| ComrEntrez II | O Gene Name | Prediction | Sorted Category |
|---------------|--|---|-------------------------|
| | 69/2-C-methyl-D-erythritol 4-phosphate cytidylyltransferase | Tededoff | Temporarly Up Regulated |
| | 191-deoxy-D-xylulose 5-phosphate reductoisomerase | | Down Regulated |
| | 31 D-ribose-binding periplasmic protein | | Up Regulated |
| | 22 Ribose transport system permease protein rbsC | | Ambiguous/Unsorted |
| | 33 Uncharacterized HTH-type transcriptional regulator yieP | n | Down Regulated |
| | | U. | |
| | 12 L-fucose mutarotase | | Up Regulated |
| | 17 Delta-aminolevulinic acid dehydratase | | Ambiguous/Unsorted |
| | 3 Protein tonB | | Ambiguous/Unsorted |
| | 64 Ribose import ATP-binding protein rbsA; Ribose import ATP-bin | | Ambiguous/Unsorted |
| | Pseudouridine synthase; Ribosomal large subunit pseudouridine | | Down Regulated |
| | Ribose operon repressor | | Down Regulated |
| | Putative potassium channel protein | | Ambiguous/Unsorted |
| rbsD 94826 | 67 D-ribose pyranase | | Ambiguous/Unsorted |
| | 64 DNA replication protein dnaC | | Ambiguous/Unsorted |
| ydcl 94886 | 65 Uncharacterized HTH-type transcriptional regulator ydcl | [Transcription', 'RNA related', 'Type of regulation', 'Genetic unit regulated', 'transcription', 'regulation of transcription DNA depende | Down Regulated |
| btuR 94583 | 39 Cob(I)yrinic acid a,c-diamide adenosyltransferase | | Ambiguous/Unsorted |
| hsdS 94886 | Type-1 restriction enzyme EcoKI specificity protein | | Down Regulated |
| | 37 Aldehyde-alcohol dehydrogenase | | Down Regulated |
| | 80 Ribokinase | | Ambiguous/Unsorted |
| | 69 Nicotinate-nucleotide pyrophosphorylase [carboxylating] | | Ambiguous/Unsorted |
| | 36 UPF0056 membrane protein yhcE | | Temporarly Up Regulated |
| | 35 Diguanylate cyclase yddV | | Up Regulated |
| | 35 Diguanylate cyclase yddV 35 Diguanylate cyclase yddV | | Up Regulated |
| | | | |
| | 21 Alpha-ketoglutarate-dependent taurine dioxygenase | | Down Regulated |
| | 22 DNA-damage-inducible protein I | | Down Regulated |
| | 23 Sulfurtransferase tusE | | Ambiguous/Unsorted |
| | 99 DNA mismatch repair protein mutH | | Up Regulated |
| | 72 Uncharacterized protein yieL | [Carbohydrate transport and metabolism'] | Temporarly Up Regulated |
| | 98 Uncharacterized protein yahF | | Temporarly Up Regulated |
| | Thymidine kinase | | Ambiguous/Unsorted |
| | 78 Branched-chain-amino-acid aminotransferase | | Ambiguous/Unsorted |
| | 96 SsrA-binding protein | | Down Regulated |
| ilvM 9482 | 79 Acetolactate synthase isozyme 2 small subunit | | Down Regulated |
| | N-acetylmuramoyl-L-alanine amidase amiC | | Down Regulated |
| tauB 94502 | 27 Taurine import ATP-binding protein tauB | | Ambiguous/Unsorted |
| recC 94729 | 94 Exodeoxyribonuclease V gamma chain | | Down Regulated |
| hns 94582 | 29 DNA-binding protein H-NS | | Up Regulated |
| | 76 Inner membrane protein ylaC | | Down Regulated |
| | 01 Uncharacterized protein ygdL | | Down Regulated |
| | 21 Uncharacterized protein ygdL | | Down Regulated |
| | 28 UPF0225 protein ychJ | | Down Regulated |
| | 77 1,6-anhydro-N-acetylmuramyl-L-alanine amidase ampD | | Temporarly Up Regulated |
| | 74 Aerobic respiration control protein arcA | | Ambiguous/Unsorted |
| | 78 Type I restriction enzyme EcoKI R protein | | Down Regulated |
| | 26 Uncharacterized protein ygeW | [' Central intermediary metabolism'] | Temporarly Down Regula |
| | 72 Type I restriction enzyme EcoKI M protein | Contral intermediary frictabolishing | Down Regulated |
| | | | |
| | 70 Sensor protein; Sensor protein rstB | | Down Regulated |
| | Taurine-binding periplasmic protein | | Down Regulated |
| | 47 ATP synthase subunit b; ATP synthase B chain | | Up Regulated |
| | 33 UPF0178 protein yail | | Temporarly Up Regulated |
| | 18 tRNA uridine 5-carboxymethylaminomethyl modification enzyme | | Down Regulated |
| | S1 Shikimate kinase 2 | | Ambiguous/Unsorted |
| | Protein mioC | | Ambiguous/Unsorted |
| | | 376; ECED1_3259; ECUMN_3135; E2348C_3073; APECO1_3725; ECS88_3075; EC55989_3085; ECDH10B_2975 | Down Regulated |
| 70 | | | Down Regulated |
| | 14 ATP synthase subunit beta | | Up Regulated |
| yaiW 94503 | 38 Uncharacterized protein yaiW | Carbon compound utilization', 'RNA related', 'Type of regulation', 'Genetic unit regulated', 'transcription', 'regulation of transcriptio | Temporarly Up Regulated |
| | Protease 3 | | Ambiguous/Unsorted |
| suhB 94728 | 35 Inositol-1-monophosphatase | | Temporarly Down Regula |
| recB 94728 | B6 Exodeoxyribonuclease V beta chain | | Up Regulated |
| | 37 Exodeoxyribonuclease V alpha chain | | Down Regulated |
| pstB 94824 | Phosphate import ATP-binding protein pstB | | Ambiguous/Unsorted |
| | 23 Oligopeptide transport system permease protein oppB | | Down Regulated |
| | 21 Cardiolipin synthetase | | Up Regulated |
| 3.00 | 1 | | |

| 0.470 | 000 | | A mateix manual transmit and |
|------------|---|---|------------------------------|
| | 89 Amino-acid acetyltransferase | | Ambiguous/Unsorted |
| | 16 Ribosomal RNA small subunit methyltransferase E | | Ambiguous/Unsorted |
| | 15 Heme-regulated cyclic di-GMP, cyclic AMP phosphodiesterase, | | Ambiguous/Unsorted |
| | 15 Heme-regulated cyclic di-GMP, cyclic AMP phosphodiesterase, | | Ambiguous/Unsorted |
| | 14 Uncharacterized HTH-type transcriptional regulator ygiT | [' Carbon compound utilization', 'metabolic process'] | Up Regulated |
| | 14 Uncharacterized HTH-type transcriptional regulator ygiT | | Up Regulated |
| flgD 9458 | 13 Basal-body rod modification protein flgD | | Ambiguous/Unsorted |
| oppF 9458 | 18 Oligopeptide transport ATP-binding protein oppF | | Ambiguous/Unsorted |
| flhC 94728 | 80 Flagellar transcriptional activator flhC | | Down Regulated |
| sgcX 94884 | 40 Putative sgc region protein sgcX | | Up Regulated |
| | 40 Phosphate starvation-inducible protein psiF | | Up Regulated |
| phoA 94504 | 41 Alkaline phosphatase | | Ambiguous/Unsorted |
| | 43 Uncharacterized protein yaiA | ['Amino acid transport and metabolism', 'Building block biosynthesis', 'aromatic amino acid family biosynthetic process'] | Up Regulated |
| | 58 Aspartateammonia ligase | | Ambiguous/Unsorted |
| | 44 Sensor protein; Phosphate regulon sensor protein phoR | | Ambiguous/Unsorted |
| | 45Protein aroM | | Down Regulated |
| | 59 Asparaginyl-tRNA synthetase; Regulatory protein asnC | | Down Regulated |
| | 56 ATPase ravA | | Temporarly Down Regula |
| | 46 Phosphate regulon transcriptional regulatory protein phoB | | Temporarly Up Regulated |
| | 74 Uncharacterized sufE-like protein ygdK | | Ambiguous/Unsorted |
| | 48 UPF0345 protein yaiE | | Down Regulated |
| - | 71 Protein syd | | Down Regulated |
| | 49Nuclease sbcCD subunit D | | Temporarly Up Regulated |
| | | | Ambiguous/Unsorted |
| | Low affinity potassium transport system protein kup | | |
| | 52 ATP synthase subunit a | | Down Regulated |
| gcvA 9472 | 78 Glycine cleavage system transcriptional activator | | Temporarly Up Regulated |
| | 76 Uncharacterized lipoprotein ygdl | Carbon compound utilization', 'RNA related', 'Type of regulation', 'Genetic unit regulated'] | Up Regulated |
| | 49 Uxu operon transcriptional regulator | | Temporarly Up Regulated |
| | 02 Oligopeptide transport ATP-binding protein oppD | | Ambiguous/Unsorted |
| | 04 UPF0260 protein ycgN | | Down Regulated |
| | 57 Malate synthase G | | Up Regulated |
| | 07 Respiratory nitrate reductase 1 delta chain | | Up Regulated |
| | 65 Probable glucarate transporter | | Temporarly Down Regula |
| | Acetolactate synthase isozyme 3 small subunit | | Ambiguous/Unsorted |
| | 20 Low affinity tryptophan permease | | Temporarly Up Regulated |
| | 22tRNA modification GTPase mnmE | | Temporarly Down Regula |
| | 60 23S rRNA (uracil-5-)-methyltransferase rumB | | Ambiguous/Unsorted |
| | 63 Flavoprotein wrbA; Putative flavoprotein wrbA | | Up Regulated |
| | 62 L-serine dehydratase 2 | | Up Regulated |
| | 26 Uncharacterized protein yieE | ['Amino acid transport and metabolism'] | Temporarly Down Regula |
| | 25 Uncharacterized protein yieF | | Up Regulated |
| | 27 HTH-type transcriptional regulator yidZ | | Up Regulated |
| | 29 Uncharacterized protein yieK | ['Carbohydrate transport and metabolism'] | Temporarly Up Regulated |
| | 56 Uncharacterized exonuclease xni | | Temporarly Up Regulated |
| | 55 Sensor protein; Signal transduction histidine-protein kinase bar | A | Ambiguous/Unsorted |
| | 54 Protein mazG | | Down Regulated |
| | 53 Aminopeptidase N | | Ambiguous/Unsorted |
| | 52 PemK-like protein 1 | | Temporarly Up Regulated |
| nhoA 9472 | 51 N-hydroxyarylamine O-acetyltransferase | | Ambiguous/Unsorted |
| | 346-phospho-beta-glucosidase bglB | | Ambiguous/Unsorted |
| | 33 Phosphate transport system protein phoU | | Ambiguous/Unsorted |
| yieH 94823 | 32 Phosphatase yieH | | Down Regulated |
| pstS 94823 | 37 Phosphate-binding protein pstS | | Up Regulated |
| gudD 9472 | 58 Glucarate dehydratase | | Ambiguous/Unsorted |
| qmcA 9472 | 57 Protein qmcA | | Temporarly Up Regulated |
| | 00 Small heat shock protein ibpA | | Down Regulated |
| | 41 Cytochrome b561 | | Down Regulated |
| | 40 Putative electron transfer flavoprotein subunit ygcR | | Down Regulated |
| | 43 23S rRNA (uracil-5-)-methyltransferase rumA | | Temporarly Down Regula |
| | 45 Peml-like protein 1 | | Ambiguous/Unsorted |
| | 44GTP pyrophosphokinase | | Ambiguous/Unsorted |
| | 46 Carboxylate-amine ligase ybdK | | Up Regulated |
| | 49 Uncharacterized protein ygiC | in | Down Regulated |
| | 09 DNA replication and repair protein recF | <u></u> | Temporarly Up Regulated |
| | 03 Uncharacterized lipoprotein ydcL | ['Amino acid transport and metabolism', 'amino acid transport'] | Up Regulated |
| V | | F | 1 1 23 2 2 2 2 |

| yjiA 948882 Uncharacterized GTP-binding protein yjiA | In | Ambiguous/Unsorted |
|---|--|--|
| opgB 94888 Phosphoglycerol transferase I | | Down Regulated |
| yjjA 94887 Uncharacterized protein yjjA | n | Down Regulated |
| cusR 945003 Transcriptional regulatory protein cusR | | Temporarly Down Regula |
| yidB 948212 Uncharacterized protein yidB | n | Up Regulated |
| pal 945004 Peptidoglycan-associated lipoprotein | | Up Regulated |
| | n | |
| | U | Temporarly Up Regulated Ambiguous/Unsorted |
| | | |
| | | Down Regulated |
| hisQ 947235 Histidine transport system permease protein hisQ | | Down Regulated |
| dnaN 948218 DNA polymerase III subunit beta | | Down Regulated |
| dnaA 948217 Chromosomal replication initiator protein dnaA | | Up Regulated |
| rpmH 948216 50S ribosomal protein L34 | | Up Regulated |
| rnpA 948215 Ribonuclease P protein component | | Temporarly Down Regula |
| cynR 945001 HTH-type transcriptional regulator cynR | | Up Regulated |
| cysJ 947239 Sulfite reductase [NADPH] flavoprotein alpha-component | | Down Regulated |
| osmY 948895 Osmotically-inducible protein Y | | Up Regulated |
| riml 948894 Ribosomal-protein-alanine acetyltransferase | | Ambiguous/Unsorted |
| fhuF 948891 Ferric iron reductase protein fhuF | | Temporarly Up Regulated |
| holD 948890 DNA polymerase III subunit psi | | Temporarly Up Regulated |
| yjjG 9488995'-nucleotidase yjjG | | Down Regulated |
| mrr 948898 Mrr restriction system protein | | Ambiguous/Unsorted |
| prfC 948897 Peptide chain release factor 3 | | Temporarly Down Regula |
| lpp 946175 Major outer membrane lipoprotein | | Up Regulated |
| entB 946178 Isochorismatase | | Up Regulated |
| ydhR 946177 Putative monooxygenase ydhR | | Down Regulated |
| ynhG 946174 Uncharacterized protein ynhG | П | Ambiguous/Unsorted |
| slyA 946167 Transcriptional regulator slyA | | Temporarly Down Regula |
| plsX 946165 Fatty acid/phospholipid synthesis protein plsX | | Ambiguous/Unsorted |
| pdxY 946162 Pyridoxamine kinase | | Down Regulated |
| rsxG 946158 Electron transport complex protein rnfG | | Ambiguous/Unsorted |
| rnt 946159 Ribonuclease T | | Ambiguous/Unsorted |
| uidR 946150 HTH-type transcriptional regulator uidR | | Down Regulated |
| cnu 946152 OriC-binding nucleoid-associated protein | | Up Regulated |
| hdhA 9461517-alpha-hydroxysteroid dehydrogenase | | Down Regulated |
| Ihr 946156 Probable ATP-dependent helicase Ihr | | Down Regulated |
| uidA 946149Beta-glucuronidase | | Ambiguous/Unsorted |
| fumC 946147 Fumarate hydratase class II | | Temporarly Down Regula |
| rsxB 946146 Electron transport complex protein rnfB | | Down Regulated |
| | | |
| topB 946141 DNA topoisomerase 3 | | Temporarly Up Regulated |
| ycel 946145 Protein ycel | | Ambiguous/Unsorted |
| pntB 946144 NAD(P) transhydrogenase subunit beta | | Ambiguous/Unsorted |
| fecC 948826 Fe(3+) dicitrate transport system permease protein fecC | | Temporarly Up Regulated |
| fecE 948819 Fe(3+) dicitrate transport ATP-binding protein fecE | | Up Regulated |
| yfcZ 948817 UPF0381 protein yfcZ | | Up Regulated |
| sspA 944744 Stringent starvation protein A | | Ambiguous/Unsorted |
| | y['Amino acid transport and metabolism', ' Carbon compound utilization', 'metabolic process'] | Up Regulated |
| ydil 946190 Esterase ydil | | Up Regulated |
| ybaQ 948800 Uncharacterized HTH-type transcriptional regulator ybaQ | [Energy production and conversion', 'Energy production transport', 'electron transport', 'amino acid metabolic process'] | Temporarly Down Regula |
| yddH 946192 Uncharacterized protein yddH | <u> </u> | Up Regulated |
| sapB 946191 Peptide transport system permease protein sapB | | Temporarly Down Regula |
| ygcE 946193 Uncharacterized sugar kinase ygcE | [' Central intermediary metabolism', 'carbohydrate metabolic process'] | Temporarly Up Regulated |
| trxA 948289 Thioredoxin-1 | | Up Regulated |
| ydiH 946194 Uncharacterized protein ydiH | ['proteolysis'] | Up Regulated |
| fucl 946195 L-fucose isomerase | | Up Regulated |
| ilvA 948287 Threonine dehydratase biosynthetic | | Ambiguous/Unsorted |
| ppiC 948285 Peptidyl-prolyl cis-trans isomerase C | | Ambiguous/Unsorted |
| rstA 946199 Transcriptional regulatory protein rstA | | Temporarly Up Regulated |
| gpp 948291 Guanosine-5'-triphosphate,3'-diphosphate pyrophosphatase | | Down Regulated |
| rep 948292 ATP-dependent DNA helicase rep | | Down Regulated |
| wzyE 948293 Putative ECA polymerase | | Down Regulated |
| ydiL 946181 Uncharacterized protein ydiL | in . | Temporarly Up Regulated |
| ydhZ 946180 Uncharacterized protein ydhZ | in The Control of the | Up Regulated |
| rffH 948299 Glucose-1-phosphate thymidylyltransferase; Glucose-1-phosph | nu nate thymidylytransferase 2 | Temporarly Down Regula |
| sufS 946185 Cysteine desulfurase | | Up Regulated |
| para 5 to toolo yatema desumardas | | OP I TOGUIATEU |

