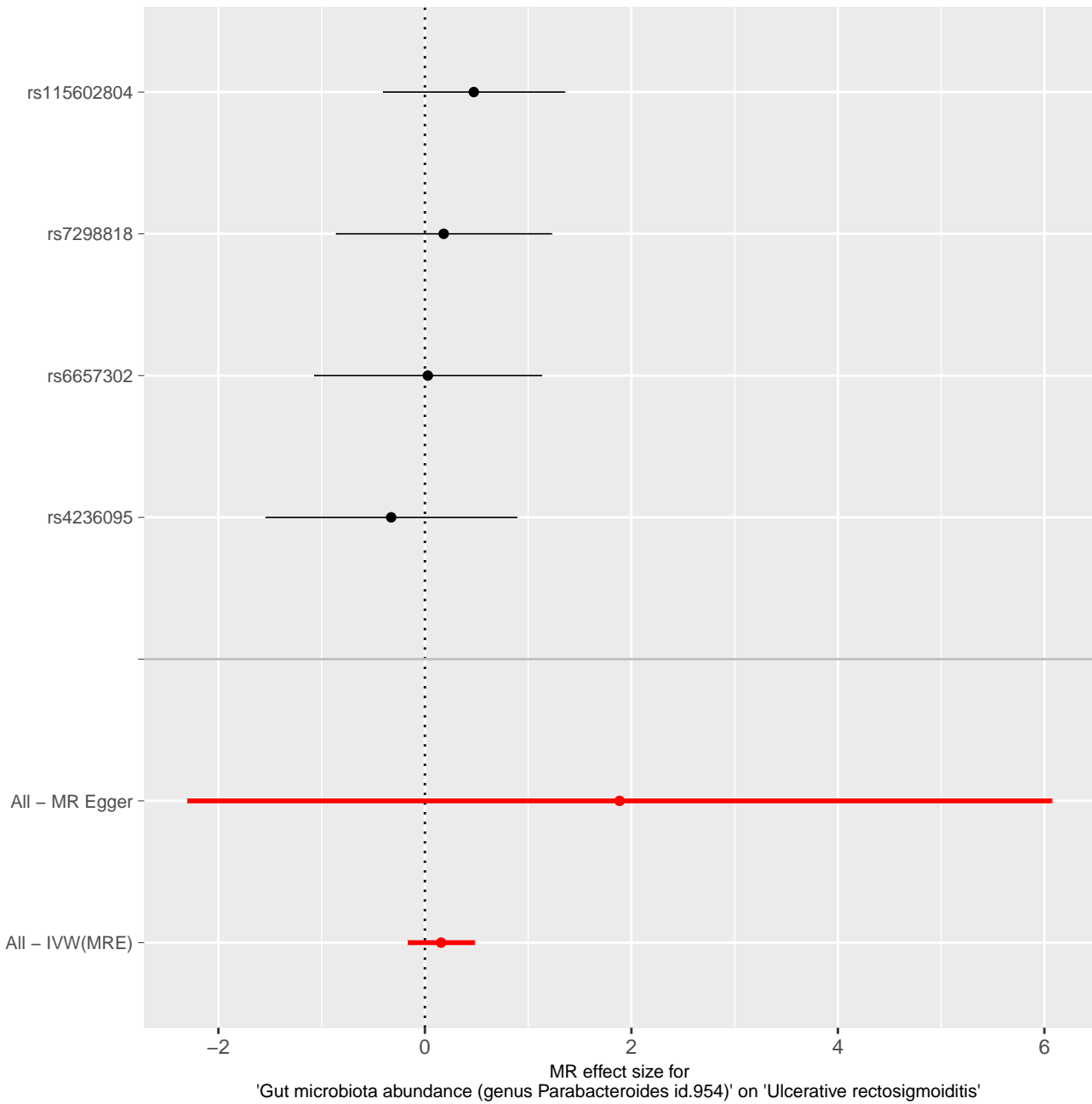


Batch 1610 : Gut microbiota abundance (genus Oscillibacter id.2063) on Ulcerative rectosigmoiditis

