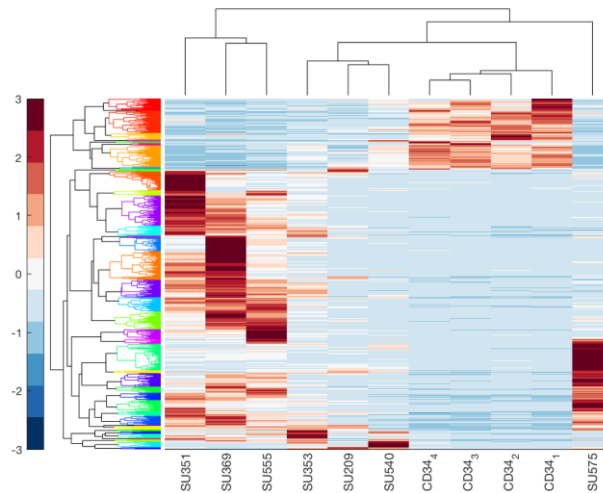
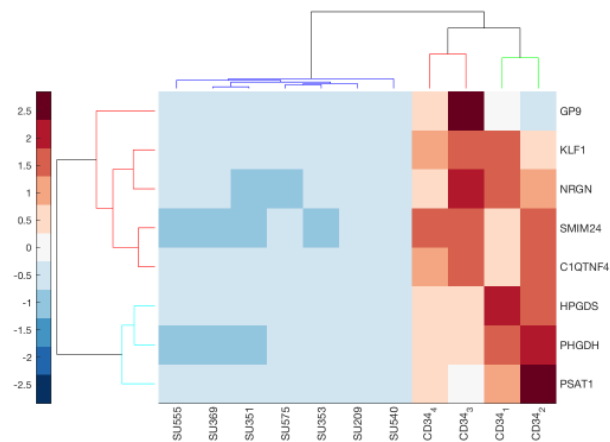


Fig. S17.



B.



RNA-Seq of purified populations of human AML blasts and CB-CD34⁺ cells shows an inflammatory transcriptome signature in AML. (A) Heatmap representation of unsupervised hierarchical clustering of the differentially expressed transcripts between purified populations of primary human AML blasts (n=7) and normal CB-CD34⁺ cells (n=4). (B) Heatmap of the transcripts whose expression was down regulated at least 8 fold ($p < 0.00005$) in AML blasts compared to normal CB-CD34⁺ cells.