							Lege	nd	
				Difference order of m	in E-value			ches E-valu	e E-value
				of best hit		Graphical representation of E-values of best hits	RNA sear	ches E-valu	
Ribnucleo- protein complex	Protein name and length	RNA name and length	Query (Q) and Target (T)	Protein: BLASTP- BLASTN	RNA: Infernal - BLASTN	1 1e-50	1e-100	1e-1	50 1e-200
						BLASTN E=6e-43			BLASTP E=6e-159
Signal Recognition Peptide (SRP)	ffh 453aa 1362nt	SRP RNA 100nt	E. coli (Q) B.subtilis (T)	116	5	Exde-8 BLASTN Covvil.00			cov=0.95
Ribosome	rpl30p 158aa 477nt	5S rRNA 115nt	S.solfataricus (Q) T.kodakarensis (T)	33	1	BLASTN BLASTP Ede-33 cov=0.04		•	·
						BLASTN E=8e-79		•	BLASTP E=3e-180
Lysine riboswitch	lysC 409aa 1230nt	Lysine riboswitch 187nt	B.cereus (Q) B.subtilis (T)	101	>=18	PLASTN Infernal Not found E-16-18 cov-1.00			cov=0.97
RNase P	rpp29	RNase P	M.jannaschii (Q)	17	25	BLASTN BLASTP E=6e-5 E=2e-22 cov=0.18 cov=0.94		•	
	95aa 288nt	RNA 258nt	P.horikoshii (T)			BLASTN Infemal E=2e-9 E=5e-34 cov=0.19 cov=1.00			
glmS	glmS	glmS	B.subtilis (Q)	163	13	BLASTN E=8e-11 cov=0.07			BLASTP E=3e-174 cov=1.00
ribozyme	600aa 1803nt	riboswitch 168nt	C.acetobutylicum (10	BLASTN Infernal E=4e-8 E=4e-21 cov=0.28 cov=1.00			
			- "			BLASTN E=1e-28 cov=0.15			BLASTP E=~0 cov:1.00
Cobalamin riboswitch	btuB 614aa 1845nt	Cobalamin 191nt	E. coli (Q) Y.enterocolitica (T)	>=152	9	BLASTN Infernal E=2e-13 E=1e-22 cov=0.54 cov=1.00			
						BLASTN BLASTP E=8e-5 E=2e-42 cov=0.04 cov=0.97	•		
Cobalamin riboswitch	btuB 614aa 1845nt	Cobalamin 191nt	E. coli (Q) A.baumannii (T)	37	>=4	niformal P=8e-4 BLASTN cov=0.72 Not found			