1. Data is sourced from Figure 2 of the paper. The data is presented as a categorical heatmap of 3 T cell response classes, roughly corresponding to our categories, so we record, with index\_peptide\_activity=1,
   1. White pixels, >50% of index peptide response, as 0.75, to classify them as SB
   2. Yellow pixels, 5-50% of index peptide response, as 0.25, to classify them as WB
   3. Red pixels, <5% of index peptide response, as 0.02, to classify them as NB
2. TCR sequences are taken from Supplementary Table 1 of “Interface-disrupting amino acids establish specificity between T cell receptors and complexes of major histocompatibility complex and peptide” (PMID: 17041605) and Supplemental Experimental Procedures of the source paper.