



## Corrigendum: Machine learning identifies activation of RUNX/AP-1 as drivers of mesenchymal and fibrotic regulatory programs in gastric cancer

Milad Razavi-Mohseni, Weitai Huang, Yu A. Guo, et al.

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## Corrigendum

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**Corrigendum: Machine learning identifies activation of RUNX/AP-1 as drivers of mesenchymal and fibrotic regulatory programs in gastric cancer**

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Following genotyping analysis from ATAC-seq and RNA-seq reads, the authors found the ATAC-seq sample labeled SNU484 was mislabeled, and it was a duplicate of AGS ATAC-seq, which is now relabeled AGS\_r2 in the corrected text, figures (Figs. 1–3, 6, 7), and Supplemental Material (Supplemental Figs. S6–S12, S14–S16, S21–S23; Supplemental Tables S1, S2). This relabeling resolves the inconsistency between SNU484 ATAC-seq and RNA-seq noted in the original article, and strengthens some *P*-value and RNA correlations slightly, as noted in the corrected text. SNU484 is now classified as a Mes-like cell line based on RNA-seq. None of the main conclusions of the article are changed. The only notable exception is in Figure 2A, where SNU484 had been an outlier but now is consistent with the rest of the data.

Both the PDF and full-text HTML files online have been corrected.

The authors apologize for any confusion this may have caused.

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