

Figure S1. Peptide $2F_o - F_c$ electron density in the native and anchor modified HuD/HLA-A2 structures. Mol1 (chain C) is for the first molecule in each asymmetric unit; Mol2 (chain F) is for the second molecule in each asymmetric unit.

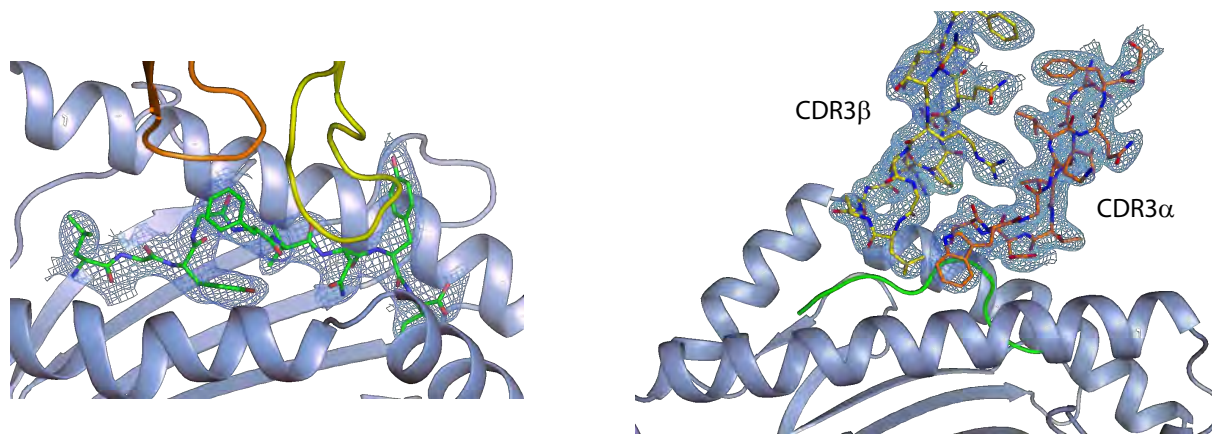


Figure S2. Peptide (left) and CDR3 α/β loop (right) $2F_o - F_c$ electron density in the A6-HuD/HLA-A2 structure.

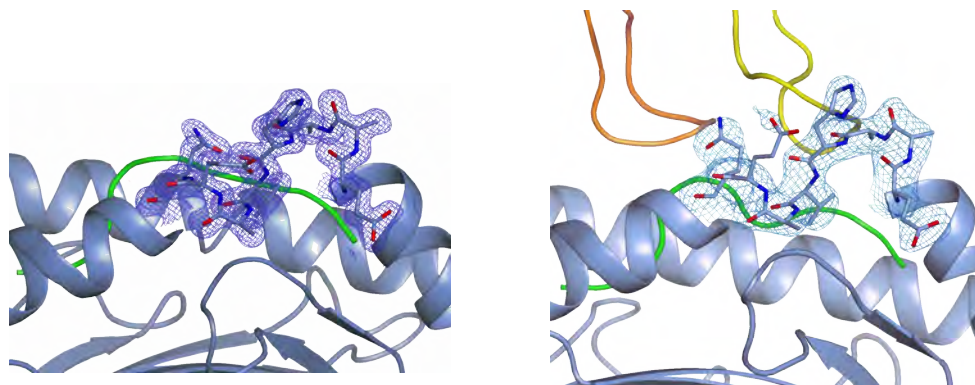


Figure S3. $2F_o - F_c$ electron density for the $\alpha 2$ helix linker region in the free HuD/HLA-A2 complex (left; for Mol 1) and in the A6-HuD/HLA-A2 complex (center).

