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		

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

30S ribosomal protein S4	malate dehydrogenase (acceptor)
30S ribosomal protein S5	oligoribonuclease
30S ribosomal protein S7 30S ribosomal protein S9	purine-nucleoside phosphorylase
4,4'- diaponeurosporenoat e glycosyltransferase	
50S ribosomal protein L11 50S ribosomal protein L13 50S ribosomal protein L14 50S ribosomal protein L15	
50S ribosomal protein L21	
50S ribosomal	
50S ribosomal protein L23	
50S ribosomal protein L25	

30S ribosomal protein S7 5'(3')deoxyribonucleotida se 50S ribosomal protein L10 50S ribosomal protein L13 50S ribosomal protein L19 50S ribosomal protein L24 50S ribosomal protein L25 50S ribosomal protein L29 50S ribosomal protein L7/L12 ABC transporter ATP-binding protein ATP synthase F0F1 CDPglycerol:poly(glycer ophosphate) D-alanine aminotransferase

50S ribosomal protein L27 50S ribosomal protein L30 50S ribosomal 50S ribosomal protein L33 50S ribosomal protein L4 50S ribosomal protein L6 ABC transporter ATP-binding protein ATP-dependent Clp protease proteolytic subunit DNA-directed RNA polymerase subunit beta F0F1 ATP synthase subunit beta

IMP dehydrogenase

DNA-binding protein DNA-directed RNA polymerase subunit beta' F0F1 ATP synthase Fe-S cluster assembly protein SufD **GTPase** Era GTPase ObgE NADH dehydrogenase Oligoendopeptidase F PTS glucose transporter subunit IIA ThiJ/PfpI family protein UDP-glucose 4epimerase

UDP-Nacetylglucosamine 1carboxyvinyltransfer ase beta-ketoacyl-ACP reductase beta-ketoacyl-[acylcarrier-protein] synthase II branched-chain alpha-keto acid dehydrogenase subunit E2 catabolite control protein A catalase cell division protein FtsZ cell division protein GpsB citrate synthase cysteine synthase dihydrolipoyl dehydrogenase

acetyltransferase

aldehyde dehydrogenase

alkaline shock protein 23

aspartate aminotransferase

bifunctional 5,10methylenetetrahydrofolate dehydrogenase/5,10cell division protein DivIVA

citrate synthase

class I fructosebisphosphate aldolase

cold-shock protein

cysteine synthase dihydroxyacetone 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 peroxidase

malate:quinone oxidoreductase

molecular chaperone DnaK

ornithine--oxo-acid aminotransferase

pyruvate kinase

serine hydroxymethyltransf erase single-stranded DNA-binding protein stage V sporulation protein G

succinate--CoA ligase subunit alpha

succinate--CoA ligase subunit beta hypothetical protein A4A82_09265

hypothetical protein A4A82_12170

hypothetical protein AXY36_02415

hypothetical protein AXY36_02580

hypothetical protein AXY36_03955

hypothetical protein AXY36_06600

hypothetical protein AXY36_06780

hypoxanthine phosphoribosyltransf erase malate dehydrogenase (acceptor) malate:quinone oxidoreductase

transaldolase

transcription elongation factor GreA transglycosylase transketolase

translation initiation factor IF-3

trigger factor triose-phosphate isomerase

type I glutamate-ammonia ligase

type I glyceraldehyde-3universal stress protein UspA manganesedependent inorganic pyrophosphatase

metal-dependent hydrolase

methionine sulfoxide molecular chaperone DnaK

oligoribonuclease

peptidase

peptidase M28

phosphoenolpyruvat e-protein phosphotransferase

phosphonate ABC transporter ATP-

pyruvate kinase

pyruvate oxidase pyruvate, phosphate ribosome recycling factor

sigma-54 modulation protein

		stage V sporulation protein G
		succinate dehydrogenase
		flavoprotein subunit
		succinateCoA ligase subunit alpha
		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	
200 mg L-1	protease ATP- binding subunit	
8	ClpX	

