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# Erratum: Defining Mononuclear Phagocyte Subset Homology across Several Distant Warm-Blooded Vertebrates through Comparative Transcriptomics

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**Reason for Erratum:**
Due to a typesetting error, a misalignment in **Table 1** lead to the publication of incorrect information. The publisher apologizes for this error and the correct version of **Table 1** appears below.

This error does not change the scientific conclusions of the article in any way.

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**TABLE 1 | Conserved gene signatures for mammalian mononuclear phagocytic cell subsets.**

| Cell subset gene signatures | Genes conserved in 3/3 or 4/4 species[a] | Genes conserved in 2/3 or 3/4 species[b] |
|---|---|---|
| B cell | *TRAF5; SP140; RAD17; MEF2C; MBD4;* ***FCRL1***[c,d,e]*;* ***CD19*** | ***VPREB3****; RFX5;* ***PAX5****; BACH2; AFF3; SWAP70; PLEKHA2;* ***MS4A1****; DMXL1;* ***CR2****;* ***CD79B****; CD22;* ***BLK****; ELL3; STRBP;* ***EBF1*** |
| cDC vs. Mo/MP | *NAV1; MSI2;* ***HLA-DMB****;* ***FLT3****; BCL11A* | *RFX5; PLEKHA5;* ***HLA-DOA****; BCAT2; AFF3; FAM149A; APOBEC3H; UVRAG; SPINT2; PDXP;* ***HLA-DOB****;* ***CD74****; CD5; AP1S3;* ***HLA-DRA*** |
| cDC vs. pDC | *WDR41; WDFY3; TPM4;* ***TLR2****; SPI1; SNX14; SNX10; SERPINB1;* ***SAMHD1****; RIN3; REL; RAB32; NHSL1; NCOR2; NAV1; MARCKS; LYZ; LGALS3; KLF3;* ***JAK2****; ITGA5; IL4I1;* ***IL1B****;* ***IFNGR1****;* ***IFI30****;* ***ID2****;* ***ICAM1****;* ***HLA-DMB****; GCA; FGL2; F11R; ETV3; DOCK7; DENND4A; CXCL16; CLEC7A; CHSY1;* ***BATF3****; ATP2B1; ARRB1; ARHGAP22;* ***ANPEP****; AIM1; AIF1; AHR; ADAM8* | *YWHAH; TPCN1; TDRD7; SNX21; SLC7A10; SIPA1L3; RGS12; MYO1D; MRC2; METRNL; MEA1; LRRK2; LRRC8C; LOXL3;* ***HLA-DQB2****; HAVCR2; FGF17; EHF; DOK1; DGKH; ATXN1; ASB2; ARHGAP26; ACTR3; RNF144B; PLEKHO2; MYOF; LPCAT2; KANK1; FAM114A1; DENND5A; ZNF524; VASP; SULT1A1; SPRED1; SNX8; SH3BP1; SH3BGRL;* ***RELB****; RALB; RAC1; PTPN12; PLEKHO1; PIK3CB; PAK1; NR4A1; NAB2; LFNG; JUNB;* ***IFNGR2****; IER2; HFE; FAM49B; EPSTI1; EGR1; EFHD2; DHRS3; CTBP2; COTL1; CD74; CD63; CBFB; C9ORF72; C1ORF21;* ***BCL6****; BASP1; ANXA5; SR140; PKM2;* ***HLA-DRA****; RGS4; TMSB4X; GMIP; MAST2; CXCL9; DNAJA4; KIF14; MTUS1; RABGGTA; RTN1; SYNJ1; TBX3* |
| DCs vs. [Mo/MP and MoDC] | *MSI2; BCL11A* | |
| MoDC | *TPI1; NDUFV2;* ***FCGR2B****; CD200R1; ALDOA* | *RAB34;* ***PDCD1LG2****; CHCHD7; CCL17; CARM1; AUH;* ***VEGFA****; UBA3; TUBA1A; TSKU; TMEM159; SLC48A1; SIGMAR1; RNF181; PTGR1;* ***NOS2****; IKBIP; FAM162A; BHLHE40* |