| | 946182 | TorCAD operon transcriptional regulatory protein torR | | Temporarly Up Regulated |
|---|--|--|--|---|
| torR vdiJ | | 9 Uncharacterized protein ydiJ | ['Energy production and conversion'] | Ambiguous/Unsorted |
| rffA | | 6 Lipopolysaccharide biosynthesis protein rffA | [Energy production and conversion] | Down Regulated |
| ybaS | | Glutaminase 1 | | Up Regulated |
| puuA | | 2 Gamma-glutamylputrescine synthetase | | Down Regulated |
| Int | | Apolipoprotein N-acyltransferase | | Ambiguous/Unsorted |
| sapD | | Peptide transport system ATP-binding protein sapD | | Down Regulated |
| yadD | | Uncharacterized protein yadD | [Replication recombination and repair', 'DNA recombination'] | Down Regulated |
| appA | | Periplasmic appA protein | [Technolities and Technolities and Techn | Up Regulated |
| ydiA | | Uncharacterized protein yuaU; Putative phosphotransferase ydi | | Down Regulated |
| ppsR | | Uncharacterized protein yuaU; Putative phosphotransferase ydi | | Down Regulated |
| mdtB | | Multidrug resistance protein mdtB | | Up Regulated |
| mdtC | | Multidrug resistance protein mdtC | | Ambiguous/Unsorted |
| rffE | | UDP-N-acetylglucosamine 2-epimerase | | Ambiguous/Unsorted |
| yegQ | 946609 | Uncharacterized protease yegQ | | Down Regulated |
| ydiB | | Quinate/shikimate dehydrogenase | | Temporarly Up Regulated |
| ftsZ | 944786 | Cell division protein ftsZ | | Down Regulated |
| baeS | 946611 | Sensor protein; Signal transduction histidine-protein kinase baes | S | Down Regulated |
| ybiS | 945441 | 1 Uncharacterized protein ybiS | ['phosphoenolpyruvate dependent sugar phosphotransferase system'] | Ambiguous/Unsorted |
| ybiT | | Uncharacterized ABC transporter ATP-binding protein ybiT | | Ambiguous/Unsorted |
| flhD | | 2 Transcriptional activator flhD | | Down Regulated |
| puuE | | 4-aminobutyrate aminotransferase | | Down Regulated |
| ydcF | | Protein ydcF | | Ambiguous/Unsorted |
| fadK | | 3 Short-chain-fatty-acidCoA ligase | | Ambiguous/Unsorted |
| mltC | | Membrane-bound lytic murein transglycosylase C | | Temporarly Up Regulated |
| ydiS | | Probable electron transfer flavoprotein-quinone oxidoreductase | ydiS | Temporarly Down Regula |
| folA | | Dihydrofolate reductase; Dihydrofolate reductase type 1 | | Ambiguous/Unsorted |
| ydiU | | UPF0061 protein ydiU | | Down Regulated |
| ydiY | | BUncharacterized protein ydiY | IJ. | Up Regulated |
| mur⊟ | | 1 UDP-N-acetylmuramoyl-L-alanyl-D-glutamate2,6-diaminopime | iate rigase | Ambiguous/Unsorted |
| lpxL | | SLipid A biosynthesis lauroyl acyltransferase Non-specific ribonucleoside hydrolase rihC | | Up Regulated Up Regulated |
| rihC leuB | | 33-isopropylmalate dehydrogenase | | Temporarly Down Regula |
| aroD | | 03-dehydroquinate dehydratase | | Down Regulated |
| aceK | | | | Down Regulated |
| | | /llsocitrate.dehydrogenase.kinase/nhosnhatase | | Down Regulated |
| lftsI | | 7 Isocitrate dehydrogenase kinase/phosphatase | | Down Regulated |
| ftsI veaE | 944799 | Peptidoglycan synthetase ftsl | [Signal transduction mechanisms', 'Type of regulation', 'two component signal transduction system phosphorelay', 'regulation of tr | Down Regulated |
| yegE | 944799 946600 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE | [Signal transduction mechanisms', 'Type of regulation', 'two component signal transduction system phosphorelay', 'regulation of tr | Down Regulated aTemporarly Up Regulated |
| yegE mdtA | 944799 946600 946604 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA | | Down Regulated aTemporarly Up Regulated Up Regulated |
| yegE mdtA ybil | 944799 946600 946604 945434 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil | ['Signal transduction mechanisms', 'Type of regulation', 'two component signal transduction system phosphorelay', 'regulation of tr [' Adaptation to stress'] | Down Regulated aTemporarly Up Regulated Up Regulated Up Regulated |
| yegE mdtA ybil ybiJ | 944799 946600 946604 945434 945433 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybiJ | | Down Regulated aTemporarly Up Regulated Up Regulated |
| yegE mdtA ybil | 944799 946600 946604 945434 945433 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybiJ | | Down Regulated aTemporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated |
| yegE mdtA ybiI ybiJ pps | 944799 946600 946604 945434 945433 946209 946209 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil | | Down Regulated aTemporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted |
| yegE mdtA ybiI ybiJ pps ppsA | 944799 946600 946604 945434 945433 946209 946209 945431 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil 3 Uncharacterized protein ybiJ 3 Phosphoenolpyruvate synthase 9 Phosphoenolpyruvate synthase | | Down Regulated aTemporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted |
| yegE mdtA ybil ybiJ pps ppsA dinG | 944799 946600 946604 945434 945433 946209 946209 945431 945435 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ | | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Down Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB | 944799 946600 946604 945434 945433 946209 946209 945431 945435 944762 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase | | Down Regulated aTemporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog | 944799 946600 946604 945434 945433 946209 946209 945431 945437 945435 944762 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein | | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT | 944799 946600 946604 945434 945433 946209 945431 945437 945435 944762 944762 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Deptide protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 | | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR | 944798 946600 946604 945434 945433 946208 945431 945437 945436 944760 944758 944759 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein mdtA Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR | | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE | 944798 946600 946604 945434 945433 946209 945431 945437 945438 944766 944759 944757 945428 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhIE | [Adaptation to stress'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB | 944799 946600 946604 945434 945436 946209 946209 945431 945436 944762 944760 944759 944759 944759 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG 7Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 7Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB | [Adaptation to stress'] [] [] [[Energy metabolism carbon ', 'nitrate assimilation'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB | 944799 946604 945434 945433 946209 94523 945431 945433 945436 944762 944762 944759 944752 945422 945422 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 330S ribosomal protein S20 Transcriptional activator protein nhaR Sputative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH | [Adaptation to stress'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB | 944799 946604 945434 945433 946209 945433 945437 945435 944760 944759 944759 945422 945424 945424 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH Biopolymer transport protein exbB | [Adaptation to stress'] [Energy metabolism carbon ', 'nitrate assimilation'] [Defense mechanisms', ' Primary Active Transporters'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB ybiH exbB apaH | 944799 946604 945434 945433 946209 945437 945435 945435 944766 944759 944759 945422 945424 945424 945424 945424 945424 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH Disiopolymer transport protein exbB Bis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl) | [Adaptation to stress'] [Energy metabolism carbon ', 'nitrate assimilation'] [Defense mechanisms', ' Primary Active Transporters'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Temporarly Up Regulated Ambiguous/Unsorted |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB ybiB ybiB ybiB | 944799 946600 946604 945433 945433 946209 945431 945435 944762 944762 944759 944759 944759 944759 945424 945421 945427 945427 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhIE Uncharacterized HTH-type transcriptional regulator ybiH Dispolymer transport protein exbB Disis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'- | [Adaptation to stress'] [Energy metabolism carbon ', 'nitrate assimilation'] [Defense mechanisms', ' Primary Active Transporters'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted |
| yegE mdtA ybil ybiJ pps ppsA dinG mntB glnQ dapB Mog rpsT nhaR rhIE ybiB ybiH exbB apaH ftsA ispH | 944799 946604 946604 945434 945433 946209 945431 945437 944766 944759 944759 945422 945422 945427 945427 945427 945427 945427 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhlE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH Diopolymer transport protein exbB Disi(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleoss) Cell division protein ftsA 74-hydroxy-3-methylbut-2-enyl diphosphate reductase | [Adaptation to stress'] [Energy metabolism carbon ', 'nitrate assimilation'] [Defense mechanisms', ' Primary Active Transporters'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhlE ybiB ybiH exbB apaH ftsA ispH apaG | 944799 946604 945434 945433 946209 94523 945437 945436 944762 944775 944775 945422 945421 945421 945427 945427 945427 945427 945427 945427 945427 945427 945427 945427 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhlE Uncharacterized protein ybilB Uncharacterized HTH-type transcriptional regulator ybiH DBiopolymer transport protein exbB DBis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl-archydroxy-3-methylbut-2-enyl diphosphate reductase | [Adaptation to stress'] [Energy metabolism carbon ', 'nitrate assimilation'] [Defense mechanisms', ' Primary Active Transporters'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB ybiH exbB apaH ftsA ispH apaG sspB | 944799 946604 945434 945433 946209 945437 945436 945436 944760 944760 944759 945422 945422 945421 945427 945427 945427 945427 945427 945427 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH Disiopolymer transport protein exbB Disis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl Protein apaG Stringent starvation protein B | [Adaptation to stress'] [Energy metabolism carbon ', 'nitrate assimilation'] [Defense mechanisms', ' Primary Active Transporters'] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated |
| yegE mdtA ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB exbB apaH ftsA ispH apaG sspB rpmD | 944799 946604 945434 945433 946209 945433 945436 945436 944760 944759 944759 945422 945422 945424 945424 945427 945427 945427 945427 945427 945427 945427 945427 945427 945427 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH Disipolymer transport protein exbB Disis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl) Religious protein ftsA 74-hydroxy-3-methylbut-2-enyl diphosphate reductase Protein apaG Stringent starvation protein B 750S ribosomal protein L30 | [' Adaptation to stress'] [] [] [] [] [] [] [] [] [] [] [] [] [] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Down Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB ybiH exbB apaH ftsA ispH apaG sspB rpmD secY | 944799 946600 946604 946604 945434 945434 945434 945435 944776 944776 944776 944777 944777 944777 944777 944777 947779 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhlE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH Disipolymer transport protein exbB Disis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl)-tetrapho | [' Adaptation to stress'] [] [] [] [] [] [] [] [] [] [] [] [] [] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Temporarly Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Down Regulated Down Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB ybiH exbB apaH ftsA ispH apaG sspB rpmD secty rplO | 944799 946600 946604 945434 945433 946209 945437 945437 944766 944766 944769 9447767 945422 945422 945422 945427 945427 947779 944777 944777 947779 947799 947798 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhlE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH Biopolymer transport protein exbB Disiof'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl)-Cled division protein ftsA 74-hydroxy-3-methylbut-2-enyl diphosphate reductase Protein apaG Stringent starvation protein B Foos ribosomal protein L30 Preprotein translocase subunit secY; Preprotein translocase sec | [' Adaptation to stress'] [] [] [] [] [] [] [] [] [] [] [] [] [] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB ybiH exbB apaH ftsA ispH apaG sspB rpmD secY rpsD | 944799 946604 945434 945433 946209 945437 945437 945436 944775 944775 944777 944777 944777 947799 947799 947799 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybiJ Uncharacterized protein ybiJ Uncharacterized protein ybiJ Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Sputative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH DBiopolymer transport protein exbB DBis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl-retrophosphatase) Protein apaG 4 Stringent starvation protein B 750S ribosomal protein L15 30S ribosomal protein L15 | [' Adaptation to stress'] [] [] [] [] [] [] [] [] [] [] [] [] [] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Up Regulated Down Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB apaH ftsA ispH apaG sspB rpmD secY rplO rpsK | 944799 946604 945434 945433 946209 945433 945433 945436 944760 944755 944776 945422 945422 945422 945422 945422 945422 945427 94778 94779 94779 94779 94779 94779 94779 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Uncharacterized protein ybil Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Putative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH Disiopolymer transport protein exbB Disis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl)-tetraphosphatase, symmetrical; Pisic(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl)-tetraphosphatase, symmetr | [' Adaptation to stress'] [] [] [] [] [] [] [] [] [] [] [] [] [] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Up Regulated Down Regulated |
| yegE mdtA ybil ybiJ pps ppsA dinG mntR glnQ dapB Mog rpsT nhaR rhIE ybiB apaH ftsA ispH apaG sspB rpmD secY rpIO rpsC | 944799 946604 945434 945433 946209 945437 945436 944760 944776 944776 944777 944777 94779 94779 94779 94779 94779 94779 94779 | Peptidoglycan synthetase ftsl Uncharacterized protein yegE Multidrug resistance protein mdtA Uncharacterized protein ybiJ Uncharacterized protein ybiJ Uncharacterized protein ybiJ Phosphoenolpyruvate synthase Phosphoenolpyruvate synthase Probable ATP-dependent helicase dinG Transcriptional regulator mntR Glutamine transport ATP-binding protein glnQ Dihydrodipicolinate reductase Molybdopterin biosynthesis mog protein 30S ribosomal protein S20 Transcriptional activator protein nhaR Sputative ATP-dependent RNA helicase rhIE Uncharacterized protein ybiB Uncharacterized HTH-type transcriptional regulator ybiH DBiopolymer transport protein exbB DBis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl)-tetraphosphatase, symmetrical; Bis(5'-nucleosyl-retrophosphatase) Protein apaG 4 Stringent starvation protein B 750S ribosomal protein L15 30S ribosomal protein L15 | [' Adaptation to stress'] [] [] [] [] [] [] [] [] [] [] [] [] [] | Down Regulated a Temporarly Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Up Regulated Down Regulated |

| ybhF | 945413 | Uncharacterized ABC transporter ATP-binding protein ybhF | | Temporarly Up Regulated |
|--------------|--------|---|--|---------------------------------|
| rpsM | | 30S ribosomal protein S13; 30S ribosomal protein | | Down Regulated |
| vehl | | Uncharacterized protein yehl | "Translation ribosomal structure and biogenesis', 'translation', 'ion transport'] | Up Regulated |
| metC | | Cystathionine beta-lyase metC | Translation indescribe and dispersess, translation, for transport | Ambiguous/Unsorted |
| ybdG | | Uncharacterized protein ybdG | n | Ambiguous/Unsorted |
| fiu | | Catecholate siderophore receptor fiu | U | Up Regulated |
| rlmN | | Ribosomal RNA large subunit methyltransferase N | | Down Regulated |
| rimN | | Putative ribosome maturation factor rimN | | Up Regulated |
| SDH | | Succinate dehydrogenase flavoprotein subunit | | Ambiguous/Unsorted |
| ybhR | | Inner membrane transport permease ybhR | | Temporarly Up Regulated |
| rplQ | | 50S ribosomal protein L17 | | Temporarly Down Regula |
| yrdD | | Uncharacterized protein yrdD | ['Replication recombination and repair', 'amino acid biosynthetic process', 'aromatic amino acid family biosynthetic process'] | Up Regulated |
| dld | | D-lactate dehydrogenase | , | Ambiguous/Unsorted |
| mscL | | Large-conductance mechanosensitive channel | | Temporarly Up Regulated |
| trkA | | Trk system potassium uptake protein trkA | | Up Regulated |
| yhdN | | Uncharacterized protein yhdN | П | Down Regulated |
| zntR | 947786 | HTH-type transcriptional regulator zntR | | Down Regulated |
| yeiE | | Uncharacterized HTH-type transcriptional regulator yeiE | ['Transcription', ' RNA related', ' Type of regulation', 'transcription', 'regulation of transcription DNA dependent'] | Down Regulated |
| yegW | 946639 | Uncharacterized HTH-type transcriptional regulator yegW | | Temporarly Up Regulated |
| astC | 946255 | Succinylornithine transaminase | | Up Regulated |
| spy | | Spheroplast protein Y | | Temporarly Up Regulated |
| astB | | N-succinylarginine dihydrolase | | Up Regulated |
| lptE | | LPS-assembly lipoprotein lptE | | Ambiguous/Unsorted |
| yrdB | | Uncharacterized protein yrdB | [' Building block biosynthesis', 'purine nucleotide biosynthetic process', 'aromatic amino acid family biosynthetic process'] | Temporarly Down Regula |
| yrdA | | Protein yrdA | | Ambiguous/Unsorted |
| aroE | 947776 | Shikimate dehydrogenase; Quinate/shikimate dehydrogenase | | Up Regulated |
| pncB | | Nicotinate phosphoribosyltransferase | | Down Regulated |
| rihB | | Pyrimidine-specific ribonucleoside hydrolase rihB | | Temporarly Up Regulated |
| mrp | | Protein mrp | | Down Regulated |
| pntA | | NAD(P) transhydrogenase subunit alpha | | Up Regulated |
| helD | | Helicase IV | | Down Regulated |
| pheS | | Phenylalanyl-tRNA synthetase alpha chain | | Ambiguous/Unsorted |
| ycgL | | UPF0745 protein ycgL | | Ambiguous/Unsorted |
| aaeR | | HTH-type transcriptional activator aaeR | | Up Regulated |
| acrF | | Acriflavine resistance protein F Putative amino-acid ABC transporter-binding protein yhdW | | Temporarly Up Regulated |
| yhdW | | Uncharacterized protein yegU | n | Up Regulated Ambiguous/Unsorted |
| yegU yhdZ | | Uncharacterized protein yego Uncharacterized amino-acid ABC transporter ATP-binding prote | IU sin yhd7 | Down Regulated |
| yhdY | | Inner membrane amino-acid ABC transporter permease protein | | Up Regulated |
| fbaB | | Fructose-bisphosphate aldolase class 1 | ynu | Ambiguous/Unsorted |
| pfkB | | 6-phosphofructokinase isozyme 2 | | Ambiguous/Unsorted |
| chbG | | UPF0249 protein chbG | | Temporarly Up Regulated |
| tldD | | Protein tldD | | Down Regulated |
| ydjN | | Uncharacterized symporter ydjN | n | Down Regulated |
| yniA | | Uncharacterized protein yniA | <u> </u> | Up Regulated |
| katE | | Catalase HPII; Catalase | <u> </u> | Ambiguous/Unsorted |
| cedA | | Cell division activator cedA | | Down Regulated |
| yhcH | | Uncharacterized protein yhcH | ['Carbohydrate transport and metabolism', 'carbohydrate metabolic process'] | Up Regulated |
| tolA | | Protein tolA | | Down Regulated |
| accB | 947758 | Biotin carboxyl carrier protein of acetyl-CoA carboxylase | | Ambiguous/Unsorted |
| hpt | | Hypoxanthine phosphoribosyltransferase | | Ambiguous/Unsorted |
| cyaY | 947754 | Protein cyaY | | Up Regulated |
| cyaA | 947755 | Adenylate cyclase | | Down Regulated |
| nanA | 947742 | N-acetylneuraminate lyase 1; N-acetylneuraminate lyase | | Up Regulated |
| | 946697 | Cytochrome c-type biogenesis protein ccmE | | Temporarly Up Regulated |
| | | p-hydroxybenzoic acid efflux pump subunit aaeA | | Ambiguous/Unsorted |
| yejK | | Nucleoid-associated protein ndpA | | Ambiguous/Unsorted |
| rlml | | Ribosomal RNA large subunit methyltransferase I | | Ambiguous/Unsorted |
| yejL | | UPF0352 protein yejL | | Temporarly Up Regulated |
| rtn | | Protein rtn | | Up Regulated |
| lpxT | | Inner membrane protein yeiU | | Down Regulated |
| recQ | 948318 | ATP-dependent DNA helicase recQ | | Down Regulated |
| | | Succinyl-diaminopimelate desuccinylase | | Ambiguous/Unsorted |
| | | Lysophospholipase L2 HTH-type transcriptional regulator metR | | Down Regulated |
| metR | 948310 | ı п ı п-ıype transcriptionai regulator metк | I . | Down Regulated |

| hom | 049211 | Drotoin homV | | Ambiguous/Uncorted |
|--------------|--------|--|--|---|
| hem\ ushA | | Protein hemY Protein ushA | | Ambiguous/Unsorted Ambiguous/Unsorted |
| yejM | | Inner membrane protein yejM | | Down Regulated |
| mlaE | | UPF0393 inner membrane protein yrbE | | Temporarly Down Regula |
| yrbE | | UPF0393 inner membrane protein yrbE | | Temporarly Down Regula |
| flgC | | Flagellar basal-body rod protein flgC | | Down Regulated |
| yejF | 946689 | Uncharacterized ABC transporter ATP-binding protein yeiF | | Ambiguous/Unsorted |
| cueO | | Blue copper oxidase cueO | | Temporarly Up Regulated |
| kdsD | | Arabinose 5-phosphate isomerase | | Ambiguous/Unsorted |
| osmF | | Putative osmoprotectant uptake system substrate-binding protei | | Up Regulated |
| flgM | | Negative regulator of flagellin synthesis | | Ambiguous/Unsorted |
| pldA | 948307 | Phospholipase A1 | | Up Regulated |
| rffG | | dTDP-glucose 4,6-dehydratase | | Down Regulated |
| ydhP | | Inner membrane transport protein ydhP | | Temporarly Down Regula |
| rffM | | Probable UDP-N-acetyl-D-mannosaminuronic acid transferase | | Temporarly Up Regulated |
| gltB | 947724 | Glutamate synthase [NADPH] large chain | | Ambiguous/Unsorted |
| gltD | | Glutamate synthase [NADPH] small chain | | Down Regulated |
| ampE | 946678 | Protein ampE | | Temporarly Up Regulated |
| speE | | Spermidine synthase | | Ambiguous/Unsorted |
| secG | | Protein-export membrane protein secG | | Up Regulated |
| fruK | | 1-phosphofructokinase | | Down Regulated |
| fruB | | Multiphosphoryl transfer protein | | Ambiguous/Unsorted |
| yeiP | | Elongation factor P-like protein | | Ambiguous/Unsorted |
| yejA | | Uncharacterized protein yejA | | Down Regulated |
| fruA | | PTS system fructose-specific EIIBC component | | Down Regulated |
| fixA | | Protein fixA | | Ambiguous/Unsorted |
| pcnB | | Poly(A) polymerase | | Down Regulated |
| lysA | | Diaminopimelate decarboxylase | | Ambiguous/Unsorted |
| galR | | HTH-type transcriptional regulator galR | | Temporarly Down Regula |
| aas | | Bifunctional protein aas | | Temporarly Up Regulated Down Regulated |
| hpf | 045903 | Probable sigma(54) modulation protein Phage shock protein B | | Temporarly Up Regulated |
| pspB speD | | S-adenosylmethionine decarboxylase proenzyme | | Ambiguous/Unsorted |
| speB | | Agmatinase | | Down Regulated |
| lysP | | Lysine-specific permease | | Down Regulated |
| amiD | | N-acetylmuramoyl-L-alanine amidase amiD | | Temporarly Up Regulated |
| rpoN | | RNA polymerase sigma-54 factor | | Down Regulated |
| ybjQ | | UPF0145 protein ybjQ | | Temporarly Down Regula |
| ybjT | | Uncharacterized protein ybjT | [Cell wall membrane envelope biogenesis] | Temporarly Down Regula |
| nfo | | Endonuclease 4; Probable endonuclease 4 | | Down Regulated |
| ybjS | | Uncharacterized protein ybjS | n | Ambiguous/Unsorted |
| aat | | Leucyl/phenylalanyl-tRNAprotein transferase | | Down Regulated |
| cdd | | Cytidine deaminase | | Up Regulated |
| ybjP | 945491 | Uncharacterized lipoprotein ybjP | ['Amino acid transport and metabolism', ' Building block biosynthesis'] | Up Regulated |
| pbpG | 946662 | D-alanyl-D-alanine endopeptidase | | Down Regulated |
| ybjD | | Uncharacterized protein ybjD | ['SOS response'] | Up Regulated |
| pspC | | Phage shock protein C | | Up Regulated |
| kduD | 947323 | 2-deoxy-D-gluconate 3-dehydrogenase | | Down Regulated |
| yqeF | 947324 | Probable acetyl-CoA acetyltransferase | | Temporarly Up Regulated |
| ymd(| | Uncharacterized protein ymdC | <u>[0 </u> | Temporarly Up Regulated |
| macA | | Macrolide-specific efflux protein macA | | Ambiguous/Unsorted |
| iraP | | Anti-adapter protein iraP | | Temporarly Up Regulated |
| ygeV | 947320 | Uncharacterized sigma-54-dependent transcriptional regulator y | [Signal transduction mechanisms', 'Genetic unit regulated', 'two component signal transduction system phosphorelay ', 'regulation of | |
| dusB | | tRNA-dihydrouridine synthase B | | Temporarly Up Regulated |
| prmA | | Ribosomal protein L11 methyltransferase | | Ambiguous/Unsorted |
| artP | | Arginine transport ATP-binding protein artP | | Temporarly Down Regula Ambiguous/Unsorted |
| potF | | Putrescine-binding periplasmic protein Protein gmr | | Temporarly Up Regulated |
| gmr yciT | | Uncharacterized HTH-type transcriptional regulator yciT | | Ambiguous/Unsorted |
| rimK | | Ribosomal protein S6 modification protein | | Ambiguous/Unsorted |
| csrD | | Uncharacterized protein yhdA | | Temporarly Up Regulated |
| ybjN | | Uncharacterized protein yhdx Uncharacterized protein ybjN | [' Energy metabolism carbon', 'electron transport', 'translation'] | Down Regulated |
| murA | | UDP-N-acetylglucosamine 1-carboxyvinyltransferase | t many many many many many many many many | Ambiguous/Unsorted |
| nfsA | 945483 | Oxygen-insensitive NADPH nitroreductase | | Ambiguous/Unsorted |
| | | Peptide transport periplasmic protein sapA | | Up Regulated |
| | | | | |

| ycjF | 945878 | UPF0283 membrane protein ycjF | | Temporarly Down Regula |
|------|--------|--|--|-------------------------|
| prr | | Gamma-aminobutyraldehyde dehydrogenase | | Up Regulated |
| ycjW | | Uncharacterized HTH-type transcriptional regulator ycjW | "Transcription', 'RNA related', 'Type of regulation', 'Genetic unit regulated', 'transcription', 'regulation of transcription DNA depend | |
| potG | | Putrescine transport ATP-binding protein potG | [Transcription, Traversacion, Type of regulation, Content and regulation, regulation of automption Draved point | Ambiguous/Unsorted |
| grxA | | Glutaredoxin-1 | | Ambiguous/Unsorted |
| tyrR | | Transcriptional regulatory protein tyrR | | Ambiguous/Unsorted |
| ybjl | | Phosphatase ybil | | Down Regulated |
| ihfA | | Integration host factor subunit alpha | | Ambiguous/Unsorted |
| ybjL | | Putative transport protein ybjL | | Ambiguous/Unsorted |
| puuD | | Gamma-glutamyl-gamma-aminobutyrate hydrolase | | Ambiguous/Unsorted |
| rppH | | RNA pyrophosphohydrolase | | Temporarly Up Regulated |
| pspA | | Phage shock protein A | | Down Regulated |
| puuR | | HTH-type transcriptional regulator puuR | | Up Regulated |
| omp(| | Outer membrane protein G | | Ambiguous/Unsorted |
| ygdR | | Uncharacterized lipoprotein ygdR | ['Amino acid transport and metabolism', ' Building block biosynthesis', 'amino acid metabolic process'] | Temporarly Up Regulated |
| lgt | | Prolipoprotein diacylglyceryl transferase | | Ambiguous/Unsorted |
| yfhM | 947302 | Uncharacterized lipoprotein yfhM | П | Up Regulated |
| rhmD | 945881 | L-rhamnonate dehydratase | | Ambiguous/Unsorted |
| yfaW | | L-rhamnonate dehydratase | | Ambiguous/Unsorted |
| tas | 947306 | Protein tas | | Temporarly Up Regulated |
| ascG | 947305 | HTH-type transcriptional regulator ascG | | Ambiguous/Unsorted |
| acrD | 945464 | Probable aminoglycoside efflux pump | | Ambiguous/Unsorted |
| rimO | 945465 | Ribosomal protein S12 methylthiotransferase rimO | | Down Regulated |
| yliE | 945462 | Uncharacterized protein yliE | ['Signal transduction mechanisms', 'Type of regulation', 'two component signal transduction system phosphorelay', 'transcription', 'I | Up Regulated |
| rssB | 945855 | Protein hnr | | Temporarly Down Regula |
| yciO | 945854 | Uncharacterized protein yciO | | Ambiguous/Unsorted |
| Ogt | 945853 | Methylated-DNAprotein-cysteine methyltransferase | | Down Regulated |
| add | | Adenosine deaminase | | Ambiguous/Unsorted |
| gsiA | 945457 | Glutathione import ATP-binding protein gsiA | | Down Regulated |
| gsiB | 945459 | Glutathione-binding protein gsiB | | Ambiguous/Unsorted |
| deoR | | Deoxyribose operon repressor | | Down Regulated |
| moe/ | | Molybdopterin biosynthesis protein moeA | | Ambiguous/Unsorted |
| dacC | 945455 | D-alanyl-D-alanine carboxypeptidase dacC | | Ambiguous/Unsorted |
| folC | | Bifunctional protein folC | | Ambiguous/Unsorted |
| moeE | | Molybdopterin biosynthesis protein moeB | | Temporarly Down Regula |
| yciV | | Protein trpH | | Down Regulated |
| yciW | | Uncharacterized protein yciW | ['Lipid transport and metabolism'] | Down Regulated |
| tktB | | Transketolase 2 | | Up Regulated |
| rnb | | Exoribonuclease 2 | | Temporarly Down Regula |
| osmE | | Osmotically-inducible lipoprotein B | | Up Regulated |
| ydfl | | Uncharacterized oxidoreductase ydfl | [' Carbon compound utilization'] | Up Regulated |
| yciN | | Protein yciN | | Up Regulated |
| dapF | | Diaminopimelate epimerase | | Down Regulated |
| ybcJ | | Uncharacterized protein ybcJ | <u>D</u> | Ambiguous/Unsorted |
| yihN | | Inner membrane protein yihN | | Ambiguous/Unsorted |
| glnL | 948360 | Nitrogen regulation protein NR(II); Sensor protein | | Down Regulated |
| proX | | Glycine betaine-binding periplasmic protein | | Temporarly Up Regulated |
| glnG | | Nitrogen regulation protein NR(I) | | Temporarly Up Regulated |
| emrA | | Multidrug resistance protein A | | Down Regulated |
| hemi | | Oxygen-independent coproporphyrinogen-III oxidase | | Down Regulated |
| yihl | | UPF0241 protein yihl | | Ambiguous/Unsorted |
| luxS | | S-ribosylhomocysteine lyase | | Ambiguous/Unsorted |
| allC | 945150 | Allantoate amidohydrolase | | Down Regulated |
| rfbA | 945154 | Glucose-1-phosphate thymidylyltransferase; Glucose-1-phosphate | ate tnymicylytransterase 1 | Down Regulated |
| yihL | | Uncharacterized HTH-type transcriptional regulator yihL | ['Amino acid transport and metabolism', ' Carbon compound utilization', ' Central intermediary metabolism', ' Primary Active Transpo | |
| typA | | GTP-binding protein typA/BipA; GTP-binding protein typA/bipA | | Ambiguous/Unsorted |
| rpIT | | 50S ribosomal protein L20 | | Temporarly Up Regulated |
| ytfE | | Regulator of cell morphogenesis and NO signaling | | Ambiguous/Unsorted |
| malY | | Protein malY | | Up Regulated |
| Cyca | 948725 | D-serine/D-alanine/glycine transporter | Positioning appropriation and apprint UNIA propositional UNIA interesting | Down Regulated |
| ytfB | | Uncharacterized protein ytfB | ['Replication recombination and repair', 'DNA recombination', 'DNA integration'] | Up Regulated |
| rpll | | 50S ribosomal protein L9 | | Ambiguous/Unsorted |
| rpsR | | 30S ribosomal protein S18 | | Down Regulated |
| | | Mhp operon transcriptional activator | | Temporarly Up Regulated |
| Papa | 94/152 | Penicillin-binding protein 1C | I . | Temporarly Down Regula |

