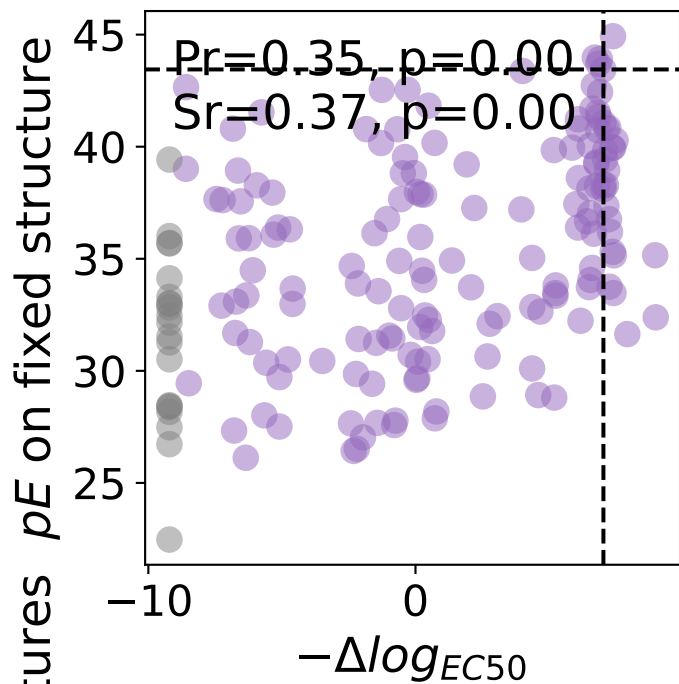
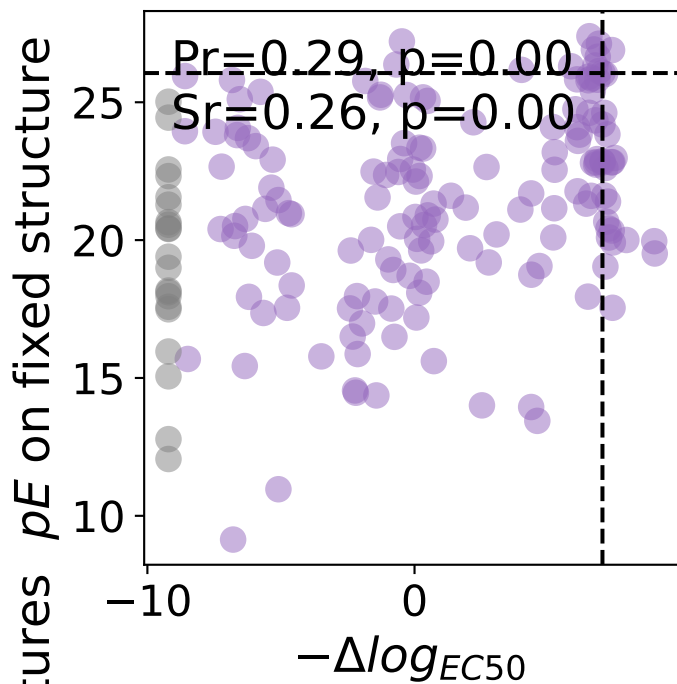


