Supplementary Table 1. Protein-protein interaction data reported in this study

Bait	Prey	MALDI-ToF	LC-MS/MS
abc	abc	Y	Υ
abc	b0817		Υ
abc	gyrB		Υ
abc	hycG		Υ
abc	ilvD		Υ
abc	perR		Υ
abc	tufB	Υ	
abc	wcaC		Υ
abc	yfeA		Υ
abc	yhjB		Υ
abc	yjhC		Υ
accA	accA	Υ	Y
accA	accD	Υ	Υ
accA	dnaK		Υ
accA	infB		Υ
accA	lysU		Υ
accA	rplC	Υ	
accA	rplR		Υ
accA	rplV		Υ
accA	rplW		Υ
accC	accC	Y	Υ
accC	accA		Υ
accC	ассВ	Υ	Υ
accC	accD		Υ
accC	cysM		Υ
accC	dnaK	Υ	
accC	dnaN	Υ	
accC	rplC	Υ	
accC	yhdY -		Y
accD	accD	Y	Y
accD	accA	Υ	Y
accD	dnaK		Y
accD	pepD		Y
accD	pgm		Y
accD	rfal		Y
accD	rpIU		Y
accD	rplV		Y
accD	rpmG		Y
accD	yegN		Y
accD	yhdA	V	Y
acpP	асрР	Y	Υ

acpP	aas	Υ	
acpP	accA		Υ
acpP	aceE	Y	
acpP	aceF	Υ	
асрР	acpS	Υ	Υ
асрР	aidB	Υ	Υ
acpP	b0968	Υ	
acpP	b4285	Υ	
асрР	fabB	Υ	
асрР	fabF	Υ	Υ
асрР	fabG	Υ	Υ
acpP	fabH	Υ	
асрР	fabZ	Υ	
асрР	glmU	Υ	Υ
acpP	hupB		Υ
acpP	IpdA	Υ	Υ
асрР	lpxD	Υ	Υ
acpP	malT	Υ	
асрР	mukB	Υ	
асрР	plsB	Υ	
acpP	ribF	Υ	
acpP	rplO		Υ
acpP	rplV		Υ
acpP	rpmB		Υ
acpP	rpsF		Υ
acpP	rpsG	Υ	Υ
acpP	rpsH		Υ
acpP	rpsM		Υ
асрР	secA	Υ	
асрР	spoT	Υ	
асрР	ybgC	Υ	Υ
acpP	ybgJ	Υ	
acpP	yhbY		Υ
асрР	yiiD	Υ	Υ
ada	ada	Υ	
adhC	adhC	Υ	Υ
adhE	adhE	Υ	Υ
adhE	rpsB		Υ
adiA	adiA	Υ	Υ
adiA	ftsJ		Υ
adiA	rplB	Υ	
adiA	rplO	Υ	
adiA	sucA		Υ
adiA	yahF		Υ
•			

adi A	ybal		Y
adiA	ybgl	V	Ϋ́
adk	adk	Y Y	Y
adk	rpoC	ľ	V
adk	ybdL	V	Y Y
agaY	agaY	Υ	Ϋ́
agaY	rplL	V	Y
ahpC	ahpC	Y Y	V
aidB	aidB		Y
aidB	aceF	Y	Y
aidB	dnaK	Y	Y
aidB	mopA	Y	Y
aidB	rfaD	Υ	V
aidB	rplA	V	Y
aidB	rpIC	Y	Y
aidB	rplM		Y
aidB	rpIS		Y
aidB	rplU		Y
aidB	rplV		Y
aidB	rpmB		Y
aidB	rpsB	V	Y
aidB	rpsD	Y	V
aidB	rpsE		Y
aidB	rpsF		Y
aidB	rpsG		Y Y
aidB	rpsL		Ϋ́
aidB	rpsN		
aidB aidB	rpsS		Y Y
aidB	rpsT		Ϋ́
	yhbY		
aidB <i>alaS</i>	yjgD <i>alaS</i>	V	Y
alaS	b1555	Y Y	
alaS	dnaK	Y	
alaS	rpsB	Ϋ́	
alaS	tufA	Ϋ́	
alaS	tufB	Y	
alaS alkA	alkA	Y	
alkA	aceF	Ϋ́	
ankA	acer apt	Y	Υ
apı argB	арі argB	Y	1
argR	argB argR	Y	
aroE	argit aroE	Y	Υ
aroE	yedE	,	Ϋ́
artP	yeu∟ artP	Y	Y
aiti	arti	ı	1

1			
artP	ppk		Υ
artP	topA		Υ
artP	yccA		Υ
asnS	asnS	Υ	
aspA	aspA	Y	Y
aspA	dnaK	Υ	Υ
aspA	secA	Υ	
aspS	aspS	Y	Y
aspS	asnS	Υ	
aspS	gltD	Υ	
atpA	atpA	Y	Y
atpA	atpD	Υ	
atpG	atpG	Y	Y Y
atpG	atpA	Υ	
atpG	atpC	Υ	Υ
atpG	atpD	Υ	
atpG	rplW		Υ
b0822	b0822	Y	Y
b0844	b0844	Y	
b0866	b0866	Υ	
b0866	rfaD	Υ	
b0947	b0947	Υ	Υ
b0947	cysl	Υ	
b0947	cysJ	Υ	Υ
b0959	b0959	Υ	
b0959	cobB	Υ	
b0959	prsA	Υ	Υ
b0965	b0965	Υ	Y
b0965	rpIL		Υ
b0968	b0968	Υ	
b1134	b1134	Υ	Υ
b1163	b1163	Υ	Υ
b1163	aceE	Υ	Υ
b1163	aceF		Υ
b1163	lpdA	Υ	Υ
b1163	mopA		Υ
b1163	rpIL		Υ
b1163	yfiD		Υ
b1192	b1192	Y	Υ
b1200	b1200	Y	Y
b1200	guaC	Υ	
b1200	yhiH		Υ
b1248	b1248		Y
b1248	aceE	Υ	Υ

