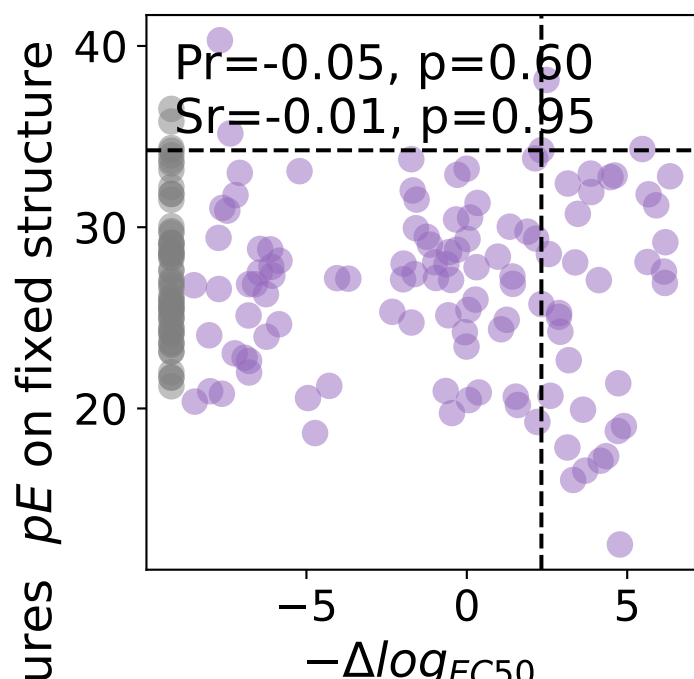
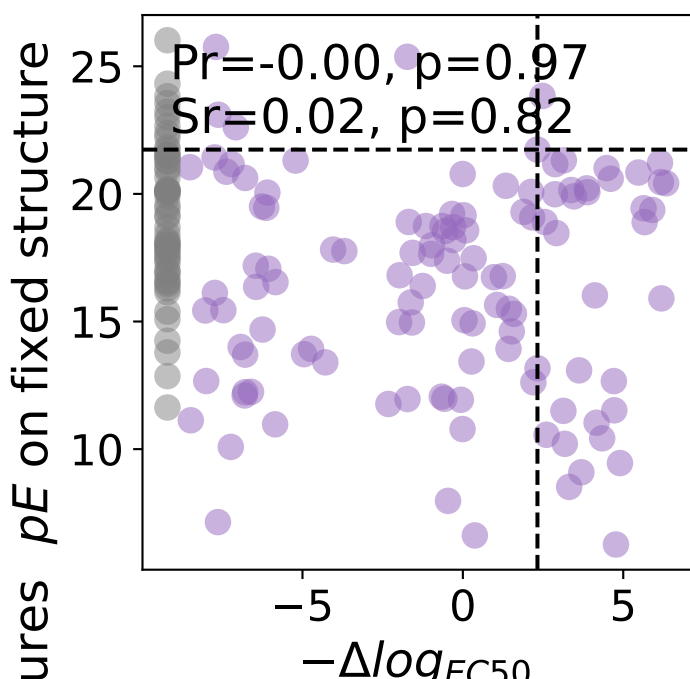


