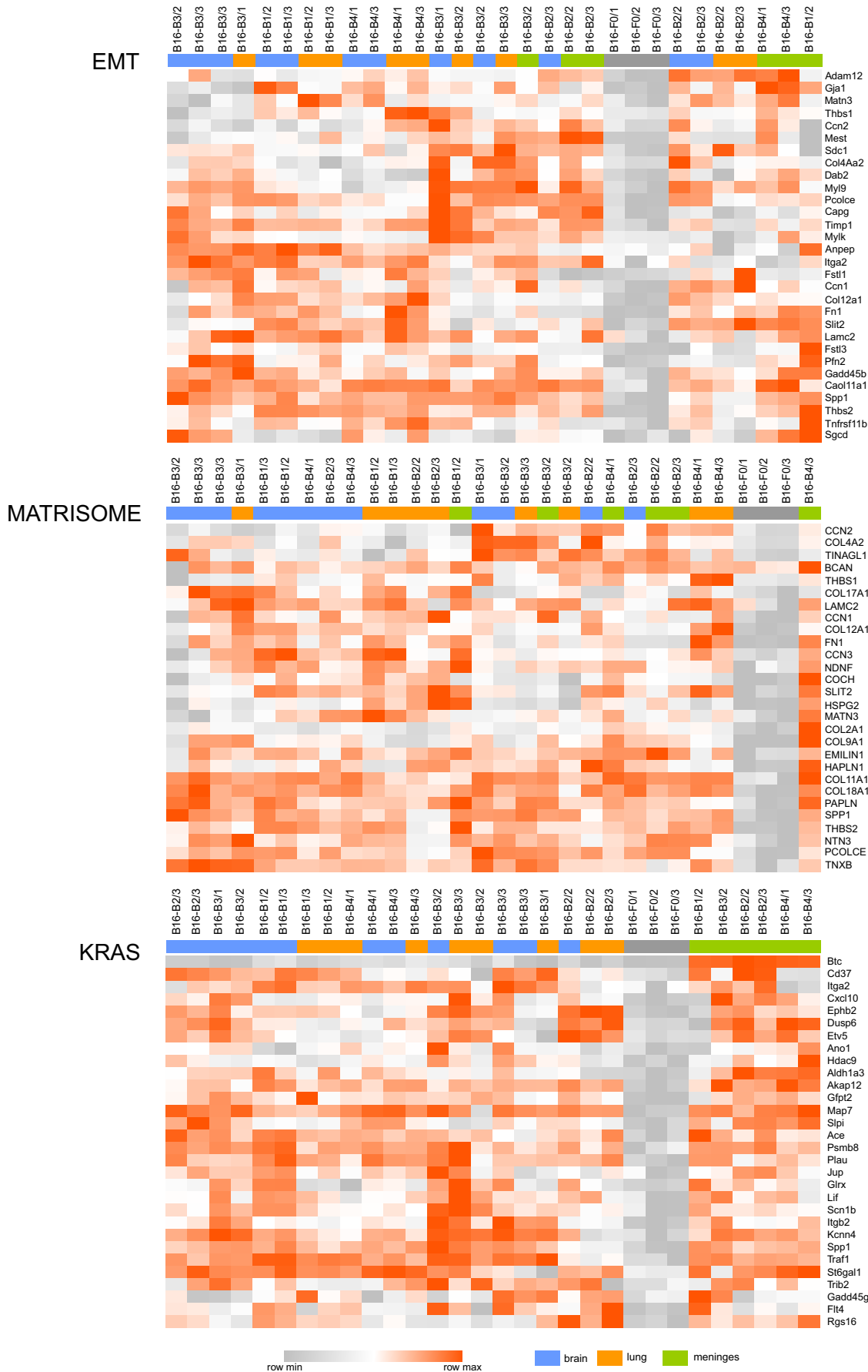# Additional file 3: Heat maps

Heat maps of log2 normalized counts of top core genes with a rank metric score >1 from the GSEA.
Hierarchical clustering of columns and rows using a 1- Pearson correlation coefficients as the distance metric
and clustering using the average linkage method.