| المقام | 040070D toward tDNA/Town describes | | Davis Dagulated |
|--|--|---|--|
| dtd | 948378 D-tyrosyl-tRNA(Tyr) deacylase | | Down Regulated |
| dbpA | 947153 ATP-independent RNA helicase dbpA | | Down Regulated |
| yqcA | 947151 Uncharacterized protein yqcA | | Ambiguous/Unsorted |
| ygaM | 947157 Uncharacterized protein ygaM | | Up Regulated |
| prpD | 945931 2-methylcitrate dehydratase | | Ambiguous/Unsorted |
| cpdB | | 4750; Z5824; EcoIC_3793; ECS88_4807; E2348C_4544; EC55989_4773; ECDH10B_4408; ECED1_5072; ECH74115_5730; ECIAI | |
| yihP | 948371 Inner membrane symporter yihP | | Down Regulated |
| nrdE | 947155 Ribonucleoside-diphosphate reductase; Ribonucleoside-diphos | | Temporarly Up Regulated |
| cysQ | 948728 3'(2'),5'-bisphosphate nucleotidase cysQ | | Up Regulated |
| sfmD | 945160 Outer membrane usher protein sfmD | | Ambiguous/Unsorted |
| yegl | 947159 Uncharacterized protein yegl | | Ambiguous/Unsorted |
| macE | 945164 Macrolide export ATP-binding/permease protein macB | | Up Regulated |
| glnA | 948370 Glutamine synthetase | | Temporarly Down Regula |
| ytfJ | 948737 Uncharacterized protein ytfJ | [Energy production and conversion', 'Secondary metabolites biosynthesis transport and catabolism'] | Temporarly Up Regulated |
| ytfL | 948735 UPF0053 inner membrane protein ytfL | | Temporarly Down Regula |
| ytfM | 948733 Uncharacterized protein ytfM | | Up Regulated |
| msrA | 948734 Peptide methionine sulfoxide reductase msrA | | Ambiguous/Unsorted |
| qorB | 948731 Uncharacterized oxidoreductase ytfG | | Up Regulated |
| fdhE | 948384 Protein fdhE | | Down Regulated |
| ydjA | 945964 Putative NAD(P)H nitroreductase ydjA | | Ambiguous/Unsorted |
| yddB | 945961 Uncharacterized protein yddB | [Cell wall membrane envelope biogenesis', 'translation', 'ion transport'] | Up Regulated |
| MItB | 947184 Membrane-bound lytic murein transglycosylase B | | Down Regulated |
| allB | 945134 Allantoinase | | Down Regulated |
| tus | 945135 DNA replication terminus site-binding protein | | Down Regulated |
| norR | 947186 Anaerobic nitric oxide reductase transcription regulator norR | | Temporarly Down Regula |
| kefA | 945132 Potassium efflux system kefA | | Ambiguous/Unsorted |
| pmrD | 945130 Signal transduction protein pmrD | | Temporarly Up Regulated |
| ytfN | 948742 Uncharacterized protein ytfN | | Down Regulated |
| fhIA | 947181 Formate hydrogenlyase transcriptional activator | | Ambiguous/Unsorted |
| ytfQ | 948746 ABC transporter periplasmic-binding protein ytfQ | | Up Regulated |
| рра | 948748 Inorganic pyrophosphatase | | Ambiguous/Unsorted |
| ydeN | 945957 Uncharacterized sulfatase ydeN | | Up Regulated |
| yihW | 948381 Uncharacterized HTH-type transcriptional regulator yihW | [Transcription', 'RNA related', 'Type of regulation', 'Genetic unit regulated', 'transcription', 'regulation of transcription DNA dependent | Temporarly Down Regula |
| yihX | 948380 Phosphatase yihX | | Up Regulated |
| fdoH | 948395 Formate dehydrogenase-O iron-sulfur subunit | | Down Regulated |
| ylbA | 945149 Uncharacterized protein ylbA | | Ambiguous/Unsorted |
| ydaL | 945953 Uncharacterized protein ydaL | | Up Regulated |
| clcB | 947179 Voltage-gated CIC-type chloride channel clcB | | Temporarly Down Regula |
| csrA | 947176 Carbon storage regulator | | Temporarly Up Regulated |
| fdhD | 948393 Protein fdhD | | Ambiguous/Unsorted |
| fdoG | 948394 Formate dehydrogenase-O major subunit | | Ambiguous/ Onsonicu |
| dgt | | | Down Regulated |
| | 947177 Deoxyguanosinetriphosphate triphosphohydrolase | | Down Regulated |
| qseC | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein | | |
| qseC xdhC | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein | | Down Regulated Down Regulated Down Regulated |
| | 947177 Deoxyguanosinetriphosphate triphosphohydrolase | | Down Regulated Down Regulated |
| xdhC | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit | | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted |
| xdhC mpp# | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein | | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated |
| xdhC mppA ThiM | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase | | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted |
| xdhC mpp# ThiM lpxM yniB | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase | | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted |
| xdhC mppA ThiM lpxM | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB | | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated |
| xdhC mppA ThiM lpxM yniB yjgA nrdD | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Kanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase | | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated |
| xdhC mppA ThiM lpxM yniB yjgA nrdD ydbH | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yigA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH | | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated |
| xdhC mpp/ ThiM lpxM yniB yjgA nrdD ydbH ydhV | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydhV | [] [] [] [Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Down Regulated Temporarly Up Regulated |
| xdhC mppA ThiM lpxM yniB yjgA nrdD ydbH ydhV fbp | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydhV 948753 Fructose-1,6-bisphosphatase class 1 | [] [] Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Down Regulated Temporarly Up Regulated Ambiguous/Unsorted |
| xdhC mppA ThiM lpxM yniB yjgA nrdD ydbH ydhV fbp yddA | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbH 948755 Fructose-1,6-bisphosphatase class 1 945945 Inner membrane ABC transporter ATP-binding protein yddA | [] [] Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Down Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted |
| xdhC mppA ThiM lpxM yniB yjgA nrdD ydbH ydhV fbp yddA uspF | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbH 948753 Fructose-1,6-bisphosphatase class 1 945948 Universal stress protein F | [] [] [' Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Down Regulated Temporarly Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated |
| xdhC mppA ThiM lpxM yniB yjgA nrdD ydbH ydhV fbp yddA uspF fre | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945947 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbV 948753 Fructose-1,6-bisphosphatase class 1 945945 Inner membrane ABC transporter ATP-binding protein yddA 948948 Universal stress protein F 948325 NAD(P)H-flavin reductase | [] [] [Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated |
| xdhC mppA ThiM lpxM yniB yigA nrdD ydbH ydhV fbp yddA uspF fre fadA | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbH 948753 Fructose-1,6-bisphosphatase class 1 945948 Universal stress protein F 948325 NAD(P)H-flavin reductase 948324 3-ketoacyl-CoA thiolase | Central intermediary metabolism" | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated |
| xdhC mppA ThiM lpxM yniB yigA nrdD ydbH ydhV fbp yddA uspF fre fadA raiA | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yigA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbH 948757 Fructose-1,6-bisphosphatase class 1 945948 Universal stress protein F 948325 NAD(P)H-flavin reductase 948324 3-ketoacyl-CoA thiolase 947129 Ribosome-associated inhibitor A | Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated |
| xdhC mppA ThiM lpxM yniB yjgA nrdD ydbH ydhV fbp yddA uspF fre fadA raiA | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbV 948753 Fructose-1,6-bisphosphatase class 1 945945 Inner membrane ABC transporter ATP-binding protein yddA 945948 Universal stress protein F 948325 NAD(P)H-flavin reductase 948324 3-ketoacyl-CoA thiolase 947129 Ribosome-associated inhibitor A 948327 Transcriptional activator rfaH | [] [] Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Down Regulated Temporarly Up Regulated Ambiguous/Unsorted |
| xdhC mppA ThiM lpxM yniB yjgA nrdD ydbH ydhV fbp yddA uspF fre fadA raiA rfaH ubiD | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945947 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yigA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbV 948753 Fructose-1,6-bisphosphatase class 1 945945 Universal stress protein F 948325 NAD(P)H-flavin reductase 948324 3-ketoacyl-CoA thiolase 947129 Ribosome-associated inhibitor A 948326 3-octaprenyl-4-hydroxybenzoate carboxy-lyase | [] [] Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Down Regulated Temporarly Up Regulated Ambiguous/Unsorted Up Regulated Down Regulated |
| xdhC mppA ThiM lpxM yniB yigA nrdD ydbH ydhV fbp yddA uspF fre fadA raiA rfaH ubiD tatA | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945947 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yigA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbH 948756 Fructose-1,6-bisphosphatase class 1 945945 Inner membrane ABC transporter ATP-binding protein yddA 945948 Universal stress protein F 948325 NAD(P)H-flavin reductase 948324 3-ketoacyl-CoA thiolase 947129 Ribosome-associated inhibitor A 948327 Transcriptional activator rfaH 948326 3-octaprenyl-4-hydroxybenzoate carboxy-lyase 948321 Sec-independent protein translocase protein tatA | [] [] [] Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated |
| xdhC mppA ThiM lpxM yniB yigA nrdD ydbH ydhV fbp yddA uspF fre fadA raiA rfaH ubiD tatA ysgA | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945947 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbH 948757 Fructose-1,6-bisphosphatase class 1 945945 Inner membrane ABC transporter ATP-binding protein yddA 945948 Universal stress protein F 948325 NAD(P)H-flavin reductase 948324 3-ketoacyl-CoA thiolase 947129 Ribosome-associated inhibitor A 948327 Transcriptional activator rfaH 948326 3-octaprenyl-4-hydroxybenzoate carboxy-lyase 948321 Sec-independent protein translocase protein tatA 948320 Putative carboxymethylenebutenolidase | Central intermediary metabolism' | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated |
| xdhC mppA ThiM lpxM lpxM yniB yjgA nrdD ydbH ydhV fbp yddA uspF fre fadA raiA rfaH ubiD tatA ysgA | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945951 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yigA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbH 948753 Fructose-1,6-bisphosphatase class 1 945948 Universal stress protein F 948325 NAD(P)H-flavin reductase 948324 3-ketoacyl-CoA thiolase 947129 Ribosome-associated inhibitor A 948327 Transcriptional activator rfaH 948326 3-octaprenyl-4-hydroxybenzoate carboxy-lyase 948321 Sec-independent protein translocase protein tatA 948320 Putative carboxymethylenebutenolidase 947120 Putrescine aminotransferase | Central intermediary metabolism' | Down Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Up Regulated Up Regulated Up Regulated Down Regulated Up Regulated |
| xdhC mppA ThiM lpxM yniB yigA nrdD ydbH ydhV fbp yddA uspF fre fadA raiA rfaH ubiD tatA ysgA | 947177 Deoxyguanosinetriphosphate triphosphohydrolase 947174 Sensor protein qseC; Sensor protein 945148 Xanthine dehydrogenase iron-sulfur-binding subunit 945947 Periplasmic murein peptide-binding protein 945142 Hydroxyethylthiazole kinase 945143 Lipid A biosynthesis (KDO)2-(lauroyl)-lipid IVA acyltransferase 945140 Uncharacterized protein yniB 948751 UPF0307 protein yjgA 948755 Anaerobic ribonucleoside-triphosphate reductase 945949 Uncharacterized protein ydbH 948756 Uncharacterized protein ydbH 948757 Fructose-1,6-bisphosphatase class 1 945945 Inner membrane ABC transporter ATP-binding protein yddA 945948 Universal stress protein F 948325 NAD(P)H-flavin reductase 948324 3-ketoacyl-CoA thiolase 947129 Ribosome-associated inhibitor A 948327 Transcriptional activator rfaH 948326 3-octaprenyl-4-hydroxybenzoate carboxy-lyase 948321 Sec-independent protein translocase protein tatA 948320 Putative carboxymethylenebutenolidase | [] [] Central intermediary metabolism'] | Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated |

| | 04544011 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 | | Tu B |
|------|---|---|-------------------------|
| ybbP | 945118 Uncharacterized ABC transporter permease ybbP | | Up Regulated |
| ydgH | 945117 Protein ydgH | | Down Regulated |
| yjgl | 948765 Uncharacterized oxidoreductase yjgl | | Down Regulated |
| pyrB | 948767 Aspartate carbamoyltransferase catalytic chain; Aspartate carba | ımoyltransferase | Ambiguous/Unsorted |
| treR | 948760 HTH-type transcriptional regulator treR | | Down Regulated |
| pyrl | 948763 Aspartate carbamoyltransferase regulatory chain | | Down Regulated |
| treC | 948762 Trehalose-6-phosphate hydrolase | | Ambiguous/Unsorted |
| xdhA | 947116 Xanthine dehydrogenase molybdenum-binding subunit | | Ambiguous/Unsorted |
| | 948338 Uncharacterized protein yigl | n | Down Regulated |
| yigl | | Ш | |
| ybbA | 945122 Uncharacterized ABC transporter ATP-binding protein ybbA | | Ambiguous/Unsorted |
| fadB | 948336 Fatty acid oxidation complex subunit alpha | | Up Regulated |
| pepQ | 948335 Xaa-Pro dipeptidase | | Ambiguous/Unsorted |
| queD | 945123 Putative 6-pyruvoyl tetrahydrobiopterin synthase | | Up Regulated |
| hem(| 948331 Protoporphyrinogen oxidase | | Ambiguous/Unsorted |
| ompV | 945128 Outer membrane protein W | | Up Regulated |
| tesA | 945127 Acyl-CoA thioesterase I | | Down Regulated |
| tabA | 948777 Uncharacterized protein yigK | | Down Regulated |
| | 948776 Uncharacterized protein yigh | Type of regulation | Up Regulated |
| yjgL | | | |
| yjgJ | 948775 Uncharacterized HTH-type transcriptional regulator yjgJ | [Cell wall membrane envelope biogenesis', 'Intracellular trafficking secretion and vesicular transport', 'Cell division', 'protein transp | |
| argl | 948774 Ornithine carbamoyltransferase chain I; Ornithine carbamoyltran | sferase | Down Regulated |
| rraB | 948773 Uncharacterized protein yjgD | | Up Regulated |
| yjgM | 948772 Uncharacterized N-acetyltransferase yjgM | | Temporarly Up Regulated |
| uvrD | 948347 DNA helicase II | | Temporarly Down Regula |
| rdoA | 948346 Protein rdoA | | Temporarly Down Regula |
| mob/ | 948349 Molybdopterin-guanine dinucleotide biosynthesis protein A | | Down Regulated |
| proV | 947148 Glycine betaine/L-proline transport ATP-binding protein proV | | Temporarly Down Regula |
| | | | |
| nrdF | 947149 Ribonucleoside-diphosphate reductase 2 subunit beta | | Up Regulated |
| ygaU | 947144 Uncharacterized protein ygaU | Ш | Ambiguous/Unsorted |
| ygiS | 947140 Putative binding protein ygiS | | Up Regulated |
| allD | 948342 Ureidoglycolate dehydrogenase | | Temporarly Down Regula |
| holC | 948787 DNA polymerase III subunit chi | | Ambiguous/Unsorted |
| recE | 945918 Exodeoxyribonuclease 8 | | Temporarly Down Regula |
| yccU | 949008 Uncharacterized protein yccU | П | Up Regulated |
| fadE | 949007 Acyl-coenzyme A dehydrogenase | | Up Regulated |
| btuE | 945915 Vitamin B12 transport periplasmic protein btuE; Glutathione perc | L. J. | Ambiguous/Unsorted |
| | | DAIGUSSE | |
| abgT | 945912 Aminobenzoyl-glutamate transport protein | | Ambiguous/Unsorted |
| rpsl | 949000 30S ribosomal protein S9 | | Down Regulated |
| yjgN | 948784 Inner membrane protein yjgN | | Ambiguous/Unsorted |
| prfA | 949002 Peptide chain release factor 1 | | Ambiguous/Unsorted |
| YigA | 948359 Uncharacterized protein yigA | | Temporarly Up Regulated |
| recR | 945100 Recombination protein recR | | Down Regulated |
| degP | 947139 Protease do | | Ambiguous/Unsorted |
| yihF | 948352 Uncharacterized protein yihF | [Replication recombination and repair', ' Adaptation to stress'] | Down Regulated |
| | | [Nephication recombination and repair, Adaptation to stress] | |
| corA | 948351 Magnesium transport protein corA | | Temporarly Down Regula |
| yahE | 947134 Uncharacterized protein yahE | Ш | Temporarly Up Regulated |
| dnaX | 945105 DNA polymerase III subunit tau | | Ambiguous/Unsorted |
| ydaN | 945910 Zinc transport protein zntB | | Ambiguous/Unsorted |
| ftsK | 945102 DNA translocase ftsK | | Down Regulated |
| polA | 948356 DNA polymerase I | | Ambiguous/Unsorted |
| dps | 945101 DNA protection during starvation protein | | Up Regulated |
| xerC | 948355 Tyrosine recombinase xerC | | Temporarly Up Regulated |
| | | | |
| stpA | 947130 DNA-binding protein stpA | | Ambiguous/Unsorted |
| ybaB | 945104 UPF0133 protein ybaB | | Ambiguous/Unsorted |
| dsbA | 948353 Thiol:disulfide interchange protein dsbA | | Up Regulated |
| ykgG | 945906 UPF0707 protein ykgG | | Up Regulated |
| fnr | 945908 Fumarate and nitrate reduction regulatory protein | | Down Regulated |
| uspE | 945904 Universal stress protein E | | Up Regulated |
| folK | 948792 2-amino-4-hydroxy-6-hydroxymethyldihydropteridine pyrophospl | hokinase | Up Regulated |
| lptF | 948795 Lipopolysaccharide export system permease protein lptF | | Down Regulated |
| | 949024 Uncharacterized response regulatory protein yehT | "transcription," regulation of transcription, DNA dependent" | |
| yehT | | ['transcription', 'regulation of transcription DNA dependent'] | Down Regulated |
| CAR | 949025 Carbamoyl-phosphate synthase small chain | | Ambiguous/Unsorted |
| yacL | 944809 UPF0231 protein yacL | | Up Regulated |
| yehS | 949026 Uncharacterized protein yehS | [Replication recombination and repair'] | Temporarly Up Regulated |
| idi | 949020 Isopentenyl-diphosphate Delta-isomerase | | Ambiguous/Unsorted |
| ftsL | 944803 Cell division protein ftsL | | Down Regulated |
| | | | <u> </u> |

| fruD | 044904 Eruptoco repressor | | Tomporarly Down Pogula |
|--|---|---------------------------------|---|
| fruR mlrA | 944804 Fructose repressor 949029 HTH-type transcriptional regulator mlrA | | Temporarly Down Regula |
| | | | Ambiguous/Unsorted |
| acpP | 944805 Acyl carrier protein | | Down Regulated |
| | 946031 NAD-dependent malic enzyme | | Ambiguous/Unsorted |
| cfa | 944811 Cyclopropane-fatty-acyl-phospholipid synthase | | Temporarly Up Regulated |
| mraZ | 944810 Protein mraZ | | Up Regulated |
| | 946035 Formate dehydrogenase, nitrate-inducible, major subunit | | Down Regulated |
| hyfB | 946961 Hydrogenase-4 component B | | Temporarly Up Regulated |
| | 946960 Oxidoreductase ydhF | | Up Regulated |
| | 946036 Alcohol dehydrogenase, propanol-preferring | | Up Regulated |
| ycgK | 946963 Uncharacterized protein ycgK | μ | Up Regulated |
| frsA | 946039 Esterase frsA | | Ambiguous/Unsorted |
| sufA | 949014 Protein sufA | | Up Regulated |
| 70 | 949018 Uncharacterized protein ygfT | [electron transport'] | Ambiguous/Unsorted |
| yncH | 946024 Uncharacterized protein yncH | Ш | Ambiguous/Unsorted |
| ррх | 946970 Exopolyphosphatase | | Down Regulated |
| fucK | 946022 L-fuculokinase | | Ambiguous/Unsorted |
| yfgD | 946974 Uncharacterized protein yfgD | | Ambiguous/Unsorted |
| yqcC | 946027 Uncharacterized protein yqcC | μ | Ambiguous/Unsorted |
| rhsE | 946026 ECDH10B_1586; EC55989_0238; b1456 | | Down Regulated |
| ybjX | 946025 Uncharacterized protein ybjX | Ц | Down Regulated |
| | 946977 DnaA-homolog protein hda | | Down Regulated |
| | 949042 Colicin I receptor | | Ambiguous/Unsorted |
| galS | 949043 HTH-type transcriptional regulator galS | | Up Regulated |
| yeiG | 949045 S-formylglutathione hydrolase yeiG | | Down Regulated |
| artQ | 949046 Arginine transport system permease protein artQ | | Ambiguous/Unsorted |
| | 949048 3-deoxy-D-manno-octulosonic-acid transferase | | Down Regulated |
| gltl | 946938 Glutamate/aspartate periplasmic-binding protein | | Ambiguous/Unsorted |
| yeiT | 949049 Uncharacterized oxidoreductase yeiT | [Primary Active Transporters'] | Temporarly Up Regulated |
| | 944827 Pyruvate dehydrogenase complex repressor | | Ambiguous/Unsorted |
| | 944823 Cell division protein ftsQ | | Down Regulated |
| | 944824 Mutator mutT protein | | Down Regulated |
| | 946055 Protein ydeP | | Down Regulated |
| | 944833 Uncharacterized ABC transporter ATP-binding protein yadG | | Ambiguous/Unsorted |
| | 946054 Protein gnsB | | Temporarly Up Regulated |
| | 946057 Probable glutamate/gamma-aminobutyrate antiporter | | Ambiguous/Unsorted |
| gcd | 944830 Quinoprotein glucose dehydrogenase | | Ambiguous/Unsorted |
| ygiW | 946051 Protein ygiW | | Temporarly Up Regulated |
| ydeQ | 946050 Uncharacterized fimbrial-like protein ydeQ | Ц | Temporarly Up Regulated |
| | 946053 Uncharacterized protein ypfl | | Down Regulated |
| | 946052 Probable D,D-dipeptide-binding periplasmic protein ddpA | | Up Regulated |
| PqqL | 946059 Probable zinc protease pqqL | | Up Regulated |
| | 949041 D-galactose-binding periplasmic protein | | Up Regulated |
| | 946941 50S ribosomal protein L28 | | Down Regulated |
| eutD | 946940 Ethanolamine utilization protein eutD | | Down Regulated |
| | 949033 3-octaprenyl-4-hydroxybenzoate carboxy-lyase | | Ambiguous/Unsorted |
| leuO | 949034 Probable HTH-type transcriptional regulator leuO | | Up Regulated |
| | 946948 Nitrate/nitrite sensor protein narQ | | Up Regulated |
| bcp | 946949 Putative peroxiredoxin bcp | | Temporarly Down Regula |
| ppiB | 949038 Peptidyl-prolyl cis-trans isomerase B; Peptidyl-prolyl cis-trans is | omerase | Up Regulated |
| thyA | 949035 Thymidylate synthase | | Down Regulated |
| mglA | 949036 Galactose/methyl galactoside import ATP-binding protein mglA | | Up Regulated |
| maeE | 946947 NADP-dependent malic enzyme | | Ambiguous/Unsorted |
| linv(, i | | | |
| | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylase | e T | Ambiguous/Unsorted |
| Gpt | 944816 UDP-3-0-[3-hydroxymyristoyl] N-acetylglucosamine deacetylast 944817 Xanthine phosphoribosyltransferase | e | Ambiguous/Unsorted |
| Gpt murD | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylase 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase | e | Ambiguous/Unsorted Ambiguous/Unsorted |
| Gpt murD mgIC | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylast 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase 949039 Galactoside transport system permease protein mglC | e | Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated |
| Gpt murD mglC murF | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylast 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase 949039 Galactoside transport system permease protein mglC 944813 UDP-N-acetylmuramoyl-tripeptideD-alanyl-D-alanine ligase | e | Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted |
| Gpt murD mglC murF wzzE | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylase 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase 949039 Galactoside transport system permease protein mglC 944813 UDP-N-acetylmuramoyl-tripeptideD-alanyi-D-alanine ligase 944815 Lipopolysaccharide biosynthesis protein wzzE | e | Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted |
| murD mglC murF wzzE nhaB | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylast 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase 949039 Galactoside transport system permease protein mglC 944813 UDP-N-acetylmuramoyl-tripeptideD-alanyl-D-alanine ligase 944815 Lipopolysaccharide biosynthesis protein wzzE 944822 Na(+)/H(+) antiporter nhaB | e | Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated |
| Gpt murD mglC murF wzzE nhaB osmC | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylast 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase 949039 Galactoside transport system permease protein mglC 944813 UDP-N-acetylmuramoyl-tripeptideD-alanyl-D-alanine ligase 944815 Lipopolysaccharide biosynthesis protein wzzE 944822 Na(+)/H(+) antiporter nhaB 946043 Peroxiredoxin osmC | | Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Up Regulated |
| Gpt murD mgIC murF wzzE nhaB osmC ypfJ | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylast 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase 949039 Galactoside transport system permease protein mglC 944813 UDP-N-acetylmuramoyl-tripeptideD-alanyl-D-alanine ligase 944815 Lipopolysaccharide biosynthesis protein wzzE 944822 Na(+)/H(+) antiporter nhaB 946043 Peroxiredoxin osmC 946956 Uncharacterized protein ypfJ | © | Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Up Regulated Up Regulated |
| Gpt murD mglC murF wzzE nhaB osm(ypfJ rnhA | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylast 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase 949039 Galactoside transport system permease protein mglC 944813 UDP-N-acetylmuramoyl-tripeptideD-alanyl-D-alanine ligase 944815 Lipopolysaccharide biosynthesis protein wzzE 944812 Na(+)/H(+) antiporter nhaB 946043 Peroxiredoxin osmC 946956 Uncharacterized protein ypfJ 946955 Ribonuclease HI; Ribonuclease H | | Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Up Regulated Up Regulated Up Regulated Down Regulated |
| Gpt murD mglC murF wzzE nhaB osmC ypfJ mhA yffB | 944816 UDP-3-O-[3-hydroxymyristoyl] N-acetylglucosamine deacetylast 944817 Xanthine phosphoribosyltransferase 944818 UDP-N-acetylmuramoylalanineD-glutamate ligase 949039 Galactoside transport system permease protein mglC 944813 UDP-N-acetylmuramoyl-tripeptideD-alanyl-D-alanine ligase 944815 Lipopolysaccharide biosynthesis protein wzzE 944822 Na(+)/H(+) antiporter nhaB 946043 Peroxiredoxin osmC 946956 Uncharacterized protein ypfJ | | Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Up Regulated Up Regulated |

