

locus	log2ratio	lowerCI	upperCI	quantified_peptic	description
sp_P0A6X3_HFQ_ECOLI	4.5	4	5	7	Protein hfq OS=Escherichia coli
sp_P0AG67_RS1_ECOLI	3.4	3.1	3.7	22	30S ribosomal protein S16
sp_P0A7V8_RS4_ECOLI	3.1	2.5	3.8	4	30S ribosomal protein S16
sp_P0A6P9_ENO_ECOLI	2.3	1.7	2.9	7	Enolase OS=Escherichia coli
sp_P30850_RNB_ECOLI	2.2	1.5	3	3	Exoribonuclease 2 OS=Escherichia coli
sp_P02359_RS7_ECOLI	2.1	1.5	2.7	6	30S ribosomal protein S16
sp_P0A7D4_PURA_ECOLI	2	1.4	2.5	9	Adenylosuccinate synthetase
sp_P0A7L0_RL1_ECOLI	2	0.8	2.8	4	50S ribosomal protein L16

chia coli (strain K12) GN=hfq PE=1 SV=2 [MASS=11166]
51 OS=Escherichia coli (strain K12) GN=rpsA PE=1 SV=1 [MASS=61158]
54 OS=Escherichia coli (strain K12) GN=rpsD PE=1 SV=2 [MASS=23469]
i coli (strain K12) GN=eno PE=1 SV=2 [MASS=45655]
Escherichia coli (strain K12) GN=rnb PE=1 SV=3 [MASS=72491]
57 OS=Escherichia coli (strain K12) GN=rpsG PE=1 SV=3 [MASS=20019]
etase OS=Escherichia coli (strain K12) GN=purA PE=1 SV=2 [MASS=47345]
.1 OS=Escherichia coli (strain K12) GN=rplA PE=1 SV=2 [MASS=24730]