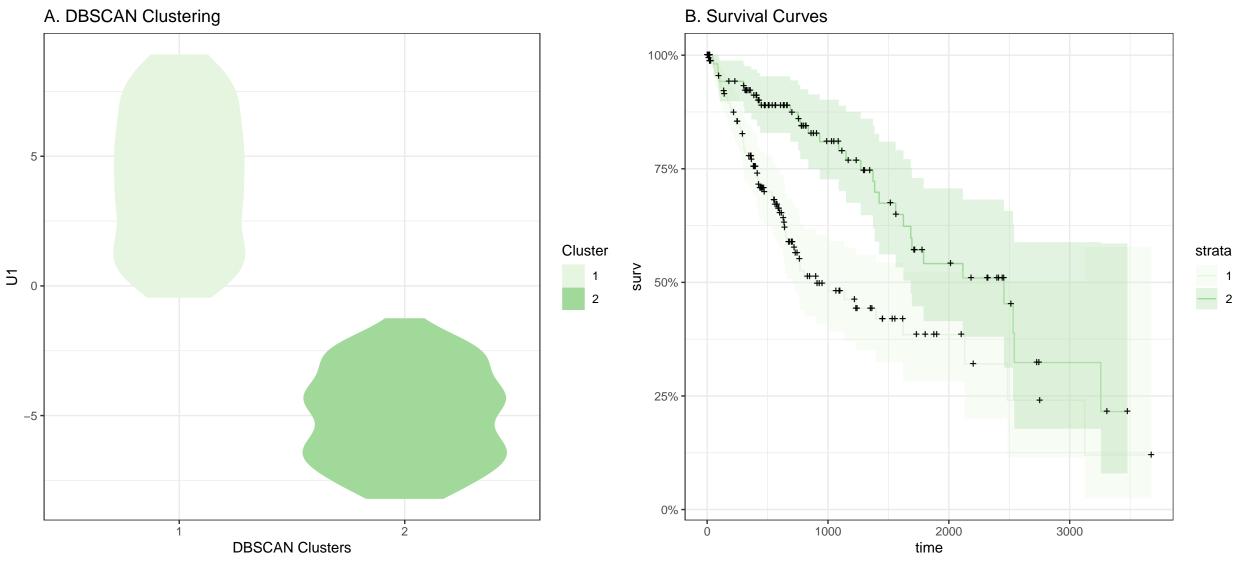
TCGA-LIHC Clustering and Survival, UMAP Embedding of:

GO_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION GO_MACROPHAGE_INFLAMMATORY_PROTEIN_1_ALPHA_PRODUCTION



A. DBSCAN clustering 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).