| gcvR | 946950 | Glycine cleavage system transcriptional repressor | | Ambiguous/Unsorted |
|------|--------|--|---|-------------------------|
| ligT | | 2'-5'-RNA ligase | | Down Regulated |
| yadB | | Glutamyl-Q tRNA(Asp) synthetase; Insertion element IS150 und | haracterized 14 kDa protein | Temporarly Up Regulated |
| uxaC | | Uronate isomerase | | Up Regulated |
| hrpB | | ATP-dependent RNA helicase hrpB | | Ambiguous/Unsorted |
| ycgF | | Uncharacterized protein ycgF | Signal transduction mechanisms', 'two component signal transduction system phosphorelay ', 'regulation of transcription DNA dep | |
| ygfK | | Uncharacterized protein yegfK | Togeth an addition medianion, two component agran an addition by seem proprietary, regulation of adries paid. | Temporarly Down Regula |
| rna | | Ribonuclease I | | Temporarly Down Regula |
| fadH | | 2,4-dienoyl-CoA reductase [NADPH] | | Up Regulated |
| coaE | | Dephospho-CoA kinase | | Ambiguous/Unsorted |
| dksA | | DnaK suppressor protein | | Up Regulated |
| ydhQ | | Uncharacterized protein ydhQ | ['Translation ribosomal structure and biogenesis', 'Adaptation to stress', 'translation'] | Temporarly Down Regula |
| tam | | Trans-aconitate 2-methyltransferase | | Up Regulated |
| HtpX | | Probable protease htpX | | Down Regulated |
| sfsA | | Sugar fermentation stimulation protein A | | Up Regulated |
| rnhB | | Ribonuclease HII | | Down Regulated |
| yneJ | | Uncharacterized HTH-type transcriptional regulator yneJ | [Transcription', 'RNA related', 'Type of regulation', 'Genetic unit regulated', 'transcription', 'regulation of transcription DNA depende | |
| panB | 944839 | 3-methyl-2-oxobutanoate hydroxymethyltransferase | | Down Regulated |
| lpxB | | Lipid-A-disaccharide synthase | | Temporarly Down Regula |
| rpIU | | 50S ribosomal protein L21 | | Temporarly Up Regulated |
| Irp | | Leucine-responsive regulatory protein | | Ambiguous/Unsorted |
| cydD | | ATP-binding/permease protein cydD | | Ambiguous/Unsorted |
| НурВ | | Hydrogenase isoenzymes nickel incorporation protein hypB | | Down Regulated |
| hipA | | Protein hipA | | Temporarly Up Regulated |
| ygbK | | Uncharacterized protein ygbK | ['Carbohydrate transport and metabolism'] | Down Regulated |
| yadN | | Uncharacterized fimbrial-like protein yadN | [' Building block biosynthesis'] | Ambiguous/Unsorted |
| vdel | 946068 | Uncharacterized protein ydel | ['Signal transduction mechanisms', 'response to stress'] | Up Regulated |
| hipB | 946065 | HTH-type transcriptional regulator hipB | | Temporarly Up Regulated |
| ykgC | | Probable pyridine nucleotide-disulfide oxidoreductase ykgC | | Ambiguous/Unsorted |
| ulaG | | Probable L-ascorbate-6-phosphate lactonase ulaG | | Temporarly Up Regulated |
| argP | | Chromosome initiation inhibitor | | Down Regulated |
| mak | | Fructokinase | | Up Regulated |
| symE | 949088 | Endoribonuclease symE | | Ambiguous/Unsorted |
| ymdE | | UPF0189 protein ymdB | | Up Regulated |
| xseA | | Exodeoxyribonuclease 7 large subunit | | Ambiguous/Unsorted |
| narP | 949081 | Nitrate/nitrite response regulator protein narP | | Temporarly Down Regula |
| der | 946983 | GTP-binding protein engA | | Down Regulated |
| yfgJ | | Uncharacterized protein yfgJ | [' Type of regulation'] | Down Regulated |
| ycil | 949085 | Protein ycil | | Up Regulated |
| dnaE | 944877 | DNA polymerase III subunit alpha | | Down Regulated |
| ispU | 944874 | Undecaprenyl pyrophosphate synthetase | | Up Regulated |
| yfgM | 946981 | UPF0070 protein yfgM | | Ambiguous/Unsorted |
| yaeR | 944875 | Uncharacterized protein yaeR | | Ambiguous/Unsorted |
| flhE | 946094 | Flagellar protein flhE | | Down Regulated |
| glnH | 944872 | Glutamine-binding periplasmic protein | | Ambiguous/Unsorted |
| yacF | | UPF0289 protein yacF | | Temporarly Down Regula |
| prc | 946096 | Tail-specific protease | | Down Regulated |
| rseP | | Regulator of sigma E protease | | Down Regulated |
| yneK | 946080 | Uncharacterized protein yneK | | Ambiguous/Unsorted |
| ulaF | 948711 | L-ribulose-5-phosphate 4-epimerase ulaF | | Temporarly Down Regula |
| erpA | 944857 | Iron-sulfur cluster insertion protein erpA | | Ambiguous/Unsorted |
| ulaD | 948714 | 3-keto-L-gulonate-6-phosphate decarboxylase ulaD | | Temporarly Down Regula |
| fhuA | | Ferrichrome-iron receptor | | Up Regulated |
| yciM | 944858 | Uncharacterized protein yciM | | Temporarly Up Regulated |
| sbcC | 949076 | Nuclease sbcCD subunit C | | Temporarly Down Regula |
| | | Probable hypoxanthine oxidase xdhD | | Ambiguous/Unsorted |
| iscA | | Iron-binding protein iscA | | Ambiguous/Unsorted |
| hscB | | Co-chaperone protein hscB | | Down Regulated |
| | | Thiol:disulfide interchange protein dsbE | | Temporarly Up Regulated |
| flgH | | Flagellar L-ring protein | | Down Regulated |
| sseA | | 3-mercaptopyruvate sulfurtransferase | | Up Regulated |
| | | Uncharacterized protein ygfM | | Ambiguous/Unsorted |
| | | Protein sseB | | Temporarly Down Regula |
| ytjB | | Protein smp | | Down Regulated |
| ulaR | 948706 | HTH-type transcriptional regulator ulaR | | Up Regulated |
| | | | | |

| ispG | 946991 4-hydroxy-3-methylbut-2-en-1-yl diphosphate synthase | | Down Regulated |
|--------------|---|---|--|
| glnD | 944863 [Protein-PII] uridylyltransferase | | Down Regulated |
| yifP | 948707 Esterase yifP | | Up Regulated |
| ydfH | 946087 Uncharacterized HTH-type transcriptional regulator ydfH | n | Temporarly Up Regulated |
| ybbL | 946990 Uncharacterized ABC transporter ATP-binding protein ybbL | u Macromolecules cellular constituent biosynthesis', ' Primary Active Transporters', 'DNA repair', 'response to DNA damage stimulu | |
| tsf | 944866 Elongation factor Ts | I macromolecules central constituent biosynthesis, i filmary Active fransporters, biva repair, response to biva damage sumulo | Down Regulated |
| ydfG | 946085 NADP-dependent L-serine/L-allo-threonine dehydrogenase ydf0 | | Ambiguous/Unsorted |
| yebR | 946086 UPF0067 protein yebR | | Down Regulated |
| skp | 944861 Chaperone protein skp | | Up Regulated |
| dapD | 944862 2,3,4,5-tetrahydropyridine-2,6-dicarboxylate N-succinyltransfera | Se | Down Regulated |
| caiC | 944886 Probable crotonobetaine/carnitine-CoA ligase | | Temporarly Up Regulated |
| hscA | 944885 Chaperone protein hscA | | Temporarly Down Regula |
| lpxD | 944882 UDP-3-O-[3-hydroxymyristoyl] glucosamine N-acyltransferase | | Ambiguous/Unsorted |
| kdsB | 945539 3-deoxy-manno-octulosonate cytidylyltransferase | | Down Regulated |
| folM | 949096 Dihydrofolate reductase folM | | Temporarly Down Regula |
| hupB | 949095 DNA-binding protein HU-beta | | Ambiguous/Unsorted |
| fadJ | 949097 Fatty acid oxidation complex subunit alpha | | Up Regulated |
| ycbC | 945546 Uncharacterized protein ycbC | [regulation of cell shape', 'peptidoglycan biosynthetic process'] | Temporarly Up Regulated |
| mukF | 945548 Chromosome partition protein mukF | | Down Regulated |
| smtA | 945547 Protein smtA | | Up Regulated |
| ycbB | 945541 Uncharacterized protein ycbB | ['DNA replication'] | Temporarly Up Regulated |
| rimP | 947680 b3170 | | Temporarly Down Regula |
| rlmE | 947689 b3179 | | Ambiguous/Unsorted |
| yhbY | 947688 RNA-binding protein yhbY | | Ambiguous/Unsorted |
| sufD | 944878 Protein sufD | | Up Regulated |
| truB | 947687 tRNA pseudouridine synthase B | | Down Regulated |
| gmhE | 944879 D,D-heptose 1,7-bisphosphate phosphatase | | Down Regulated |
| rpsO | 947686 30S ribosomal protein S15 | | Ambiguous/Unsorted |
| yafL | 944899 Uncharacterized lipoprotein yafL | [Cell wall membrane envelope biogenesis', 'lipid biosynthetic process', 'response to antibiotic'] | Down Regulated |
| metl | 944894 D-methionine transport system permease protein metl | | Up Regulated |
| rof | 944891 Protein rof | I Duilding blook bigg with soil Jaming soil bigg with the suppose! | Ambiguous/Unsorted |
| yejH | 945529 Uncharacterized protein yejH 945528 3-phosphoshikimate 1-carboxyvinyltransferase | [' Building block biosynthesis', 'amino acid biosynthetic process'] | Temporarly Up Regulated Ambiguous/Unsorted |
| aroA ycaR | 945537 UPF0434 protein ycaR | | Ambiguous/Unsorted |
| rpsA | 945536 30S ribosomal protein S1 | | Ambiguous/Unsorted |
| deaD | 947674 Cold-shock DEAD box protein A | | Down Regulated |
| cmk | 945535 Cytidylate kinase | | Ambiguous/Unsorted |
| ycaL | 945534 Uncharacterized metalloprotease ycaL | п | Up Regulated |
| nlpl | 947673 Lipoprotein nlpl | u . | Temporarly Up Regulated |
| ihfB | 945533 Integration host factor subunit beta | | Ambiguous/Unsorted |
| yhbS | 947670 Uncharacterized N-acetyltransferase yhbS | Building block biosynthesis'] | Up Regulated |
| msb/ | 945530 Lipid A export ATP-binding/permease protein msbA | | Down Regulated |
| tilS | 944889 tRNA(IIe)-lysidine synthase | | Ambiguous/Unsorted |
| yhbW | 947677 Uncharacterized protein yhbW | n | Ambiguous/Unsorted |
| rlmL | 945564 Ribosomal RNA large subunit methyltransferase L | | Ambiguous/Unsorted |
| uup | 945566 ABC transporter ATP-binding protein uup | | Down Regulated |
| fabA | 945568 3-hydroxydecanoyl-[acyl-carrier-protein] dehydratase | | Ambiguous/Unsorted |
| ycbZ | 945569 Putative protease La homolog | | Ambiguous/Unsorted |
| mukE | 945549 Chromosome partition protein mukB | | Down Regulated |
| ftsH | 947690 Cell division protease ftsH | | Ambiguous/Unsorted |
| ompF | 945554 Outer membrane protein F | | Down Regulated |
| obgE | | ;; ECH74115_4505; ECUMN_3663; EcolC_0517; c3940; EcHS_A3376; EC55989_3601; ECSE_3467; ECED1_3841; ECDH10B_33 | |
| dacB | 947693 D-alanyl-D-alanine carboxypeptidase dacB | | Ambiguous/Unsorted |
| greA | 947696 Transcription elongation factor; Transcription elongation factor g | reA | Ambiguous/Unsorted |
| pyrD | 945556 Dihydroorotate dehydrogenase | | Ambiguous/Unsorted |
| fis | 947697 DNA-binding protein fis | | Ambiguous/Unsorted |
| yhbE | 947699 Uncharacterized inner membrane transporter yhbE | [' Type of regulation'] | Temporarly Up Regulated |
| mukE | 945550 Chromosome partition protein mukE | | Ambiguous/Unsorted |
| copA | 946106 Copper-exporting P-type ATPase A | PColl motified Untracellular tentfolding appraising and ungiquestors and University University and University | Ambiguous/Unsorted |
| ycbS | 946934 Uncharacterized outer membrane usher protein ycbS | [Cell motility', 'Intracellular trafficking secretion and vesicular transport', 'Macromolecules cellular constituent biosynthesis'] | Ambiguous/Unsorted |
| yeeO agaD | 946506 Uncharacterized transporter yeeO 947649 N-acetylgalactosamine permease IID component | <u> </u> | Up Regulated Ambiguous/Unsorted |
| hisF | 946516 midazole glycerol phosphate synthase subunit hisF | | Ambiguous/Unsorted |
| hisl | 946515 Histidine biosynthesis bifunctional protein hislE | | Down Regulated |
| yeeU | 946510 Uncharacterized protein yeeU | | Up Regulated |
| yeeu | OTOO TO OTO I OTO | I. | op regulated |

| D | 040540 | Inner membrone protein veeD | | Un Domilated |
|--------------|--------|---|--|--------------------------------|
| yeeR | | Inner membrane protein yeeR | | Up Regulated |
| garD | | D-galactarate dehydratase | | Temporarly Up Regulated |
| rihA | | Pyrimidine-specific ribonucleoside hydrolase rihA | | Up Regulated |
| cydC | | ATP-binding/permease protein cydC | | Down Regulated |
| infA | | Translation initiation factor IF-1 | | Up Regulated |
| grxB | | Glutaredoxin-2 | | Ambiguous/Unsorted |
| ynfB | | UPF0482 protein ynfB | | Ambiguous/Unsorted |
| speG | | Spermidine N(1)-acetyltransferase | | Down Regulated |
| ycgM | | Uncharacterized protein ycgM | | Up Regulated |
| pyrG | | CTP synthase | | Temporarly Down Regula |
| ydfE | | b1577; ECDH10B_1709 | | Temporarly Down Regula |
| ynjE | | Putative thiosulfate sulfurtransferase ynjE | | Temporarly Up Regulated |
| sxy | | Protein sxy | | Down Regulated |
| agaR | 947636 | Putative aga operon transcriptional repressor | | Temporarly Down Regula |
| cbl | | HTH-type transcriptional regulator cbl | | Down Regulated |
| garR | | 2-hydroxy-3-oxopropionate reductase | | Up Regulated |
| efeB | | Peroxidase ycdB | | Down Regulated |
| garK | | Glycerate kinase 2 | | Up Regulated |
| yfeY | | Uncharacterized protein yfeY | | Ambiguous/Unsorted |
| ypeA | | Acetyltransferase ypeA | | Down Regulated |
| amiA | | Probable N-acetylmuramoyl-L-alanine amidase amiA | | Down Regulated |
| bisC | | Biotin sulfoxide reductase | | Down Regulated |
| eutL | 946914 | Ethanolamine utilization protein eutL | | Down Regulated |
| yfeX | | Uncharacterized protein yfeX | [' Building block biosynthesis'] | Ambiguous/Unsorted |
| pflA | 945517 | Pyruvate formate-lyase 1-activating enzyme | | Ambiguous/Unsorted |
| acrR | | HTH-type transcriptional regulator acrR | | Up Regulated |
| trpC | | Tryptophan biosynthesis protein trpCF | | Down Regulated |
| sufC | | Probable ATP-dependent transporter sufC | | Ambiguous/Unsorted |
| kdgR | 946129 | Transcriptional regulator kdgR | | Ambiguous/Unsorted |
| rimJ | | Ribosomal-protein-alanine acetyltransferase | | Ambiguous/Unsorted |
| sbcB | 946529 | Exodeoxyribonuclease I | | Down Regulated |
| yeeY | 946532 | Uncharacterized HTH-type transcriptional regulator yeeY | Ր RNA related'] | Down Regulated |
| yraR | | Uncharacterized protein yraR | | Up Regulated |
| yraL | | UPF0011 protein yraL | | Ambiguous/Unsorted |
| rsml | | UPF0011 protein yraL | | Ambiguous/Unsorted |
| yeeF | | Inner membrane transport protein yeeF | | Temporarly Up Regulated |
| yhbP | | UPF0306 protein yhbP | | Ambiguous/Unsorted |
| yeeZ | | Protein yeeZ | | Ambiguous/Unsorted |
| yhbT | 947669 | Uncharacterized protein yhbT | ['Coenzyme transport and metabolism', ' Energy metabolism' carbon ', ' Building block biosynthesis'] | Down Regulated |
| yajO | 946903 | Uncharacterized oxidoreductase yajO | The state of the s | Up Regulated |
| ycaK | | Uncharacterized NAD(P)H oxidoreductase ycaK | amino acid transport'] | Temporarly Up Regulated |
| ampl | | Penicillin-binding protein ampH | [animo acid dansport] | Down Regulated |
| ycaN | | Uncharacterized HTH-type transcriptional regulator ycaN | ['Transcription', ' RNA related', ' Type of regulation', 'transcription', 'regulation of transcription DNA dependent'] | Down Regulated |
| ycaQ | | Uncharacterized protein ycaQ | ['two component signal transduction system phosphorelay'] | Ambiguous/Unsorted |
| ivy | | Inhibitor of vertebrate lysozyme | [wo component signal transduction system phosphoretay] | Up Regulated |
| diaA | | DnaA initiator-associating protein diaA | | Up Regulated |
| hemF | | Coproporphyrinogen-III oxidase, aerobic | | Down Regulated |
| lpxK | | Tetraacyldisaccharide 4'-kinase | | Down Regulated Down Regulated |
| | | Hydroxyacylglutathione hydrolase | | Down Regulated Down Regulated |
| gloB | | Electron transport complex protein rnfC | | Down Regulated Down Regulated |
| rsxC | | Anaerobic dimethyl sulfoxide reductase chain B | | Down Regulated Down Regulated |
| dmsE | | Putative dimethyl sulfoxide reductase chain ynfE | | |
| ynfE | | | | Up Regulated |
| dkgB | | 2,5-diketo-D-gluconic acid reductase B | | Ambiguous/Unsorted |
| rarA | | Replication-associated recombination protein A | | Ambiguous/Unsorted |
| yafJ | | Putative glutamine amidotransferase yafJ | | Ambiguous/Unsorted |
| - | | S-adenosylmethionine:tRNA ribosyltransferase-isomerase | | Ambiguous/Unsorted |
| ycaO | | UPF0142 protein ycaO | | Down Regulated |
| cobT | | Nicotinate-nucleotidedimethylbenzimidazole phosphoribosyltra | Insterase | Ambiguous/Unsorted |
| dacD | | D-alanyl-D-alanine carboxypeptidase dacD | | Temporarly Down Regula |
| ynfD | | Uncharacterized protein ynfD | ['Energy production and conversion', ' Electrochemical potential driven transporters', 'response to stress'] | Up Regulated |
| ynfC | | UPF0257 lipoprotein ynfC | | Down Regulated |
| poxB | | Pyruvate dehydrogenase [cytochrome] | | Up Regulated |
| mreC | 947655 | Rod shape-determining protein mreC | | Down Regulated |
| | | | ITA mains a sid transport and marketedianal ICa american transport and marketedianal I Cantral intermedian constable liquell | II In Dogulated |
| yral hslJ | | Uncharacterized fimbrial chaperone yral Heat shock protein hslJ | ['Amino acid transport and metabolism', 'Coenzyme transport and metabolism', 'Central intermediary metabolism'] | Up Regulated Up Regulated |