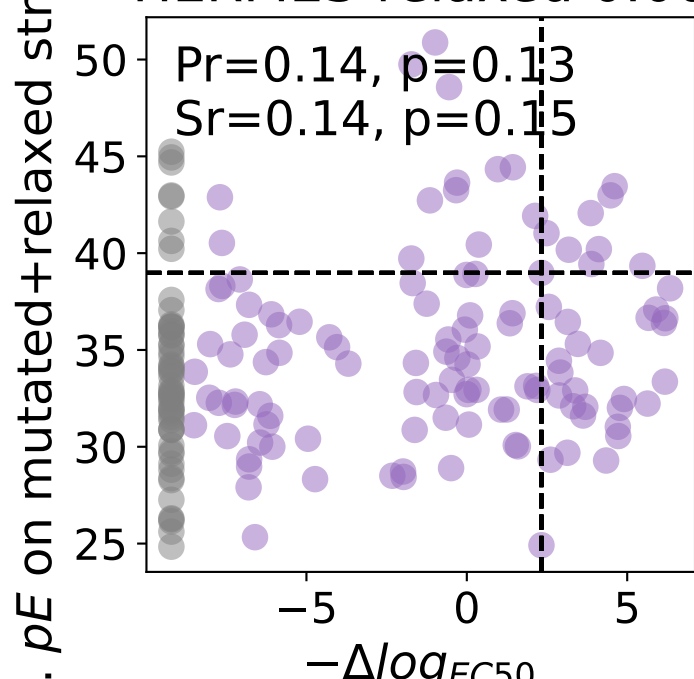
HERMES-fixed 0.00



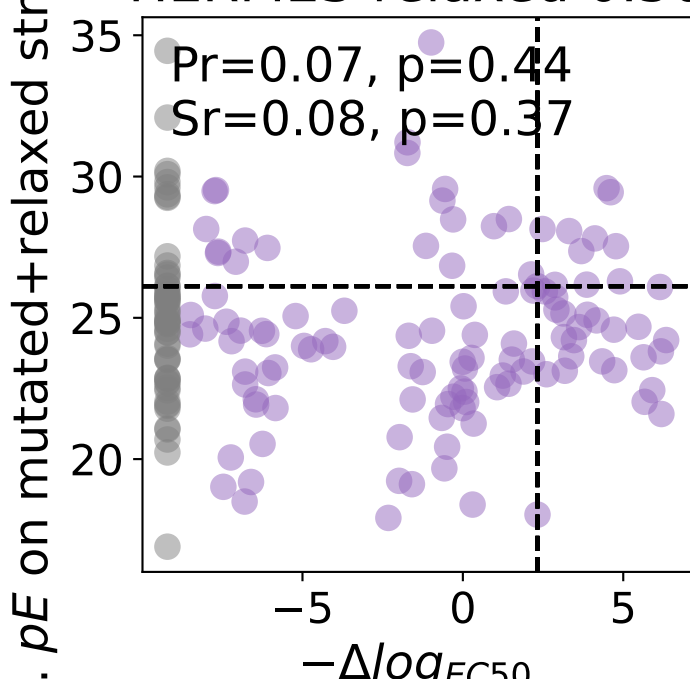
HERMES-fixed 0.50



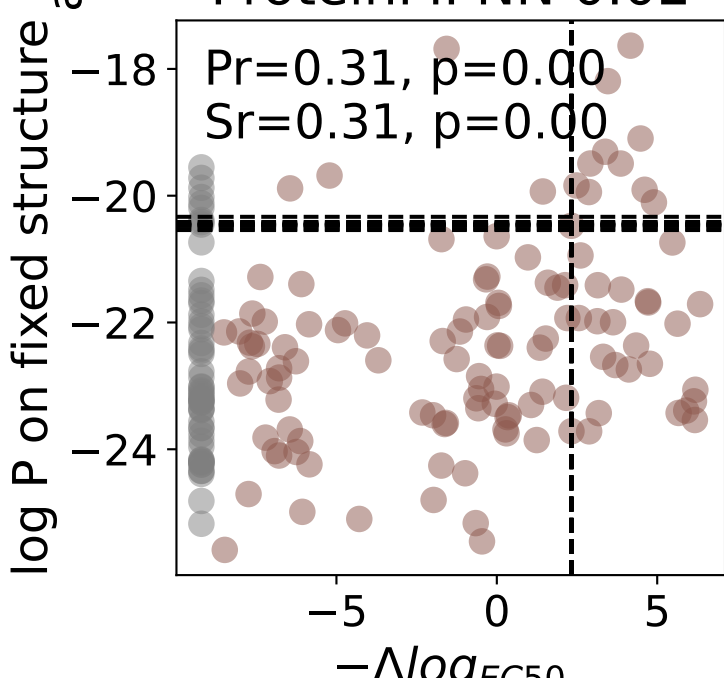