b1248 lepA Y b1248 rplQ Y b1248 rpsJ Y b1327 b1327 Y b1327 aceE Y b1327 y Y b1327 y Y b1327 y Y b1337 b1337 Y b1341 b1341 Y b1341 lpdA Y b1341 tufB Y b1342 y Y b1399 b1399	b1248	aceF		Y
b1248 rpIQ Y b1248 rpsJ Y b1327 b1327 Y b1327 aceE Y b1337 b1337 Y b1341 b1341 Y b1341 aceE Y b1341 lpdA Y b1341 tufA Y b1342 Y Y b1398 prpIC <td></td> <td></td> <td></td> <td></td>				
b1248 rpsJ Y b1327 b1327 Y b1327 aceE Y b1337 b1337 Y b1341 b1341 Y b1341 aceE Y b1341 lpdA Y b1341 tufA Y b1341 tufB Y b1342 b1398 Y b1399 pplC Y b1399 rplC Y b1399 rpsC		-		
b1327 b1327 Y Y b1327 aceE Y b1337 b1337 Y Y b1341 b1341 Y Y b1341 aceE Y Y b1341 lpdA Y Y b1341 tufB Y Y b1342 tufB Y Y b1399 prpIC Y Y b1399 rpsI Y Y b1399 rpsG Y Y <t< td=""><td></td><td>•</td><td></td><td></td></t<>		•		
b1327 aceE Y b1337 b1337 Y b1341 b1341 Y b1341 aceE Y b1341 lpdA Y b1341 lpdA Y b1341 tufB Y b1344 y Y b1398 y Y b1399 ppIC Y b1399 rpIV Y b1399 rpsG Y b1399 rpsG Y b1399 rpsG Y b1399 rpsG		•	Υ	
b1337 b1337 Y Y b1341 b1341 Y b1341 aceE Y b1341 lpdA Y b1341 tufA Y b1341 tufB Y b1374 b1374 Y b1374 lpdA Y b1374 lpdA Y b1374 lpdA Y b1374 lpdA Y b1398 b1398 Y b1399 b1399 Y b1399 rplC Y b1399 rplO Y b1399 rplU Y b1399 rplU Y b1399 rpsG Y b1399 rpsG Y b1399 rpsG Y b1399 rpsG Y b1423 b1423 Y b1583 lpdA Y b1583 lpdA Y b1583<				
b1341 aceE Y b1341 lpdA Y b1341 tufA Y b1341 tufB Y b1374 b1374 Y Y b1374 b1374 Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1398 b1398 Y Y b1399 b1399 Y Y b1399 rplC Y Y b1399 rplO Y Y b1399 rplV Y Y b1399 rplV Y Y b1399 rplV Y Y b1399 rpsG Y Y b1399 rpsG Y Y b1583 b1583 Y Y b1583 lpdA Y Y b1583 lpdA Y Y b1598	b1337	b1337	Y	Y
b1341 aceF Y b1341 tufA Y b1341 tufB Y b1374 b1374 Y Y b1374 b1374 Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1398 b1398 Y Y b1399 b1399 Y Y b1399 rplC Y Y b1399 rplO Y Y b1399 rplU Y Y b1399 rplV Y Y b1399 rplV Y Y b1399 rpsG Y Y b1399 rpsG Y Y b1399 rpsG Y Y b1399 rpsL Y Y b1583 b1683 Y Y b1583 lpdA Y Y b1583 lpdA Y Y	b1341	b1341	Y	
b1341	b1341	aceE	Υ	
b1341 tufA Y b1341 tufB Y b1374 b1374 Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1398 b1398 Y Y b1399 b1399 Y Y b1399 rplC Y Y b1399 rplD Y Y b1399 rplU Y Y b1399 rplU Y Y b1399 rplU Y Y b1399 rplV Y Y b1399 rpsG Y Y b1399 rpsG Y Y b1399 rpsG Y Y b1423 b1423 Y Y b1583 lpdA Y Y b1583 lpdA Y Y b1583 lpdA Y Y b1593 Y Y Y	b1341	aceF	Υ	
b1341 tufB Y b1374 b1374 Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1374 lpdA Y Y b1398 b1398 Y Y b1399 b1399 Y Y b1399 rplC Y Y b1399 rplO Y Y b1399 rplU Y Y b1399 rpsG Y Y b1423 b1423 Y Y b1583 lpdA Y b1583 lpdA Y b1583 lpdA Y b1598 Y Y b1598 Y Y b1624 b1624 Y	b1341	lpdA	Υ	
b1374 b1374 Y Y b1374 aceF Y Y b1374 lpdA Y Y b1398 b1398 Y Y b1399 b1399 Y Y b1399 lpdA Y Y b1399 rplC Y Y b1399 rplM Y Y b1399 rplU Y Y b1399 rplU Y Y b1399 rpsG Y Y b1583 b1583 Y Y b1583 lpdA Y Y b1583 lpdA Y Y b1593 Y Y Y b1694 b1604 Y Y	b1341	tufA	Υ	
b1374 aceF Y Y b1374 lpdA Y b1398 b1398 Y b1399 b1399 Y Y b1399 lpdA Y Y b1399 lpdA Y Y b1399 rplC Y Y b1399 rplM Y Y b1399 rplU Y Y b1399 rplV Y Y b1399 rpsG Y Y b1583 b1583 Y Y b1583 lpdA Y Y b1583 lpdA Y Y b1593 b1593 Y Y b1604 b1604 Y Y b1624 b1624 Y Y b1624 prsA Y Y	b1341	tufB	Υ	
b1374 IpdA Y b1398 b1398 Y b1399 b1399 Y b1399 aceF Y b1399 rplC Y b1399 rplC Y b1399 rplO Y b1399 rplU Y b1399 rplU Y b1399 rpsG Y b1399 rpsG Y b1399 rpsL Y b1399 rpsG Y b1399 rpsL Y b1423 b1423 Y b1583 lpdA Y b1583 lpdA Y b1583 lpdA Y b1598 Y Y b1698 b1598 Y b1624 b1624 Y b1624 prsA Y b1647	b1374	b1374	Y	
b1398 b1398 Y b1399 b1399 Y b1399 aceF Y b1399 rplC Y b1399 rplC Y b1399 rplO Y b1399 rplU Y b1399 rplU Y b1399 rpsG Y b1399 rpsG Y b1399 rpsL Y b1399 rpsL Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1593 b1593 Y b1598 Y Y b1624 b1624 Y b1624 b1624 Y b1624 prsA Y b1647 b1649 Y				Υ
b1399 b1399 Y Y b1399 lpdA Y Y b1399 rplC Y Y b1399 rplM Y Y b1399 rplO Y Y b1399 rplU Y Y b1399 rplV Y Y b1399 rpsG Y Y b1399 rpsG Y Y b1399 rpsL Y Y b1399 rpsG Y Y b1399 rpsL Y Y b1583 b1583 Y Y b1583 lpdA Y Y b1593 b1593 Y Y b1598 b1598 Y Y b1624 b1624 Y Y b1624 b1624 Y Y b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y Y		•		
b1399 aceF Y b1399 rplC Y b1399 rplM Y b1399 rplM Y b1399 rplO Y b1399 rplU Y b1399 rplV Y b1399 rpsG Y b1399 rpsG Y b1399 rpsL Y b1399 rpsL Y b1583 b1583 Y b1583 lpdA Y b1593 Y Y b1598 b1598 Y b1604 b1604 Y b1624 cobB Y b1640 b1640 Y b1647 b1647 Y b1649 b1649 Y				
b1399 lpdA Y Y b1399 rplC Y b1399 rplM Y b1399 rplT Y b1399 rplU Y b1399 rpmB Y b1399 rpsG Y b1399 rpsG Y b1399 rpsL Y b1423 b1423 Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1583 lpdA Y b1593 Y Y b1598 b1593 Y b1604 b1604 Y Y b1624 b1624 Y Y b1624 prsA Y Y b1647 b1647 Y Y b1649 b1649 Y Y				Y
b1399 rpIC Y b1399 rpIM Y b1399 rpIO Y b1399 rpIU Y b1399 rpIV Y b1399 rpsG Y b1399 rpsG Y b1399 rpsL Y b1399 rpsL Y b1399 rpsL Y b1583 b1583 Y b1583 lpdA Y b1593 b1593 Y b1598 b1598 Y b1604 b1604 Y b1624 b1624 Y b1624 prsA Y b1647 b1647 Y b1649 b1649 Y				
b1399 rpIM Y b1399 rpIO Y b1399 rpIU Y b1399 rpIV Y b1399 rpmB Y b1399 rpsG Y b1399 rpsL Y b1399 rpsL Y b1399 rpsL Y b1399 rpsL Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1593 y Y b1593 y Y b1604 b1604 Y Y b1624 b1624 Y Y b1624 prsA Y Y b1647 b1647 Y Y b1649 b1649 Y Y		•	Υ	
b1399 rpIO Y b1399 rpIT Y b1399 rpIU Y b1399 rpIV Y b1399 rpmB Y b1399 rpsG Y b1399 rpsL Y b1423 b1423 Y Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1583 lpdA Y b1593 b1593 Y Y b1598 b1598 Y Y b1624 b1624 Y Y b1624 cobB Y b1624 prsA Y b1647 b1647 Y Y b1649 b1649 Y				
b1399 rpIT Y b1399 rpIU Y b1399 rpIV Y b1399 rpmB Y b1399 rpsG Y b1399 rpsL Y b1423 b1423 Y Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1583 lpdA Y b1593 b1593 Y Y b1598 b1598 Y Y b1624 b1624 Y Y b1624 cobB Y b1624 prsA Y b1649 b1649 Y		-		
b1399 rpIU Y b1399 rpIV Y b1399 rpsG Y b1399 rpsG Y b1399 rpsL Y b1399 rpsL Y b1399 rpsL Y b1423 Y Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1583 lpdA Y b1593 b1593 Y b1598 b1598 Y b1604 b1604 Y b1624 b1624 Y b1624 prsA Y b1647 b1647 Y b1649 b1649 Y		-		
b1399 rplV Y b1399 rpmB Y b1399 rpsG Y b1399 rpsL Y b1399 rpsL Y b1399 rpsL Y b1423 Y Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1583 lpdA Y b1593 b1593 Y b1598 b1598 Y b1604 b1604 Y b1624 b1624 Y b1624 prsA Y b1640 b1640 Y b1647 b1649 Y				
b1399 rpmB Y b1399 rpsG Y b1399 rpsL Y b1399 rpsL Y b1423 Y Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1593 b1593 Y b1598 b1598 Y b1604 b1604 Y b1624 b1624 Y b1624 prsA Y b1640 b1640 Y b1647 b1649 Y		•		
b1399 rpsG Y b1399 rpsL Y b1423 b1423 Y Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1583 ygiW Y b1593 b1593 Y b1598 b1598 Y b1604 b1604 Y b1624 b1624 Y b1624 cobB Y b1640 b1640 Y b1647 b1647 Y b1649 b1649 Y				
b1399 rpsL Y b1423 b1423 Y b1583 b1583 Y b1583 aceE Y b1583 lpdA Y b1583 ygiW Y b1593 b1593 Y b1598 b1598 Y b1604 b1604 Y b1624 b1624 Y b1624 prsA Y b1640 b1640 Y b1647 b1647 Y b1649 b1649 Y		-		
b1423 b1423 Y Y b1583 b1583 Y b1583 lpdA Y b1583 lpdA Y b1583 ygiW Y b1593 b1593 Y b1598 b1598 Y b1604 b1604 Y b1624 b1624 Y b1624 cobB Y b1624 prsA Y b1640 b1640 Y b1647 b1647 Y b1649 b1649 Y		•		
b1583 b1583 Y b1583 aceE Y b1583 lpdA Y b1583 ygiW Y b1593 b1593 Y Y b1598 b1598 Y Y b1604 b1604 Y Y b1624 b1624 Y Y b1624 cobB Y b1624 prsA Y b1640 b1640 Y Y b1647 b1649 Y Y		•	V	
b1583 aceE Y b1583 lpdA Y b1583 ygiW Y b1593 Y Y b1598 b1598 Y b1604 b1604 Y b1624 b1624 Y b1624 cobB Y b1624 prsA Y b1640 b1640 Y b1647 b1647 Y b1649 b1649 Y			Y	
b1583 lpdA Y b1583 ygiW Y b1593 b1593 Y Y b1598 b1598 Y Y b1604 b1604 Y Y b1624 b1624 Y Y b1624 cobB Y Y b1624 prsA Y Y b1640 b1640 Y Y b1647 b1649 Y Y			V	Y
b1583 ygiW Y b1593 b1593 Y Y b1598 b1598 Y Y b1604 b1604 Y Y b1624 b1624 Y Y b1624 cobB Y b1624 prsA Y b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y			ı	V
b1593 b1593 Y Y b1598 Y Y Y b1604 b1604 Y Y b1624 b1624 Y Y b1624 cobB Y Y b1624 prsA Y Y b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y Y		•		
b1598 b1598 Y Y b1604 b1604 Y Y b1624 b1624 Y Y b1624 cobB Y b1624 prsA Y b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y Y			V	
b1604 b1604 Y Y b1624 b1624 Y Y b1624 cobB Y b1624 prsA Y b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y Y				
b1624 b1624 Y Y b1624 cobB Y b1624 prsA Y b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y				
b1624 cobB Y b1624 prsA Y b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y				
b1624 prsA Y b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y			•	
b1640 b1640 Y Y b1647 b1647 Y Y b1649 b1649 Y			Υ	-
b1647 b1647 Y Y b1649 b1649 Y		•		Υ
b1649 b1649 Y				Υ
b1640 kdoD V				
DIO49 KUSD Y	b1649	kdsB		Υ

