

CORRECTION

# Correction: A Newfound Association between *MDC1* Functional Polymorphism and Lung Cancer Risk in Chinese

The *PLOS ONE* Staff

There are errors in the values of the fourth and sixth sentences in the Abstract. The correct fourth sentence is: We found the SNP rs4713354A>C that is located in the 5'-untranslated region of *MDC1* was significantly associated with lung cancer risk in both populations ( $P = 0.001$ ), with an odds ratio as 1.33(95% confidence interval = 1.14–1.55) for the rs4713354C (CA+CC) genotypes compared to the rs4713354AA genotype. The correct sixth sentence is: The gene-based analysis rested with these SNPs suggested the *MDC1* as a susceptible gene for lung cancer ( $P = 0.057$ ).

## Reference

1. Wang B, Zhang L, Qiu F, Fang W, Deng J, et al. (2014) A Newfound Association between *MDC1* Functional Polymorphism and Lung Cancer Risk in Chinese. *PLoS ONE* 9(9): e106794. doi:[10.1371/journal.pone.0106794](https://doi.org/10.1371/journal.pone.0106794) PMID: [25198518](https://pubmed.ncbi.nlm.nih.gov/25198518/)



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