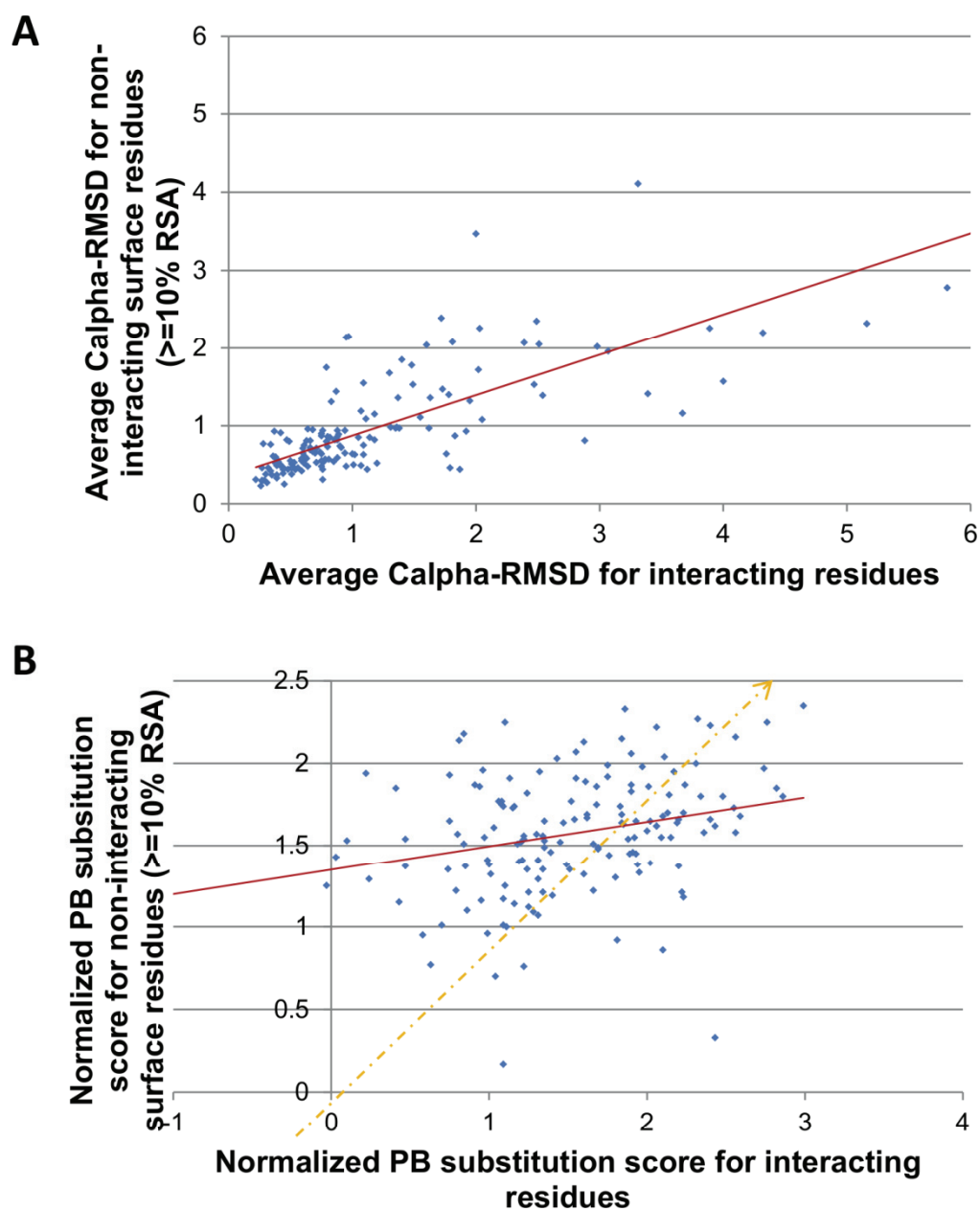


Figure S7: Distribution of parameters for interface vs. non-interacting surface regions per protein.



Scatter plot of a). C α RMSD b). PB substitution score for interacting residues vs. rest of surface residues for each of the 77 protein-protein complex pairs. The line indicating the trend of the scores is shown in brown. The line depicting an exact 1:1 correspondence in scores is shown as an yellow dotted line.