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Consolidated list of down-regulated proteins expressed at different Cr (VI) concentrations at time points

Cr (VI) concentration	6 h	9 h	12 h	18 h	24 h	36 h
	1-pyrroline-5- carboxylate dehydrogenase	1,4-dihydroxy-2- naphthoyl-CoA synthase	50S ribosomal protein L27	1,4-dihydroxy-2- naphthoyl-CoA synthase	hypothetical protein A4A82_03560	DNA-binding protein
	2-oxoglutarate dehydrogenase E1	30S ribosomal protein S1		1-pyrroline-5- carboxylate dehydrogenase		
	2-oxoisovalerate dehydrogenase	ATP-dependent Zn metallopeptidase		2-deoxyribose-5- phosphate aldolase		
	30S ribosomal protein S10	DNA-directed RNA polymerase subunit beta'		3-hexulose-6- phosphate synthase		
	30S ribosomal protein S11	UDP-N- acetylglucosamine 1- carboxyvinyltransfer ase		3-hydroxyacyl-CoA dehydrogenase		
	30S ribosomal protein S16	enoyl-ACP reductase		30S ribosomal protein S2		
	30S ribosomal protein S17	ferritin		30S ribosomal protein S3		
	30S ribosomal protein S20	fructose- bisphosphate aldolase		30S ribosomal protein S5		
	30S ribosomal protein S3	hypothetical protein BSF33_05765		30S ribosomal protein S6		

30S ribosomal protein S4 30S ribosomal protein S5 30S ribosomal protein S7 30S ribosomal protein S9	malate dehydrogenase (acceptor) oligoribonuclease purine-nucleoside phosphorylase	30S ribosomal protein S7 5'(3')- deoxyribonucleotida se 50S ribosomal protein L10 50S ribosomal protein L13
4,4'- diaponeurosporen e glycosyltransfera		50S ribosomal protein L19
50S ribosomal protein L11 50S ribosomal protein L13 50S ribosomal protein L14 50S ribosomal protein L15		50S ribosomal protein L24 50S ribosomal protein L25 50S ribosomal protein L29 50S ribosomal protein L7/L12
50S ribosomal protein L21		ABC transporter ATP-binding protein
50S ribosomal 50S ribosomal protein L23 50S ribosomal protein L25		ATP synthase F0F1 CDP- glycerol:poly(glycer ophosphate) D-alanine aminotransferase

50S ribosomal **DNA-binding** protein L27 protein DNA-directed RNA 50S ribosomal polymerase subunit protein L30 beta' 50S ribosomal F0F1 ATP synthase Fe-S cluster 50S ribosomal assembly protein protein L33 SufD 50S ribosomal GTPase Era protein L4 50S ribosomal GTPase ObgE protein L6 ABC transporter NADH ATP-binding protein dehydrogenase ATP-dependent Clp Oligoendopeptidase protease proteolytic subunit DNA-directed RNA PTS glucose polymerase subunit transporter subunit IIA beta F0F1 ATP synthase ThiJ/PfpI family subunit beta protein UDP-glucose 4-

epimerase

IMP dehydrogenase

UDP-N-

acetylglucosamine 1acetyltransferase carboxyvinyltransfer

ase

beta-ketoacyl-ACP aldehyde

reductase dehydrogenase

beta-ketoacyl-[acyl-

alkaline shock carrier-protein] protein 23

synthase II

branched-chain

alpha-keto acid aspartate

dehydrogenase aminotransferase

subunit E2

bifunctional 5,10-

catabolite control methyleneprotein A tetrahydrofolate

> dehydrogenase/5,10cell division protein

catalase DivIVA

cell division protein citrate synthase

FtsZ

class I fructosecell division protein bisphosphate GpsB

aldolase

citrate synthase cold-shock protein

cysteine synthase cysteine synthase dihydrolipoyl dihydroxyacetone dehydrogenase kinase subunit L

100 mg L-1	elongation factor G	elongation factor P
	elongation factor Tu	enolase
	fructose- bisphosphate aldolase	glutamate dehydrogenase
	glutaminefructose- 6-phosphate aminotransferase	glutaminefructose- 6-phosphate aminotransferase
	glycerol kinase	glutaryl-CoA dehydrogenase
	glycerol-3-phosphate cytidylyltransferase	glycerol kinase
	glycerol-3-phosphate dehydrogenase	glycine C- acetyltransferase
	glycine C- acetyltransferase glycine cleavage system protein T	glycinetRNA ligase hydroxyacid dehydrogenase
	hypothetical protein AXY36_00580	hypothetical protein A4A82_02240
	hypothetical protein RES9_02285	hypothetical protein A4A82_08945

lipid hydroperoxide hypothetical protein peroxidase A4A82_09265

malate:quinone hypothetical protein oxidoreductase A4A82_12170

molecular chaperone hypothetical protein DnaK AXY36_02415

ornithine--oxo-acid hypothetical protein aminotransferase AXY36_02580

pyruvate kinase hypothetical protein AXY36_03955

serine
hydroxymethyltransf
hypothetical protein
AXY36_06600

erase

single-stranded
DNA-binding
protein
AXY36_06780

stage V sporulation hypoxanthine

protein G phosphoribosyltransf

erase succinate--CoA

ligase subunit alpha

dehydrogenase
(acceptor)

succinate--CoA malate:quinone ligase subunit beta oxidoreductase

transaldolase

manganese-

dependent inorganic pyrophosphatase

transcription elongation factor

metal-dependent

GreA

hydrolase

transglycosylase

methionine sulfoxide molecular chaperone

DnaK

transketolase

oligoribonuclease

translation initiation

peptidase

factor IF-3

peptidase M28

trigger factor triose-phosphate

isomerase

phosphoenolpyruvat

e-protein

type I glutamate-ammonia ligase

phosphotransferase

type I glyceraldehyde-3-

phosphonate ABC transporter ATP-

universal stress protein UspA

pyruvate kinase

pyruvate oxidase

pyruvate, phosphate ribosome recycling

factor

sigma-54

modulation protein

		ļ
	stage V sporulation	
	protein G	
	succinate	
	dehydrogenase	
	flavoprotein subunit	
	succinateCoA	
	ligase subunit alpha	
	over a married a	
	superoxide dismutase	
	thiol reductase thioredoxin	
	thioredoxin-disulfide	
	transketolase	
	translation initiation translation initiation	
	type I	
	glyceraldehyde-3-	
	phosphate	
	dehydrogenase	
	universal stress	
	protein	
	universal stress	
	protein UspA	
	uracil	
	uridylate kinase	
	ATP-dependent Clp	•
	protease ATP-	•
200 mg L-1	binding subunit	
	ClpX	ļ

