Common	N Entro- ID	Gono Namo	Prediction	Sorted Catagory
yjeK	948662	Uncharacterized kamA family protein yjeK	Prediction ['metabolic process']	Sorted Category Down Regulated
			[metabolic process]	
dxr		1-deoxy-D-xylulose 5-phosphate reductoisomerase 60 kDa chaperonin		Temporarly Down Repown Regulated
groL				
ribF		Riboflavin biosynthesis protein ribF		Down Regulated
kch		Putative potassium channel protein	Illumposition DNA seleted Type of regulation Appropriation Institute of transcription DNA dependent	Down Regulated
ydhB fonC	945208	Uncharacterized HTH-type transcriptional regulator ydhB	[Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent']	Up Regulated
fepG	945209	Ferric enterobactin transport system permease protein fepG	n	Ambiguous/Unsorted
ybaA	948863	Uncharacterized protein ybaA	III	Up Regulated
ydcl	948865	Uncharacterized HTH-type transcriptional regulator ydcl	[Transcription', 'RNA related', 'Type of regulation', 'Genetic unit regulated', 'transcription', 'regulation of transcription DNA dependence of transcription	
btuR	945839	Cob(I)yrinic acid a,c-diamide adenosyltransferase		Down Regulated
hsdS	948867	Type-1 restriction enzyme EcoKI specificity protein Uncharacterized oxidoreductase yciK	EDNIA reneid legenere to DNIA demons etimulus!	Temporarly Down Re
yciK	945838		['DNA repair', 'response to DNA damage stimulus']	Ambiguous/Unsorted
creD	948868	Inner membrane protein creD		Ambiguous/Unsorted
cstA	945213	Carbon starvation protein A		Ambiguous/Unsorted
rbsK	948260	Ribokinase		Down Regulated
ychE	945836	UPF0056 membrane protein yhcE		Temporarly Down Re
yjiN	948860	Uncharacterized protein yjiN	['response to antibiotic']	Up Regulated
sugE	948671	Quaternary ammonium compound-resistance protein sugE		Down Regulated
tauD	945021	Alpha-ketoglutarate-dependent taurine dioxygenase		Down Regulated
dinl		DNA-damage-inducible protein I		Temporarly Up Regul
rcsF	949113	Protein rcsF		Down Regulated
yaeB	949112	UPF0066 protein yaeB		Ambiguous/Unsorted
psd	948673	Phosphatidylserine decarboxylase proenzyme		Ambiguous/Unsorted
poxA		Putative lysyl-tRNA synthetase		Up Regulated
abgR	945831	HTH-type transcriptional regulator abgR		Ambiguous/Unsorted
yqeG	945028	Inner membrane transport protein yqeG		Ambiguous/Unsorted
yhbV	949117	Uncharacterized protein yhbV	['Coenzyme transport and metabolism', ' Energy metabolism carbon ', ' Building block biosynthesis', 'ion transport', 'iron ion transport'	
yjeM	948679	Inner membrane transporter yjeM		Up Regulated
yieL	948272	Uncharacterized protein yieL	['Carbohydrate transport and metabolism']	Up Regulated
yjeP	948676	Uncharacterized mscS family protein yjeP	['purine nucleotide biosynthetic process', 'biosynthetic process']	Down Regulated
ilvE		Branched-chain-amino-acid aminotransferase		Down Regulated
il∨M	948279	Acetolactate synthase isozyme 2 small subunit		Down Regulated
tauC	945026	Taurine transport system permease protein tauC		Down Regulated
tauB	945027	Taurine import ATP-binding protein tauB		Down Regulated
ilvD	948277	Dihydroxy-acid dehydratase		Down Regulated
оррА	945830	Periplasmic oligopeptide-binding protein		Down Regulated
thrB	947498	Homoserine kinase		Down Regulated
ylaC	948876	Inner membrane protein ylaC		Up Regulated
ampD	948877	1,6-anhydro-N-acetylmuramyl-L-alanine amidase ampD		Ambiguous/Unsorted
leuS	947497	Leucyl-tRNA synthetase		Temporarly Down Re
dkgA	947495	2,5-diketo-D-gluconic acid reductase A		Temporarly Up Regul
tatE	945228	Sec-independent protein translocase protein tatE		Down Regulated
marR	945825	Multiple antibiotic resistance protein marR		Up Regulated
ybgl	945824	UPF0135 protein ybgl		Ambiguous/Unsorted
dacA	945222	D-alanyl-D-alanine carboxypeptidase dacA		Down Regulated
ahpC	945225	Alkyl hydroperoxide reductase subunit C		Temporarly Up Regul
yqhC	947491	Uncharacterized HTH-type transcriptional regulator yqhC	['RNA related', 'Type of regulation', 'Genetic unit regulated', 'Protection', 'regulation of transcription DNA dependent']	Up Regulated
dsbG	945224	Thiol:disulfide interchange protein dsbG		Down Regulated
yjjL	948879	Inner membrane transport protein yjjL		Up Regulated
ybeD	945220	UPF0250 protein ybeD		Ambiguous/Unsorted
hsdM	948872	Type I restriction enzyme EcoKI M protein		Down Regulated
fecR	949104	Protein fecR		Ambiguous/Unsorted
tauA	945030	Taurine-binding periplasmic protein		Down Regulated
dsdX	949103	DsdX permease		Up Regulated
fumB		Fumarate hydratase class I, anaerobic		Up Regulated
yail		UPF0178 protein yail		Down Regulated
atpF		ATP synthase subunit b; ATP synthase B chain		Down Regulated
mnmG	948248	tRNA uridine 5-carboxymethylaminomethyl modification enzy	me mnmG	Ambiguous/Unsorted
atpG		ATP synthase gamma chain		Down Regulated
atpD		ATP synthase subunit beta		Down Regulated
ptrA		Protease 3		Temporarly Down Re
atpC		ATP synthase epsilon chain		Down Regulated
glmU		Bifunctional protein glmU		Down Regulated
recB		Exodeoxyribonuclease V beta chain		Down Regulated
. 505	10 17 200	=		- Strii i togulatea

ynfH	945822	Anaerobic dimethyl sulfoxide reductase chain ynfH	Down Regulated
RecD	947287	Exodeoxyribonuclease V alpha chain	Down Regulated
oppB	945823	Oligopeptide transport system permease protein oppB	Down Regulated
fimA	948838	Type-1 fimbrial protein, A chain	Up Regulated
nanM	949106	N-acety/neuraminate epimerase	Up Regulated
glmS	948241	Glucosaminefructose-6-phosphate aminotransferase [isomerizing]	Down Regulated
_	947289	Amino-acid acetyltransferase	Down Regulated
argA atpA	948242	ATP synthase subunit alpha	Down Regulated
Dos	945815	Heme-regulated cyclic di-GMP, cyclic AMP phosphodiesterase,	Temporarly Up Regul
dosP	945815	Heme-regulated cyclic di-GMP, cyclic AMP phosphodiesterase,	Temporarly Up Regul
	947480	Uncharacterized protein yghZ	Temporarly Up Regul
gpr	947480	Uncharacterized protein yghZ	
yghZ fimU	948847	Orionaracterized protein ygriz. Protein fimH	Temporarly Up Regul Temporarly Down Regul
fimH ygiT	945814	Uncharacterized HTH-type transcriptional regulator ygiT [' Carbon compound utilization', 'metabolic process']	Up Regulated
	945814		
mqsA gntP	948848	Uncharacterized HTH-type transcriptional regulator ygiT	Up Regulated
		High-affinity gluconate transporter Recal hady and modification protein flat.	Up Regulated
flgD	945813	Basal-body rod modification protein flgD	Down Regulated
rfbC	947482	dTDP-4-dehydrorhamnose 3,5-epimerase	Down Regulated
yfdY	948842	Uncharacterized protein yfdY []	Up Regulated
uidB	947484	ECIAI39_1442; EcHS_A1691; APECO1_699; b1616; ECDH10B_1749; c2008; UTI89_C1804; EcSMS35_1583; EcE24377A_1824; Z2620; ECS88_1662; ECED1_1815; ECUMN_1905; ECH7	
yciU	945819	UPF0263 protein yciU	Down Regulated
rlmH	945239	Ribosomal RNA large subunit methyltransferase H	Temporarly Down Re
mrdB	945238	Rod shape-determining protein rodA	Down Regulated
flhC	947280	Flagellar transcriptional activator flhC	Temporarly Up Regul
ybeB	945237	Uncharacterized protein ybeB []	Down Regulated
sgcX	948840	Putative sgc region protein sgcX	Up Regulated
fadR	948652	Fatty acid metabolism regulator protein	Down Regulated
psiF	945040	Phosphate starvation-inducible protein psiF	Up Regulated
ygdD	947279	UPF0382 inner membrane protein ygdD	Temporarly Up Regul
yjdC	948650	HTH-type transcriptional regulator yjdC	Temporarly Up Regul
phoR	945044	Sensor protein; Phosphate regulon sensor protein phoR	Temporarly Down Re
aroM	945045	Protein aroM Protein aroM	Temporarly Down Re
asnC	948259	Asparaginyl-tRNA synthetase; Regulatory protein asnC	Down Regulated
fucO	947273	Lactaldehyde reductase Lactaldehyde reductase	Temporarly Up Regul
csdE	947274	Uncharacterized sufE-like protein ygdK	Ambiguous/Unsorted
yaiE	945048	UPF0345 protein yaiE	Down Regulated
atpH	948254	ATP synthase subunit delta	Down Regulated
aspA	948658	Aspartate ammonia-lyase	Temporarly Up Regul
atpB	948252	ATP synthase subunit a	Down Regulated
fxsA	948657	UPF0716 protein fxsA	Temporarly Up Regul
yjeH	948656	Inner membrane protein yjeH	Down Regulated
atpE	948253	ATP synthase subunit c	Down Regulated
oppC	945810	Oligopeptide transport system permease protein oppC	Down Regulated
rsmG	948250	Ribosomal RNA small subunit methyltransferase G	Down Regulated
groS	948655	10 kDa chaperonin	Down Regulated
uxuR	948849	Uxu operon transcriptional regulator	Ambiguous/Unsorted
thrA	945803	Bifunctional aspartokinase/homoserine dehydrogenase 1	Temporarly Up Regul
oppD	945802	Oligopeptide transport ATP-binding protein oppD	Down Regulated
holE	947471	DNA polymerase III subunit theta	Temporarly Down Re
nadD	945248	Nicotinate-nucleotide adenylyltransferase; Probable nicotinate-nucleotide adenylyltransferase	Down Regulated
gss	947474	Bifunctional glutathionylspermidine synthetase/amidase	Temporarly Down Re
	947474	Bifunctional glutathionylspermidine synthetase/amidase	Temporarly Down Re
gsp yjiE	948852	Uncharacterized HTH-type transcriptional regulator yjiE ['Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent']	Ambiguous/Unsorted
glcG	947473	Protein glcG	Temporarly Up Regul
yqhA	947479	UPF0114 protein yqhA	Down Regulated
rlpA	945241	Rare lipoprotein A	Down Regulated
mrdA	945240	Penicillin-binding protein 2	Down Regulated
ygdH	947266	UPF0717 protein ygdH	Down Regulated
gudX	947261	Glucarate dehydratase-related protein	Up Regulated
rumB	947260	23S rRNA (uracil-5-)-methyltransferase rumB	Temporarly Down Re
		Flavoprotein wrbA; Putative flavoprotein wrbA	Temporarly Up Regul
lw/rhΔ		i idvoprotoin wibri, i ututivo navoprotoin wibri	In comporating up regul
wrbA	947263	Uncharacterized protein viel	Tomporarly Down Da
yieE	948226	Uncharacterized protein yieE ['Amino acid transport and metabolism']	Temporarly Down Re
yieE yieF	948226 948225	Uncharacterized protein yieF	Temporarly Down Re
yieE	948226		

djlC	945253	Uncharacterized J domain-containing protein djlC	Up Regulated
fhuC	945250	Iron(3+)-hydroxamate import ATP-binding protein fhuC	Temporarly Up Regul
serA	945258	D-3-phosphoglycerate dehydrogenase	Down Regulated
adiA	948638	Biodegradative arginine decarboxylase	Up Regulated
dcuS	948639	Sensor protein dcuS	Temporarly Down Re
phoU	948233	Phosphate transport system protein phoU	Down Regulated
melR	948637	Melibiose operon regulatory protein	Up Regulated
glcA	947259	Glycolate permease glcA	Temporarly Up Regul
yhjA	948038	Probable cytochrome c peroxidase	Up Regulated
miaB	945260	(Dimethylallyl)adenosine tRNA methylthiotransferase miaB	Down Regulated
ubif	945261	2-octaprenyl-3-methyl-6-methoxy-1,4-benzoquinol hydroxylase	Up Regulated
yhjE	948032	Inner membrane metabolite transport protein yhjE	Temporarly Up Regul
yhjD	948033	Inner membrane protein yhjD	Temporarly Up Regul
eptA	948629	ECIAI1_4344; E2348C_4442; ECUMN_4646; ECS88_4616; b4114; ECED1_4848; ECIAI39_4538; EC55989_4605; ECDH10B_4306	Ambiguous/Unsorted
yhjC	948035	Uncharacterized HTH-type transcriptional regulator yhjC ['Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent']	Up Regulated
ibpA	948200	Small heat shock protein ibpA	Up Regulated
yecJ	946388	Uncharacterized protein yecJ	Down Regulated
yidX CYBB	948202 947241	Uncharacterized protein yidX [' Type of regulation', 'proteolysis', 'signal transduction'] Cytochrome b561	Up Regulated
			Down Regulated
torZ	946389	Trimethylamine-N-oxide reductase 2	Ambiguous/Unsorted
rumA	947243	23S rRNA (uracil-5-)-methyltransferase rumA	Temporarly Down Re
nudB	946383	Dihydroneopterin triphosphate pyrophosphatase	Down Regulated
recF	948209	DNA replication and repair protein recF	Temporarly Down Re
yecR	946386	Uncharacterized protein yecR ['Inorganic ion transport and metabolism', 'Adaptation to stress', 'iron ion transport']	Down Regulated
cmoB	946387	tRNA (mo5U34)-methyltransferase	Down Regulated
pgm	945271	Phosphoglucomutase	Ambiguous/Unsorted
pitA	948009	Low-affinity inorganic phosphate transporter 1	Down Regulated
seqA	945272	Protein seqA	Temporarly Down Re
tsr	948884	Methyl-accepting chemotaxis protein I	Up Regulated
yhiQ	948005	UPF0341 protein yhiQ	Temporarly Up Regul
rfbB	945276		Down Regulated
erfK	945273	Protein erfK/srfK	Temporarly Up Regul
uspA	948007	Universal stress protein A	Up Regulated
yhil	948003	Uncharacterized protein yhil [' Protection']	Up Regulated
yahB	945278	Uncharacterized HTH-type transcriptional regulator yahB [Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent']	Temporarly Down Re
cusR	945003	Transcriptional regulatory protein cusR	Down Regulated
yael	945002		Down Regulated
opgG	945005	Glucans biosynthesis protein G	Down Regulated
gyrB	948211	DNA gyrase subunit B	Down Regulated
pal	945004	Peptidoglycan-associated lipoprotein	Down Regulated
lacl	945007	Lactose operon repressor	Up Regulated
trmJ	948610	tRNA (cytidine/uridine-2'-O-)-methyltransferase trmJ	Temporarly Down Re
cysl	947231	Sulfite reductase [NADPH] hemoprotein beta-component	Down Regulated
hisQ	947235	Histidine transport system permease protein hisQ	Down Regulated
dnaN	948218	DNA polymerase III subunit beta	Down Regulated
dnaA	948217		Down Regulated
rpmH	948216	50S ribosomal protein L34	Down Regulated
yidC	948214		Down Regulated
osmY	948895	Osmotically-inducible protein Y	Temporarly Up Regul
asnB	945281	Asparagine synthetase B [glutamine-hydrolyzing]	Down Regulated
fhuF	948891	Ferric iron reductase protein fhuF	Temporarly Up Regul
flhA	946390	Flagellar biosynthesis protein flhA	Down Regulated
ybfF	945288	Esterase ybfF	Ambiguous/Unsorted
nagA	945289	N-acetylglucosamine-6-phosphate deacetylase	Ambiguous/Unsorted
yjjĞ	948899	5'-nucleotidase yjjG	Temporarly Down Re
mrr	948898	Mrr restriction system protein	Temporarly Up Regul
prfC	948897		Down Regulated
	948066	Inner membrane protein yhjX	Up Regulated
VIII		Dipeptide transport system permease protein dppC	Down Regulated
yhjX dppC	948064		
dppC	948064 946170		Temporarly Down Re
dppC valV		c5541; Z2692; UTI89_C1856; ECDH10B_1799; b1665; E2348C_tRNA31; ECs5461	Temporarly Down Re Temporarly Down Re
dppC valV eptB	946170 948068	c5541; Z2692; UTI89_C1856; ECDH10B_1799; b1665; E2348C_tRNA31; ECs5461 b3546; ECS88_3965; ECIAI39_4054; EC55989_3997; E2348C_3793; ECED1_4224; ECUMN_4058; ECDH10B_3725; ECIAI1_3705	Temporarly Down Re
dppC valV eptB yfeA	946170 948068 946864	c5541; Z2692; UTI89_C1856; ECDH10B_1799; b1665; E2348C_tRNA31; ECs5461 b3546; ECS88_3965; ECIAl39_4054; EC55989_3997; E2348C_3793; ECED1_4224; ECUMN_4058; ECDH10B_3725; ECIAl1_3705 Uncharacterized protein yfeA [Signal transduction mechanisms', 'Type of regulation', 'two component signal transduction system phosphorelay', 'regulation of t	Temporarly Down Re Temporarly Down Re
dppC valV eptB	946170 948068	c5541; Z2692; UTI89_C1856; ECDH10B_1799; b1665; E2348C_tRNA31; ECs5461 b3546; ECS88_3965; ECIAI39_4054; EC55989_3997; E2348C_3793; ECED1_4224; ECUMN_4058; ECDH10B_3725; ECIAI1_3705	Temporarly Down Re