HERMES-relaxed 0.00



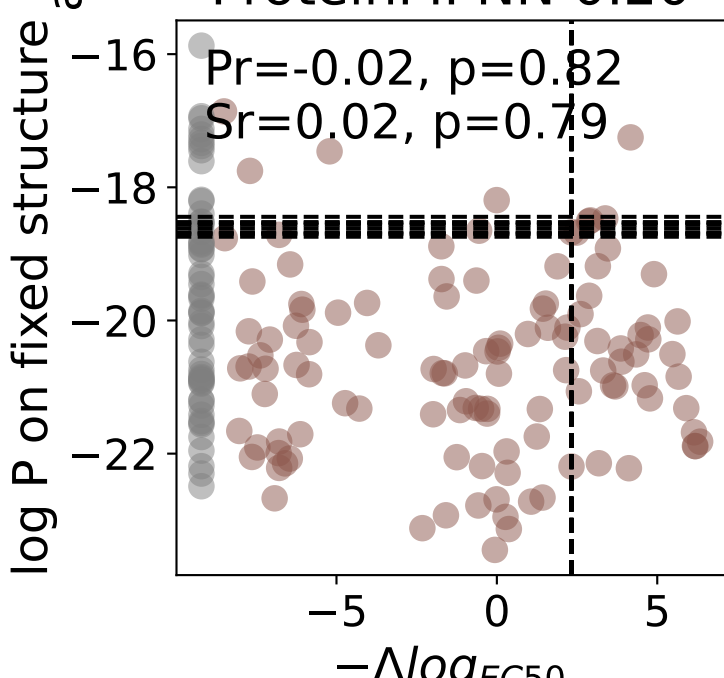
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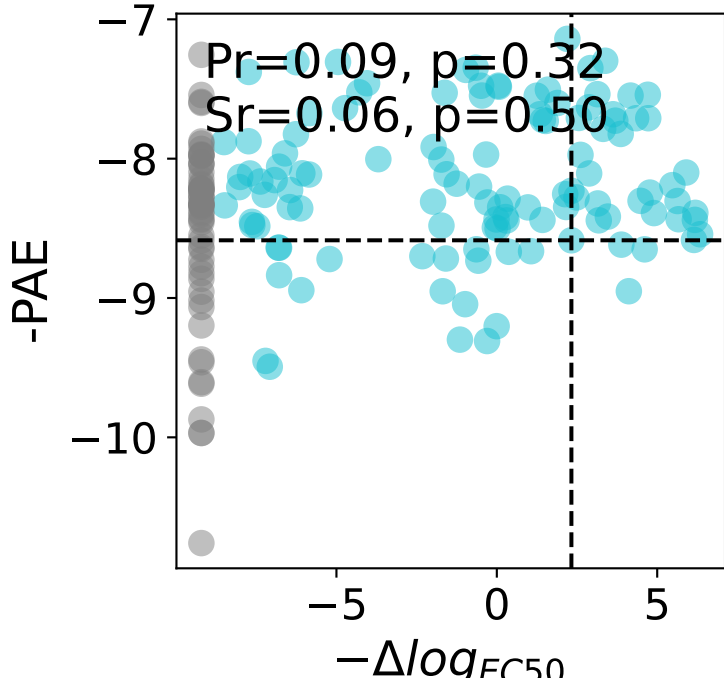
ProteinMPNN 0.02



