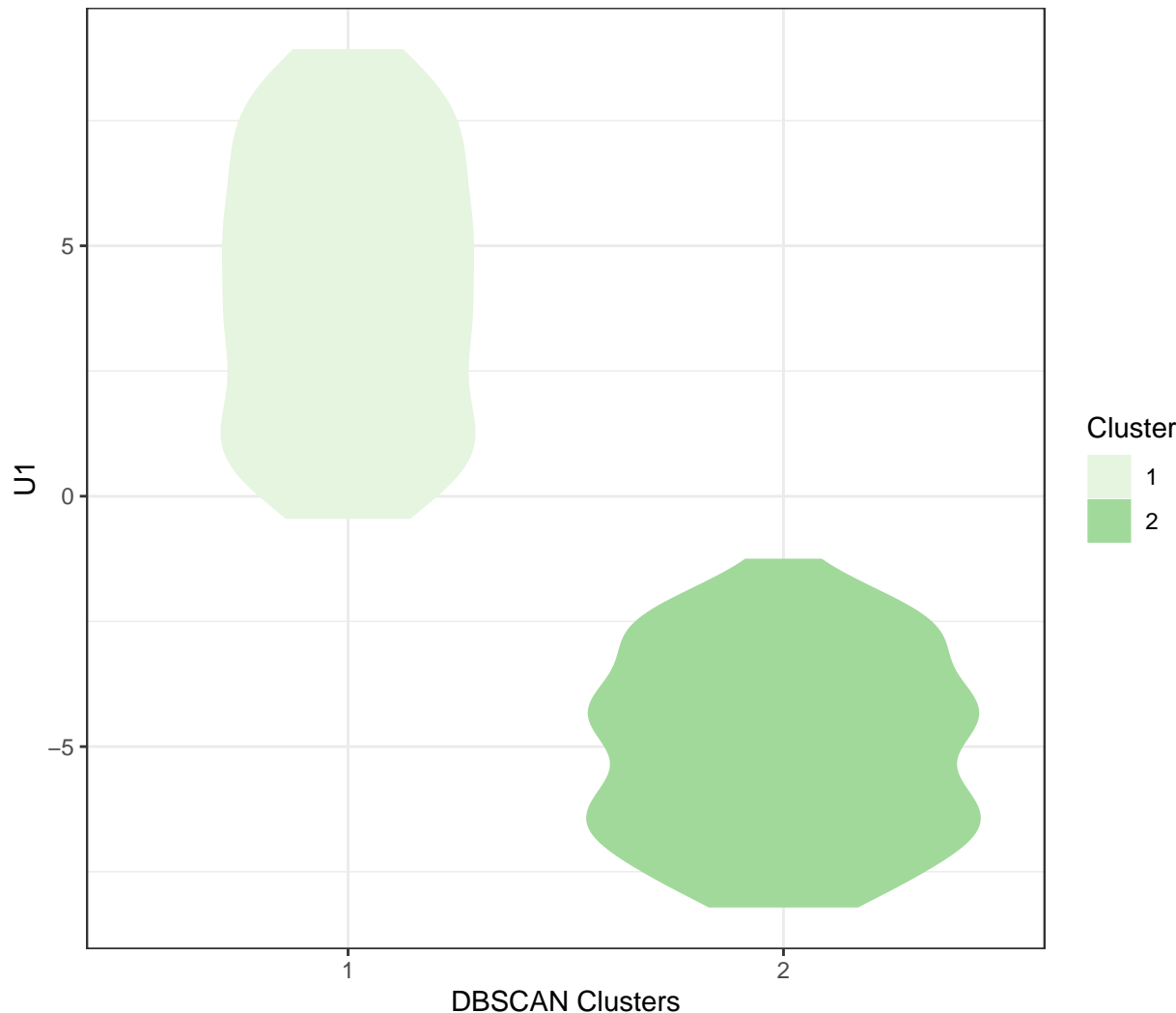
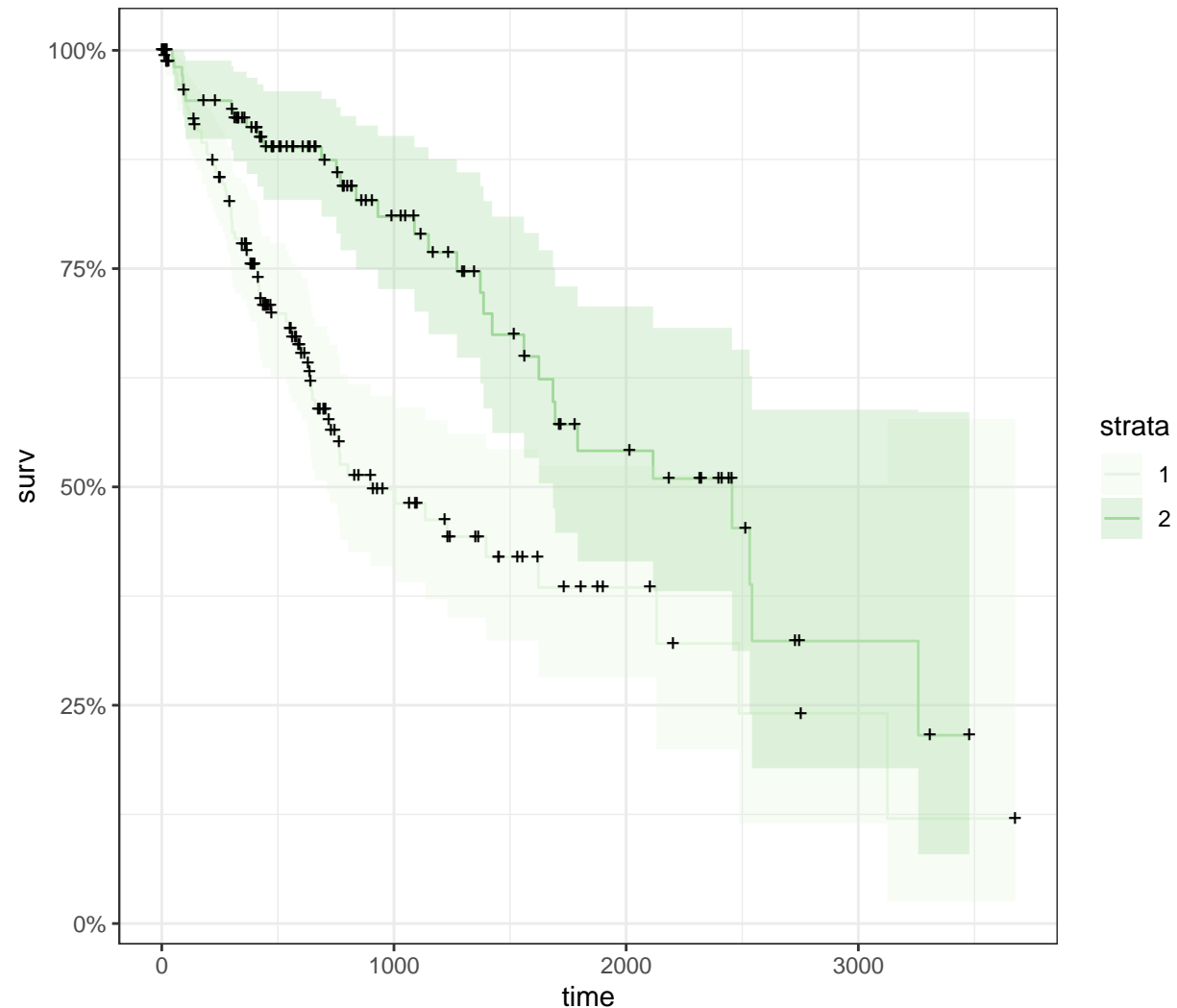


TCGA-LIHC Clustering and Survival, UMAP Embedding of:
GO_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION
GO_MACROPHAGE_INFLAMMATORY_PROTEIN_1_ALPHA_PRODUCTION

A. DBSCAN Clustering



B. Survival Curves



A. DBSCAN clustering of LIHC using gene ontology terms indicative of EMT and INFLAMMATION signatures (min group size = 30).

B. Survival plots corresponding to the clustering on EMT and INFLAMMATION ($p = 8.555e-05$).