ا ماله ا	046477	Dutativa managa wanaga udh D		Davis Dagid-t
ydhR		Putative monooxygenase ydhR		Down Regulated
ydgA	946172	Protein ydgA	1	Temporarly Up Regul
ynhG	946174	Uncharacterized protein ynhG		Temporarly Up Regul
sufE		Cysteine desulfuration protein sufE		Temporarly Up Regul
gabD		Succinate-semialdehyde dehydrogenase [NADP+]		Temporarly Up Regul
yiaD	948075	Inner membrane lipoprotein yiaD		Temporarly Up Regul
glyQ		Glycyl-tRNA synthetase alpha subunit		Down Regulated
yiaA	948078	Inner membrane protein yiaA		Up Regulated
ptsl		Phosphoenolpyruvate-protein phosphotransferase	The force and beginned I Drivery Active Topper dated I Destroited	Down Regulated
yfeK	946876		'Defense mechanisms', ' Primary Active Transporters', ' Protection']	Up Regulated
cysK		Cysteine synthase A; Cysteine synthase	The table is a first of the company	Down Regulated
ydhL	946168	Uncharacterized protein ydhL	Metabolism of other compounds', 'purine nucleotide biosynthetic process']	Up Regulated
plsX		Fatty acid/phospholipid synthesis protein plsX		Down Regulated
nemA		N-ethylmaleimide reductase		Ambiguous/Unsorted
pdxY		Pyridoxamine kinase	UDNA related UTura of acculation LAdoptation to strong Managaritical Jacquistics of transposition DNA dependent Jacquistics	Down Regulated
yiaG		Uncharacterized HTH-type transcriptional regulator yiaG	'RNA related', 'Type of regulation', 'Adaptation to stress', 'transcription', 'regulation of transcription DNA dependent', 'response to	
yiaF	948072		'Carbohydrate transport and metabolism', 'Carbon compound utilization', 'phosphoenolpyruvate dependent sugar phosphotransfel	
yhjK			CIAI1_3680; b3529; ECIAI39_4032; ECED1_4207; ECS88_3943; ECUMN_4036; UTI89_C4061; Z4943; E2348C_3771	Ambiguous/Unsorted
bcsZ		Endoglucanase		Down Regulated
bcsB	948045	Cyclic di-GMP-binding protein		Down Regulated
ybeZ		PhoH-like protein		Temporarly Up Regul
pdxK		Pyridoxine kinase		Up Regulated
kdgK		2-dehydro-3-deoxygluconokinase		Temporarly Up Regul
cysU		Sulfate transport system permease protein cysT		Down Regulated
cysP		Thiosulfate-binding protein		Down Regulated
ptsH		Phosphocarrier protein HPr		Down Regulated
yfeS	946887	Uncharacterized protein yfeS	<u>I</u>	Down Regulated
cysM		Cysteine synthase B; Cysteine synthase		Down Regulated
rsxG		Electron transport complex protein rnfG		Down Regulated
leuE	946157	Leucine efflux protein		Down Regulated
yhjJ	948040	Protein yhjJ		Down Regulated
hdhA	946151	7-alpha-hydroxysteroid dehydrogenase		Temporarly Up Regul
murC	946153	UDP-N-acetylmuramateL-alanine ligase		Down Regulated
yhjV	948057	Inner membrane transport protein yhjV		Temporarly Down Re
csiR	948055	Uncharacterized HTH-type transcriptional regulator ygaE		Ambiguous/Unsorted
prpE		PropionateCoA ligase		Up Regulated
cyoC		Cytochrome o ubiquinol oxidase subunit 3		Down Regulated
mntH		Manganese transport protein mntH		Up Regulated
uidA		Beta-glucuronidase		Up Regulated
rsxB		Electron transport complex protein rnfB		Temporarly Down Re
topB		DNA topoisomerase 3		Down Regulated
pntB		NAD(P) transhydrogenase subunit beta		Down Regulated
tqsA		Al-2 transport protein tqsA		Temporarly Down Re
fadL	946820	Long-chain fatty acid transport protein		Temporarly Up Regul
fecE		Fe(3+) dicitrate transport ATP-binding protein fecE		Down Regulated
dnaJ		Chaperone protein dnaJ		Down Regulated
dnaK		Chaperone protein dnaK		Temporarly Up Regul
aroC		Chorismate synthase		Down Regulated
sixA		Phosphohistidine phosphatase sixA		Up Regulated
fecD		Fe(3+) dicitrate transport system permease protein fecD	2014 avatain in A. Inspetion alamost IC4 avatain in A. Inspetion also at IC4 describing D. Inspetion also at IC4 4 170	Down Regulated
insB	944743		2/3/4 protein insA; Insertion element IS1 protein insA; Insertion element IS1 4 protein insB; Insertion element IS1 1/5/6 protein insB	
sspA		Stringent starvation protein A		Down Regulated
nfrA		Bacteriophage N4 adsorption protein A		Ambiguous/Unsorted
efeO		UPF0409 protein ycdO		Temporarly Up Regul
putP		Sodium/proline symporter		Down Regulated
yceA		UPF0176 protein yceA	Transfer I DNA state of I Transfer and Francisco I and	Down Regulated
ypfG	946828		'Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent']	Temporarly Down Re
evgS		Sensor protein evgS; Sensor protein		Temporarly Down Re
BAX	948089	Protein bax		Temporarly Down Re
ydil		Esterase ydil		Down Regulated
malS		Alpha-amylase		Down Regulated
sapB		Peptide transport system permease protein sapB		Temporarly Up Regul
xylR		Xylose operon regulatory protein		Up Regulated
ygcE		Uncharacterized sugar kinase ygcE	" Central intermediary metabolism', 'carbohydrate metabolic process']	Temporarly Up Regul
yiaJ	948084	HTH-type transcriptional regulator yiaJ		Ambiguous/Unsorted

Selder Processing Process	ydiH	946194	Uncharacterized protein ydiH	['proteolysis']	Temporarly Down Re
Metalog Metalog Ambiguoscharged protein yell Ambiguoscharged protein Ambiguoscharged Ambiguoscharged Ambiguoscharged Donn Regulated Donn Regulated New York Metalog Metalog Donn Regulated Preparative	_			[procession]	
98 Seldon G. Stycy IR-Not symbotise den suburit V. 19429 N. 19429				['Amino acid transport and metabolism', ' Building block biosynthesis', 'amino acid biosynthetic process']	
Methods Miles Mi	-	_		, , , , , , , , , , , , , , ,	
984199 Transcriptional registatory proteins Doon Regulated Temporary by Pequided Temporary by Pequided Temporary by Pequided Personal P					
948837 D-seem dehydratese:	rstA	946199			
See330 Fe/13 (distrate binding periplayants potent factor (Fig. RNA polymerase signification of the periplayant factor (Fig. RNA polymerase) Fig.	ilvL	948283			Temporarly Up Regul
See330 Fe/13 (distrate binding periplayants potent factor (Fig. RNA polymerase signification of the periplayant factor (Fig. RNA polymerase) Fig.	dsdA	946837	D-serine dehydratase; D-serine dehydratase 1		Ambiguous/Unsorted
yiPP 44955 Uncharacterized protein yiPP (Regiozation recombination and rigidary) Vic. S48655 Uncharacterized protein yiPC 1 Vic. S48655 Uncharacterized protein yiPC 2 Vic. S48655 Uncharacterized	fecB	946838	Fe(3+) dicitrate-binding periplasmic protein		
MeSS7 Uncharacterated protein yeC 0 Depressive		946839	Probable RNA polymerase sigma factor fecl; RNA polymeras	se sigma factor	Up Regulated
Section Sect	yjhP		Uncharacterized protein yjhP	['Replication recombination and repair']	Up Regulated
Metabook AFF-dependent FNA helicose mile Down Regulated Down Regul	yfeC				
September Sept				['Energy production and conversion']	
yiP 948912 Inner membrane protein yiB Ambiguous Unsorted Down Regulated Work E 94829 Putative EAA polymerase Down Regulated Work EAA polymerase Down Regulated Down Regulat					
Medical Medical Arthogenedin DNA Indicase rep Down Regulated Dow					
March Section Down Regulated Dow		_			
## H 94229 Guesse-t-phosphate thymidy/transferase 2 Down Regulated					
Suff Schedule Sufficience				U state the wright through a second	
jork 94153 PFO118 inner membrane protein ydlk Ambiguoust/Insorted Protein yasses				priate triymioyiytiransierase z	
Post					
yine E 946188 Protein yine E Down Regulated Down Regulated Protein yine E Down Regulated Protein yine E Down Regulated Protein yine E Down Regulated Down Regulated Protein yine E Down Regulated Down Regulated Protein yine E Down Regulated Down Re		_	Transposase insG for insertion sequence element IS4		
Wex 94824 Protein wax Protein wax National Content Protein P			I IDE0187 protein yeaF		
Section Sect					
Season					
FifA S48296 Lipopolysacchande biosynthesis protein rffA Down Regulated appl S49132 Phosphotephotes isomerase Ambiguous/Unsorded appl S49132 Phosphotephotes isomerase Up Regulated appl S49132 Phosphotephotes isomerase Up Regulated S49130 Protein-sept membrane protein spcD Down Regulated S49130 Oueune iRNA-ribosyntrasiferase Down Regulated					
IpcA 949134 Phosphoheplose isomerase Ambiguous/Unsorted appH 949132 Acyl carrier protein phosphodiesterase Up Regulated SecD 949133 Protein-export membrane protein secD Down Regulated Down					
sept 949132 Ayd carrier protein phosphodiesterase Up Regulated					
SecD 949133 Protein-export membrane protein secD Down Regulated					
Down Regulated Down					
malZ 949131 Maltodextrin glucosidase					
Dum Dum Dum Dum Dum Dum Dum Dum Regulated Euro Dum Regulated Puri Dum Regulated Puri Dum Regulated Puri Pu					
Euro					
yadD 94781 Uncharacterized protein yadD [Replication recombination] [Temporarly Up Regul ydM 945644 Incremembrane protein ydM Up Regulated araC 94780	trpA	946204			Down Regulated
ydjM 945844 Inner membrane protein ydjM Up Regulated araC 94780 Arbinose open negulatory protein Uncharacterized protein ydiE (Amino acid transport and metabolism', 'Building block biosynthesis', 'metabolic process', 'biosynthetic process', 'aromatic amino a Up Regulated BlgL 945864 Juncharacterized protein yau0.' Putative phosphotransferase [] Ambiguous/Unsorted pps8 946207 Uncharacterized protein yau0.' Putative phosphotransferase gldA Ambiguous/Unsorted BaseR 946207 Uncharacterized protein yau0.' Putative phosphotransferase ydiA Ambiguous/Unsorted BaseR 946607 Uncharacterized protein yau0.' Putative phosphotransferase ydiA Ambiguous/Unsorted BaseR 946607 Uncharacterized protein yau0.' Putative phosphotransferase ydiA Ambiguous/Unsorted BaseR 946607 Uncharacterized protein yau0.' Putative phosphotransferase ydiA Ambiguous/Unsorted BaseR 946607 Uncharacterized protein yau0.' Putative phosphotransferase ydiA Ambiguous/Unsorted BaseR 946607 Uncharacterized protein yau0.' Putative phosphotransferase ydiA Dawn Regulated Up Regulated WibiG 946607 architecture 93 -methythythinane 93 -methythythinane 93 -methythythinane 94 Dawn Regulated Up Regulated UpP-N-acetylglucosamine 2-epimerase [] Emporary Down Regulated YbiG 946607 architecture ydia ydia ydia ydia ydia ydia ydia ydia	leuD	945642	3-isopropylmalate dehydratase small subunit		Down Regulated
Arabinose operon regulatory protein Up Regulated Arabinose operon regulatory protein Up Regulated Arabinose operon regulatory protein Up Regulated Arabinose operon regulatory protein Arabinose operon regulatory Arabinose		944781	Uncharacterized protein yadD	['Replication recombination and repair', 'DNA recombination']	Temporarly Up Regul
Valie	ydjM	945644	Inner membrane protein ydjM		Up Regulated
figL 945646 Flagellar hook-associated protein 3 Down Regulated ydiA 945207 Uncharacterized protein yuaU; Putative phosphotransferase [] Ambiguous/Unsorted pps 945207 Uncharacterized protein yuaU; Putative phosphotransferase diA Ambiguous/Unsorted fligG 945647 Flagellar basal-body rod protein fligG Down Regulated Pack Pack Pack Pack Pack Pack Pack Pack					Up Regulated
ydia 946207 Uncharacterized protein yuaU; Putative phosphotransferase [] Ambiguous/Unsorted ppsR 946207 Uncharacterized protein yuaU; Putative phosphotransferase ydiA Ambiguous/Unsorted DaeR 946607 Flagellar basal-body rod protein figG Down Regulated DaeR 946607 Transcriptional regulatory protein baeR Up Regulated Up Regulated Up Regulated Up Regulated (ffE 944789 UpP-N-acetylglucosamine 2-epimerase UpD-N-acetylglucosamine	-			['Amino acid transport and metabolism', ' Building block biosynthesis', 'metabolic process', 'biosynthetic process', 'aromatic amino a	
ppsR 946207 Uncharacterized protein yuaU; Putative phosphotransferase ydiA Ambiguous/Unsorted baeR 946605 Flagellar basal-body rod protein figG Down Regulated Up Regula					
Figg 945647 Fiagellar basal-body rod protein ftg Down Regulated					
baeR 946605 Transcriptional regulatory protein baeR Up Regulated				ydiA	
LibiG 946607 3-demethylubiquinone-9 3-methylytransferase Down Regulated Temporarly Down Regulated Temporarly Down Regulated September			Flagellar basal-body rod protein flgG		
fffE 944789 UDP-N-acetylglucosamine 2-epimerase Temporarly Down Re ydiB 946200 Quinate/shikimate dehydrogenase Temporarly Down Re mrdA 946612 Ribonucleoside-diphosphate reductase; Ribonucleoside-diphosphate reductase 1 subunit alpha Down Regulated ybiS 945441 Uncharacterized protein ybiS ['phosphoenolpyruvate dependent sugar phosphotransferase system'] Ambiguous/Unsorted bya601 Uncharacterized HTH-type transcriptional regulator ydcQ Up Regulated ydcQ 946011 Uncharacterized HTH-type transcriptional regulator ydcQ ['Coenzyme transport and metabolism'] Up Regulated figH 945033 Soxoacyl-[acyl-carrier-protein] synthase 3 Down Regulated ydcR 946004 Uncharacterized HTH-type transcriptional regulator ydcR ['Coenzyme transport and metabolism'] Down Regulated ydcR 945034 Glutamate/aspartate transport system permease protein gitJ Down Regulated ydcR 945049 Uncharacterized HTH-type transcriptional regulator ydcR ['Degulated June All All All All All All All All All Al					
ydiB 946200 Quinate/shikimate dehydrogenase Temporarly Down Re nrdA 946612 Ribonucleoside-diphosphate reductase; Ribonucleoside-diphosphate reductase; Ribonucleoside-diphosphate reductase; Ribonucleoside-diphosphate reductase 1 subunit alpha Down Regulated SpyrA 946414 DnA gyrase subunit A Down Regulated Uncharacterized HTH-type transcriptional regulator ydcQ Uncharacterized HTH-type transcriptional regulator ydcQ Up Regulated Up Regu					
nrdA 946612 Ribonucleoside-diphosphate reductase; Ribonucleoside-diphosphate reductase 1 subunit alpha ybiS 945441 Uncharacterized protein ybiS ['phosphoenolpyruvate dependent sugar phosphotransferase system'] Ambiguous/Unsorted gyrA 946614 DNA gyrase subunit A hicB 946010 Uncharacterized HTH-type transcriptional regulator ydcQ ydcQ 946001 Uncharacterized HTH-type transcriptional regulator ydcQ ['Coenzyme transport and metabolism'] Up Regulated gltJ 945433 Glutamate/aspartate transport system permease protein gltJ ydcR 946004 Uncharacterized HTH-type transcriptional regulator ydcR [] flgF 945639 Flagellar basal-body rod protein flgF yceF 945631 Maf-like protein yceF 1; Maf-like protein yceF yniC 945632 Phosphatase yniC tolB 945429 Protein tolB 945635 Phage shock protein D Down Regulated pspD 945635 Phage shock protein D Down Regulated pspD 945635 Flagellar hook protein flgE 945636 Flagellar hook protein flgE 945637 Page shock protein D Down Regulated pspD 945638 Flagellar hook protein flgE 945636 Flagellar hook protein flgE 945637 Phosphatase yniC Down Regulated pspD 945638 Phage shock protein D Down Regulated pspD 945635 Phage shock protein D Down Regulated pspD 945636 Flagellar hook protein flgE 945636 Flagellar hook protein flgE 945637 Pown Regulated					
ybiS 945441 Uncharacterized protein ybiS [phosphoenolpyruvate dependent sugar phosphotransferase system'] Ambiguous/Unsorted gyrA 946614 DNA gyrase subunit A Down Regulated bydGQ 946001 Uncharacterized HTH-type transcriptional regulator ydcQ ydGQ 946001 Uncharacterized HTH-type transcriptional regulator ydcQ [Coenzyme transport and metabolism'] Up Regulated ydcQ yd6001 Uncharacterized HTH-type transcriptional regulator ydcQ [Coenzyme transport and metabolism'] Up Regulated ydcQ yd6001 Up Regulated ydcQ yd6001 Uncharacterized HTH-type transcriptional regulator ydcQ ydcQ ydcQ ydcQ ydcQ ydcQ ydcQ ydcQ		_		posphata reductore 1 autumit alaba	
gyrA 946614 DNA gyrase subunit A hicB 946001 Uncharacterized HTH-type transcriptional regulator ydcQ ydcQ 946001 Uncharacterized HTH-type transcriptional regulator ydcQ ['Coenzyme transport and metabolism'] Up Regulated ydcQ 946001 Uncharacterized HTH-type transcriptional regulator ydcQ ['Coenzyme transport and metabolism'] Up Regulated gltJ 945403 Glutamate/aspartate transport system permease protein gltJ ydcR 946004 Uncharacterized HTH-type transcriptional regulator ydcR [] Up Regulated ydcR 946004 Uncharacterized HTH-type transcriptional regulator ydcR [] Up Regulated flgF 945639 Flagellar basal-body rod protein flgF yoeF 945631 Maf-like protein yceF 1; Maf-like protein yceF yniC 945632 Phosphatase yniC tolB 945429 Protein tolB ydiU 946219 UPF0061 protein ydiU pown Regulated pspD 945635 Phage shock protein D floIA 944790 Dihydrofolate reductase; Dihydrofolate reductase type 1 flgE 945636 Flagellar hook protein flgE Down Regulated flgE 945636 Flagellar hook protein flgE Down Regulated					
hicB 946001 Uncharacterized HTH-type transcriptional regulator ydcQ ydcQ 946001 Uncharacterized HTH-type transcriptional regulator ydcQ [Coenzyme transport and metabolism] Up Regulated 946003 3-oxoacyl-[acyl-carrier-protein] synthase 3 Up Regulated gltU 945443 Glutamate/aspartate transport system permease protein gltU ydcR 946004 Uncharacterized HTH-type transcriptional regulator ydcR [] Up Regulated ydcR 946004 Uncharacterized HTH-type transcriptional regulator ydcR [] Up Regulated ydcR 94639 Flagellar basal-body rod protein flgF Down Regulated yceF 945631 Maf-like protein yceF I; Maf-like protein yceF Protein tolB Protein tolB 945429 Protein tolB 945429 Protein tolB UpF0061 protein ydiU 946219 UPF0061 protein ydiU Down Regulated pspD 945635 Phage shock protein D Down Regulated pspD 945636 Phage shock protein D Up Regulated Ip Protein ydiV Down Regulated Protein ydiV Dihydrofolate reductase type 1 Down Regulated Ip Protein ydiV Dihydrofolate reductase type 1 Down Regulated Ip Down Regul				[priosprioenorpyruvate dependent sugar priospriotransierase system]	
ydcQ946001Uncharacterized HTH-type transcriptional regulator ydcQ['Coenzyme transport and metabolism']Up RegulatedfabH9460033-oxoacyl-[acyl-carrier-protein] synthase 3Down RegulatedgltJ945443Glutamate/aspartate transport system permease protein gltJDown RegulatedydcR946004Uncharacterized HTH-type transcriptional regulator ydcRUp RegulatedflgF945639Flagellar basal-body rod protein flgFDown RegulatedyceF945631Maf-like protein yceF 1; Maf-like protein yceFUp RegulatedyniC945632Phosphatase yniCTemporarly Down RegulatedtolB945429Protein tolBDown RegulatedydiU946219UPF0061 protein ydiUDown RegulatedpspD945635Phage shock protein DDown RegulatedfolA944790Dihydrofolate reductase; Dihydrofolate reductase type 1Temporarly UP RegulatedflgE945636Flagellar hook protein flgEDown Regulated					
fabH 946003 3-oxoacyl-[acyl-carrier-protein] synthase 3 gltJ 945443 Glutamate/aspartate transport system permease protein gltJ ydcR 946004 Uncharacterized HTH-type transcriptional regulator ydcR [] Up Regulated ydcR 945639 Flagellar basal-body rod protein flgF yceF 945631 Maf-like protein yceF 1; Maf-like protein yceF yniC 945632 Phosphatase yniC Up Regulated ydiU 945429 Protein tolB ydiU 946219 UPF0061 protein ydiU pspD 945635 Phage shock protein D folA 944790 Dihydrofolate reductase; Dihydrofolate reductase type 1 flgE 945636 Flagellar hook protein flgE Down Regulated				[Coenzyme transport and metabolism]	
gltJ 945443 Glutamate/aspartate transport system permease protein gltJ ydcR 946004 Uncharacterized HTH-type transcriptional regulator ydcR [] Up Regulated flgF 945639 Flagellar basal-body rod protein flgF Down Regulated ynic 945631 Maf-like protein yceF 1; Maf-like protein yceF Up Regulated ynic 945632 Phosphatase ynic Temporary Down Regulated ynic 945639 Protein tolB UpF0061 protein ydiu 946219 UPF0061 protein ydiu Down Regulated pspD 945635 Phage shock protein D Up Regulated pspD 945635 Phage shock protein D Up Regulated pspD 945636 Flagellar hook protein flgE Down Regulated Down Regulated PflogE P45636 Flagellar hook protein flgE Down Regulated Down Regul				Coonstyling transport and metabolism [
ydcR 94604 Uncharacterized HTH-type transcriptional regulator ydcR [] Up Regulated flgF 945639 Flagellar basal-body rod protein flgF Down Regulated yceF 945631 Maf-like protein yceF 1; Maf-like protein yceF Up Regulated yceF 945632 Phosphatase yniC Temporarly Down Retoll 945429 Protein tolB 945429 Protein tolB Down Regulated ydiU 946219 UPF0061 protein ydiU Down Regulated pspD 945635 Phage shock protein D Up Regulated Up Regulated pspD 945635 Phage shock protein D Up Regulated Poil Poil Poil Poil Poil Poil Poil Poil			, , , , , ,		
figF 945639 Flagellar basal-body rod protein flgF Down Regulated yceF 945631 Maf-like protein yceF 1; Maf-like protein yceF Up Regulated yniC 945632 Phosphatase yniC Temporarly Down Re tolB 945429 Protein tolB Down Regulated ydiU 946219 UPF0061 protein ydiU Down Regulated pspD 945635 Phage shock protein D Up Regulated folA 944700 Dihydrofolate reductase; Dihydrofolate reductase type 1 Temporarly UP Regulated flgE 945636 Flagellar hook protein flgE Down Regulated				п	
yceF945631Maf-like protein yceF 1; Maf-like protein yceFUp RegulatedyniC945632Phosphatase yniCTemporarly Down RetolB945429Protein tolBDown RegulatedydiU946219UPF0061 protein ydiUDown RegulatedpspD945635Phage shock protein DUp RegulatedfolA94470Dihydrofolate reductase; Dihydrofolate reductase type 1Temporarly Up RegulatedflgE945636Flagellar hook protein flgEDown Regulated				<u>u</u>	
yniC 945632 Phosphatase yniC Temporarly Down Re tolB 945429 Protein tolB Down Regulated ydiU 946219 UPF0061 protein ydiU Down Regulated pspD 945635 Phage shock protein D Up Regulated folA 944790 Dihydrofolate reductase; Dihydrofolate reductase type 1 Temporarly Up Regulated flgE 945636 Flagellar hook protein flgE Down Regulated					
toIB 945429 Protein toIB ydiU 946219 UPF0061 protein ydiU pspD 945635 Phage shock protein D foIA 944790 Dihydrofolate reductase; Dihydrofolate reductase type 1 flgE 945636 Flagellar hook protein flgE Down Regulated Down Regulated Temporarity D Down Regulated Down Regulated Down Regulated					
ydiU946219UPF0061 protein ydiUDown RegulatedpspD945635Phage shock protein DUp RegulatedfolA944790Dihydrofolate reductase; Dihydrofolate reductase type 1Temporarly Up RegulflgE945636Flagellar hook protein flgEDown Regulated					
pspD 945635 Phage shock protein D Up Regulated folA 944790 Dihydrofolate reductase; Dihydrofolate reductase type 1 Temporarly Up Regulated flgE 945636 Flagellar hook protein flgE Down Regulated					
folA944790Dihydrofolate reductase; Dihydrofolate reductase type 1Temporarly Up RegulflgE945636Flagellar hook protein flgEDown Regulated					
flgE 945636 Flagellar hook protein flgE Down Regulated	folA				
yaaH 944792 Inner membrane protein yaaH Ambiguous/Unsorted					
	yaaH	944792	Inner membrane protein yaaH		