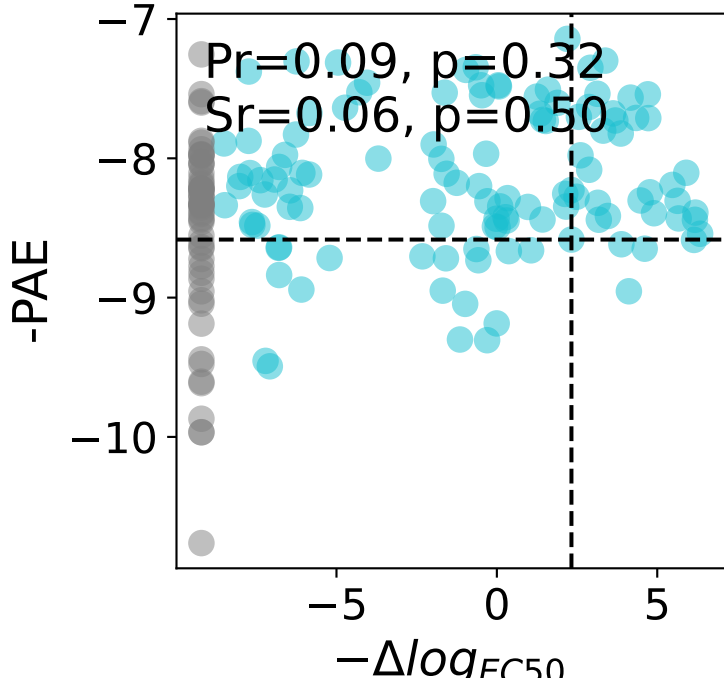
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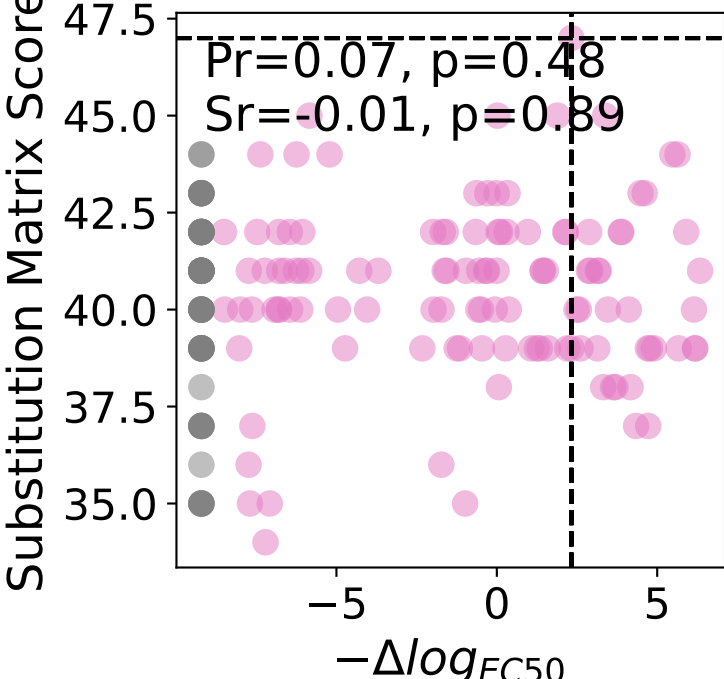
TCRdock



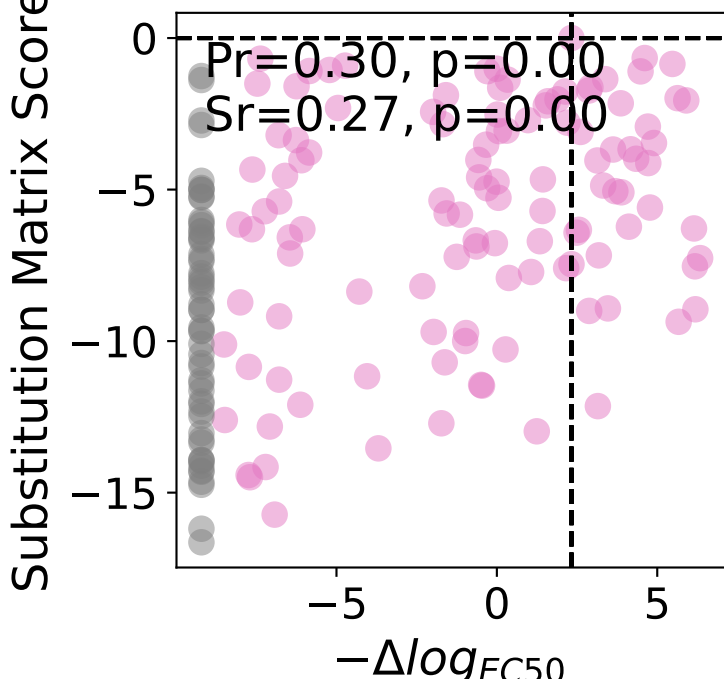
TCRdock benchmark



BLOSUM62



Luksza et al. C



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