



Extended data Figure 6. Individual TRAV and TRBV scTCR-seq counts on unexpanded cells of CD8+ Tconv cluster 3 across samples from a single SJS/TEN patient. Individual counts for all identified TRAV and TRBV genes across samples from unaffected skin, affected skin, and blister from three anatomical sites. The dominant TRAV and TRBV of the dominantly-expanded cells in CD8+ Tconv cluster 3 in affected skin and blister fluid are highlighted in red. Figure created using VGAS. TCR, T-cell receptor; TRAV, TCR alpha variable; TRBV, TCR beta variable; Tconv, T conventional cell.