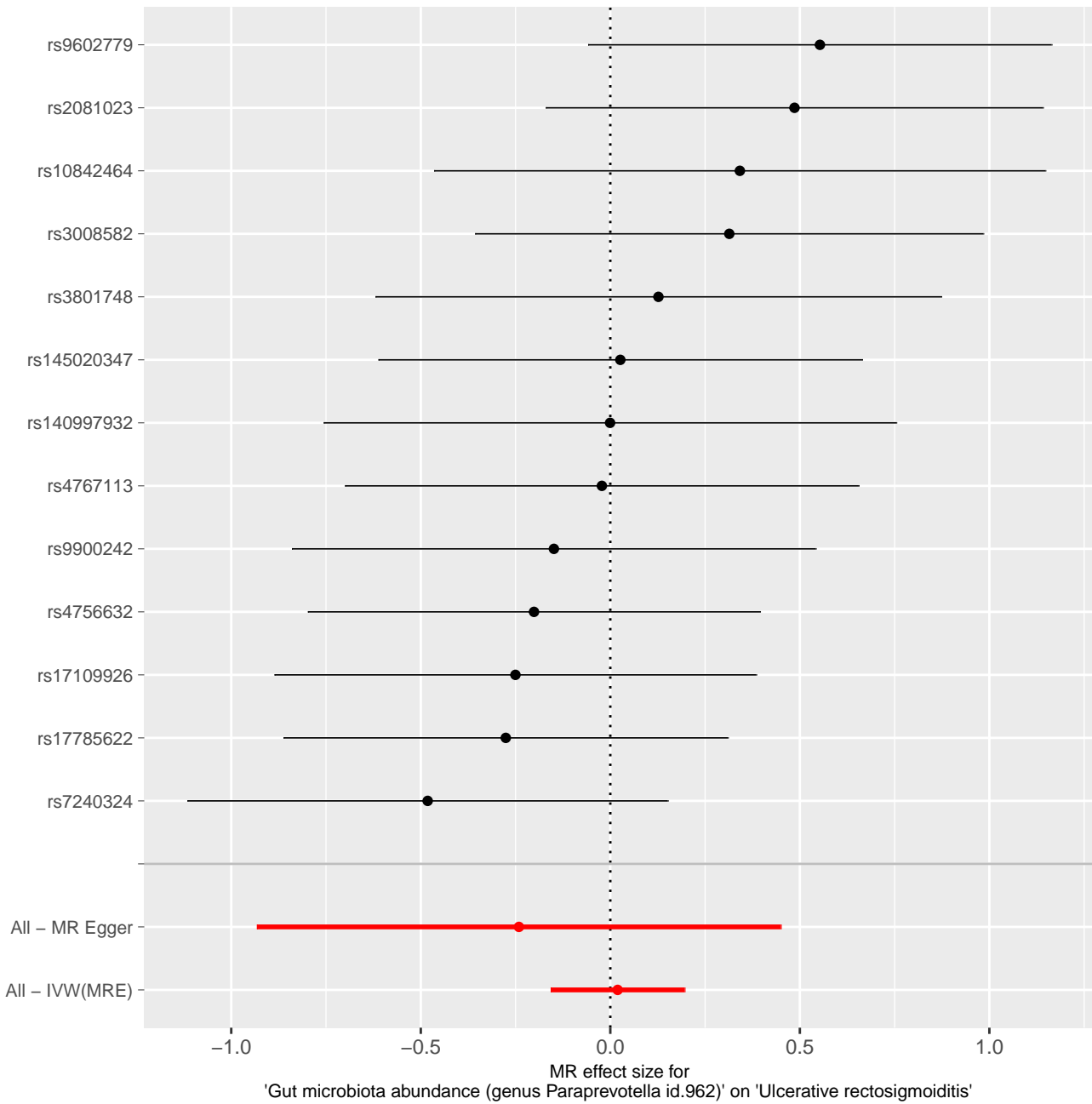


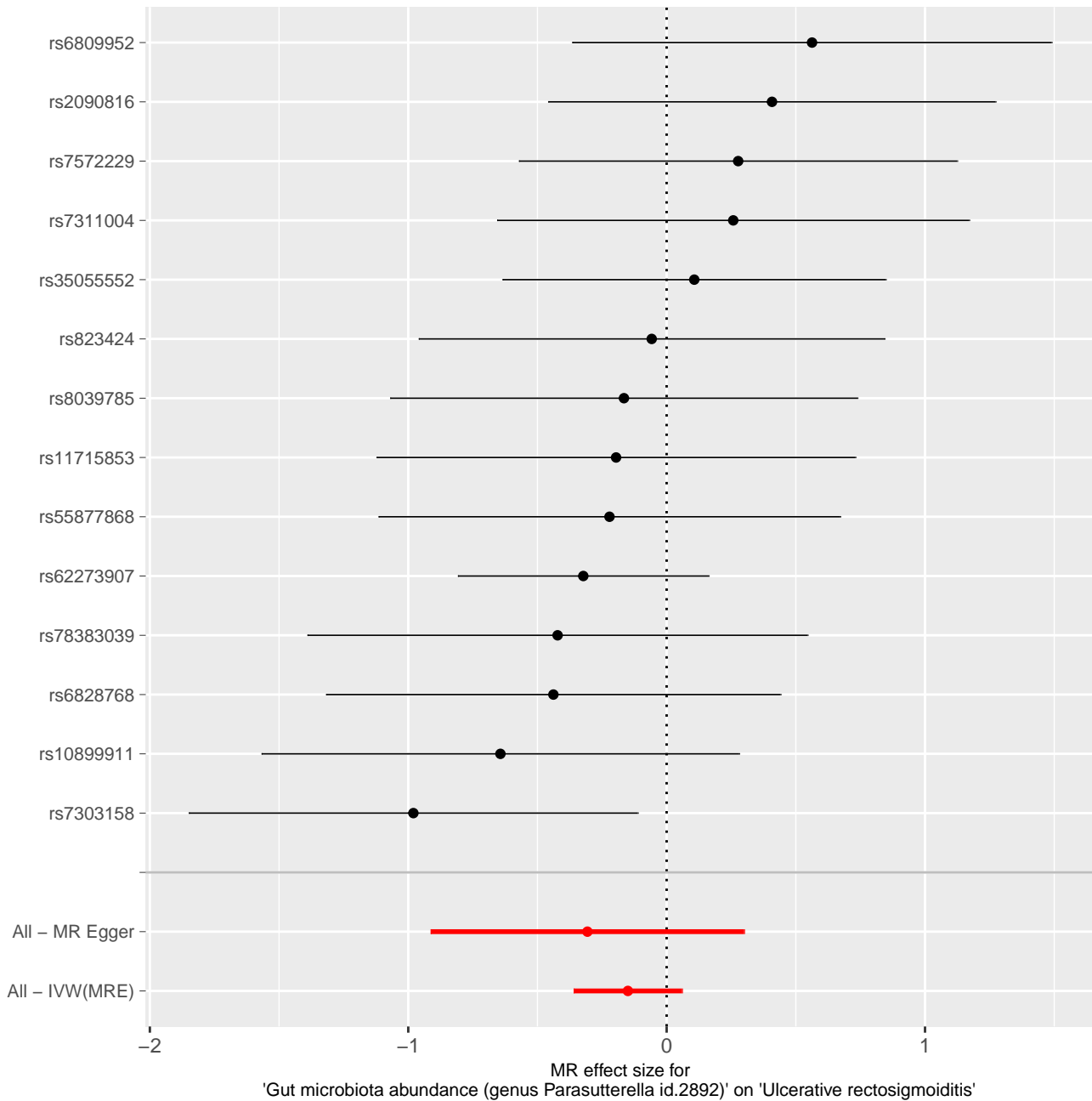


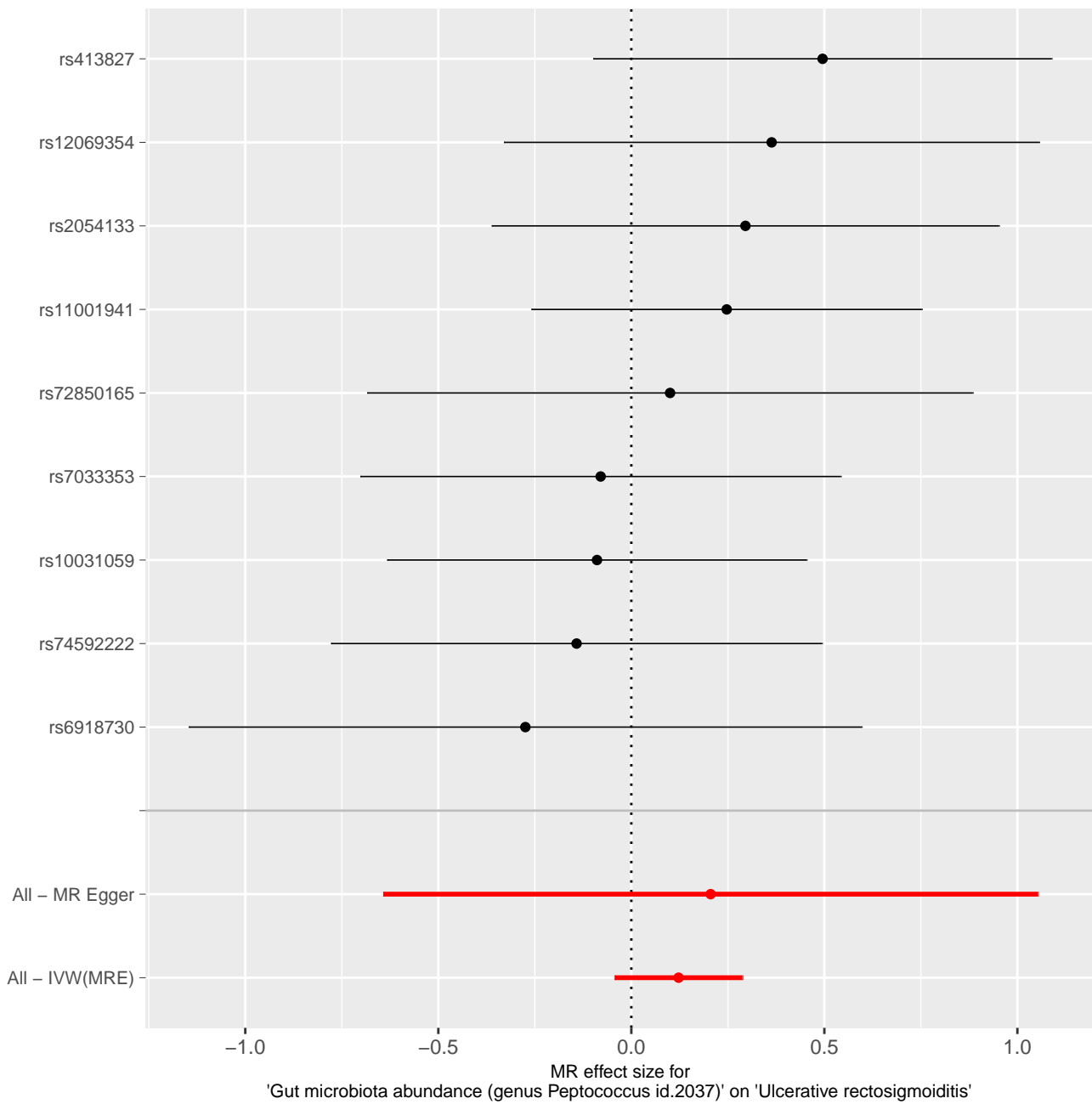


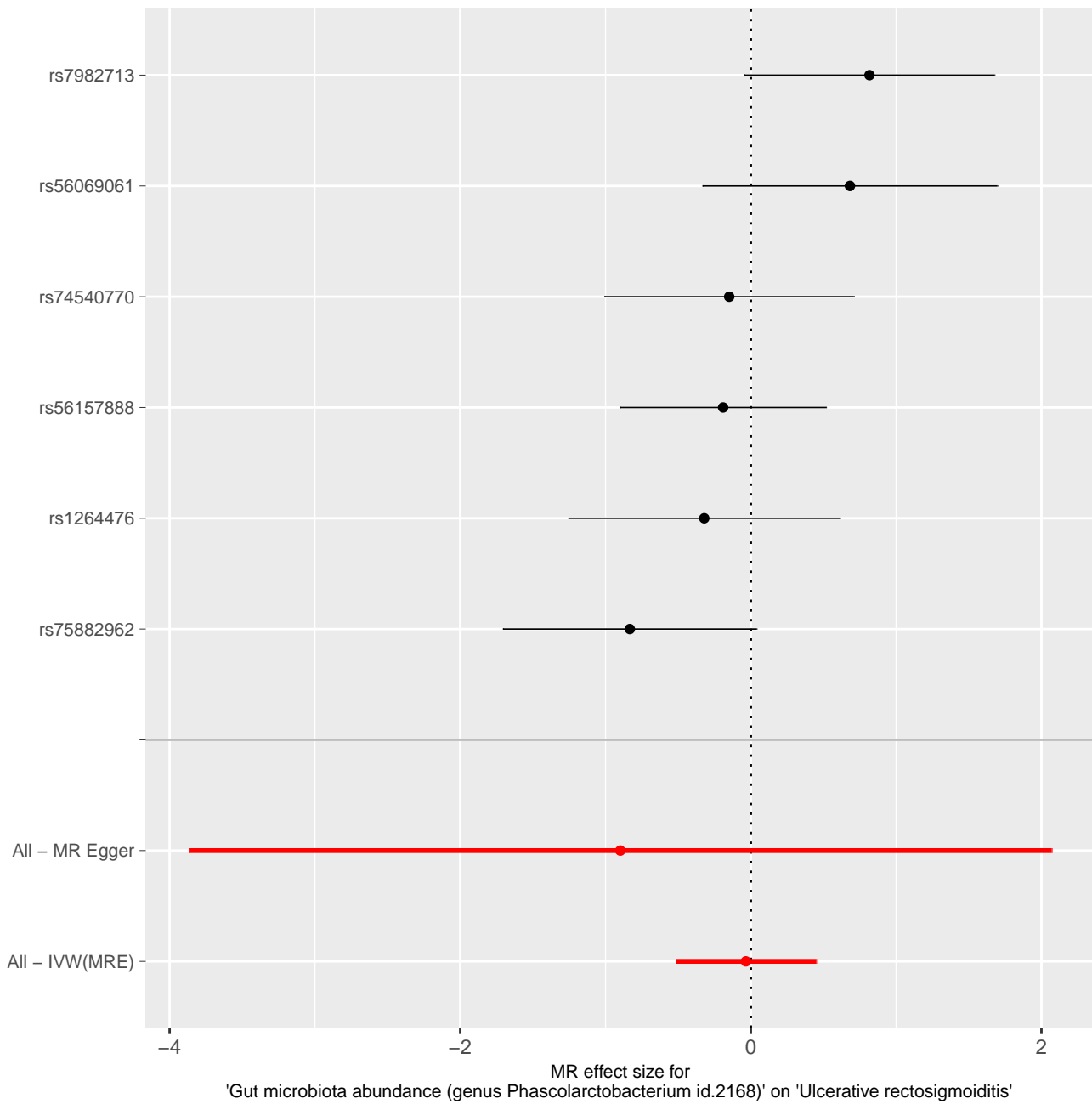


Batch 1614 : Gut microbiota abundance (genus Paraprevotella id.962) on Ulcerative rectosigmoiditis