flaN	945634	Flagella synthesis protein flgN		Down Regulated
flgN murE	944791	UDP-N-acetylmuramoyl-L-alanyl-D-glutamate2,6-diaminopimel	lata linasa	Down Regulated
aceF	944794	Dihydrolipoyllysine-residue acetyltransferase component of pyrus		Down Regulated
setA	944793	Sugar efflux transporter A	wate denydrogenase complex	Temporarly Down Re
leuB	944798	3-isopropylmalate dehydrogenase		Down Regulated
otsB	946406	Trehalose-phosphate phosphatase		Up Regulated
ftsI	944799	Peptidoglycan synthetase ftsl		Down Regulated
yncB	946012	Putative NADP-dependent oxidoreductase yncB		Ambiguous/Unsorted
ycjG	946013		UTI89_C1596; E2348C_1518; ECIAI1_1350; c1797; APECO1_478; ECS88_1467; ECIAI39_1677; EC55989_1489	Temporarly Down Re
	946016		01109_C1090, E2040C_1010, ECIAIT_1000, C1797, AFECOT_470, EC000_1407, ECIAID9_1077, EC00009_1400	Down Regulated
yccA ybiJ	945433	Inner membrane protein yccA Uncharacterized protein ybiJ		Up Regulated
araB	946017	Ribulokinase		
				Up Regulated
glnQ	945435 946808	Glutamine transport ATP-binding protein glnQ UPF0721 transmembrane protein vfcA		Down Regulated
yfcA				Down Regulated
csgC	945623	Putative curli production protein csgC	la combrana angulara his canacia! Isasina asid his conthatic areasas! Isasina asid family his conthatic areasas!	Down Regulated
yfcL	946804		ell wall membrane envelope biogenesis', 'amino acid biosynthetic process', 'aromatic amino acid family biosynthetic process']	Down Regulated
ylbG	946803	b0502; ECDH10B_0458		Down Regulated
ribD	945620	Riboflavin biosynthesis protein ribD		Down Regulated
glnP	945621	Glutamine transport system permease protein glnP		Temporarly Down Re
dapB	944762	Dihydrodipicolinate reductase		Down Regulated
ileS	944761	Isoleucyl-tRNA synthetase	VA. L.	Down Regulated
hcaD	945427	3-phenylpropionate/cinnamic acid dioxygenase ferredoxinNAD(n(+) reductase component	Temporarly Up Regul
mepA	946812	Penicillin-insensitive murein endopeptidase		Down Regulated
ybiH	945421		efense mechanisms', ' Primary Active Transporters']	Ambiguous/Unsorted
exbB	945420	Biopolymer transport protein exbB		Down Regulated
cyoB	945615	Ubiquinol oxidase subunit 1		Down Regulated
ndk	945611	Nucleoside diphosphate kinase		Temporarly Down Re
msyB	945612	Acidic protein msyB		Temporarly Up Regul
rbbA	947998	Uncharacterized ABC transporter ATP-binding protein yhiH		Temporarly Up Regul
carB	944775	Carbamoyl-phosphate synthase large chain		Down Regulated
ftsA	944778	Cell division protein ftsA		Down Regulated
nikR	947995	Nickel-responsive regulator		Temporarly Up Regul
sspB	944774	Stringent starvation protein B		Down Regulated
kefC	944773	Glutathione-regulated potassium-efflux system protein kefC		Ambiguous/Unsorted
rpmD	947797	50S ribosomal protein L30		Down Regulated
secY	947799	Preprotein translocase subunit secY; Preprotein translocase sec	cY subunit	Down Regulated
rplO	947798	50S ribosomal protein L15		Down Regulated
rpsD	947793	30S ribosomal protein S4		Down Regulated
rpsK	947792	30S ribosomal protein S11		Down Regulated
rpsE	947795	30S ribosomal protein S5		Down Regulated
rpoA	947794	DNA-directed RNA polymerase subunit alpha		Down Regulated
rpsM	947791	30S ribosomal protein S13; 30S ribosomal protein		Down Regulated
metC	946240	Cystathionine beta-lyase metC		Down Regulated
ybdG	946243	Uncharacterized protein ybdG		Down Regulated
pspF	945683	Psp operon transcriptional activator		Down Regulated
potD	945682	Spermidine/putrescine-binding periplasmic protein		Down Regulated
fiu	946246	Catecholate siderophore receptor fiu		Temporarly Up Regul
cobB	945687	NAD-dependent deacetylase		Up Regulated
panD	945686	Aspartate 1-decarboxylase		Down Regulated
nudJ	945689	Phosphatase nudJ		Down Regulated Down Regulated
ybhL	945401	Inner membrane protein ybhL		Temporarly Up Regul
rplQ	947784	50S ribosomal protein L17		
trkA	947788	Trk system potassium uptake protein trkA		Down Regulated
yhdN	947785	Uncharacterized protein vhdN		Up Regulated
	947785		2005- 62422- 72205- ECEDA 2479- ECHMN 2456- HTIOL C2204- ECNMD 2277- ECMM 2400	Ambiguous/Unsorted
yehQ			2265; b2122; Z3295; ECED1_2478; ECUMN_2456; UTI89_C2394; ECDH10B_2277; ECIAI1_2198	Up Regulated
spy	946253	Spheroplast protein Y		Up Regulated
ycfP	945676	UPF0227 protein ycfP		Temporarly Up Regul
astB	946259	N-succinylarginine dihydrolase		Ambiguous/Unsorted
lptE	946257	LPS-assembly lipoprotein lptE		Down Regulated
lolC	945673	Lipoprotein-releasing system transmembrane protein IoIC		Down Regulated
flgB	945678	Flagellar basal-body rod protein flgB		Down Regulated
parE	947501	DNA topoisomerase 4 subunit B		Down Regulated
ybiX	947502	PKHD-type hydroxylase ybiX		Ambiguous/Unsorted
yrdB	947774		uilding block biosynthesis', 'purine nucleotide biosynthetic process', 'aromatic amino acid family biosynthetic process']	Temporarly Down Re
yrdA	947775	Protein yrdA		Down Regulated

	0.47770	hu iii i i i i i i i i i i i i i i i i i		IA 1: #1 (1)
fmt	947779	Methionyl-tRNA formyltransferase		Ambiguous/Unsorted
nlpC	946220	Probable lipoprotein nlpC		Up Regulated
holB	945661	DNA polymerase III subunit delta'		Down Regulated
thrS	946222	Threonyl-tRNA synthetase		Down Regulated
mrp	946627	Protein mrp		Ambiguous/Unsorted
pntA	946628	NAD(P) transhydrogenase subunit alpha	n.	Down Regulated
ycfS	945666	Uncharacterized protein ycfS Phospho-2-dehydro-3-deoxyheptonate aldolase, Trp-sensitiv	IU	Down Regulated
aroH	946229 946225	Translation initiation factor IF-3	le	Down Regulated
infC				Down Regulated
accC aaeR	947761 947760	Biotin carboxylase HTH-type transcriptional activator aaeR		Down Regulated Up Regulated
yhdV	947767	Uncharacterized protein yhdV	[' Electrochemical potential driven transporters', ' Protection', 'cell cycle', 'cell division']	Up Regulated
acrF	947768	Acriflavine resistance protein F	Electrochemical potential universitalisporters , Protection , Celi cycle , Celi division j	Up Regulated
yagV	946631	Uncharacterized protein yagV	n	Up Regulated
yegU	946630	Uncharacterized protein yegU	<u>u</u> n	Down Regulated
yhdY	947764	Inner membrane amino-acid ABC transporter permease protein	IU ein vhdY	Down Regulated
yceD	945650	Uncharacterized protein yceD	en yna'i	Down Regulated
ydiZ	946232	Uncharacterized protein ydiZ	 Carbon compound utilization', ' Energy metabolism carbon ', 'translation'	Temporarly Up Regul
yfaA	946618	Uncharacterized protein yfaA	[Replication recombination and repair, 'RNA related', 'Type of regulation']	Ambiguous/Unsorted
rpmF	945657	50S ribosomal protein L32	(Down Regulated
ycfH	945656	Uncharacterized deoxyribonuclease ycfH	ln	Temporarly Up Regul
ydjN	946238	Uncharacterized symporter ydjN		Down Regulated
yhcH	947750	Uncharacterized protein yhcH	[Carbohydrate transport and metabolism', 'carbohydrate metabolic process']	Up Regulated
flgK	945648	Flagellar hook-associated protein 1	(Entering and Entering and Annual An	Down Regulated
fhuE	945649	FhuE receptor		Ambiguous/Unsorted
yegS	946626	Probable lipid kinase yegS; Lipid kinase yegS		Up Regulated
mreD	947756	Rod shape-determining protein mreD		Temporarly Down Re
nanK	947757	N-acetylmannosamine kinase		Temporarly Up Regul
tolA	946625	Protein tolA		Down Regulated
ассВ	947758	Biotin carboxyl carrier protein of acetyl-CoA carboxylase		Down Regulated
hpt	946624	Hypoxanthine phosphoribosyltransferase		Down Regulated
ccmH	946623	Cytochrome c-type biogenesis protein ccmH		Down Regulated
cyaA	947755	Adenylate cyclase		Temporarly Down Re
nanE	947745	Putative N-acetylmannosamine-6-phosphate 2-epimerase		Up Regulated
aaeA	947748	p-hydroxybenzoic acid efflux pump subunit aaeA		Up Regulated
aaeB	947747	p-hydroxybenzoic acid efflux pump subunit aaeB		Up Regulated
glgS	947533	Glycogen synthesis protein glgS		Up Regulated
hybC	945182	Hydrogenase-2 large chain		Up Regulated
flgl	947534	Flagellar P-ring protein		Down Regulated
yejL	946694	UPF0352 protein yejL		Ambiguous/Unsorted
yqiK	947537	Inner membrane protein yqiK		Up Regulated
rtn	946695	Protein rtn		Temporarly Down Re
nanT	947740	Putative sialic acid transporter; Putative sialic acid transporte	er 1	Up Regulated
lpxT	946693	Inner membrane protein yeiU		Down Regulated
fepB	947538	Ferrienterobactin-binding periplasmic protein		Ambiguous/Unsorted
yeeN	946493	UPF0082 protein yeeN		Temporarly Down Re
guaA	947334	GMP synthase [glutamine-hydrolyzing]		Down Regulated
rhtC	948317	Threonine efflux protein		Down Regulated
sulA	947335	Cell division inhibitor sulA		Up Regulated
tatB	948319	Sec-independent protein translocase protein tatB homolog; S	Sec-independent protein translocase protein tatB	Down Regulated
dapE	948313	Succinyl-diaminopimelate desuccinylase		Ambiguous/Unsorted
pldB	948314	Lysophospholipase L2		Temporarly Down Re
shiA	946495	Shikimate transporter		Down Regulated
RhtB	948316	Homoserine/homoserine lactone efflux protein		Down Regulated
yhcG	947739		[Macromolecule degradation']	Down Regulated
glgB	947940	1,4-alpha-glucan-branching enzyme		Down Regulated
metR	948310	HTH-type transcriptional regulator metR		Up Regulated
hemY	948311	Protein hemY		Down Regulated
glgX	947941	Glycogen debranching enzyme		Up Regulated
yeeJ	946498	Uncharacterized protein yeeJ		Up Regulated
ushA	947331	Protein ushA		Down Regulated
yejM	946685	Inner membrane protein yejM		Down Regulated
aroB	947927	3-dehydroquinate synthase		Down Regulated
		Classifier hand hady rad protein fino		
flgC nuoM	946687 947731	Flagellar basal-body rod protein flgC NADH-quinone oxidoreductase subunit M		Down Regulated Down Regulated

