

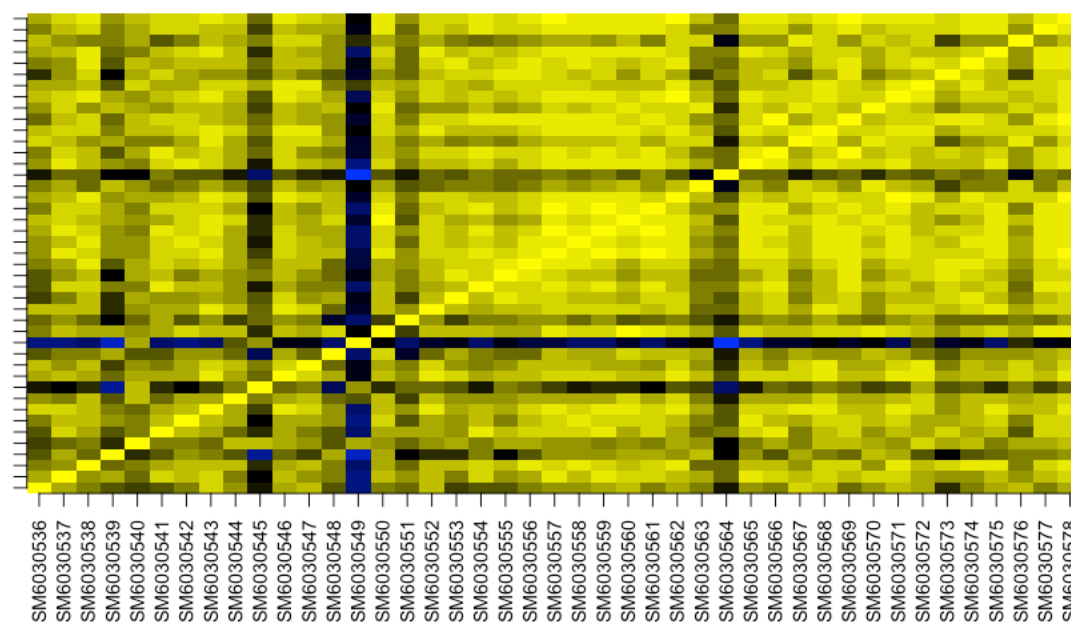
Figures and data from simple gene expression analysis

Citation for data source:

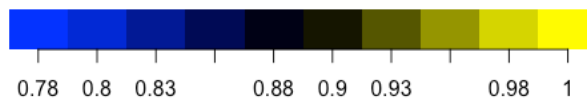
Shao L, Pelayo A, Shi R, Ma J, Liu H, Cai Y, Prochazkova M, Somerville RP, Panch SR, Shah NN, Stroncek DF, Jin P. Identification of genomic determinants contributing to cytokine release in immunotherapies and human diseases. *J Transl Med.* 2022 Jul 28;20(1):338. doi: 10.1186/s12967-022-03531-3. PMID: 35902861

Link to GEO page with data: <https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE200296>