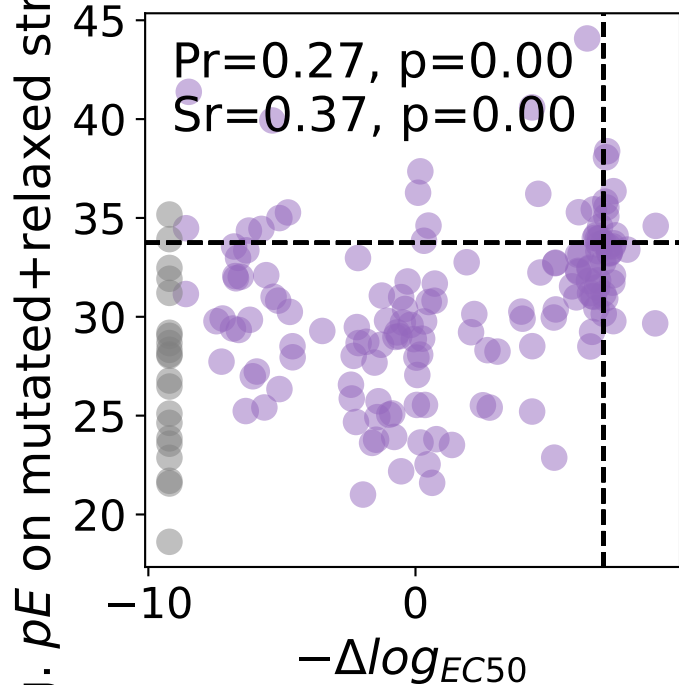
HERMES-fixed 0.00



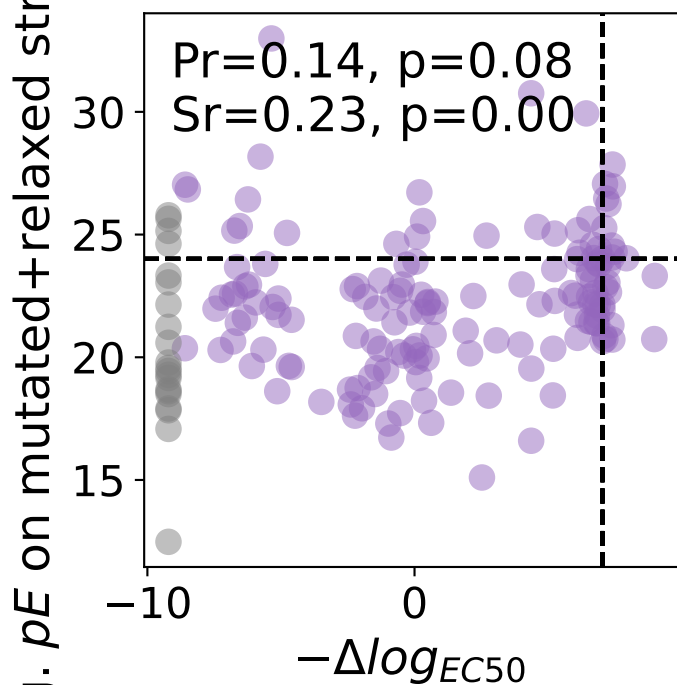
HERMES-fixed 0.50



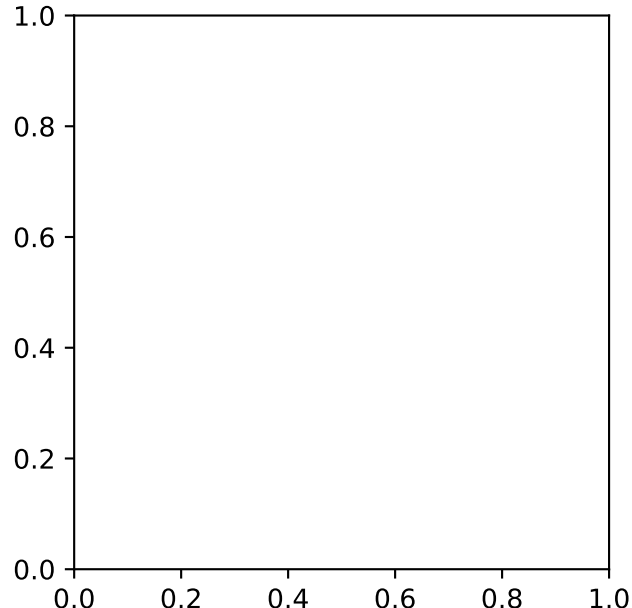
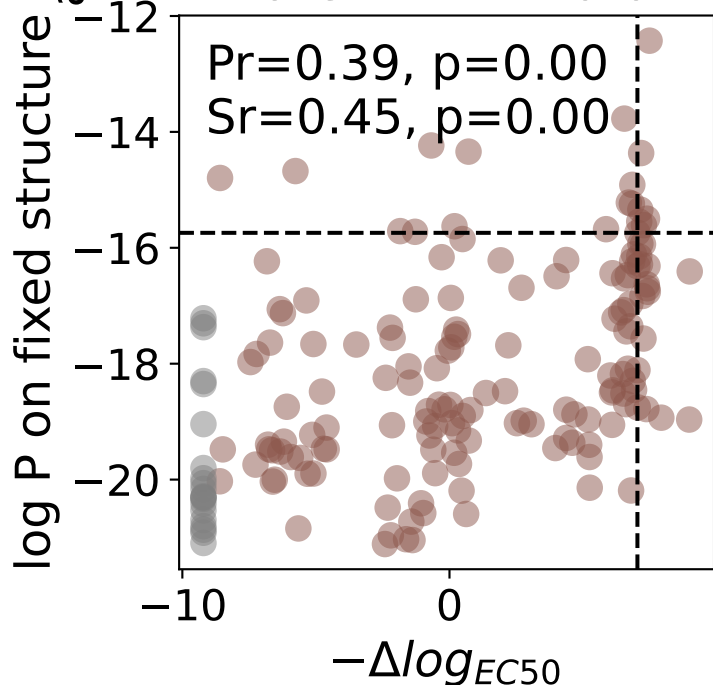
HERMES-relaxed 0.00