yrbG	947730	Inner membrane protein yrbG	Ambiguous/Unsorted
yejF	946689	Uncharacterized ABC transporter ATP-binding protein yejF	Down Regulated
pgpA	947542	Phosphatidylglycerophosphatase A	Up Regulated
ahpF	947540	Alkyl hydroperoxide reductase subunit F	Down Regulated
kdsD	947734	Arabinose 5-phosphate isomerase	Down Regulated
rtcB	947929	Protein rtcB	Temporarly Up Regul
yggG	945173	Uncharacterized metalloprotease yggG	Up Regulated
osmF	946681	Putative osmoprotectant uptake system substrate-binding protein osmF	Temporarly Up Regul
bglX	946682	Periplasmic beta-glucosidase	Temporarly Down Re
rfaE	947548	Bifunctional protein hIdE	Down Regulated
flgM	946684 946480	Negative regulator of flagellin synthesis	Down Regulated Up Regulated
yodA zinT	946480	Metal-binding protein yodA Metal-binding protein yodA	Up Regulated
ygeA	947348	Uncharacterized protein your	Up Regulated
hchA	946481	Chaperone protein hchA	Ambiguous/Unsorted
pldA	948307	Phospholipase A1	Down Regulated
yedW	946486	Probable transcriptional regulatory protein yedW	Temporarly Down Re
damX	947930	Protein damX	Down Regulated
rffG	948300	dTDP-glucose 4,6-dehydratase	Down Regulated
glgP	947931	Phosphorylase; Glycogen phosphorylase	Down Regulated
rffM	948301	Probable UDP-N-acetyl-D-mannosaminuronic acid transferase	Down Regulated
yhbJ	947727	UPF0042 protein yhbJ	Down Regulated
glpE	947935	Thiosulfate sulfurtransferase glpE	Temporarly Down Re
glgA	947932	Glycogen synthase	Temporarly Up Regul
gltB	947724	Glutamate synthase [NADPH] large chain	Down Regulated
gltD	947723 947726	Glutamate synthase [NADPH] small chain Spermidine synthase	Down Regulated Down Regulated
speE ampE	946678	Protein ampE	Ambiguous/Unsorted
lptB	947725	Lipopolysaccharide export system ATP-binding protein lptB	Down Regulated
fruK	946676	I-phosphofructokinase	Temporarly Up Regul
secG	947720	Protein-export membrane protein secG	Down Regulated
npr	947914	Phosphocarrier protein NPr	Ambiguous/Unsorted
yeiP	946674	Elongation factor P-like protein	Down Regulated
lptC	947722	Lipopolysaccharide export system protein lptC	Down Regulated
bioH	947916	Carboxylesterase bioH	Down Regulated
fruA	946672	PTS system fructose-specific EIIBC component	Temporarly Up Regul
aes	947514	Acetyl esterase	Up Regulated
araA	947511	L-arabinose isomerase	Up Regulated
lplT	947317	Lysophospholipid transporter lpIT	Temporarly Up Regul
kdul yafC	947319 947507	4-deoxy-L-threo-5-hexosulose-uronate ketol-isomerase Uncharacterized HTH-type transcriptional regulator yafC ['Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent']	Down Regulated Up Regulated
rsxE	947507	Uncharacterized HTH-type transcriptional regulator yafC ['Transcription', 'RNA related', 'Type of regulation', 'transcription', 'regulation of transcription DNA dependent'] Electron transport complex protein rnfE	Down Regulated
galR	947314	HTH-type transcriptional regulator galR	Temporarly Down Re
malP	947922	Phosphorylase; Maltodextrin phosphorylase	Down Regulated
malQ	947923	4-alpha-qlucanotransferase	Down Regulated
gntT	947924	High-affinity gluconate transporter	Down Regulated
kdsC	947717	3-deoxy-D-manno-octulosonate 8-phosphate phosphatase	Ambiguous/Unsorted
hpf	947718	Probable sigma(54) modulation protein	Down Regulated
pspB	945893	Phage shock protein B	Up Regulated
speD	947719	S-adenosylmethionine decarboxylase proenzyme	Down Regulated
IptA	947920	Lipopolysaccharide export system protein lptA	Down Regulated
speB	947715	Agmatinase	Down Regulated
yeiH	946668 945493	UPF0324 inner membrane protein yeiH; Probable endonuclease 4 UPF0145 protein ybjQ	Up Regulated
ybjQ rpoN	945493	RNA polymerase sigma-54 factor	Up Regulated Down Regulated
nfo	947714	Endonuclease 4; Probable endonuclease 4	Down Regulated Down Regulated
yrbD	947712	Uncharacterized protein yrbD ['Amino acid transport and metabolism']	Down Regulated
mlaD	947712	Uncharacterized protein yrbD	Down Regulated
ybjS	945495	Uncharacterized protein ybjS []	Ambiguous/Unsorted
aat	945490	Leucyl/phenylalanyl-tRNAprotein transferase	Down Regulated
nudE	947906	ADP compounds hydrolase nudE	Down Regulated
mlaC	947710	Protein yrbC	Down Regulated
yrbC	947710	Protein yrbC	Down Regulated
yohD	946661	Inner membrane protein yohD	Ambiguous/Unsorted
thrC	945198	Threonine synthase	Down Regulated

pbpG	946662	D-alanyl-D-alanine endopeptidase		Ambiguous/Unsorted
ybjD	945498		"SOS response"	Temporarly Up Regul
entD		4'-phosphopantetheinyl transferase entD		Temporarly Up Regul
rpmA		50S ribosomal protein L27		Down Regulated
cusA	945191	Cation efflux system protein cusA		Down Regulated
phoP	945697	Transcriptional regulatory protein phoP		Ambiguous/Unsorted
yiaC	946460	Uncharacterized N-acetyltransferase yiaC	"amino acid biosynthetic process"	Temporarly Down Re
nudF	947519	ADP-ribose pyrophosphatase		Down Regulated
hsIR	947912	Heat shock protein 15		Ambiguous/Unsorted
potC	945691	Spermidine/putrescine transport system permease protein po	tC	Down Regulated
ymdC	947321	Uncharacterized protein ymdC		Down Regulated
ompR	947913	Transcriptional regulatory protein ompR	u	Temporarly Up Regul
hflD	945693		39 2005; ECED1 1276; E2348C 1273; ECUMN 1376; EC55989 1245	Temporarly Down Re
dusB	947707	tRNA-dihydrouridine synthase B		Down Regulated
prmA	947708	Ribosomal protein L11 methyltransferase		Down Regulated
artP	945489	Arginine transport ATP-binding protein artP		Ambiguous/Unsorted
ygjD	947578	Probable O-sialoglycoprotein endopeptidase		Ambiguous/Unsorted
murJ	945487	Virulence factor mviN homolog		Down Regulated
potF	945480	Putrescine-binding periplasmic protein		Temporarly Up Regul
ybjC	945481	Uncharacterized protein ybjC	['Energy production and conversion', 'Translation ribosomal structure and biogenesis', 'Energy metabolism carbon', 'Building blo	Ambiguous/Unsorted
dnaG	947570	DNA primase		Up Regulated
gmr	945868	Protein gmr		Down Regulated
rimK	945484	Ribosomal protein S6 modification protein		Down Regulated
holA	947573	DNA polymerase III subunit delta		Down Regulated
potl	945485	Putrescine transport system permease protein potl		Temporarly Up Regul
csrD	947702	Uncharacterized protein yhdA		Temporarly Down Re
murA	947703	UDP-N-acetylglucosamine 1-carboxyvinyltransferase		Down Regulated
btuF	947574	Vitamin B12-binding protein		Down Regulated
srlA	947575	Glucitol/sorbitol permease IIC component		Temporarly Up Regul
yhhQ	947984	Inner membrane protein yhhQ		Ambiguous/Unsorted
ymjA	945874	Uncharacterized protein ymjA	['Amino acid transport and metabolism', ' Electrochemical potential driven transporters']	Ambiguous/Unsorted
ribE	946453	6,7-dimethyl-8-ribityllumazine synthase		Down Regulated
yhhS	947982	UPF0226 protein yhhS		Temporarly Down Re
potG	945476	Putrescine transport ATP-binding protein potG		Temporarly Up Regul
tyrR	945879	Transcriptional regulatory protein tyrR		Up Regulated
ftsX	947969	Cell division protein ftsX		Down Regulated
ybjl	945470	Phosphatase ybjl		Up Regulated
ybjJ	945471	Inner membrane protein ybjJ		Temporarly Down Re
ihfA	945472	Integration host factor subunit alpha		Down Regulated
ybjK	945473		['translation', 'ion transport']	Up Regulated
puuD	945882	Gamma-glutamyl-gamma-aminobutyrate hydrolase		Down Regulated
yhhN	947973	Uncharacterized membrane protein yhhN	[Energy metabolism carbon]	Ambiguous/Unsorted
ybaV	945884	Uncharacterized protein ybaV		Up Regulated
pspA	945887	Phage shock protein A		Up Regulated
puuR	945886	HTH-type transcriptional regulator puuR		Ambiguous/Unsorted
ygdR	947304	Uncharacterized lipoprotein ygdR	['Amino acid transport and metabolism', ' Building block biosynthesis', 'amino acid metabolic process']	Temporarly Up Regul
hemD	948587	Uroporphyrinogen-III synthase		Down Regulated
livJ	947971	Leu/Ile/Val-binding protein		Down Regulated
rpoH	947970	RNA polymerase sigma-32 factor; RNA polymerase sigma fa	ctor	Up Regulated
ascG	947305	HTH-type transcriptional regulator ascG		Temporarly Down Re
ygiF	947554	Uncharacterized protein ygiF	['Amino acid transport and metabolism']	Temporarly Down Re
ylil		Soluble aldose sugar dehydrogenase ylil		Up Regulated
ygiM	947555	Uncharacterized protein ygiM		Down Regulated
rimO	945465	Ribosomal protein S12 methylthiotransferase rimO		Down Regulated
yliF	945463		"Signal transduction mechanisms', 'Type of regulation', 'two component signal transduction system phosphorelay', 'transcription',	
bacA	947551	Undecaprenyl-diphosphatase		Down Regulated
gsiC	945460	Glutathione transport system permease protein gsiC		Ambiguous/Unsorted
gsiD	945461	Glutathione transport system permease protein gsiD		Down Regulated
ribC	945848	Riboflavin synthase alpha chain	Wasing add transport and analohaliant Uniddies blads bigger to the country of the	Ambiguous/Unsorted
yrbA troE	947958		['Amino acid transport and metabolism', ' Building block biosynthesis', 'amino acid biosynthetic process']	Down Regulated
trpE	945846	Anthranilate synthase component 1	magaa protein lii.M.	Down Regulated
livM	947966	High-affinity branched-chain amino acid transport system per Protein ampG	півазе ріоцені нумі	Temporarly Up Regul
ampG	946438		[PDNA processing]	Down Regulated
yjcE	948577		[rRNA processing]	Down Regulated
livH	947965	High-affinity branched-chain amino acid transport system per	mease protein iivi i	Temporarly Up Regul

ydgD	946436	Uncharacterized serine protease ydgD	[response to stress']	Up Regulated
livG	947967	High-affinity branched-chain amino acid transport ATP-bindir		Temporarly Up Regul
livF	947961	High-affinity branched-chain amino acid transport ATP-bindir		Down Regulated
yhhK	947963	Uncharacterized protein yhhK	[' Carbon compound utilization', ' Metabolism of other compounds']	Down Regulated
ybeY	946430	Putative metalloprotease ybeY		Temporarly Up Regul
ghra	946431	Glyoxylate/hydroxypyruvate reductase A		Down Regulated
ујсН	948574	Inner membrane protein yjcH		Ambiguous/Unsorted
gsiA	945457	Glutathione import ATP-binding protein gsiA		Down Regulated
gsiB	945459	Glutathione-binding protein gsiB		Down Regulated
rpoD	947567	RNA polymerase sigma factor rpoD; RNA polymerase sigma	factor	Temporarly Down Re
moeA	945454	Molybdopterin biosynthesis protein moeA		Down Regulated
dacC	945455	D-alanyl-D-alanine carboxypeptidase dacC		Temporarly Up Regul
iaaA	945456	Isoaspartyl peptidase		Down Regulated
folC	945451	Bifunctional protein folC		Down Regulated
moeB	945452	Molybdopterin biosynthesis protein moeB		Ambiguous/Unsorted
mug	947560	G/U mismatch-specific DNA glycosylase		Ambiguous/Unsorted
	947947			
ggt		Gamma-glutamyltranspeptidase	Ill pid transport and matchelians!	Up Regulated
yciW	945859	Uncharacterized protein yciW	[Lipid transport and metabolism']	Down Regulated
proB	946425	Glutamate 5-kinase	n	Ambiguous/Unsorted
yjcB	948569	Uncharacterized protein yjcB	Ш	Down Regulated
rnb	945864	Exoribonuclease 2		Down Regulated
soxR	948566	Redox-sensitive transcriptional activator soxR		Up Regulated
fecA	946427	Fe(3+) dicitrate transport protein fecA		Down Regulated
soxS	948567	Regulatory protein soxS		Temporarly Up Regul
yrbB	947954	Uncharacterized protein yrbB	[Cell wall membrane envelope biogenesis', ' Macromolecules cellular constituent biosynthesis', 'cell cycle', 'cell division']	Down Regulated
mlaB	947954	Uncharacterized protein yrbB		Down Regulated
ugpC	947953	sn-glycerol-3-phosphate import ATP-binding protein ugpC		Temporarly Up Regul
ydfl	945861	Uncharacterized oxidoreductase ydfl	[' Carbon compound utilization']	Temporarly Down Re
yciN	945860	Protein yciN		Temporarly Up Regul
рдрВ	945863	Phosphatidylglycerophosphatase B		Down Regulated
topA	945862	DNA topoisomerase 1		Down Regulated
uvrY	946424	Response regulator uvrY		Ambiguous/Unsorted
yjbR	948560	Uncharacterized protein yibR	[Amino acid transport and metabolism', ' Central intermediary metabolism', 'amino acid biosynthetic process', 'biosynthetic process	
alr	948564	Alanine racemase, biosynthetic; Alanine racemase	,	Down Regulated
yjcD	948565	Putative permease yjcD		Temporarly Up Regul
dapF	948364	Diaminopimelate epimerase		Down Regulated
ydgC	945156	Inner membrane protein ydgC		Temporarly Down Re
ydgJ	947164	Uncharacterized oxidoreductase ydgJ	П	Temporarly Down Re
glnL	948360	Nitrogen regulation protein NR(II); Sensor protein	<u>u</u>	Down Regulated
glnG	948361	Nitrogen regulation protein NR(I)		Down Regulated
hemN	948362	Oxygen-independent coproporphyrinogen-III oxidase		Down Regulated
feaR	945941	Transcriptional activator feaR		Ambiguous/Unsorted
emrB	947167	Multidrug resistance protein B		Ambiguous/Unsorted
	948363			
yihl	947168	UPF0241 protein yihl		Up Regulated
luxS iclR		S-ribosylhomocysteine lyase		Down Regulated
	948524	Acetate operon repressor	PAmino gold transport and metabolism! ! Corbon compound utilization! ! Corbon distributions of the corporation and transport and metabolism! ! Corbon compound utilization! ! Corbon compound utilization!	Down Regulated
yihL	948368		['Amino acid transport and metabolism', ' Carbon compound utilization', ' Central intermediary metabolism', ' Primary Active Transport	
rfbA	945154	Glucose-1-phosphate thymidylyltransferase; Glucose-1-phos		Down Regulated
ubiE	948926	Ubiquinone/menaquinone biosynthesis methyltransferase ub		Down Regulated
typA	948369	GTP-binding protein typA/BipA; GTP-binding protein typA/bip)A	Down Regulated
prfB	947369	Peptide chain release factor 2		Down Regulated
yccJ	948930	Uncharacterized protein yccJ	[Macromolecules cellular constituent biosynthesis']	Temporarly Up Regul
malY	945937	Protein malY		Temporarly Up Regul
ybgS	945356	Uncharacterized protein ybgS	['Inorganic ion transport and metabolism', ' Electrochemical potential driven transporters', 'ion transport']	Temporarly Up Regul
Суса	948725	D-serine/D-alanine/glycine transporter		Down Regulated
fkIB	948726	FKBP-type 22 kDa peptidyl-prolyl cis-trans isomerase		Down Regulated
ybiO	945935	Uncharacterized mscS family protein ybiO		Temporarly Up Regul
ytfB	948727	Uncharacterized protein ytfB	['Replication recombination and repair', 'DNA recombination', 'DNA integration']	Ambiguous/Unsorted
galE	945354	UDP-glucose 4-epimerase		Ambiguous/Unsorted
rpsR		30S ribosomal protein S18		Down Regulated
priB	948722	Primosomal replication protein n		Down Regulated
fdx	947160	2Fe-2S ferredoxin		Temporarly Down Re
mngB	945359	Alpha-mannosidase mngB		Up Regulated
rpsF		30S ribosomal protein S6		Down Regulated
ygfA	945167	Uncharacterized protein ygfA	['cell cycle', 'cell division']	Up Regulated
J J., .	10 .0.		K aan ayara, aan annarang	1-F