L4007			
b1667	b1667	Y	Υ
b1667	dnaK	Υ	Υ
b1667	dnaN	Υ	
b1668	b1668	Υ	Y
b1673	b1673	Y	
b1675	b1675	Y	
b1685	b1685	Y	
b1685	b1410	Υ	
b1685	rfaD	Υ	
b1685	rfaF	Υ	
b1685	rpsC	Υ	
b1685	rpsE	Υ	
b1685	rpsG	Υ	
b1685	spoT	Y	
b1685	tig	Υ	
b1685	tufA	Y	
b1731	b1731	Υ	Y
b1731	aceE		Υ
b1731	aceF	Υ	Υ
b1731	hns		Υ
b1731	rplC		Υ
b1731	rpoA	Y	Υ
b1731	rpoB	Y	Υ
b1731	rpoC	Y	Υ
b1731	rpoD	Y	Y
b1731	rpoZ	Υ	Y
b1731	rpsO		Y
b1731	tufA		Υ
b1741	b1741	Υ	
	_		
b1741	mopA	Y	
b1746	b1746	Y Y	
<i>b1746</i> b1746	<i>b1746</i> mutH	Y	Y
b1746 b1746 b1759	b1746 mutH b1759	Y Y	Y Y
b1746 b1746 b1759 b1759	b1746 mutH b1759 ygeA	Y Y Y	
b1746 b1746 b1759 b1759 b1770	b1746 mutH b1759 ygeA b1770	Y Y Y Y	
b1746 b1746 b1759 b1759 b1770 b1770	b1746 mutH b1759 ygeA b1770 lpdA	Y Y Y Y	Υ
b1746 b1746 b1759 b1759 b1770 b1770 b1773	b1746 mutH b1759 ygeA b1770 lpdA b1773	Y Y Y Y	Y
b1746 b1746 b1759 b1759 b1770 b1770 b1773 b1773	b1746 mutH b1759 ygeA b1770 lpdA b1773 b1501	Y Y Y Y	Y Y Y
b1746 b1746 b1759 b1759 b1770 b1770 b1773 b1773	b1746 mutH b1759 ygeA b1770 lpdA b1773 b1501 dsbG	Y Y Y Y	Y Y Y Y
b1746 b1746 b1759 b1759 b1770 b1770 b1773 b1773 b1773	b1746 mutH b1759 ygeA b1770 lpdA b1773 b1501 dsbG prsA	Y Y Y Y	Y Y Y Y
b1746 b1746 b1759 b1759 b1770 b1770 b1773 b1773 b1773 b1773	b1746 mutH b1759 ygeA b1770 lpdA b1773 b1501 dsbG prsA rplL	Y Y Y Y	Y Y Y Y Y
b1746 b1746 b1759 b1759 b1770 b1770 b1773 b1773 b1773 b1773 b1773	b1746 mutH b1759 ygeA b1770 lpdA b1773 b1501 dsbG prsA rplL yidS	Y Y Y Y Y Y	Y Y Y Y
b1746 b1746 b1759 b1759 b1770 b1770 b1773 b1773 b1773 b1773	b1746 mutH b1759 ygeA b1770 lpdA b1773 b1501 dsbG prsA rplL	Y Y Y Y	Y Y Y Y Y

