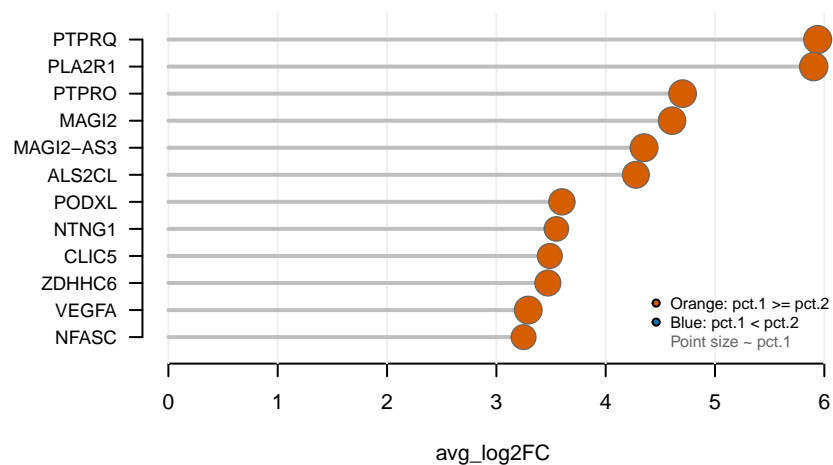
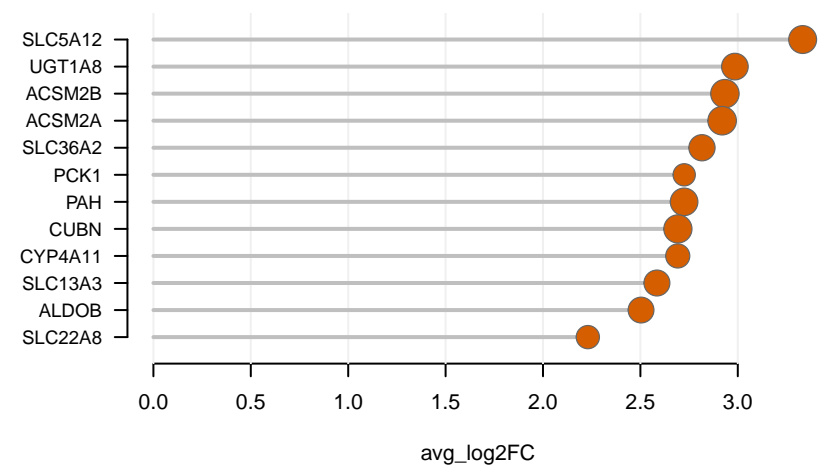


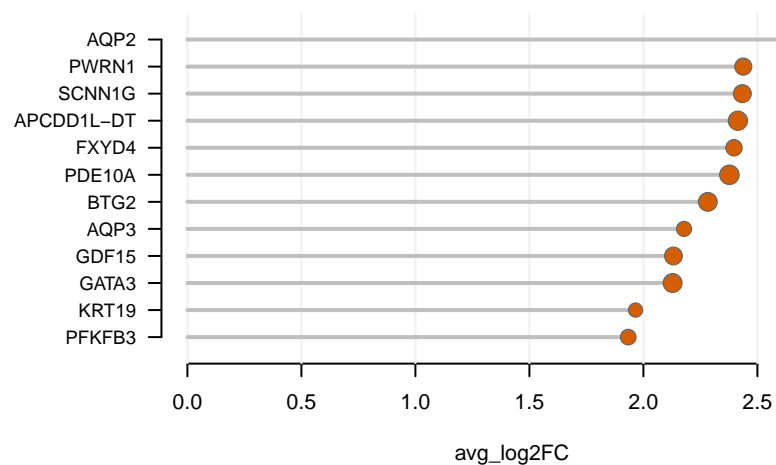
A Top markers (Cluster 12 (Podocyte (glomerular) / Podocyte program))



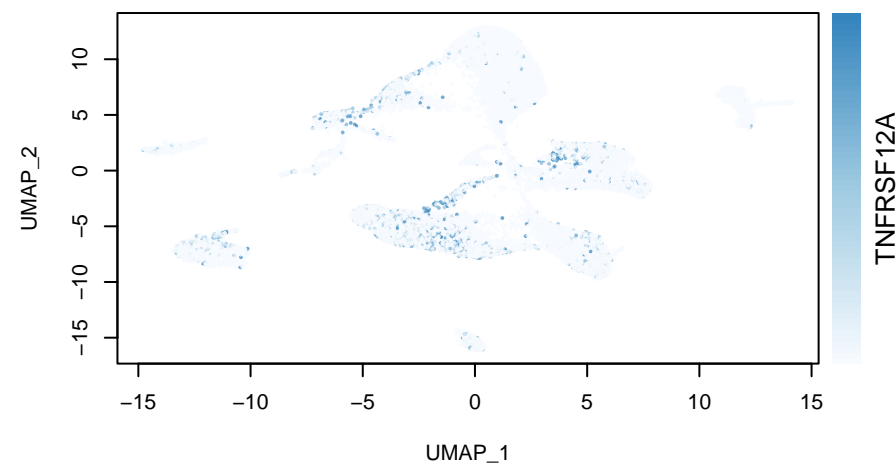
B Top markers (Cluster 3 (Proximal tubule (S3-like) / Metabolic/detox program))



