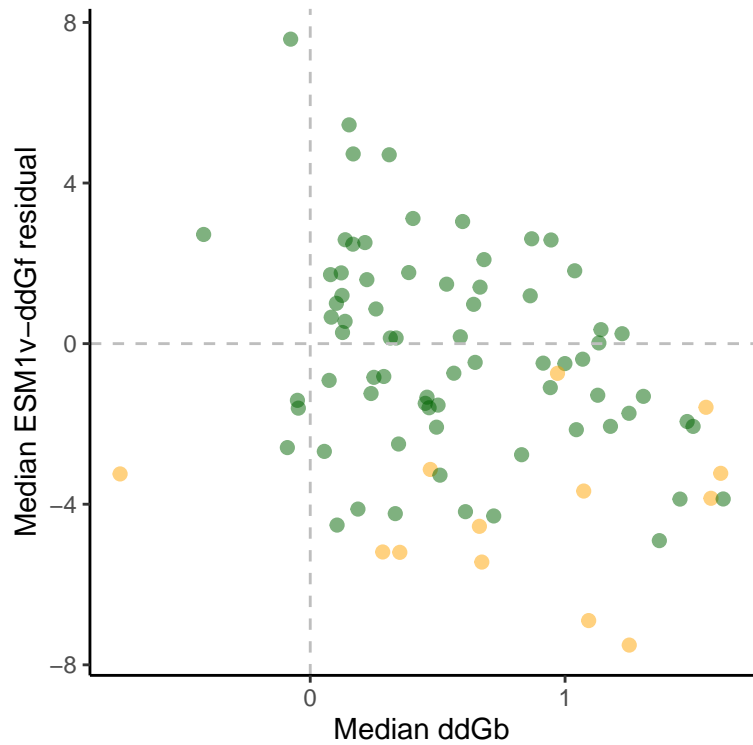


PSD-95-PDZ3: 84 residues (all 1587 mutat

Spearman's rho = -0.31



Binding site Non-orthosteric site