dtd	948378	D-tyrosyl-tRNA(Tyr) deacylase		Down Regulated
dbpA	947153	ATP-independent RNA helicase dbpA		Down Regulated
ycfL	948539	Uncharacterized protein ycfL	['Nucleotide transport and metabolism', 'Cell wall membrane envelope biogenesis', ' Building block biosynthesis']	Down Regulated
yfgl	945168	Uncharacterized protein yfgl	[Lipid transport and metabolism', 'Building block biosynthesis']	Temporarly Up Regul
malE	948538	Maltose-binding periplasmic protein	Elpid transport and metabolism, Dunding block blosynthesis j	Temporarly Up Regul
malK	948537	Maltose/maltodextrin import ATP-binding protein malK		Down Regulated
ygaC	947156	Uncharacterized protein ygaC	п	Up Regulated
cpdB		EC-5101: ECD 4467: ECSE 4510: E-SMS35 4603: ECLIM	III N 4750; Z5824; EcolC 3793; ECS88 4807; E2348C 4544; EC55989 4773; ECDH10B 4408; ECED1 5072; ECH74115 5730; I	Tomporarly Down Box
trkG	945932	Trk system potassium uptake protein trkG	N_4130, 23024, E0010_3193, E0300_4001, E23400_4344, E033908_4113, E0DITIOE_4400, E0EDI_3012, E0TIT4113_3130, E	Up Regulated
ascF	945952	PTS system arbutin-, cellobiose-, and salicin-specific EIIBC of	nome and the second sec	
feaB	947134	Phenylacetaldehyde dehydrogenase	amponent	Up Regulated Up Regulated
	948532			
malF		Maltose transport system permease protein malF		Down Regulated
lysC	948531 944999	Aspartokinase; Lysine-sensitive aspartokinase 3 Paraquat-inducible protein A		Down Regulated
pqiA				Ambiguous/Unsorted
malG		Maltose transport system permease protein malG		Temporarly Up Regul
ytjC	948918 945164	Probable phosphoglycerate mutase gpmB		Temporarly Down Re
macB		Macrolide export ATP-binding/permease protein macB		Temporarly Down Re
ybcL	945165	UPF0098 protein ybcL	Providing a granting II	Up Regulated
yigP	948915	Uncharacterized protein yigP	['protein secretion']	Down Regulated
yjtD	948924	Uncharacterized tRNA/rRNA methyltransferase lasT		Ambiguous/Unsorted
ytfH	948730	Uncharacterized HTH-type transcriptional regulator ytfH	Ш	Up Regulated
yqeA	947358	Carbamate kinase-like protein yqeA		Temporarly Up Regul
mocA	947356	Uncharacterized protein ygfJ		Temporarly Up Regul
ygfJ	947356	Uncharacterized protein ygfJ	['electron transport']	Temporarly Up Regul
glnA	948370	Glutamine synthetase		Down Regulated
ytfJ	948737	Uncharacterized protein ytfJ	['Energy production and conversion', 'Secondary metabolites biosynthesis transport and catabolism']	Up Regulated
ytfL	948735	UPF0053 inner membrane protein ytfL		Temporarly Down Re
modF	945368	Putative molybdenum transport ATP-binding protein modF		Down Regulated
yaiZ	947355	Uncharacterized protein yaiZ		Ambiguous/Unsorted
gltK	947354	Glutamate/aspartate transport system permease protein gltK		Down Regulated
qorB	948731	Uncharacterized oxidoreductase ytfG		Up Regulated
ygeR	947352	Uncharacterized lipoprotein ygeR		Ambiguous/Unsorted
ygbA	947187	Uncharacterized protein ygbA	['Energy production and conversion', ' Primary Active Transporters']	Up Regulated
msrB	947188	Peptide methionine sulfoxide reductase msrB		Ambiguous/Unsorted
fdol	948383	Formate dehydrogenase, cytochrome b556(fdo) subunit		Down Regulated
malM	948547	Maltose operon periplasmic protein		Temporarly Up Regul
allS	945139	HTH-type transcriptional activator allS		Up Regulated
ydjA	945964	Putative NAD(P)H nitroreductase ydjA		Down Regulated
yddB	945961	Uncharacterized protein yddB	['Cell wall membrane envelope biogenesis', 'translation', 'ion transport']	Temporarly Up Regul
hinT	948549	HIT-like protein hinT		Down Regulated
ybdJ	947183	Uncharacterized protein ybdJ	[' Protection', 'ion transport', 'iron ion transport']	Up Regulated
nuoN	945136	NADH-quinone oxidoreductase subunit N		Down Regulated
MItB	947184	Membrane-bound lytic murein transglycosylase B		Temporarly Down Re
lamB	948548	Maltoporin		Down Regulated
frvX	948388	Putative frv operon protein frvX		Ambiguous/Unsorted
yfiL	947106	Uncharacterized protein yfiL	['Replication recombination and repair', 'DNA recombination']	Temporarly Up Regul
norR	947186	Anaerobic nitric oxide reductase transcription regulator norR	•	Up Regulated
tus	945135	DNA replication terminus site-binding protein		Up Regulated
recN	947105	DNA repair protein recN		Ambiguous/Unsorted
pmrD	945130	Signal transduction protein pmrD		Ambiguous/Unsorted
ffh	947102	Signal recognition particle protein		Temporarly Down Re
rimM	947101	Ribosome maturation factor rimM		Down Regulated
dgkA	948543	Diacylglycerol kinase		Ambiguous/Unsorted
mtn		5'-methylthioadenosine/S-adenosylhomocysteine nucleosida	se	Down Regulated
ylaB			['Signal transduction mechanisms', 'Type of regulation', 'two component signal transduction system phosphorelay ', 'transcription',	
yibA		Protein yibA	(-1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3 1.3.	Up Regulated
visC	947389	Protein visC		Down Regulated
efeU	948956		89; EC55989 1127; b4490; ECED1 1172; ECIAI1 1062; E2348C 1067	Up Regulated
hypE	947182	Hydrogenase isoenzymes formation protein hypE		Temporarly Up Regul
ubiH	947388	2-octaprenyl-6-methoxyphenol hydroxylase		Down Regulated
ybbO	945337	Uncharacterized oxidoreductase vbbO	п	Down Regulated
fhIA	947181	Formate hydrogenlyase transcriptional activator	и	Up Regulated
thiL	947387	Thiamine-monophosphate kinase		Down Regulated
lyxK	948101	L-xylulose/3-keto-L-gulonate kinase		Ambiguous/Unsorted
sapF	945335	Peptide transport system ATP-binding protein sapF		Down Regulated
3api	37 0000	i chara a anahori ayatem A ri -binung biotem aabi-		Powii i regulated

рра	948748	Inorganic pyrophosphatase		Down Regulated
allR		HTH-type transcriptional repressor allR		Up Regulated
ygfZ		tRNA-modifying protein ygfZ		Temporarly Down Re
selB		Selenocysteine-specific elongation factor		Down Regulated
yihW	948381	Uncharacterized HTH-type transcriptional regulator yihW	['Transcription', 'RNA related', 'Type of regulation', 'Genetic unit regulated', 'transcription', 'regulation of transcription DNA depen	
ydeN	945957	Uncharacterized sulfatase ydeN	['Signal transduction mechanisms', ' Metabolism of other compounds']	Temporarly Up Regul
cueR		HTH-type transcriptional regulator cueR		Up Regulated
aegA	947383	Protein aegA		Temporarly Up Regul
fdoH	948395	Formate dehydrogenase-O iron-sulfur subunit		Down Regulated
dusA	948558	tRNA-dihydrouridine synthase A		Down Regulated
csrA	947176	Carbon storage regulator		Up Regulated
qorA	948556	Quinone oxidoreductase		Temporarly Up Regul
dnaB	948555	Replicative DNA helicase		Down Regulated
rhaT	948398	L-rhamnose-proton symporter		Up Regulated
mppA	945951	Periplasmic murein peptide-binding protein		Up Regulated
allA	945141	Ureidoglycolate hydrolase		Up Regulated
ThiM	945142	Hydroxyethylthiazole kinase		Temporarly Down Re
dinF	948554	DNA-damage-inducible protein F		Ambiguous/Unsorted
yjbJ	948553	UPF0337 protein yjbJ		Temporarly Up Regul
mtlA	948118	PTS system mannitol-specific EIICBA component		Temporarly Up Regul
mtIR	948116	Mannitol operon repressor		Ambiguous/Unsorted
cydA	945341	Cytochrome d ubiquinol oxidase subunit 1		Up Regulated
mpl	948752	UDP-N-acetylmuramate:L-alanyl-gamma-D-glutamyl-meso-d	iaminopimelate ligase	Down Regulated
yibK	948119	Uncharacterized tRNA/rRNA methyltransferase yibK	· · · · ·	Temporarly Down Re
yibH	948110	Inner membrane protein yibH		Up Regulated
zitB	945348	Zinc transporter zitB		Ambiguous/Unsorted
yceB	947377	Uncharacterized lipoprotein yceB	['electron transport']	Down Regulated
betB		Betaine aldehyde dehydrogenase		Down Regulated
ydhV	948756	Uncharacterized protein ydhV	[' Central intermediary metabolism']	Up Regulated
cydB		Cytochrome d ubiquinol oxidase subunit 2		Ambiguous/Unsorted
fbp		Fructose-1.6-bisphosphatase class 1		Ambiguous/Unsorted
ygfF	947373	Uncharacterized oxidoreductase ygfF	['tricarboxylic acid cycle']	Ambiguous/Unsorted
lysS	947372	Lysyl-tRNA synthetase; Lysyl-tRNA synthetase, heat inducible	e	Down Regulated
yibF	948113	Uncharacterized GST-like protein yibF	П	Temporarly Down Re
yddA	945945	Inner membrane ABC transporter ATP-binding protein yddA		Up Regulated
yibJ	948112	b3595; ECDH10B 3776; Z5017		Up Regulated
alkA		DNA-3-methyladenine glycosylase 2		Temporarly Down Re
yobB	948329	Uncharacterized protein yobB	['Replication recombination and repair']	Up Regulated
rluA		Ribosomal large subunit pseudouridine synthase A; Pseudou		Ambiguous/Unsorted
ygaH	945111	Uncharacterized protein ygaH	[Transcription', 'RNA related', 'Type of regulation', 'Protection', 'transcription', 'regulation of transcription DNA dependent']	Down Regulated
fre		NAD(P)H-flavin reductase		Down Regulated
fadA		3-ketoacyl-CoA thiolase		Up Regulated
ubiD		3-octaprenyl-4-hydroxybenzoate carboxy-lyase		Ambiguous/Unsorted
tatA		Sec-independent protein translocase protein tatA		Down Regulated
ysgA		Putative carboxymethylenebutenolidase		Down Regulated
patA		Putrescine aminotransferase		Up Regulated
torA	946267	Trimethylamine-N-oxide reductase 1		Up Regulated
apt		Adenine phosphoribosyltransferase		Down Regulated
metE		5-methyltetrahydropteroyltriglutamatehomocysteine methylt	ransferase	Ambiguous/Unsorted
acrA		Acriflavine resistance protein A		Down Regulated
ybbP	945118	Uncharacterized ABC transporter permease ybbP		Temporarly Down Re
pyrF	947121	Orotidine 5'-phosphate decarboxylase		Down Regulated
rhsD		Protein rhsD		Up Regulated
yjgH	948769	UPF0076 protein yjgH		Up Regulated
secB	948123	Protein-export protein secB		Down Regulated
trpD		Anthranilate synthase component II		Down Regulated
gpsA		Glycerol-3-phosphate dehydrogenase [NAD(P)+]		Down Regulated
moaE	945399	Molybdopterin-converting factor subunit 2		Temporarly Up Regul
pyrB	945399	Aspartate carbamoyltransferase catalytic chain; Aspartate ca	rhamoultransferase	Down Regulated
rhsA		Protein rhsA	nouncy) autoloruse	Up Regulated
pepB		Peptidase B		Down Regulated
treR		HTH-type transcriptional regulator treR		Ambiguous/Unsorted
ybhK	948760	UPF0052 protein vbhK		Temporarly Down Re
IVDIII	1940390		_	
		Uncharacterized protein vcfD		
ycfD cysE	945391	Uncharacterized protein ycfD Serine acetyltransferase		Up Regulated Down Regulated

yibQ	948128	Uncharacterized protein yibQ	['proteolysis']	Temporarly Down Re
ybbN	947119	Uncharacterized protein ybbN	[Nucleotide transport and metabolism']	Down Regulated
pepQ	948335	Xaa-Pro dipeptidase		Down Regulated
ynjH	946279	Uncharacterized protein ynjH	['Amino acid transport and metabolism']	Up Regulated
ybbJ	945126	Inner membrane protein ybbJ		Temporarly Up Regul
ompW	945128	Outer membrane protein W		Temporarly Up Regul
gcvT	947390	Aminomethyltransferase		Down Regulated
rfaF	948135	ADP-heptoseLPS heptosyltransferase 2		Down Regulated
gcvP	947394 948136	Glycine dehydrogenase [decarboxylating]		Down Regulated
rfaC ypdB	947395	Lipopolysaccharide heptosyltransferase 1 Uncharacterized response regulatory protein ypdB	n	Down Regulated Temporarly Up Regul
rfaD	948134	ADP-L-glycero-D-manno-heptose-6-epimerase	<u>U</u>	Down Regulated
gcvH	947393	Glycine cleavage system H protein		Down Regulated
yibN	948131	Uncharacterized protein yibN	n	Down Regulated
grxC	948132	Glutaredoxin-3	<u> </u>	Down Regulated
argl	948774	Ornithine carbamoyltransferase chain I; Ornithine carbamoyl	I Itransferase	Temporarly Down Re
rraB	948773	Uncharacterized protein yjgD		Up Regulated
yjgM	948772	Uncharacterized N-acetyltransferase yigM	П	Temporarly Up Regul
tdh	948139	L-threonine 3-dehydrogenase		Ambiguous/Unsorted
kbl	948138	2-amino-3-ketobutyrate coenzyme A ligase		Temporarly Up Regul
uvrD	948347	DNA helicase II		Down Regulated
rdoA	948346	Protein rdoA		Down Regulated
mobA	948349	Molybdopterin-guanine dinucleotide biosynthesis protein A		Temporarly Up Regul
ybgC	948907	Acyl-CoA thioester hydrolase ybgC		Temporarly Down Re
yihD	948348	Protein yihD		Down Regulated
yafD	946286	UPF0294 protein yafD		Up Regulated
ygaW	947147	Uncharacterized protein ygaW	<u>D</u>	Down Regulated
sppA	946281	Protease 4		Down Regulated
nupG	946282	Nucleoside permease nupG		Down Regulated
ygaU	947144	Uncharacterized protein ygaU	μ	Ambiguous/Unsorted
trmA	947143	tRNA (uracil-5-)-methyltransferase Uncharacterized protein yjaG		Temporarly Down Re
yjaG nfi	948501 948502	Endonuclease V	N .	Down Regulated Down Regulated
proW	947145	Glycine betaine/L-proline transport system permease proteir	a proW	Ambiguous/Unsorted
purH	948503	Bifunctional purine biosynthesis protein purH	TPIOV	Down Regulated
ygiS	947140	Putative binding protein ygiS		Temporarly Down Re
purD	948504	Phosphoribosylamineglycine ligase		Down Regulated
zraS	948506	Sensor protein; Sensor protein zraS		Ambiguous/Unsorted
ybjE	947142	Uncharacterized protein ybjE	[' Building block biosynthesis', 'metabolic process']	Temporarly Down Re
zraP	948507	Zinc resistance-associated protein		Temporarly Up Regul
holC	948787	DNA polymerase III subunit chi		Down Regulated
yfcD	946783	Uncharacterized Nudix hydrolase yfcD	['tricarboxylic acid cycle', 'response to antibiotic']	Down Regulated
yqhG	945919	Uncharacterized protein yqhG	['Replication recombination and repair', ' Carbon compound utilization', 'DNA recombination']	Up Regulated
yccU	949008	Uncharacterized protein yccU		Temporarly Up Regul
folX	946781	D-erythro-7,8-dihydroneopterin triphosphate epimerase		Ambiguous/Unsorted
rfe	948789	Undecaprenyl-phosphate alpha-N-acetylglucosaminyl 1-pho	sphate transferase	Temporarly Down Re
rfaY	948145	Lipopolysaccharide core biosynthesis protein rfaY		Down Regulated
yajC	945374	UPF0092 membrane protein yajC		Down Regulated
rfaL	948148	O-antigen ligase		Down Regulated
rpsl	949000	30S ribosomal protein S9		Down Regulated
DeoB ybhA	948910 945372	Phosphopentomutase Phosphatase ybhA		Temporarly Up Regul Down Regulated
yfcl	945372	Uncharacterized protein yfcl	['Amino acid transport and metabolism', ' Building block biosynthesis', 'amino acid metabolic process', 'aromatic amino acid family to	b I b Dogulated
radA	946787	DNA repair protein radA	Paristic and transport and metabolism, building block biosynthesis, affilio acid metabolic process, aformatic affilio acid family to	Temporarly Up Regul
mngR	945371	Mannosyl-D-glycerate transport/metabolism system repressor	L OF MACK	Ambiguous/Unsorted
intB	948782	ECDH10B_4464; ECIAI39_4743; UTI89_C4878; b4271; ECI	· U	Temporarly Up Regul
nadR	948911	Transcriptional regulator nadR	==	Down Regulated
valS	948785	Valyl-tRNA synthetase		Down Regulated
serB	948913	Phosphoserine phosphatase		Temporarly Down Re
yfcH	946784	Epimerase family protein yfcH		Up Regulated
yeaG	946297	Uncharacterized protein yeaG		Up Regulated
YigA	948359	Uncharacterized protein yigA		Down Regulated
recR	945100	Recombination protein recR		Down Regulated
yihA	948358	Probable GTP-binding protein engB		Down Regulated
yigB	948357	Uncharacterized protein yigB		Down Regulated

