

**Supplemental Figure 1. The level of ACE2 mRNA in common blood cells in the transcriptome datasets from Monaco scaled dataset and Schmiedel dataset.**

(A) Bar plot of the transcript abundance of ACE2 across common blood cells obtained from the quantification of *Monaco* scaled dataset. The level of ACE2 mRNA was represented by the mean pTPM; The data no labelling meant no ACE2 expression in corresponding samples;

(B) Bar plot of the transcript abundance of ACE2 across common blood cells obtained from the quantification of *Schmiedel* dataset. The level of ACE2 mRNA was represented by the mean pTPM; The data no labelling meant no ACE2 expression in corresponding samples.