b1808	tufA	Υ	
b1808	tufB	Υ	
b1844	b1844	Y	
b1983	b1983	Y	Υ
b1983	cca	Υ	
b1983	nrdA	Υ	
b1983	ydaY		Υ
b2097	b2097	Y	Υ
b2097	dnaJ		Υ
b2097	dnaK		Υ
b2097	mopA	Υ	Υ
b2097	ybbN	Υ	Υ
b2255	b2255	Y	
b2255	b1410	Υ	
b2290	b2290	Υ	Υ
b2290	tufB	Υ	
b2299	b2299		Υ
b2299	panC	Υ	Υ
b2324	b2324	Υ	
b2341	b2341	Y	
b2341	aceF	Υ	
b2341	b2342	Υ	
b2341	b2351		Υ
b2341	lpdA	Υ	
b2341	pepN	Υ	
b2342	b2342	Υ	
b2342	b2341	Υ	
b2342	lpdA	Υ	
b2342	rfaD	Υ	
b2342	rpIS		Υ
b2342	rpIU		Υ
b2342	rpsE		Υ
b2342	ycaO	Υ	
b2383	b2383	Υ	
b2383	aceE	Υ	
b2463	b2463	Y	Υ
b2463	b1409		Υ
b2463	cca	Υ	
b2463	hrpA	Υ	
b2463	nrdA		Υ
b2463	ycjZ		Υ
b2496	b2496	Y	
b2496	dnaN	Υ	
b2496	nfrA		Υ

b2496 b2496 b2496	rho rpIL rpmG		Y Y Y
b2496	rpsB		Ϋ́
b2496	rpsJ		Υ
b2496	yaiP		Υ
b2496	yejE		Υ
b2511	b2511	Υ	
b2511	b1410	Y	
b2511	rpIB	Y	
b2511	rpID	Y Y	
b2511 b2511	rpIN rpsB	Ϋ́	
b2511	rpsC	Y	
b2511	b2520	Y	Y
b2520	rplL	,	Ϋ́
b2520	rpmB		Y
b2520	rpsB		Υ
b2520	rpsE		Υ
b2520	rpsJ		Υ
b2710	b2710	Y	
b2810	b2810	Y	
b2810	gcd		Υ
bglG	bglG	Υ	
bglG	b1690		Y
bglG	folP		Y
bglG	yahB	V	Υ
bioH birA	bioH birA	Y Y	
birA	glpB	7	Y
birA	nfrB		Y
birA	rplU		Ϋ́
birA	rplV		Y
birA	rpsC		Υ
birA	rpsN		Υ
birA	yhcJ	Υ	
bolA	bolA	Y	
bolA	dppB		Υ
bolA	ydhD	Υ	
bolA	yehY	37	Υ
bolA	yleA	Y	
cadA	<i>cadA</i>	Y Y	
cadA cadA	aceE	Y Y	
CauA	cobB	ſ	

