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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

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## KEYWORDS

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

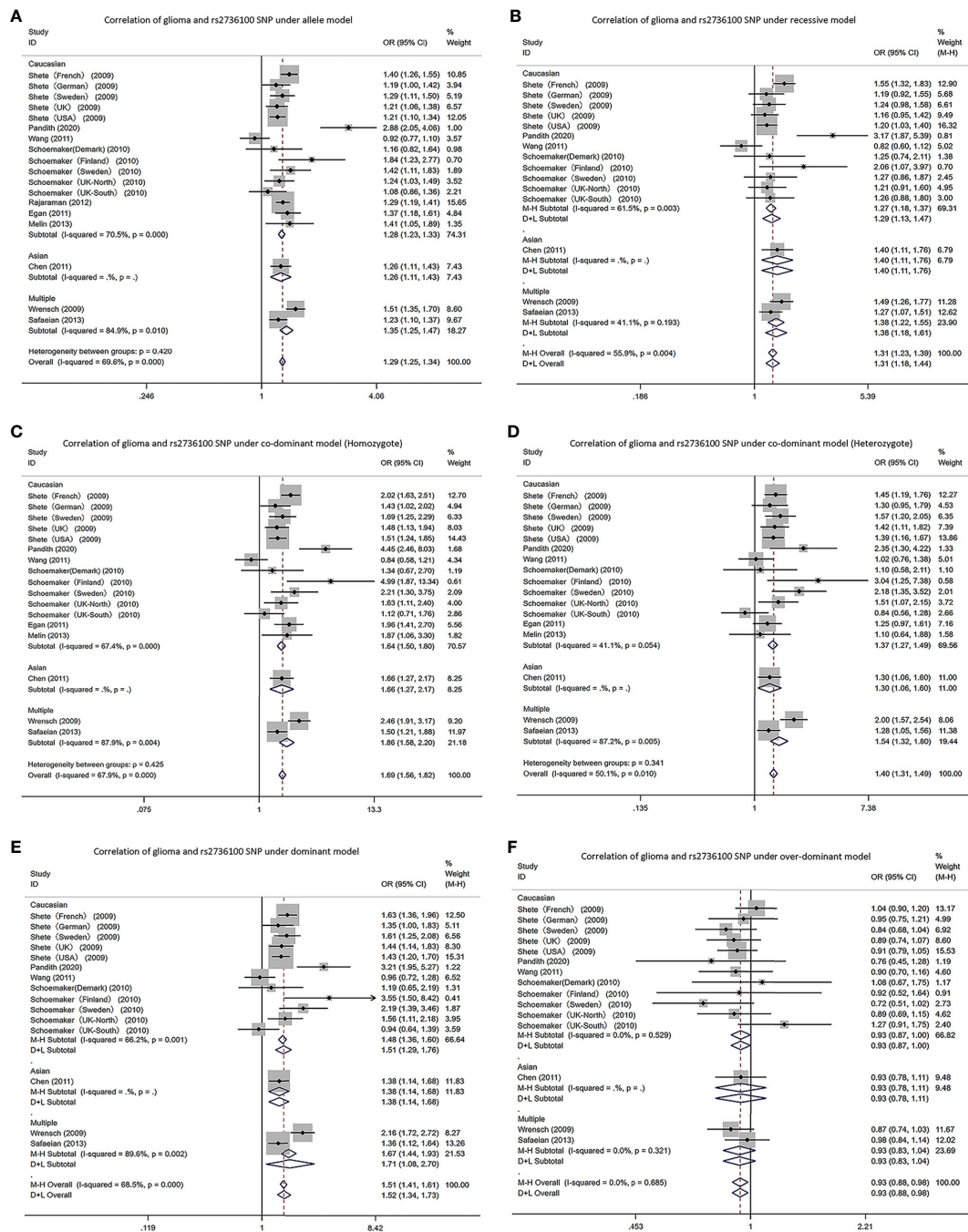
## An Erratum on:

[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.