vas\/	047136	Probable HTH-type transcriptional regulator ygaV		Lin Populated
ygaV tomB	947136 945106	Uncharacterized protein ybaJ		Up Regulated Up Regulated
	948513			
metA		Homoserine O-succinyltransferase	n	Temporarly Up Regul
yjaB	948514	Uncharacterized N-acetyltransferase yjaB	U .	Up Regulated
dnaX	945105	DNA polymerase III subunit tau		Down Regulated
polA	948356	DNA polymerase I		Down Regulated
ftsK	945102	DNA translocase ftsK		Temporarly Down Re
aceA	948517	Isocitrate lyase		Temporarly Up Regul
pagB	948518	Uncharacterized protein yjbD		Ambiguous/Unsorted
yjbD	948518	Uncharacterized protein yjbD		Ambiguous/Unsorted
xerC	948355	Tyrosine recombinase xerC		Down Regulated
ybaB	945104	UPF0133 protein ybaB		Down Regulated
asr	945103	Acid shock protein		Temporarly Up Regul
dsbA	948353	Thiol:disulfide interchange protein dsbA		Down Regulated
yicC	948154	UPF0701 protein yicC		Down Regulated
idnT	948798	Gnt-II system L-idonate transporter		Ambiguous/Unsorted
purF	946794	Amidophosphoribosyltransferase		Down Regulated
truA	946793	tRNA pseudouridine synthase A		Down Regulated
ујаН	948508	Uncharacterized protein yjaH	[Cell wall membrane envelope biogenesis', 'lipopolysaccharide biosynthetic process']	Down Regulated
pyrE	948157	Orotate phosphoribosyltransferase		Down Regulated
ynaJ	945901	Uncharacterized protein ynaJ	['Replication recombination and repair', 'DNA recombination']	Temporarly Down Re
bioC	945388	Biotin synthesis protein bioC		Down Regulated
slmA	948158	HTH-type protein slmA		Temporarly Down Re
uspE	945904	Universal stress protein E		Temporarly Up Regul
rph	948156	Ribonuclease PH		Temporarly Down Re
folK	948792	2-amino-4-hydroxy-6-hydroxymethyldihydropteridine pyropho	sphokinase	Down Regulated
accD	946796	Acetyl-coenzyme A carboxylase carboxyl transferase subunit	beta	Down Regulated
ydaM	945909	Uncharacterized protein ydaM		Up Regulated
uxuB	946795	D-mannonate oxidoreductase		Up Regulated
ybhJ	945380	Uncharacterized protein ybhJ	['Energy production and conversion', ' Energy metabolism carbon ', 'glycolysis', 'tricarboxylic acid cycle']	Up Regulated
deoC	948902	Deoxyribose-phosphate aldolase		Down Regulated
dedA	946798	Protein dedA		Down Regulated
deoA	948901	Thymidine phosphorylase		Down Regulated
toIQ	948900	Protein tolQ		Down Regulated
pheT	945382	Phenylalanyl-tRNA synthetase beta chain		Down Regulated
nuoG	946762	NADH-quinone oxidoreductase; NADH-quinone oxidoreducta	ise subunit G	Down Regulated
ydcD	946958	Uncharacterized protein ydcD	['Cell wall membrane envelope biogenesis']	Up Regulated
CARA	949025	Carbamoyl-phosphate synthase small chain		Temporarly Up Regul
yacL	944809	UPF0231 protein yacL		Temporarly Up Regul
nuoA	946764	NADH-quinone oxidoreductase subunit; NADH-quinone oxido	preductase subunit A	Down Regulated
yehU	949027	Inner membrane protein yehU		Ambiguous/Unsorted
mutM	946765	Formamidopyrimidine-DNA glycosylase		Temporarly Up Regul
idi	949020	Isopentenyl-diphosphate Delta-isomerase		Up Regulated
yfbS	946766	Uncharacterized transporter yfbS	n	Temporarly Down Re
yfbU	946767	UPF0304 protein yfbU	М	Down Regulated
selD	946768	Selenide, water dikinase		Down Regulated
IrhA	946769	Probable HTH-type transcriptional regulator IrhA		Up Regulated
ftsL	944803	Cell division protein ftsL		Ambiguous/Unsorted
yicl	948169	Alpha-xylosidase		Temporarly Up Regul
fruR	944804	Fructose repressor		Ambiguous/Unsorted
yfdC	944801	Inner membrane protein yfdC		Up Regulated
gltS	948166	Sodium/glutamate symport carrier protein		Ambiguous/Unsorted
ydfJ	944802		_1671; ECDH10B_1674; ECH74115_2159; EC55989_1684; ECS88_1623; EcE24377A_1752; EcSMS35_1621; ECED1_1674; EC	
fkpB	944807	FKBP-type 16 kDa peptidyl-prolyl cis-trans isomerase		Down Regulated
nuoH	946761	NADH-quinone oxidoreductase subunit H		Down Regulated Down Regulated
recG	948162	ATP-dependent DNA helicase recG		Down Regulated Down Regulated
	948160	DNA-directed RNA polymerase subunit omega		
rpoZ trmH				Temporarly Down Re
	948161	tRNA guanosine-2'-O-methyltransferase		Temporarly Down Re
maeA	946031	NAD-dependent malic enzyme		Down Regulated
ydhF	946960	Oxidoreductase ydhF		Up Regulated
adhP	946036	Alcohol dehydrogenase, propanol-preferring		Temporarly Up Regul
nuoC	946759	NADH-quinone oxidoreductase subunit C/D		Down Regulated
ackA	946775	Acetate kinase		Down Regulated
yfbV	946774	UPF0208 membrane protein yfbV		Up Regulated
ymdA	949012	Uncharacterized protein ymdA	U	Down Regulated

yfbT	946777	Phosphatase yfbT		Down Regulated
pta	946778	Phosphate acetyltransferase		Down Regulated
yicN	948173	Uncharacterized protein yicN	[' Carbon compound utilization', 'regulation of transcription']	Temporarly Down Re
yicL	948176	Uncharacterized inner membrane transporter yicL	[regulation of cell shape']	Ambiguous/Unsorted
nlpA	948175	Lipoprotein 28		Down Regulated
ygfT	949018	Uncharacterized protein ygfT	[relectron transport']	Temporarly Up Regul
yicH	948171	Uncharacterized protein yicH	[Energy production and conversion']	Up Regulated
IspA	944800	Lipoprotein signal peptidase		Down Regulated
yfgD	946974	Uncharacterized protein yfgD	0	Temporarly Down Re
rhsE	946026	ECDH10B_1586; EC55989_0238; b1456		Ambiguous/Unsorted
ymbA	946972	Uncharacterized lipoprotein ymbA	['Amino acid transport and metabolism']	Temporarly Up Regul
uraA	946978	Uracil permease		Down Regulated
galS	949043	HTH-type transcriptional regulator galS		Temporarly Up Regul
nuoE	946746	NADH-quinone oxidoreductase subunit E	The construction of the control of t	Down Regulated
yeiB	949044	Uncharacterized protein yeiB	['two component signal transduction system phosphorelay ']	Down Regulated
yeiG	949045 949046	S-formylglutathione hydrolase yeiG		Down Regulated
artQ	949046	Arginine transport system permease protein artQ Coenzyme A biosynthesis bifunctional protein coaBC		Down Regulated Down Regulated
dfp menE	949047	2-succinylbenzoateCoA ligase		Temporarly Up Regul
gltl	946938	Glutamate/aspartate periplasmic-binding protein		Temporarly Up Regul
ftsQ	944823	Cell division protein ftsQ		Down Regulated
yadG	944833	Uncharacterized ABC transporter ATP-binding protein yadG		Down Regulated Down Regulated
can	944832	Carbonic anhydrase 2		Down Regulated Down Regulated
gnsb	946054	Protein gnsB		Down Regulated
secM	944831	Secretion monitor		Down Regulated
gcd	944830	Quinoprotein glucose dehydrogenase		Down Regulated
tmcA	946053	Uncharacterized protein ypfl		Ambiguous/Unsorted
ivbL	948181	IIvBN operon leader peptide		Temporarly Up Regul
nuoB	946738	NADH-quinone oxidoreductase subunit B		Down Regulated
rpmB	946941	50S ribosomal protein L28		Down Regulated
yfaZ	946739	Uncharacterized protein yfaZ		Temporarly Up Regul
nuol	946757	NADH-quinone oxidoreductase subunit I		Down Regulated
ubiX	949033	3-octaprenyl-4-hydroxybenzoate carboxy-lyase		Down Regulated
purE	949031	Phosphoribosylaminoimidazole carboxylase catalytic subunit		Down Regulated
yfcE	946755	Phosphodiesterase yfcE		Down Regulated
nuoJ	946756	NADH-quinone oxidoreductase subunit J		Down Regulated
nuoF	946753	NADH-quinone oxidoreductase subunit F		Down Regulated
yeiA	949037	Uncharacterized protein yeiA	[[Nucleotide transport and metabolism']	Up Regulated
ppiB	949038	Peptidyl-prolyl cis-trans isomerase B; Peptidyl-prolyl cis-trans	s isomerase	Down Regulated
nadE	946946	NH(3)-dependent NAD(+) synthetase		Down Regulated
thyA	949035	Thymidylate synthase		Down Regulated
murD	944818	UDP-N-acetylmuramoylalanineD-glutamate ligase		Down Regulated
surA	944812	Chaperone surA		Down Regulated
murF	944813	UDP-N-acetylmuramoyl-tripeptideD-alanyl-D-alanine ligase		Down Regulated
mraY wzzE	944814 944815	Phospho-N-acetylmuramoyl-pentapeptide-transferase		Down Regulated
		Lipopolysaccharide biosynthesis protein wzzE		Down Regulated
uidC ddpB	944820 946044	Membrane-associated protein uidC Probable D,D-dipeptide transport system permease protein of	L doB	Up Regulated Down Regulated
secA	946044	Protein translocase subunit secA	лары 	
ibpB	948192	Small heat shock protein ibpB		Down Regulated Up Regulated
yidE	948193	Putative transport protein yidE		Up Regulated
ypfJ	946956	Uncharacterized protein ypfJ		Temporarly Up Regul
bamC	946954	Lipoprotein 34	<u> </u>	Down Regulated
gcvR	946950	Glycine cleavage system transcriptional repressor		Temporarly Up Regul
vebQ	946048	Uncharacterized transporter vebQ		Temporarly Down Re
yfcN	944847	UPF0115 protein yfcN		Temporarly Down Re
ybgJ	945311	Uncharacterized protein ybgJ	ln	Down Regulated
sucC	945312	Succinyl-CoA ligase [ADP-forming] subunit beta	M .	Down Regulated
yadB	944846	Glutamyl-Q tRNA(Asp) synthetase; Insertion element IS150	uncharacterized 14 kDa protein	Ambiguous/Unsorted
hrpB	944845	ATP-dependent RNA helicase hrpB	· F · · · ·	Temporarly Down Re
ybgK	945317	Uncharacterized protein ybgK	in .	Down Regulated
lpxA	944849	Acyl-[acyl-carrier-protein]UDP-N-acetylglucosamine O-acyl	transferase	Down Regulated
	948978	Uncharacterized protein yigR	ln	Temporarly Up Regul
yjgR rffD	948977	UDP-N-acetyl-D-mannosamine dehydrogenase		Down Regulated
glpQ	946725	Glycerophosphoryl diester phosphodiesterase		Temporarly Up Regul

nin D	040072	Dutativa DNA isyantasa from lambdaid prophaga Da	Lin Domilated
pinR	948973	Putative DNA-invertase from lambdoid prophage Rac	Up Regulated
coaE	949060 946715	Dephospho-CoA kinase	Up Regulated
clcA		H(+)/Cl(-) exchange transporter clcA	Down Regulated
dksA	944850	DnaK suppressor protein	Temporarly Up Regul
sucB	945307	Dihydrolipoyllysine-residue succinyltransferase component of 2-oxoglutarate dehydrogenase complex	Down Regulated
hisJ	945309	Histidine-binding periplasmic protein	Temporarly Down Re
lpd Lttn V	944854	Dihydrolipoyl dehydrogenase	Down Regulated
HtpX	946076	Probable protease htpX	Ambiguous/Unsorted
ydhC	946077	Inner membrane transport protein ydhC	Down Regulated
rnhB aceE	944852 944834	Ribonuclease HII	Down Regulated Down Regulated
	945324	Pyruvate dehydrogenase E1 component	
lptG		Lipopolysaccharide export system permease protein lptG	Ambiguous/Unsorted
sdhD	945322 944839	Succinate dehydrogenase hydrophobic membrane anchor subunit 3-methyl-2-oxobutanoate hydroxymethyltransferase	Down Regulated Down Regulated
panB	945328	Protein toIR	
toIR			Down Regulated
yjhQ	949059		Up Regulated
csgA	949055	Major curlin subunit	Up Regulated
glpC trxB	946735 949054	Anaerobic glycerol-3-phosphate dehydrogenase subunit C Thioredoxin reductase	Up Regulated
			Down Regulated
menC lpxH	946734 949053	o-succinylbenzoate synthase	Down Regulated
•	949053	UDP-2,3-diacylglucosamine hydrolase	Down Regulated
lrp	949051	Leucine-responsive regulatory protein	Down Regulated
cydD		ATP-binding/permease protein cydD	Down Regulated
rluE	945701	Pseudouridine synthase; Ribosomal large subunit pseudouridine synthase E	Down Regulated
hipA	946064	Protein hipA	Temporarly Up Regul
ygbL	947197	Putative aldolase class 2 protein ygbL	Temporarly Up Regul
manA	944840	Mannose-6-phosphate isomerase	Temporarly Down Re
ybgL	945318	UPF0271 protein ybgL	Down Regulated
pphB	947196	Serine/threonine-protein phosphatase 2	Temporarly Up Regul
ygbK	947199	Uncharacterized protein ygbK ["Carbohydrate transport and metabolism"]	Up Regulated
ydeJ	946067	Protein ydeJ Uncharacterized protein yacH ["Signal transduction mechanisms", 'Type of regulation', 'signal transduction']	Up Regulated
yacH	944868		Up Regulated
argP	944867 948997	Chromosome initiation inhibitor	Ambiguous/Unsorted
caiB	946700	Crotonobetainyl-CoA:carnitine CoA-transferase Ecotin	Up Regulated Down Regulated
eco	949088	Endoribonuclease symE	
symE ymdB	946987	UPF0189 protein ymdB	Up Regulated Ambiguous/Unsorted
bamB	946982	Lipoprotein yfgL	Down Regulated
	946983	GTP-binding protein engA	Down Regulated Down Regulated
der yfgJ	946984	Uncharacterized protein rigg 'Type of regulation'	Ambiguous/Unsorted
cdsA	944876	Phosphatidate cytidylytransferase	Temporarly Down Re
dnaE	944877	DNA polymerase III subunit alpha	Down Regulated
yfgG	946980	Uncharacterized protein yfgG ["Nucleotide transport and metabolism"]	Up Regulated
	946981	UPF0070 protein yfgM	Down Regulated
yfgM yaeR	944875	Uncharacterized protein yaeR	Ambiguous/Unsorted
glnH	944872	Glutamine-binding periplasmic protein	Down Regulated
flhE	946094	Flagellar protein flhE	Down Regulated Down Regulated
prc	946096	Trajespecific protease	Down Regulated
bamA	944870	Outer membrane protein assembly factor yaeT	Down Regulated
rseP	944871	Regulator of sigma E protease	Down Regulated
ynfN	946097	Uncharacterized protein ynfN []	Up Regulated
osmE	945305	Osmotically-inducible lipoprotein E	Temporarly Up Regul
sucA	945303	2-oxoglutarate dehydrogenase E1 component	Down Regulated
ulaF	948711	L-ribulose-5-phosphate 4-epimerase ulaF	Up Regulated
aer	945301	Aerotaxis receptor	Up Regulated
erpA	944857	Iron-sulfur cluster insertion protein erpA	Up Regulated
yjfY	948713	Uncharacterized protein yif'	Temporarly Up Regul
kdpE	945302	KDP operon transcriptional regulatory protein kdpE	Temporarly Up Regul
yciM	944858	Month operant varies introduced by Protein Rup.	Ambiguous/Unsorted
glpA	946713	Anaerobic glycerol-3-phosphate dehydrogenase subunit A	Up Regulated
menF	946712	Menaquinone-specific isochorismate synthase	Ambiguous/Unsorted
sbcC	949076	Nuclease sbcCD subunit C	Down Regulated
apbE	946711	Thiamine biosynthesis lipoprotein apbE	Temporarly Down Re
IoIA	948989	Outer-membrane lipoprotein carrier protein	Down Regulated
betA	945716	Choline dehydrogenase	Down Regulated
2001	10 107 10	onomio donyarogonaco	- Committee Galatea

xdhD	949079	Probable hypoxanthine oxidase xdhD		Up Regulated
flgH	946996	Flagellar L-ring protein		Down Regulated
sseA	946993	3-mercaptopyruvate sulfurtransferase		Temporarly Up Regul
glpT	946704	Glycerol-3-phosphate transporter		Up Regulated
sseB	946994	Protein sseB		Down Regulated
ulaR	948706	HTH-type transcriptional regulator ulaR		Up Regulated
rodZ	946992	Cytoskeleton protein rodZ		Temporarly Down Re
acnB	944864	Aconitate hydratase 2		Down Regulated
yjfP	948707	Esterase yjfP		Ambiguous/Unsorted
yjfN	948709	Uncharacterized protein yjfN	[' Building block biosynthesis', 'ion transport']	Up Regulated
tsf	944866	Elongation factor Ts		Down Regulated
pinQ	946088	Putative DNA-invertase from lambdoid prophage Qin		Up Regulated
yebR	946086	UPF0067 protein yebR		Down Regulated
skp	944861	Chaperone protein skp		Down Regulated
dcp	946084	Peptidyl-dipeptidase dcp		Ambiguous/Unsorted
fabZ	944888	(3R)-hydroxymyristoyl-[acyl-carrier-protein] dehydratase		Down Regulated
yhfY	947888	Uncharacterized protein yhfY	[Replication recombination and repair, ' Carbon compound utilization']	Temporarly Up Regul
yhfX	947889	Uncharacterized protein yhfX	[[Replication recombination and repair', ' Carbon compound utilization']	Up Regulated
lpxD	944882	UDP-3-O-[3-hydroxymyristoyl] glucosamine N-acyltransferas	te	Down Regulated
map	947882	Methionine aminopeptidase	nestein.	Ambiguous/Unsorted
DHAR	945743	PTS-dependent dihydroxyacetone kinase operon regulatory		Up Regulated
yjiX	949090	Uncharacterized protein yjiX	['Signal transduction mechanisms', 'Adaptation to stress', 'response to stress']	Down Regulated
cysG	947880 945746	Siroheme synthase Protein umuD		Down Regulated
umuD				Up Regulated
minE	945740	Cell division topological specificity factor		Down Regulated Down Regulated
kdsB folM	945539 949096	3-deoxy-manno-octulosonate cytidylyltransferase		
hupB	949096	Dihydrofolate reductase folM DNA-binding protein HU-beta		Temporarly Up Regul Ambiguous/Unsorted
rbfA	949095	Ribosome-binding factor A		Down Regulated
ycbC	9475546	Uncharacterized protein ycbC	['regulation of cell shape', 'peptidoglycan biosynthetic process']	Ambiguous/Unsorted
infB	947684	Translation initiation factor IF-2	[regulation of cell shape, peptidogycan biosynthetic process]	Down Regulated
arnA	947683	Bifunctional polymyxin resistance protein arnA		Up Regulated
mukF	945548	Chromosome partition protein mukF		Down Regulated
nusA	947682	Transcription elongation protein nusA		Down Regulated
smtA	945547	Protein smtA		Down Regulated
uxaB	945542	Altronate oxidoreductase		Up Regulated
ycbB	945541	Uncharacterized protein ycbB	['DNA replication']	Temporarly Up Regul
rimP	947680	b3170	[Div represent]	Down Regulated
yegX	945543	Uncharacterized protein yegX	['translation']	Up Regulated
nuoL	945540	NADH-quinone oxidoreductase subunit L		Down Regulated
rlmE	947689	b3179		Down Regulated
yhbY	947688	RNA-binding protein yhbY		Down Regulated
truB	947687	tRNA pseudouridine synthase B		Down Regulated
sufD	944878	Protein sufD		Up Regulated
rpsO	947686	30S ribosomal protein S15		Down Regulated
PabA	947873	Para-aminobenzoate synthase glutamine amidotransferase	component II	Ambiguous/Unsorted
metQ	944893	D-methionine-binding lipoprotein metQ		Down Regulated
rpsB	947874	30S ribosomal protein S2		Down Regulated
accA	944895	Acetyl-coenzyme A carboxylase carboxyl transferase subuni	it alpha	Down Regulated
yejH	945529	Uncharacterized protein yejH	[' Building block biosynthesis', 'amino acid biosynthetic process']	Temporarly Down Re
serC	945527	Phosphoserine aminotransferase		Down Regulated
pnp	947672	Polyribonucleotide nucleotidyltransferase		Down Regulated
yhbQ	947671	UPF0213 protein yhbQ		Up Regulated
deaD	947674	Cold-shock DEAD box protein A		Down Regulated
cmk	945535	Cytidylate kinase		Temporarly Down Re
nlpl	947673	Lipoprotein nlpl		Temporarly Down Re
ihfB	945533	Integration host factor subunit beta		Down Regulated
ycal	945531	Uncharacterized protein ycal	<u>[D</u>	Up Regulated
rssA	945725	NTE family protein rssA		Temporarly Down Re
mtr	947675	Tryptophan-specific transport protein		Temporarly Up Regul
yhbW	947677	Uncharacterized protein yhbW	μ	Down Regulated
ymgE	945760	UPF0410 protein ymge		Up Regulated
clpA	945764	ATP-dependent Clp protease ATP-binding subunit clpA		Temporarly Up Regul
pth	945765	Peptidyl-tRNA hydrolase		Ambiguous/Unsorted
fabD	945766	Malonyl CoA-acyl carrier protein transacylase		Down Regulated

