

TAILS *Lm* supernatant (proteins found in 3 replicates)

Master Protein Descriptions

Phosphatidylcholine cholinephosphohydrolase
Quinol oxidase subunit 2
NusG_II domain-containing protein
Cysteine synthase
Uncharacterized protein
Glycerophosphoryl diester phosphodiesterase
Lipoprotein
Internalin B
Tagatose 1,6-diphosphate aldolase
Spermidine/putrescine transport system permease
Sulfatase domain-containing protein
Uncharacterized protein
GW domain-containing protein
Uncharacterized protein
ABC transport system
Signal peptidase I
DNA gyrase subunit B
Universal stress protein
Protease IV
Uncharacterized protein
Peptide/nickel transport system substrate-binding protein
G5 domain-containing protein
Putative septation protein SpoVG
Aspartyl/glutamyl-tRNA(Asn/Gln) amidotransferase subunit B
50S ribosomal protein L25
YycH protein
Thiol-activated cytolysin
DivIC protein
Fumarate reductase flavoprotein subunit
Probable transcriptional regulatory protein LMRG_00061
Peptidoglycan-N-acetylglucosamine deacetylase PgdA
Probable succinyl-diaminopimelate desuccinylase
LytR_cpsA_psr domain-containing protein
10 kDa chaperonin
Manganese-binding lipoprotein mntA
DD-transpeptidase
4-hydroxy-tetrahydrodipicolinate reductase
Histidine kinase
Flagellar hook protein FlgE
Phage capsid protein
Aminotransferase
Peptidoglycan bound protein
CBM6 domain-containing protein
30S ribosomal protein S5
Phosphoglycerol transferase
FeS assembly protein SufD
Uncharacterized protein

Accessions

A0A0H3G8P1
A0A0H3G8T0
A0A0H3G991
A0A0H3G9A7
A0A0H3G9Q4
A0A0H3G9S8
A0A0H3G9T8
A0A0H3G9Y9
A0A0H3GA75
A0A0H3GAF1
A0A0H3GAG2
A0A0H3GBD8
A0A0H3GBE7
A0A0H3GBP9
A0A0H3GBZ4
A0A0H3GC68
A0A0H3GCB7
A0A0H3GCF8
A0A0H3GCG3
A0A0H3GCN8
A0A0H3GCP2
A0A0H3GCT9
A0A0H3GCV1
A0A0H3GCW2
A0A0H3GCW8
A0A0H3GD67
A0A0H3GD84
A0A0H3GD99
A0A0H3GDC3
A0A0H3GDD6
A0A0H3GDH9
A0A0H3GDI0
A0A0H3GDK2
A0A0H3GDS3
A0A0H3GDS9
A0A0H3GDX6
A0A0H3GDZ1
A0A0H3GE39
A0A0H3GEC2
A0A0H3GEG6
A0A0H3GEK9
A0A0H3GES0
A0A0H3GET7
A0A0H3GFB3
A0A0H3GFC6
A0A0H3GFE5
A0A0H3GFH4

