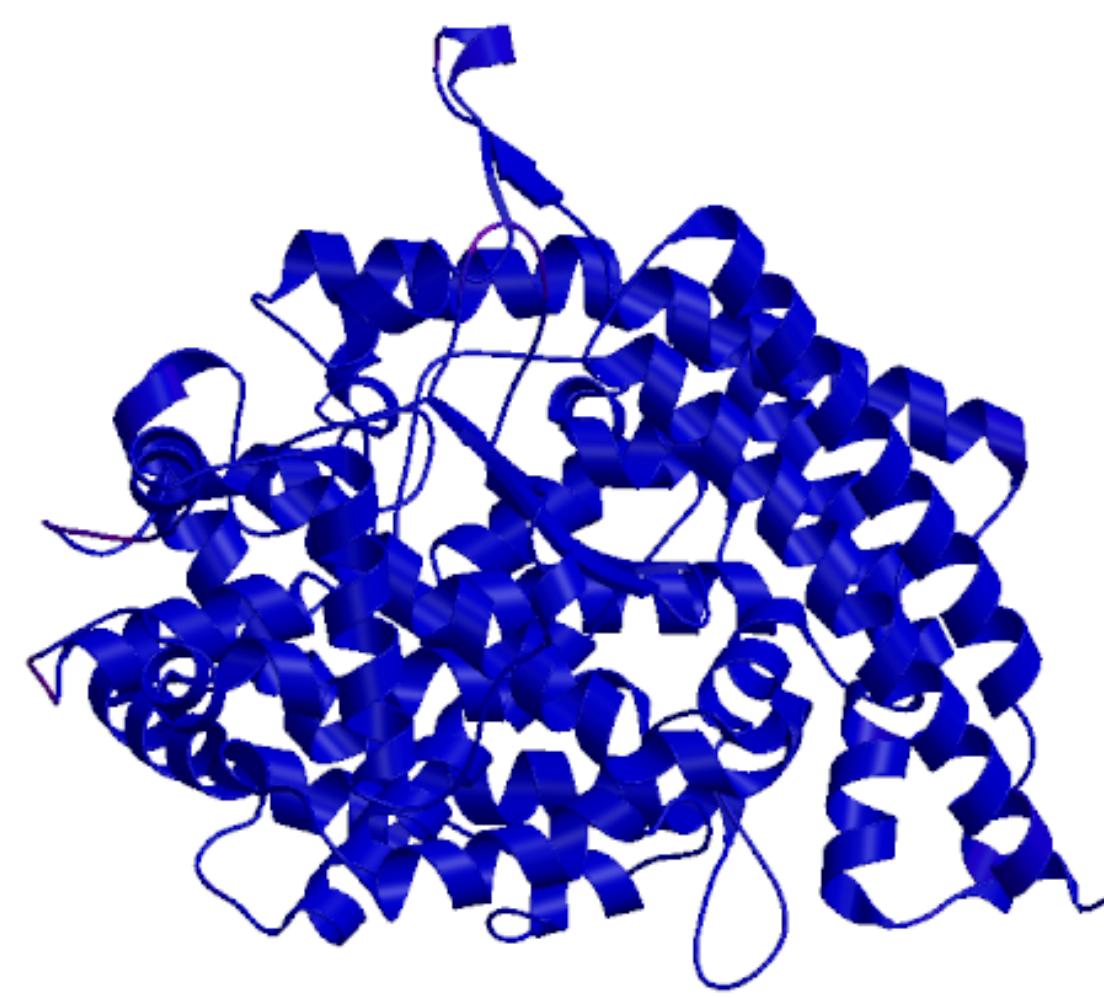
