

Accession	Description	MW	calc. pI
P68206	UPF0337 protein YbjJ	8,325	5.44
P0A7M2	50S ribosomal protein L28	9,006	11.42
P0ADZ4	30S ribosomal protein S15	10,269	10.4
P0A6X3	RNA-binding protein Hfq	11,166	6.97
P0AEN8	L-fucose mutarotase	15,473	5.57
P39177	Universal stress protein G	15,935	6.03
P0A9A9	Ferric uptake regulation protein	16,795	5.68
P0AGD1	Superoxide dismutase [Cu-Zn]	17,681	5.95
P0AEZ9	Molybdenum cofactor biosynthesis protein B	18,665	5.73
P0ABT2	DNA protection during starvation protein	18,695	5.7
P0A9K9	FKBP-type peptidyl-prolyl cis-trans isomerase SlyD	20,853	4.86
P0A8E1	UPF0227 protein YcfP	21,226	6.13
P07464	Galactoside O-acetyltransferase	22,799	6.23
P0ACJ8	cAMP-activated global transcriptional regulator CRP	23,640	8.38
P76344	Metal-binding protein ZinT	24,762	5.91
P61517	Carbonic anhydrase 2	25,097	6.16
P62577	Chloramphenicol acetyltransferase	25,663	5.91
P0AA43	Ribosomal small subunit pseudouridine synthase A	25,865	5.75
P0AEQ3	Glutamine-binding periplasmic protein	27,190	8.44
P39831	NADP-dependent 3-hydroxy acid dehydrogenase YdfG	27,249	5.65
P0AA16	Transcriptional regulatory protein OmpR	27,354	6.04
P62707	2,3-bisphosphoglycerate-dependent phosphoglycerate mutase	28,556	5.85
P60422	50S ribosomal protein L2	29,860	10.93
P62593	Beta-lactamase TEM	31,515	5.69
P0A910	Outer membrane protein A	37,200	5.99
P21151	3-ketoacyl-CoA thiolase	40,876	6.31
P23908	Acetylornithine deacetylase	42,347	5.54
P77581	Succinylornithine transaminase	43,665	5.91
P0AFG6	Dihydrolipoyllysine-residue succinyltransferase component of 2-oxoglutarate dehydrogenase complex	44,011	5.58
P00393	NADH dehydrogenase	47,359	8.96
P0A6U8	Glycogen synthase	52,822	6.14
P0AC53	Glucose-6-phosphate 1-dehydrogenase	55,704	5.55
P0A6F5	60 kDa chaperonin	57,329	4.85
P17169	Glutamine--fructose-6-phosphate aminotransferase [isomerizing]	66,894	5.56
P77398	Bifunctional polymyxin resistance protein ArnA	74,289	6.39
P0AFG3	2-oxoglutarate dehydrogenase E1 component	105,062	6.04