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Erratum: Pertinence of glioma and single nucleotide polymorphism of TERT, CCDC26, CDKN2A/B, and RTEL1 genes in glioma: a meta-analysis

Frontiers Production Office*

Frontiers Media SA, Lausanne, Switzerland

KEYWORDS

glioma, single nucleotide polymorphism, risk, meta-analysis, genetic model

An Erratum on:

Pertinence of glioma and single nucleotide polymorphism of TERT, CCDC26, CDKN2A/B and RTEL1 genes in glioma: a meta-analysis

by Wu Y, Zhou J, Zhang J, Tang Z, Chen X, Huang L, Liu S, Chen H and Wang Y (2023) *Front. Oncol.* 13:1180099. doi: 10.3389/fonc.2023.1180099

Due to a production error, there was a mistake in the name of Figure 2 as published. The figure part labels read as ‘Correlation of glioma and rs6010620 SNP’. The correct text is ‘Correlation of glioma and rs2736100 SNP’. The corrected Figure 2 appears below.

Due to a production error, there was a mistake in the name of Figure 4 as published. The figure part labels read as ‘Correlation of glioma and rs2736100 SNP’. The correct text is ‘Correlation of glioma and rs4977756 SNP.’ The corrected Figure 4 appears below.

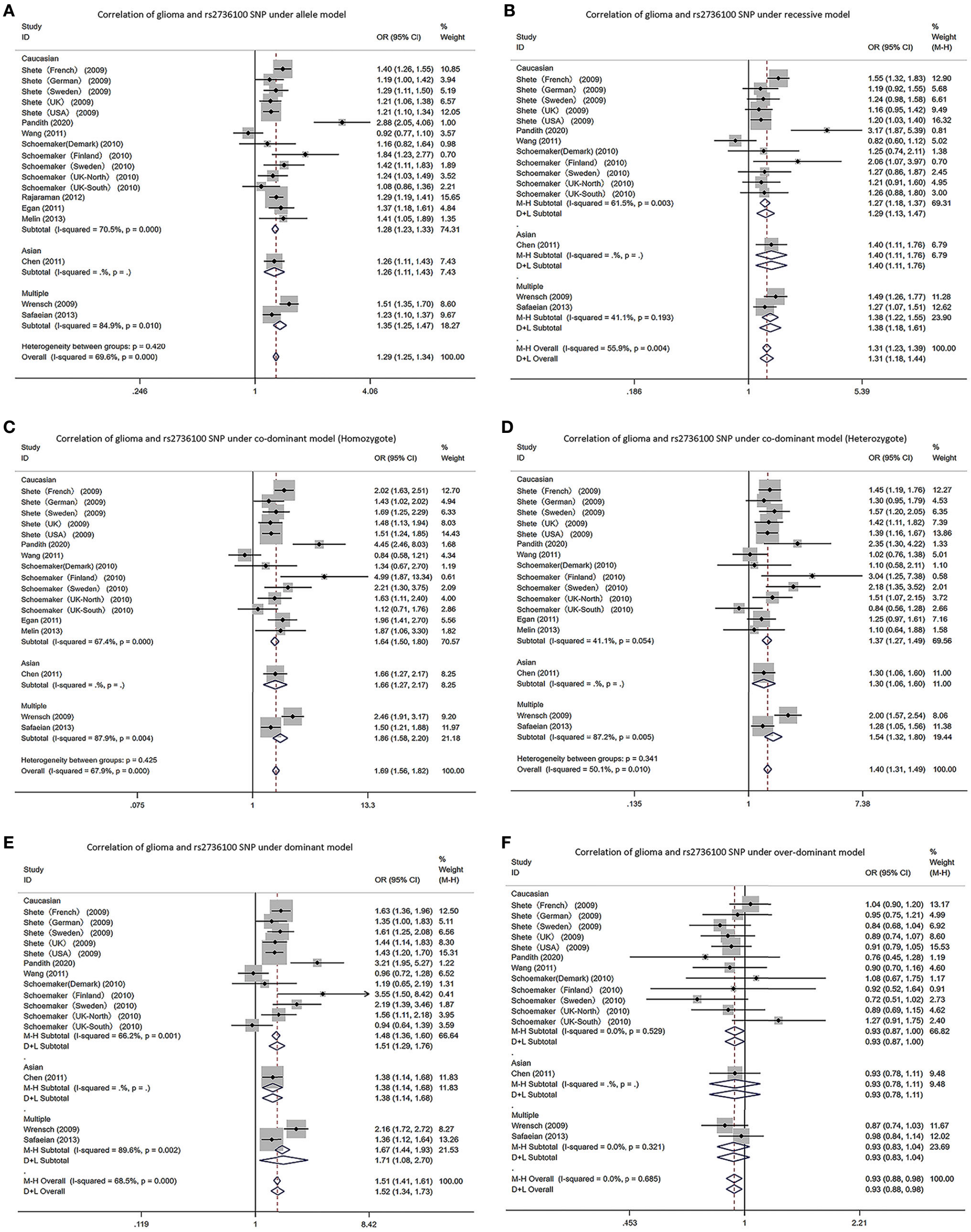

FIGURE 2
Forest plots of meta-analyses for correlation of glioma and rs2736100 SNP under all models. **(A)** allele model; **(B)** recessive model; **(C)** co-dominant model (Homozygote); **(D)** co-dominant model (Heterozygote); **(E)** dominant model; **(F)** over-dominant model.

Due to a production error, there was a mistake in the name of Figure 5 as published. The figure part labels read as 'Correlation of glioma and rs4977756 SNP'. The correct text is 'Correlation of glioma and rs6010620 SNP.' The corrected Figure 5 appears below.

The publisher apologizes for this mistake.

The original version of this article has been updated.

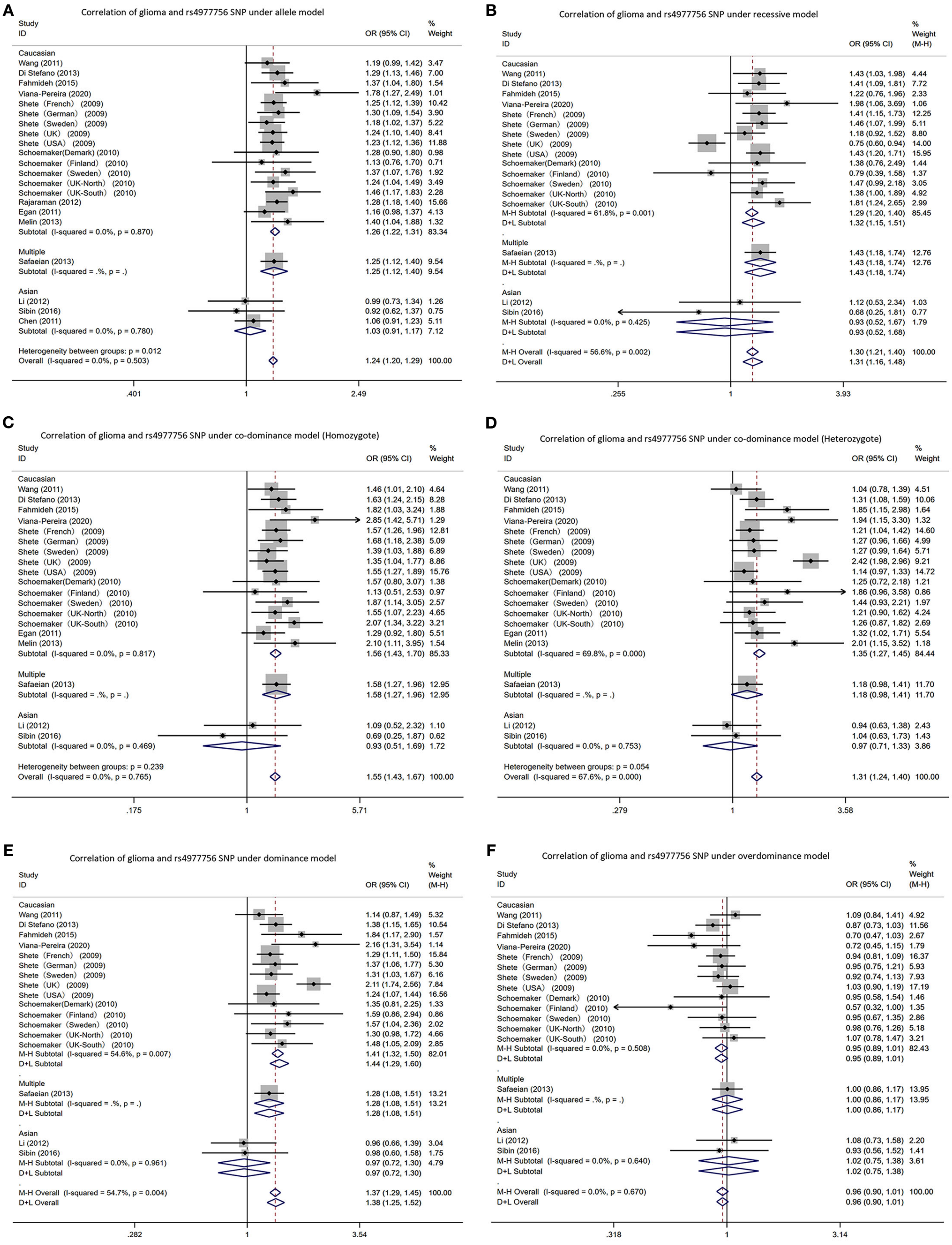

FIGURE 4

Forest plots of meta-analyses for correlation of glioma and rs4977756 SNP under all models. **(A)** allele model; **(B)** recessive model; **(C)** co-dominant model (Homozygote); **(D)** co-dominant model (Heterozygote); **(E)** dominant model; **(F)** over-dominant model.