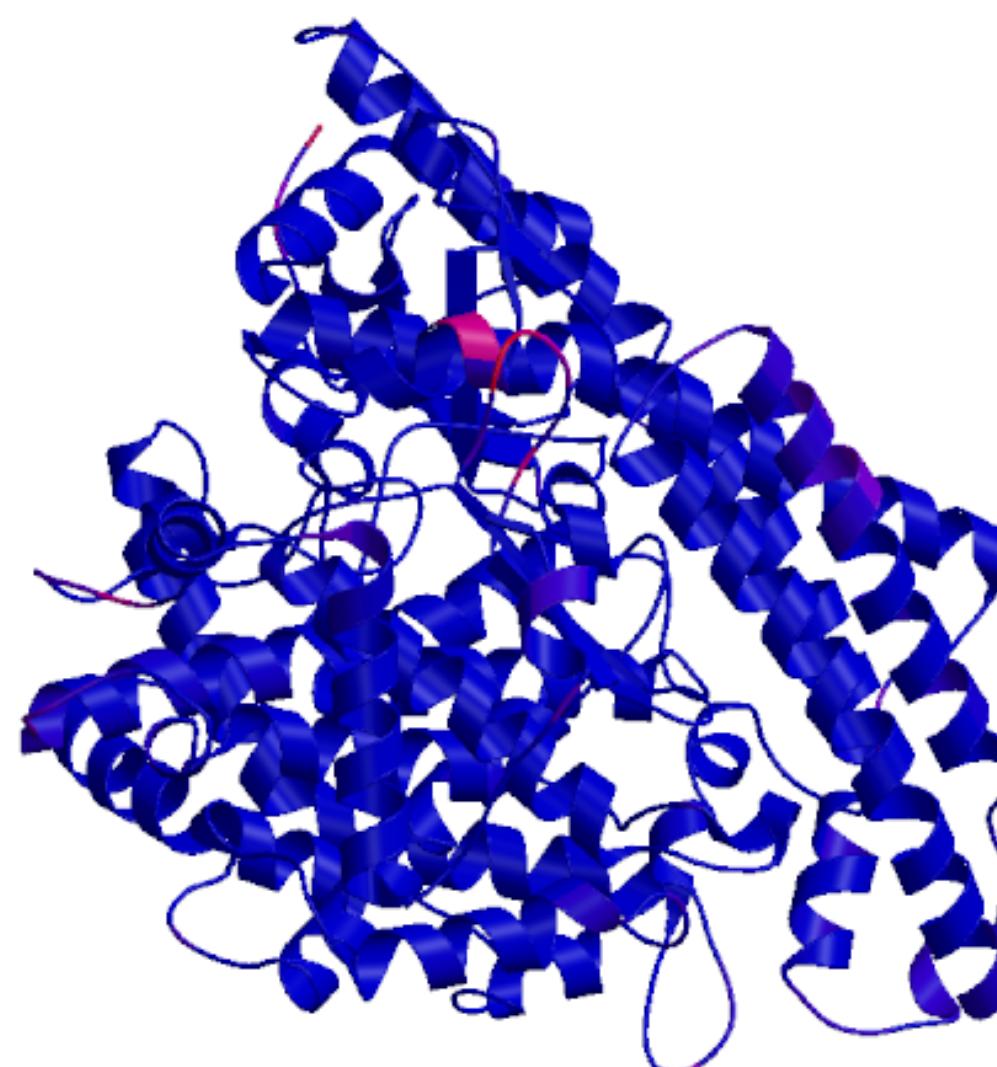