	sulfurtransferase 30S ribosomal	30S ribosomal	30S ribosomal	30S ribosomal	2-oxoisovalerate
peptidase M23	protein S10	protein S16	protein S10	protein S17	dehydrogenase
	30S ribosomal	30S ribosomal	30S ribosomal	30S ribosomal	30S ribosomal
	protein S16	protein S19	protein S18	protein S19	protein S19
	30S ribosomal	beta-ketoacyl-[acyl-	30S ribosomal	50S ribosomal	30S ribosomal
	protein S9	carrier-protein]	protein S9	protein L1	protein S4
	protein 37	synthase II	protein 57	protein L1	protein 54
	50S ribosomal	class I fructose-	50S ribosomal	50S ribosomal	30S ribosomal
	protein L1	bisphosphate	protein L15	protein L11	protein S8
	protein L1	aldolase	protein L13	protein L11	protein 38
	50S ribosomal	universal stress	50S ribosomal	50S ribosomal	30S ribosomal
	protein L14	protein	protein L18	protein L14	protein S9
	50S ribosomal		50S ribosomal	50S ribosomal	50S ribosomal
	protein L18		protein L27	protein L15	protein L1
	50S ribosomal		50S ribosomal	50S ribosomal	50S ribosomal
	protein L23		protein L28	protein L17	protein L14
	50S ribosomal		50S ribosomal	50S ribosomal	50S ribosomal
	protein L30		protein L3	protein L18	protein L15
	50S ribosomal		50S ribosomal	50S ribosomal	50S ribosomal
	protein L33		protein L30	protein L23	protein L17
	DNA-directed RNA		50S ribosomal	50S ribosomal	50S ribosomal
	polymerase subunit		protein L4	protein L3	protein L23
	alpha		protein L4	protein L3	protein L23
	glutamyl-tRNA		6-phosphogluconate	50S ribosomal	50S ribosomal
	synthetase		dehydrogenase	protein L30	protein L30
	glycinetRNA		alanine racemase	50S ribosomal	50S ribosomal
	ligase		aranine racentase	protein L33	protein L4

	octanoyltransferase	aminoacyltransferase	50S ribosomal protein L4	branched-chain alpha-keto acid dehydrogenase subunit E2
	preprotein translocase subunit YajC	asparaginetRNA ligase	aminoacyltransferase	cell division protein FtsZ
	pyruvate kinase	ATP-dependent Clp protease ATP-binding subunit ClpX	ATP-dependent Clp protease ATP- binding subunit ClpX	DNA-directed RNA polymerase subunit beta
300 mg L-1	succinateCoA ligase subunit beta	ATP-dependent Zn metallopeptidase	beta-ketoacyl-[acyl- carrier-protein] synthase II	elongation factor Ts
300 mg L-1	transcription elongation factor GreA	beta-ketoacyl-[acyl- carrier-protein] synthase II	cell division protein FtsZ	fructose- bisphosphate aldolase
	type I glyceraldehyde-3- phosphate dehydrogenase	beta-ketoacyl-ACP reductase	DNA-directed RNA polymerase subunit alpha	hsp90-like protein
	den ja o genade	glucokinase	fructose- bisphosphate aldolase	malate:quinone oxidoreductase
		glucosamine-6- phosphate isomerase	glutamyl-tRNA amidotransferase	pyridoxamine 5- phosphate oxidase
		glutamyl-tRNA amidotransferase	hypothetical protein	trigger factor

		glutamyl-tRNA synthetase	hypothetical protein RES9_02285	type I glyceraldehyde-3- phosphate dehydrogenase
		hypothetical protein AXY36_09360	lysinetRNA ligase	
		IMP dehydrogenase	NAD(P)-dependent oxidoreductase	
		isocitrate dehydrogenase	preprotein translocase subunit YajC	
		lysinetRNA ligase	pyridoxamine 5- phosphate oxidase	
		methioninetRNA ligase peptidase M23 preprotein translocase subunit YajC ribonuclease R single-stranded DNA-binding protein		
		urease subunit alpha Fis family		
2-deoxyribose-5- phosphate aldolase	2-deoxyribose-5- phosphate aldolase	transcriptional regulator	alpha/beta hydrolase	

1	
30S ribosomal protein S1	histidine kinase
30S ribosomal protein S15	hypothetical protein AXY36_01150
30S ribosomal protein S19	hypothetical protein BSF33_08575
30S ribosomal protein S2	phosphopyruvate hydratase
30S ribosomal protein S6	
30S ribosomal protein S8	
3-oxoacyl-[acyl-carrier protein] reductase	
50S ribosomal protein L1	
50S ribosomal protein L10 50S ribosomal protein L17 50S ribosomal protein L19	

L-serine dehydratase, ironsulfur-dependent subunit alpha adenylosuccinate synthetase alkaline-shock protein alpha/beta hydrolase aspartatesemialdehyde dehydrogenase branched-chain amino acid aminotransferase elastin-binding protein malonyl CoA-acyl carrier protein transacylase purine-nucleoside phosphorylase

50S ribosomal

protein L24

50S ribosomal

protein L29

50S ribosomal

protein L3

50S ribosomal

protein L31

50S ribosomal

protein L5

50S ribosomal

protein L7/L12

aconitate hydratase

acyl carrier protein

adenylate kinase

alanine

dehydrogenase

alkaline shock

protein 23

anti-anti-sigma

factor

ATP synthase F0F1

subunit alpha

ATP synthase F0F1

subunit delta

ATP-dependent Zn metallopeptidase

CDP-

glycerol:poly(glycer ophosphate) glycerophosphotrans ferase chorismate mutase

cold-shock protein

delta-aminolevulinic acid dehydratase

dihydrolipoamide succinyltransferase

dihydroxyacetone kinase subunit L DNA-binding protein

elongation factor P

endoribonuclease L-PSP enolase

enoyl-ACP reductase

ferritin

Fe-S cluster assembly protein SufB Fe-S cluster assembly protein SufD glucose-6-phosphate isomerase glutamate dehydrogenase glutaminehydrolyzing GMP synthase glycine--tRNA 100 + 200 mg Lheme-binding protein hydroxymethylpyrim idine/phosphomethyl pyrimidine kinase hypothetical protein A4A82_02240 hypothetical protein A4A82_03560 hypothetical protein A4A82_07220