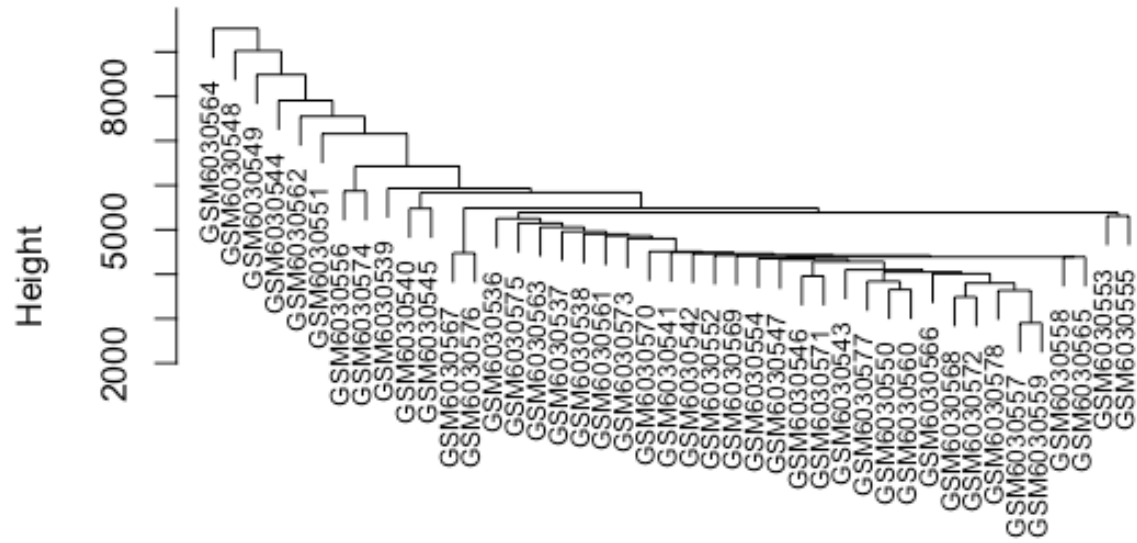
Correlation heat map of samples in CD22 CAR T-cell product dataset



(y-axis is the same as x-axis)

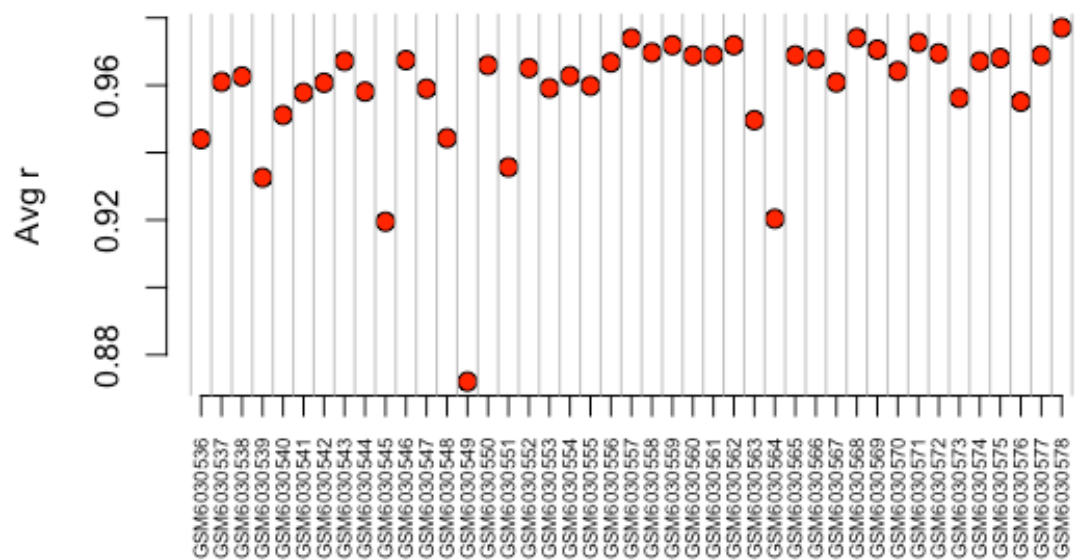


Cluster Dendrogram



dat.dist
hclust (*, "single")

Avg correlation of CD22 CAR T-cell product samples



Differentially expressed genes with p-value < 0.05

```
> pv[pv < 0.05]
      CCL5      CD8A      CTSD      NKG7      PFKFB4      TNFSF9      TRAV19      TRBV7-4
0.028857025 0.014039904 0.023625708 0.030171948 0.001353091 0.029974824 0.024916979 0.044381774
```

Fold change of those genes:

```
> fold[names(pv[pv<0.05])]
      CCL5      CD8A      CTSD      NKG7      PFKFB4      TNFSF9      TRAV19      TRBV7-4
0.6888487 0.5667767 0.2315774 0.5473447 -1.2178337 0.5464836 0.3301870 -0.3548841
```

Conclusion:

This relatively simple analysis turned up the main gene this paper is about PFKFB4. It also turned up some other genes that would be interesting to explore further with laboratory experiments.