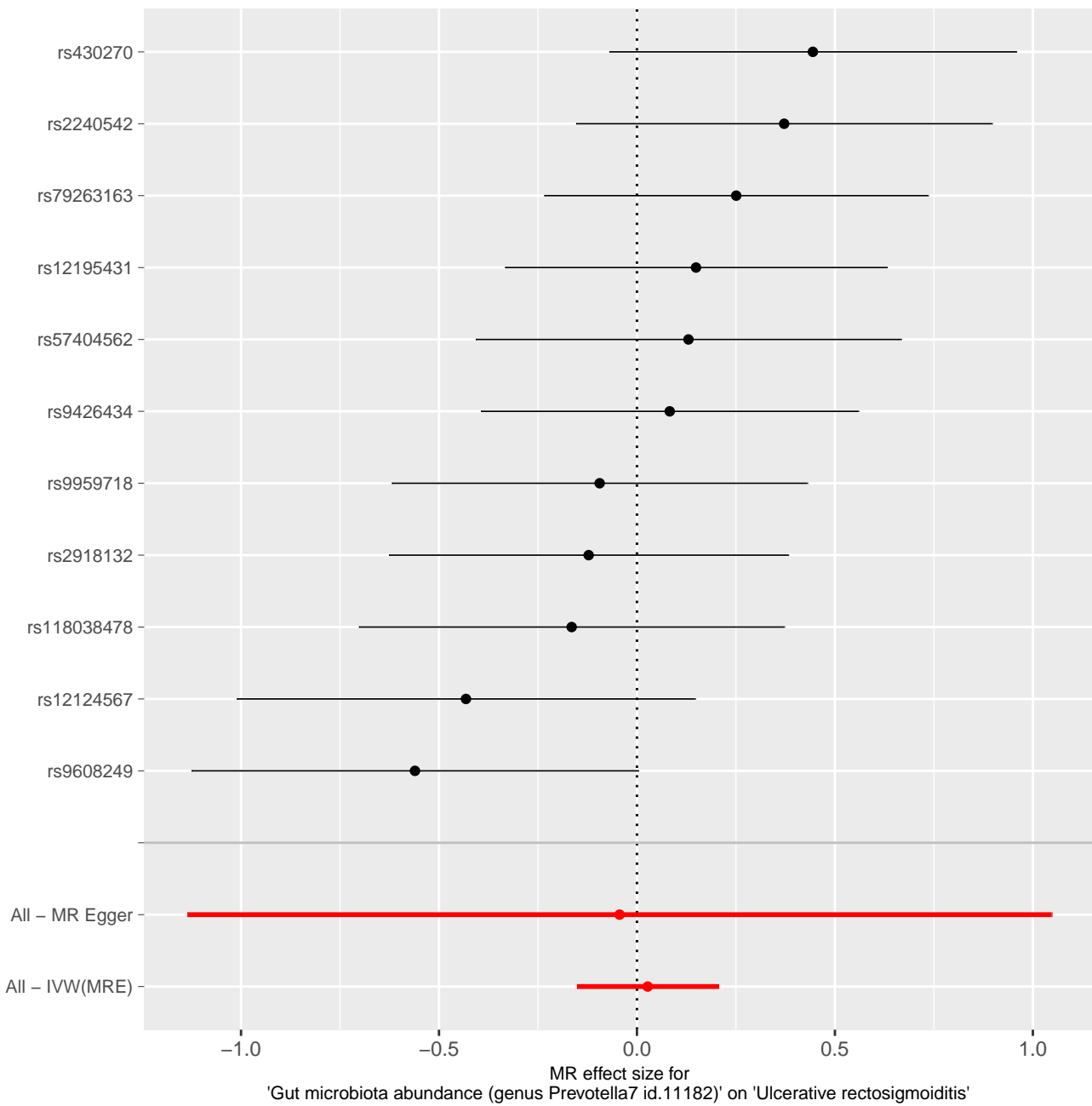




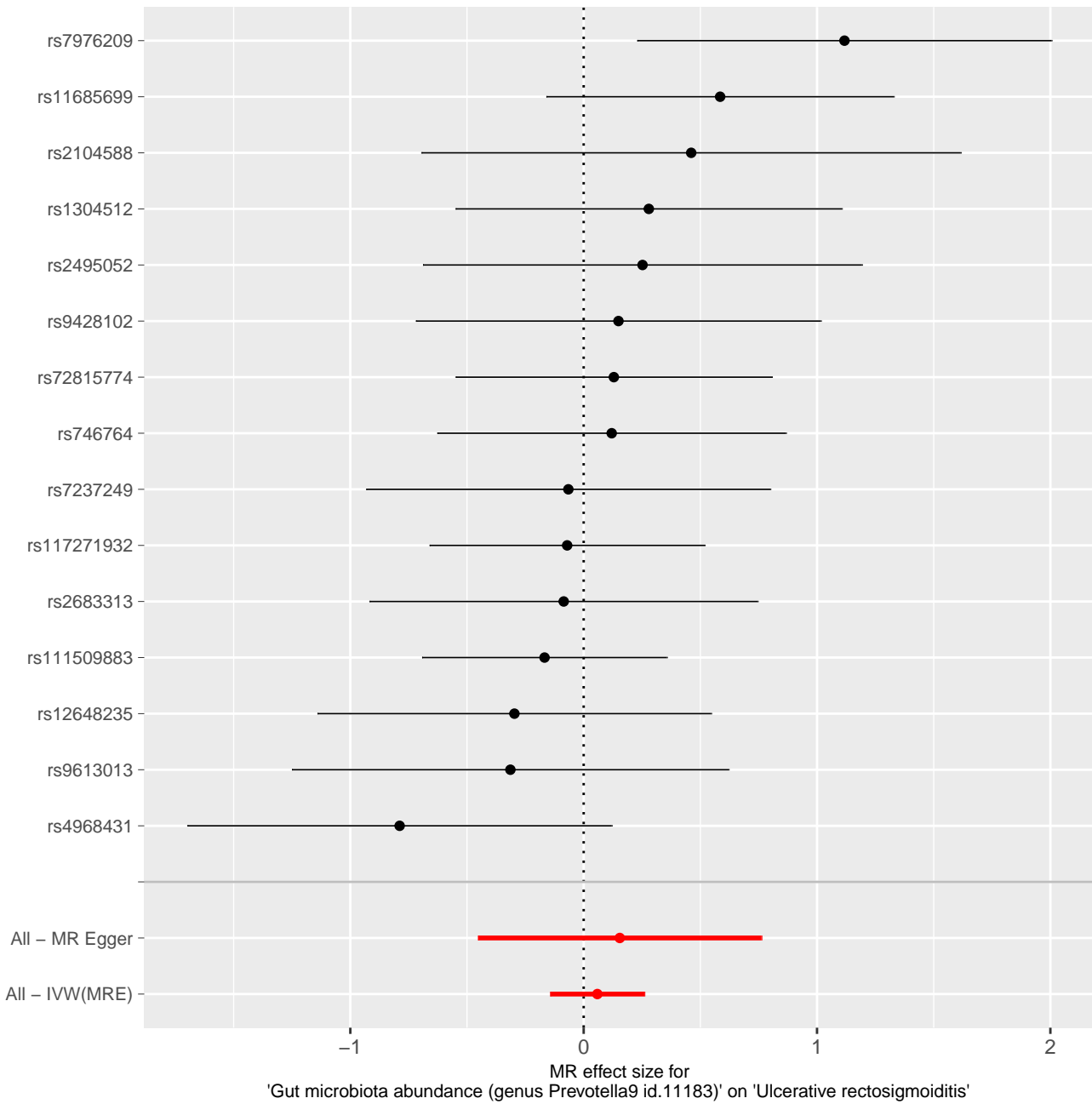


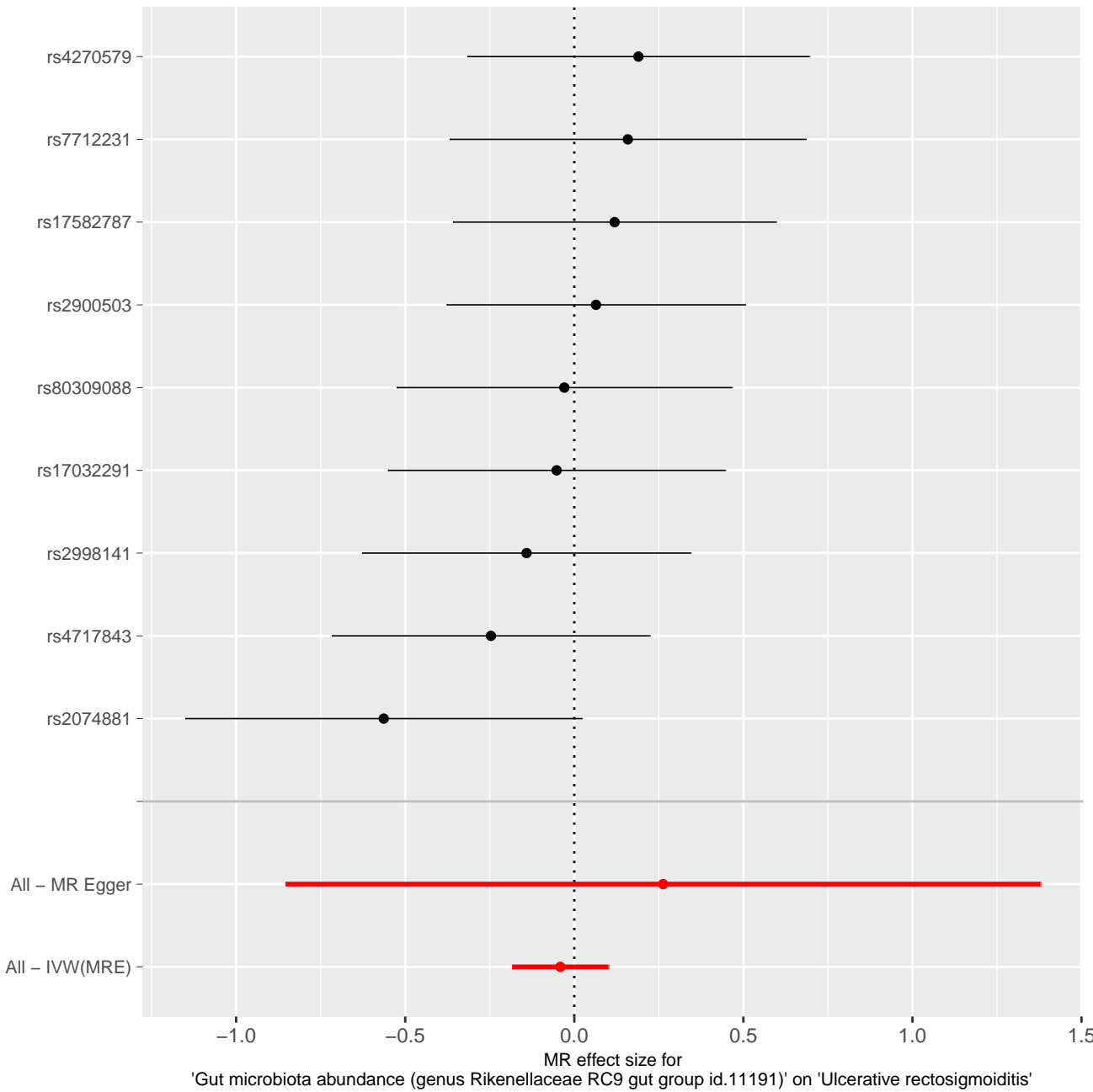


Batch 1618 : Gut microbiota abundance (genus Prevotella7 id.11182) on Ulcerative rectosigmoiditis

