

sRNA	L	G (Kcal/mol)	V	P	R_m	E	R_e
C0067	125	-22.99	5.48	0.380	0.911	-0.095	0.822
C0293	73	-22.97	4.06	0.279	0.743	0.134	0.681
C0299	79	-13.34	2.80	0.312	0.889	-0.034	0.858
C0343	75	-7.82	2.54	0.434	0.864	0.255	0.847
C0362	386	-174.51	16.28	0.303	0.965	-0.084	0.768
C0465	78	-20.26	3.96	0.299	0.876	-0.052	0.851
C0614	87	-22.84	4.13	0.551	0.859	0.176	0.822
C0664	113	-56.41	4.91	0.609	0.755	0.215	0.577
C0719	222	-62.47	9.43	0.574	0.895	0.111	0.674
csrB	360	-134.57	19.20	0.292	0.946	0.025	0.754
csrC	245	-97.61	9.62	0.255	0.934	-0.008	0.783
dicF	53	-17.74	2.79	0.159	0.889	-0.214	0.907
dsrA	87	-33.99	4.98	0.284	0.909	0.003	0.873
gadY	105	-30.71	6.71	0.365	0.850	0.123	0.782
gcvB	205	-67.66	11.12	0.349	0.916	0.117	0.836
IS128	209	-57.16	8.49	0.254	0.961	0.073	0.883
isrA	158	-49.23	7.01	0.506	0.868	0.164	0.762
isrB	160	-46.13	7.78	0.438	0.922	0.084	0.831
isrC	204	-65.24	10.14	0.246	0.951	0.128	0.884
micA	72	-23.71	4.68	0.187	0.904	0.037	0.939
micC	109	-28.46	3.93	0.214	0.908	-0.035	0.866
micF	93	-19.30	3.07	0.078	0.955	-0.020	0.961
omrA	88	-31.64	4.92	0.315	0.861	0.142	0.868
omrB	76	-27.96	4.00	0.327	0.854	0.136	0.778
oxyS	110	-35.12	4.33	0.257	0.915	0.065	0.867
psrD	169	-47.37	8.03	0.526	0.873	0.144	0.765
psrN	188	-93.36	9.18	0.299	0.940	0.111	0.879
psrO	174	-56.26	7.41	0.363	0.946	-0.033	0.852
rdIA	67	-28.15	2.25	0.122	0.921	0.055	0.935
rdIB	66	-28.18	2.39	0.160	0.928	-0.101	0.920
rdIC	68	-28.66	2.42	0.157	0.923	0.070	0.930
rdID	64	-28.76	2.66	0.147	0.919	0.002	0.941
rprA	105	-29.23	4.45	0.318	0.887	0.109	0.892
rseX	91	-18.77	3.77	0.348	0.892	-0.134	0.856
rttR	171	-59.60	6.56	0.449	0.886	0.252	0.794
rybA	89	-21.15	3.81	0.518	0.746	0.227	0.674
rybB	81	-26.62	4.17	0.216	0.908	0.108	0.894
rydB	68	-18.25	2.28	0.364	0.814	0.287	0.817
rydC	64	-15.94	2.50	0.609	0.705	0.355	0.762
ryeA	249	-87.97	13.62	0.121	0.916	0.048	0.699
ryeB	121	-32.85	6.26	0.304	0.922	-0.112	0.849
ryeC	143	-58.60	6.93	0.280	0.949	0.020	0.927
ryeD	136	-63.46	7.11	0.167	0.944	0.024	0.895
ryeE	86	-21.17	3.78	0.180	0.913	0.180	0.898
ryfA	304	-140.46	14.71	0.147	0.942	-0.054	0.731
ryfB	319	-115.30	12.54	0.460	0.913	0.148	0.724
ryfC	77	-30.71	3.12	0.122	0.926	0.152	0.950
ryfD	143	-56.43	5.88	0.229	0.897	0.040	0.741
rygC	140	-61.63	8.48	0.088	0.933	-0.227	0.875
rygD	145	-57.88	6.46	0.227	0.940	-0.115	0.839
rygE	142	-58.40	5.95	0.198	0.918	0.103	0.865

ryhA	108	-32.75	4.60	0.682	0.846	0.117	0.714
ryhB	90	-26.39	4.71	0.408	0.824	0.260	0.812
ryjA	140	-54.40	7.72	0.281	0.929	0.146	0.842
ryjB	90	-33.58	3.17	0.385	0.757	0.227	0.785
sgrS	227	-87.79	8.90	0.341	0.919	0.126	0.793
sokB	56	-24.94	1.19	0.090	0.901	0.124	0.963
sokC	55	-22.45	1.99	0.186	0.921	0.088	0.956
spf	109	-45.29	5.42	0.290	0.856	0.191	0.831
sraA	57	-12.89	2.13	0.336	0.896	0.090	0.869
sroA	93	-33.88	5.36	0.365	0.851	0.173	0.814
sroB	82	-25.30	4.65	0.221	0.932	-0.089	0.917
sroC	163	-70.61	8.34	0.146	0.954	-0.095	0.908
sroD	86	-32.47	4.28	0.196	0.889	0.177	0.871
sroE	92	-40.66	3.69	0.131	0.934	0.041	0.942
sroG	149	-56.90	8.11	0.424	0.860	0.189	0.710
sroH	161	-64.20	8.56	0.188	0.934	0.010	0.882
symR	77	-40.65	2.24	0.097	0.941	-0.124	0.969
tff	136	-49.32	4.75	0.111	0.943	-0.188	0.869
tp2	161	-52.89	8.87	0.533	0.893	0.065	0.714
tpke11	89	-37.09	3.96	0.142	0.952	-0.088	0.929
tpke70	436	-150.42	16.60	0.384	0.940	-0.095	0.609
istR-1	75	-33.22	3.15	0.112	0.899	0.181	0.908
istR-2	140	-52.87	7.18	0.493	0.889	0.058	0.739
ssrA	363	-136.43	18.05	0.234	0.925	-0.147	0.529
ssrS	183	-86.10	7.57	0.322	0.919	0.105	0.819
glmZ	172	-47.86	7.61	0.473	0.903	0.119	0.780
glmY	184	-70.41	10.26	0.210	0.931	-0.050	0.795
ffs	114	-63.39	4.21	0.198	0.918	0.011	0.841