RNA Function

ycgV	945767	Uncharacterized protein ycgV ['Amino acid transport and metabolism', ' Building block biosynthesis']	Up Regulated
trpB	945768	Tryptophan synthase beta chain	, million and manager, and metabolism, Danism group stock st	Down Regulated
rlmL	945564	Ribosomal RNA large subunit methyltransferase L		Down Regulated
gst	945758	Glutathione S-transferase		Up Regulated
ycbX	945563	Uncharacterized protein ycbX		Temporarly Down Re
rmf	945567	Ribosome modulation factor		Temporarly Up Regul
gph	947895	Phosphoglycolate phosphatase		Down Regulated
rpe	947896	Ribulose-phosphate 3-epimerase		Down Regulated
yhfZ	947897	Uncharacterized protein yhfZ	'Replication recombination and repair', ' Carbon compound utilization', 'DNA replication']	Up Regulated
dadA	945752	D-amino acid dehydrogenase small subunit		Temporarly Down Re
sufB	945753	Protein sufB		Temporarly Up Regul
mukB	945549	Chromosome partition protein mukB		Down Regulated
IdcA	945756	Muramoyltetrapeptide carboxypeptidase		Temporarly Down Re
treA	945757	Periplasmic trehalase; Putative periplasmic trehalase		Up Regulated
dam	947893	DNA adenine methylase		Down Regulated
trpS	947894	Tryptophanyl-tRNA synthetase		Down Regulated
ompF	945554	Outer membrane protein F		Down Regulated
dhaM	945749	PTS-dependent dihydroxyacetone kinase, phosphotransferas		Down Regulated
dhaL	945748	PTS-dependent dihydroxyacetone kinase, ADP-binding subur	ill driat.	Down Regulated
glmM	947692	Phosphoglucosamine mutase		Down Regulated
aspC	945553	Aspartate aminotransferase		Down Regulated
foIP	947691 945552	Dihydropteroate synthase	"Nucle of ide transport and metabolisms! Duilding blook bigs unthe sign."	Down Regulated
ycbW	945552		"Nucleotide transport and metabolism", ' Building block biosynthesis']	Ambiguous/Unsorted Down Regulated
dhaK	947693	PTS-dependent dihydroxyacetone kinase, dihydroxyacetone-	DITUING SUDUING UNIAN	
dacB	947693	D-alanyl-D-alanine carboxypeptidase dacB		Down Regulated
ybaN greA	945558	Inner membrane protein ybaN Transcription elongation factor; Transcription elongation facto	or groß	Down Regulated
yhdJ	947695	Uncharacterized adenine-specific methylase yhdJ	I GIEA	Down Regulated Up Regulated
yhbE	947699	Uncharacterized inner membrane transporter yhbE	Type of regulation'	Down Regulated
ycbL	945551	Uncharacterized protein ycbL	Type of regulation j	Temporarly Down Re
mukE	945550	Chromosome partition protein mukE		Down Regulated
pyrC	945787	Dihydroorotase		Down Regulated
yjiM	945789	Uncharacterized protein yiiM		Up Regulated
ychA	945784	Protein sirB1		Down Regulated
yeeO	946506	Uncharacterized transporter yeeO	1	Temporarly Up Regul
hisF	946516	Imidazole glycerol phosphate synthase subunit hisF		Down Regulated
agaA	947646		9; ECIAI1 3285; ECDH10B 3308; ECIAI39 3636; b3135; ECS88 3523	Ambiguous/Unsorted
garP	947642	Probable galactarate transporter		Temporarly Up Regul
yeeR	946512	Inner membrane protein yeeR		Down Regulated
dmsC	945502	Anaerobic dimethyl sulfoxide reductase chain C		Up Regulated
rihA	945503	Pyrimidine-specific ribonucleoside hydrolase rihA		Down Regulated
cydC	945504	ATP-binding/permease protein cydC		Temporarly Up Regul
grxB	946926	Glutaredoxin-2		Temporarly Up Regul
yfiH	947089	UPF0124 protein yfiH		Down Regulated
hemA	945777	Glutamyl-tRNA reductase		Temporarly Down Re
ycgM	946115	Uncharacterized protein ycgM		Ambiguous/Unsorted
aroF	947084	Phospho-2-dehydro-3-deoxyheptonate aldolase, Tyr-sensitive		Down Regulated
IoIB	945775	Outer-membrane lipoprotein IoIB		Down Regulated
ispE	945774	4-diphosphocytidyl-2-C-methyl-D-erythritol kinase		Down Regulated
bamD	947086	UPF0169 lipoprotein yfiO		Down Regulated
agp	945773	Glucose-1-phosphatase		Up Regulated
prs	945772	Ribose-phosphate pyrophosphokinase	Es author D	Down Regulated
rluD	947087	Pseudouridine synthase; Ribosomal large subunit pseudourid	line synthase D	Ambiguous/Unsorted
cysB	945771	HTH-type transcriptional regulator cysB		Ambiguous/Unsorted
ychM	945770	Putative sulfate transporter ychM		Ambiguous/Unsorted
agaR	947636	Putative aga operon transcriptional repressor		Up Regulated
yhaV	947638	Uncharacterized protein yhaV		Up Regulated
cbl	946502	HTH-type transcriptional regulator cbl		Down Regulated
nac	946501 946500	Nitrogen assimilation regulatory protein nac Peroxidase vcdB		Down Regulated
efeB	946500	P-protein		Down Regulated Down Regulated
pheA pheL	947081	Phe operon leader peptide		Up Regulated
yfeZ	947080	Inner membrane protein yfeZ		Down Regulated
uxuA	947082	Mannonate dehydratase		Temporarly Up Regul
уреА	946917	Acetyltransferase ypeA		Down Regulated
JPON	0 100 11	, iostylia alloror doo yport		- Strii i togalatea

bisC	946915	Biotin sulfoxide reductase		Down Regulated
yfeX	946913	Uncharacterized protein yfeX	[Building block biosynthesis']	Temporarly Down Re
pflA		Pyruvate formate-lyase 1-activating enzyme		Temporarly Up Regul
ypjD	947098	Inner membrane protein ypjD		Down Regulated
trpC	945519	Tryptophan biosynthesis protein trpCF		Down Regulated
rspB		Starvation-sensing protein rspB		Up Regulated
rpIS		50S ribosomal protein L19		Down Regulated
sufC		Probable ATP-dependent transporter sufC		Temporarly Up Regul
trmD	947099	tRNA (guanine-N(1)-)-methyltransferase		Down Regulated
ydhK	946528	Uncharacterized transporter ydhK	[' Type of regulation']	Temporarly Down Re
sbcB		Exodeoxyribonuclease I		Temporarly Down Re
hisD		Histidinol dehydrogenase		Temporarly Up Regul
yraL	947664	UPF0011 protein yraL		Ambiguous/Unsorted
rsml	947664	UPF0011 protein yraL	TO all well prompting a pugling higgs point. His problem appended higgs point in a problem.	Ambiguous/Unsorted
yadE yraQ	946536 947668	Uncharacterized protein yadE UPF0718 protein yraQ	[Cell wall membrane envelope biogenesis', 'lipopolysaccharide biosynthetic process']	Up Regulated Up Regulated
yajO	946903	Uncharacterized oxidoreductase yaiO	n	Temporarly Up Regul
ycaK	945520	Uncharacterized NAD(P)H oxidoreductase ycaK	III 'amino acid transport'	Up Regulated
yajG	945521	Uncharacterized lipoprotein yajG	[Posttranslational modification protein turnover chaperones', 'Electrochemical potential driven transporters']	Down Regulated
yraN	947662	UPF0102 protein yraN	Troductional modification protein turnover draperories, Electrodrientical potential driven transporters	Down Regulated
yraM	947663	Uncharacterized protein yraM	['DNA replication']	Down Regulated
ivy	946530	Inhibitor of vertebrate lysozyme	<u> </u>	Temporarly Up Regul
ycaP	945525	UPF0702 transmembrane protein ycaP		Ambiguous/Unsorted
hcr		NADH oxidoreductase hcr		Up Regulated
diaA	947661	DnaA initiator-associating protein diaA		Down Regulated
rsxC		Electron transport complex protein rnfC		Down Regulated
rhmT	945796	Inner membrane transport protein yfaV		Ambiguous/Unsorted
yfaV	945796	Inner membrane transport protein yfaV		Ambiguous/Unsorted
yafS	944903	Uncharacterized protein yafS	['DNA replication']	Down Regulated
serS	945506	Seryl-tRNA synthetase		Temporarly Down Re
rarA	945505	Replication-associated recombination protein A		Down Regulated
ychN	945794	Protein ychN		Temporarly Up Regul
dkgB		2,5-diketo-D-gluconic acid reductase B		Up Regulated
yafJ		Putative glutamine amidotransferase yafJ		Ambiguous/Unsorted
queA		S-adenosylmethionine:tRNA ribosyltransferase-isomerase		Down Regulated
cobU		Bifunctional adenosylcobalamin biosynthesis protein cobU		Up Regulated
ynfD	946133	Uncharacterized protein ynfD	[Energy production and conversion', ' Electrochemical potential driven transporters', 'response to stress']	Up Regulated
rsxD		Electron transport complex protein rnfD		Down Regulated
chaB		Cation transport regulator chaB		Temporarly Up Regul
chaA		Calcium/proton antiporter		Down Regulated
poxB		Pyruvate dehydrogenase [cytochrome]		Up Regulated
mreC hisA	947655 946521	Rod shape-determining protein mreC	paminal imidazala A carbovamida isomorana	Down Regulated Down Regulated
cobS		1-(5-phosphoribosyl)-5-[(5-phosphoribosylamino)methylidene Cobalamin synthase	earlinoj iriluazbie-4-carboxarniue isornerase	Temporarly Up Regul
	947656	Uncharacterized outer membrane usher protein yraJ	['Amino acid transport and metabolism', 'Coenzyme transport and metabolism', 'Central intermediary metabolism', 'translation']	Up Regulated
yraJ yeeE	946526	UPF0394 inner membrane protein yeeE	Chimic and transport and metabolism, Coenzyme transport and metabolism, Central intermediaty metabolism, translation	Down Regulated
yraH	947658	Uncharacterized fimbrial-like protein yraH		Up Regulated
hslJ	946525	Heat shock protein hslJ	<u>u</u>	Temporarly Up Regul
yafK	944910	Uncharacterized protein yafK	in	Temporarly Down Re
ycaD	945515	Uncharacterized MFS-type transporter ycaD	u ['translation']	Temporarly Up Regul
yhgE	947651	Uncharacterized protein yhgE		Temporarly Up Regul
focA	945513	Type-1 fimbrial protein, C chain; Probable formate transporte	™ er 1	Temporarly Down Re
hisC	946551	Histidinol-phosphate aminotransferase		Down Regulated
hisB	946552	Histidine biosynthesis bifunctional protein hisB		Down Regulated
mutL		DNA mismatch repair protein mutL		Temporarly Down Re
yjeT	948696	Uncharacterized protein yjeT	[Posttranslational modification protein turnover chaperones', 'purine nucleotide biosynthetic process']	Temporarly Down Re
rplJ		50S ribosomal protein L10		Down Regulated
hflK	948698	Protein hflK		Down Regulated
yjfC	948699	Uncharacterized protein yjfC	['electron transport', 'metabolic process']	Up Regulated
exuT		Hexuronate transporter		Up Regulated
rpIE		50S ribosomal protein L5		Down Regulated
rpsN		30S ribosomal protein S14		Down Regulated
rpsH		30S ribosomal protein S8		Down Regulated
umuC		Protein umuC		Up Regulated
rpIF	947803	50S ribosomal protein L6		Down Regulated

ntrD	046250	Protoco 2		Ambiguous/Unsorted
ptrB	946358 946357	Protease 2 UPF0076 protein yoaB		
yoaB				Down Regulated
pphA	946356	Serine/threonine-protein phosphatase 1		Up Regulated
cdh	948410	CDP-diacylglycerol pyrophosphatase		Temporarly Down Re
pdxJ	947039	Pyridoxine 5'-phosphate synthase		Temporarly Down Re
pdxA	944919	4-hydroxythreonine-4-phosphate dehydrogenase		Down Regulated
cyoD	944918	Cytochrome o ubiquinol oxidase protein cyoD		Down Regulated
yebW	946354	Uncharacterized protein yebW	Π	Up Regulated
queC	947034	Queuosine biosynthesis protein queC		Up Regulated
pfkA	948412	6-phosphofructokinase; 6-phosphofructokinase isozyme 1		Down Regulated
fieF	948413	Ferrous-iron efflux pump fieF		Temporarly Down Re
fpr	948414	FerredoxinNADP reductase		Temporarly Down Re
uspD	948415	Universal stress protein D		Temporarly Up Regul
yiiS	948416	UPF0381 protein yiiS		Temporarly Up Regul
hemE	948497	Uroporphyrinogen decarboxylase		Down Regulated
yebO	946350	Uncharacterized protein yebO		Down Regulated
yiiR	948417	Uncharacterized protein yiiR	['Amino acid transport and metabolism', ' Building block biosynthesis']	Up Regulated
hupA	948499	DNA-binding protein HU-alpha		Down Regulated
menA	948418	1,4-dihydroxy-2-naphthoate octaprenyltransferase		Down Regulated
nudC	948498	NADH pyrophosphatase		Ambiguous/Unsorted
yafQ	944911	Uncharacterized protein yafQ		Up Regulated
flu	946540	Antigen 43		Down Regulated
frdC	948680	Fumarate reductase subunit C		Temporarly Up Regul
zapA	947404	Cell division protein zapA		Temporarly Up Regul
yeeX	946541	UPF0265 protein yeeX		Ambiguous/Unsorted
hfq	948689	Protein hfg		Down Regulated
sbmC	946546	DNA gyrase inhibitory protein		Temporarly Up Regul
hisL	946547	His operon leader peptide		Ambiguous/Unsorted
hflX	948688	GTP-binding protein hflX		Down Regulated
hisG	946549	ATP phosphoribosyltransferase		Down Regulated
hisH	946544	Imidazole glycerol phosphate synthase subunit hisH		Down Regulated
yeeA	946545	Inner membrane protein yeeA		Up Regulated
ygbJ	947200	Uncharacterized oxidoreductase yqbJ	n	Up Regulated
rpml	946349	50S ribosomal protein L35	U	Down Regulated
rsmF	946348	Ribosomal RNA small subunit methyltransferase F		Temporarly Down Re
rplL	948489	50S ribosomal protein L7/L12		Down Regulated
	946539	UPF0033 protein yeeD		
yeeD dsbB	946344			Temporarly Up Regul
		Disulfide bond formation protein B	Wantington receptionation and renaid!	Down Regulated
yebV	946347	Uncharacterized protein yebV	[Replication recombination and repair]	Up Regulated
yoaH	946346	UPF0181 protein yoaH	1 0050 F0000 0740 F00140D 0740 F0144D 0000 F0140D 0700 HT00 00007 F055000 0004 F045 4+ 000 70057 H0	Down Regulated
yfiC	947047		01_3956; ECS88_2749; ECDH10B_2743; ECUMN_2898; ECIAI39_2780; UTI89_C2897; EC55989_2864; pC15-1a_066; Z3857; b2	
rplK	948484	50S ribosomal protein L11		Down Regulated
rpIA	948483	50S ribosomal protein L1		Down Regulated
rlmA	946340	b1822		Ambiguous/Unsorted
cpxR	948404	Transcriptional regulatory protein cpxR		Down Regulated
yobD	946343	UPF0266 membrane protein yobD		Down Regulated
rhtA	947045	Inner membrane transporter rhtA		Temporarly Up Regul
ygbM	947207	Protein ygbM		Temporarly Up Regul
manZ	946342	Mannose permease IID component		Temporarly Up Regul
rpoB	948488	DNA-directed RNA polymerase subunit beta		Down Regulated
bolA	947043	Protein bolA		Temporarly Up Regul
dinB	944922	DNA polymerase IV		Ambiguous/Unsorted
nusG	948485	Transcription antitermination protein nusG		Temporarly Down Re
yiiM	948406	Protein yiiM		Down Regulated
lерВ	947040	Signal peptidase I		Down Regulated
pgk	947414	Phosphoglycerate kinase		Down Regulated
mscS	947416	Small-conductance mechanosensitive channel		Down Regulated
cpsG	946574	Phosphomannomutase		Temporarly Down Re
fbaA	947415	Fructose-bisphosphate aldolase class 2		Down Regulated
ugd	946571	UDP-glucose 6-dehydrogenase		Temporarly Up Regul
argO	947418	Arginine exporter protein argO		Up Regulated
rpoE	947050	RNA polymerase sigma factor; RNA polymerase sigma-E fac	I ctor	Up Regulated
rpID	947818	50S ribosomal protein L4		Down Regulated
rpIC	947817	50S ribosomal protein L3		Down Regulated
	947410	Uncharacterized protein yggF	['Replication_recombination and repair', ' Central intermediary metabolism', 'DNA recombination']	Temporarly Down Re
yggF	341410	ononaracienzeu protein yggr	I nepiloation recombination and repair, Central intermediary metabolism, DNA recombination;	Tremporariy Down Re

