



Fig. S2 A simplified representation of the estimated overlap between peptide binding repertoire of HLA molecules. Each circle represents the peptide binding repertoire of a single HLA molecule, i.e. the top 1000 best binding peptides, as shown in the dash-dotted circle. By predicting the peptide binding repertoire of all HLA molecules, we can subsequently investigate the overlap between each pair of HLA molecules by looking at their intersection.