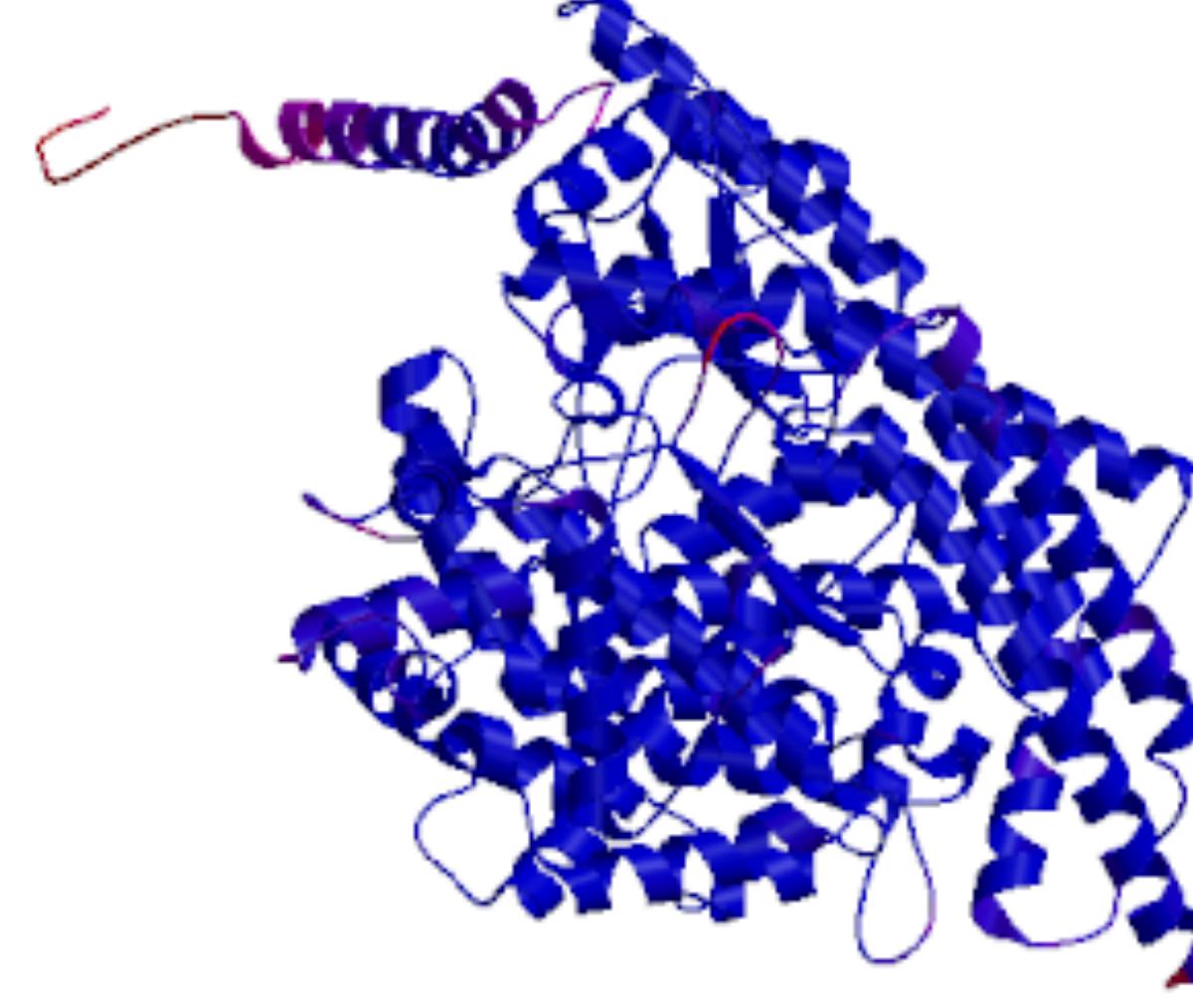


