

Score: -89		P value: 9.81557e-05	
sRNA3	41	GUUUUUCUUCGCA-UUC-UUUUUUAUC	65
		: :	
STM2284 (<i>glpA</i>)	-2	CAAAAAGGAGUGUCAAGCAAAAAAUAG	-28
Score: -87		P value: 4.25713e-05	
sRNA5	40	AGCUUUUAUUGACUAUUCUUUUU	62
		:	
STM1875 (<i>yobA</i>)	6	UCGGUAUAAGUGAUAAGGAAAAA	-17
Score: -87		P value: 0.000239255	
sRNA7	3	GCGAUUAAGC-UGUUAUAGCUUGAUUAUCUGUGACGUAUAGAGCGCU	49
		: : : : : : : : : : : : :	
STM0687 (<i>ybfM</i>)	18	CGGUGAUUUGCAUGCGUAUUCAAUUAGGAGAAACUGCAUAUUUAACGA	-30
Score: -86		P value: 0.000540291	
sRNA11	242	AUAUUUUUCUCGGAUUAUAUUGAG	265
		:	
STM3984 (<i>yigG</i>)	2	UAUAAAAA-AGUCUAAUAUAAAUUC	-21
Score: -78		P value: 0.000659353	
sRNA16	26	GAAACGAGACAGUGAACAGUACCGUUUUCUUCGGUUUUUA-UGCCAUAGUA	75
		: : : : :	
STM0225 (<i>ynaI</i>)	19	CUUUUUUAAGUCACUUGUAAUAACAAAAG--GCGAGGAAGGAUGCUAUUGU	-30
Score: -86		P value: 0.000760695	
sRNA17	291	ACAAGGAC-GCCAGCCACUUUUUAGUUGUUUGAA	323
		: :	
STM0938 (<i>ybjE</i>)	4	UGUACUUGUCGGCCCGGAAAAAAUAACAAACUU	-30

Score: -79	<i>P</i> value: 0.000144163		
sRNA18	8	AUUUUCUGUAGCACUAUAAGUAUUAAUUUAUUGUUUUUUU : : ::	47
STM1403 (<i>sscB</i>)	18	UAGAAGAAA--GUAGUAGU-AUGGAUAACUAACAAUAAAA	-19
Score: -89	<i>P</i> value: 4.07964e-05		
sRNA19	28	AUAACUUCAUUAUUGAAAUAAAUAUAUGC :	57
STM1426 (<i>ribE</i>)	2	UAUAGAAGUUGAGGACUUUAUUUAUUUUCG	-28
Score: -98	<i>P</i> value: 2.24624e-05		
sRNA20	68	AUU-UUCGUUCAUUAACGCGCCUCUUGUC-UCCUGAAAAGUA : :	107
STM2154 (<i>mrp</i>)	13	UAACAAGCAAGUAGGGCUGAGGAAAAACAUAAGACUUUUCAU	-29
Score: -250	<i>P</i> value: 4.66294e-15		
sRNA21	226	AGGGGAGGGGAAGCAUUCAAUCGUAUGGUUGGCUUUAUGAAAAACGGUGU 	275
STM4316	20	UCCCCUCCCCUUCGUAAGUUAGCAUACCAACCGAAAUAUUUUUGCCACA	-30
Score: -74	<i>P</i> value: 0.000359706		
sRNA22	13	CGCUGUCGUUUAAAAAAUAAAAAGA : : : :	36
STM3773	-3	GUGAUGGCGAAUGCUUUUUUUUCU	-26
Score: -67	<i>P</i> value: 0.00299573		
sRNA23	5	UUCGUGAAGGAAUGUUCUAA 	24
STM0224 (<i>rcsF</i>)	-6	AAGGAGUUACUACAAGAUU	-25

Score: -87	<i>P</i> value: 5.17551e-05		
sRNA24	29	UUGUUUUU-A-UUAUUUCAUUUUUACAUUUUUGUUAGUG	65
		: :: :	
STM4370 (<i>hemE</i>)	20	AGCAAAACUCAAGCCAGUAAAAUGUAAGGA-AAUCGC	-18

Score: -84	<i>P</i> value: 0.000741386		
sRNA25	91	AUCCGGCACCUGAGGAAUA	109
STM3766	-5	UAGGCAGUGGACUCCUUAU	-23

Score: -88	<i>P</i> value: 2.17524e-05		
sRNA26	2	AUUUUUAAAAUUUUUUGUCGUUAAACUUUCUUGAUUUUGAAUUUUUAUGC	51
		:: : :	
STM1379 (<i>orf48</i>)	19	UAAAAACUGAAAAAAGUA-CUAUUUAUGGAAUUA---CCGAAAAUACG	-27

Score: -77	<i>P</i> value: 0.000272471		
sRNA27	39	UUUAUCAUUUAUUUAUCAUUA	60
STM4206	20	AAAUUUUAUUAAAAUAGUAAU	-2

Score: -94	<i>P</i> value: 9.99577e-05		
sRNA28	67	AAAAAAUGUAAUUUAGCAUUUUUGUUUUUUUAAUGAAUUUAUACCAAAAA	120
		: :	
STM0276	20	UUUUUUUACAGUAU-UCGGUAAAAAUAGGAA--UAAUUCAAA-AUGAAUUUU	-30

Score: -107	<i>P</i> value: 2.12733e-05		
sRNA29	108	UUUUUAUCAUUUUA-AUAGUUCUAUACAUUCUUUUUUCUAUGGAAAAC	156
		: : : :	
STM0335	16	AAUAAUAGUAAAGUAUAUCAAUUAUCUGAGG-UAG----UUUCUUUUG	-29

Score: -87	<i>P</i> value: 0.00057811		

sRNA30	68	CUUUGUGCUAUUGUAAUCAUA-ACCAUUCUCAUUUACACUUUGUGCGGAAAU : : : : :	118
STM3138	20	GGAUUAUGAAAAAAUUGUAUAUGGUAAGAGGAA---UAAUAUACACCUGUA	-29

Score: -85	P value: 0.000315897		
sRNA31	4	UAAAAUCAUAGGGUUAUUUAUUGUCGGC : : : : : :	31
STM4162 (<i>thiF</i>)	19	AUUUUAGUA--CCAGUAAGUAGCGGCUG	-7

Score: -95	P value: 5.20344e-05		
sRNA32	94	CAGGGAGGU-CUCAUUCUUUAUUUAUUGCUCACACUGUGUGGGUGUGU : : : : : :	137
STM3630 (<i>dppA</i>)	15	GUUCCUUUAUGAGUAAGAAG-----ACGAGGUUA-ACACUACAUA	-24

Score: -89	P value: 0.000195753		
sRNA33	46	UAAUCACGCGAUUAAUUAUACCACGCUAAUUGUUUCU : :	83
STM1417 (<i>ssaP</i>)	10	AUUA-UGCG-UAGCGAAUUAUGGUAUCAUUA AAAAAGA	-26