A picture containing many, person, bunch, person

Description automatically generated

**Supplemental Figure 1** Workflow for SCRS of isolated lymphoid and myeloid immune cell populations from ccRCC subjects and computational analysis. SCRS data across ccRCC and data derived from 10x Genomics and EGAS00001002325 were integrated using SCTtransform approach to form a UMAP of 37,055, acting as a basis for subclustering and further analysis by cell type.