

# Monitoring CAR T cell generation with a CD8-targeted lentiviral vector by single-cell transcriptomics

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In the originally published version of this article, the authors noticed a misspelling error in the R environment script used to generate the left Venn plot in Figure 3C. This mistake affected some of the numbers listed. The corrected Figure 3 is shown here. The two sentences describing Figure 3C in the main text should read as follows: “The majority of the differentially expressed genes, 130 out of 155 either up- or downregulated, were identified when comparing untransduced with  $CAR^{high}$  cells (Figure 3C, left). From these, 104 genes were shared between the comparisons of untransduced with either  $CAR^{high}$  or  $CAR^{neg/low}$  cells, 33 genes were shared in all of the comparisons, including  $CAR^{neg/low}$  with  $CAR^{high}$ .” The overall conclusions drawn are not affected. The authors regret this error.



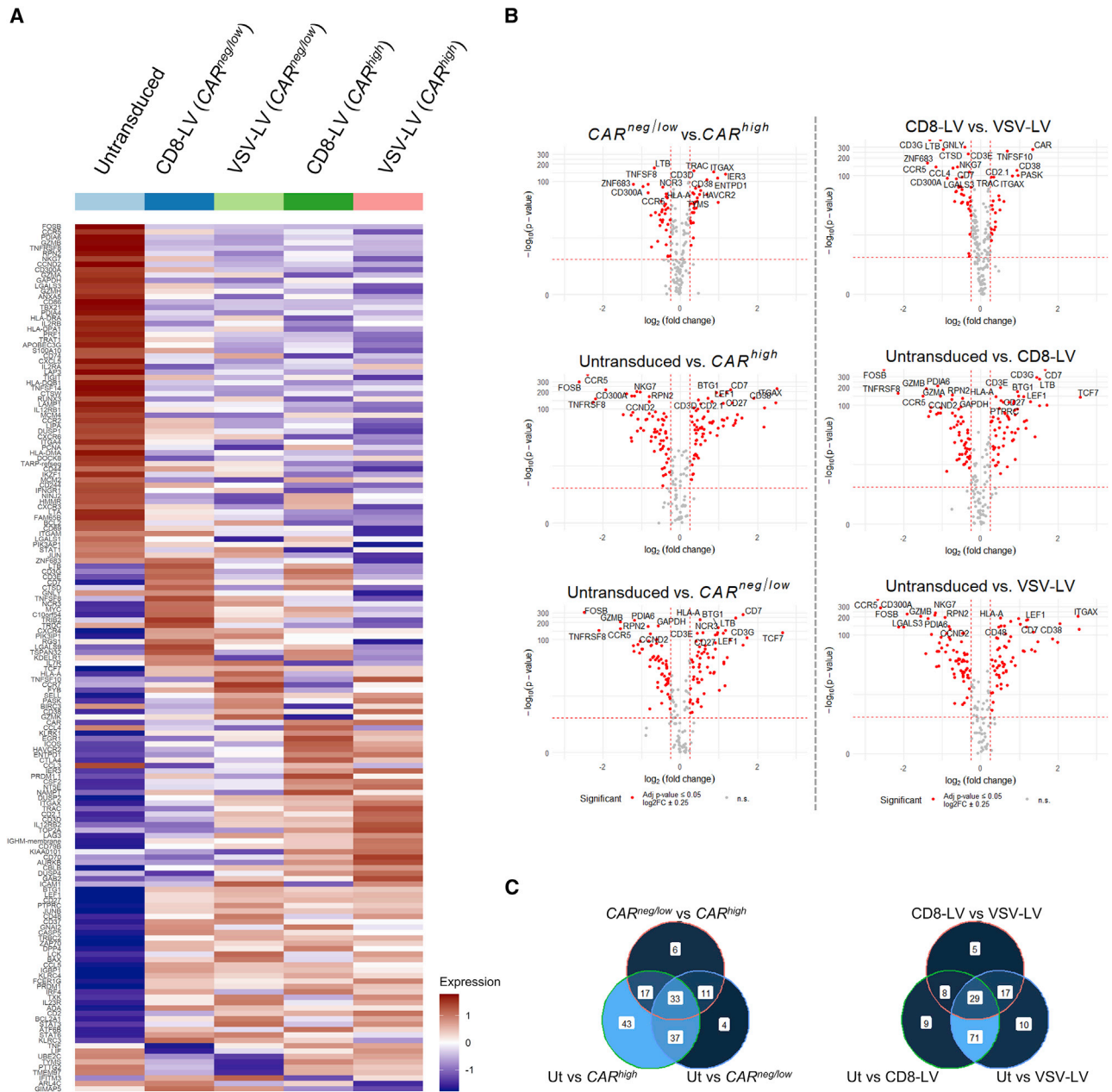
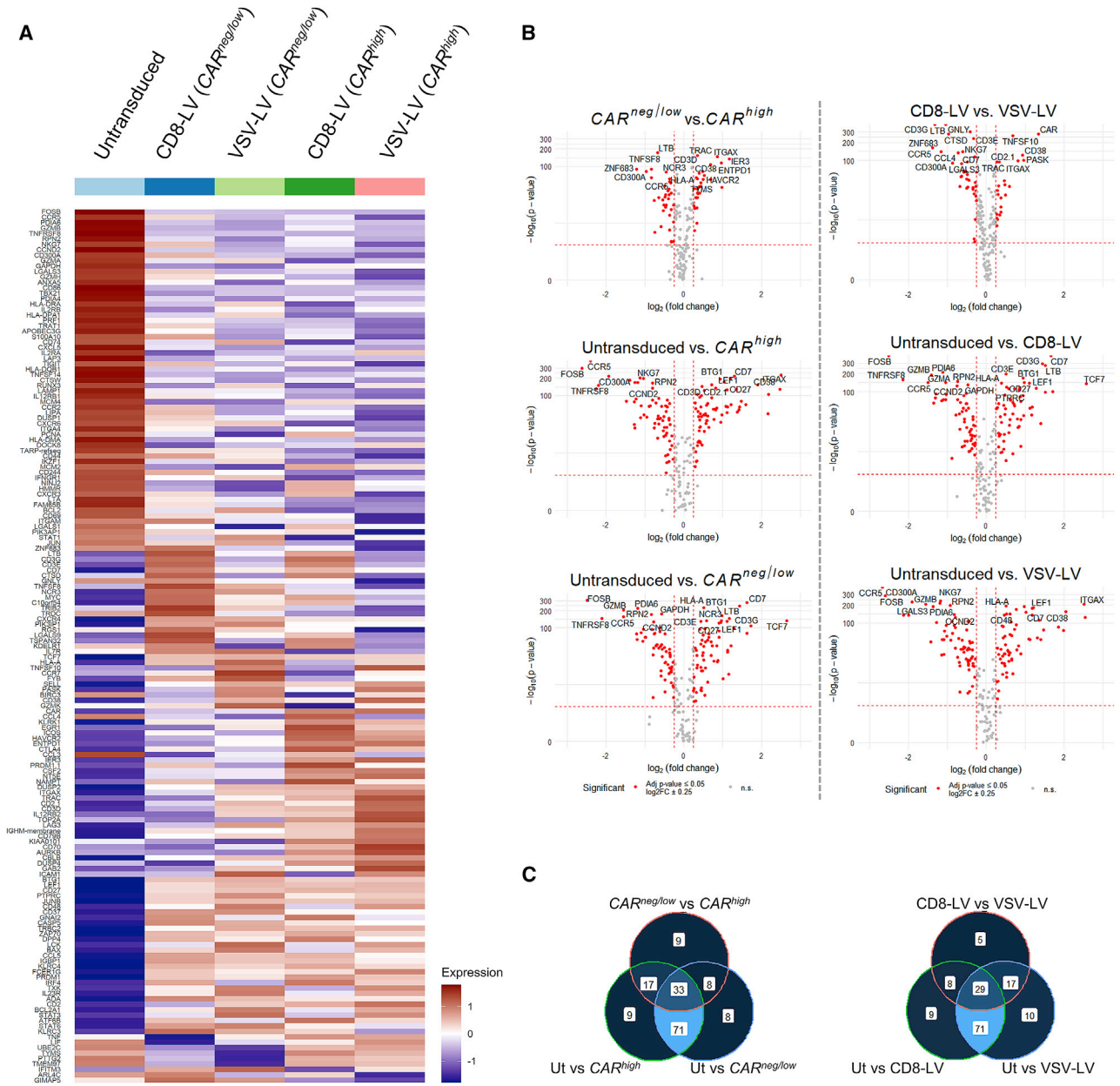


Figure 3. Differentially expressed genes in the subpopulations (original)



**Figure 3. Differentially expressed genes in the subpopulations (corrected)**