peptidase M23	30S ribosomal protein S10 30S ribosomal protein S16	30S ribosomal protein S16 30S ribosomal protein S19	30S ribosomal protein S10 30S ribosomal protein S18	30S ribosomal protein S17 30S ribosomal protein S19	2-oxoisovalerate dehydrogenase 30S ribosomal protein S19
	30S ribosomal protein S9	beta-ketoacyl-[acyl- carrier-protein] synthase II	30S ribosomal protein S9	50S ribosomal protein L1	30S ribosomal protein S4
	50S ribosomal protein L1	class I fructose- bisphosphate aldolase	50S ribosomal protein L15	50S ribosomal protein L11	30S ribosomal protein S8
	50S ribosomal protein L14 50S ribosomal protein L18 50S ribosomal protein L23 50S ribosomal protein L30 50S ribosomal protein L33 DNA-directed RNA polymerase subunit alpha	universal stress protein	50S ribosomal protein L18 50S ribosomal protein L27 50S ribosomal protein L28 50S ribosomal protein L3 50S ribosomal protein L30 50S ribosomal protein L4	50S ribosomal protein L14 50S ribosomal protein L15 50S ribosomal protein L17 50S ribosomal protein L18 50S ribosomal protein L23 50S ribosomal protein L3	30S ribosomal protein S9 50S ribosomal protein L1 50S ribosomal protein L14 50S ribosomal protein L15 50S ribosomal protein L17 50S ribosomal protein L23
	glutamyl-tRNA synthetase		6-phosphogluconate dehydrogenase	50S ribosomal protein L30	50S ribosomal protein L30
	glycinetRNA ligase		alanine racemase	50S ribosomal protein L33	50S ribosomal 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
200 I 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
		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		
2-deoxyribose-5- phosphate aldolase	2-deoxyribose-5- phosphate aldolase	Fis family transcriptional regulator	alpha/beta hydrolase	

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 CDPglycerol: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

100 + 200 mg L-1	Fe-S cluster assembly protein SufB Fe-S cluster assembly protein SufD glucose-6-phosphate isomerase glutamate dehydrogenase glutamine- hydrolyzing GMP synthase glycinetRNA ligase heme-binding protein hydroxymethylpyrim idine/phosphomethyl pyrimidine kinase hypothetical protein A4A82_02240 hypothetical protein A4A82_03560 hypothetical protein A4A82_07220
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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 IIA putative transporter 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				
30S ribosomal protein S14	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
50S ribosomal protein L36	2-oxoisovalerate dehydrogenase	3-hexulose-6- phosphate synthase	2-oxoglutarate dehydrogenase E1	2-deoxyribose-5- phosphate aldolase	30S ribosomal protein S20
acetyltransferase	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
elongation factor Ts	30S ribosomal protein S11	30S ribosomal protein S2	oxobutanoate hydroxymethyltransf erase	2-oxoisovalerate dehydrogenase	30S ribosomal protein S7
hypothetical protein AXY36_09360	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 F0F1 subunit alpha

30S ribosomal protein S5	30S ribosomal protein S6	30S ribosomal protein S16	30S ribosomal protein S16
30S ribosomal protein S6	30S ribosomal protein S8	50S ribosomal protein L1	30S ribosomal protein S18
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
50S ribosomal protein L10	50S ribosomal protein L15	50S ribosomal protein L36	30S ribosomal protein S3
50S ribosomal protein L11	50S ribosomal protein L17	8-amino-7- oxononanoate synthase	30S ribosomal protein S4
50S ribosomal protein L13	50S ribosomal protein L18	ATP synthase F0F1 subunit B	30S ribosomal protein S5
50S ribosomal protein L17	50S ribosomal protein L21	ATP synthase F0F1 subunit delta	30S ribosomal protein S6
50S ribosomal protein L21	50S ribosomal protein L22	DNA polymerase III subunit beta	30S ribosomal protein S7
50S ribosomal protein L22	50S ribosomal protein L23	DNA primase	30S ribosomal protein S8
50S ribosomal protein L25	50S ribosomal protein L24	LysR family transcriptional regulator	30S ribosomal protein S9

ATP-dependent Clp protease proteolytic subunit F0F1 ATP synthase

subunit beta

IMP dehydrogenase

NADH dehydrogenase PTS glucose transporter subunit IIA

aconitate hydratase

alpha/beta hydrolase

beta-ketoacyl-ACP reductase

class I fructosebisphosphate aldolase

dihydrolipoamide succinyltransferase

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
50S ribosomal protein L31	50S ribosomal protein L3	UDP-N-acetyl-D- galactosamine dehydrogenase	50S ribosomal protein L16	elongation factor
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 isomera
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 prote A4A82_02240
ATP-dependent Clp protease proteolytic subunit CDP-	6-phosphogluconate dehydrogenase	catabolite control protein A	50S ribosomal protein L27	hypothetical prote A4A82_07220
glycerol:poly(glycer ophosphate) glycerophosphotrans ferase	ABC transporter ATP-binding protein	chorismate mutase	50S ribosomal protein L29	hypothetical prote AXY36_00580
D-alanine aminotransferase	ATP-dependent Clp protease proteolytic subunit	cysteine desulfurase	50S ribosomal protein L31	hypothetical prote BSF33_05765