| MoDC vs. Mo/MP | *TPI1; SLC2A1; SLAMF1; PRNP; PPA2; POLR1D; PLAU; PALLD; NDUFV2; NARF; MRPL4; IL1R2;* ***FCGR2B****; EGLN3; DGKA; CSNK2B; CISH; CD200R1; AVPI1; ALDOA; ADAMTSL4* | *ZNF747; ZNF219; WIBG; VDR; SLC45A4; ROGDI; RASSF7; RAB34; RAB33A; PDE6D;* ***PDCD1LG2****; PBX2; NAGS; KCNK6;* ***ICOSLG****; HRH1; GOLGA8B; GOLGA8A; ETHE1; ERCC6; DVL2; DGUOK; CLEC10A; CHN2; CHCHD7;* ***CD209****; CCNG2; CCL17; CARM1; C1ORF122; AUH; ANKRD37; ZEB1;* ***VEGFA****; UBA3; TUBA1A; TSKU; TMEM159; TCTEX1D2; STRA13; SPATA24; SNRNP27; SLC48A1; SIGMAR1; S1PR3; RNF181; RMND1; RAB7A; PTGR1; PIGU; PI4K2A; OST4; NSL1;* ***NOS2****; NAE1; MT1A; MORN4; LMF2; JKAMP; IKBIP; IFT46; HAUS4; GLTPD1; GATC; FAM162A; FAM13A; FAM134A; ESYT1; ERI2; EEPD1; DNLZ; DHRS11; DCTPP1; CENPW; BHLHE40; APOO; AKIP1; CD1B; CGREF1; NOSTRIN; OLFM4;* ***GAS6****; SLC27A3* |
| [Mo/MP and MoDC] vs. DCs | ***CEBPB****; CCDC93;* ***C5AR1*** | ***TLR8****; FTL; DOK3;* ***CD68*** |
| Mo/MP vs. cDC | ***TLR4****;* ***SOD2****; RBMS1;* ***LAMP2****; GLUL; FNDC3B; CYBB;* ***CEBPB****; CCPG1; CCDC93;* ***C5AR1*** | ***TLR8****; SNX27; RHOQ; OSTM1; KIF1B; FTL; DUSP6; DOK3;* ***CTSD****;* ***CTSB****;* ***CD68****; HERC5; IPMK; DPYD* |
| Mo/MP vs. MoDC | *WDR33; VPS13D; UBE2D2; TRA2A; STAG2; SFPQ; NSD1;* ***NFKB1****; NADK; ITPR1; CFLAR; ARFGEF1* | *ZNF407; VPS13C; USP31; SLC16A4; SKAP2; PRKCH; PPFIA1; PIAS2; MDN1; MAP3K5; LRRC8D; CHM; AKAP13; ACTR3; SFRS2IP; RAD51L1; NAT12; MYST3; CDC2L5; ZNF830; ZBED5; TPPP3; TMEM164; TGS1; TBC1D8B; SNRNP35; SMEK1; SLC38A10; SHISA2; RSRC2; REV1; RALGAPB; PWWP2A; PRRC2B; PBRM1; NLRC5; MOGS; MAP7D1; LUC7L3; LIMCH1; KDM4C; ISY1; IP6K1; HNRNPUL2; HNRNPU; HNRNPK; HNRNPH2; HNRNPH1; HNRNPD; HNRNPA1; FOXN3; FAM173B; FAM159B; ERVW-1; CELF2; C9; NUP210L; PDZK1; ALMS1; LAMB1; METTL3; PAIP1* |
| pDC | ***RUNX2*** | *LRP8; INPP4A; TSPAN13; SLC30A5; GPM6B* |
| pDC vs. cDC | *UBR2; UBE2H; TMED3;* ***TCF4****; TARBP1;* ***SYK****; STT3B; SPCS3; SNX5; SLC39A7; SIT1; SEMA4D; SEC61A1; SCYL3; SCAMP2; SAP30BP;* ***RUNX2****; RDH11; RASGRP2; RABAC1; PPAPDC1B; PGM3; PARN; PAG1; OGT; NUCB2; MSI2; MEF2C; LMAN2; IQCB1; IFT52; HBS1L; GPAM; GORASP2; FKBP2; FAM3C; EIF2AK3; DERL1; DDOST; DAD1; CYBB; COPA; CDC42SE2; CD4; CD164; BTRC;* ***BLNK****; BCL7A; ATP2A3; ATG5* | *ZXDC; VPS13A; UEVLD; TNRC6B; TMEM63A; TAF9B; TAF1A; SUSD1; STOML1; ST6GALNAC4; SSR2; SRPRB; SPG20; SLC38A6; SLC38A1; SLC25A36; SGCB; SERPINI1; SEC24C; SAP130; RAPGEF2; RALGPS1; RAB28; RAB11FIP2; PTAR1; PIK3AP1; OSTM1; NRP1; MYB; MGAT4A; MCOLN2; MCOLN1; LRP8; KIF13B; KIAA0226;* ***IRF7****; INPP4A; IMPACT; HIVEP1; FKBP8; FANCD2; FAM122B; DMTF1; CSTF1; CREB3L2; COBLL1; CBX4; CANX; ATG4D; ANKRD28; ANKIB1; AGBL3; AFF3; TPRG1L; RNF144A; IFI27L1; FAM65B; ELMOD3; DCAF7; CARS2; ZMYND11; YPEL3; USP24; TUBGCP6; TSPAN13; TRAM1; TOE1; TMEM138; TM9SF1; TCTA; SURF4; STAMBPL1; SSR3; SPCS2; SPATA13; SNX9; SLC7A5; SLC44A2; SLC30A5; SEPP1; SCAND1; SCAMP3; RHOH; RHBDF2; RHBDD1; REXO2; QDPR; PYCR2; PTPRCAP; PRMT7; POLD1; PEX5; NSUN3; MTMR9; LPGAT1; INTS7;* ***IFNAR1****; HM13; GRAP; GANAB; FNDC3A; FASTK; EXOC7; ELOF1; ELMOD2; CTCF; COPE; COMMD6; CNP; CIRBP; CDS2; CD79B;* ***CARD11****; C19ORF10; C16ORF80; C10ORF88; BTD; BET1; ARHGAP12; AHI1; WDR51B; SAPS3; MLF1IP; KIAA1370; CYBASC3; CEP110; CCDC111; ANUBL1; MME; PTPRS; ATF2; GPM6B; MON2; PPM1A; TM7SF3; TMCO1; UGCG; ZDHHC14; ZNF521; TMED10; PAIP1* |