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|---|------------|
| Potassium-transporting ATPase KdpC subunit | A0A0H3GFI3 |
| Glyceraldehyde-3-phosphate dehydrogenase | A0A0H3GFI9 |
| HTH lacI-type domain-containing protein | A0A0H3GFM6 |
| Peptidase_M23 domain-containing protein | A0A0H3GFN8 |
| SSD domain-containing protein | A0A0H3GFS0 |
| Peptide/nickel transport system substrate-binding protein | A0A0H3GFV1 |
| Signal peptidase I | A0A0H3GFV8 |
| 30S ribosomal protein S19 | A0A0H3GG04 |
| Elongation factor Tu | A0A0H3GG29 |
| Peptidoglycan bound protein | A0A0H3GGA1 |
| Transcription elongation factor GreA | A0A0H3GGF0 |
| N-acetylmuramoyl-L-alanine amidase | A0A0H3GGH0 |
| Transketolase | A0A0H3GGH6 |
| Glutamyl-tRNA(Gln) amidotransferase subunit A | A0A0H3GH17 |
| Multifunctional fusion protein | A0A0H3GH33 |
| Cell shape-determining protein MreC | A0A0H3GH53 |
| Actin-assembly inducing protein ActA | A0A0H3GH64 |
| 30S ribosomal protein S2 | A0A0H3GHH8 |
| Carboxyl-terminal processing protease | A0A0H3GI96 |
| Flagellin | A0A0H3GID7 |
| UPF0302 protein LMRG_01068 | A0A0H3GIF4 |
| Uncharacterized protein | A0A0H3GJ46 |
| Peptide/nickel transport system substrate-binding protein | A0A0H3GJB6 |
| Pyruvate dehydrogenase E1 component subunit alpha | A0A0H3GJB7 |
| 50S ribosomal protein L18 | A0A0H3GJD7 |
| FAD:protein FMN transferase | A0A0H3GJF7 |
| 30S ribosomal protein S12 | A0A0H3GJH5 |
| Polar amino acid transport system ATP-binding protein | A0A0H3GJI6 |
| Dihydroxyacetone kinase L subunit | A0A0H3GJL3 |
| dITP/XTP pyrophosphatase | A0A0H3GJR4 |
| Multiple sugar transport system substrate-binding protein | A0A0H3GJY9 |
| Uncharacterized protein | A0A0H3GK85 |
| Transcriptional regulator LytR | A0A0H3GKG2 |
| N-acetylmuramoyl-L-alanine amidase | A0A0H3GKJ7 |
| 50S ribosomal protein L14 | A0A0H3GKP3 |
| Peptidoglycan bound protein | A0A0H3GKR6 |
| Polar amino acid transport system substrate-binding protein | A0A0H3GKX3 |
| 60 kDa chaperonin | A0A0H3GLZ0 |
| Polar amino acid transport system substrate-binding protein | A0A0H3GMT4 |
| Enolase | A0A0H3GN27 |
| D-glutamyl-L-m-Dpm peptidase P45 | A0A0H3GN80 |

2D SDS-PAGE *Lm* supernatant (proteins found in 3 replicates)

| Master Protein Descriptions | Accessions |
|--|------------|
| Phosphatidylcholine cholinephosphohydrolase | A0A0H3G8P1 |
| Thiol-activated cytolysin | A0A0H3GD84 |
| Fumarate reductase flavoprotein subunit | A0A0H3GDC3 |
| Peptidase_M23 domain-containing protein | A0A0H3GFN8 |
| Peptidoglycan bound protein | A0A0H3GGA1 |
| Peptide/nickel transport system substrate-binding protein | A0A0H3GJB6 |
| Transcriptional regulator LytR | A0A0H3GKG2 |
| 60 kDa chaperonin | A0A0H3GLZ0 |
| D-glutamyl-L-m-Dpm peptidase P45 | A0A0H3GN80 |
| Internalin D | A0A0H3G8X7 |
| Flagellar hook capping protein | A0A0H3GA06 |
| Peptidoglycan bound protein | A0A0H3GAK6 |
| DUF5105 domain-containing protein | A0A0H3GAX9 |
| Multiple sugar transport system substrate-binding protein | A0A0H3GCT3 |
| Uncharacterized protein (PepSY domain-containing protein) | A0A0H3GCV0 |
| 1-phosphatidylinositol phosphodiesterase | A0A0H3GCV8 |
| Invasion-associated protein p60 (Probable endopeptidase p60) | A0A0H3GDY4 |
| Thioredoxin-like_fold domain-containing protein | A0A0H3GFB7 |
| Phosphocarrier protein HPr | A0A0H3GFK1 |
| Antigen A | A0A0H3GGY8 |
| Beta-lactamase domain-containing protein | A0A0H3GI09 |
| N-acetylmuramoyl-L-alanine amidase | A0A0H3GJB3 |
| DNA-directed RNA polymerase subunit alpha | A0A0H3GJC8 |
| Competence protein ComEA | A0A0H3GKB9 |
| Uncharacterized protein | A0A0H3GMD3 |
| Putative septation protein SpoVG | A0A0H3GD80 |
| Peptide/nickel transport system substrate-binding protein | A0A0H3GH20 |
| Penicillin binding protein 2B | A0A0H3GLV8 |
| FeS assembly ATPase SufC | A0A0H3GMZ0 |