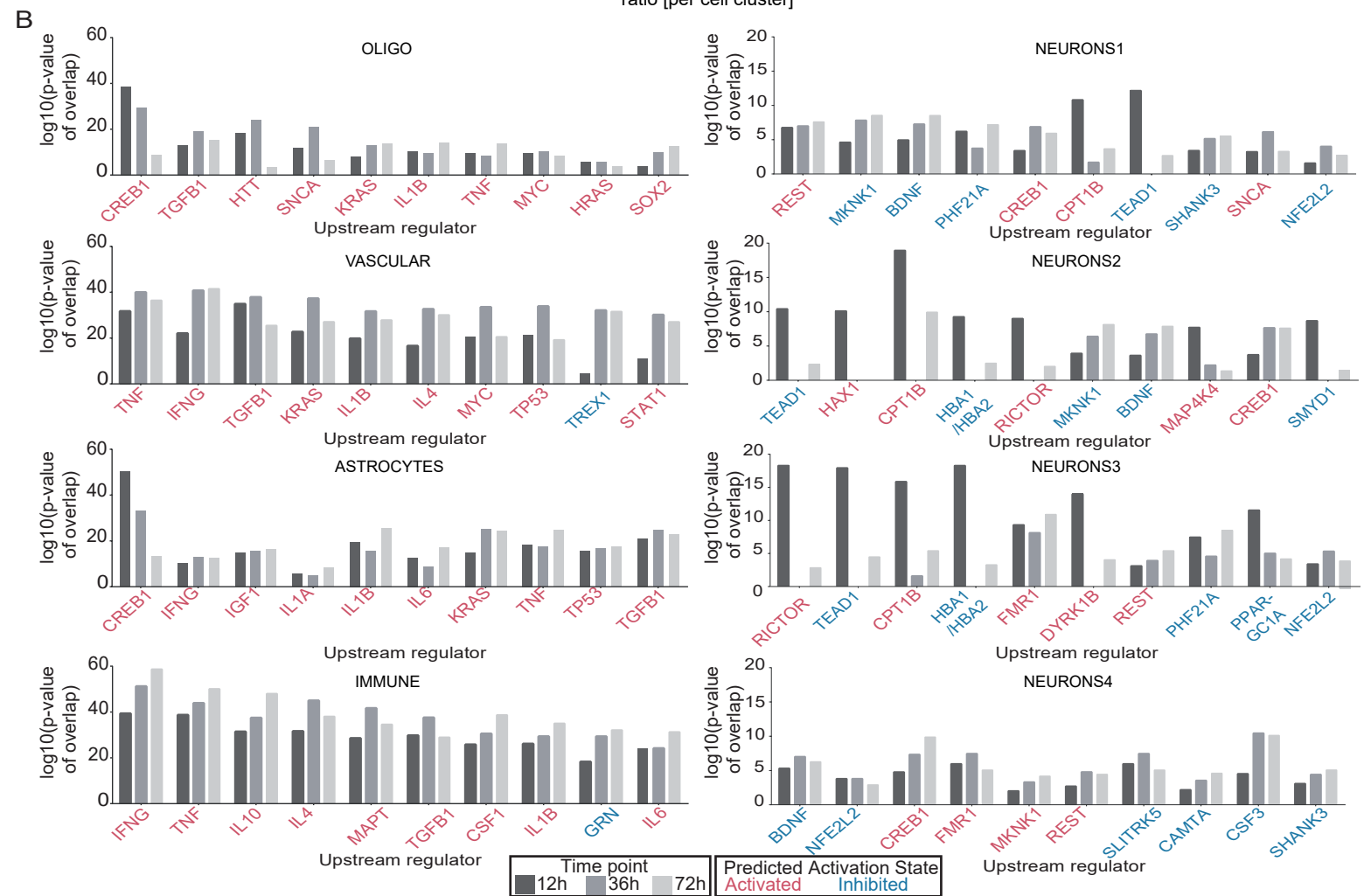
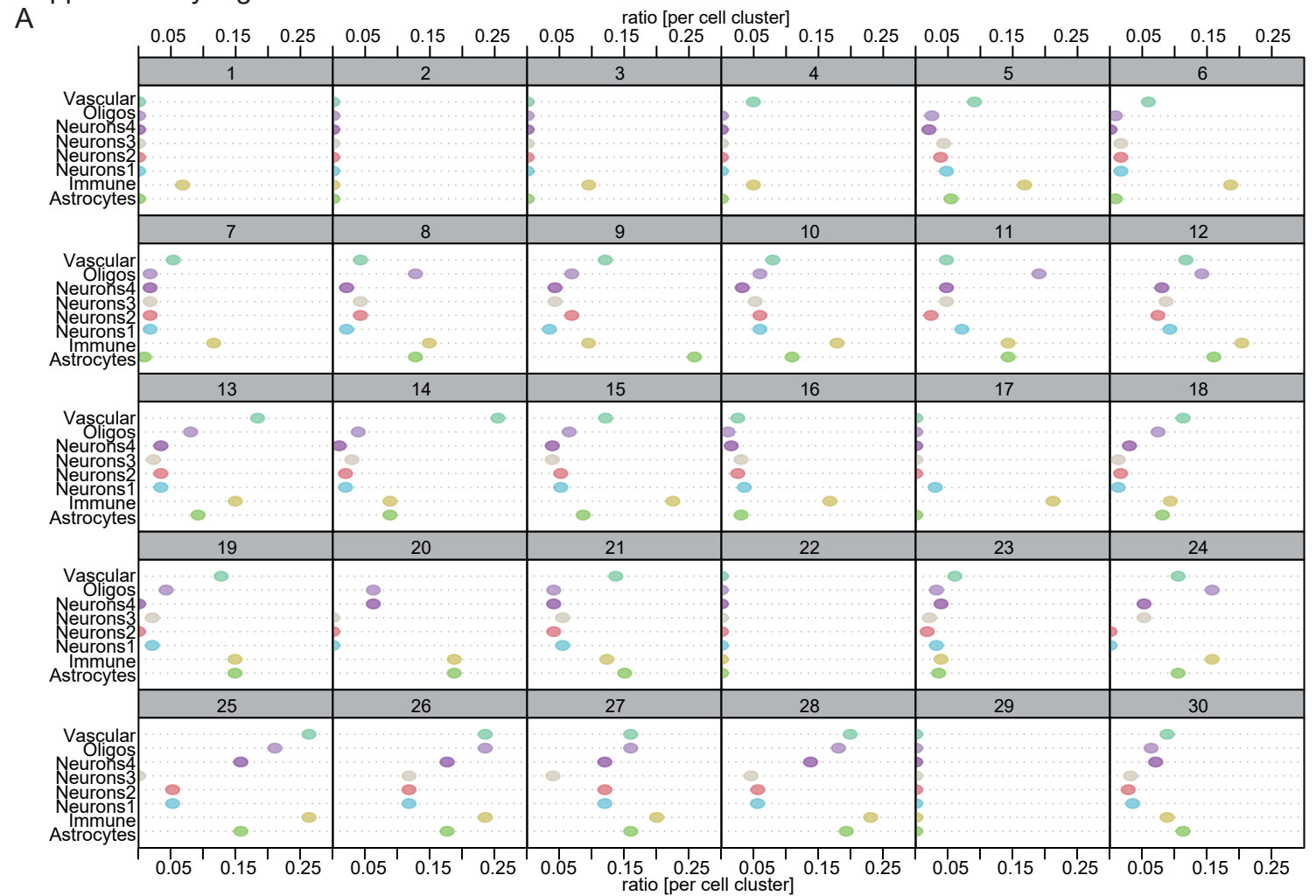


Supplementary Figure 4



Supplement Fig. 4. Clustering analysis and cell-type associated clusters reveal a predominant immune cell type population associated to specific upstream regulators (A) BHC clustered genes were associated with specific CA1 cell types. The ratio represents the proportion of related cell type genes that are significant markers for each cluster ($p < 0.05$). (B) Comparison of the most relevant upstream regulators for each cell populations at different time points. (p -value of overlap < 0.05)