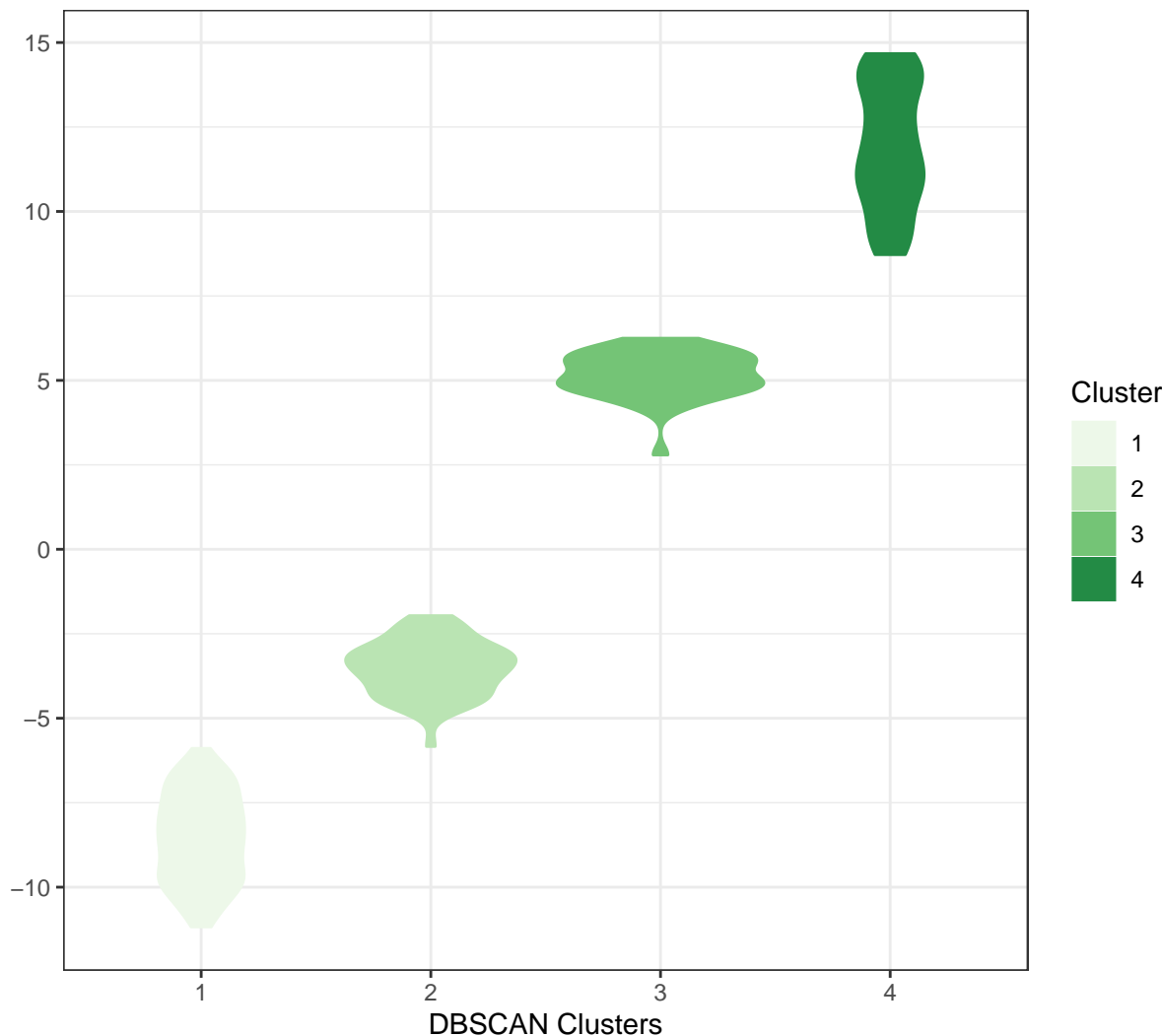
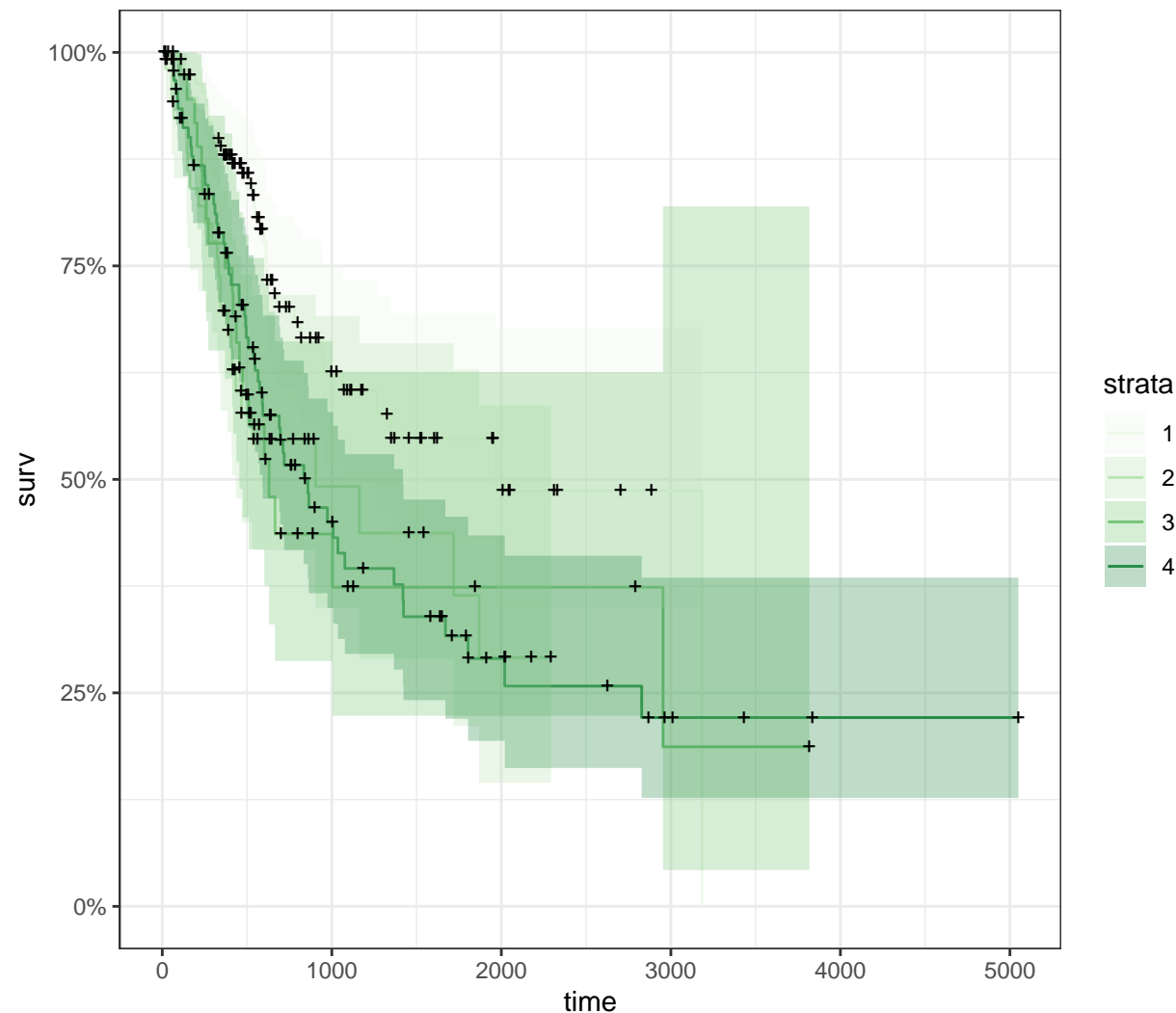


TCGA-BLCA Clustering and Survival, UMAP Embedding of:  
GOTZMANN\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION\_UP  
GO\_POSITIVE\_REGULATION\_OF\_ACUTE\_INFLAMMATORY\_RESPONSE\_TO\_ANTIGENIC\_STIMULUS

A. DBSCAN Clustering



B. Survival Curves



A. DBSCAN clustering of BLCA 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 ( $p = 0.00823$ ).