hypothetical protein A4A82_09265

hypothetical protein AXY36_02580

hypothetical protein BSF33_05765

hypothetical protein RES13_12170, partial

hypothetical protein XA22_10040

inosine-5monophosphate dehydrogenase

iron ABC transporter substrate-binding protein

lactate dehydrogenase late competence protein ComEC

methionine sulfoxide reductase

NADH

dehydrogenase nitric oxide dioxygenase

nucleosidediphosphate kinase

ornithine--oxo-acid transaminase peroxiredoxin Phosphate acetyltransferase phosphoglycerate kinase PTS glucose transporter subunit

putative transporter

IIA

pyruvate dehydrogenase (acetyl-transferring) E1 component subunit alpha

pyruvate, phosphate dikinase

ribose-phosphate pyrophosphokinase

ribosome recycling factor short-chain dehydrogenase succinate dehydrogenase flavoprotein subunit superoxide dismutase thiol reductase thioredoxin thioredoxin thymidine phosphorylase transcriptional repressor CodY Translation elongation factor Ts translation initiation factor IF-1 UDP-glucose 4epimerase UDP-N-acetyl-Dgalactosamine dehydrogenase uracil phosphoribosyltransf

erase

	pathogenicity island protein	30S ribosomal protein S14	transcription elongation factor GreA	DNA polymerase III subunit alpha	DNA polymerase III subunit alpha	Short-chain dehydrogenase/reduc tase SDR
		catabolite control protein A		VOC family protein	hypothetical protein AXY36_09360	alkyl hydroperoxide reductase
		cell division protein GpsB		hypothetical protein AXY36_04620	isocitrate dehydrogenase	hypothetical protein AXY36_09360
		glucosamine-6- phosphate isomerase		phosphogluconate dehydrogenase (NADP(+)- dependent, decarboxylating)		isocitrate dehydrogenase
200 + 300 mg L-1		glycine C- acetyltransferase		pyrimidine- nucleoside phosphorylase		transcriptional regulator
		hypothetical protein A4A82_08725				
		hypothetical protein BSF33_04470				
		isocitrate dehydrogenase				
		manganese- dependent inorganic pyrophosphatase				
		membrane protein				

	sufurtransferase FdhD transglycosylase				
	2-oxoglutarate dehydrogenase E1	2,3- bisphosphoglycerate- independent phosphoglycerate mutase	2,3- bisphosphoglycerate- independent phosphoglycerate mutase	2,3- bisphosphoglycerate- independent phosphoglycerate mutase	1-pyrroline-5- carboxylate dehydrogenase
	2-oxoisovalerate dehydrogenase	3-hexulose-6- phosphate synthase	2-oxoglutarate dehydrogenase E1	2-deoxyribose-5- phosphate aldolase	30S ribosomal protein S20
lacetyltransferase	3-hexulose-6- phosphate synthase	30S ribosomal protein S1	2-oxoglutarate ferredoxin oxidoreductase subunit alpha 3-methyl-2-	2-oxoglutarate dehydrogenase E1	30S ribosomal protein S5
lelongation factor Te	30S ribosomal protein S11	30S ribosomal protein S2	oxobutanoate	2-oxoisovalerate dehydrogenase	30S ribosomal protein S7
Jr · · · · · · · · · · · · · ·	30S ribosomal protein S17	30S ribosomal protein S20	30S ribosomal protein S1	3-hexulose-6- phosphate synthase	50S ribosomal protein L16
	30S ribosomal protein S19 30S ribosomal protein S3	30S ribosomal protein S3 30S ribosomal protein S4	30S ribosomal protein S11 30S ribosomal protein S13	30S ribosomal protein S11 30S ribosomal protein S13	50S ribosomal protein L2 50S ribosomal protein L7/L12
	30S ribosomal protein S4	30S ribosomal protein S5	30S ribosomal protein S15	30S ribosomal protein S15	ATP synthase F0 subunit alpha

30S ribosomal protein S5	30S ribosomal protein S6	30S ribosomal protein S16	30S ribosomal protein S16	ATP-dependent Clp protease proteolytic subunit
30S ribosomal protein S6	30S ribosomal protein S8	50S ribosomal protein L1	30S ribosomal protein S18	F0F1 ATP synthase subunit beta
30S ribosomal protein S7 30S ribosomal protein S8	50S ribosomal protein L10 50S ribosomal protein L13	50S ribosomal protein L16 50S ribosomal protein L21	30S ribosomal protein S2 30S ribosomal protein S20	IMP dehydrogenase NADH dehydrogenase
50S ribosomal protein L10	50S ribosomal protein L15	50S ribosomal protein L36	30S ribosomal protein S3	PTS glucose transporter subunit IIA
50S ribosomal protein L11	50S ribosomal protein L17	8-amino-7- oxononanoate synthase	30S ribosomal protein S4	aconitate hydratase
50S ribosomal protein L13	50S ribosomal protein L18	ATP synthase F0F1 subunit B	30S ribosomal protein S5	alpha/beta hydrolase
50S ribosomal protein L17	50S ribosomal protein L21	ATP synthase F0F1 subunit delta	30S ribosomal protein S6	beta-ketoacyl-ACP reductase
50S ribosomal protein L21	50S ribosomal protein L22	DNA polymerase III subunit beta	30S ribosomal protein S7	class I fructose- bisphosphate aldolase
50S ribosomal protein L22	50S ribosomal protein L23	DNA primase	30S ribosomal protein S8	dihydrolipoamide succinyltransferase
50S ribosomal protein L25	50S ribosomal protein L24	LysR family transcriptional regulator	30S ribosomal protein S9	dihydrolipoyl dehydrogenase

50S ribosomal protein L27	50S ribosomal protein L25	NAD kinase	50S ribosomal protein L10	dihydroxyacetone kinase subunit L
50S ribosomal protein L3	50S ribosomal protein L29	Translation elongation factor Ts	50S ribosomal protein L13	elongation factor G
50S ribosomal protein L31	50S ribosomal protein L3	UDP-N-acetyl-D- galactosamine dehydrogenase	50S ribosomal protein L16	elongation factor Tu
50S ribosomal protein L4	50S ribosomal protein L30	acetate kinase	50S ribosomal protein L19	ferritin
50S ribosomal protein L5	50S ribosomal protein L31	alanine dehydrogenase	50S ribosomal protein L21	glucosamine-6- phosphate isomerase
50S ribosomal protein L6 50S ribosomal protein L7/L12	50S ribosomal protein L5 50S ribosomal protein L6	alcohol dehydrogenase anti-anti-sigma factor	50S ribosomal protein L22 50S ribosomal protein L24	glutamyl-tRNA synthetase hydroxyacid dehydrogenase
ABC transporter ATP-binding protein	50S ribosomal protein L7/L12	anti-sigma B factor RsbW	50S ribosomal protein L25	hypothetical protein A4A82_02240
ATP-dependent Clp protease proteolytic subunit CDP-	6-phosphogluconate dehydrogenase	catabolite control protein A	50S ribosomal protein L27	hypothetical protein A4A82_07220
glycerol:poly(glycer ophosphate) glycerophosphotrans ferase	ABC transporter ATP-binding protein	chorismate mutase	50S ribosomal protein L29	hypothetical protein AXY36_00580
D-alanine aminotransferase	ATP-dependent Clp protease proteolytic subunit	cysteine desulfurase	50S ribosomal protein L31	hypothetical protein BSF33_05765