cadA	dnaN	Y	
cadA	IdcC	Y	
cadA	lpdA	Y	
cadA	rplC	Y	
cadA	rplE	Ϋ́	
cadA	rpsB	Y	
cadA	rpsb	Y	
cadA	secA	Y	
cadA	tufA	Y	
cadA	ycdQ	Y	
cadA	cafA	Y	Υ
cafA	dnaK	,	Ϋ́
cafA	eno		Ϋ́
cafA	rpe		Ϋ́
cafA	rpsD	Υ	•
cafA	rpsE	•	Υ
cafA	rpsL		Ϋ́
cafA	rpsM		Ϋ́
cafA	rpsivi		Ϋ́
cafA	rpsS		Ϋ́
cafA	rpsT		Ϋ́
cafA	rpsU		Ϋ́
carA	carA	Υ	Y
carA	carB	Ϋ́	Ϋ́
cbpA	cbpA	Y	Y
cbpA	aidB	Ý	•
cbpA	clpA	Ý	
cbpA	clpB	Ý	
cbpA	deaD	Ϋ́	
cbpA	dnaK	Ϋ́	Υ
cbpA	fabZ	Y	-
cbpA	hscA	Y	
cbpA	infC	Υ	
cbpA	metK	Υ	
cbpA	mukB	Υ	
cbpA	prsA	Υ	
cbpA	rbsK		Υ
cbpA	rho	Υ	
cbpA	rplA	Υ	
cbpA	rplB	Y	
cbpA	rplC	Υ	
cbpA	rpID	Y	
cbpA	rplE	Υ	
cbpA	rplF	Υ	

cbpA	rpll	Y	
cbpA	rplL	•	Υ
-	rpIS	Υ	ī
cbpA		Y	
cbpA	rpIV rpIX	Y	
cbpA	rpmC	1	~
cbpA cbpA	rpmG		Y Y
cbpA	rpoC	Υ	'
cbpA	rpsA	Y	
cbpA	rpsB	Ϋ́	Υ
cbpA	rpsC	Ϋ́	•
cbpA	rpsD	Ϋ́	
cbpA	rpsE	Ϋ́	
cbpA	rpsG	Ϋ́	Υ
cbpA	rpsM	Ϋ́	•
cbpA	rpsP	•	Υ
cbpA	srmB	Υ	•
cbpA	tufA	Ϋ́	
cbpA	ugpB	•	Υ
cbpA	yfbT		Ϋ́
cbpA	ygiF	Υ	•
cbpA	yheM	•	Υ
cbpA	yjfK		Y Y
cca	cca	Υ	Υ
cca	pflB	Υ	
cca	rpIC	Υ	
cca	rpsB	Υ	
cca	tufB	Υ	
cheA	cheA		Y
cheW	cheW	Υ	Y
cheY	cheY	Υ	Y
cheY	aceE	Υ	Υ
cheY	aceF	Υ	Υ
cheY	IpdA	Υ	
cheZ	cheZ	Υ	Y
cheZ	gapA		Υ
citB	citB	Y	
clpA	clpA	Y	Υ
clpA	barA		Υ
clpA	dnaK		Υ
clpA	mukB	Υ	
clpA	proV		Υ
clpA	purL		Y
clpA	rho	Υ	Υ

clpA	rhsC		Υ
clpA	rplC		Υ
clpA	rplD		Υ
clpA	rplE	Υ	
clpA	rpll		Υ
clpA	rplJ	Υ	
clpA	rpIL		Υ
clpA	rplM		Υ
clpA	rplR		Υ
clpA	rplU		Υ
clpA	rpmA		Υ
clpA	rpmC		Υ
clpA	rpmG		Υ
clpA	rpoB	Υ	
clpA	rpoC	Υ	
clpA	rpsA	Υ	Υ
clpA	rpsB	Υ	Υ
clpA	rpsD	Υ	
clpA	rpsF		Υ
clpA	rpsG	Υ	Υ
clpA	rpsJ		Υ
clpA	rpsN		Υ
clpA	rpsP		Υ
clpA	rpsU		Υ
clpA	sbcC		Υ
clpA	secA	Υ	
clpA	spoT	Υ	
clpA	tufB	Υ	
clpA	ybjE		Υ
clpA	ygiF		Υ
clpA	yhbY		Υ
clpA	yljA		Υ
clpB	clpB	Y	Y
clpB	b1501		Υ
clpB	ccmB		Υ
clpB	elaC		Υ
clpB	evgS		Υ
clpB	glcB		Υ
clpB	lasT		Υ
clpB	ptsN		Υ
clpB	rplC		Υ
clpB	rplM		Υ
clpB	rplU		Υ
clpB	rplV		Υ

clpB	rnmC		Υ
-	rpmC		I
clpB	rpsB	Υ	
clpB	rpsG		Y
clpB	rpsH		Υ
clpB	rpsJ		Υ
clpB	rpsN		Υ
clpB	sapC		Ϋ́
clpB	thiG		Ϋ́
		V	•
clpB	tufA	Y	
clpB	yagW		Y
clpB	ygaU		Υ
clpB	yhfT		Υ
clpB	ymcC		Υ
clpP	clpP	Y	Y
clpP	aceE	Υ	
clpP	agal	Υ	
clpP	b1685		Υ
clpP	b2073	Υ	·
clpP	clpA	Ϋ́	
clpP	dnaK	Ϋ́	
	dsbG	Y	
clpP			
clpP	greA	Y	
clpP	gyrA	Y	
clpP	hsdR	Y	
clpP	htrA	Υ	
clpP	kefC		Υ
clpP	lon	Υ	
clpP	metK	Υ	
clpP	mopA	Υ	Υ
clpP	mtlA		Υ
clpP	narG	Υ	
clpP	narZ	Υ	
clpP	parE	Υ	
clpP	recA	Υ	
clpP	rfaD	Y	Y
clpP	rho	Y	-
clpP	rplA	•	Υ
clpP	rpIC		· V
•	-		Y Y
clpP	rpID	V	Ĭ
clpP	rplE	Y	
clpP	rplJ	Υ	
clpP	rplK		Y
clpP	rplL		Υ
clpP	rplM	Υ	