| vroD 047650 Incharacterized protein vroD | IDNA conjugation! | Un Dogulated |
|---|---|-------------------------------------|
| yraP 947659 Uncharacterized protein yraP ddsA 945510 Protein mlc | ['DNA replication'] | Up Regulated Temporarly Down Regula |
| dgsA 945510 Protein mlc entC 945511 Isochorismate synthase entC | | Up Regulated |
| yafK 944910 Uncharacterized protein yafK | n | Down Regulated |
| ycaC 945512 Uncharacterized protein yark | lu l | Up Regulated |
| miaA 948690 tRNA Delta(2)-isopentenylpyrophosphate transferase | | Down Regulated |
| yafN 944920Uncharacterized protein yafN | ['Replication recombination and repair', ' RNA related', ' Type of regulation', ' Protection', 'transcription', 'regulation of transcription' | |
| tpiA 948409 Triosephosphate isomerase | [Neplication recombination and repair, New related, Type of regulation, Trotection, transcription, regulation of transcription | Temporarly Up Regulated |
| mutL 948691 DNA mismatch repair protein mutL | | Temporarly Down Regula |
| rnr 948692 Ribonuclease R | | Ambiguous/Unsorted |
| nsrR 948693HTH-type transcriptional repressor nsrR | | Down Regulated |
| rlmB 948694 23S rRNA (guanosine-2'-O-)-methyltransferase rlmB | | Ambiguous/Unsorted |
| uxaA 947603 Altronate hydrolase | | Up Regulated |
| exuR 947602 Exu regulon transcriptional regulator | | Ambiguous/Unsorted |
| yjfC 948699 Uncharacterized protein yjfC | ['electron transport', 'metabolic process'] | Temporarly Up Regulated |
| sstT 947605 Serine/threonine transporter sstT | | Down Regulated |
| wza 946558 Putative polysaccharide export protein wza | | Ambiguous/Unsorted |
| ygjR 947600 Uncharacterized oxidoreductase ygjR | ['Carbohydrate transport and metabolism'] | Up Regulated |
| cdh 948410 CDP-diacylglycerol pyrophosphatase | | Up Regulated |
| sbp 948411 Sulfate-binding protein | | Temporarly Down Regula |
| fieF 948413 Ferrous-iron efflux pump fieF | | Down Regulated |
| fpr 948414 FerredoxinNADP reductase | | Up Regulated |
| uspD 948415 Universal stress protein D | | Up Regulated |
| dinJ 944914 DNA-damage-inducible protein J | | Temporarly Up Regulated |
| yiiS 948416 UPF0381 protein yiiS | | Up Regulated |
| rraA 948419 Regulator of ribonuclease activity A | | Down Regulated |
| flu 946540 Antigen 43 | | Up Regulated |
| yjeF 948685 Uncharacterized protein yjeF | | Up Regulated |
| amiB 948683 N-acetylmuramoyl-L-alanine amidase amiB | | Down Regulated |
| yjeE 948684 UPF0079 ATP-binding protein yjeE | | Ambiguous/Unsorted |
| sbm(946546 DNA gyrase inhibitory protein | | Up Regulated |
| hflX 948688 GTP-binding protein hflX yeeD 946539 UPF0033 protein yeeD | | Ambiguous/Unsorted |
| | | Down Regulated Up Regulated |
| sodA 948403 Superoxide dismutase [Mn]; Superoxide dismutase phoE 944926 Outer membrane pore protein E | | Temporarly Down Regula |
| cpxR 948404 Transcriptional regulatory protein cpxR | | Temporarly Down Regula |
| rhaD 948401 Rhamnulose-1-phosphate aldolase | | Down Regulated |
| mutS 947206 DNA mismatch repair protein mutS | | Ambiguous/Unsorted |
| pcm 944923 Protein-L-isoaspartate O-methyltransferase | | Ambiguous/Unsorted |
| yiiQ 948408Uncharacterized protein yiiQ | п | Down Regulated |
| cpxA 948405 Sensor protein; Sensor protein cpxA | | Down Regulated |
| yiiM 948406 Protein yiiM | | Up Regulated |
| uvrC 947203 UvrABC system protein C | | Temporarly Up Regulated |
| wcaA 946570 Putative colanic acid biosynthesis glycosyl transferase wo | caA | Temporarly Down Regula |
| cpsG 946574Phosphomannomutase | | Temporarly Down Regula |
| yhaJ 947621 Uncharacterized HTH-type transcriptional regulator yhaJ | ['Posttranslational modification protein turnover chaperones'] | Down Regulated |
| yhaK 947620 Pirin-like protein yhaK | | Temporarly Down Regula |
| tdcE 947623 Keto-acid formate acetyltransferase | | Temporarly Down Regula |
| ybaL 946576 Inner membrane protein ybaL | | Ambiguous/Unsorted |
| tdcF 947624 Protein tdcF | | Up Regulated |
| metL 948433 Bifunctional aspartokinase/homoserine dehydrogenase 2 | | Down Regulated |
| rpoS 947210 RNA polymerase sigma factor rpoS; RNA polymerase sig | ma factor | Up Regulated |
| surE 947211 Multifunctional protein surE | | Ambiguous/Unsorted |
| yciS 944936 Inner membrane protein yciS | | Temporarly Down Regula |
| fsaB 948439 Fructose-6-phosphate aldolase 2 | | Ambiguous/Unsorted |
| truD 947214 tRNA pseudouridine synthase D | | Ambiguous/Unsorted |
| cysD 947217 Sulfate adenylyltransferase subunit 2 | | Down Regulated |
| metB 948434 Cystathionine gamma-synthase | | Ambiguous/Unsorted |
| metJ 948435 Met repressor | | Up Regulated |
| rsmA 944939 b0051 | | Ambiguous/Unsorted |
| cysN 947219 Sulfate adenylyltransferase subunit 1 | | Down Regulated |
| rlmF 944938 Ribosomal RNA large subunit methyltransferase F ptsA 948437 Multiphosphoryl transfer protein 2 | | Down Regulated Ambiguous/Unsorted |
| galF 946560 UTPglucose-1-phosphate uridylyltransferase | | Up Regulated |
| mdIA 945092 Multidrug resistance-like ATP-binding protein mdIA | | Down Regulated |
| muraj ==================================== | l . | Down Regulated |

| wcaL | 946565 | Putative colanic acid biosynthesis glycosyltransferase wcaL | | Down Regulated |
|------|--------|--|---|-------------------------|
| yegH | | UPF0053 protein yegH | | Temporarly Up Regulated |
| yqjG | | Uncharacterized protein yqjG | | Up Regulated |
| yqjD | | Uncharacterized protein ygiD | | Ambiguous/Unsorted |
| ycgR | | Uncharacterized protein ycgR | | Ambiguous/Unsorted |
| glpF | | Glycerol uptake facilitator protein | | Up Regulated |
| yagX | | Uncharacterized protein yagX | Carbohydrate transport and metabolism', Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcriptio | |
| zapB | | Cell division protein zapB | | Up Regulated |
| cysC | | Adenylyl-sulfate kinase | | Down Regulated |
| hslV | | ATP-dependent protease hslV | | Ambiguous/Unsorted |
| ybgF | | Uncharacterized protein ybgF | | Ambiguous/Unsorted |
| cytR | | HTH-type transcriptional repressor cytR | | Temporarly Down Regula |
| ftsN | | Cell division protein ftsN | | Down Regulated |
| maa | | Maltose O-acetyltransferase | | Down Regulated |
| rpmE | | 50S ribosomal protein L31 | | Down Regulated |
| hha | | Hemolysin expression-modulating protein | | Down Regulated |
| priA | | Primosomal protein N' | | Temporarly Up Regulated |
| glpK | | Glycerol kinase | | Up Regulated |
| glpX | | Fructose-1,6-bisphosphatase class 2 | | Down Regulated |
| smp/ | | Small protein A | | Ambiguous/Unsorted |
| bamE | | Small protein A | | Ambiguous/Unsorted |
| vacJ | | Lipoprotein vacJ | | Ambiguous/Unsorted |
| mlaA | | Lipoprotein vacJ | | Ambiguous/Unsorted |
| mpa/ | | Protein mpaA | | Ambiguous/Unsorted |
| hyaB | 945580 | Hydrogenase-1 large chain | | Down Regulated |
| udk | 946597 | Uridine kinase | | Down Regulated |
| cyoA | 945080 | Ubiquinol oxidase subunit 2 | | Temporarly Down Regula |
| vfaL | 946595 | Uncharacterized protein yfaL | | Ambiguous/Unsorted |
| mltA | 944964 | Membrane-bound lytic murein transglycosylase A | | Down Regulated |
| hypF | 944963 | Carbamoyltransferase hypF | | Up Regulated |
| dcd | 946593 | Deoxycytidine triphosphate deaminase | | Up Regulated |
| rutE | | Putative NADH dehydrogenase/NAD(P)H nitroreductase rutE | | Temporarly Up Regulated |
| gfcD | | Uncharacterized lipoprotein gfcD | | Temporarly Down Regula |
| ykgM | 944960 | 50S ribosomal protein L31 type B | | Up Regulated |
| cof | 945089 | HMP-PP phosphatase | | Temporarly Down Regula |
| mdIB | 945088 | Multidrug resistance-like ATP-binding protein mdlB | | Temporarly Up Regulated |
| ygiB | 945086 | UPF0441 protein ygiB | | Down Regulated |
| lon | 945085 | ATP-dependent protease La | | Up Regulated |
| clpX | 945083 | ATP-dependent Clp protease ATP-binding subunit clpX | | Down Regulated |
| ItaE | 944955 | Low specificity L-threonine aldolase | | Ambiguous/Unsorted |
| clpP | 945082 | ATP-dependent Clp protease proteolytic subunit | | Ambiguous/Unsorted |
| ydcU | 945976 | Inner membrane ABC transporter permease protein ydcU | | Ambiguous/Unsorted |
| yddW | 945975 | Uncharacterized lipoprotein yddW | | Up Regulated |
| ycbG | 945570 | UPF0268 protein ycbG | | Temporarly Up Regulated |
| omp/ | | Outer membrane protein A | | Up Regulated |
| cusS | | Sensor kinase cusS; Sensor protein | | Temporarly Up Regulated |
| ydbK | | Probable pyruvate-flavodoxin oxidoreductase | | Temporarly Down Regula |
| tehB | | Tellurite resistance protein tehB | | Up Regulated |
| hspQ | | Heat shock protein hspQ | | Ambiguous/Unsorted |
| yneH | | Glutaminase 2 | | Down Regulated |
| yafV | | UPF0012 hydrolase yafV | | Up Regulated |
| yahJ | | Uncharacterized protein yahJ | [Amino acid transport and metabolism', 'Inorganic ion transport and metabolism', 'Carbon compound utilization', 'Primary Active Tra | |
| yahK | 944975 | Uncharacterized zinc-type alcohol dehydrogenase-like protein y | | Up Regulated |
| | | Uncharacterized lipoprotein ybaY | | Up Regulated |
| gsk | | Inosine-guanosine kinase | | Down Regulated |
| yahL | | Uncharacterized protein yahL | | Up Regulated |
| mgsA | | Methylglyoxal synthase | | Ambiguous/Unsorted |
| scpA | | Protein sbm | | Temporarly Down Regula |
| cpsB | | Mannose-1-phosphate guanylyltransferase 1; Mannose-1-phosp | | Ambiguous/Unsorted |
| dedD | | Protein dedD | | Down Regulated |
| LeuC | | 3-isopropylmalate dehydratase large subunit | | Ambiguous/Unsorted |
| rutR | | HTH-type transcriptional regulator rutR | | Down Regulated |
| yfjF | | UPF0125 protein yfjF; Protein rnfH | | Ambiguous/Unsorted |
| crl | | Sigma factor-binding protein crl | | Down Regulated |
| puuB | 945072 | Gamma-glutamylputrescine oxidoreductase | | Down Regulated |

| omp | 944967 Outer membrane protein X | T | Up Regulated |
|------|--|--|-------------------------|
| priC | 945071 Primosomal replication protein N" | | Temporarly Down Regula |
| tesB | 945074 Acyl-CoA thioesterase 2 | | Down Regulated |
| ycjY | 945988 Uncharacterized protein ycjY | ['Amino acid transport and metabolism'] | Up Regulated |
| dmsE | 945987 Twin-arginine leader-binding protein dmsD | [Allino acid transport and metabolism] | Down Regulated |
| ydbD | 945983 Uncharacterized protein ydbD | п | Ambiguous/Unsorted |
| ftsP | 944982 Protein sufl | U . | Down Regulated |
| ykgF | 944980 Uncharacterized electron transport protein ykgF | n | Temporarly Up Regulated |
| frmR | 944986 Transcriptional repressor frmR | U . | Down Regulated |
| yahO | 944985 Uncharacterized protein yahO | [' Building block biosynthesis'] | Up Regulated |
| eutH | 944979 Ethanolamine utilization protein eutH | [Building block blosynthesis] | Up Regulated |
| yecA | 945061 Uncharacterized protein yecA | п | Down Regulated |
| dxs | 945060 1-deoxy-D-xylulose-5-phosphate synthase | U . | Ambiguous/Unsorted |
| thil | 945067 Thiamine biosynthesis protein thil | | Ambiguous/Unsorted |
| ydcj | 945991 Uncharacterized protein vdcJ | | Up Regulated |
| panE | 945065 2-dehydropantoate 2-reductase | | Temporarly Down Regula |
| ispA | 945064 Geranyltranstransferase | | Ambiguous/Unsorted |
| xseB | 945069 Exodeoxyribonuclease 7 small subunit | | Ambiguous/Unsorted |
| ydcP | 945993 Uncharacterized protease ydcP | ['amino acid biosynthetic process'] | Temporarly Up Regulate |
| narZ | 945999 Respiratory nitrate reductase 2 alpha chain | [animo dola biodynthicute process] | Ambiguous/Unsorted |
| ydcH | 945997 Uncharacterized protein ydcH | [' Channel type Transporters'] | Up Regulated |
| cspG | 945591 Cold shock-like protein cspG | [Shames type Transporters] | Down Regulated |
| yccF | 945594 Inner membrane protein yccF | | Down Regulated |
| prpB | 944990 Methylisocitrate lyase | | Temporarly Up Regulated |
| torS | 945595 Sensor protein torS; Sensor protein | | Down Regulated |
| codA | 944996 Cytosine deaminase; Protein ccdA | | Down Regulated |
| frmA | 944988 S-(hydroxymethyl)glutathione dehydrogenase | | Temporarly Down Regula |
| pyrH | 944989 Uridylate kinase | | Ambiguous/Unsorted |
| yajD | 945051 Uncharacterized protein yajD | [' Primary Active Transporters'] | Down Regulated |
| yajR | 945058 Inner membrane transport protein yajR | [Timaly Active Transporters] | Temporarly Down Regula |
| ispF | 945057 2-C-methyl-D-erythritol 2,4-cyclodiphosphate synthase | | Ambiguous/Unsorted |
| McrB | 949122 5-methylcytosine-specific restriction enzyme B | | Temporarly Up Regulate |
| secF | 949120 Protein-export membrane protein secF | | Temporarly Down Regula |
| cutA | 948660 Divalent-cation tolerance protein cutA | | Up Regulated |
| yjeK | 948662 Uncharacterized kamA family protein yjeK | ['metabolic process'] | Down Regulated |
| efp | 948661 Elongation factor P | [metabolio process] | Ambiguous/Unsorted |
| yjel | 948664 Uncharacterized protein yjel | ['cell cycle', 'cell division'] | Ambiguous/Unsorted |
| frdB | 948666 Fumarate reductase iron-sulfur subunit | [con cycle; con division] | Up Regulated |
| groL | 948665 60 kDa chaperonin | | Temporarly Up Regulated |
| ribF | 949129 Riboflavin biosynthesis protein ribF | | Ambiguous/Unsorted |
| frdA | 948667 Fumarate reductase flavoprotein subunit | | Up Regulated |
| amp | 948669 Beta-lactamase | | Down Regulated |
| lipB | 945217 Octanoyltransferase | | Down Regulated |
| entH | 945215 Esterase ybdB | | Up Regulated |
| ybdH | 945212 Uncharacterized oxidoreductase ybdH | П | Down Regulated |
| ybdL | 945211 Aminotransferase ybdL | | Down Regulated |
| blc | 948670 Outer membrane lipoprotein blc | | Up Regulated |
| orn | 948675 Oligoribonuclease | | Ambiguous/Unsorted |
| rcsF | 949113 Protein rcsF | | Ambiguous/Unsorted |
| yaeB | 949112 UPF0066 protein yaeB | | Ambiguous/Unsorted |
| psd | 948673 Phosphatidylserine decarboxylase proenzyme | | Ambiguous/Unsorted |
| poxA | 948672 Putative lysyl-tRNA synthetase | | Ambiguous/Unsorted |
| yjeM | 948679 Inner membrane transporter yieM | | Up Regulated |
| yjeP | 948676 Uncharacterized mscS family protein yjeP | ['purine nucleotide biosynthetic process', 'biosynthetic process'] | Ambiguous/Unsorted |
| lipA | 945227 Lipoyl synthase | , | Ambiguous/Unsorted |
| plsC | 947496 1-acyl-sn-glycerol-3-phosphate acyltransferase | | Temporarly Down Regula |
| uspG | 945229 Universal stress protein G | | Up Regulated |
| dkgA | 947495 2,5-diketo-D-gluconic acid reductase A | | Up Regulated |
| tatE | 945228 Sec-independent protein translocase protein tatE | | Ambiguous/Unsorted |
| tdcA | 947494 HTH-type transcriptional regulator tdcA | | Up Regulated |
| yqhD | 947493 Alcohol dehydrogenase yghD | | Up Regulated |
| dacA | 945222 D-alanyl-D-alanine carboxypeptidase dacA | | Down Regulated |
| ahpC | 945225 Alkyl hydroperoxide reductase subunit C | | Down Regulated |
| dsbG | 945224 Thiol:disulfide interchange protein dsbG | | Down Regulated |
| folD | 945221 Bifunctional protein folD | | Up Regulated |
| | , | 1 | 124 - 123 - 104 |

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|-------|---|---|-------------------------|
| ybeD | 945220 UPF0250 protein ybeD | | Down Regulated |
| parC | 947499 DNA topoisomerase 4 subunit A | | Ambiguous/Unsorted |
| dsdX | 949103 DsdX permease | | Temporarly Down Regula |
| dcuB | 948641 Anaerobic C4-dicarboxylate transporter dcuB | | Ambiguous/Unsorted |
| dsbD | 948649 Thiol:disulfide interchange protein dsbD | | Temporarly Down Regula |
| cadA | 948643 Lysine decarboxylase, inducible | | Up Regulated |
| nanM | 949106 N-acetylneuraminate epimerase | | Up Regulated |
| lysU | 948645 Lysyl-tRNA synthetase; Lysyl-tRNA synthetase, heat inducible | | Down Regulated |
| gpr | 947480 Uncharacterized protein yghZ | | Up Regulated |
| yghZ | 947480 Uncharacterized protein yghZ | | Up Regulated |
| dpiB | 945233 Sensor kinase dpiB | | Temporarly Up Regulated |
| rfbC | 947482 dTDP-4-dehydrorhamnose 3,5-epimerase | | Down Regulated |
| rlmH | 945239 Ribosomal RNA large subunit methyltransferase H | | Up Regulated |
| ybeB | 945237 Uncharacterized protein ybeB | | Up Regulated |
| citC | 945231 [Citrate [pro-3S]-lyase] ligase | | Ambiguous/Unsorted |
| fadR | 948652 Fatty acid metabolism regulator protein | | Down Regulated |
| yjdC | 948650 HTH-type transcriptional regulator yjdC | | Up Regulated |
| aspA | 948658 Aspartate ammonia-lyase | | Up Regulated |
| fxsA | 948657 UPF0716 protein fxsA | | Up Regulated |
| yghU | 947472 Uncharacterized GST-like protein yghU | | Ambiguous/Unsorted |
| holE | 947471 DNA polymerase III subunit theta | | Temporarly Down Regula |
| nadD | 945248 Nicotinate-nucleotide adenylyltransferase; Probable nicotinate-r | nucleotide adenylyltransferase | Ambiguous/Unsorted |
| gss | 947474 Bifunctional glutathionylspermidine synthetase/amidase | | Down Regulated |
| gsp | 947474 Bifunctional glutathionylspermidine synthetase/amidase | | Down Regulated |
| glcG | 947473 Protein glcG | | Up Regulated |
| yqhA | 947479 UPF0114 protein yghA | | Up Regulated |
| yghA | 947478 Uncharacterized oxidoreductase yghA | | Up Regulated |
| rlpA | 945241 Rare lipoprotein A | U | Temporarly Down Regula |
| mrdA | 945240 Penicillin-binding protein 2 | | Down Regulated |
| yjdM | 948621 Protein phnA | | Up Regulated |
| yjdN | 948622 Protein phnB | | Up Regulated |
| proP | 948626 Proline/betaine transporter | | Down Regulated |
| vbeL | 945252 Uncharacterized protein ybeL | n | Up Regulated |
| fhuC | 945250 Iron(3+)-hydroxamate import ATP-binding protein fhuC | | Ambiguous/Unsorted |
| slp | 948022 Outer membrane protein slp | | Up Regulated |
| serA | 945258 D-3-phosphoglycerate dehydrogenase | | Ambiguous/Unsorted |
| | 945256 PTS system mannitol-specific cryptic EIICB component | | |
| cmtA | | | Ambiguous/Unsorted |
| adiA | 948638 Biodegradative arginine decarboxylase | | Ambiguous/Unsorted |
| melR | 948637 Melibiose operon regulatory protein | | Ambiguous/Unsorted |
| dctA | 948039 C4-dicarboxylate transport protein; Aerobic C4-dicarboxylate tra | insport protein | Up Regulated |
| miaB | 945260 (Dimethylallyl)adenosine tRNA methylthiotransferase miaB | | Down Regulated |
| ubif | 945261 2-octaprenyl-3-methyl-6-methoxy-1,4-benzoquinol hydroxylase | | Down Regulated |
| yhjE | 948032 Inner membrane metabolite transport protein yhjE | | Down Regulated |
| yhjD | 948033 Inner membrane protein yhjD | | Ambiguous/Unsorted |
| treF | 948037 Cytoplasmic trehalase | | Up Regulated |
| yhjC | 948035 Uncharacterized HTH-type transcriptional regulator yhjC | [Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent'] | Ambiguous/Unsorted |
| rpiR | 948603 HTH-type transcriptional regulator rpiR | | Down Regulated |
| yecJ | 946388 Uncharacterized protein yecJ | <u>[D</u> | Ambiguous/Unsorted |
| torZ | 946389 Trimethylamine-N-oxide reductase 2 | | Temporarly Up Regulated |
| phnJ | 948606 Protein phnJ | | Ambiguous/Unsorted |
| phnP | 948600 Protein phnP | | Ambiguous/Unsorted |
| rpiB | 948602 Ribose-5-phosphate isomerase B | | Down Regulated |
| cmo/ | 946380 tRNA (cmo5U34)-methyltransferase | | Ambiguous/Unsorted |
| yecR | 946386 Uncharacterized protein yecR | ['Inorganic ion transport and metabolism', ' Adaptation to stress', 'iron ion transport'] | Down Regulated |
| cmoE | | | Down Regulated |
| yecD | 946384 Isochorismatase family protein yecD; Uncharacterized isochoris | 10 | Temporarly Up Regulated |
| pitA | 948009 Low-affinity inorganic phosphate transporter 1 | | Down Regulated |
| seqA | 945272 Protein segA | | Ambiguous/Unsorted |
| yhiQ | 948005 UPF0341 protein yhiQ | | Ambiguous/Unsorted |
| rfbB | 945276 dTDP-glucose 4,6-dehydratase | | Down Regulated |
| erfK | 945273 Protein erfK/srfK | | Ambiguous/Unsorted |
| uspA | | | Ambiguous/Unsorted |
| iscR | 945279 HTH-type transcriptional regulator iscR | | Down Regulated |
| yhil | 948003 Uncharacterized protein yhil | [' Protection'] | Temporarly Up Regulated |
| trmJ | 948610 tRNA (cytidine/uridine-2'-O-)-methyltransferase trmJ | [| Down Regulated |
| uiiio | OTOO TO LE TAN (CYTICHIE ALICHE TO -)-HIEUTYILI ALISIELASE LITTIJ | I . | Powii ivegulated |