DNA gyrase subunit	DNA starvation/stationary	delta-aminolevulinic	509 ribosomal	iron ABC transporter
A	phase protection protein	acid dehydratase	protein L5	substrate-binding protein
DNA	•			
starvation/stationary	DNA-binding	dephospho-CoA	50S ribosomal	nitric oxide
phase protection protein	protein	kinase	protein L6	dioxygenase
± •	DNA-directed RNA polymerase subunit alpha	dihydrolipoamide succinyltransferase	50S ribosomal protein L7/L12	ornithineoxo-acid aminotransferase
FOF1 ATP synthase	DNA-directed RNA polymerase subunit beta	dihydrolipoyl dehydrogenase	ABC transporter ATP-binding protein	peroxiredoxin
assembly protein	DNA-directed RNA polymerase subunit beta'	elongation factor Tu	ATP synthase F0F1 subunit alpha	phosphoglycerate kinase
assembly protein	Fe-S cluster assembly protein SufD	endopeptidase Clp ATP-binding subunit C	ATP-dependent Clp protease proteolytic subunit	phosphopyruvate hydratase
NADH dehydrogenase	IMP dehydrogenase	endoribonuclease L- PSP	ATP-dependent Zn metallopeptidase	pyruvate dehydrogenase (acetyl-transferring) E1 component subunit alpha
transporter subunit	NADH dehydrogenase	ferritin	D-alanine aminotransferase	pyruvate kinase

UDP-N-acetyl-D-galactosamine dehydrogenase	PTS glucose transporter subunit IIA	formate tetrahydrofolate ligase	DNA starvation/stationary phase protection protein	serine hydroxymethyltransf erase
acyl carrier protein	Translation elongation factor Ts	general stress protein	DNA-binding protein	succinateCoA ligase subunit alpha
adenylate kinase	UDP-N- acetylglucosamine 1- carboxyvinyltransfer ase	glucose-6-phosphate isomerase	DNA-directed RNA polymerase subunit beta	succinateCoA ligase subunit beta
alkaline shock protein 23	UDP-glucose 4- epimerase	glycerol-3-phosphate dehydrogenase	DNA-directed RNA polymerase subunit beta'	transketolase
branched-chain alpha-keto acid dehydrogenase subunit E2	aconitate hydratase	glycine cleavage system protein T	F0F1 ATP synthase subunit beta	universal stress protein UspA
cell division protein FtsZ	acyl carrier protein	glycine dehydrogenase	Fe-S cluster assembly protein SufB	uracil phosphoribosyltransf erase
class I fructose- bisphosphate aldolase	adenylate kinase	heme-binding protein	Fe-S cluster assembly protein SufD	
cysteine synthase	beta-ketoacyl-ACP reductase	hydroxymethylpyrim idine/phosphomethyl pyrimidine kinase		

dihydrolipoyl dehydrogenase	branched-chain alpha-keto acid dehydrogenase subunit E2	hypothetical protein A4A82_07220	NADH dehydrogenase
elongation factor G	cell division protein FtsZ	hypothetical protein AXY36_00580	Oligoendopeptidase F
elongation factor P	chorismate mutase	hypothetical protein AXY36_05930	PTS glucose transporter subunit IIA
elongation factor Tu	cold-shock protein	hypothetical protein AXY36_07855	Phosphate acetyltransferase
enolase	elongation factor G	hypothetical protein BSF33_00215	RNA-binding protein
glycerol-3-phosphate dehydrogenase	elongation factor Tu	hypothetical protein BSF33_00540	Ribonucleotide reductase of class Ib (aerobic), alpha subunit
glycine cleavage system protein H	enolase	hypothetical protein BSF33_00755	Translation elongation factor Ts
glycine cleavage system protein T	enoyl-ACP reductase	hypothetical protein BSF33_05765	UDP-N- acetylglucosamine 1- carboxyvinyltransfer ase

hydroxymethylpyrim idine/phosphomethyl pyrimidine kinase		hypothetical protein RES10_11420	UDP-glucose 4- epimerase
hypothetical protein XA22_10040	glucose-6-phosphate isomerase	hypothetical protein XA22_10040	acetate kinase
lipid hydroperoxide peroxidase	glycerol-3-phosphate dehydrogenase	iron ABC transporter substrate-binding protein	acetyltransferase
malate:quinone oxidoreductase	glycine C- acetyltransferase	isoleucyl-tRNA synthetase, partial	adenylate kinase
mobilization protein C (plasmid)	glycinetRNA ligase	late competence protein ComEC	alaninetRNA ligase
molecular chaperone DnaK	glyoxalase	lipoateprotein ligase	aldehyde dehydrogenase
peptidase M24	hydroxymethylglutar yl-CoA synthase	malate dehydrogenase	alkaline shock protein 23
phosphoglycerate kinase	hypothetical protein A4A82_06020	manganese ABC transporter substrate- binding protein	aminopeptidase
pseudouridine synthase	hypothetical protein A4A82_07220	nucleoside- diphosphate kinase	aspartyl/glutamyl- tRNA amidotransferase subunit A

		pyridoxamine 5- phosphate oxidase	hypothetical protein A4A82_11275	peroxiredoxin	bifunctional 5,10- methylene- tetrahydrofolate dehydrogenase/5,10- methylene- tetrahydrofolate cyclohydrolase
		pyruvate dehydrogenase (acetyl-transferring) E1 component subunit alpha	malate dehydrogenase (acceptor)	phosphoenolpyruvat e carboxykinase	branched-chain alpha-keto acid dehydrogenase subunit E2
		recombinase A	malate:quinone oxidoreductase	phosphoenolpyruvat e synthase regulatory protein	catabolite control protein A
		ribose-phosphate pyrophosphokinase	malonyl CoA-acyl carrier protein transacylase	phosphoglycerate kinase	cell division protein DivIVA
100 + 300 mg L-	1	ribosome recycling factor	manganese- dependent inorganic pyrophosphatase	phosphopentomutase	cell division protein GpsB
		serine hydroxymethyltransf erase	metal-dependent hydrolase	polynucleotide phosphorylase	chorismate mutase
		succinateCoA ligase subunit alpha	peroxiredoxin	preprotein translocase subunit SecA	class I fructose- bisphosphate aldolase
		superoxide dismutase	phosphoglycerate kinase	proline dehydrogenase	cold-shock protein

