



**Supplemental Figure 13. Expression of LTB, BLK, CXCL12, CCL4, XCL1 and LMNA within T cells and NK cells.**

Two-dimensional UMAP plots displaying single cell transcriptomics of 14,242 T cells and 2,820 NK cells from 13 OPSCC patients. Each dot represents a single cell. Expression levels of LTB, BLK, CXCL12 and CCL4 are depicted in color code. Dotted borders denote where CD4, CD8 and NK cell clusters are located.