| flhB | 046204 Elegallar biocypthetic protein flbD | | Down Bogulated |
|--------------|---|---|-----------------------------|
| - | 946391 Flagellar biosynthetic protein flhB 946392 Chemotaxis protein cheZ | | Down Regulated |
| cheZ | | | Up Regulated |
| cheB | 946394 Chemotaxis response regulator protein-glutamate methylestera 946396 Chemotaxis protein methyltransferase | SE | Down Regulated |
| cheR | | | Temporarly Up Regulated |
| tap | 946397 Methyl-accepting chemotaxis protein IV | | Ambiguous/Unsorted |
| mprA | 945282 Transcriptional repressor mprA | | Ambiguous/Unsorted |
| arsC | 948018 Arsenate reductase | | Temporarly Up Regulated |
| yhiD | 948019 Uncharacterized protein yhiD | Ц | Ambiguous/Unsorted |
| nagC | 945285 N-acetylglucosamine repressor | | Down Regulated |
| flhA | 946390 Flagellar biosynthesis protein flhA | | Down Regulated |
| yhiR | 948012 Uncharacterized protein yhiR | Ц | Ambiguous/Unsorted |
| phoH | 948010 Protein phoH | | Temporarly Up Regulated |
| dut | 948607 Deoxyuridine 5'-triphosphate nucleotidohydrolase | | Ambiguous/Unsorted |
| dppD | 948065 Dipeptide transport ATP-binding protein dppD | | Down Regulated |
| dppB | 948063 Dipeptide transport system permease protein dppB | | Ambiguous/Unsorted |
| eptB | 948068 b3546; ECS88_3965; ECIAI39_4054; EC55989_3997; E2348C | _3793; ECED1_4224; ECUMN_4058; ECDH10B_3725; ECIAI1_3705 | Temporarly Down Regula |
| gabT | 948067 4-aminobutyrate aminotransferase | | Up Regulated |
| yfeD | 946865 Uncharacterized protein yfeD | IU | Up Regulated |
| yejO | | ; APECO1_4367; UTI89_C2469; EC55989_2444; EC55989_2445; ECUMN_2527; ECIAI1_2272; ECIAI1_2273 | Ambiguous/Unsorted |
| yfeA | 946864 Uncharacterized protein yfeA | [Signal transduction mechanisms', 'Type of regulation', 'two component signal transduction system phosphorelay', 'regulation of t | |
| gabD | 948060 Succinate-semialdehyde dehydrogenase [NADP+] | | Up Regulated |
| yiaD | 948075 Inner membrane lipoprotein yiaD | | Up Regulated |
| csiD | 948076 Protein csiD | | Ambiguous/Unsorted |
| yfeK | 946876 Uncharacterized protein yfeK | ['Defense mechanisms', ' Primary Active Transporters', ' Protection'] | Down Regulated |
| cspA | 948070 Cold shock protein cspA | | Down Regulated |
| yiaF | 948072 Uncharacterized protein yiaF | ['Carbohydrate transport and metabolism', ' Carbon compound utilization', 'phosphoenolpyruvate dependent sugar phosphotransfer | |
| yhjK | | AI1_3680; b3529; ECIAI39_4032; ECED1_4207; ECS88_3943; ECUMN_4036; UTI89_C4061; Z4943; E2348C_3771 | Temporarly Up Regulated |
| bcsZ | 948046 Endoglucanase | | Up Regulated |
| bcsB | 948045 Cyclic di-GMP-binding protein | | Ambiguous/Unsorted |
| crr | 946880 Glucose-specific phosphotransferase enzyme IIA component | | Up Regulated |
| yhjG | 948043 Uncharacterized protein yhjG | ['Lipid transport and metabolism'] | Temporarly Down Regula |
| pdxK | 946881 Pyridoxine kinase | | Down Regulated |
| kdgK | 948041 2-dehydro-3-deoxygluconokinase | | Ambiguous/Unsorted |
| ptsH | 946886 Phosphocarrier protein HPr | | Temporarly Down Regula |
| yfeS | 946887 Uncharacterized protein yfeS | | Ambiguous/Unsorted |
| cysM | 946888 Cysteine synthase B; Cysteine synthase | | Down Regulated |
| cysA | 946889 Sulfate/thiosulfate import ATP-binding protein cysA | | Down Regulated |
| yhjJ | 948040 Protein yhjJ | | Ambiguous/Unsorted |
| dppF | 948056 Dipeptide transport ATP-binding protein dppF | | Down Regulated |
| bcsG | 948058 Uncharacterized protein yhjU | | Temporarly Up Regulated |
| murQ | 946893 N-acetylmuramic acid 6-phosphate etherase | | Ambiguous/Unsorted |
| csiR | 948055 Uncharacterized HTH-type transcriptional regulator ygaE | | Temporarly Up Regulated |
| nupC | 946895 Nucleoside permease nupC | | Down Regulated |
| ucpA | 946898 Oxidoreductase ucpA | | Up Regulated |
| bcsE | 948050 Uncharacterized protein yhjS | | Ambiguous/Unsorted |
| fadL | 946820 Long-chain fatty acid transport protein | | Up Regulated |
| sixA | 946815 Phosphohistidine phosphatase sixA | | Down Regulated |
| fecD | 946816 Fe(3+) dicitrate transport system permease protein fecD | | Down Regulated |
| emrY | 946835 Multidrug resistance protein Y | | Down Regulated |
| efeO | 945603 UPF0409 protein ycdO | | Up Regulated |
| putP | 945602 Sodium/proline symporter | | Ambiguous/Unsorted |
| yceA | 945601 UPF0176 protein yceA | | Down Regulated |
| putA | 945600 Bifunctional protein putA | | Down Regulated |
| frc | 946842 Formyl-coenzyme A transferase | | Down Regulated |
| evgS | 946844 Sensor protein evgS; Sensor protein | | Ambiguous/Unsorted |
| malS | 948088 Alpha-amylase | | Down Regulated |
| xylR | 948086 Xylose operon regulatory protein | | Ambiguous/Unsorted |
| yiaJ | 948084 HTH-type transcriptional regulator yiaJ | | Down Regulated |
| dsdA | 946837 D-serine dehydratase; D-serine dehydratase 1 | | Up Regulated |
| fecB | 946838 Fe(3+) dicitrate-binding periplasmic protein | | Temporarly Up Regulated |
| fect | 946839 Probable RNA polymerase sigma factor fecl; RNA polymerase s | l signed factor | Ambiguous/Unsorted |
| | 946852 Multiphosphoryl transfer protein 1 | signia racion | Up Regulated |
| frya yfdZ | 946850 Uncharacterized aminotransferase yfdZ | | |
| | 946857 Uncharacterized aminotransferase ytdz | <u>U</u> | Down Regulated Up Regulated |
| yfeC | 945201 Ferric enterobactin transport ATP-binding protein fepC | U . | |
| fepC | 94920 Iperiic enteropactin transport ATP-binding protein iepC | I . | Up Regulated |

| L(D) | 0.45007 | 10:1 (15:1) | | A 1: (1) |
|------|---------|--|--|-------------------------|
| stfR | | Side tail fiber protein homolog from lambdoid prophage Rac | | Ambiguous/Unsorted |
| yceM | | Virulence factor mviM homolog | | Temporarly Down Regula |
| xylF | | D-xylose-binding periplasmic protein | | Up Regulated |
| lpcA | | Phosphoheptose isomerase | | Ambiguous/Unsorted |
| lpxP | | Protein ddg | | Up Regulated |
| secD | | Protein-export membrane protein secD | | Ambiguous/Unsorted |
| tgt | | Queuine tRNA-ribosyltransferase | | Ambiguous/Unsorted |
| malZ | | Maltodextrin glucosidase | | Ambiguous/Unsorted |
| leuD | | 3-isopropylmalate dehydratase small subunit | | Ambiguous/Unsorted |
| flgL | | Flagellar hook-associated protein 3 | | Down Regulated |
| flgG | | Flagellar basal-body rod protein flgG | | Down Regulated |
| ftnA | | Ferritin-1 | | Temporarly Down Regula |
| yecM | 946415 | Protein yecM | | Temporarly Up Regulated |
| ybeX | | Magnesium and cobalt efflux protein corC | | Down Regulated |
| ftnB | 946407 | Ferritin-like protein 2 | | Ambiguous/Unsorted |
| araG | 946408 | Arabinose import ATP-binding protein araG | | Ambiguous/Unsorted |
| araF | 946409 | L-arabinose-binding periplasmic protein | | Up Regulated |
| rluC | | Pseudouridine synthase; Ribosomal large subunit pseudouridin | e synthase C | Ambiguous/Unsorted |
| ydcS | 946005 | Putative ABC transporter periplasmic-binding protein ydcS | | Up Regulated |
| flgF | | Flagellar basal-body rod protein flgF | | Down Regulated |
| vncE | | Uncharacterized protein yncE | ['Translation ribosomal structure and biogenesis', 'rRNA processing'] | Up Regulated |
| vdcT | | Uncharacterized ABC transporter ATP-binding protein ydcT | [| Temporarly Up Regulated |
| yceF | | Maf-like protein yceF 1; Maf-like protein yceF | | Ambiguous/Unsorted |
| yacG | | UPF0243 zinc-binding protein yacG | | Up Regulated |
| flgE | | Flagellar hook protein figE | | Down Regulated |
| | | UPF0502 protein yceH | | Up Regulated |
| yceH | | | | |
| flgN | | Flagella synthesis protein flgN | | Ambiguous/Unsorted |
| otsA | | Alpha,alpha-trehalose-phosphate synthase [UDP-forming] | | Up Regulated |
| uspC | | Universal stress protein C | | Up Regulated |
| yncB | 946012 | Putative NADP-dependent oxidoreductase yncB | | Up Regulated |
| ycjG | 946013 | Z2450; ECUMN_1620; ECED1_1533; ECDH10B_1444; b1325 | ; UTI89_C1596; E2348C_1518; ECIAI1_1350; c1797; APECO1_478; ECS88_1467; ECIAI39_1677; EC55989_1489 | Up Regulated |
| araB | | Ribulokinase | | Up Regulated |
| yncD | | Probable tonB-dependent receptor yncD | | Down Regulated |
| aroP | | Aromatic amino acid transport protein aroP | | Temporarly Down Regula |
| ansP | | L-asparagine permease | | Temporarly Down Regula |
| yfcM | | Uncharacterized protein yfcM | ['Cell wall membrane envelope biogenesis'] | Down Regulated |
| opgH | | Glucans biosynthesis glucosyltransferase H | | Ambiguous/Unsorted |
| yaeQ | | Uncharacterized protein yaeQ | ['Inorganic ion transport and metabolism', 'DNA repair', 'DNA recombination', 'response to DNA damage stimulus'] | Down Regulated |
| yfcL | 946804 | Uncharacterized protein yfcL | ['Cell wall membrane envelope biogenesis', 'amino acid biosynthetic process', 'aromatic amino acid family biosynthetic process'] | Down Regulated |
| ylbG | 946803 | b0502; ECDH10B_0458 | | Up Regulated |
| ribD | 945620 | Riboflavin biosynthesis protein ribD | | Up Regulated |
| glnP | 945621 | Glutamine transport system permease protein glnP | | Down Regulated |
| mep/ | | Penicillin-insensitive murein endopeptidase | | Down Regulated |
| суоВ | | Ubiquinol oxidase subunit 1 | | Temporarly Down Regula |
| rplY | | 50S ribosomal protein L25 | | Down Regulated |
| yfjG | | UPF0083 protein yfjG | | Down Regulated |
| yhhJ | | Inner membrane transport permease yhhJ | | Temporarly Down Regula |
| msyB | | Acidic protein msyB | | Up Regulated |
| rbbA | | Uncharacterized ABC transporter ATP-binding protein yhiH | | Down Regulated |
| rhsB | | Protein rhsB | | Down Regulated |
| | | | 1 | |
| nikR | | Nickel-responsive regulator | n n | Temporarly Down Regula |
| yhiN | | Uncharacterized protein yhiN | Ψ | Up Regulated |
| slyB | | Outer membrane lipoprotein slyB | | Ambiguous/Unsorted |
| mnm | | tRNA 5-methylaminomethyl-2-thiouridine biosynthesis bifunction | nai protein mnmC | Temporarly Up Regulated |
| mfd | | Transcription-repair-coupling factor | | Ambiguous/Unsorted |
| pspF | | Psp operon transcriptional activator | | Temporarly Up Regulated |
| potD | | Spermidine/putrescine-binding periplasmic protein | | Up Regulated |
| cobB | | NAD-dependent deacetylase | | Up Regulated |
| panD | | Aspartate 1-decarboxylase | | Down Regulated |
| nudJ | | Phosphatase nudJ | | Down Regulated |
| aldA | 945672 | Lactaldehyde dehydrogenase | | Up Regulated |
| nagZ | 945671 | Beta-hexosaminidase | | Temporarly Up Regulated |
| IoID | | Lipoprotein-releasing system ATP-binding protein IoID | | Ambiguous/Unsorted |
| | | UPF0227 protein ycfP | | Ambiguous/Unsorted |
| ycfP | 943070 | | | |
| | | Uncharacterized protein yeaO | l' Primary Active Transporters', 'electron transport'] | Up Regulated |

| IniC | 045672 Linearotain releasing avetem transmembrane protein IolC | | Ambiguous/Unsorted |
|------|---|--|-------------------------|
| lolC | 945673 Lipoprotein-releasing system transmembrane protein IoIC | | Ŭ |
| flgB | 945678 Flagellar basal-body rod protein flgB | | Down Regulated |
| yagU | 945677 Inner membrane protein yagU | | Ambiguous/Unsorted |
| parE | 947501 DNA topoisomerase 4 subunit B | | Down Regulated |
| ybiX | 947502 PKHD-type hydroxylase ybiX | | Up Regulated |
| ygiN | 947506 Probable quinol monooxygenase ygiN | | Up Regulated |
| holB | 945661 DNA polymerase III subunit delta' | | Ambiguous/Unsorted |
| yceG | 945660 Uncharacterized protein yceG | Д | Temporarly Up Regulated |
| pck | 945667 Phosphoenolpyruvate carboxykinase [ATP] | | Up Regulated |
| ycfS | 945666 Uncharacterized protein ycfS | Ц | Down Regulated |
| cspD | 945669 Cold shock-like protein cspD | | Temporarly Up Regulated |
| tmk | 945663 Thymidylate kinase | | Down Regulated |
| IoIE | 945665 Lipoprotein-releasing system transmembrane protein IolE | | Ambiguous/Unsorted |
| | 945657 50S ribosomal protein L32 | - | Down Regulated |
| ycfH | 945656 Uncharacterized deoxyribonuclease ycfH | | Down Regulated |
| emtA | 945655 Endo-type membrane-bound lytic murein transglycosylase A | | Down Regulated |
| deoD | 945654 Purine nucleoside phosphorylase deoD-type | | Up Regulated |
| pqiB | 945653 Paraquat-inducible protein B | | Temporarly Down Regula |
| pspE | 945652 Phage shock protein E | | Temporarly Up Regulated |
| flgK | 945648 Flagellar hook-associated protein 1 | | Down Regulated |
| fhuE | 945649 FhuE receptor | | Temporarly Up Regulated |
| gntK | 947937 Thermoresistant gluconokinase | | Temporarly Up Regulated |
| glpG | 947936 Rhomboid protease glpG | | Ambiguous/Unsorted |
| cusB | 945189 Cation efflux system protein cusB | | Down Regulated |
| glgS | 947533 Glycogen synthesis protein glgS | | Ambiguous/Unsorted |
| entF | 945184 Enterobactin synthetase component F | | Up Regulated |
| heml | 947532 Ferrochelatase | | Up Regulated |
| fes | 945181 Enterochelin esterase | | Ambiguous/Unsorted |
| flgl | 947534 Flagellar P-ring protein | | Down Regulated |
| cusF | 945188 Cation efflux system protein cusF | | Down Regulated |
| ompT | 945185 Protease 7 | | Up Regulated |
| fepB | 947538 Ferrienterobactin-binding periplasmic protein | | Up Regulated |
| veeN | 946493 UPF0082 protein yeeN | | Down Regulated |
| fepE | 945180 Ferric enterobactin transport protein fepE | | Temporarly Up Regulated |
| shiA | 946495 Shikimate transporter | | Temporarly Up Regulated |
| glgB | 947940 1,4-alpha-glucan-branching enzyme | | Ambiguous/Unsorted |
| ydhJ | 946499 Uncharacterized protein ydhJ | | Ambiguous/Unsorted |
| glgX | 947941 Glycogen debranching enzyme | | Temporarly Up Regulated |
| yeeJ | 946498 Uncharacterized protein yeeJ | | Temporarly Up Regulated |
| | 947942 Glucose-1-phosphate adenylyltransferase | | Up Regulated |
| glgC | 947944 Uncharacterized oxidoreductase yhhX | | Ambiguous/Unsorted |
| yhhX | | ll la l | |
| yhhW | 947945 Protein yhhW | | Ambiguous/Unsorted |
| gntR | 947946 HTH-type transcriptional regulator gntR | | Ambiguous/Unsorted |
| rtcR | 947928 Transcriptional regulatory protein rtcR | | Ambiguous/Unsorted |
| glpR | 947926 Glycerol-3-phosphate regulon repressor | | Ambiguous/Unsorted |
| ahpF | 947540 Alkyl hydroperoxide reductase subunit F | | Ambiguous/Unsorted |
| rtcB | 947929 Protein rtcB | | Temporarly Up Regulated |
| rnk | 947546 Regulator of nucleoside diphosphate kinase | | Ambiguous/Unsorted |
| folB | 947544 Dihydroneopterin aldolase | I . | Ambiguous/Unsorted |
| quuD | 945177 Antitermination protein Q homolog from lambdoid prophage DLF | 712 | Temporarly Down Regula |
| yodA | 946480 Metal-binding protein yodA | | Up Regulated |
| zinT | 946480 Metal-binding protein yodA | | Up Regulated |
| ydiQ | 946482 Putative electron transfer flavoprotein subunit ydiQ | | Temporarly Up Regulated |
| | 946484 UPF0190 protein yedY | | Down Regulated |
| hiuH | 946485 5-hydroxyisourate hydrolase | | Ambiguous/Unsorted |
| dam | 947930 Protein damX | | Ambiguous/Unsorted |
| yedV | 946487 Sensor protein; Putative sensor-like histidine kinase yedV | | Ambiguous/Unsorted |
| glgP | 947931 Phosphorylase; Glycogen phosphorylase | | Ambiguous/Unsorted |
| mtfA | 946489 Protein mtfA | | Up Regulated |
| glpD | | | Up Regulated |
| | 947935 Thiosulfate sulfurtransferase glpE | | Down Regulated |
| glgA | 947932 Glycogen synthase | | Ambiguous/Unsorted |
| feoB | 947919 Ferrous iron transport protein B | | Temporarly Up Regulated |
| | 947515 Protein icc | | Down Regulated |
| | 947511 L-arabinose isomerase | | Up Regulated |
| | 1 | 1 | |

