

Investigating the dissociation process and binding free energy of the p53-DBD/DNA complex

by PaCS-MD and MSM

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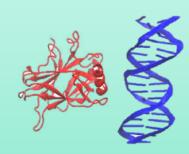
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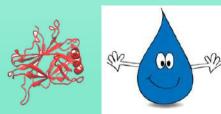


Dissociation simulation

poster number

1P-37

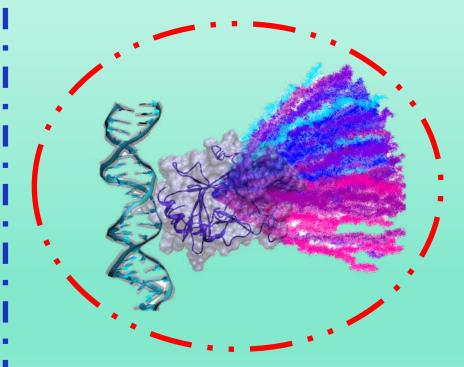






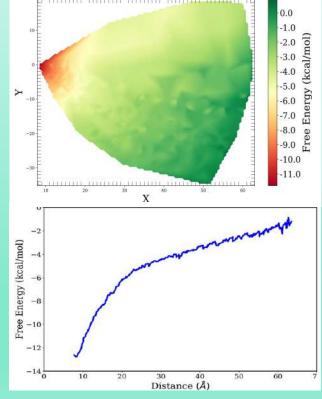
Parallel cascade selection molecular dynamics

Dissociation pathways



75 different dissociation pathways obtained by ran PaCS-MD for five different starting conformations

Free energy analysis by MSM



Binding free energy

- Predicted value: 11.8 kcal/mol
- Experimental value: 11.1 kcal/mol