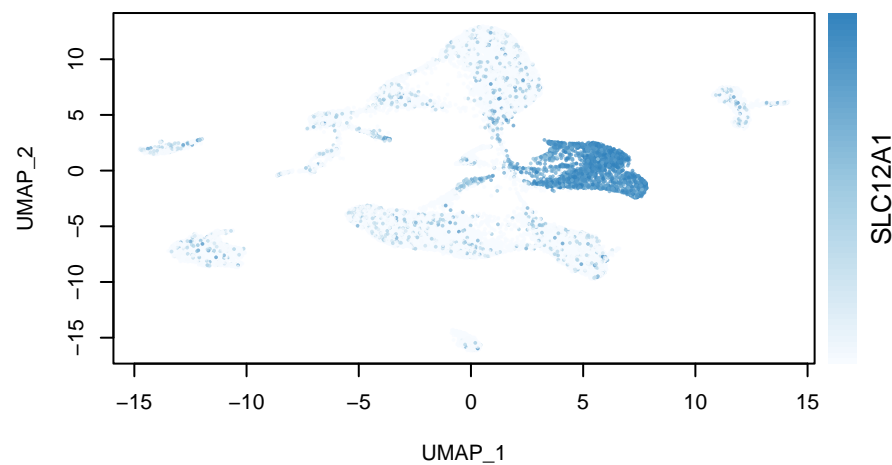
C Top markers (Cluster 0 (Collecting duct principal cell / Principal/CD program))



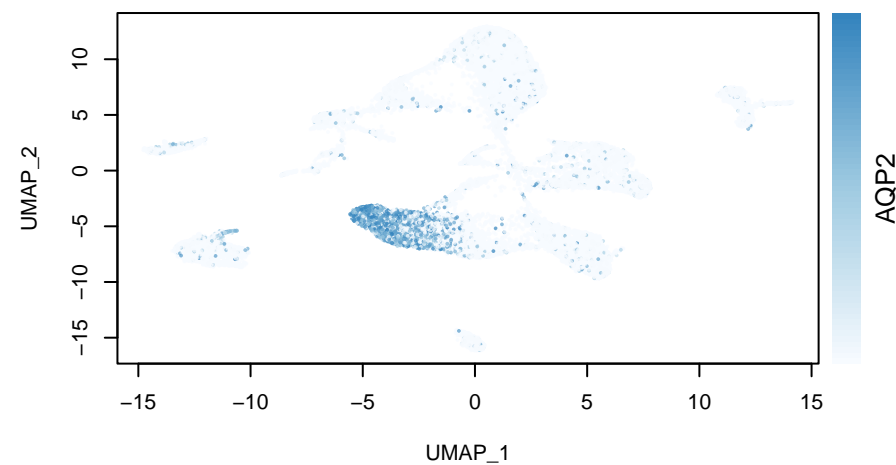
D UMAP feature: TNFRSF12A



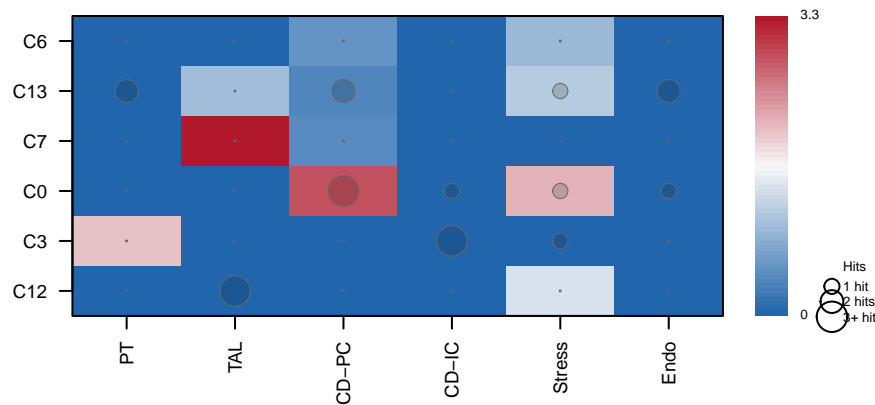
E UMAP feature: SLC12A1



F UMAP feature: AQP2



G Canonical marker support (max avg_log2FC)



H Top annotated contexts