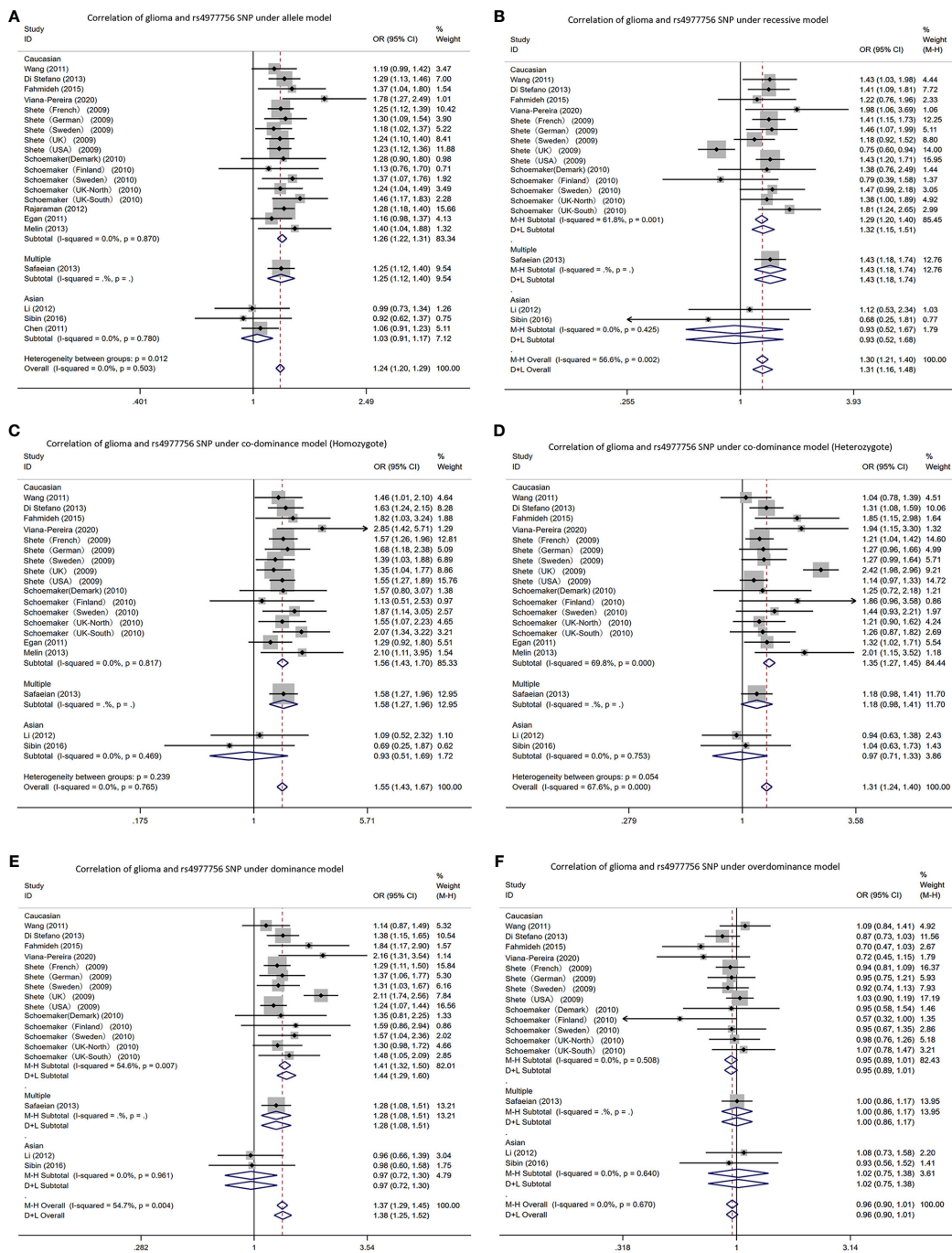


FIGURE 4 Forest plots of meta-analyses for correlation of glioma and rs4977756 SNP under all models. (A) allele model; (B) recessive model; (C) co-dominant model (Homozygote); (D) co-dominant model (Heterozygote); (E) dominant model; (F) over-dominant model.

