

CORRECTION

Correction: Bioinformatics Analysis of the Effects of Tobacco Smoke on Gene Expression

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There is an error in the fifth sentence of the Abstract. The correct sentence is: In total, 288 up-regulated DEGs and 106 down-regulated DEGs were identified.

There is an error in the third sentence of the Results section under the heading “GO and pathway enrichment analyses of DEGs.” The correct sentence is: Specifically, the DEG distribution in the pathway of Metabolism of xenobiotics by cytochrome P450 was shown in Fig 2.

Reference

1. Cao C, Chen J, Lyu C, Yu J, Zhao W, Wang Y, et al. (2015) Bioinformatics Analysis of the Effects of Tobacco Smoke on Gene Expression. PLoS ONE 10(12): e0143377. doi: [10.1371/journal.pone.0143377](https://doi.org/10.1371/journal.pone.0143377) PMID: [26629988](https://pubmed.ncbi.nlm.nih.gov/26629988/)



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