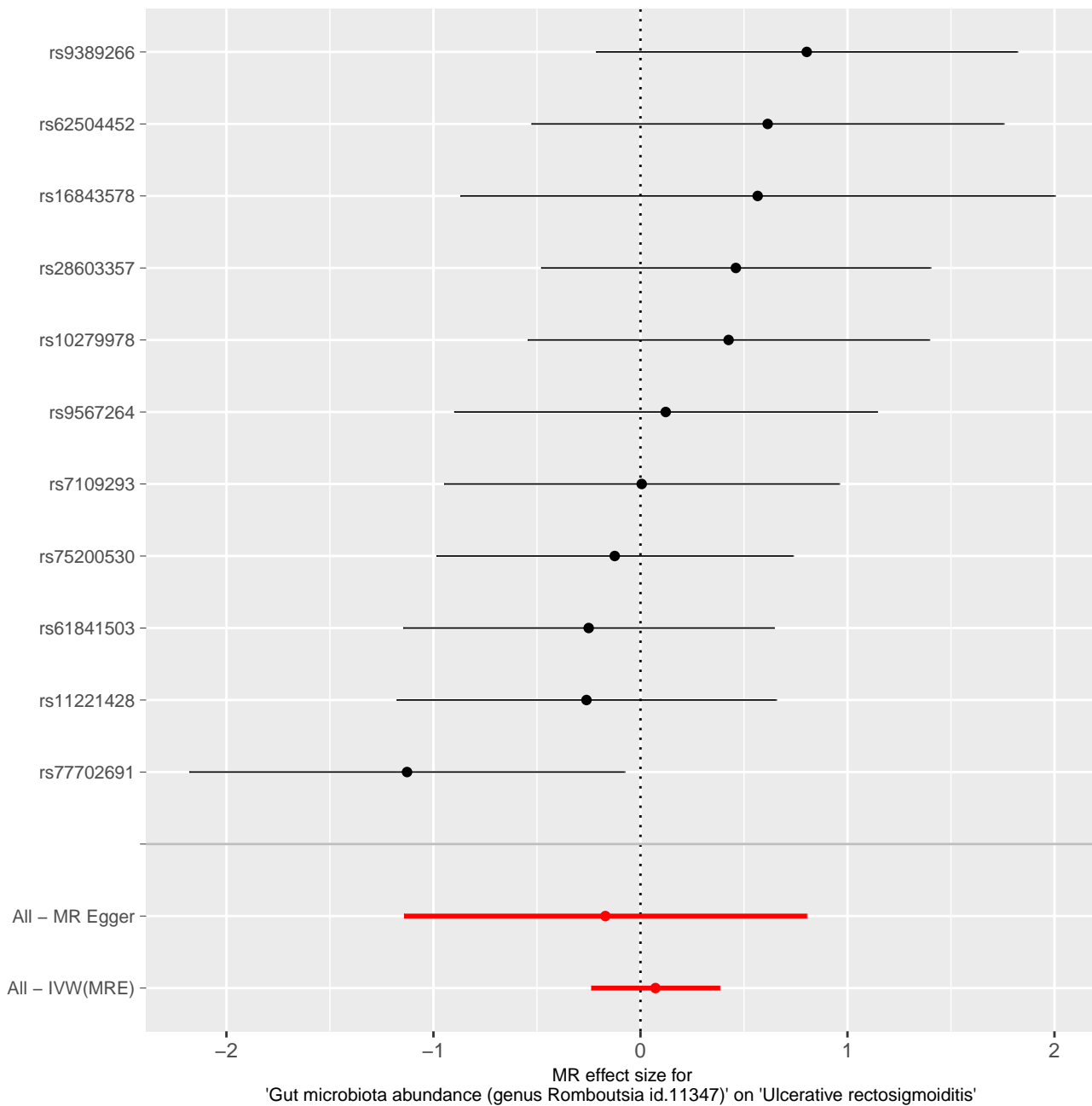


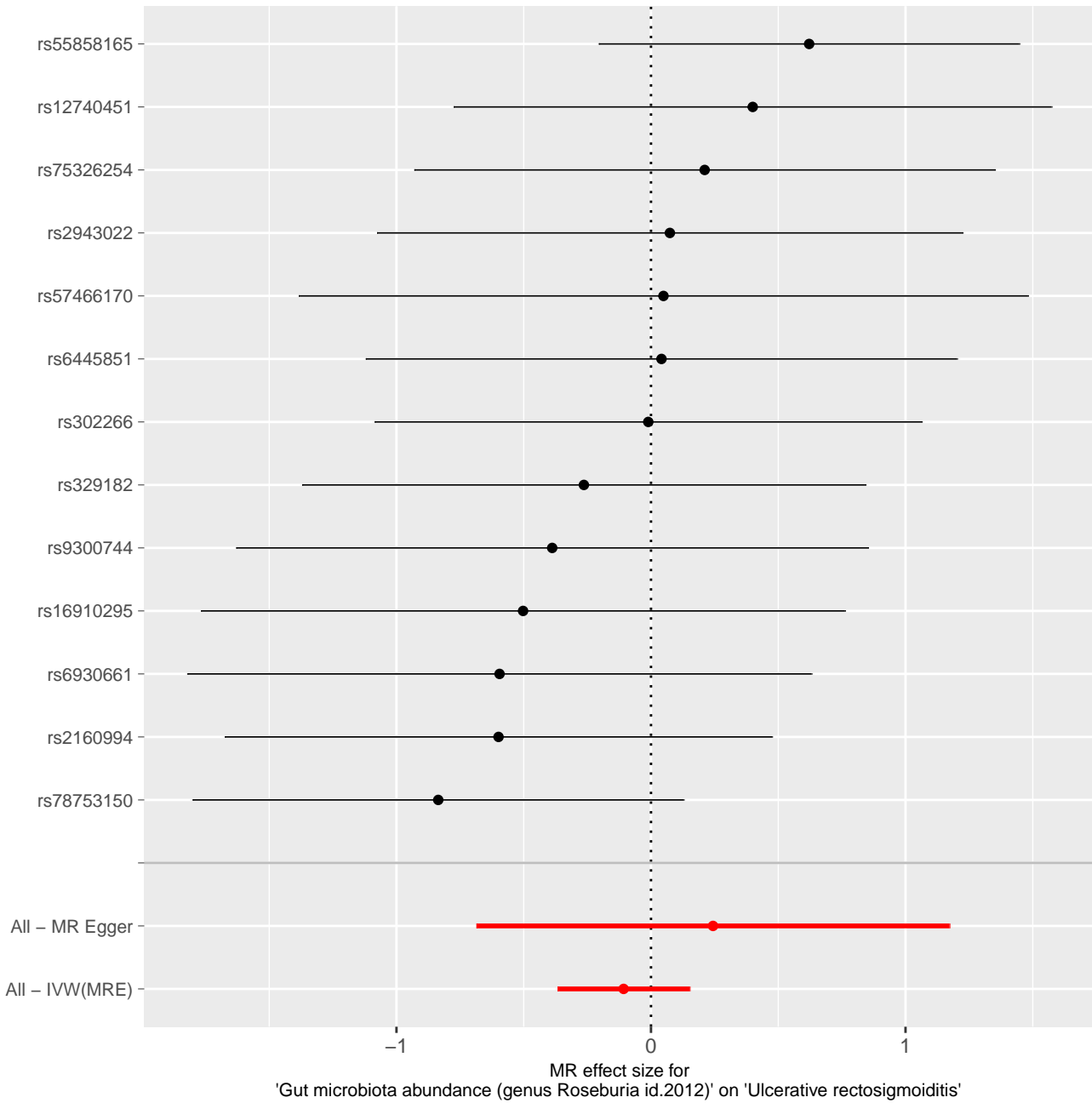
Batch 1619 : Gut microbiota abundance (genus Prevotella9 id.11183) on Ulcerative rectosigmoiditis