*(Continued)*

**TABLE 1** | **Continued**

| Cell subset gene signatures | Genes conserved in 3/3 or 4/4 species[a] | Genes conserved in 2/3 or 3/4 species[b] |
|---|---|---|
| cDC2 | | *FCER1A* |
| cDC1 | ***XCR1***; *WDFY4; FNBP1; FLT3;* ***CADM1*** | *SNX22; GCET2* |
| cDC2 vs. [pDC and cDC1] | *TRPS1; STK24; SLC16A3;* ***SIRPA***; *SIGLEC8; S100A4; RIN2;* ***REL***; ***PILRA***; *NFAM1; NCF2; MAFB; LRP1; ITGAM;* ***IL1R2***; ***IL1B***; *IGSF6;* ***IFI30***; *FHL3; EPB41L3; DOCK4; DHRS3; CSF3R;* ***CSF1R***; *CLEC4A;* ***CD300A***; *C19ORF59; ADRBK2; TREM1* | *TNFRSF1B;* ***TLR8***; *TICAM2; STK10; SP2; SLFN12; SIGLEC9; SIGLEC7; RNASE2; PHF21A; LST1; LIMD2;* ***LILRB2***; ***LILRB1***; ***LILRA6***; ***LILRA3***; *IFITM2; GNGT2; GBP4; FAM111A; EMR1; DPP10; DENND1A;* ***DDX58***; *CDKN2B; CD300LF; CD300LB;* ***CD209***; *C10ORF11; ADAP1; CLEC6A; DAGLB; WDR45L; SIGLEC5; SFRS5; S100A12; PLEC1; MYST1;* ***MX2***; *MS4A8B; LRRC33; HSPA6; GK3P; GAPDH; FAM45B; CEBPD;* ***CD1E***; ***CD1B***; ***FCER1A***; *KSR1;* ***OAS2***; *PTGER3* |
| cDC1 vs. [pDC and cDC2] | ***XCR1***; *WDFY4; ST3GAL5; RAB32; PPT1; PPA1; LRRC1; KIAA1598; FNBP1;* ***FLT3***; *CALM1;* ***CADM1*** | *SNX22; PPAP2A; PLEKHA5; GRAMD2; DENND1B; CLEC1A; ATXN1; FAM114A1; HEPACAM2; PI4K2A; PLEKHO2; WDR91; TRIO; RALB; PKP4; PDLIM7; G3BP2;* ***BCL6***; *ATPIF1; GCET2; BRWD2; FGD6; MYO9A* |
| ncMo vs. cMo[f] | *ACAT2; ACE; ACOT9; ADRBK2; ANKRD42; APOA2; ASB2; BDKRB2; BGLAP; C1ORF112; C1ORF56; C20ORF112; CAPZB; CBX4; CD4;* ***CD83***; *CDH24; CHD5;* ***CSF1R***; *CYP2R1; DCBLD1; DDB2; DDIT4; DLGAP4; FBP1; GABBR1; GLMN; GNE; GNPNAT1; GPT; GRHPR; HEY1; HN1;* ***IL12RB1***; *IL17A; IL2RG; KCNMA1; KCTD11; KNDC1; LMX1B; LUZP1; MAFF; MPZL1; MUTYH; MYOD1; NCAPH2; NCOR2; NFKBIA; NPAS2; NUB1; PCK1; PDCD4; PGR; PITPNM1; PLEKHH1; PMF1; PMVK; POLR3H; RAB25; RAD52; RFC5; RHOF; RSAD1; RWDD3; SECISBP2; SERPINA1; SH2D3C; SIRT5; SLC37A1; SMS; ST3GAL1; ST3GAL5; TBC1D8; TCF7L2; TNNC1; U2AF1L4; UNG; WDR76* | |