tail protein	purine-nucleoside phosphorylase	pyruvate dehydrogenase (acetyl-transferring) E1 component subunit alpha	cysteine synthase
transaldolase	pyruvate dehydrogenase (acetyl-transferring) E1 component subunit alpha	quinone oxidoreductase	dihydrolipoyl dehydrogenase
transcription termination/antiterm ination protein NusA		ribose-phosphate pyrophosphokinase	elongation factor G
transcriptional repressor CodY	pyruvate oxidase	serine hydroxymethyltransf erase	elongation factor P
translation initiation factor IF-1	ribose-phosphate pyrophosphokinase	signal recognition particle	elongation factor Tu
trigger factor	serine hydroxymethyltransf erase	succinyl- diaminopimelate desuccinylase	endoribonuclease L- PSP
triose-phosphate isomerase	superoxide dismutase	thioredoxin	enolase
universal stress protein UspA	thiol reductase thioredoxin	thymidine phosphorylase	enoyl-ACP reductase

uracil phosphoribosyltransf erase	thioredoxin-disulfide reductase	transaldolase	ferritin
uridylate kinase	transaldolase	transcription termination/antiterm ination protein NusA	general stress protein
	transcriptional regulator	transcriptional regulator	glucosamine-6- phosphate isomerase
	transketolase	transcriptional repressor CodY	glucose-6-phosphate isomerase
	translation initiation factor IF-2	triose-phosphate isomerase	glutamate dehydrogenase
	translation initiation factor IF-3	type I glutamate ammonia ligase	glutamyl-tRNA synthetase
	trigger factor		glycerol kinase
	type I glyceraldehyde-3- phosphate dehydrogenase		glycerol-3-phosphate cytidylyltransferase
			glycerol-3-phosphate dehydrogenase
			glycinetRNA ligase glyoxalase

hydroxymethylpyrim idine/phosphomethyl pyrimidine kinase

hypothetical protein A4A82_02240

hypothetical protein A4A82_09265

hypothetical protein A4A82_11275

hypothetical protein AXY36_00580

hypothetical protein AXY36_01150

hypothetical protein AXY36_06600

hypothetical protein AXY36_06780

hypothetical protein BSF33_00215

hypothetical protein BSF33_05765

hypothetical protein BSF33_08575

hypothetical protein XA22_10040

hypoxanthine phosphoribosyltransf erase

iron ABC transporter substrate-binding protein

isoleucyl-tRNA synthetase, partial

lipid hydroperoxide peroxidase

malate dehydrogenase (acceptor) malate:quinone oxidoreductase malonyl CoA-acyl carrier protein transacylase

manganesedependent inorganic pyrophosphatase

membrane protein

metal-dependent hydrolase molecular chaperone DnaK molecular chaperone GrpE

nucleosidediphosphate kinase

oligoribonuclease

ornithine--oxo-acid aminotransferase

peptidase M24 peptidase M28 peroxiredoxin

phosphoenolpyruvat e-protein phosphotransferase

phosphoglycerate kinase preprotein translocase subunit SecA

pyruvate dehydrogenase (acetyl-transferring) E1 component subunit alpha

pyruvate kinase pyruvate oxidase

ribose-phosphate pyrophosphokinase

ribosome recycling factor serine hydroxymethyltransf erase signal recognition particle single-stranded DNA-binding protein

stage V sporulation protein G

succinate--CoA ligase subunit alpha

succinate--CoA ligase subunit beta superoxide dismutase thiol reductase thioredoxin thioredoxin thymidine phosphorylase transaldolase transcription termination/antiterm ination protein NusA transcriptional regulator transcriptional repressor CodY transketolase translation initiation factor IF-1 translation initiation factor IF-3 transposon DNAinvertase trigger factor triose-phosphate isomerase type I glutamate-ammonia ligase type I glyceraldehyde-3phosphate dehydrogenase uracil phosphoribosyltransf erase

uridylate kinase

50S ribosomal protein L16	1-pyrroline-5- carboxylate dehydrogenase	1,4-dihydroxy-2- naphthoyl-CoA synthase	2-heptaprenyl-1,4- naphthoquinone methyltransferase MenG	1-pyrroline-5- carboxylate dehydrogenase	2-oxoglutarate dehydrogenase E1
adenylosuccinate synthetase	2,3- bisphosphoglycerate- independent phosphoglycerate mutase	1-pyrroline-5- carboxylate dehydrogenase	30S ribosomal protein S12	2-oxoglutarate dehydrogenase E1 component	30S ribosomal protein S1
alkyl hydroperoxide reductase subunit F	2-succinylbenzoate- CoA ligase	2-oxoisovalerate dehydrogenase	50S ribosomal protein L2	30S ribosomal protein S1	30S ribosomal protein S12
anti-sigma B factor RsbW	30S ribosomal protein S12	30S ribosomal protein S10	50S ribosomal protein L20	30S ribosomal protein S12	50S ribosomal protein L28
aryldialkylphosphata se	30S ribosomal protein S13	30S ribosomal protein S11	50S ribosomal protein L32	3-hydroxy-3- methylglutaryl-CoA reductase	acetateCoA ligase
glutamine amidotransferase	30S ribosomal protein S15	30S ribosomal protein S13	6,7-dimethyl-8- ribityllumazine synthase	3-hydroxyacyl-CoA dehydrogenase	acetyl-CoA acetyltransferase
glutamine-binding periplasmic protein	30S ribosomal protein S18	30S ribosomal protein S18	aconitate hydratase	3-oxoacyl-[acyl-carrier protein] reductase	alanine dehydrogenase
glutamyl-tRNA amidotransferase	30S ribosomal protein S2	30S ribosomal protein S7	adenine methyltransferase	50S ribosomal protein L2	alcohol dehydrogenase
heme ABC transporter ATP- binding protein	30S ribosomal protein S20	30S ribosomal protein S9	alkyl hydroperoxide reductase subunit F	50S ribosomal protein L20	autolysin

