

CORRECTION

Correction: Preferential Allele Expression Analysis Identifies Shared Germline and Somatic Driver Genes in Advanced Ovarian Cancer

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Reference

1. Halabi NM, Martinez A, Al-Farsi H, Mery E, Puydenus L, Pujol P, et al. (2016) Preferential Allele Expression Analysis Identifies Shared Germline and Somatic Driver Genes in Advanced Ovarian Cancer. *PLoS Genet* 12(1): e1005755. doi:[10.1371/journal.pgen.1005755](https://doi.org/10.1371/journal.pgen.1005755) PMID: [26735499](https://pubmed.ncbi.nlm.nih.gov/26735499/)



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