yhaH	947617	Inner membrane protein yhaH	Temporarly Up Regul
hsIU	948430	ATP-dependent hsl protease ATP-binding subunit hslU	Down Regulated
yebC	946379	UPF0082 protein yebC	Down Regulated
ruvC	946378	Crossover junction endodeoxyribonuclease ruvC	Down Regulated
rpIB	947820	50S ribosomal protein L2	Down Regulated
metF	948432	Methylenetetrahydrofolate reductase; 5,10-methylenetetrahydrofolate reductase	Temporarly Up Regul
surE	947211	Multifunctional protein surE	Down Regulated
yciS	944936	Inner membrane protein yciS	Down Regulated
ruvB	946371	Holliday junction ATP-dependent DNA helicase ruvB	Temporarly Down Re
truD	947214	tRNA pseudouridine synthase D	Down Regulated
rseB	947054	Sigma-E factor regulatory protein rseB	Temporarly Down Re
rseA	947053	Sigma-E factor negative regulatory protein	Ambiguous/Unsorted
metB	948434	Cystathionine gamma-synthase	Down Regulated
znuA	946375	High-affinity zinc uptake system protein znuA	Ambiguous/Unsorted
ydeA	947218	UPF0401 protein yubL; Probable sugar efflux transporter; Sugar efflux transporter	Up Regulated
metJ	948435	Met repressor	Temporarly Up Regul
srmB	947055	ATP-dependent RNA helicase srmB	Down Regulated
rsmA	944939	b0051	Down Regulated
cysN	947219	Sulfate adenylyltransferase subunit 1	Down Regulated
yciH	947058	Uncharacterized protein yciH ['Translation ribosomal structure and biogenesis', 'translation']	Down Regulated
nagE	945292	PTS system N-acetylglucosamine-specific EIICBA component	Up Regulated
ycfQ	947425	Uncharacterized HTH-type transcriptional regulator ycfQ	Up Regulated
FldA	945293	Flavodoxin-1	Down Regulated
rdgB	947429	Nucleoside-triphosphatase; Nucleoside-triphosphatase rdgB	Down Regulated
insE	944952	Transposase insE for insertion sequence IS3A/B/C/D/E/fA/fit; Transposase insE for insertion sequence IS3	Up Regulated
fur	945295	Ferric uptake regulation protein	Ambiguous/Unsorted
entE	947426	Enterobactin synthetase component E	Temporarly Up Regul
ybaO	945091	Uncharacterized HTH-type transcriptional regulator ybaO Metabolism of other compounds'	Up Regulated
rpmC	947807	50S ribosomal protein L29	Down Regulated
rpIP	947806	50S ribosomal protein L16	Down Regulated Down Regulated
	947805	50S ribosomal protein L36 2; 50S ribosomal protein L36; 50S ribosomal protein L36 1	
rpmJ			Down Regulated
yegH	946566	UPF0053 protein yegH	Ambiguous/Unsorted
rpIR	947804	50S ribosomal protein L18	Down Regulated
murR	946568	b2427	Down Regulated
tktA	947420	Transketolase 1	Down Regulated
rplN	947809	50S ribosomal protein L14	Down Regulated
rpsQ	947808	30S ribosomal protein S17	Down Regulated
marA	947613	Multiple antibiotic resistance protein marA	Up Regulated
yqjF	947608	Inner membrane protein yqjF	Up Regulated
eda	946367	KHG/KDPG aldolase	Down Regulated
glpF	948422	Glycerol uptake facilitator protein	Up Regulated
zapB	948420	Cell division protein zapB	Up Regulated
purT	946368	Phosphoribosylglycinamide formyltransferase 2	Down Regulated
rpIV	947813	50S ribosomal protein L22	Down Regulated
rpsC	947814	30S ribosomal protein S3	Down Regulated
rpsS	947811	30S ribosomal protein S19	Down Regulated
cysC	947221	Adenylyl-sulfate kinase	Down Regulated
hsIV	948429	ATP-dependent protease hslV	Down Regulated
eamB	947065	Cysteine/O-acetylserine efflux protein	Temporarly Up Regul
ybgF	947227	Uncharacterized protein ybgF	Down Regulated
cytR	948427	HTH-type transcriptional repressor cytR	Down Regulated
exoX	946361	Exodeoxyribonuclease 10	Down Regulated
selU	947063	tRNA 2-selenouridine synthase	Up Regulated
maa	945096	Maltose O-acetyltransferase	Up Regulated
pagP	946360	Protein crcA	Up Regulated
kgtP	947069	Alpha-ketoqlutarate permease	Temporarly Up Regul
yebF	946363	Protein yebF	Up Regulated
adk	945097	Adenvlate kinase	Down Regulated
edd	946362	Phosphogluconate dehydratase	Temporarly Down Re
hha	945098	Hemolysin expression-modulating protein	Up Regulated
yfiD	947068	Autonomous glycyl radical cofactor	Temporarly Down Re
insF	944948	Transposase insF for insertion sequence IS3A/B/C/D/E/fA	Up Regulated
ung	947067	Uracil-DNA glycosylase	Down Regulated
htpG	945099	Chaperone protein htpG	Down Regulated Down Regulated
glpK	948423	Glycerol kinase	Up Regulated
Rihu	340423	Orycerol milase	op negulated

yebG	946364	Uncharacterized protein yebG	[Translation_ribosomal structure and biogenesis', 'Replication_recombination and repair']	Up Regulated
glpX	948424	Fructose-1,6-bisphosphatase class 2		Temporarly Up Regul
yfiF	947066	Uncharacterized tRNA/rRNA methyltransferase yfiF		Down Regulated
speA	947432	Biosynthetic arginine decarboxylase; Arginine decarboxylase		Down Regulated
vacJ	945582	Lipoprotein vacJ		Down Regulated
mlaA	945582	Lipoprotein vacJ		Down Regulated
CHIA	947837	Probable bifunctional chitinase/lysozyme		Up Regulated
mpaA	945969	Protein mpaA		Up Regulated
tufA	947838	Elongation factor Tu 2; Elongation factor Tu; Elongation facto	rTu1	Down Regulated
bfr	947839	Bacterioferritin		Up Regulated
cyoA	945080	Ubiquinol oxidase subunit 2 Uncharacterized protein yfaL		Down Regulated
yfaL	946595 947436	' '		Up Regulated
dtpA topB	947436	Tripeptide permease tppB Tripeptide permease tppB		Ambiguous/Unsorted Ambiguous/Unsorted
tppB cof	945089	HMP-PP phosphatase		Ambiguous/Unsorted
mdlB	945088	Multidrug resistance-like ATP-binding protein mdlB		Temporarly Down Re
glnK	945087	Nitrogen regulatory protein P-II 2		Down Regulated
yijO	948451	Uncharacterized HTH-type transcriptional regulator yijO	[' RNA related', ' Type of regulation', ' Genetic unit regulated', 'regulation of transcription DNA dependent']	Down Regulated
lon	945085	ATP-dependent protease La	Traversiated, Type or regulation, Scriptic unit regulated, regulation or transcription Drive approaching	Ambiguous/Unsorted
clpX	945083	ATP-dependent Clp protease ATP-binding subunit clpX		Ambiguous/Unsorted
argC	948455	N-acetyl-gamma-glutamyl-phosphate reductase		Temporarly Down Re
ykgA	944956		; c0416; ECIAI39 0385; ECIAI1 0298; ECUMN 0338; b0300; EC55989 0303; Z0376; ECDH10B 0289	Ambiguous/Unsorted
ycfJ	945977	Uncharacterized protein ycfJ		Temporarly Up Regul
argE	948456	Acetylornithine deacetylase		Down Regulated
ydcU	945976	Inner membrane ABC transporter permease protein ydcU		Temporarly Up Regul
rpmG	946318	50S ribosomal protein L33		Down Regulated
ррс	948457	Phosphoenolpyruvate carboxylase		Down Regulated
fusA	947847	Elongation factor G		Down Regulated
yijP	948458		I0B_4143; ECIAI1_4163; EC55989_4437; Z5513; ECS88_4410; UTI89_C4546; c4914; ECIAI39_3034	Down Regulated
rpsG	947846	30S ribosomal protein S7		Down Regulated
yeaQ	946311	UPF0410 protein yeaQ		Temporarly Up Regul
yeaY	946312		['Cell wall membrane envelope biogenesis', 'metabolic process']	Up Regulated
yqgE	947442	UPF0301 protein yqgE		Down Regulated
ompA	945571	Outer membrane protein A		Temporarly Up Regul
cusS	945978	Sensor kinase cusS; Sensor protein		Down Regulated
rpIM	947828	50S ribosomal protein L13		Down Regulated
ydbK	946587	Probable pyruvate-flavodoxin oxidoreductase Tellurite resistance protein tehB		Up Regulated
tehB	945979			Temporarly Down Re
gspB hspQ	947826 945578	Protein pioO Heat shock protein hspQ		Ambiguous/Unsorted Temporarly Up Regul
asmA	946582	Protein asmA		Down Regulated
trml	947448	tRNA (guanine-N(7)-)-methyltransferase		Up Regulated
mgsA	945574	Methylglyoxal synthase		Ambiguous/Unsorted
yggW	947446	Oxygen-independent coproporphyrinogen-III oxidase-like prof	tein vaaW	Down Regulated
mutY	947447	A/G-specific adenine glycosylase	199.	Down Regulated
scpA	945576	Protein sbm		Up Regulated
gshB	947445	Glutathione synthetase		Temporarly Down Re
iscU	947002	NifU-like protein		Temporarly Up Regul
LeuC	945076	3-isopropylmalate dehydratase large subunit		Down Regulated
gldA	948440	Glycerol dehydrogenase		Ambiguous/Unsorted
puuC	947003	Gamma-glutamyl-gamma-aminobutyraldehyde dehydrogenas	se	Down Regulated
ydbC	945980	Putative oxidoreductase ydbC		Ambiguous/Unsorted
hcaR	947000	Hca operon transcriptional activator		Down Regulated
yccS	947001	Inner membrane protein yccS		Down Regulated
puuB	945072	Gamma-glutamylputrescine oxidoreductase		Down Regulated
clpS	948443	ATP-dependent Clp protease adapter protein clpS		Up Regulated
ompX	944967	Outer membrane protein X		Up Regulated
yahN	944968	Uncharacterized membrane protein yahN		Up Regulated
tesB	945074	Acyl-CoA thioesterase 2		Temporarly Up Regul
yahM	944969	Uncharacterized protein yahM	0	Temporarly Up Regul
emrE	948442	Multidrug transporter emrE		Up Regulated
cyoE	945073	Protoheme IX farnesyltransferase		Down Regulated
ydcA	945986		[Energy production and conversion', 'Energy production transport', 'Macromolecules cellular constituent biosynthesis', 'electron	
yoaA	946305	Probable ATP-dependent helicase yoaA	Transportation Duilding blook biggs with a sid DNA related Transfer and State IDNA related IDNA relat	Down Regulated
yhcN	947835	Uncharacterized protein yhcN	['Transcription', 'Building block biosynthesis', 'RNA related', 'Type of regulation', 'DNA recombination', 'transcription', 'regulation o	IJUP Regulated

thiD	946306	Thismine hinding peripleanie protein		Tomporarly Un Doguil
thiB	948446	Thiamine-binding periplasmic protein		Temporarly Up Regul
hemX	946307	Putative uroporphyrinogen-III C-methyltransferase Uncharacterized protein yeaL	п	Down Regulated
yeaL			μ	Temporarly Down Report Down Regulated
flgA	946300	Flagella basal body P-ring formation protein flgA		
mipA	946301	MITA-interacting protein		Down Regulated
ydeM	945981	Anaerobic sulfatase-maturating enzyme homolog ydeM		Up Regulated
iscS	947004	Cysteine desulfurase		Ambiguous/Unsorted
yeaE	946302	Uncharacterized protein yeaE	n	Temporarly Down Re
ydfK	947451	Uncharacterized protein ydfK		Up Regulated
yggN	947453	Uncharacterized protein yggN	[' Central intermediary metabolism']	Ambiguous/Unsorted
solA	944983	N-methyl-L-tryptophan oxidase		Down Regulated
flgJ	947456	Peptidoglycan hydrolase flgJ		Down Regulated
betl	944981	HTH-type transcriptional regulator betl		Up Regulated
prpR	944987	Propionate catabolism operon regulatory protein		Up Regulated
frmR	944986	Transcriptional repressor frmR		Down Regulated
ycdX	948477	Putative hydrolase ycdX		Down Regulated
sdaA	946331	L-serine dehydratase 1		Up Regulated
murB	948470	UDP-N-acetylenolpyruvoylglucosamine reductase		Down Regulated
manY	946332	Mannose permease IIC component		Temporarly Down Re
yncJ	945990	Uncharacterized protein yncJ		Ambiguous/Unsorted
qseE	947013	Putative sensor-like histidine kinase yfhK; Sensor protein		Down Regulated
glrK	947013	Putative sensor-like histidine kinase yfhK; Sensor protein		Down Regulated
panE	945065	2-dehydropantoate 2-reductase		Up Regulated
yebY	947014	Uncharacterized protein yebY	['translation']	Down Regulated
nudL	946330	Uncharacterized Nudix hydrolase nudL		Up Regulated
yoaE	946335	UPF0053 inner membrane protein yoaE		Temporarly Down Re
yheU	947863	UPF0270 protein yheU		Temporarly Down Re
trg	945995	Methyl-accepting chemotaxis protein III		Up Regulated
glnB	947016	Nitrogen regulatory protein P-II 1		Ambiguous/Unsorted
xseB	945069	Exodeoxyribonuclease 7 small subunit		Ambiguous/Unsorted
degS	947865	Protease degS		Down Regulated
pepT	946333	Peptidase T		Ambiguous/Unsorted
argD	947864	Succinylornithine transaminase; Acetylornithine/succinyldian	ninopimelate aminotransferase	Ambiguous/Unsorted
manX	946334	PTS system mannose-specific EIIAB component		Ambiguous/Unsorted
cspC	946339	Cold shock-like protein cspC		Down Regulated
crp	947867	Catabolite gene activator		Temporarly Down Re
rimL	945998	Ribosomal-protein-serine acetyltransferase		Up Regulated
coaA	948479	Pantothenate kinase		Temporarly Down Re
ydcH	945997	Uncharacterized protein ydcH	[' Channel type Transporters']	Temporarly Up Regul
tsgA	947869	Protein tsgA	[Similar type Transportate]	Ambiguous/Unsorted
yobF	946338	Uncharacterized protein yobF	RNA related', 'Type of regulation', 'Adaptation to stress', 'transcription', 'regulation of transcription DNA dependent'	Down Regulated
yhdH	947848	Putative quinone oxidoreductase yhdH	(*************************************	Temporarly Up Regul
slyX	947849	Protein slyX		Temporarly Up Regul
	947462	Uncharacterized protein yqgA	n	Up Regulated
yqgA	944990	Methylisocitrate lyase	<u>U</u>	Up Regulated
prpB torS	944990	Sensor protein torS; Sensor protein		Temporarly Down Re
frmB	943393	S-formylglutathione hydrolase frmB		Temporarly Down Re
cbpM	944991	Chaperone modulatory protein cbpM		Up Regulated
	945597	Cytosine permease		
codB				Down Regulated
glcC	947466	Glc operon transcriptional activator		Temporarly Down Ro
proY	945050	Proline-specific permease proY		Temporarly Down Re
frmA	944988	S-(hydroxymethyl)glutathione dehydrogenase		Down Regulated
argH	948463	Argininosuccinate lyase		Ambiguous/Unsorted
sthA	948461	Soluble pyridine nucleotide transhydrogenase		Down Regulated
nusB	945054	N utilization substance protein B homolog; N utilization subs	Tance protein B	Down Regulated
oxyR	948462	Hydrogen peroxide-inducible genes activator		Temporarly Down Re
glyA	947022	Serine hydroxymethyltransferase		Down Regulated
yphH	947023	Uncharacterized protein yphH	[Carbohydrate transport and metabolism', 'Transcription', 'carbohydrate metabolic process']	Ambiguous/Unsorted
murG	946321		pyrophosphoryl-undecaprenol N-acetylglucosamine transferase	Down Regulated
mdh	947854	Malate dehydrogenase		Ambiguous/Unsorted
ftsW	946322	Cell division protein ftsW		Temporarly Down Re
yajR	945058	Inner membrane transport protein yajR		Up Regulated
ddlB	946324	D-alanineD-alanine ligase B		Down Regulated
yheO yfhR	947851 945059	Uncharacterized protein yheO Uncharacterized protein yfhR	['protein folding']	Down Regulated Ambiguous/Unsorted

RNA Function

kefG	947857	Glutathione-regulated potassium-efflux system ancillary prote	ein kefG	Up Regulated
yheS	947856	Uncharacterized ABC transporter ATP-binding protein yheS		Temporarly Down Re
rnd	946328	Ribonuclease D		Ambiguous/Unsorted