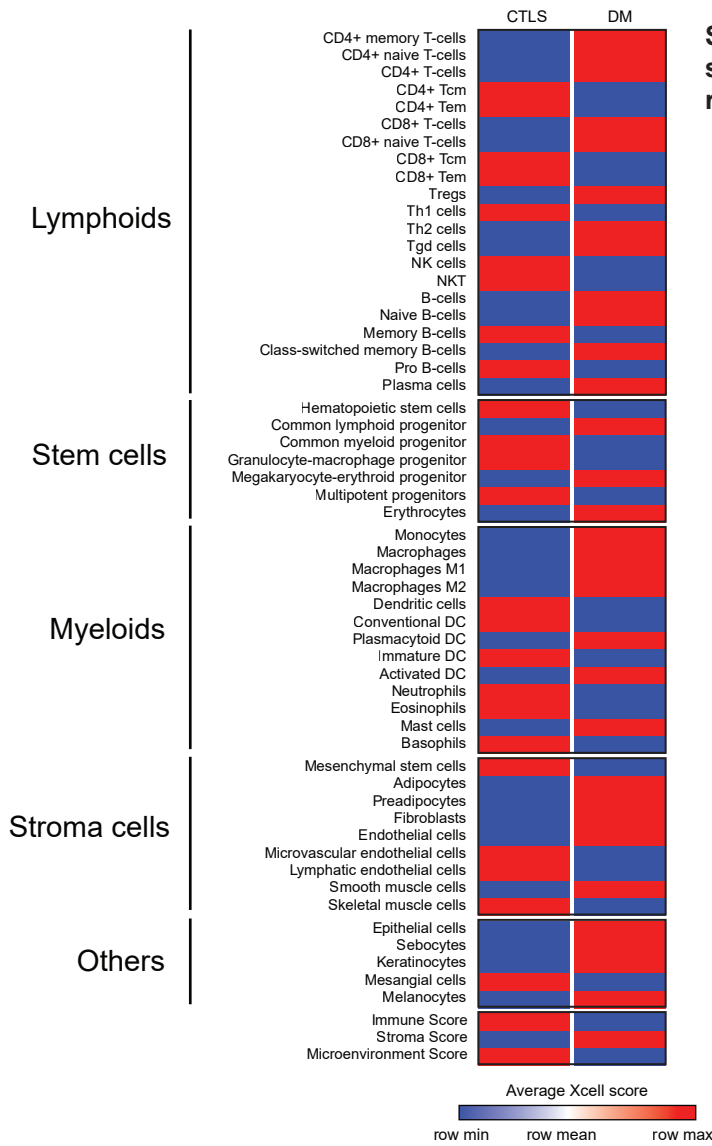
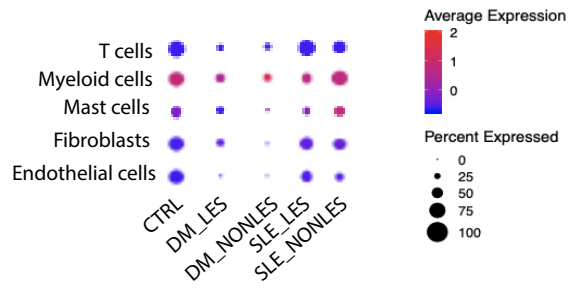


Supplemental Figure 1: xCell analysis of HC vs. DM showing stromal and inflammatory cell populations responsible for gene expression.





Supplemental Figure 2. Single cell RNA-sequencing analysis of IL-18 expression in inflammatory cell populations in control (CTRL), DM (lesional and non-lesional) and SCLE (lesional and non-lesional) skin. Graph represents gene expression in indicated cells for % of cells expressing IL-18 (by size of circle) and degree of expression (by color).