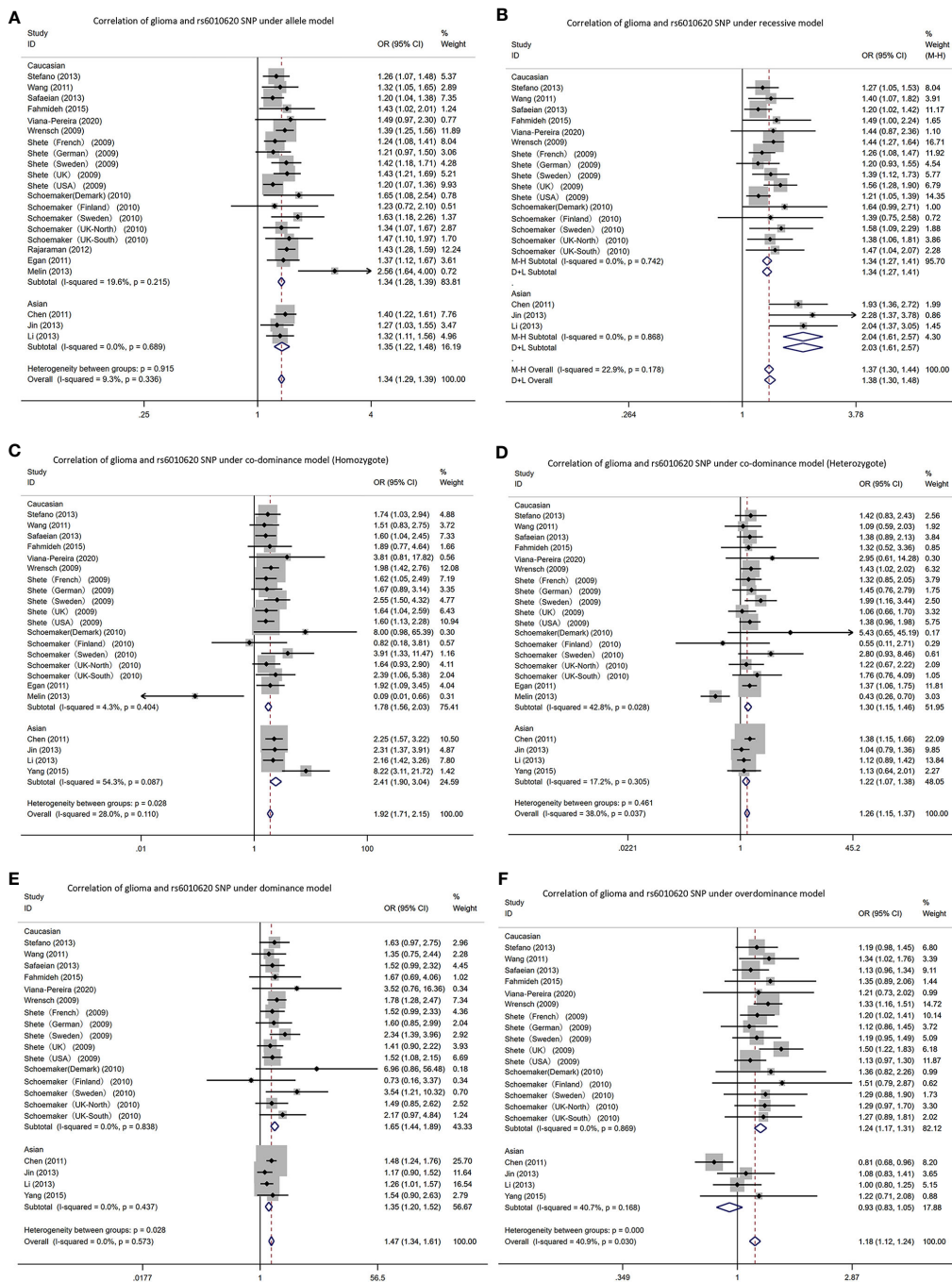


FIGURE 5

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.