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 TRADITION 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).