

scoring function	S(1)	S(3)	S(5)
RF-Score-VS v2	8.60 ± 3.27	6.94 ± 3.18	6.27 ± 3.29
rDock (dock)	6.81 ± 3.41	5.60 ± 3.01	5.23 ± 2.82
rDock (dock_solv)	6.92 ± 3.33	5.80 ± 2.94	5.12 ± 2.69
LigandRNA (2013)	6.67 ± 2.82	5.51 ± 2.31	4.92 ± 2.38
LigandRNA (updated)	6.37 ± 2.74	5.49 ± 2.23	5.00 ± 2.36
AnnapuRNA DL (2013)	5.39 ± 2.85	4.58 ± 2.47	4.37 ± 2.21
AnnapuRNA DL (2016)	6.68 ± 3.04	4.82 ± 2.66	4.15 ± 2.24
AnnapuRNA kNN (2013)	6.06 ± 2.92	4.86 ± 2.56	4.14 ± 2.18
AnnapuRNA kNN (2016)	5.76 ± 2.87	5.00 ± 2.64	4.35 ± 2.32
minimum RMSD	2.90 ± 1.76	2.90 ± 1.76	2.90 ± 1.76
median RMSD	7.61 ± 2.70	7.61 ± 2.70	7.61 ± 2.70