I			
clpP	rpIS	Υ	
clpP	rplU		Υ
clpP	rpmB		Υ
clpP	rpmC		Υ
clpP	rpmG		Υ
clpP	rpoC	Υ	
clpP	rpsB		Υ
clpP	rpsF		Υ
clpP	rpsG	Υ	Υ
clpP	rpsJ	Υ	
clpP	rpsP		Υ
clpP	secA	Υ	
clpP	tufA	Y Y	Υ
clpP	tufB	Υ	
clpP	ugpB		Υ
clpP	yfiF		Υ
clpP	yjgD		Υ
clpX	clpX	Υ	Υ
clpX	ahpC		Υ
clpX	clpA	Υ	
clpX	clpB	Υ	
clpX	dnaK		Υ
clpX	grpE		Υ
clpX	metK	Υ	
clpX	mreB	Y Y Y	
clpX	pstB	Υ	
clpX	rfbB		Υ
clpX	rplL		Υ
clpX	rpmC		Υ
clpX	rpmG		Υ
clpX	tufB	Υ	
clpX	yfhE	Υ	
cobB	cobB	Υ	Υ
cobB	hsdR	Υ	
cobB	prsA	Υ	
cobB	rpsC	Υ	
cobB	rpsD	Y	
cobB	rpsN		Υ
cobB	rpsT		Ϋ́
crp	crp	Y	-
crp	rplC	Ϋ́	
crp	rpIL	-	Υ
crp	rpsE	Υ	-
crp	rpsJ	Ϋ́	
1	- 12-30	· ·	

cspA	cspA	Υ	Y
cspA	accA	Υ	
cspA	atpG		Υ
cspA	cca	Υ	
cspA	cspB		Υ
cspA	cspG		Υ
cspA	deaD	Υ	
cspA	dnaN	Υ	
cspA	dsbG	Υ	
cspA	hfq		Υ
cspA	hupA		Υ
cspA	hupB		Υ
cspA	rplA	Υ	
cspA	rplC	Y Y	Υ
cspA	rplD	Υ	
cspA	rplL		Υ
cspA	rplR		Υ
cspA	rpIS		Υ
cspA	rpoA		Υ
cspA	rpsC		Υ
cspA	rpsD	Υ	
cspA	rpsE		Υ
cspA	rpsG		Υ
cspA	rpsJ		Υ
cspA	rpsN		Υ
cspA	ybhF	Υ	
cspA	ycbY	Υ	
cspA	yceC	Υ	
cspA	yciL	Υ	
cspA	ydaY		Υ
cspA	ygcG		Υ
cspB	cspB	Υ	Y
cspB	aceE	Υ	
cspB	aceF	Υ	
cspB	b2998		Υ
cspB	cspA	Υ	Y
cspB	cspE		Y
cspB	cspG		Υ
cspB	dnaN	Υ	
cspB	eno		Υ
cspB	gltA		Υ
cspB	hfq		Υ
cspB	lpdA	Υ	Υ
cspB	rplC		Υ

cspB	rplL		Υ
cspB	rpIV		Υ
cspB	rpsB		Υ
cspB	rpsJ		Y
cspC	cspC	Υ	Ϋ́
cspC	accA	Ϋ́	Ϋ́
cspC	accD	•	Ϋ́
cspC	aidB	Y	Ϋ́
cspC	b1978	ı	Y
-	creC		Y
cspC			Y
cspC	cspD		Ϋ́
cspC	cspE	V	
cspC	deaD	Y	Y
cspC	dnaJ		Y
cspC	entB		Y
cspC	glpK		Y
cspC	hfq		Υ
cspC	hrpA	Y	
cspC	hupA		Υ
cspC	hupB		Υ
cspC	metR		Y
cspC	pnp	Υ	
cspC	prpD		Υ
cspC	putA		Υ
cspC	rcsB		Υ
cspC	recN		Υ
cspC	rfaJ		Υ
cspC	rhIE		Υ
cspC	rho	Υ	Υ
cspC	rplA	Υ	Υ
cspC	rplB	Υ	Υ
cspC	rpIC	Υ	Υ
cspC	rplD		Υ
cspC	rplE	Υ	Y
cspC	rpll		Υ
cspC	rplM	Υ	Y
cspC	rplR		Υ
cspC	rpIS		Υ
cspC	rpIU		Υ
cspC	rpIV		Υ
cspC	rplW		Υ
cspC	rplX		Υ
cspC	rpsA	Υ	Υ
cspC	rpsB	Y	Y
	r	- -	=

cspC	rpsC		Y
cspC	rpsD	Υ	Y
cspC	rpsE	Ý	Ϋ́
cspC	rpsF	-	Y
cspC	rpsG	Υ	Y
cspC	rpsl	-	Y
cspC	rpsJ		Ϋ́
cspC	rpsM	Υ	•
cspC	rpsN	-	Υ
cspC	rpsT		Y
cspC	sun		Ϋ́
cspC	vacB	Υ	-
cspC	ycbY	Y	
cspC	yceC	Ϋ́	
cspC	yciL	Ϋ́	
cspC	yciR		Υ
cspC	ydaY		Ϋ́
cspC	yfiF	Υ	•
cspC	yhbY	•	Υ
cspC	yhiP		Ϋ́
cspC	yhiR	Υ	Ϋ́
cspC	yjdB	'	Ϋ́
cspC	ymfC		Ϋ́
cspD	cspD	V	Y
cspD	accA	Y Y	,
cspD	aceE	Ϋ́	
cspD	accD	'	Υ
cspD	aspA		Y
cspD	b1486		Ϋ́
cspD	cspC	Υ	Ϋ́
cspD	cspE	Ϋ́	•
cspD	ddlA	ı	Y
cspD	deaD	Υ	I
cspD	dppD	ı	Υ
cspD	hlpA		Y
cspD	hupA		Y
cspD	pnp	Υ	I
cspD cspD	rho	Ϋ́	
cspD	rplA	Y	
cspD	rpIB	Y	
cspD cspD	rplC	Ϋ́	Υ
cspD	rplM	•	Ϋ́
cspD	rpiN		Y
cspD cspD	rpiX		Y
cahn	ιριν		Ĭ