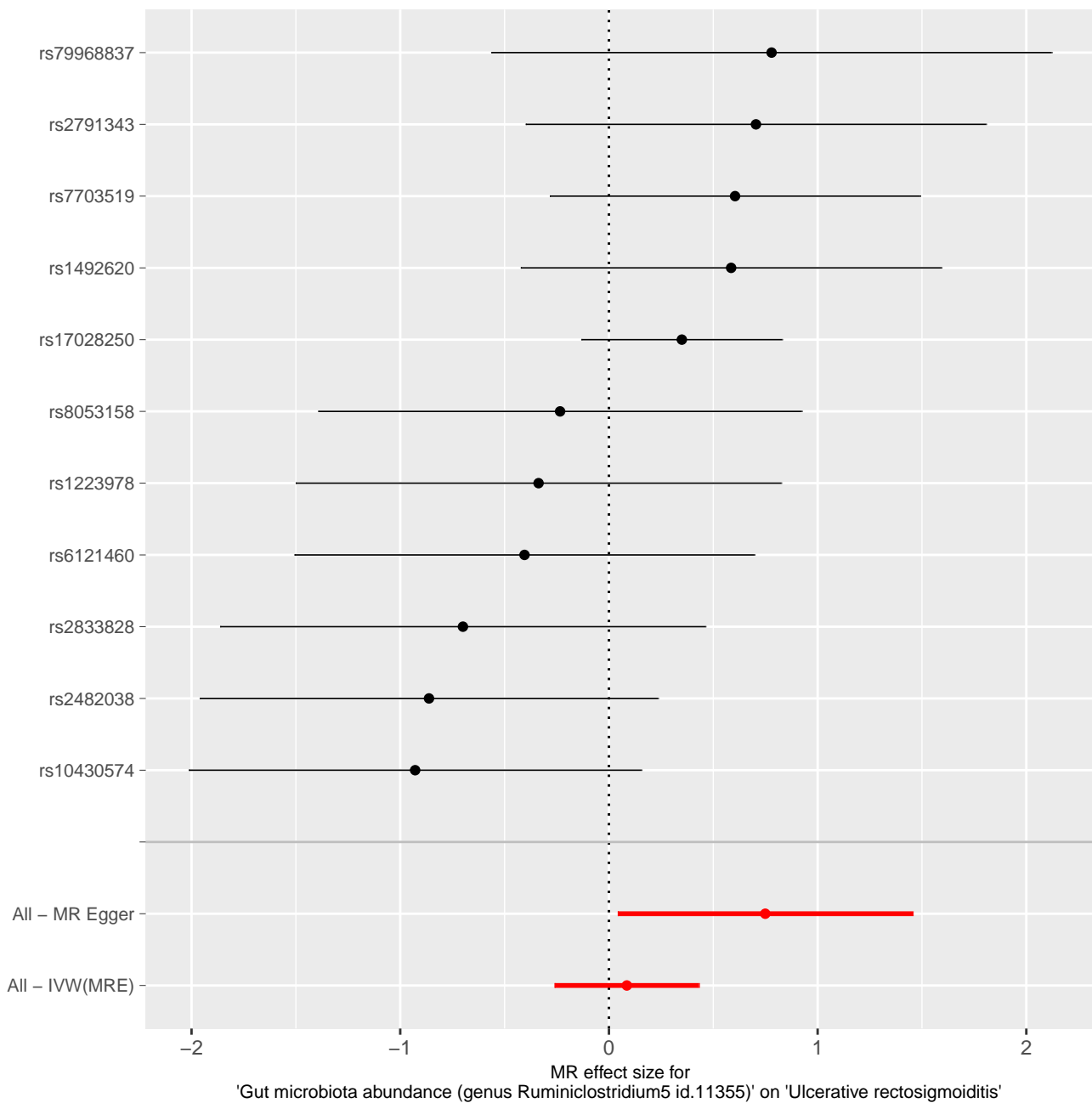


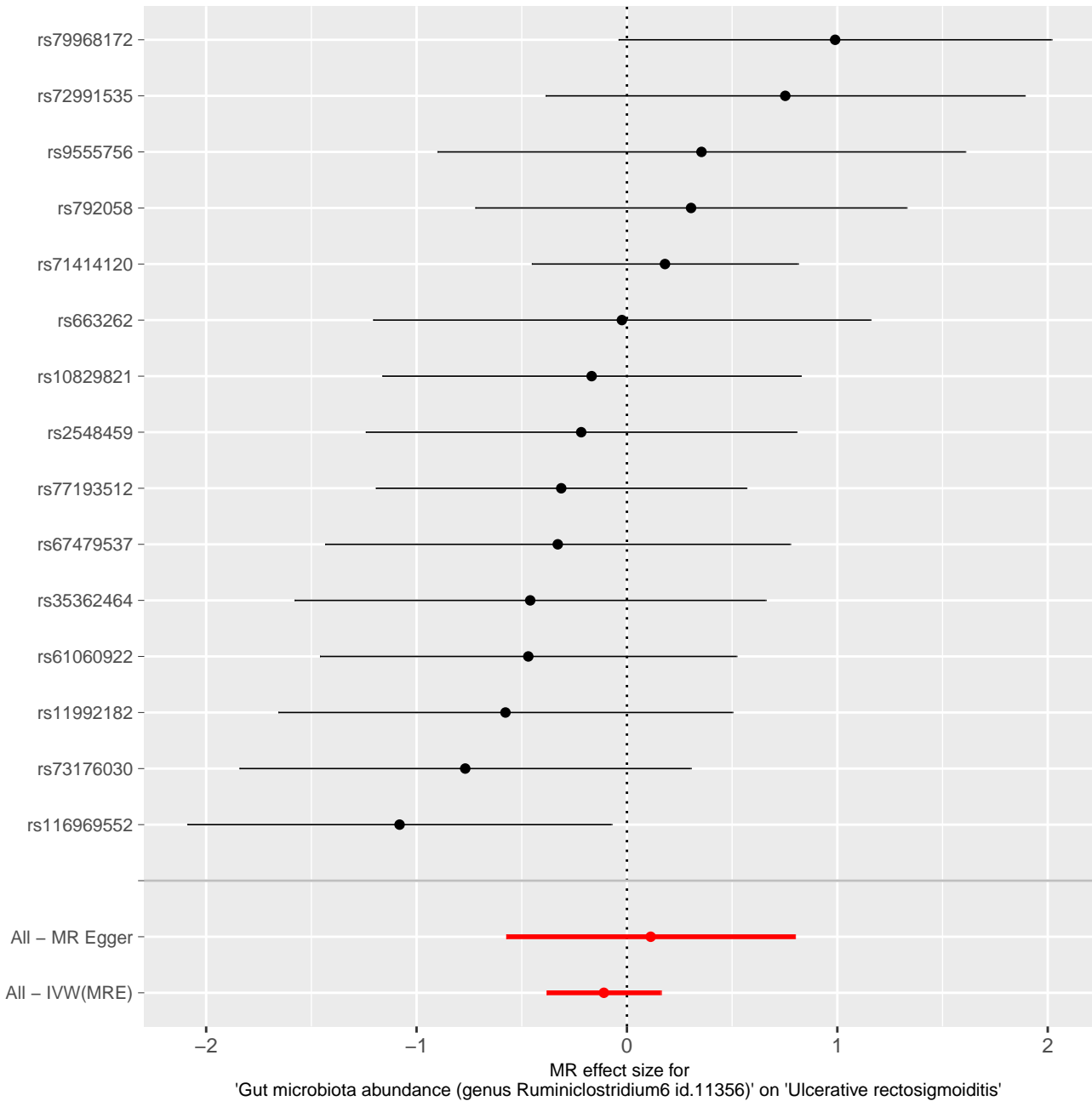


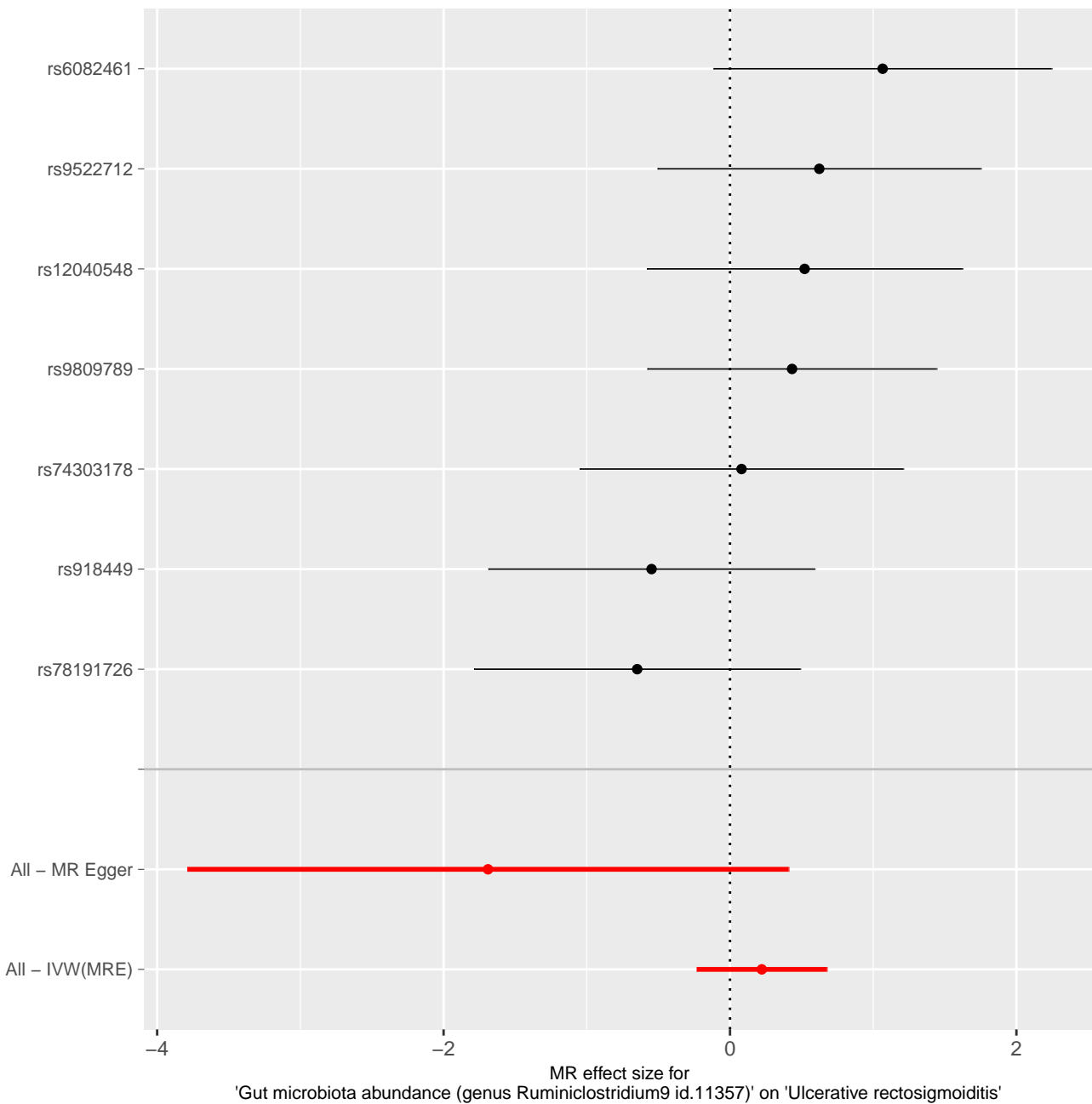
Batch 1621 : Gut microbiota abundance (genus Romboutsia id.11347) on Ulcerative rectosigmoiditis