hypothetical protein A4A82_05120	30S ribosomal protein S21	3-hydroxyacyl-CoA dehydrogenase	arginine repressor	50S ribosomal protein L28	betaine- dehydro
hypothetical protein A4A82_07100	3-hydroxy-3- methylglutaryl-CoA reductase	3-oxoacyl-[acyl-carrier protein] reductase	aspartyl/glutamyl- tRNA amidotransferase subunit A	50S ribosomal protein L35	catalase
hypothetical protein A4A82_08360	3-oxoacyl-[acyl-carrier protein] reductase	5'(3')- deoxyribonucleotida se	C4-dicarboxylate ABC transporter	50S ribosomal protein L36	compet ComF
hypothetical protein A4A82_08725	50S ribosomal protein L15	50S ribosomal protein L1	catalase	6,7-dimethyl-8- ribityllumazine synthase	glycero
hypothetical protein A4A82_11275	50S ribosomal protein L16	50S ribosomal protein L11	Chemotaxis- inhibiting protein CHIPS, phage associated	acetateCoA ligase	glycine system
hypothetical protein AXY36_01150	50S ribosomal protein L19	50S ribosomal protein L14	choline dehydrogenase	acetyl-CoA acetyltransferase	hypothe A4A82
hypothetical protein AXY36_04620	50S ribosomal protein L2	50S ribosomal protein L16	CTP synthase	aconitate hydratase	hypothe XA21_
hypothetical protein AXY36_08620	50S ribosomal protein L20	50S ribosomal protein L19	cytochrome aa3 quinol oxidase subunit II	adenylosuccinate synthetase	hypothe XA22_
hypothetical protein AXY36_09850	50S ribosomal protein L24	50S ribosomal protein L2	diaminopimelate decarboxylase	alanine dehydrogenase	hypothe

hypothetical protein BSF33_04470	50S ribosomal protein L29	50S ribosomal protein L4	dihydroorotase	alcohol dehydrogenase	lactate dehydrogenase
hypothetical protein RES9_02800	50S ribosomal protein L32	6- phosphogluconolact onase	DNA polymerase I	aldehyde dehydrogenase family protein	magnesium transporter CorA
integrase	50S ribosomal protein L36	acetate kinase	DNA-binding response regulator	alkaline-shock protein	mercuric reductase
L-lactate dehydrogenase	methyltetrahydropter oyltriglutamate homocysteine methyltransferase	acetateCoA ligase	exported protein	alkyl hydroperoxide reductase subunit F	MFS transporter
malonyl CoA-acyl carrier protein transacylase	6,7-dimethyl-8- ribityllumazine synthase	acetyltransferase	F0F1 ATP synthase subunit epsilon	argininetRNA ligase	serinetRNA ligase
membrane protein	8-amino-7- oxononanoate synthase	alanine dehydrogenase	F0F1 ATP synthase subunit gamma	ArsC family protein	succinate dehydrogenase
methionyl-tRNA	acetyl-CoA	alcohol	formate C-	aspartate	sufurtransferase
formyltransferase	acetyltransferase	dehydrogenase	acetyltransferase	aminotransferase	FdhD TatP family
molecular chaperone GroEL	aconitate hydratase	aldehyde dehydrogenase family protein	glutamate-1- semialdehyde aminotransferase	aspartate- semialdehyde dehydrogenase	TetR family transcriptional regulator
molecular chaperone GrpE	adenylosuccinate synthetase	alkaline shock protein 23	glutamate-1- semialdehyde-2,1- aminomutase	ATP synthase F0F1 subunit B	
pectate lyase	alanine dehydrogenase	aminopeptidase T	glutathione S- transferase	ATP synthase F0F1 subunit delta	

	phosphopyruvate hydratase	alcohol dehydrogenase	anti-sigma B factor RsbW	glutathione- dependent formaldehyde dehydrogenase	ATPase P
	plasmid partitioning protein ParB	alkyl hydroperoxide reductase subunit F	AraC family transcriptional regulator	glycine cleavage system protein H	autolysin
	pyridoxal biosynthesis protein	anti-anti-sigma factor	arylamine N-acetyltransferase	glycine dehydrogenase subunit 2	bacteriophage resistance protein
	redox-regulated ATPase YchF	anti-sigma B factor RsbW	aspartate aminotransferase	glycine/betaine ABC transporter permease	branched-chain amino acid aminotransferase
	serinetRNA ligase	ArsC family protein	ATP synthase F0F1 subunit alpha	histidine kinase	catalase
	sodium/sulfate symporter family protein	aryldialkylphosphata se	ATP synthase F0F1 subunit B	Histidine transport protein (permease)	choline dehydrogenase
	succinate dehydrogenase	aspartate- semialdehyde dehydrogenase	ATP-dependent 6- phosphofructokinase	HslUHslV peptidase ATPase subunit	citrate synthase
		aspartatetRNA ligase	ATP-dependent Clp protease ATP-binding subunit ClpC	hypothetical protein	class II fumarate hydratase
		aspartyl/glutamyl- tRNA amidotransferase subunit A	ATP-dependent Clp protease ATP-binding subunit ClpX	hypothetical protein A4A82_03310	cobalt ABC transporter ATPase

ATP synthase F0F1 subunit alpha	ATP-dependent DNA helicase RecG	hypothetical protein A4A82_05660	cytochrome aa3 quinol oxidase subunit II
ATP synthase F0F1 subunit delta	ATP-dependent Zn metallopeptidase	hypothetical protein A4A82_06485	D-alanineD-alanine ligase
ATP synthase F0F1 subunit gamma	autolysin	hypothetical protein A4A82_08360	dihydrolipoamide succinyltransferase
ATPase P	bifunctional 5,10- methylene- tetrahydrofolate dehydrogenase/5,10- methylene- tetrahydrofolate cyclohydrolase	hypothetical protein A4A82_09985	dihydroxyacetone kinase
ATP-dependent Clp protease ATP- binding subunit ClpC	branched-chain amino acid aminotransferase	hypothetical protein A4A82_10705	dihydroxyacetone kinase subunit L
ATP-dependent DNA helicase RecG	catalase	hypothetical protein A4A82_11275	DNA gyrase subunit A
autolysin	CBS domain- containing protein	hypothetical protein AXY36_03305	DNA primase

