

Fig. S1

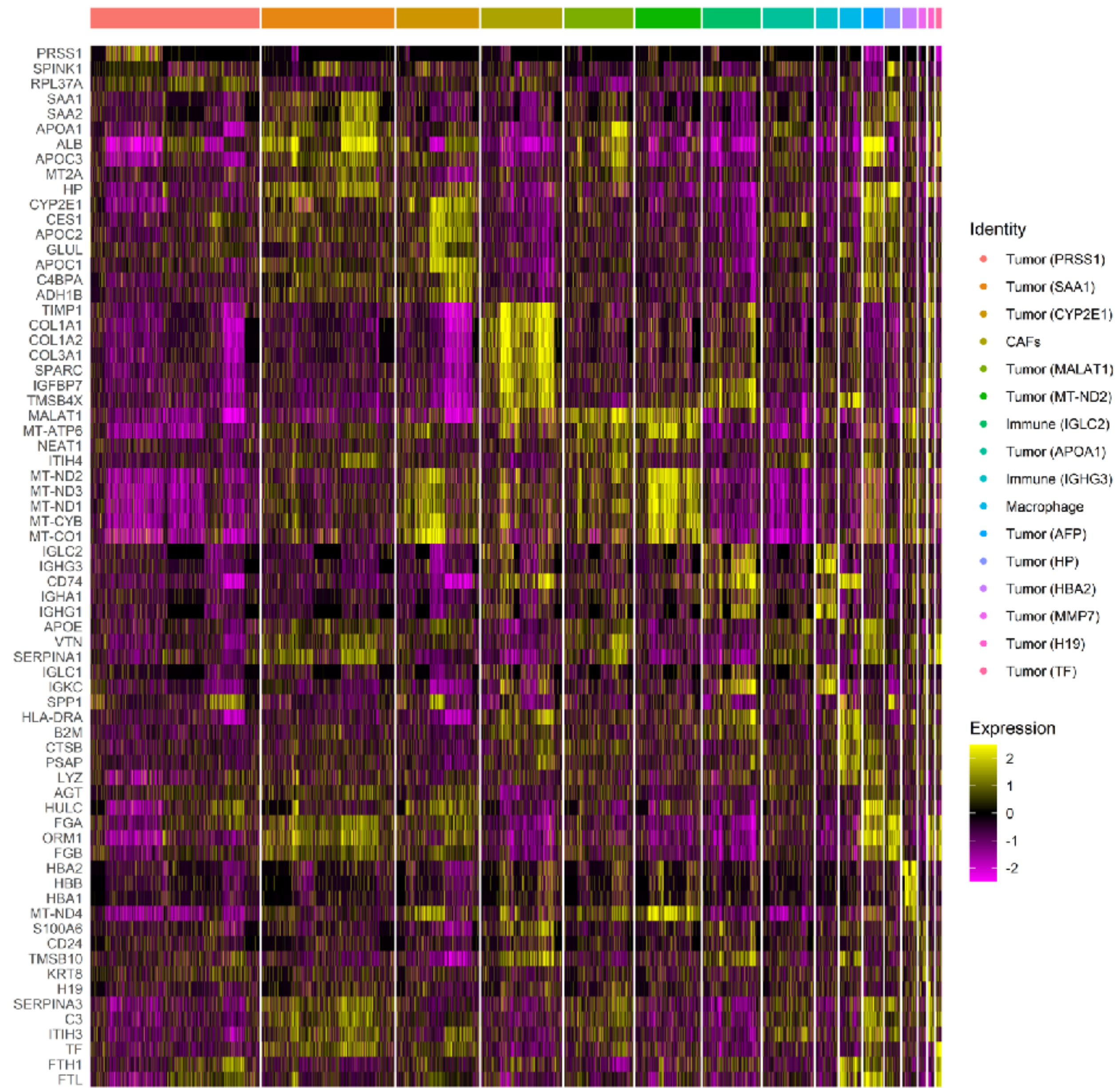


Fig. S1 - Heatmap of cluster markers across samples. Expression of highly expressed genes in each of the clusters identified in the HCC cohort. Clusters that map to the same cell type were identified by the cell type (tumor, immune or CAF) and by the highest expressed gene (in parenthesis).