

# Identification of CD15 as a Marker for Tumor-Propagating Cells in a Mouse Model of Medulloblastoma

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Following publication of this manuscript, the authors discovered some text that had been erroneously included in the Supplemental Experimental Procedures. The section “Microarray Preprocessing and CD15+ Signature Development” (pp. 3 to 4) was adapted from a description of the methods used in another study, and the following text was mistakenly included in the current description:

“To identify differentially expressed genes, Student’s t tests were applied to the mean expression of samples after at least 12 hours of androgen exposure as compared to the mean expression of samples without androgen exposure. To control for the false discovery rate (FDR) of multiple tests, the Benjamini-Hochberg step-up procedure was applied to the nominal p values (Benjamini and Hochberg, 1995).”

Such analysis was not used for the current data set (which did not include androgen treatment) and, thus, should have been deleted. A corrected version of the online supplemental data is now available at [www.cell.com/cancer-cell/supplemental/S1535-6108\(08\)00435-2](http://www.cell.com/cancer-cell/supplemental/S1535-6108(08)00435-2).