DNA gyrase subunit	DNA starvation/stationary	delta-aminolevulinic	50S ribosomal	iron ABC transporte
	phase protection protein	acid dehydratase	protein L5	substrate-binding protein
DNA	-			
starvation/stationary	DNA-binding	dephospho-CoA	50S ribosomal	nitric oxide
phase protection	protein	kinase	protein L6	dioxygenase
protein				
DNA-directed RNA		dihydrolipoamide	50S ribosomal	ornithineoxo-acid
	polymerase subunit	succinyltransferase	protein L7/L12	aminotransferase
beta	alpha	,	I	
F0F1 ATP synthase	DNA-directed RNA	dihydrolipoyl	ABC transporter	
subunit beta	polymerase subunit	dehydrogenase	ATP-binding protein	peroxiredoxin
Fe-S cluster	beta DNA-directed RNA			
assembly protein		elongation factor Tu	ATP synthase F0F1	phosphoglycerate
• •	beta'	cioligation factor fu	subunit alpha	kinase
	Fe-S cluster	endopeptidase Clp	ATP-dependent Clp	
	assembly protein	ATP-binding subunit		phosphopyruvate
• •	SufD	C	subunit	hydratase
~ ~~~		-		
				pyruvate
NADH		endoribonuclease L-	ATP-dependent Zn	dehydrogenase
dehydrogenase	IMP dehydrogenase	PSP	metallopeptidase	(acetyl-transferring)
			1 1	E1 component
				subunit alpha
PTS glucose	NADU		Dalanina	
transporter subunit	NADH	ferritin	D-alanine	pyruvate kinase
IIA	dehydrogenase		aminotransferase	

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	• • •	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	IMP dehydrogenase	

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	pyruvate kinase	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 type I		glycerol kinase
	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 DNA-
	invertase
	trigger factor
	triose-phosphate
	isomerase
	type I glutamate
	ammonia ligase
	type I
	glyceraldehyde-3-
	phosphate
	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-aldehyde dehydrogenase
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	competence protein ComF
hypothetical protein A4A82_08725	50S ribosomal protein L15	50S ribosomal protein L1	catalase	6,7-dimethyl-8- ribityllumazine synthase	glycerol kinase
hypothetical protein A4A82_11275	50S ribosomal protein L16	50S ribosomal protein L11	Chemotaxis- inhibiting protein CHIPS, phage associated	acetateCoA ligase	glycine cleavage system protein T
hypothetical protein AXY36_01150	50S ribosomal protein L19	50S ribosomal protein L14	choline dehydrogenase	acetyl-CoA acetyltransferase	hypothetical protein A4A82_08945
hypothetical protein AXY36_04620	50S ribosomal protein L2	50S ribosomal protein L16	CTP synthase	aconitate hydratase	hypothetical protein XA21_11195
hypothetical protein AXY36_08620	50S ribosomal protein L20	50S ribosomal protein L19	cytochrome aa3 quinol oxidase subunit II	adenylosuccinate synthetase	hypothetical protein XA22_10040
hypothetical protein AXY36_09850	50S ribosomal protein L24	50S ribosomal protein L2	diaminopimelate decarboxylase	alanine dehydrogenase	hypothetical protein XA22_11240

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 5-	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 formyltransferase	acetyl-CoA acetyltransferase	alcohol dehydrogenase	formate C- acetyltransferase	aspartate aminotransferase	sufurtransferase FdhD
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 cytidylate kinase	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 n		hypothetical protein A4A82_03720	Glycine cleavage system H protein	signal transduction protein TRAP	hypothetical protein BSF33_06845
+ 200 + 300 h	lg	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	0	xylulose kinase	inosine-5- monophosphate dehydrogenase

hypothetical protein A4A82_11275	Homoserine dehydrogenase
hypothetical protein AXY36_00980	Homoserine kinase
hypothetical protein AXY36_02580	hydroxyacid dehydrogenase
hypothetical protein AXY36_06600	hypothetical protein
hypothetical protein AXY36_06780	hypothetical protein A4A82_02240
hypothetical protein AXY36_07550	hypothetical protein A4A82_03645
hypothetical protein AXY36_07855	hypothetical protein A4A82_03720
hypothetical protein AXY36_09360	hypothetical protein A4A82_07425
hypothetical protein BSF33_00525	hypothetical protein A4A82_12125
hypothetical protein BSF33_00755	hypothetical protein A4A82_12320