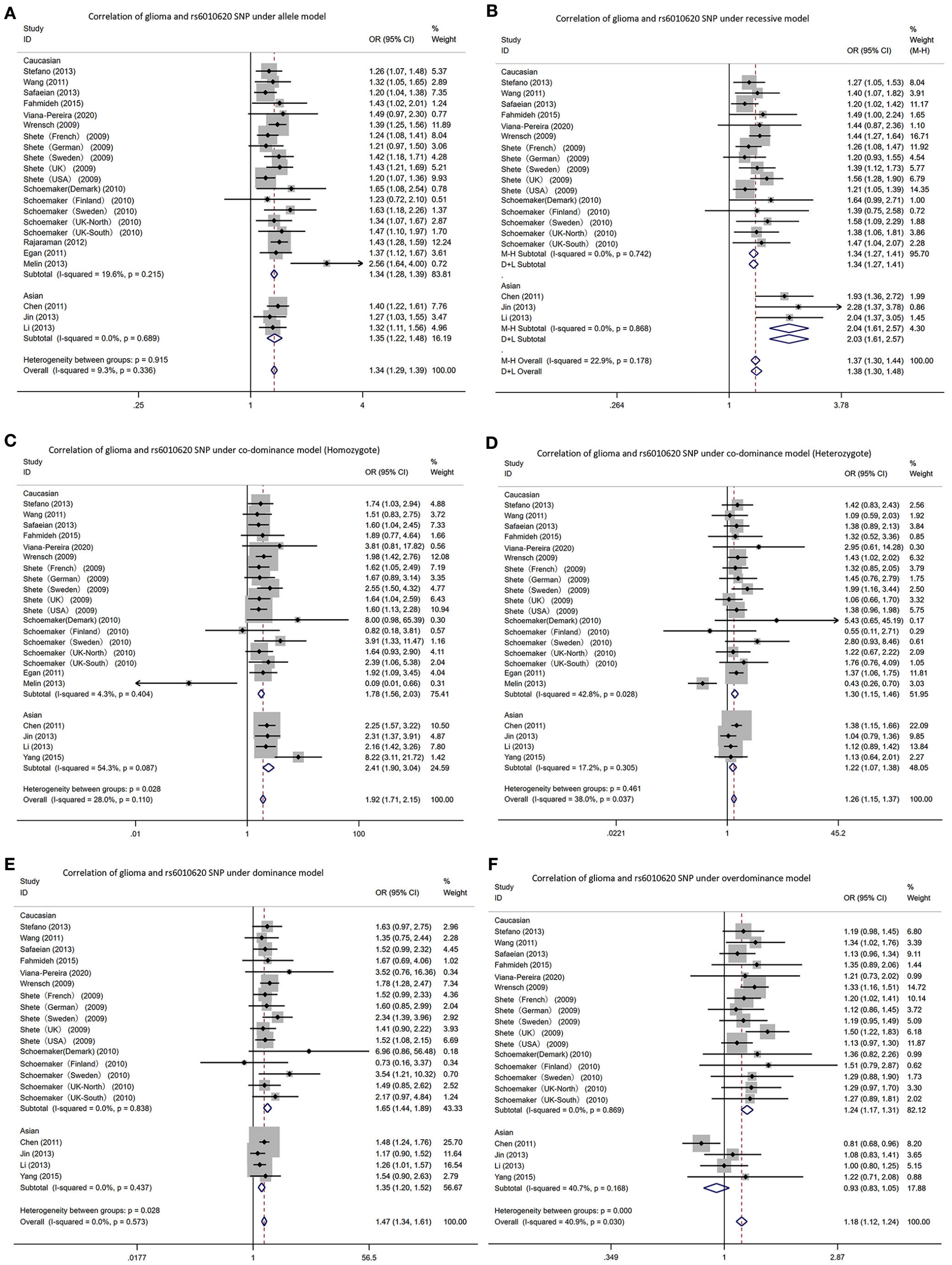

FIGURE 5

Forest plots of meta-analyses for correlation of glioma and rs6010620 SNP under all models. **(A)** allele model; **(B)** recessive model; **(C)** co-dominant model (Homozygote); **(D)** co-dominant model (Heterozygote); **(E)** dominant model; **(F)**. over-dominant model.