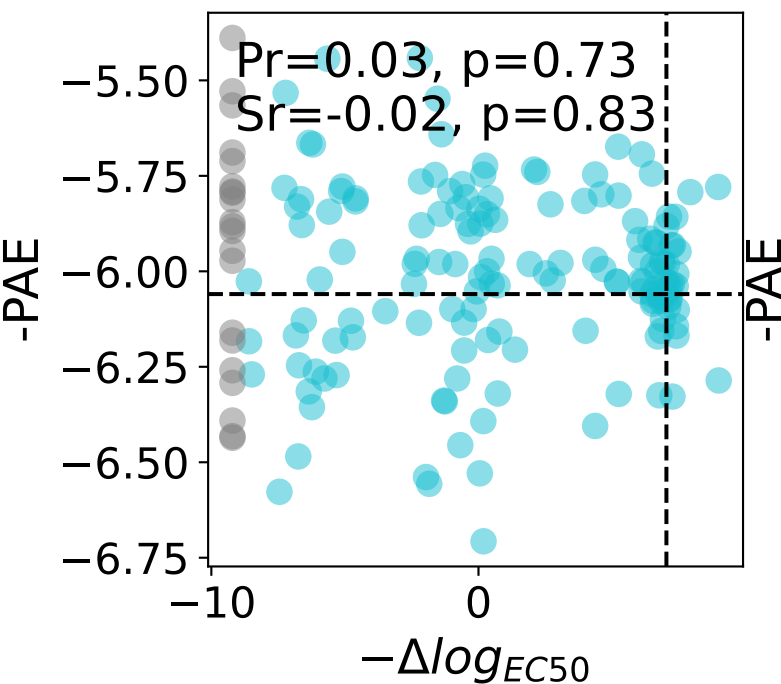
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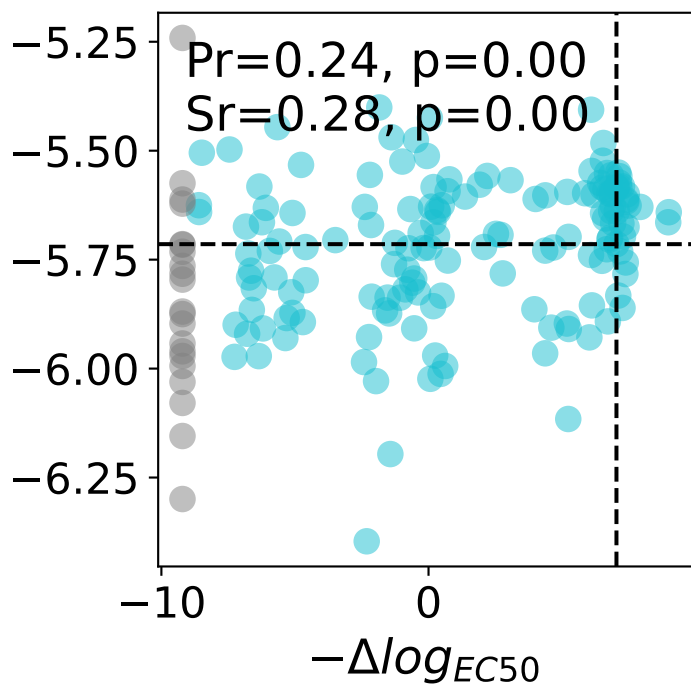
ProteinMPNN 0.02



TCRdock



TCRdock benchmark



BLOSUM62

