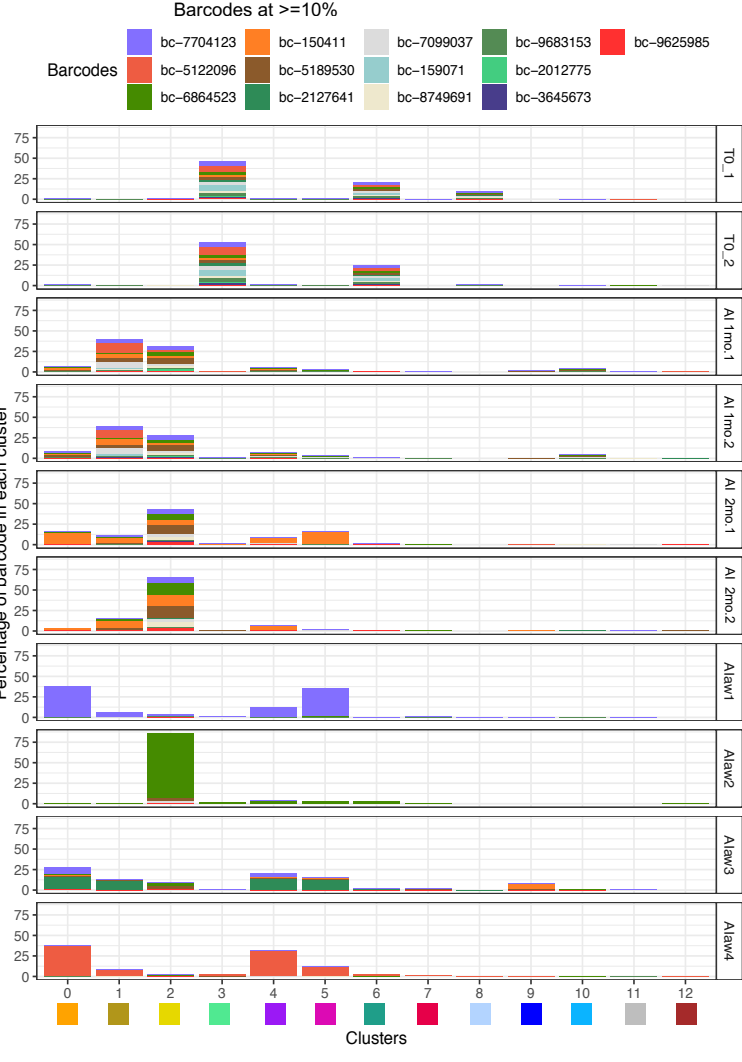
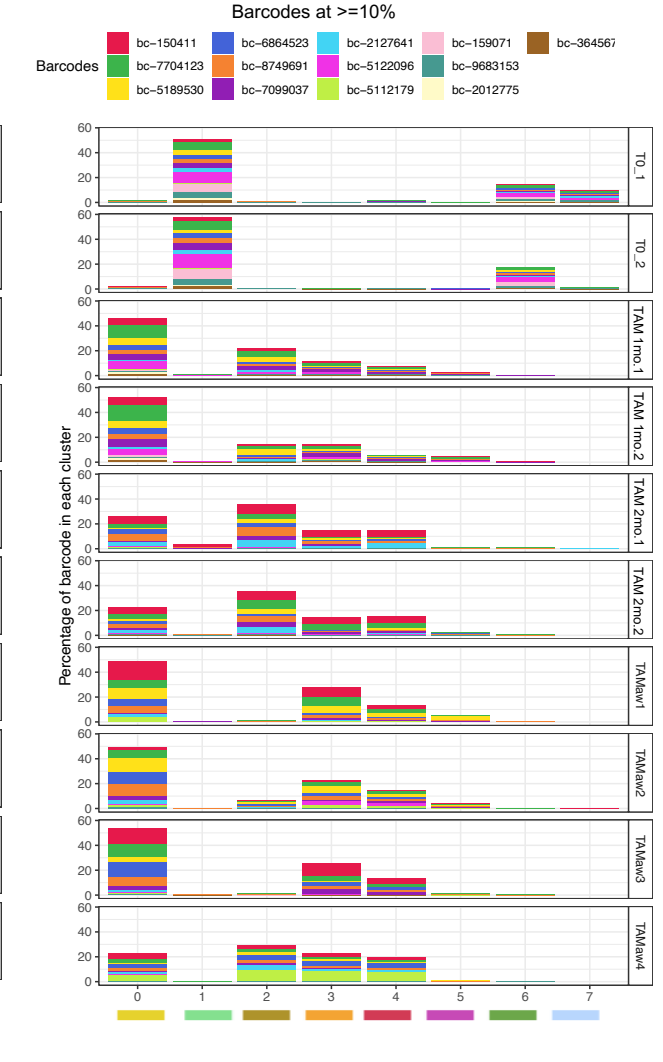


Supplementary Figure S23. High frequency barcode occupancy of scRNA-seq clusters

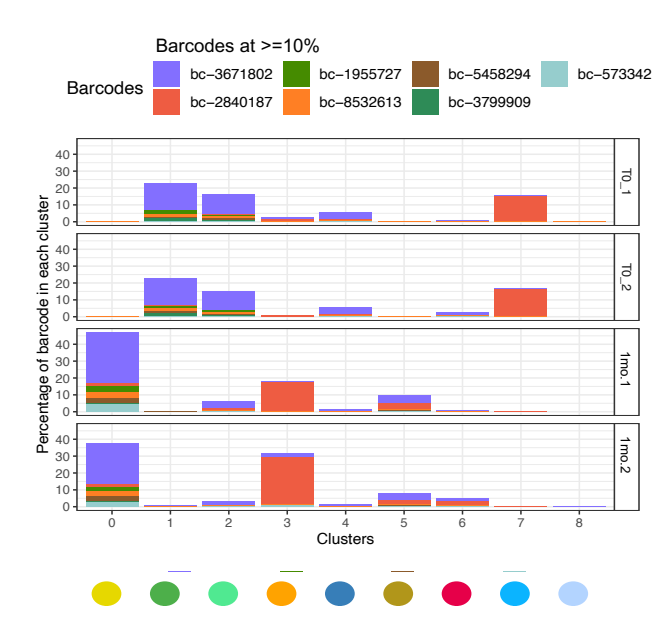
a MCF7 -E2 (AI)



b MCF7 TAM



c T47D -E2 (AI)



Supplementary Figure S23. High frequency barcode occupancy of scRNA-seq clusters. Bar plots depict the percentage of each lineage (barcode frequency >10%) in distinct UMAP clusters for TRADITIOM LSC (live single cell) MCF7 -E2 (AI) (a), MCF7 TAM (b) from T0, early dormancy (1month, 1mo), late dormancy (2month, 2mo) to awakening and T47D -E2 (AI) (c) from T0 to dormancy (1month, 1mo).