

Author Correction: Epigenetic scarring of exhausted T cells hinders memory differentiation upon eliminating chronic antigenic stimulation

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In the originally published version of this Article, there were errors in Fig. 6b, main text and Extended Data Fig. 8a caption.

Specifically, in Fig. 6b, the labels at top of the heatmap, now reading left-to-right as “REC-T_{EX}” and “T_{EX}” were in reverse order. In the Results section, under heading “Epigenetic scars in REC-T_{EX} limit core memory circuits,” second paragraph, fourth sentence, the text now reading “Of the 140 OCRs that were more accessible in REC-T_{EX} compared to T_{MEM} and T_{EX} (Fig. 5c), only a handful (for example, *Zfp36l2*) showed increased expression of the associated gene in REC-T_{EX} compared to T_{EX} (Extended Data Fig. 8a)” overwrote the previous sentence (Of the 140 OCRs that were more accessible in REC-T_{EX} compared to T_{MEM} (Fig. 5c), only a handful (for example, *Zfp362*) showed increased expression of the associated gene (Extended Data Fig. 8a)). In Extended Data Fig. 8a caption, the sentence now reading “Genes from scRNAseq with significantly different expression between REC-T_{EX} and T_{EX} that are associated with more accessible OCRs in REC-T_{EX} compared to both T_{EX} and T_{MEM}” overwrote the previous caption (a, Genes from scRNAseq with significantly different expression between REC-T_{EX} and T_{MEM} that also had associated significant differences in the chromatin accessibility).

The changes have been made to the online version of the Article.

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