| fhuD | 947510 | Iron(3+)-hydroxamate-binding protein fhuD | | Up Regulated |
|--|---|---|--|--|
| ykgH | | Uncharacterized protein ykgH | | Ambiguous/Unsorted |
| vafC | | Uncharacterized HTH-type transcriptional regulator yafC | "Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent'] | Down Regulated |
| malT | | HTH-type transcriptional regulator malT | , , , , , , , , , , , , , , , , , , , | Down Regulated |
| malQ | | 4-alpha-glucanotransferase | | Temporarly Down Regula |
| gntT | | High-affinity gluconate transporter | | Down Regulated |
| lptA | | Lipopolysaccharide export system protein lptA | | Down Regulated |
| feoA | | Ferrous iron transport protein A | | Ambiguous/Unsorted |
| mrcA | 947907 | Penicillin-binding protein 1A | | Down Regulated |
| nudE | 947906 | ADP compounds hydrolase nudE | | Ambiguous/Unsorted |
| yrfF | 947905 | Putative membrane protein igaA homolog | | Down Regulated |
| yrfG | | Uncharacterized protein yrfG | [' Adaptation to stress', 'protein folding'] | Down Regulated |
| elbB | 947903 | Enhancing lycopene biosynthesis protein 2 | | Ambiguous/Unsorted |
| mhp/ | 945197 | 3-(3-hydroxy-phenyl)propionate/3-hydroxycinnamic acid hydroxy | ylase | Ambiguous/Unsorted |
| prpC | 947528 | 2-methylcitrate synthase | | Ambiguous/Unsorted |
| ygcF | | Uncharacterized protein ygcF | | Down Regulated |
| iscX | | Protein iscX | | Down Regulated |
| ribB | | 3,4-dihydroxy-2-butanone 4-phosphate synthase | | Ambiguous/Unsorted |
| pheP | | Phenylalanine-specific permease | | Down Regulated |
| atoD | | Acetate CoA-transferase subunit alpha | | Up Regulated |
| thrC | | Threonine synthase | | Down Regulated |
| fepA | 945193 | Ferrienterobactin receptor | | Up Regulated |
| yqiC | | Uncharacterized protein yqiC | 0 | Up Regulated |
| ygiL | | Uncharacterized fimbrial-like protein ygiL | [Replication recombination and repair'] | Ambiguous/Unsorted |
| rpmA | | 50S ribosomal protein L27 | | Down Regulated |
| cusA | | Cation efflux system protein cusA | | Temporarly Up Regulated |
| ycjN | | Putative ABC transporter periplasmic-binding protein ycjN | | Ambiguous/Unsorted |
| yqiB | | Uncharacterized protein yqiB | | Down Regulated |
| nudF | | ADP-ribose pyrophosphatase | | Down Regulated |
| hslR | | Heat shock protein 15 | | Ambiguous/Unsorted |
| potB | | Spermidine/putrescine transport system permease protein potB | | Down Regulated Ambiguous/Unsorted |
| ompF | | Transcriptional regulatory protein ompR Uncharacterized protein yrbL | ['Carbohydrate transport and metabolism', ' Macromolecules cellular constituent biosynthesis', ' Central intermediary metabolism'] | Down Regulated |
| yrbL hflD | | Uncharacterized protein yrbL | | Down Regulated |
| | 045603 | ECIAI1 1170: ECS08 1147: ECDU10B 1204: 61122: ECIAI20 | 2005: ECED1 1276: E2249C 1272: ECHMN 1276: EC55090 1245 | |
| | | | _2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regula |
| yhgF | 947911 | Protein yhgF | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulation Down Regulated |
| yhgF rpsU | 947911 947577 | Protein yhgF 30S ribosomal protein S21 | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regula Down Regulated Down Regulated |
| yhgF rpsU dnaG | 947911 947577 947570 | Protein yhgF 30S ribosomal protein S21 DNA primase | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regula Down Regulated Down Regulated Up Regulated |
| yhgF rpsU dnaG ypfH | 947911 947577 947570 947571 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regula Down Regulated Down Regulated Up Regulated Up Regulated |
| yhgF rpsU dnaG ypfH cbpA | 947911 947577 947570 947571 947572 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated |
| yhgF rpsU dnaG ypfH cbpA holA | 947911 947577 947570 947571 947572 947573 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Down Regulated |
| yhgF rpsU dnaG ypfH cbpA hoIA btuF | 947911 947577 947570 947571 947572 947573 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted |
| yhgF rpsU dnaG ypfH cbpA holA | 947911 947577 947570 947571 947572 947573 947574 947983 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Down Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB | 947911 947577 947570 947571 947572 947573 947574 947983 946459 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD | 947911 947577 947570 947571 947572 947573 947574 947983 946459 947989 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted |
| yhgF rpsU dnaG ypfH cbpA hoIA btuF dcrB thiD nikD | 947911 947577 947570 947571 947572 947573 947574 947983 946459 947989 948595 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted |
| yhgF rpsU dnaG ypfH cbpA hoIA btuF dcrB thiD nikD alsE | 947911 947577 947570 947571 947572 947573 947574 947983 946459 948595 948595 948597 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated Ambiguous/Unsorted Down Regulated Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated |
| yhgF rpsU dnaG ypfH cbpA hoIA btuF dcrB thiD nikD alsE yjcS | 947911 947577 947570 947571 947572 947573 947574 947598 946459 947989 948595 948597 948591 946452 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD nikD alsE yjcS gltP | 947911 947577 947570 947571 947572 947573 947574 947983 946459 947899 948595 948595 948591 946452 946452 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD nikD alsE yjcS gltP argS | 947911 947577 947570 947571 947572 947573 947574 947983 946459 948595 948597 948597 948593 946452 946453 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase Ribosomal RNA large subunit methyltransferase G | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD nikD alsE yjcS gltP argS ribE | 947911 947577 947570 947571 947572 947573 947574 947983 946459 947989 948595 948597 948591 946452 946452 946452 947589 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase Ribosomal RNA large subunit methyltransferase G Protein gutQ | | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Up Regulated Down Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD nikD alsE yjcS gltP argS ribE rlmG gutQ yqjH | 947911 947577 947570 947571 947572 947573 947574 947983 947989 947989 948595 948595 948597 946452 946452 946453 947589 947587 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dorB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase Ribosomal RNA large subunit methyltransferase G Protein gutQ Uncharacterized protein yqjH | 2005; ECED1_1276; E2348C_1273; ECUMN_1376; EC55989_1245 [] [] [] [] [] [] [] [] [] [] [] [] [] | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Up Regulated Down Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Up Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD nikD alsE yjcS gltP argS ribE rlmG gutQ yqjH ftsX | 947911 947577 947570 947571 947572 947573 947574 947983 946459 947899 948595 948597 946452 946453 947589 947582 947582 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yicS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase Ribosomal RNA large subunit methyltransferase G Protein gutQ Uncharacterized protein ygjH Cell division protein ftsX | [] [] [] [] [] [] Adaptation to stress', 'ion transport', 'iron ion transport'] | Temporarly Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Up Regulated Down Regulated Down Regulated Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Up Regulated Ambiguous/Unsorted |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD nikD alsE yjcS gltP argS ribE rlmG gutQ yqjH ftsX | 947911 947577 947577 947570 947571 947572 947573 947574 94783 946459 948595 948597 948597 948597 947587 947587 947587 947582 947582 947582 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase Ribosomal RNA large subunit methyltransferase G Protein gutQ Uncharacterized protein ygjH Cell division protein ftsX NADH-quinone oxidoreductase subunit K | [] [] [] [] [] [] Adaptation to stress', 'ion transport', 'iron ion transport'] | Temporarly Down Regulated Down Regulated Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Up Regulated Down Regulated Up Regulated Down Regulated Down Regulated Down Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD alsE yjcS gltP argS ribE rlmG gutQ yqjH ftsX nuoK ebgR | 947911 947577 947570 947571 947572 947573 947574 947983 946459 947889 948591 946452 947689 947582 947582 947582 947580 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase Ribosomal RNA large subunit methyltransferase G Protein gutQ Uncharacterized protein ygjH Cell division protein ftsX NADH-quinone oxidoreductase subunit K HTH-type transcriptional regulator ebgR | [] [] [] [] [] [] Adaptation to stress', 'ion transport', 'iron ion transport'] | Temporarly Down Regulated Down Regulated Up Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Up Regulated Down Regulated Down Regulated Temporarly Up Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD nikD alsE yjcS gltP args ribE rlmG gutQ yqjH ftsx nuoK ebgR ebgA | 947911 947577 947570 947571 947572 947573 947574 947983 946459 947889 948595 948597 946452 947582 947582 947582 947582 947582 947582 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase Ribosomal RNA large subunit methyltransferase G Protein gutQ Uncharacterized protein tygH Cell division protein ftsX NADH-quinone oxidoreductase subunit K HTH-type transcriptional regulator ebgR Evolved beta-galactosidase subunit alpha | [] [] [] [] [] [] Adaptation to stress', 'ion transport', 'iron ion transport'] | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Up Regulated Down Regulated Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Temporarly Up Regulated |
| yhgF rpsU dnaG ypfH cbpA holA btuF dcrB thiD nikD alsE yjcS gltP argS ribE rlmG gutQ yqjH ftsX nuok ebgA yqjl | 947911 947577 947570 947571 947572 947573 947574 947983 946459 947989 948595 948595 948595 947589 947589 947580 947580 947580 947580 947583 | Protein yhgF 30S ribosomal protein S21 DNA primase Esterase ypfH Curved DNA-binding protein DNA polymerase III subunit delta Vitamin B12-binding protein Protein dcrB Phosphomethylpyrimidine kinase Nickel import ATP-binding protein nikD D-allulose-6-phosphate 3-epimerase Uncharacterized protein yjcS Proton glutamate symport protein Arginyl-tRNA synthetase 6,7-dimethyl-8-ribityllumazine synthase Ribosomal RNA large subunit methyltransferase G Protein gutQ Uncharacterized protein yqjH Cell division protein ftsX NADH-quinone oxidoreductase subunit K HTH-type transcriptional regulator ebgR Evolved beta-galactosidase subunit alpha Uncharacterized protein yqjl | [] [] [] [] [] [] Adaptation to stress', 'ion transport', 'iron ion transport'] | Temporarly Down Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Up Regulated Down Regulated Up Regulated Down Regulated Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Temporarly Up Regulated Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Temporarly Up Regulated |
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| oroP 049022 Transprintional regulatory protein oroP | | Temporarly Up Regulated |
|--|---|---------------------------------------|
| creB 948922 Transcriptional regulatory protein creB mocA 947356 Uncharacterized protein ygfJ | | Ambiguous/Unsorted |
| | [lala atran transport] | |
| ygfJ 947356 Uncharacterized protein ygfJ creA 948920 Protein creA | ['electron transport'] | Ambiguous/Unsorted |
| | | Ambiguous/Unsorted |
| mod(945362 Molybdenum import ATP-binding protein modC | | Temporarly Up Regulated Up Regulated |
| mod/ 945364 Molybdate-binding periplasmic protein | | Temporarly Down Regula |
| modf 945368 Putative molybdenum transport ATP-binding protein modF | | |
| qseB 945369 Transcriptional regulatory protein qseB ubiC 948545 Chorismatepyruvate lyase | | Down Regulated Down Regulated |
| | | Ambiguous/Unsorted |
| lexA 948544 LexA repressor malM 948547 Maltose operon periplasmic protein | | Temporarly Up Regulated |
| yqfB 947380UPF0267 protein yqfB | | Temporarly Up Regulated |
| hinT 948549HIT-like protein hinT | | Up Regulated |
| rpsP 947103 30S ribosomal protein S16 | | Down Regulated |
| lamB 948548 Maltoporin | | Up Regulated |
| yfiL 947106 Uncharacterized protein yfiL | ['Replication recombination and repair', 'DNA recombination'] | Up Regulated |
| recN 947105 DNA repair protein recN | [Replication recombination and repair ; DNA recombination] | Up Regulated |
| mtn 948542 5'-methylthioadenosine/S-adenosylhomocysteine nucleosida | Se l | Ambiguous/Unsorted |
| aldB 948104 Aldehyde dehydrogenase B | | Up Regulated |
| yagT 945330 Putative xanthine dehydrogenase yagT iron-sulfur-binding su | bunit | Up Regulated |
| paoA 945330 Putative xanthine dehydrogenase yagT iron-sulfur-binding su | | Up Regulated |
| yiaU 948107Uncharacterized HTH-type transcriptional regulator yiaU | П | Temporarly Up Regulated |
| visC 947389 Protein visC | | Temporarly Up Regulated |
| chbB 945339N.N'-diacetylchitobiose-specific phosphotransferase enzyme | IIB component | Up Regulated |
| ubiH 947388 2-octaprenyl-6-methoxyphenol hydroxylase | | Down Regulated |
| ybbO 945337 Uncharacterized oxidoreductase ybbO | П | Ambiguous/Unsorted |
| thiL 947387 Thiamine-monophosphate kinase | | Down Regulated |
| ygfY 947382UPF0350 protein ygfY | | Ambiguous/Unsorted |
| sapF 945335 Peptide transport system ATP-binding protein sapF | | Ambiguous/Unsorted |
| lyxK 948101L-xylulose/3-keto-L-gulonate kinase | | Up Regulated |
| allR 945333 HTH-type transcriptional repressor allR | | Ambiguous/Unsorted |
| selB 948103 Selenocysteine-specific elongation factor | | Down Regulated |
| cueR 945332HTH-type transcriptional regulator cueR | | Ambiguous/Unsorted |
| aegA 947383 Protein aegA | | Ambiguous/Unsorted |
| dusA 948558tRNA-dihydrouridine synthase A | | Ambiguous/Unsorted |
| qorA 948556 Quinone oxidoreductase | | Ambiguous/Unsorted |
| dnaB 948555 Replicative DNA helicase | | Ambiguous/Unsorted |
| yjbJ 948553 UPF0337 protein yjbJ | | Up Regulated |
| zur 948552 Zinc uptake regulation protein | | Down Regulated |
| srlR 948942 Glucitol operon repressor | | Ambiguous/Unsorted |
| mtlA 948118 PTS system mannitol-specific EIICBA component | | Up Regulated |
| mtID 948117 Mannitol-1-phosphate 5-dehydrogenase | | Ambiguous/Unsorted |
| cydA 945341 Cytochrome d ubiquinol oxidase subunit 1 | | Ambiguous/Unsorted |
| yibK 948119 Uncharacterized tRNA/rRNA methyltransferase yibK | | Temporarly Down Regula |
| yibH 948110 Inner membrane protein yibH | | Ambiguous/Unsorted |
| yceB 947377 Uncharacterized lipoprotein yceB | ['electron transport'] | Ambiguous/Unsorted |
| betB 947376 Betaine aldehyde dehydrogenase | | Temporarly Down Regula |
| lysS 947372 Lysyl-tRNA synthetase; Lysyl-tRNA synthetase, heat inducib | e | Down Regulated |
| sodC 945343 Superoxide dismutase [Cu-Zn] | | Up Regulated |
| yibF 948113 Uncharacterized GST-like protein yibF | Ш | Temporarly Up Regulated |
| alkA 947371 DNA-3-methyladenine glycosylase 2 | | Down Regulated |
| astA 946261 Arginine N-succinyltransferase | | Up Regulated |
| rluA 946262 Ribosomal large subunit pseudouridine synthase A; Pseudou | iridine syntnase | Up Regulated |
| astD 946260 N-succinylglutamate 5-semialdehyde dehydrogenase | | Up Regulated |
| chbF 946266 6-phospho-beta-glucosidase | | Ambiguous/Unsorted |
| moal 945398 Molybdopterin-converting factor subunit 1 | | Ambiguous/Unsorted |
| selA 948124 L-seryl-tRNA(Sec) selenium transferase | | Down Regulated |
| moat 945399 Molybdopterin-converting factor subunit 2 | | Up Regulated |
| gpsA 948125[Glycerol-3-phosphate dehydrogenase [NAD(P)+] rhsA 948120[Protein rhsA | | Ambiguous/Unsorted Up Regulated |
| rhsA 948120 Protein rhsA IldD 948121 L-lactate dehydrogenase [cytochrome] | | Down Regulated |
| ybhK 945390 UPF0052 protein ybhK | | Down Regulated Down Regulated |
| cysE 948126 Serine acetyltransferase | | Ambiguous/Unsorted |
| Moal 945396 Molybdenum cofactor biosynthesis protein B | | Ambiguous/Unsorted Ambiguous/Unsorted |
| moa(945397 Molybdenum cofactor biosynthesis protein C | | Ambiguous/Unsorted Ambiguous/Unsorted |
| mody 0+000/priorybuchum coracior biosynthesis protein C | | Ambiguous/Onsorted |

| vibQ 9481: | 28 Uncharacterized protein yibQ | ['proteolysis'] | Down Regulated |
|--|---|--|--|
| | 29 Uncharacterized protein yibQ | [proteorysis] | Down Regulated |
| | 70 Uncharacterized protein yibi | [' Primary Active Transporters'] | Down Regulated |
| | 71 Protein ynjB | [Filmary Active Transporters] | Up Regulated |
| | 72 Uncharacterized HTH-type transcriptional regulator ydjF | ['Transcription', 'transcription', 'regulation of transcription DNA dependent'] | Up Regulated |
| | 76 Pyrazinamidase/nicotinamidase | [mascription, transcription, regulation of transcription DNA dependent] | Down Regulated |
| | 77 CTP pyrophosphohydrolase | | Ambiguous/Unsorted |
| | 78 L-asparaginase 1 | | Ambiguous/Unsorted |
| | 35 ADP-heptoseLPS heptosyltransferase 2 | | Ambiguous/Unsorted |
| | 95 Uncharacterized response regulatory protein ypdB | n | Up Regulated |
| | 36 Lipopolysaccharide heptosyltransferase 1 | U | Ambiguous/Unsorted |
| | 33 Xylulose kinase | | Up Regulated |
| | 93 Glycine cleavage system H protein | | Ambiguous/Unsorted |
| | 98 Uncharacterized protein yggE | [Translation ribosomal structure and biogenesis'] | Up Regulated |
| | 32 Glutaredoxin-3 | · · · · · · · · · · · · · · · · · · · | Up Regulated |
| gpml 9481 | 30 2,3-bisphosphoglycerate-independent phosphoglycerate mutas | | Down Regulated |
| tdh 9481 | 39 L-threonine 3-dehydrogenase | | Ambiguous/Unsorted |
| | 38 2-amino-3-ketobutyrate coenzyme A ligase | | Down Regulated |
| cusC 9462 | 88 Cation efflux system protein cusC | | Ambiguous/Unsorted |
| | 06 Uncharacterized protein yjjU | | Down Regulated |
| ybgC 9489 | 07 Acyl-CoA thioester hydrolase ybgC | | Down Regulated |
| yafD 9462 | 36 UPF0294 protein yafD | | Down Regulated |
| slt 9489 | 08 Soluble lytic murein transglycosylase | | Down Regulated |
| | B1 Protease 4 | | Down Regulated |
| | O1 Uncharacterized protein yjaG | 0 | Down Regulated |
| | D2 Endonuclease V | | Ambiguous/Unsorted |
| | 06 Sensor protein; Sensor protein zraS | | Down Regulated |
| | Periplasmic protein torT | | Up Regulated |
| | B2 Lipoprotein nlpE | | Up Regulated |
| | 41 Xylose isomerase | | Up Regulated |
| | B1 D-erythro-7,8-dihydroneopterin triphosphate epimerase | | Up Regulated |
| | 78 Pseudouridine synthase; Ribosomal small subunit pseudouridin 79 Uncharacterized protein yahG | | Down Regulated |
| yahG 9453 | | ['Amino acid transport and metabolism', ' Central intermediary metabolism', ' Metabolism of other compounds', 'metabolic process'] | |
| | | | Ambiguous/Unsorted |
| bioA 9453 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas | | Down Regulated |
| bioA 9453 ybhH 9453 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH | | Down Regulated Temporarly Down Regula |
| bioA 9453 ybhH 9453 rfaG 9481 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoB 9489 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase | | Down Regulated Temporarly Down Regula Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 radA 9489 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 99 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 radA 9489 bioB 9453 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase | | Down Regulated Temporarly Down Regulate Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated |
| bioA 9453' ybhH 9453' rfaG 9481- ybhA 9453' DeoE 9489- hisP 9467' radA 9489- bioB 9453' mngF 9453' | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 radA 9489 bioB 9453 yjiY 9489 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY | | Down Regulated Temporarly Down Regula Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Temporarly Up Regulated |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 radA 9489 bioB 9453 mngF 9453 yjiY 9489 yfcH 9467 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated |
| bioA 9453 ybhH 9453 rfaG 9481- ybhA 9453 DeoE 9489 hisP 9467- radA 9489 bioB 9453 mngF 9453 yjiY 9489 yfcH 9467- serB 9489 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 89 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 84 Epimerase family protein yfcH | | Down Regulated Temporarly Down Regula Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Temporarly Up Regulated |
| bioA 9453' ybhH 9453' rfaG 9481- ybhA 9453' DeoE 9489 hisP 9467- radA 9489 bioB 9453' mngF 9453' yjiY 9489 yfcH 9467- serB 9489 fdrA 9462' | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 154 Epimerase family protein yfcH 15 Phosphoserine phosphatase | | Down Regulated Temporarly Down Regulate Temporarly Up Regulatev Temporarly Up Regulatev Ambiguous/Unsorted Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Temporarly Up Regulatev Up Regulated Down Regulated Down Regulated |
| bioA 9453' ybhH 9453' rfaG 9481- ybhA 9453' DeoE 9489- hisP 9467- radA 9489- bioB 9453' yjjY 9489- yfcH 9467- serB 9489- fdrA 9462- metA 9485 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 84 Epimerase family protein yfcH 13 Phosphoserine phosphatase 98 Protein fdrA | | Down Regulated Temporarly Down Regulate Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Temporarly Up Regulated Up Regulated Temporarly Up Regulated Down Regulated Down Regulated Ambiguous/Unsorted |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 radA 9489 bioB 9453 yjjY 9489 yfcH 9467 serB 9489 fdrA 9462 aceB 9485 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 34 Epimerase family protein yfcH 13 Phosphoserine phosphatase 98 Protein fdrA 13 Homoserine O-succinyltransferase | | Down Regulated Temporarly Down Regulate Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Temporarly Up Regulated Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 radA 9489 bioB 9453 mngF 9453 yjiY 9489 yfcH 9467 serB 9489 fdrA 9462 metA 9485 aceB 9485 rfaP 9481aceA 9485 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 77 Phosphatase ybhA 10 Phosphopentomutase 89 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 15 Epimerase family protein yfcH 16 Phosphoserine phosphatase 17 Phosphoserine O-succinyltransferase 18 Protein fdrA 19 Homoserine O-succinyltransferase 19 Malate synthase A; Malate synthase 10 Lipopolysaccharide core biosynthesis protein rfaP 17 Isocitrate lyase | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Down Regulated Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Up Regulated |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 radA 9489 bioB 9453 mngF 9453 yjiY 9489 yfcH 9467 serB 9489 fdrA 9462 metA 9485 aceB 9485 rfaP 9481aceA 9485 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 77 Phosphatase ybhA 10 Phosphopentomutase 89 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 84 Epimerase family protein yfcH 13 Phosphoserine phosphatase 88 Protein fdrA 19 Homoserine O-succinyltransferase 12 Malate synthase A; Malate synthase 50 Lipopolysaccharide core biosynthesis protein rfaP | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Ambiguous/Unsorted Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated |
| bioA 9453' ybhH 9453' rfaG 9481- ybhA 9453' DeoB 9489- hisP 9467- radA 9489- bioB 9453' mngF 9453' yjiY 9489- yfcH 9467- serB 9489- fdrA 9462- metA 9485- aceB 9485 rfaP 9481- aceA 9485- pagB 9485 yjbD 9485 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 72 Phosphatase ybhA 10 Phosphopentomutase 39 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 34 Epimerase family protein yfcH 13 Phosphoserine phosphatase 98 Protein fdrA 13 Homoserine O-succinyltransferase 12 Malate synthase A; Malate synthase 15 Lipopolysaccharide core biosynthesis protein rfaP 16 Isocitrate lyase 18 Uncharacterized protein yjbD 18 Uncharacterized protein yjbD | | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Down Regulated Up Regulated Up Regulated Up Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 DeoE 9489 hisP 9467 radA 9489 bioB 9453 yjjY 9489 yfcH 9467 serB 9489 fdrA 9462 metA 9485 aceB 9485 rfaP 9481 aceA 9485 pagB 9485 yjbD 9485 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 77 Phosphatase ybhA 10 Phosphopentomutase 89 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 84 Epimerase family protein yfcH 13 Phosphoserine phosphatase 89 Protein fdrA 13 Homoserine O-succinyltransferase 12 Malate synthase A; Malate synthase 50 Lipopolysaccharide core biosynthesis protein rfaP 17 Isocitrate lyase 18 Uncharacterized protein yjbD 19 INADH dehydrogenase | e [] | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Up Regulated Up Regulated Down Regulated |
| bioA 9453 ybhH 9453 rfaG 9481- ybhA 9453 DeoB 9489 hisP 9467- radA 9489 bioB 9453 yjjY 9489 yfcH 9467- serB 9489 fdrA 9462- metA 9485 aceB 9485 rfaP 9481- aceA 9485 pagB 9485 yjbD 9486- yicC 9481- | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 77 Phosphatase ybhA 10 Phosphopentomutase 89 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 84 Epimerase family protein yfcH 13 Phosphoserine phosphatase 89 Protein fdrA 13 Homoserine O-succinyltransferase 12 Malate synthase A; Malate synthase 50 Lipopolysaccharide core biosynthesis protein rfaP 17 Isocitrate lyase 18 Uncharacterized protein yjbD 19 INADH dehydrogenase 54 UPF0701 protein yicC | e [] | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Down Regulated Up Regulated Up Regulated Up Regulated Ambiguous/Unsorted Up Regulated Up Regulated Down Regulated Ambiguous/Unsorted Up Regulated Down Regulated Down Regulated Down Regulated Down Regulated Down Regulated Up Regulated Down Regulated Up Regulated Down Regulated |
| bioA 9453 ybhH 9453 rfaG 9481 ybhA 9453 TaG 9489 hisP 9467 radA 9489 bioB 9453 mngf 9453 yjiY 9489 yfcH 9467 serB 9489 fdrA 9462 metA 9485 aceB 9485 rfaP 9481 aceA 9485 pagB 9485 yjbD 9485 ndh 9467 yicC 9481 purF 9467 | 76 Adenosylmethionine-8-amino-7-oxononanoate aminotransferas 75 Uncharacterized protein ybhH 49 Lipopolysaccharide core biosynthesis protein rfaG 77 Phosphatase ybhA 10 Phosphopentomutase 89 Histidine transport ATP-binding protein hisP 12 DNA repair protein radA 70 Biotin synthase 71 Mannosyl-D-glycerate transport/metabolism system repressor n 14 Inner membrane protein yjiY 84 Epimerase family protein yfcH 13 Phosphoserine phosphatase 88 Protein fdrA 13 Homoserine O-succinyltransferase 12 Malate synthase A; Malate synthase 150 Lipopolysaccharide core biosynthesis protein rfaP 17 Isocitrate lyase 18 Uncharacterized protein yjbD 18 Uncharacterized protein yjbD 20 NADH dehydrogenase 24 UPF0701 protein yicC 24 Amidophosphoribosyltransferase | e [] | Down Regulated Temporarly Down Regulated Temporarly Up Regulated Temporarly Up Regulated Temporarly Up Regulated Ambiguous/Unsorted Down Regulated Up Regulated Down Regulated Down Regulated Down Regulated Ambiguous/Unsorted Ambiguous/Unsorted Up Regulated Down Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Up Regulated Down Regulated Up Regulated Down Regulated Down Regulated Temporarly Up Regulated |
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| ybgE 945326 Uncharacterized protein ybgE ['Energy production and conversion', 'Energy metabolism carbon', 'Energy production transport', 'electron transport'] Up Regulated ttcA 948967 tRNA 2-thiocytidine biosynthesis protein ttcA Down Regulated muC 948966 DNA recombination protein rmuC yeaP 948969 Probable diguanylate cyclase yeaP ygiQ 948963 UPF0313 protein ygiQ nei 945320 Endonuclease VIII; Endonuclease 8 glpC 946735 Anaerobic glycerol-3-phosphate dehydrogenase subunit C Down Regulated | lptG | | | | |
| ttcA 948967 tRNA 2-thiocytidine biosynthesis protein ttcA muC 948966 DNA recombination protein rmuC yeaP 948969 Probable diguanylate cyclase yeaP ygiQ 948963 UPF0313 protein ygiQ nei 945320 Endonuclease VIII; Endonuclease 8 glpC 946735 Anaerobic glycerol-3-phosphate dehydrogenase subunit C | | | | | |
| muC 948966 DNA recombination protein rmuC Up Regulated yeaP 948969 Probable diguanylate cyclase yeaP Down Regulated ygiQ 948963 UPF0313 protein ygiQ Down Regulated nei 945320 Endonuclease VIII; Endonuclease 8 Down Regulated glpC 946735 Anaerobic glycerol-3-phosphate dehydrogenase subunit C Down Regulated | ybgE | | | ['Energy production and conversion', ' Energy metabolism carbon ', ' Energy production transport', 'electron transport'] | |
| yeaP 948969 Probable diguanylate cyclase yeaP Down Regulated ygiQ 948963 UPF0313 protein ygiQ Down Regulated nei 945320 Endonuclease VIII; Endonuclease 8 Down Regulated glpC 946735 Anaerobic glycerol-3-phosphate dehydrogenase subunit C Down Regulated | | 948967 | | | Down Regulated |
| ygiQ948963 UPF0313 protein ygiQDown Regulatednei945320 Endonuclease VIII; Endonuclease 8Down RegulatedglpC946735 Anaerobic glycerol-3-phosphate dehydrogenase subunit CDown Regulated | rmuC | | | | |
| nei 945320 Endonuclease VIII; Endonuclease 8 Down Regulated glpC 946735 Anaerobic glycerol-3-phosphate dehydrogenase subunit C Down Regulated | | | | | Down Regulated |
| nei 945320 Endonuclease VIII; Endonuclease 8 Down Regulated glpC 946735 Anaerobic glycerol-3-phosphate dehydrogenase subunit C Down Regulated | | | | | |
| | nei | 945320 | | | Down Regulated |
| men(946734 o-succinylbenzoate synthase Temporarly Down Regula | | | | | |
| | men(| 946734 | o-succinylbenzoate synthase | | Temporarly Down Regula |