iron ABC transporter ATP-binding protein

isoleucine--tRNA ligase

lactate dehydrogenase

late competence protein ComEC

lipoate--protein ligase

L-Proline/Glycine betaine transporter ProP

malate dehydrogenase

manganese ABC transporter substratebinding protein

methionine sulfoxide reductase

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	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	leucinetRNA
aminotransferase	ligase

permease

peroxiredoxin

Phosphate acetyltransferase

manganese transporter

peroxidase

lipid hydroperoxide

Major myo-inositol

transporter IolT

phosphoenolpyruvat membrane protein e carboxykinase

phosphonate ABC transporter ATPbinding protein

IIA

methionine--tRNA ligase

PTS mannnose molecular chaperone transporter subunit DnaK

putative membrane molecular chaperone protein GroEL

pyridoxal nitric oxide biosynthesis protein dioxygenase proteinase

purine-nucleoside phosphorylase

putative membrane protein

pyridine nucleotidedisulfide oxidoreductase

pyridoxal biosynthesis protein

pyrimidinenucleoside phosphorylase

pyrroline-5carboxylate reductase

pyruvate phosphate dikinase

recombinase A

pyruv dikina	ate phosphate ase	non-canonical purine NTP pyrophosphatase	recombinase RecB
pyruv dikina	ate, phosphate ase	nucleoside- diphosphate kinase	ribonuclease H
	essor CsoR of opZA operon	ornithineoxo-acid aminotransferase	ribosome-associated protein Y
RNA- protei	binding n	ornithineoxo-acid transaminase	RNA polymerase sigma factor RpoD
RNA- protei	binding n S1	peptidase M28	septation ring formation regulator EzrA
serine	etRNA ligase	peptide deformylase	serinetRNA ligase
sigma modu	-54 lation protein	phosphoenolpyruvat e carboxykinase	succinate dehydrogenase flavoprotein subunit
signal partic	recognition le	phosphoenolpyruvat e-protein phosphotransferase	succinate dehydrogenase iron- sulfur subunit
-	-stranded -binding n	phospholipase D	succinate- semialdehyde dehydrogenase (NADP(+))

succinate dehydrogenase flavoprotein subunit	plasmid partitioning protein ParB
succinate dehydrogenase iron- sulfur subunit	PP family ATPase
sucrose-6-phosphate hydrolase	preprotein translocase subunit SecA
sugar dehydrogenase	prolyl-tRNA synthetase
teichoic acid biosynthesis protein	PTS mannitol transporter subunit IIA
ThiJ/PfpI family protein	putative acyltransferase
thiol reductase thioredoxin	pyridoxal biosynthesis protein
thioredoxin	pyruvate, phosphate dikinase
thymidine phosphorylase	recombinase A
TIGR00268 family protein	response regulator SirA

succinyldiaminopimelate desuccinylase

sufurtransferase FdhD

sugar dehydrogenase

sulfurtransferase

tail protein

teichoic acid biosynthesis protein

ThiJ/PfpI family protein

transcription termination/antiterm ination protein NusG

transglycosylase

translation initiation factor IF-2

transcriptional regulator

restriction endonuclease subunit M

Translationribosome recyclingelongation factor Tsfactor

translation initiation factor IF-3 tRNA and rRNA cytosine-C5methylase two-component response regulator, LytR

type I glutamate-ammonia ligase UDP-glucose 4epimerase universal stress protein UTP-glucose-1phosphate uridylyltransferase RNA pseudouridylate synthase Rrf2 family transcriptional regulator Sadenosylmethionine tRNA ribosyltransferase serine dehydratase

serine--tRNA ligase

siderophore synthetase

signal transduction protein TRAP

succinate dehydrogenase two-component response regulator, LytR undecaprenyldiphos phomuramoylpentapepti de beta-Nacetylglucosaminyltr ansferase

universal stress protein

universal stress protein UspA

Urease accessory protein UreG

uroporphyrinogen decarboxylase 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

transposase, IS605 OrfB family
transposase, partial
triose-phosphate
isomerase
tRNA-binding
protein
type I glutamate
ammonia ligase
universal stress
protein UspA
uracil
phosphoribosyltransf
erase