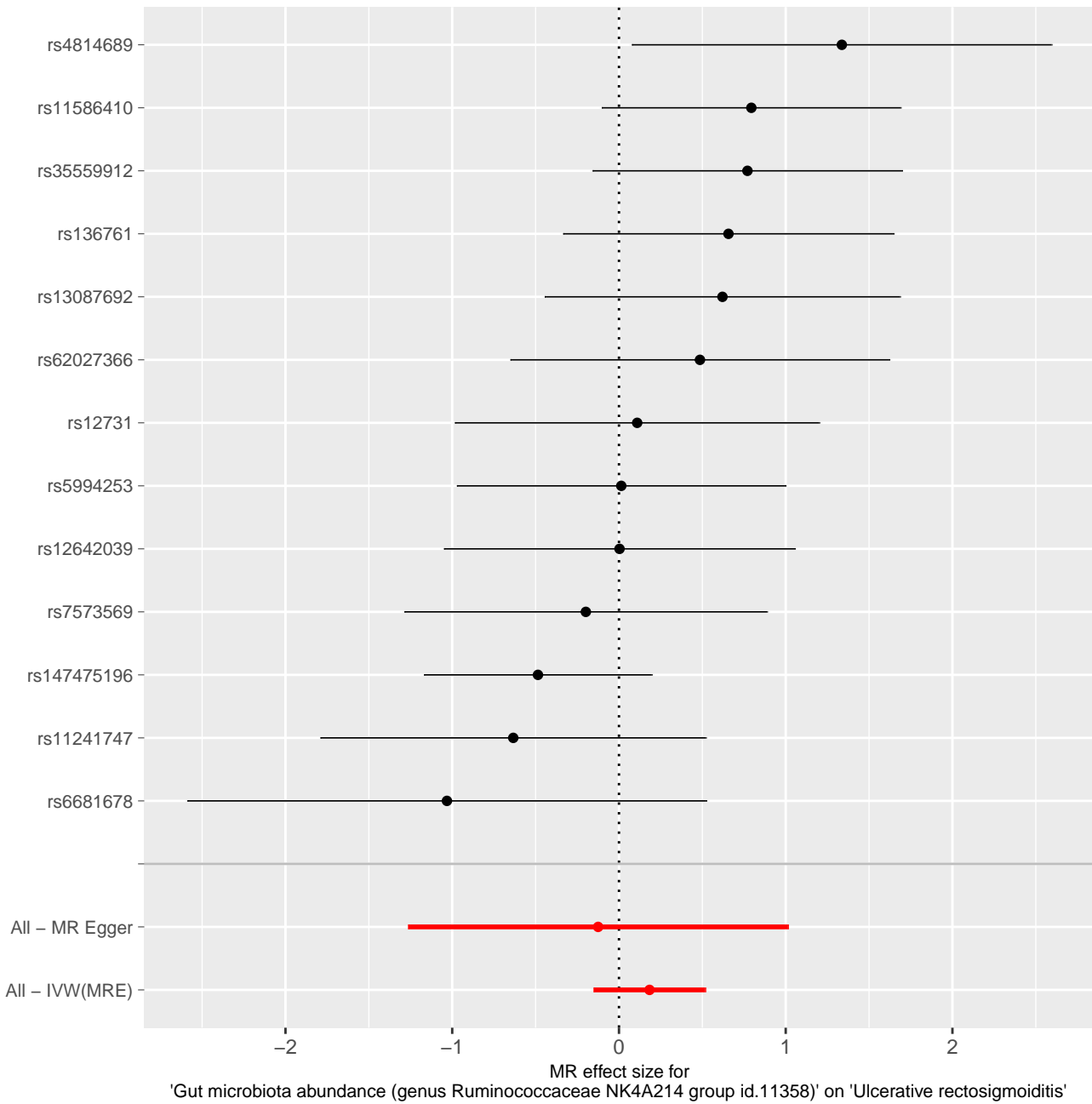


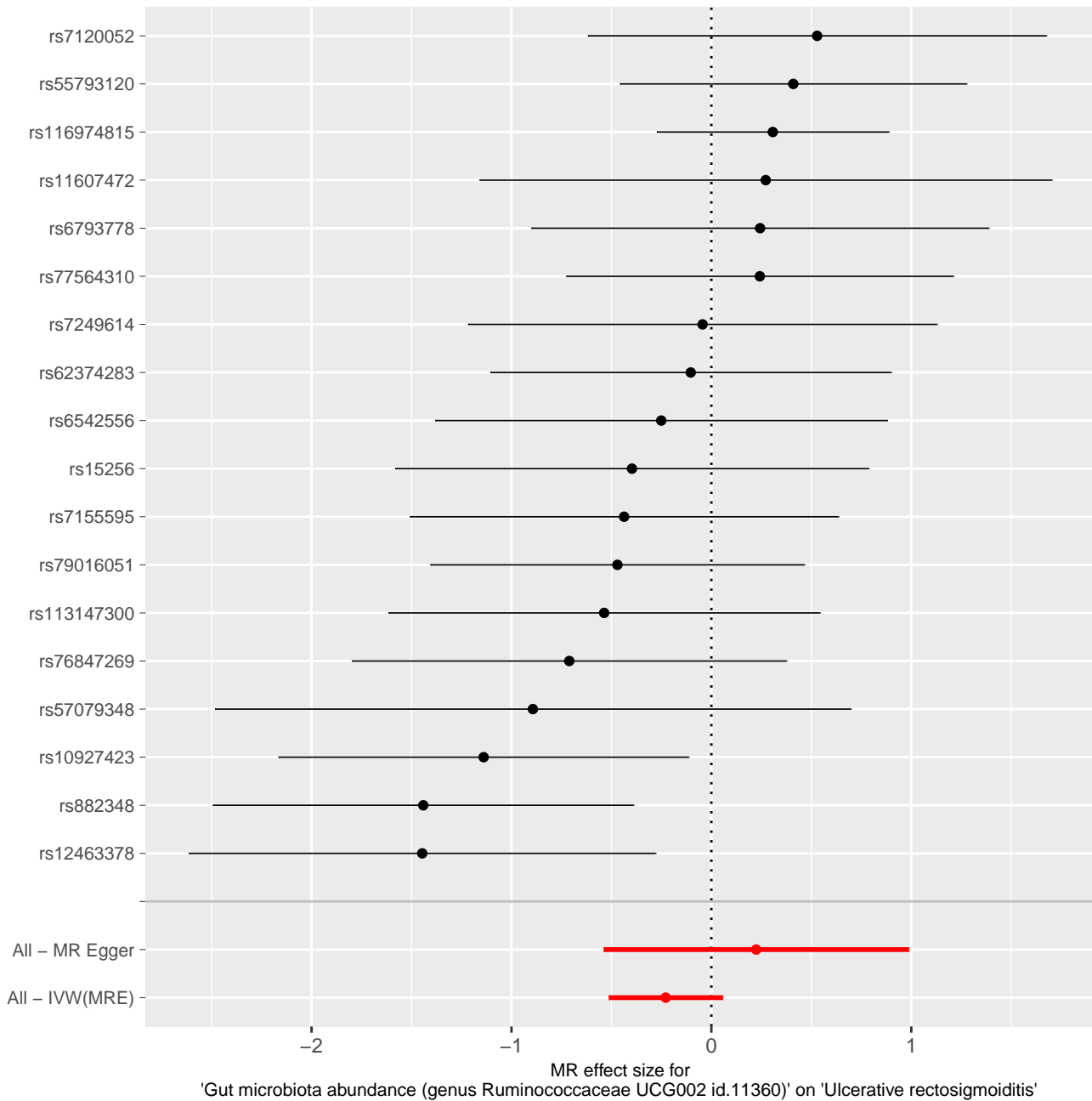




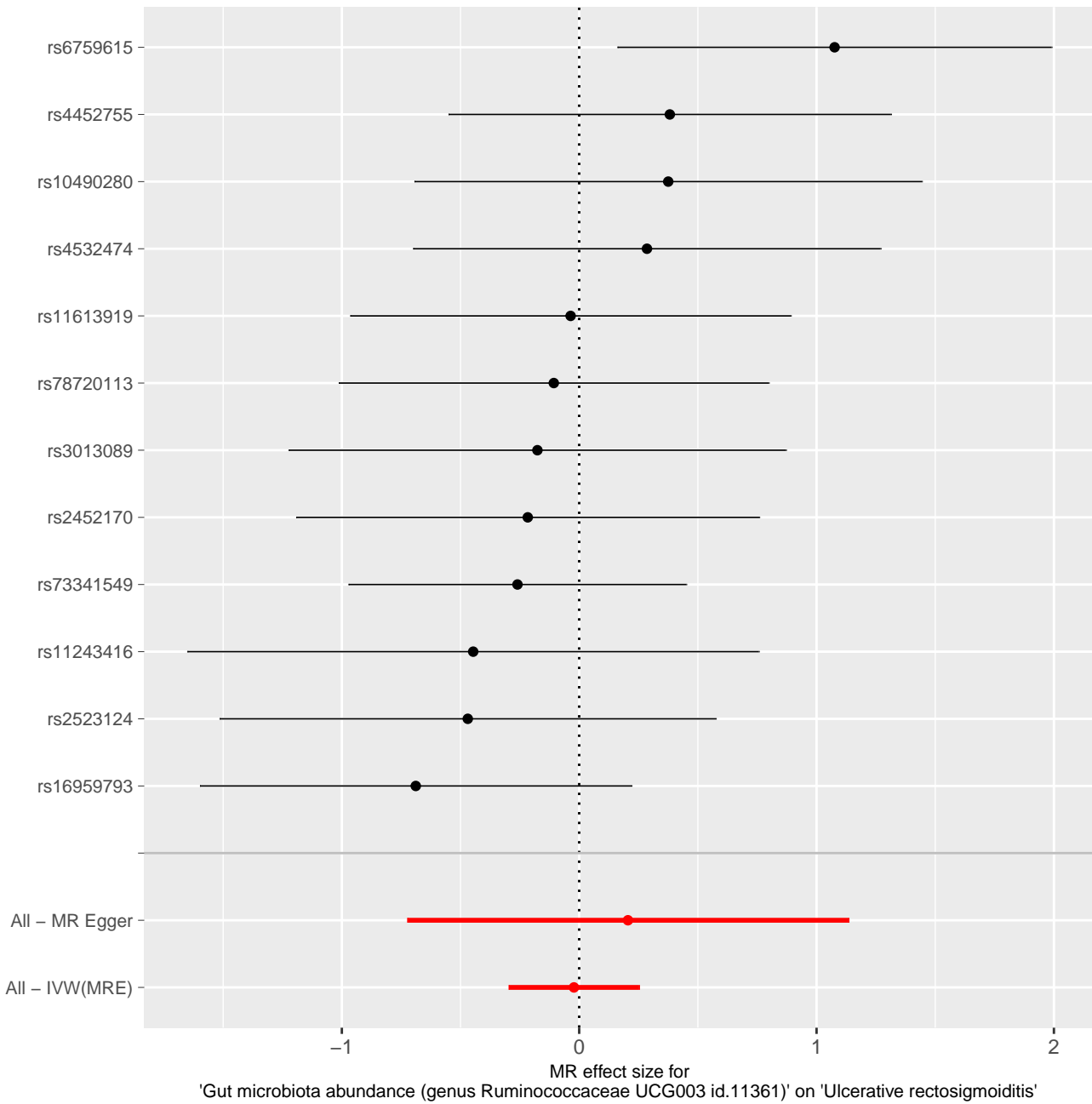


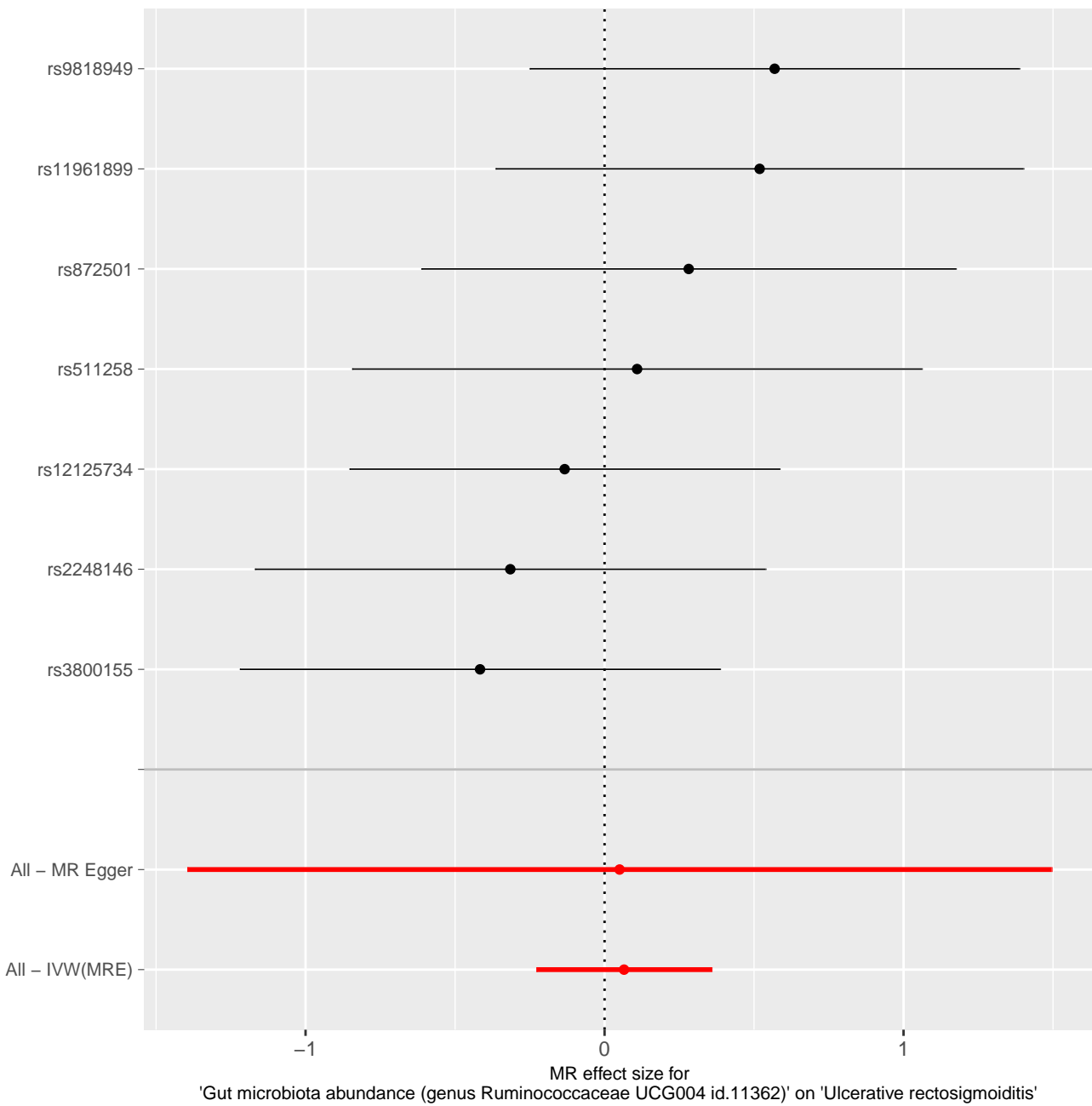


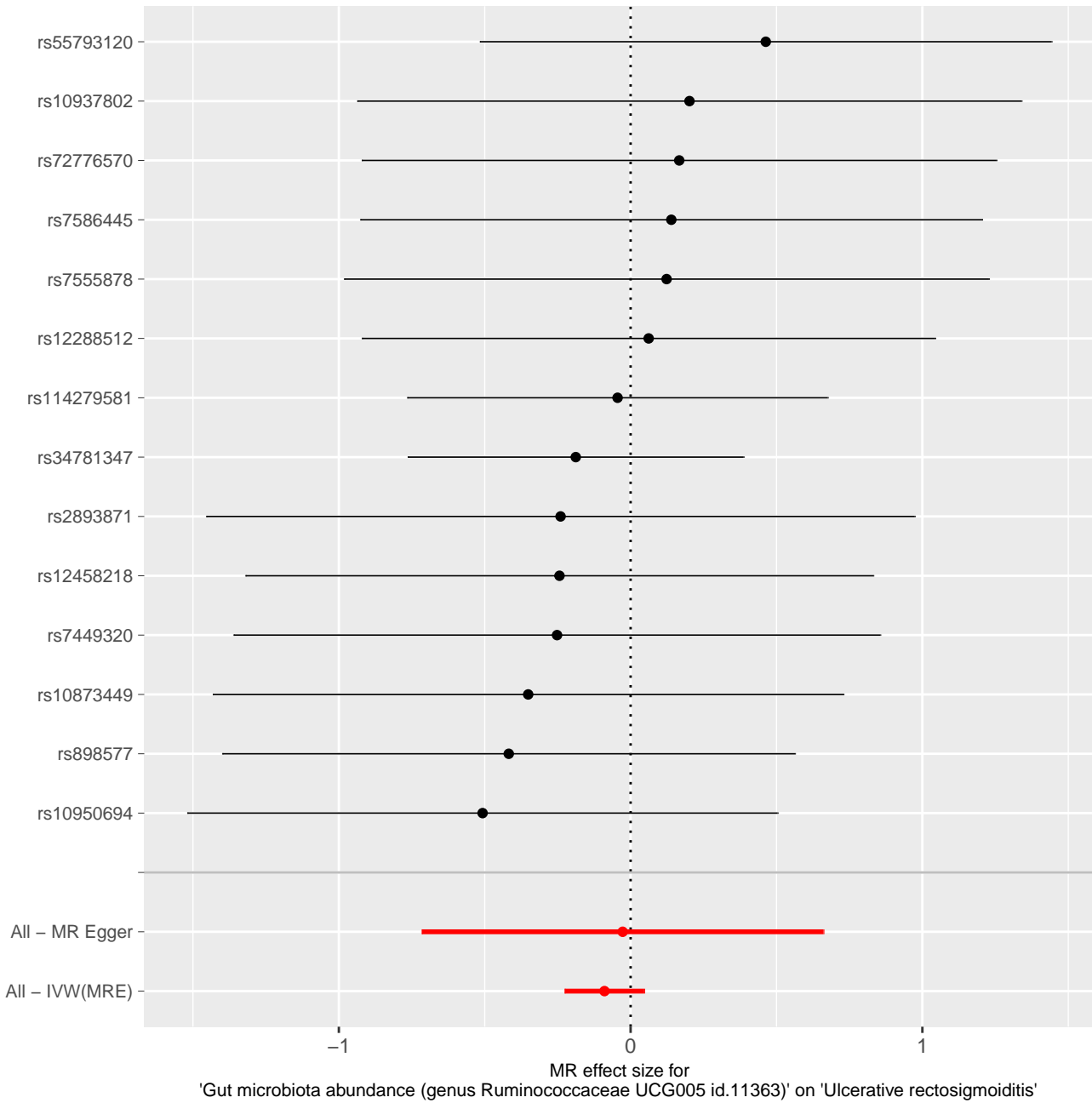


