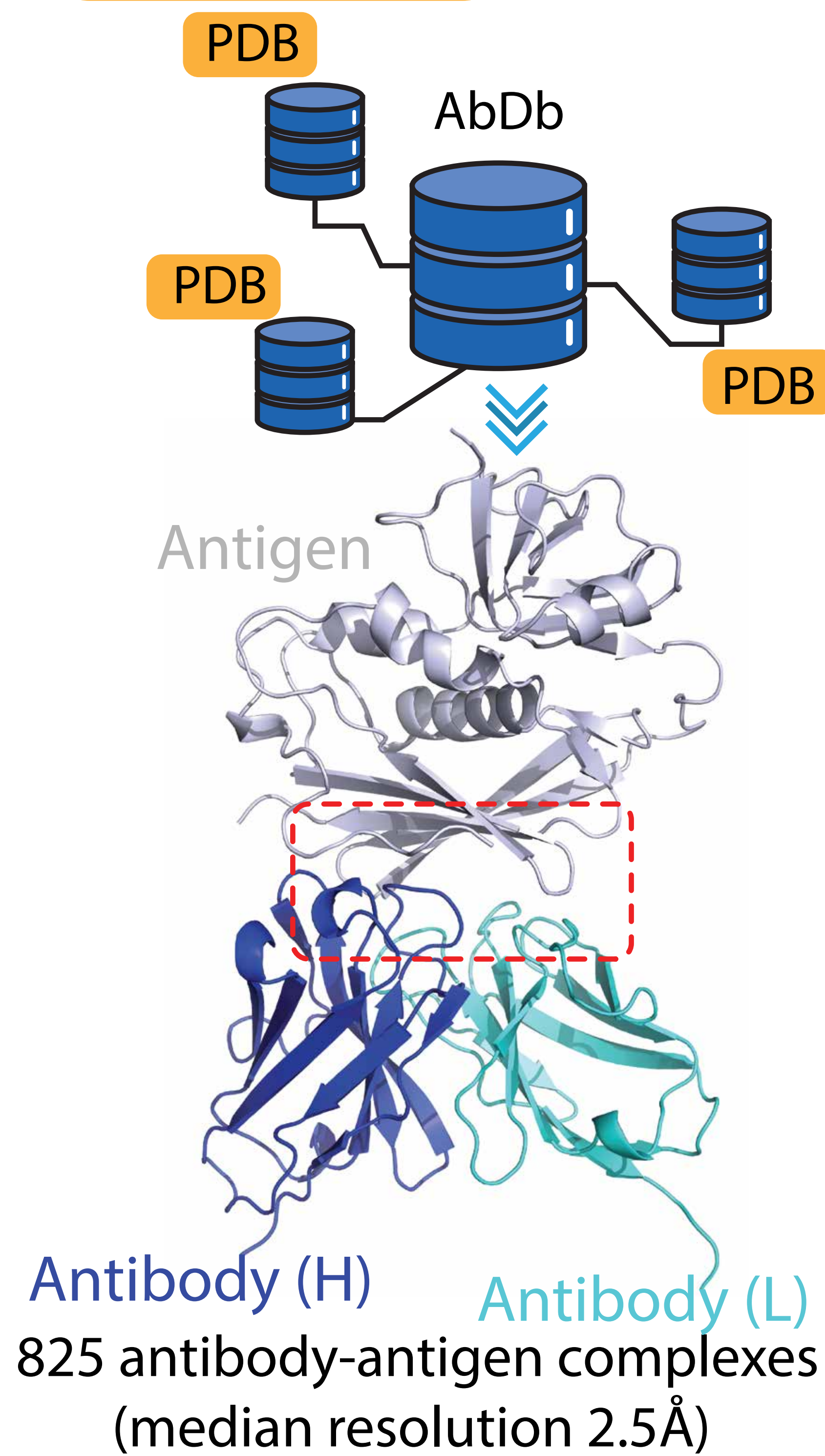
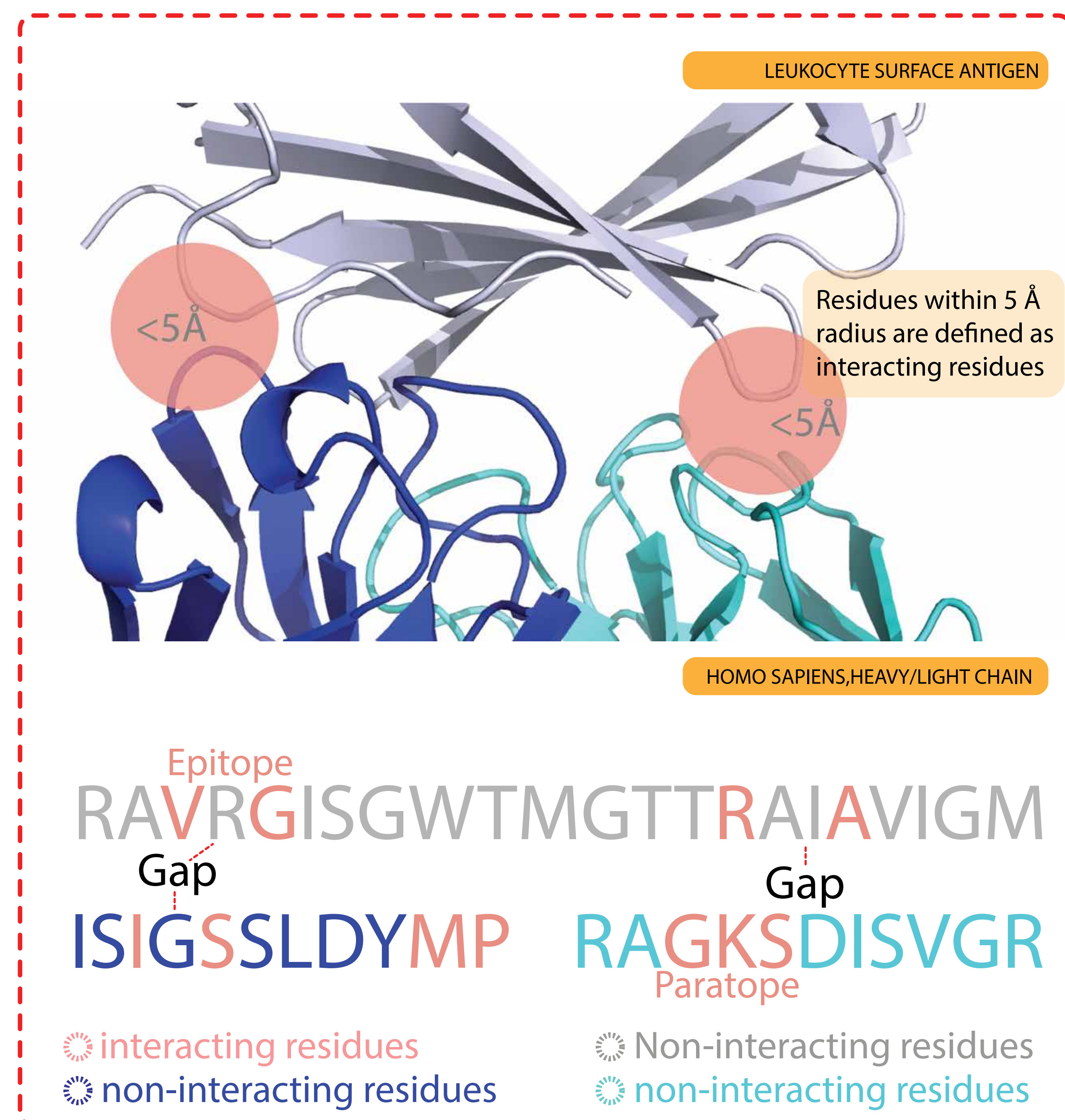