İ			
cspD	rpsA	Υ	
cspD	rpsB	Υ	Υ
cspD	rpsC	Υ	
cspD	rpsD	Υ	Υ
cspD	rpsE	Υ	Υ
cspD	rpsG	Υ	Υ
cspD	rpsJ		Υ
cspD	rpsM	Υ	
cspD	rpsN		Υ
cspD	rpsT		Υ
cspD	rpsU		Υ
cspD	tehB	Υ	•
cspD	vacB	Ϋ́	
cspD	ycbY	Ϋ́	
cspD	yceC	Ϋ́	
cspD	yccc	Ϋ́	
cspD	ydiA	•	Υ
cspD	yfiF		Y
	cspE	Y	Y
cspE		,	Ϋ́
cspE	cspl dnaN	Υ	1
cspE		Ϋ́	
cspE	gabT	Ť	V
cspE	glmU	Υ	Y
cspE	hflB	Ť	V
cspE	hfq	Υ	Υ
cspE	hrpA	Ť	V
cspE	hupA		Y
cspE	hyfG	V	Υ
cspE	pnp	Y	
cspE	rho	Y	
cspE	rplA	Y	Y
cspE	rplB	Y	
cspE	rplC	Υ	Y
cspE	rplD		Y
cspE	rpIS		Υ
cspE	rpIX		Υ
cspE	rpsA	Υ	
cspE	rpsD	Υ	
cspE	rpsE		Y
cspE	rpsG	Υ	Y
cspE	rpsJ		Υ
cspE	rpsP		Y
cspE	secA	Υ	
cspE	serS		Y

ı			
cspE	vacB	Υ	
cspE	ycbY	Υ	
cspE	yciL	Υ	
cspE	yjdG	Υ	
cspG	cspG	Υ	Y
cspG	amyA		Υ
cspG	rplA	Υ	
cspG	rplC		Υ
cspG	rplL		Υ
cspG	rplM		Υ
cspG	rplV		Υ
cspG	rpsB		Υ
cspG	rpsE		Υ
cutA	cutA	Υ	Y
cutA	b2073	Υ	
cutA	tufB	Υ	
cysB	cysB	Y	
cysS	cysS	Υ	
dam	dam	Y	
dam	aceE	Ϋ́	
dam	glpD		Υ
dam	lpdA	Υ	Ϋ́
dam	rpIS	•	Ϋ́
dam	rplV		Ϋ́
dam	serS		Ϋ́
dam	yjjl	Υ	·
dcm	dcm	Y	
dcm	aceF	Ϋ́	Υ
dcm	rplA	'	Ϋ́
dcm	rpIC		Ϋ́
dcm	rplM		Ϋ́
dcm	rpIU		Ϋ́
dcm	rplV		Ϋ́
dcm	rpsC		Ϋ́
dcm	rpsE		Ϋ́
dcm	rpsG		Ϋ́
dcm	rpsN		Ϋ́
dcm	rpsT		Ϋ́
ddlA	ddlA	Υ	Ϋ́
ddlA		Ϋ́	1
ddlA	apt b1487	Ī	Υ
	dnaK	Υ	Ī
ddIA		Ĭ	V
ddlA	ftsl	V	Y
ddlA	proA	Y	

1			
ddlA	yafC		Y
ddlA	yieM		Y
ddlA	ујсТ		Y
deaD	deaD	Υ	Y
deaD	accA	Υ	
deaD	aceE	Υ	Υ
deaD	acpP		Y
deaD	b2340		Υ
deaD	bglA		Υ
deaD	cspC		Υ
deaD	dacA		Υ
deaD	dnaJ		Υ
deaD	dppC		Υ
deaD	evgA		Υ
deaD	gsk		Υ
deaD	hrpA	Υ	
deaD	hupA		Y
deaD	hupB		Υ
deaD	rpIA	Υ	Υ
deaD	rpIB	Υ	Υ
deaD	rpIC		Υ
deaD	rpID		Y
deaD	rplF		Υ
deaD	rpll	Υ	Υ
deaD	rplK		Υ
deaD	rplM	Υ	Υ
deaD	rplP		Υ
deaD	rplQ		Υ
deaD	rplR		Υ
deaD	rpIS		Υ
deaD	rpIU		Υ
deaD	rpIV	Υ	Υ
deaD	rplW		Υ
deaD	rplX		Υ
deaD	rpmA		Υ
deaD	rpmB		Υ
deaD	rpsA	Υ	Υ
deaD	rpsB	Υ	Υ
deaD	rpsC	Υ	Y
deaD	rpsD	Υ	Υ
deaD	rpsE	Υ	Y
deaD	rpsF		Υ
deaD	rpsG	Υ	Y
deaD	rpsH		Υ

deaD	rpsl		Υ
deaD	rpsJ		Υ
deaD	rpsK		Υ
deaD	rpsM	Υ	Υ
deaD	rpsN		Υ
deaD	rpsO		Υ
deaD	rpsP		Υ
deaD	rpsR		Υ
deaD	rpsS		Υ
deaD	rpsT		Υ
deaD	rpsU		Υ
deaD	rspU	Υ	
deaD	srmB	Υ	
deaD	sucD		Υ
deaD	uraA		Υ
deaD	vacB	Υ	Y
deaD	ybjW		Υ
deaD	ycbY	Υ	
deaD	yciL	Υ	Υ
deaD	yfiF	Υ	Υ
deaD	ygiF	Υ	
deaD	yhiR		Υ
deaD	yihQ		Υ
deaD	ynhD		Υ
def	def	Υ	Y
def	rpID	Υ	
def	usg		Y
deoC	deoC	Υ	
dfp	dfp	Υ	Υ
dfp	rpIA	Υ	
dfp	rpoA		Y
dnaA	dnaA	Υ	Υ
dnaA	aspS	Υ	
dnaA	cca	Υ	
dnaA	creC		Υ
dnaA	dnaJ	Y	Υ
dnaA	lpdA	Y	
dnaA	nadE	Y	
dnaA	rplB	Υ	Y
dnaA	rplC	V	Υ
dnaA	rplD	Υ	W
dnaA	rplL		Y
dnaA	rplN		Y
dnaA	rplO		Υ