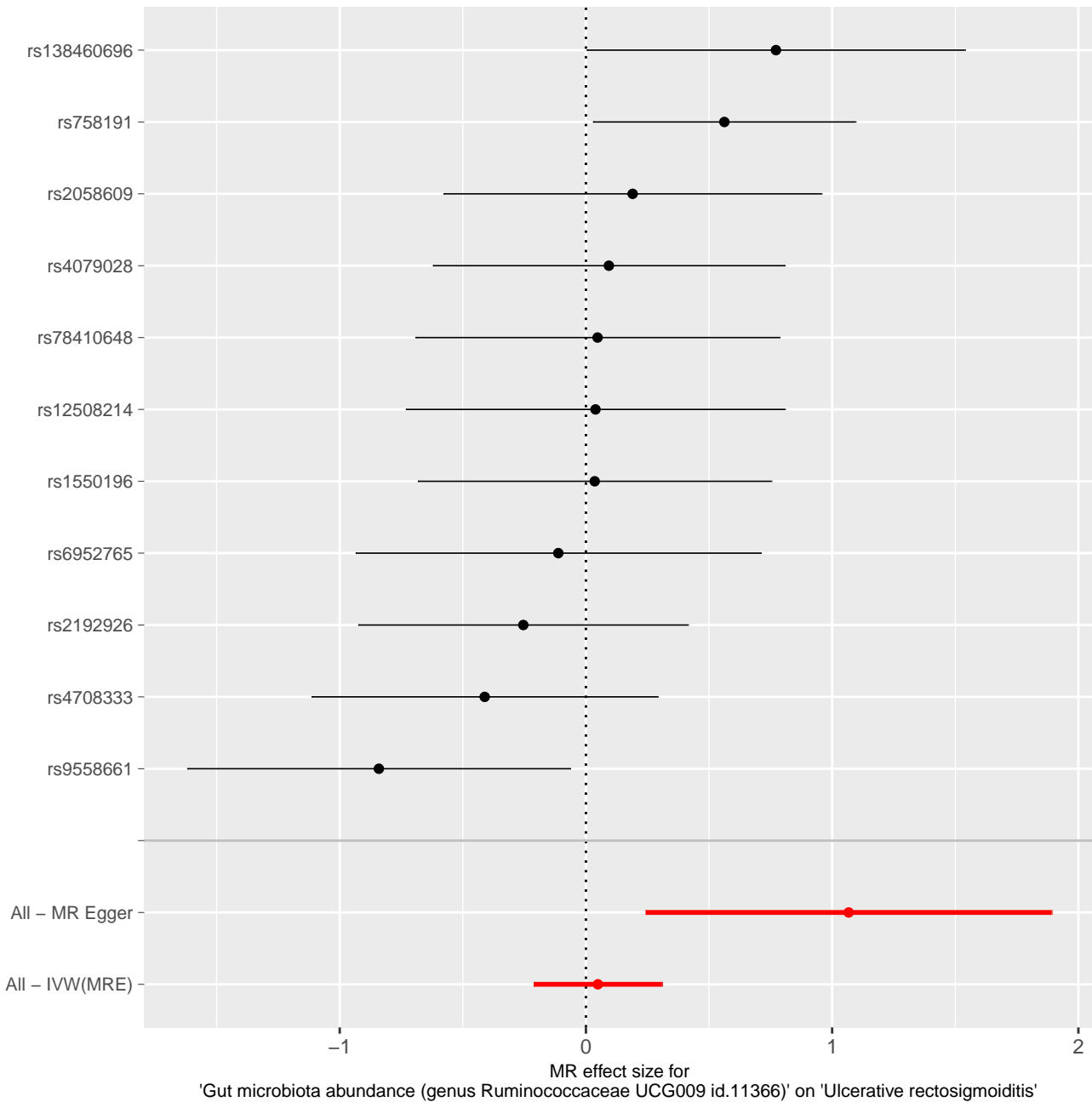


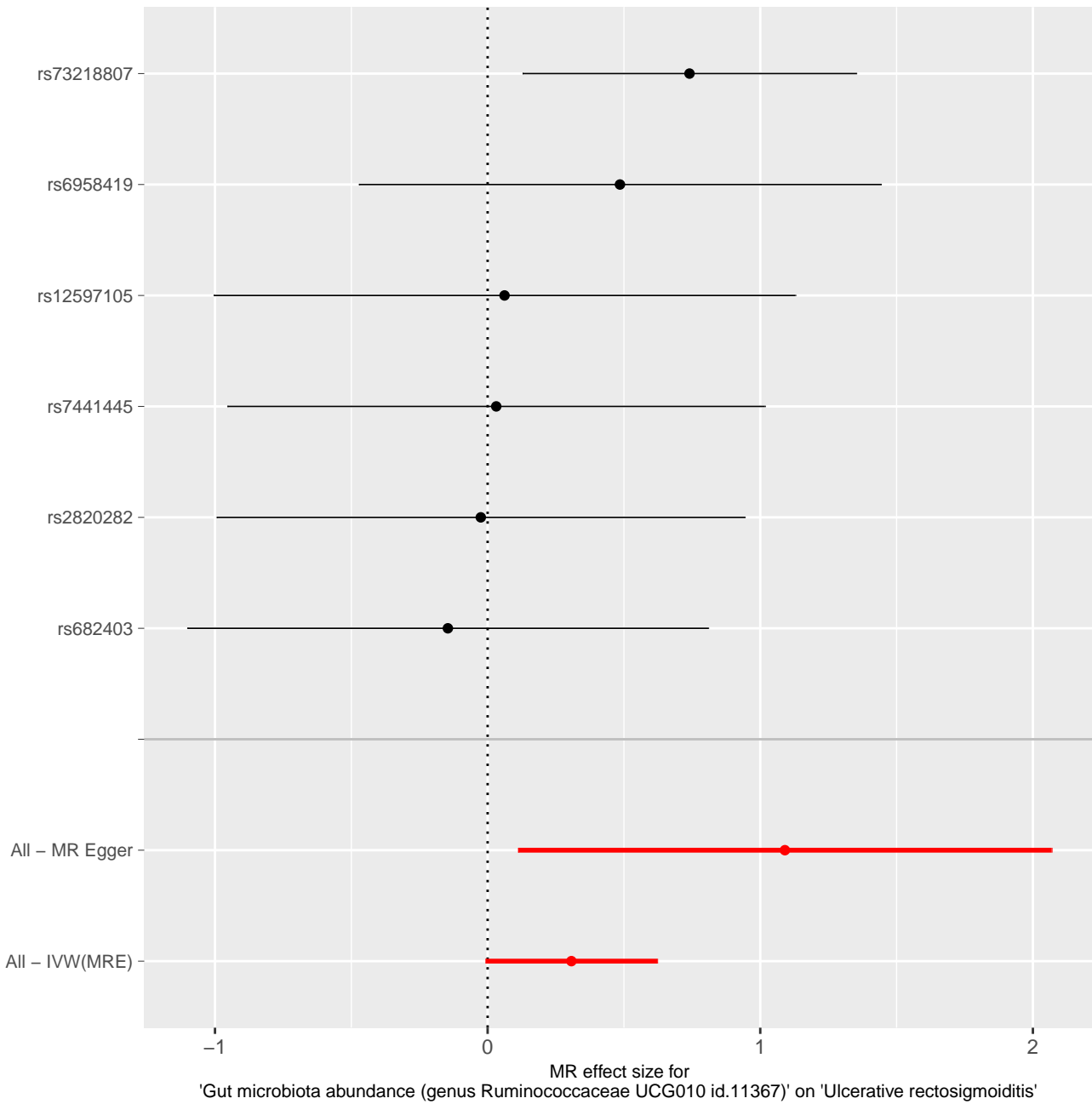
MR effect size for
'Gut microbiota abundance (genus Ruminococcaceae UCG002 id.11360)' on 'Ulcerative rectosigmoiditis'