| cMo vs. ncMo[e] | *AACS; ABHD5; AGTPBP1; ALDH2; ALOX5AP; ANXA1; AOAH; ARL8B; ATP6V1A; ATP6V1B2; ATP6V1C1; AUH; B4GALT1; C19ORF59; C5ORF15; CCR1;* ***CD164***; *CD84; CETN2; CLTA; COPB2; CSF3R; CYP27A1; DCLRE1A; DNAJC10; ECE1; EHD4;* ***EIF2AK2***; *EIF2AK3; ENSA; ENTPD7; ERP29; EXOC5; F13A1; F5; FAM102B; FAM63A; FBXL5; FBXO9; FN1; GBE1; GNA12; GNPAT; GSN; GYS1; HMGB2;* ***IL1R2***; ***IL1RN***; *ITM2B; KEAP1; LACTB; LCN2; LEO1; LMAN1; LMNB1;* ***LYZ***; *MBD5; MBIP; MGA; MPP1; NHLRC2; NISCH; NKRF; NPC1; NSF; NUCB2; PAM; PARP8; PDE2A; PGD; PLCB1; PNPLA8; PON2; PREPL; PRKAR1A; PRUNE; PSMA1; PSTPIP1; PUM2; PXK; PYGL; RAB27A; RAB3D; RABGAP1L; RARS; RHOT1; RMI1; RNF130; RPGR; RSC1A1;* ***S100A8***; *SCRN3; SCYL1; SDCBP; SEC22C; SELL; SENP5; SERPINB1; SHB; SIGLEC1; SLC16A7; SLC25A44; SLC35B3; SLC39A9; ST8SIA4; TBC1D2; TEX2; TGM1; TM6SF1; TMEM161B; TMEM71; TPCN1; TREML2; TRIP11; TSHZ1; UBE4A; UMPS; USP10; UXS1; VAPB; VNN3; VPS37B; WDTC1; XBP1; ZMYM4* | |

*[a]Genes conserved in 4/4 species, or in 3/3 species for cDC2, cMo, and ncMo since only three species could contribute to the analysis.*

*[b]Genes conserved in 3/4 species or 2/3 species.*

*[c]Genes in bold were previously demonstrated to play a significant role in the development or functions of the population of interest.*

*[d]Underlined genes were annotated as located in "plasma membrane" according to Ingenuity Pathway Analysis.*

*[e]Genes highlighted in gray have been previously identified as signatures genes for the corresponding mouse and human cell populations in our earlier study (1).*

*[f]Signature genes of the relative cMo vs. ncMo and of the ncMo vs. cMo signatures were provided only for the 3/3 species selection since the gene lists for the 2/3 species selection encompassed hundreds of genes.*

## REFERENCE

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