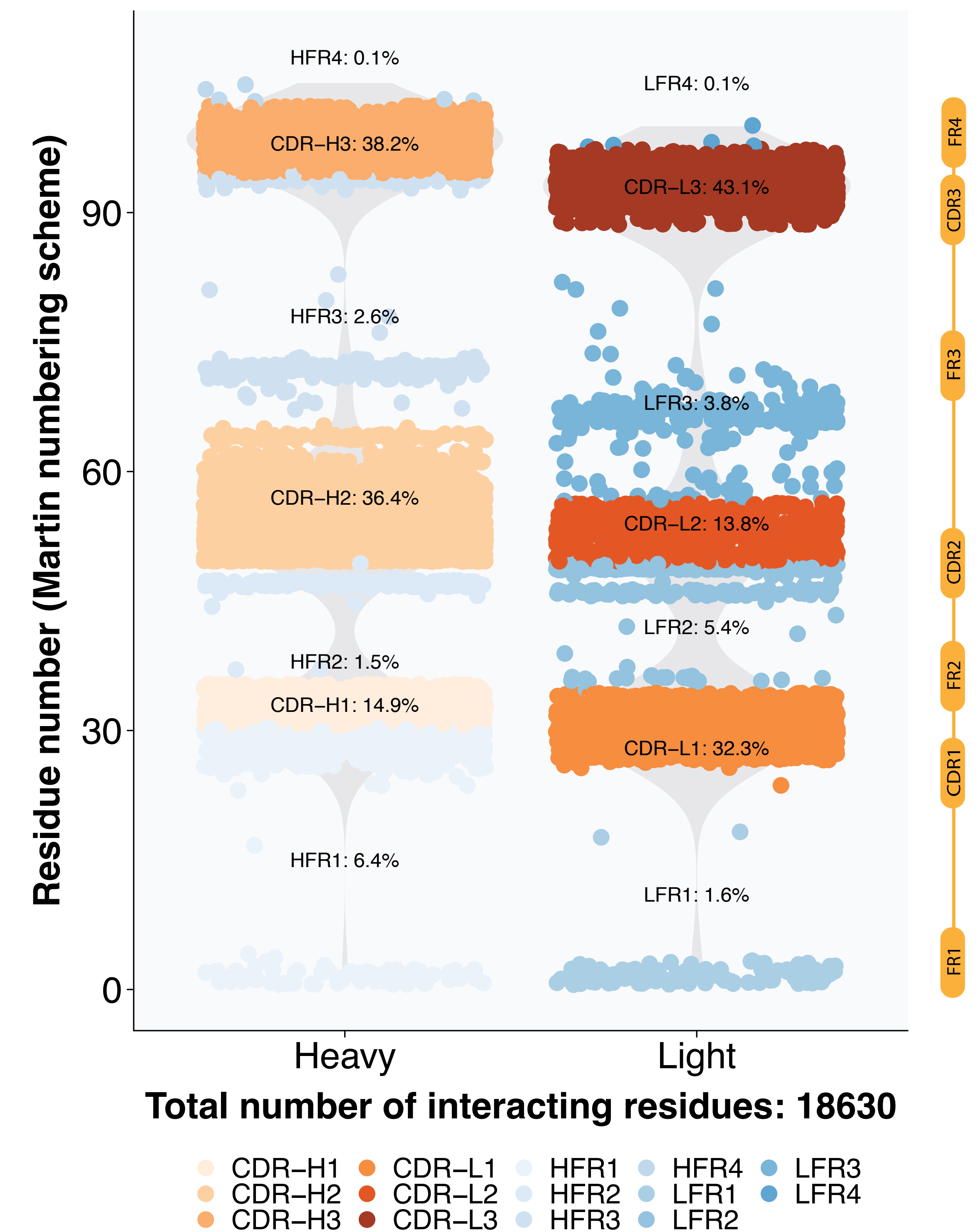
A Data source



B Interacting residue identification



C Interacting residue distribution



D Gap and paratope/epitope length distribution

