

S2 Table: Proteins identified by proteomic analysis in epimastigotes under nutritional stress.

ID	Description	Spectral count
Tc00.1047053503405.10	Glutamine synthetase, putative	7
Tc00.1047053509793.50	Hypothetical protein, conserved	7
Tc00.1047053509151.140	60S ribosomal protein L23a, putative	6
Tc00.1047053510767.20	60S ribosomal protein L36, putative	8
Tc00.1047053506977.49 Tc00.1047053509671.64	Ribosomal protein L36, putative	8
Tc00.1047053511211.160 Tc00.1047053511211.170 Tc00.1047053510439.61	Heat shock protein 70 (HSP70), putative	7
Tc00.1047053510089.110 Tc00.1047053510421.130	Hypothetical protein, conserved	7
Tc00.1047053510755.129 Tc00.1047053508413.44 Tc00.1047053511071.171 Tc00.1047053504037.30	60S ribosomal protein L12, putative	9
Tc00.1047053506297.160 Tc00.1047053506297.150 Tc00.1047053510101.170 Tc00.1047053510101.180	40S ribosomal protein S5, putative	8
Tc00.1047053508475.10 Tc00.1047053508823.120 Tc00.1047053508823.140	Ribosomal protein S20, putative	10