6LZG/6M0J

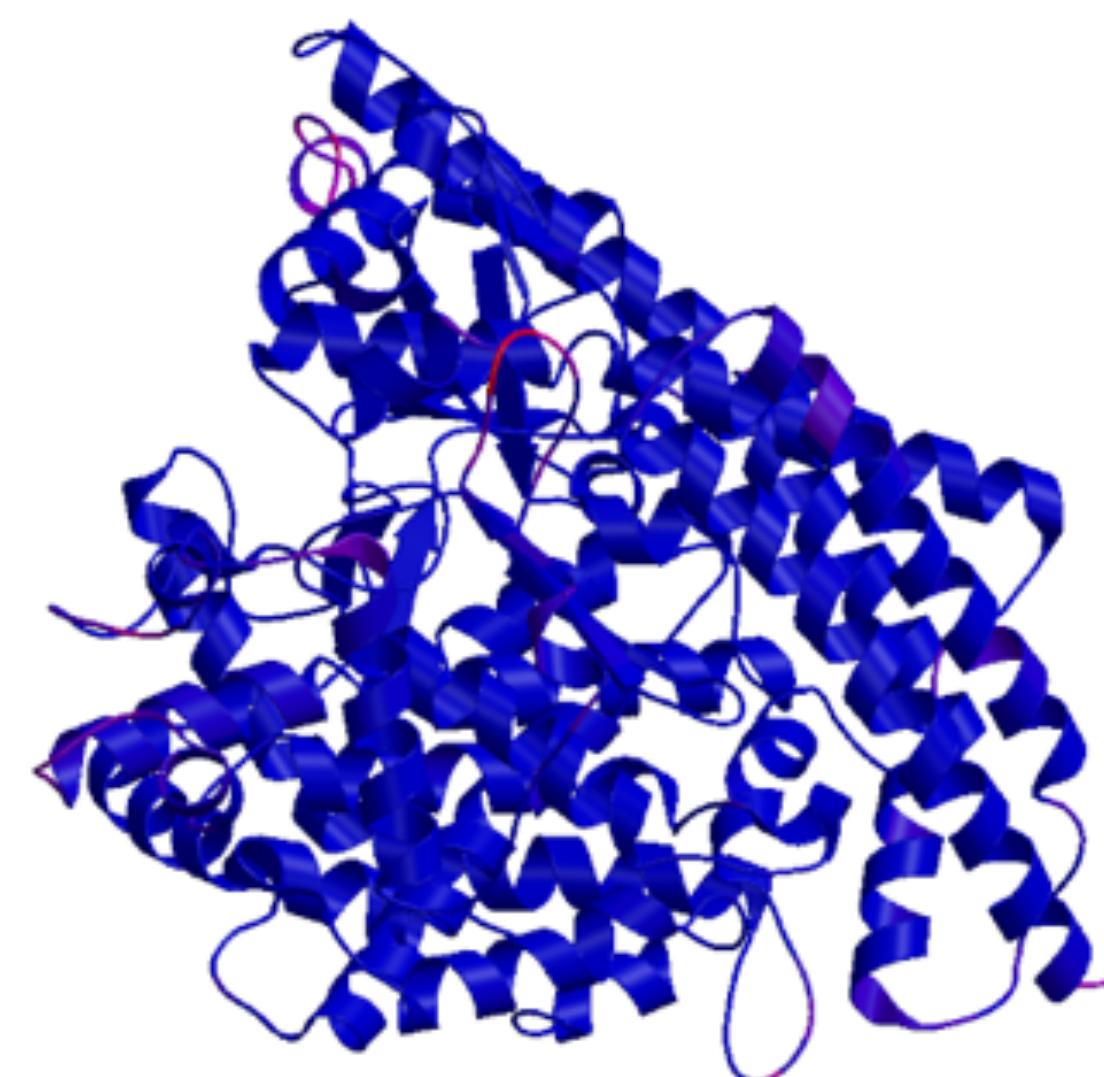
Global Score
0.88 ± 0.05

Hdock WT model

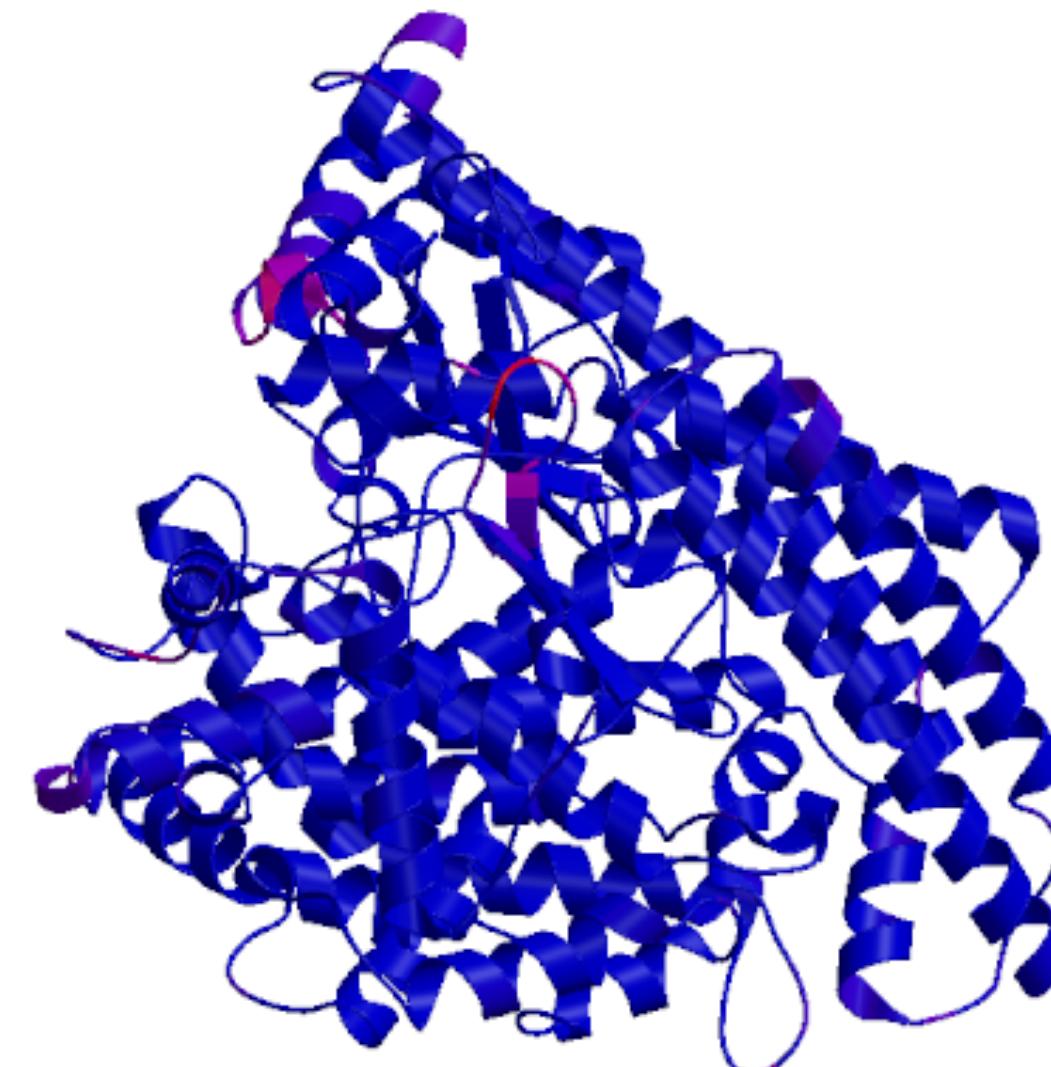
Global Score
0.84 ± 0.05

Hdock K26R model

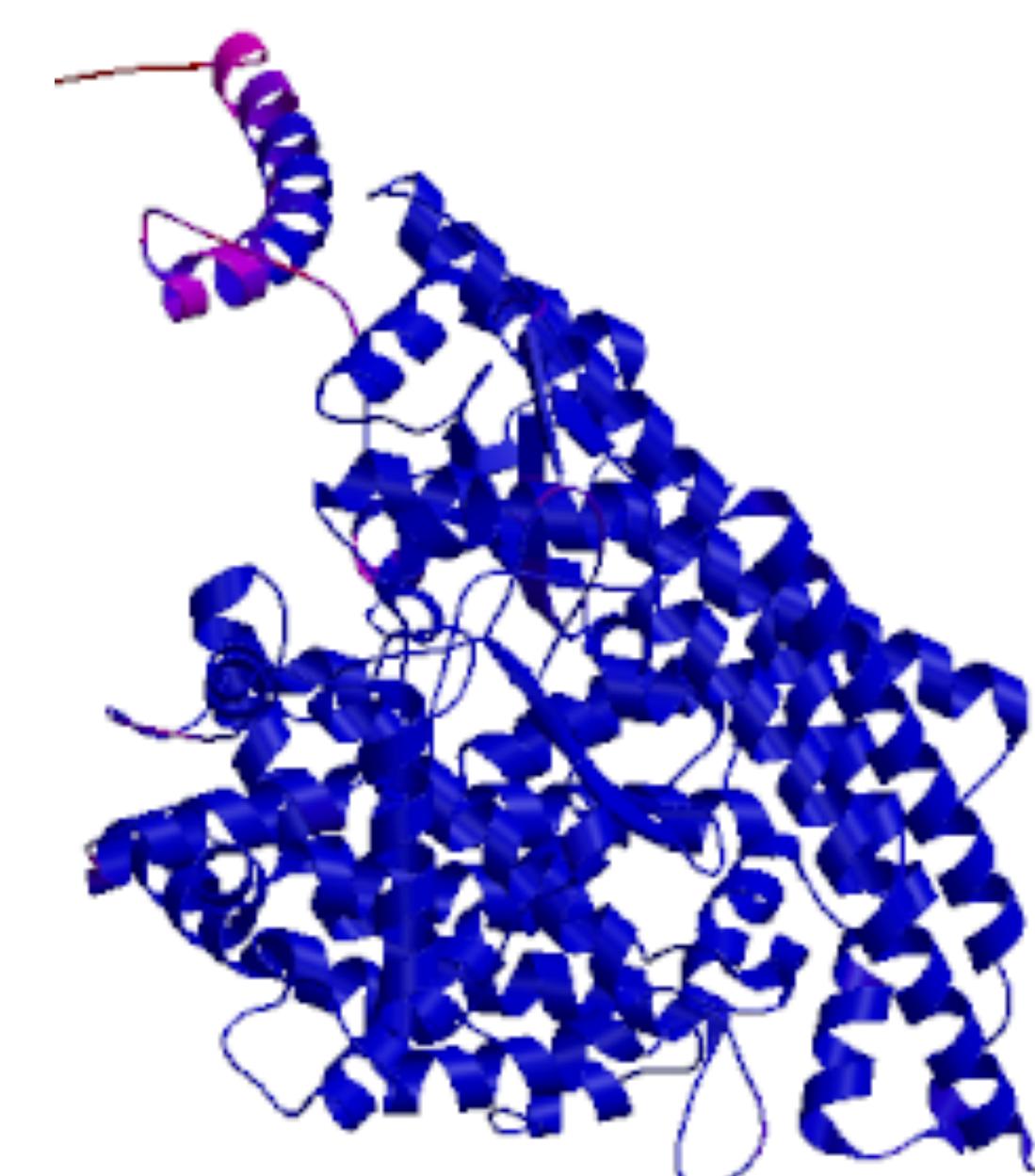
Global Score
0.83 ± 0.05

6M17

Global Score
0.83 ± 0.05

Hdock K26R model

Global Score
0.83 ± 0.05

Hdock K26R model

Global Score
0.83 ± 0.05

MolProbity

6M17	Bad bonds: 0.03%
6M17	Bad angles: 0.01%
Modeller WT	Bad bonds: 0.00%
Modeller WT	Bad angles: 1.24%
Modeller K26R	Bad bonds: 0.00%
Modeller K26R	Bad angles: 1.24%
K26R (Hdock) - Homology model	Bad bonds: 0.00%
K26R (Hdock) - Homology model	Bad angles: 0.77%
K26R (Hdock) - Docking model	Bad bonds: 0.00%
K26R (Hdock) - Docking model	Bad angles: 0.77%
WT (Hdock)	Bad bonds: 0.00%
WT (Hdock)	Bad angles: 0.80%