| atoB 94 | 6727 Acetyl-CoA acetyltransferase | | Temporarly Down Regula |
|----------|--|--|-------------------------------------|
| | 15319Uncharacterized protein yeiG | п | Down Regulated |
| , , | 5318UPF0271 protein ybgL | <u>U</u> | Ambiguous/Unsorted |
| | 8993 Sensor kinase protein rcsC; Sensor protein | | Down Regulated |
| | 8992 DnaJ-like protein djlA | | Temporarly Down Regula |
| | 6701 Uncharacterized protein yeiR | П | Down Regulated |
| | 6700 Ecotin | | Up Regulated |
| | 8996 Excinuclease cho | | Temporarly Up Regulated |
| | 6702 Malate:quinone oxidoreductase; Probable malate:quinone oxido | l preductace | Temporarly Up Regulated |
| | 15711 Curli production assembly/transport component csgE | neductase | Down Regulated |
| | 5305 Osmotically-inducible lipoprotein E | | Up Regulated |
| | 5306 Protein rhsC | | Ambiguous/Unsorted |
| yccX 94 | 5304 Acylphosphatase | | Up Regulated |
| | 8980 Lipoprotein bor homolog from lambdoid prophage DLP12 | | Ambiguous/Unsorted |
| SdhB 94 | 15300 Succinate dehydrogenase iron-sulfur subunit | | Ambiguous/Unsorted |
| | 8982 Hydrogenase 2 maturation protease | | Temporarly Down Regula |
| | 6714 Cytochrome c biogenesis ATP-binding export protein ccmA | | Temporarly Down Regula |
| | 6713 Anaerobic glycerol-3-phosphate dehydrogenase subunit A | | Up Regulated |
| | 6712 Menaquinone-specific isochorismate synthase | | Ambiguous/Unsorted |
| | 6711 Thiamine biosynthesis lipoprotein apbÉ | | Up Regulated |
| | 8986 GMP reductase | | Ambiguous/Unsorted |
| | 8989 Outer-membrane lipoprotein carrier protein | | Up Regulated |
| | 5716 Choline dehydrogenase | | Down Regulated |
| | 6705 ABC transporter ATP-binding protein yojl | | Temporarly Up Regulated |
| | 17887 Aerobic respiration control sensor protein arcB; Sensor protein | | Down Regulated |
| frIR 94 | 7885 HTH-type transcriptional regulator frIR | | Ambiguous/Unsorted |
| DHAI 94 | 5743 PTS-dependent dihydroxyacetone kinase operon regulatory pro | tein | Up Regulated |
| | 7882 Methionine aminopeptidase | | Ambiguous/Unsorted |
| minC 94 | 5744 Septum site-determining protein minC; Probable septum site-de | termining protein minC | Down Regulated |
| cysG 94 | 7880 Siroheme synthase | | Down Regulated |
| | 5740 Cell division topological specificity factor | | Down Regulated |
| PabA 94 | 7873 Para-aminobenzoate synthase glutamine amidotransferase con | nponent II | Down Regulated |
| rpsB 94 | 7874 30S ribosomal protein S2 | | Ambiguous/Unsorted |
| frlB 94 | 7875 Fructoselysine 6-phosphate deglycase | | Temporarly Up Regulated |
| | 5735 Uncharacterized HTH-type transcriptional regulator ycgE | | Down Regulated |
| | 7871 Uncharacterized protein yhfG | ['Amino acid transport and metabolism', 'protein folding', 'amino acid biosynthetic process'] | Up Regulated |
| | 17872 Cell filamentation protein fic | | Up Regulated |
| | 5725 NTE family protein rssA | | Ambiguous/Unsorted |
| | 5763 GTP cyclohydrolase-2 | | Down Regulated |
| | 5764 ATP-dependent Clp protease ATP-binding subunit clpA | | Up Regulated |
| | 5765 Peptidyl-tRNA hydrolase | | Temporarly Down Regula |
| | 5767 Uncharacterized protein ycgV | ['Amino acid transport and metabolism', ' Building block biosynthesis'] | Temporarly Up Regulated |
| | 5758 Glutathione S-transferase | | Ambiguous/Unsorted |
| | 7895 Phosphoglycolate phosphatase | | Ambiguous/Unsorted |
| | 7896 Ribulose-phosphate 3-epimerase | | Ambiguous/Unsorted |
| | 5752 D-amino acid dehydrogenase small subunit | | Down Regulated |
| | 15753 Protein sufB | Boolington recombination and renaid 1 Coulors | Up Regulated |
| | 17890 Uncharacterized protein yhfW | ['Replication recombination and repair', ' Carbon compound utilization'] | Ambiguous/Unsorted |
| | 45751 Vitamin B12 import ATP-binding protein btuD | | Temporarly Down Regula |
| | 1-5756 Muramoyltetrapeptide carboxypeptidase 1-5757 Periplasmic trehalase; Putative periplasmic trehalase | | Temporarly Down Regula Up Regulated |
| | 17893 DNA adenine methylase | | Down Regulated |
| | 17893 DNA adenine metriylase 15754 Alanine racemase, catabolic; Alanine racemase | | Up Regulated |
| | 19754 Admine racemase, catabolic, Alanine racemase 19755 Cell volume regulation protein A | | Down Regulated |
| dhaM 04 | 15749PTS-dependent dihydroxyacetone kinase, phosphotransferase | l subunit dhaM | Up Regulated |
| | 15748 PTS-dependent dinydroxyacetone kinase, phosphotiansierase | | Up Regulated |
| | 15747 PTS-dependent dinydroxyacetone kinase, Abr-binding subunit | | Ambiguous/Unsorted |
| | 15788 Nitrate/nitrite sensor protein narX | dang dabank anart | Down Regulated |
| | 17077 Chaperone protein clpB | | Ambiguous/Unsorted |
| | 15784 Protein sirB1 | | Temporarly Down Regula |
| | 5786 Uncharacterized protein ydhS | [' Building block biosynthesis'] | Up Regulated |
| | 7074 Uncharacterized protein yeaC | | Temporarly Up Regulated |
| | 5782 Respiratory nitrate reductase 1 alpha chain | IN THE PROPERTY OF THE PROPERT | Temporarly Down Regula |
| | 7072 GDP-mannose pyrophosphatase nudK | | Ambiguous/Unsorted |
| | 5779 Protein methyltransferase hemK | | Temporarly Up Regulated |
| <u> </u> | 1 | | Transparator |

| nfsB | 0/5778 | Oxygen-insensitive NAD(P)H nitroreductase | | Ambiguous/Unsorted |
|-------------|--------|--|--|---|
| yfiH | | UPF0124 protein yfiH | | Temporarly Up Regulated |
| hem/ | | Glutamyl-tRNA reductase | | Down Regulated |
| yqaB | | Phosphatase yqaB | | Up Regulated |
| lolB | | Outer-membrane lipoprotein IoIB | | Ambiguous/Unsorted |
| aroF | 947084 | Phospho-2-dehydro-3-deoxyheptonate aldolase, Tyr-sensitive | | Down Regulated |
| ispE | | 4-diphosphocytidyl-2-C-methyl-D-erythritol kinase | | Temporarly Up Regulated |
| agp | | Glucose-1-phosphatase | | Up Regulated |
| rluD | | Pseudouridine synthase; Ribosomal large subunit pseudouridine sy | winthase D | Ambiguous/Unsorted |
| cysB | | HTH-type transcriptional regulator cysB | Trial Co. D | Ambiguous/Unsorted |
| ybaK | | Uncharacterized protein ybaK | | Up Regulated |
| uxuA | | Mannonate dehydratase | | Up Regulated |
| ychF | | GTP-dependent nucleic acid-binding protein engD | | Down Regulated |
| yegP | | UPF0339 protein yegP | | Up Regulated |
| rpIS | | 50S ribosomal protein L19 | | Ambiguous/Unsorted |
| trmD | 947099 | tRNA (guanine-N(1)-)-methyltransferase | | Ambiguous/Unsorted |
| yfiB | 947094 | Putative lipoprotein yfiB | | Down Regulated |
| narL | 945795 | Nitrate/nitrite response regulator protein narL | | Temporarly Up Regulated |
| ychN | 945794 | Protein ychN | | Ambiguous/Unsorted |
| rpIE | 947800 | 50S ribosomal protein L5 | | Ambiguous/Unsorted |
| rpsN | | 30S ribosomal protein S14 | | Down Regulated |
| rpsH | | 30S ribosomal protein S8 | | Down Regulated |
| ptrB | | Protease 2 | | Ambiguous/Unsorted |
| acpS | | Holo-[acyl-carrier-protein] synthase | | Temporarly Up Regulated |
| yoaB | | UPF0076 protein yoaB | | Up Regulated |
| yebE | | Inner membrane protein yebE | | Ambiguous/Unsorted |
| pdxJ | | Pyridoxine 5'-phosphate synthase | | Ambiguous/Unsorted |
| queC | | Queuosine biosynthesis protein queC | | Down Regulated |
| thiC | | Thiamine biosynthesis protein thiC | | Temporarly Down Regula |
| rnc | | Ribonuclease 3 | | Ambiguous/Unsorted |
| yebT | | | norganic ion transport and metabolism'] | Down Regulated |
| era | | GTP-binding protein era; GTP-binding protein era homolog | | Down Regulated |
| yebO | | Uncharacterized protein yebO | and also wheater transport and market allows? | Up Regulated |
| yfhH rsd | | Uncharacterized HTH-type transcriptional regulator yfhH ['Ca Regulator of sigma D | arbohydrate transport and metabolism'] | Ambiguous/Unsorted Temporarly Down Regula |
| hupA | | DNA-binding protein HU-alpha | | Ambiguous/Unsorted |
| nudC | | NADH pyrophosphatase | | Down Regulated |
| zapA | | Cell division protein zapA | | Ambiguous/Unsorted |
| rpml | | 50S ribosomal protein L35 | | Down Regulated |
| rsmF | | Ribosomal RNA small subunit methyltransferase F | | Down Regulated |
| exbD | | Biopolymer transport protein exbD | | Down Regulated |
| dsbB | | Disulfide bond formation protein B | | Temporarly Down Regula |
| nadB | | L-aspartate oxidase | | Temporarly Up Regulated |
| yebV | | | eplication recombination and repair'] | Up Regulated |
| yfiC | | | 56; ECS88 2749; ECDH10B 2743; ECUMN 2898; ECIAI39 2780; UTI89 C2897; EC55989 2864; pC15-1a 066; Z3857; b2575 | |
| rpIA | | 50S ribosomal protein L1 | | Up Regulated |
| rlmA | 946340 | | | Ambiguous/Unsorted |
| yaeH | 947044 | UPF0325 protein yaeH | | Up Regulated |
| manz | | Mannose permease IID component | | Ambiguous/Unsorted |
| bolA | 947043 | Protein bolA | | Up Regulated |
| glrR | | Uncharacterized protein yfhA | | Down Regulated |
| qseF | | Uncharacterized protein yfhA | | Down Regulated |
| lepB | 947040 | Signal peptidase I | | Ambiguous/Unsorted |
| epd | | D-erythrose-4-phosphate dehydrogenase | | Down Regulated |
| mscS | | Small-conductance mechanosensitive channel | | Ambiguous/Unsorted |
| | | RNA polymerase sigma factor; RNA polymerase sigma-E factor | | Temporarly Up Regulated |
| yhcB | | Putative cytochrome d ubiquinol oxidase subunit 3 | | Down Regulated |
| rpIC | | 50S ribosomal protein L3 | | Temporarly Down Regula |
| | | 50S ribosomal protein L23 | | Up Regulated |
| yebC | | UPF0082 protein yebC | | Ambiguous/Unsorted |
| pssA | | CDP-diacylglycerolserine O-phosphatidyltransferase | | Down Regulated |
| rpIB | | 50S ribosomal protein L2 | | Down Regulated |
| , | | Uncharacterized protein yhcM | | Ambiguous/Unsorted |
| rseC | | Sigma-E factor regulatory protein rseC | | Down Regulated |
| etk | 947409 | Tyrosine-protein kinase etk | | Temporarly Up Regulated |

| ruvB | 046274 | Holliday junction ATP-dependent DNA helicase ruvB | | Down Regulated |
|--------------|--------|--|--|--------------------------------|
| rseB | | Sigma-E factor regulatory protein rseB | | Down Regulated Down Regulated |
| rseA | | Sigma-E factor negative regulatory protein | | Down Regulated |
| vfiQ | | SUncharacterized protein vfiQ | n | Up Regulated |
| znuA | | High-affinity zinc uptake system protein znuA | | Up Regulated |
| yciH | | Uncharacterized protein yoiH | ['Translation_ribosomal structure and biogenesis', 'translation'] | Temporarly Up Regulated |
| znuC | | Zinc import ATP-binding protein znuC | Translation modernal structure and biogenesis, translation | Up Regulated |
| vfiP | | DTW domain-containing protein yfiP | | Down Regulated |
| yebK | | BHTH-type transcriptional regulator hexR | | Up Regulated |
| ycfQ | | Uncharacterized HTH-type transcriptional regulator ycfQ | n | Up Regulated |
| nagE | | PTS system N-acetylglucosamine-specific EIICBA component | U | Up Regulated |
| yggU | | UPF0235 protein yggU | | Temporarly Down Regula |
| ybfE | | Uncharacterized protein ybfE | | Up Regulated |
| rdgB | | Nucleoside-triphosphatase; Nucleoside-triphosphatase rdgB | | Ambiguous/Unsorted |
| yfiM | | Uncharacterized protein yfiM | l' RNA related'] | Temporarly Up Regulated |
| hycl | | BHydrogenase 3 maturation protease | [| Up Regulated |
| entE | | Enterobactin synthetase component E | | Up Regulated |
| rpmC | | 750S ribosomal protein L29 | | Ambiguous/Unsorted |
| rpIP | | 50S ribosomal protein L16 | | Down Regulated |
| rpIR | | 50S ribosomal protein L18 | | Temporarly Up Regulated |
| yggT | | Uncharacterized protein yggT | П | Down Regulated |
| rpIN | | 50S ribosomal protein L14 | | Ambiguous/Unsorted |
| rpsQ | 947808 | 30S ribosomal protein S17 | | Temporarly Up Regulated |
| rpIX | 947810 | 50S ribosomal protein L24 | | Down Regulated |
| ruvA | 946369 | Holliday junction ATP-dependent DNA helicase ruvA | | Ambiguous/Unsorted |
| rpIV | 947813 | 350S ribosomal protein L22 | | Up Regulated |
| rpsS | 947811 | 30S ribosomal protein S19 | | Down Regulated |
| selU | 947063 | RNA 2-selenouridine synthase | | Down Regulated |
| exoX | 946361 | Exodeoxyribonuclease 10 | | Down Regulated |
| trxC | 947062 | Thioredoxin-2 | | Up Regulated |
| kgtP | 947069 | Alpha-ketoglutarate permease | | Temporarly Down Regula |
| yebF | | Protein yebF | | Up Regulated |
| edd | | Phosphogluconate dehydratase | | Temporarly Down Regula |
| yfiD | | Autonomous glycyl radical cofactor | | Down Regulated |
| ung | | Uracil-DNA glycosylase | | Down Regulated |
| ycgB | | Uncharacterized protein ycgB | ['metabolic process'] | Up Regulated |
| yebG | | Uncharacterized protein yebG | [Translation ribosomal structure and biogenesis', 'Replication recombination and repair'] | Down Regulated |
| nagB | | Glucosamine-6-phosphate deaminase | | Up Regulated |
| yggR | | Uncharacterized protein yggR | <u>D</u> | Temporarly Up Regulated |
| speA | | Biosynthetic arginine decarboxylase; Arginine decarboxylase | | Down Regulated |
| CHIA | | Probable bifunctional chitinase/lysozyme | | Up Regulated |
| bfr | | Bacterioferritin | | Up Regulated |
| yoaC | | Uncharacterized protein yoaC | [Replication recombination and repair', 'DNA recombination'] | Down Regulated |
| nrdR | | Transcriptional repressor nrdR | | Down Regulated |
| yqgF | | Putative Holliday junction resolvase | | Ambiguous/Unsorted |
| | | Uncharacterized protein yoaF | μ | Temporarly Down Regula |
| pflC | | Pyruvate formate-lyase 2-activating enzyme | | Down Regulated |
| pflD | | Formate acetyltransferase 2 | | Ambiguous/Unsorted |
| argC | | N-acetyl-gamma-glutamyl-phosphate reductase | | Ambiguous/Unsorted |
| argE | | Acetylornithine deacetylase | | Ambiguous/Unsorted |
| rpsL | | 530S ribosomal protein S12 50S ribosomal protein L33 | | Down Regulated Down Regulated |
| rpmG | | Protein tusB | | Up Regulated |
| yheL | 04/04/ | F2348C 4267: FCHMN 4486: FCFD1 4660: h2055: FCDH10 | B_4143; ECIAI1_4163; EC55989_4437; Z5513; ECS88_4410; UTI89_C4546; c4914; ECIAI39_3034 | Temporarly Down Regula |
| | | SE2348C_4267; ECOMN_4486; ECED1_4660; b3999; ECDH10 S30S ribosomal protein S7 | D_T170, ECIMIT_T100, EC00303_4407, 20010, EC000_4410, O1103_04040, 04314, ECIMI33_004 | Down Regulated |
| | | Outer membrane protein N | | Up Regulated |
| | | 2 Uncharacterized lipoprotein yeaY | ["Cell wall membrane envelope biogenesis', 'metabolic process'] | Up Regulated |
| yeaY sad | | Aldehyde dehydrogenase-like protein ynel | Cell wall membrane envelope biogenesis , metabolic process j | Ambiguous/Unsorted |
| | | 850S ribosomal protein L13 | | Up Regulated |
| trml | | BtRNA (guanine-N(7)-)-methyltransferase | | Temporarly Down Regula |
| | | Uncharacterized protein yggL | | Temporarly Down Regula |
| yggL yggW | | Oxygen-independent coproporphyrinogen-III oxidase-like protei | IU | Temporarly Down Regula |
| | | A/G-specific adenine glycosylase | 1934 | Temporarly Up Regulated |
| | | Acetoacetate metabolism regulatory protein atoC | | Temporarly Up Regulated |
| | | NifU-like protein | | Down Regulated |
| .000 | 3 502 | -p | I. | |

| d | 0.47003 Common allutomoral annonno ambigo brutu maldo brudo de brudos annonno | | Tamananani, Davin David |
|------|--|---|-------------------------|
| puuC | 947003 Gamma-glutamyl-gamma-aminobutyraldehyde dehydrogenase | | Temporarly Down Regula |
| gldA | 948440 Glycerol dehydrogenase | | Up Regulated |
| yccS | 947001 Inner membrane protein yccS | | Ambiguous/Unsorted |
| clpS | 948443 ATP-dependent Clp protease adapter protein clpS | | Down Regulated |
| hrpA | 948444 ATP-dependent RNA helicase hrpA | | Down Regulated |
| yeaZ | 946304 M22 peptidase homolog yeaZ | | Ambiguous/Unsorted |
| thiB | 946306 Thiamine-binding periplasmic protein | | Up Regulated |
| flgA | 946300 Flagella basal body P-ring formation protein flgA | | Down Regulated |
| talA | 947006 Transaldolase A; Transaldolase | | Ambiguous/Unsorted |
| mipA | 946301 MItA-interacting protein | | Up Regulated |
| gspK | 947831 Probable general secretion pathway protein K | | Ambiguous/Unsorted |
| yhcO | 947830 Uncharacterized protein yhcO | П | Down Regulated |
| yeaE | 946302 Uncharacterized protein yeaE | <u>u</u> | Up Regulated |
| iscS | 947004 Cysteine desulfurase | | Ambiguous/Unsorted |
| yeaK | 946303 Uncharacterized protein yeaK | | Down Regulated |
| phr | 947005 Deoxyribodipyrimidine photo-lyase | | Ambiguous/Unsorted |
| slyD | 947859 FKBP-type peptidyl-prolyl cis-trans isomerase slyD | | Ambiguous/Unsorted |
| kdpB | 947450 Potassium-transporting ATPase B chain | | Ambiguous/Unsorted |
| | 947453 Uncharacterized protein yggN | [' Central intermediary metabolism'] | Temporarly Up Regulated |
| yggN | 947455 Uncharacterized protein yggin 947454 L-asparaginase 2 | Contral intermediary metabolism; | |
| ansB | | | Up Regulated |
| flgJ | 947456 Peptidoglycan hydrolase flgJ | | Down Regulated |
| speC | 947457 Ornithine decarboxylase, constitutive | | Ambiguous/Unsorted |
| yahA | 947459 Cyclic di-GMP phosphodiesterase yahA | | Temporarly Up Regulated |
| yfhG | 947010 Uncharacterized protein yfhG | [Amino acid transport and metabolism', 'Building block biosynthesis', 'RNA related', 'Type of regulation', 'transcription', 'regulation | |
| ycdX | 948477 Putative hydrolase ycdX | | Down Regulated |
| sdaA | 946331 L-serine dehydratase 1 | | Up Regulated |
| nlpD | 947011 Lipoprotein nlpD | | Up Regulated |
| murB | 948470 UDP-N-acetylenolpyruvoylglucosamine reductase | | Ambiguous/Unsorted |
| man | 946332 Mannose permease IIC component | | Down Regulated |
| yacC | 948472 Uncharacterized protein yacC | ['Amino acid transport and metabolism'] | Down Regulated |
| argR | 947861 Arginine repressor | | Down Regulated |
| qseE | 947013 Putative sensor-like histidine kinase yfhK; Sensor protein | | Temporarly Up Regulated |
| glrK | 947013 Putative sensor-like histidine kinase yfhK; Sensor protein | | Temporarly Up Regulated |
| yhfA | 947860 Protein yhfA | | Ambiguous/Unsorted |
| yebY | 947014 Uncharacterized protein yebY | ['translation'] | Ambiguous/Unsorted |
| yoaE | 946335 UPF0053 inner membrane protein yoaE | | Up Regulated |
| prkB | 947862 Probable phosphoribulokinase; Phosphoribulokinase | | Up Regulated |
| pepT | 946333 Peptidase T | | Up Regulated |
| degS | 947865 Protease degS | | Up Regulated |
| man | 946334 PTS system mannose-specific EIIAB component | | Ambiguous/Unsorted |
| hmp | 947018 Flavohemoprotein | | Down Regulated |
| | 946339 Cold shock-like protein cspC | | Down Regulated |
| cspC | 947019 Uncharacterized zinc-type alcohol dehydrogenase-like protein y | Corban compound utilization! ! Engrav metabolism carbon !! | Ambiguous/Unsorted |
| yphC | | Carbon compound utilization, Energy metabolism carbon j | |
| crp | 947867 Catabolite gene activator | "Cianal transduction machanisma! trianshawilis acid aude! | Down Regulated |
| yhfK | 947866 Uncharacterized protein yhfK | ['Signal transduction mechanisms', 'tricarboxylic acid cycle'] | Temporarly Down Regula |
| pabB | 946337 Para-aminobenzoate synthase component 1 | | Ambiguous/Unsorted |
| ascB | 947460 6-phospho-beta-glucosidase ascB | | Ambiguous/Unsorted |
| yggX | 947461 Probable Fe(2+)-trafficking protein | | Up Regulated |
| yeaU | 946319 Probable tartrate dehydrogenase/decarboxylase | | Up Regulated |
| dmIA | 946319 Probable tartrate dehydrogenase/decarboxylase | | Up Regulated |
| yhdH | 947848 Putative quinone oxidoreductase yhdH | | Ambiguous/Unsorted |
| slyX | 947849 Protein slyX | | Up Regulated |
| leuA | 947465 2-isopropylmalate synthase | | Ambiguous/Unsorted |
| yphG | | [Carbon compound utilization', 'carbohydrate transport'] | Up Regulated |
| cspE | 947024 Cold shock-like protein cspE | | Down Regulated |
| oxyR | 948462 Hydrogen peroxide-inducible genes activator | | Ambiguous/Unsorted |
| fabR | 948460 HTH-type transcriptional repressor fabR | | Ambiguous/Unsorted |
| murG | 946321 UDP-N-acetylglucosamineN-acetylmuramyl-(pentapeptide) pyr | ophosphoryl-undecaprenol N-acetylglucosamine transferase | Ambiguous/Unsorted |
| yphH | 947023 Uncharacterized protein yphH | [Carbohydrate transport and metabolism', 'Transcription', 'carbohydrate metabolic process'] | Ambiguous/Unsorted |
| potA | 946323 Spermidine/putrescine import ATP-binding protein potA | | Down Regulated |
| yheM | 947853 Protein tusC | | Temporarly Down Regula |
| yfgC | 947029 TPR repeat-containing protein yfgC | | Ambiguous/Unsorted |
| ddlB | 946324 D-alanineD-alanine ligase B | | Down Regulated |
| yfhB | 947026 Uncharacterized protein yfhB | [Nucleotide transport and metabolism] | Down Regulated |
| tadA | 947027 tRNA-specific adenosine deaminase | | Down Regulated |
| 1007 | The state of the s | | oga.a.coa |

| yheO | 947851 Uncharacterized protein yheO | ['protein folding'] | Down Regulated |
|------|--|---------------------|-------------------------|
| phoQ | 946326 Sensor protein phoQ; Sensor protein | | Down Regulated |
| birA | 948469 Bifunctional protein birA | | Temporarly Up Regulated |
| kefB | 947858 Glutathione-regulated potassium-efflux system protein kefB | | Temporarly Down Regula |
| kefG | 947857 Glutathione-regulated potassium-efflux system ancillary protein | kefG | Up Regulated |
| Fadd | 946327 Long-chain-fatty-acidCoA ligase | | Up Regulated |
| rnd | 946328 Ribonuclease D | | Down Regulated |
| murl | 948467 Glutamate racemase | | Down Regulated |
| yheS | 947856 Uncharacterized ABC transporter ATP-binding protein yheS | | Ambiguous/Unsorted |
| yheT | 947855 Putative esterase yheT | | Temporarly Up Regulated |