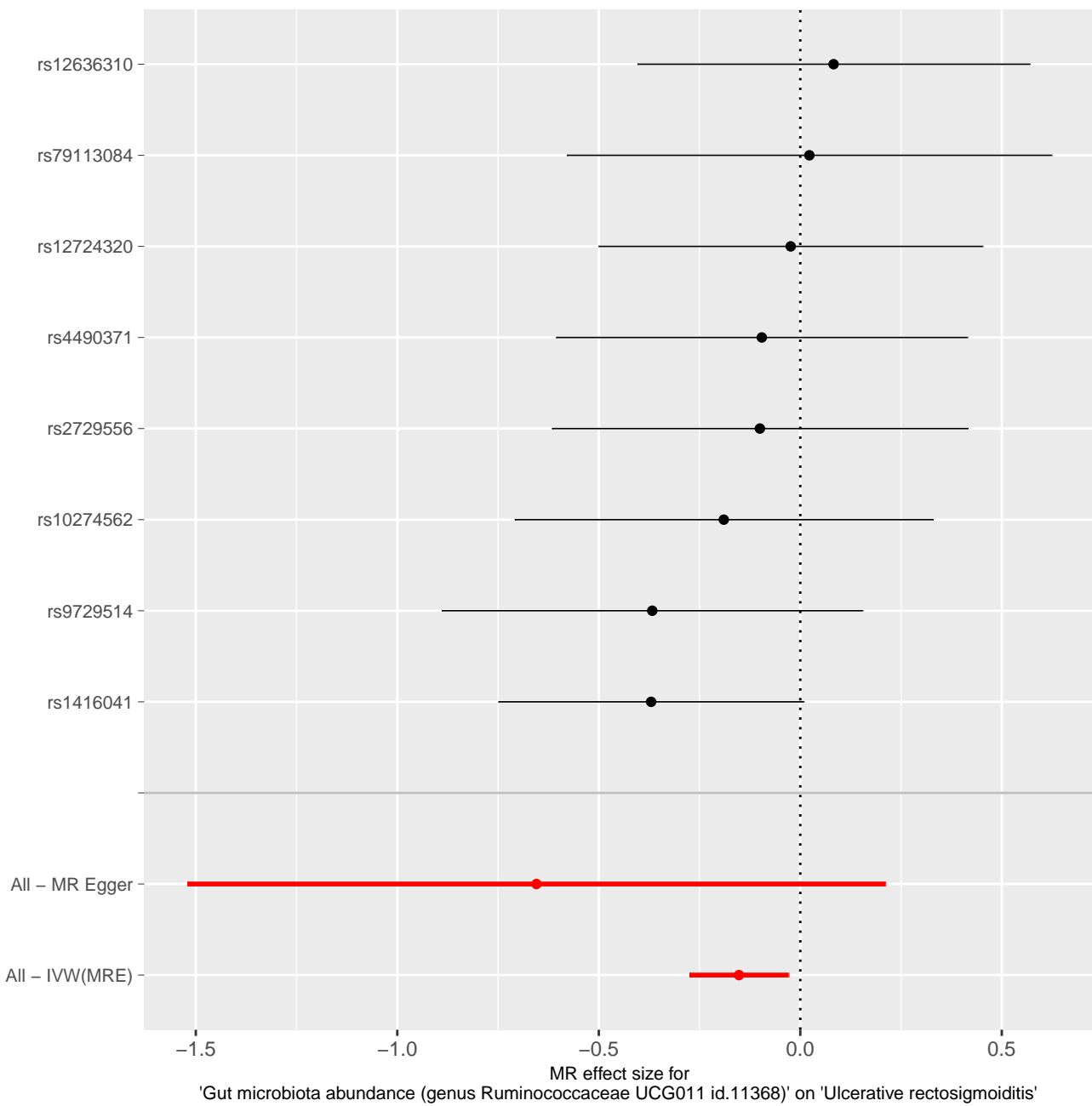












MR effect size for 'Gut microbiota abundance (genus Ruminococcaceae UCG011 id.11368)' on 'Ulcerative rectosigmoiditis'