bifunctional 5,10- methylene- tetrahydrofolate dehydrogenase/5,10- methylene- tetrahydrofolate cyclohydrolase	cell division protein FtsA	hypothetical protein AXY36_03940	DNA repair protein
bifunctional ornithine acetyltransferase/N- acetylglutamate synthase	chorismate synthase	hypothetical protein AXY36_07550	elastin-binding protein
branched-chain amino acid aminotransferase	citrate synthase	hypothetical protein BSF33_01790	endopeptidase Clp ATP-binding subunit C
Cadmium- transporting ATPase	class II fumarate hydratase	hypothetical protein BSF33_05110	excinuclease ABC subunit C
catalase	cobalamin Fe3+- siderophores ABC transporter periplasmic protein	hypothetical protein BSF33_06845	ferrichrome ABC transporter substrate- binding protein
cell division protein DivIVA	cobalt ABC transporter permease	hypothetical protein BSF33_08575	Fis family transcriptional regulator
chorismate mutase	competence protein ComG	hypothetical protein XA21_08700	formate C-acetyltransferase

citrate synthase class A beta- lactamase	cysteine synthase	hypothetical protein XA21_13400 imidazolonepropion ase	formate tetrahydrofolate ligase gas vesicle protein- protein
cold-shock protein	cytochrome aa3 quinol oxidase subunit I	lactate dehydrogenase	glutamate synthase subunit beta
coproporphyrinogen III oxidase	cytochrome aa3 quinol oxidase subunit II	leucyl aminopeptidase	glutamate-1- semialdehyde aminotransferase
cystathionine gamma synthase	D-alanineD-alanine ligase	lipoyl synthase	glutaminefructose- 6-phosphate aminotransferase
DEAD/DEAH box helicase	deoxyribose- phosphate aldolase	membrane protein	glutaryl-CoA dehydrogenase
dihydrolipoamide succinyltransferase	diaminopimelate decarboxylase	mevalonate kinase	glutathione- dependent formaldehyde dehydrogenase
dihydroxyacetone kinase subunit L	dihydrolipoamide succinyltransferase	molecular chaperone GroEL	glycerol phosphate lipoteichoic acid synthase
DNA methyltransferase	dihydrolipoyl dehydrogenase	NAD-dependent succinate- semialdehyde dehydrogenase	glycine C- acetyltransferase

DNA primase	dihydroxyacetone kinase	NADP-dependent oxidoreductase	glycine cleavage system protein H
DNA-binding protein	DNA methyltransferase	nitric oxide dioxygenase	glycine cleavage system protein T
elastin-binding protein	DNA mismatch repair protein MutS	oligopeptidase PepB	glycine dehydrogenase
endopeptidase Clp ATP-binding subunit C endoribonuclease L- PSP	omega	peptide ABC transporter ATP- binding protein phage tail tape measure protein	glycine dehydrogenase subunit 2 heme-binding protein
Fis family transcriptional regulator	elongation factor P	Phosphate acetyltransferase	HslUHslV peptidase ATPase subunit
formate acetyltransferase	endoribonuclease L- PSP	Phosphoglycerate mutase	hydroxyacid dehydrogenase
formate tetrahydrofolate ligase	F0F1 ATP synthase subunit beta	phosphotransferase	hypothetical protein A4A82_02065
gas vesicle protein- protein	F0F1 ATP synthase subunit gamma	porphobilinogen deaminase	hypothetical protein A4A82_05255
general stress protein	ferrichrome ABC transporter substrate- binding protein	pyridoxal biosynthesis protein	hypothetical protein A4A82_06485
glutamate dehydrogenase	Fis family transcriptional regulator	pyrroline-5- carboxylate reductase	hypothetical protein A4A82_07220

glutamine-binding periplasmic protein	formate C-acetyltransferase	pyruvate phosphate dikinase	hypothetical protein A4A82_08945
glutaminefructose- 6-phosphate aminotransferase	formate dehydrogenase	recombinase A	hypothetical protein A4A82_11260
glutamine- hydrolyzing GMP synthase	formate tetrahydrofolate ligase	recombinase RecB	hypothetical protein A4A82_12170
glutamyl-tRNA amidotransferase	fructose 1,6- bisphosphatase	redox-regulated ATPase YchF	hypothetical protein AXY36_01585
glutaryl-CoA dehydrogenase	gas vesicle protein- protein	restriction endonuclease subunit R	hypothetical protein AXY36_04620
glycerol-3-phosphate cytidylyltransferase	general stress protein	ribonuclease J	hypothetical protein AXY36_07245
glycine dehydrogenase subunit 2	glutamate dehydrogenase	ribonuclease Y	hypothetical protein AXY36_07550
glycosyl transferase family 2	glutamate-1- semialdehyde-2,1- aminomutase	Ribonucleotide reductase of class Ib (aerobic), alpha subunit	hypothetical protein AXY36_07855
heme-binding protein	glutaminefructose- 6-phosphate aminotransferase	ribose ABC transporter permease	hypothetical protein AXY36_08170

	hypothetical protein A4A82_02240	glutaryl-CoA dehydrogenase	S1 RNA-binding protein	hypothetical protein AXY36_11375
	hypothetical protein A4A82_03560	glycerol kinase	serinetRNA ligase	hypothetical protein BSF33_00755
+ 200 + 300 mg	hypothetical protein A4A82_03720	Glycine cleavage system H protein	signal transduction protein TRAP	hypothetical protein BSF33_06845
1 200 1 300 mg	hypothetical protein A4A82_03945	glycine cleavage system protein H	smooth muscle caldesmon	hypothetical protein RES9_00450
	hypothetical protein A4A82_05255	glycine cleavage system protein T	succinate dehydrogenase iron- sulfur subunit	hypothetical protein UF66_2387
	hypothetical protein A4A82_06485	glycine dehydrogenase	succinate- semialdehyde dehydrogenase	hypothetical protein XA21_04180
	hypothetical protein A4A82_07220	glycoprotease	tetrahydrodipicolinat e acetyltransferase	hypothetical protein XA21_13400
	hypothetical protein A4A82_08360	glyoxal reductase	thioredoxin reductase	imidazolonepropion ase
	hypothetical protein A4A82_09265	Hg(II)-responsive transcriptional regulator	uroporphyrinogen decarboxylase	inorganic pyrophosphatase
	hypothetical protein A4A82_10890	Holliday junction resolvase	xylulose kinase	inosine-5- monophosphate dehydrogenase

