Supplementary data for “Fast multiclonal clusterization of V(D)J recombinations from high-throughput sequencing”

October 9, 2013

Table 1: Statistics on the eight samples sequenced for a patient with ALL and on the results of Vidjil. On each dataset, starting from some number of reads, Vidjil detects 40-windows overlapping the CDR3. The count of unique 40-windows is smaller as clones gather several reads. The last columns shows the number of clones with at least 100 reads. All percentages are computed with respect to the number of reads of each dataset.