hypothetical protein A4A82_11275	Homoserine dehydrogenase	iron ABC transporter ATP-binding protein
hypothetical protein AXY36_00980	Homoserine kinase	isoleucinetRNA ligase
hypothetical protein AXY36_02580	hydroxyacid dehydrogenase	lactate dehydrogenase
hypothetical protein AXY36_06600	hypothetical protein	late competence protein ComEC
hypothetical protein AXY36_06780	hypothetical protein A4A82_02240	lipoateprotein ligase
hypothetical protein AXY36_07550	hypothetical protein A4A82_03645	L-Proline/Glycine betaine transporter ProP
hypothetical protein AXY36_07855	hypothetical protein A4A82_03720	malate dehydrogenase
hypothetical protein AXY36_09360	hypothetical protein A4A82_07425	manganese ABC transporter substrate- binding protein
hypothetical protein BSF33_00525	hypothetical protein A4A82_12125	methionine sulfoxide reductase
hypothetical protein BSF33_00755	hypothetical protein A4A82_12320	molecular chaperone GroEL

hypothetical protein RES9_02800	hypothetical protein AXY36_00580	monovalent cation/H+ antiporter subunit D
hypothetical protein XA22_13430	hypothetical protein AXY36_03955	N-acetyltransferase
IMP dehydrogenase	hypothetical protein AXY36_04620	NAD-dependent dehydratase
inorganic pyrophosphatase	hypothetical protein AXY36_06600	nitric oxide dioxygenase
iron ABC transporter substrate-binding protein	hypothetical protein AXY36_09360	NmrA family protein
isoleucinetRNA ligase	hypothetical protein AXY36_09565	oxidoreductase
isoleucyl-tRNA synthetase, partial	hypothetical protein BSF33_05765	peptidase M23
lactate dehydrogenase	hypothetical protein BSF33_09115	peptide ABC transporter ATP- binding protein
late competence protein ComEC	hypothetical protein BSF33_11000	peptide ABC transporter substrate- binding protein

L-lactate dehydrogenase	hypothetical protein RES10_11420	permease
luciferase	hypothetical protein RES11_11325	Phage antirepressor protein
lysinetRNA ligase	hypothetical protein UF66_0386	PhoP family transcriptional regulator
LysR family transcriptional regulator	hypothetical protein UF66_1428	phosphoenolpyruvat e carboxykinase
malonyl CoA-acyl carrier protein transacylase	hypothetical protein XA21_11830	Phosphoglycerate mutase
manganese ABC transporter substrate binding protein	iditol 2- dehydrogenase	phosphopentomutase
methionine sulfoxide reductase	e inorganic pyrophosphatase	phosphotransferase
molecular chaperone GroEL molecular chaperone GrpE nitric oxide dioxygenase	synthase	polynucleotide phosphorylase proline dehydrogenase protease
nucleoside- diphosphate kinase	lactate dehydrogenase	protein rlx

ornithineoxo-acid aminotransferase	leucinetRNA ligase	proteinase
permease	lipid hydroperoxide peroxidase	purine-nucleoside phosphorylase
peroxiredoxin	Major myo-inositol transporter IolT	putative membrane protein
Phosphate acetyltransferase	manganese transporter	pyridine nucleotide- disulfide oxidoreductase
phosphoenolpyruvat e carboxykinase	membrane protein	pyridoxal biosynthesis protein
phosphonate ABC transporter ATP- binding protein	methioninetRNA ligase	pyrimidine- nucleoside phosphorylase
PTS mannnose transporter subunit IIA	molecular chaperone DnaK	pyrroline-5- carboxylate reductase
putative membrane protein	molecular chaperone GroEL	pyruvate phosphate dikinase
pyridoxal biosynthesis protein	nitric oxide dioxygenase	recombinase A

pyruvate phosphate dikinase	non-canonical purine NTP pyrophosphatase	recombinase RecB
pyruvate, phosphate dikinase	nucleoside- diphosphate kinase	ribonuclease H
Repressor CsoR of the copZA operon	ornithineoxo-acid aminotransferase	ribosome-associated protein Y
RNA-binding protein	ornithineoxo-acid transaminase	RNA polymerase sigma factor RpoD
RNA-binding protein S1	peptidase M28	septation ring formation regulator EzrA
serinetRNA ligase	peptide deformylase	serinetRNA ligase
sigma-54 modulation protein	phosphoenolpyruvat e carboxykinase	succinate dehydrogenase flavoprotein subunit
signal recognition particle	phosphoenolpyruvat e-protein phosphotransferase	succinate dehydrogenase iron- sulfur subunit
single-stranded DNA-binding protein	phospholipase D	succinate- semialdehyde dehydrogenase (NADP(+))

succinate dehydrogenase flavoprotein subunit	plasmid partitioning protein ParB	succinyl- diaminopimelate desuccinylase
succinate dehydrogenase iron- sulfur subunit	PP family ATPase	sufurtransferase FdhD
sucrose-6-phosphate hydrolase	preprotein translocase subunit SecA	sugar dehydrogenase
sugar dehydrogenase	prolyl-tRNA synthetase	sulfurtransferase
teichoic acid biosynthesis protein	PTS mannitol transporter subunit IIA	tail protein
TELLIADO TO 11	putativa	4.1.111.1
ThiJ/PfpI family protein	putative acyltransferase	teichoic acid biosynthesis protein
	•	
protein thiol reductase	acyltransferase pyridoxal	biosynthesis protein ThiJ/PfpI family
protein thiol reductase thioredoxin	acyltransferase pyridoxal biosynthesis protein pyruvate, phosphate	biosynthesis protein ThiJ/PfpI family protein transcription termination/antiterm

transcriptional regulator	restriction endonuclease subunit M	two-component response regulator, LytR undecaprenyldiphos pho-
Translation elongation factor Ts	ribosome recycling factor	muramoylpentapepti de beta-N- acetylglucosaminyltr ansferase
translation initiation factor IF-3	RNA pseudouridylate synthase	universal stress protein
tRNA and rRNA cytosine-C5- methylase	Rrf2 family transcriptional regulator	universal stress protein UspA
two-component response regulator, LytR	S- adenosylmethionine tRNA ribosyltransferase	Urease accessory protein UreG
type I glutamate ammonia ligase UDP-glucose 4-	serine dehydratase serinetRNA ligase	uroporphyrinogen decarboxylase
epimerase universal stress protein UTP-glucose-1- phosphate uridylyltransferase	siderophore synthetase signal transduction protein TRAP succinate	
	dehydrogenase	

succinate dehydrogenase flavoprotein subunit

succinate--CoA ligase subunit alpha

succinate--CoA ligase subunit beta succinatesemialdehyde dehydrogenase sulfurtransferase TetR family transcriptional regulator

tetrahydrodipicolinat e acetyltransferase

ThiJ/PfpI family protein

Transcription regulator (contains diacylglycerol kinase catalytic domain)

transcriptional repressor CodY transglycosylase