i			
dnaA	rplP		Υ
dnaA	rplV		Υ
dnaA	rplW		Υ
dnaA	rplX		Υ
dnaA	rpmB		Υ
dnaA	rpsB	Υ	Υ
dnaA	rpsC		Υ
dnaA	rpsE		Υ
dnaA	rpsG		Υ
dnaA	rpsl		Υ
dnaA	rpsJ		Υ
dnaA	rpsM		Υ
dnaA	rpsN		Υ
dnaA	rpsT		Υ
dnaA	tufA	Υ	
dnaA	tufB	Y	
dnaA	ynhG	•	Υ
dnaB	dnaB	Υ	Y
dnaB	aceE	Ý	•
dnaB	b2248	Ϋ́	
dnaB	dnaC	Ý	
dnaB	fiml	-	Υ
dnaB	ftsY		Ϋ́
dnaB	fumA		Ϋ́
dnaB	rplA	Υ	•
dnaB	rpID	•	Υ
dnaB	rpIL		Ϋ́
dnaB	rplO		Ϋ́
dnaB	rplW		Ϋ́
dnaB	rpmC		Ϋ́
dnaB	rpsB	Υ	Ϋ́
dnaB	rpsE	•	Ϋ́
dnaB	thrA		Ϋ́
dnaB	tufA	Υ	Ý
dnaB	tufB	Ϋ́	•
dnaE	dnaE	Y	Y
dnaE	aceE	Ý	•
dnaE	asnB	•	Υ
dnaE	dnaK		Ϋ́
dnaE	dnaQ	Υ	Ϋ́
dnaE	dnaX	Ý	Ϋ́
dnaE	holA	Ϋ́	Ϋ́
dnaE	holB	Ϋ́	•
dnaE	holC	Ý	Υ
a.ia_		•	•

dnaE	holD	Y	Υ
dnaE	holE		Υ
dnaE	hrpB		Υ
dnaE	lpdA		Υ
dnaE	nikD		Υ
dnaE	rhsD		Υ
dnaE	rplA		Υ
dnaE	rplC		Υ
dnaE	rpID		Υ
dnaE	rpIL		Υ
dnaE	rplM		Υ
dnaE	rplU		Υ
dnaE	rpsA		Υ
dnaE	rpsB	Υ	Υ
dnaE	rpsP		Υ
dnaE	ssb	Υ	
dnaE	tpiA		Υ
dnaE	tufA	Υ	Υ
dnaE	tufB	Υ	
dnaE	ybdL		Y
dnaE	yhjO		Υ
dnaG	dnaG	Y	Υ
dnaG	nusG		Υ
dnaG	rplA	Υ	
dnaG	rpIL		Υ
dnaG	rpsG		Υ
dnaG	yfiF		Υ
dnaG	ygdH		Υ
dnaJ	dnaJ	Υ	
dnaJ	aceE	Υ	
dnaJ	add	Υ	
dnaJ	atpD	Υ	
dnaJ	clpA	Υ	
dnaJ	deaD	Υ	
dnaJ	dnaK	Y	
dnaJ	gatY	Y	
dnaJ	imp	Y	
dnaJ	infB	Y	
dnaJ	malT	Y	
dnaJ	metK	Y	
dnaJ	mreB	Y	
dnaJ	mukB	Y	
dnaJ	narG	Y	
dnaJ	pstB	Υ	

امما	wa a A	V	
dnaJ	recA	Y	
dnaJ	relE	Y	
dnaJ	rho	Y	
dnaJ	rplA	Y	
dnaJ	rplJ	Y	
dnaJ	rpIS	Y	
dnaJ	rplV	Y	
dnaJ	rplW		Y
dnaJ	rplX	V	Υ
dnaJ	rpoC	Y	
dnaJ	rpsB	Y	
dnaJ	rpsE	Υ	
dnaJ	rpsF		Y
dnaJ	rpsJ	Y	Υ
dnaJ	rpsM	Υ	
dnaJ	rpsN		Υ
dnaJ	srmB	Υ	
dnaJ	trxC	Υ	
dnaJ	tufB	Υ	
dnaJ	yjbJ	Υ	
dnaK	dnaK	Y	Y
dnaK	alaS	Υ	
dnaK	aphA		Υ
dnaK	b1439		Υ
dnaK	b1543		Υ
dnaK	entB		Y
dnaK	fabB		Υ
dnaK	gadA	Υ	
dnaK	gadB		Υ
dnaK	grpE	Υ	Υ
dnaK	hisC		Υ
dnaK	hscA		Υ
dnaK	lon	Υ	
dnaK	narQ		Υ
dnaK	рерВ		Υ
dnaK	proS		Υ
dnaK	rpID		Υ
dnaK	rpIF	Υ	
dnaK	rpll		Υ
dnaK	rplJ	Υ	
dnaK	rpIS	-	Υ
dnaK	rpmG		Ϋ́
dnaK	rpsB	Υ	Y
dnaK	sapA	•	Y
	P. , ,		-

I			
dnaK	tufA	Y	Υ
dnaK	tufB	Y	
dnaK	yhaP		Υ
dnaK	yhcL		Υ
dnaK	yibA	Y	
dnaK	yihM		Υ
dnaQ	dnaQ	Υ	Υ
dnaQ	atpB		Υ
dnaQ	b1685		Υ
dnaQ	cspC		Υ
dnaQ	dnaE	Υ	Υ
dnaQ	dnaX	Υ	Υ
dnaQ	fecB		Υ
dnaQ	holA	Υ	
dnaQ	holB	Υ	Υ
dnaQ	holC	Υ	Υ
dnaQ	holD	Υ	Υ
dnaQ	holE		Υ
dnaQ	lon		Υ
dnaQ	nusG		Υ
dnaQ	rplA	Y	Υ
dnaQ	rplL		Υ
dnaQ	rplM		Υ
dnaQ	rpIS		Υ
dnaQ	rpmC		Υ
dnaQ	rpmG		Υ
dnaQ	slyD		Υ
dnaQ	ssb	Υ	
dnaQ	tufA	Υ	
dnaQ	tufB	Υ	
dnaQ	yfiF		Υ
dnaT	dnaT	Υ	Y
dnaT	yheB		Υ
dnaX	dnaX	Υ	
dnaX	dnaQ		Υ
dnaX	fecB		Υ
dnaX	holE		Υ
dnaX	nusG		Υ
dnaX	rplA	Υ	
dnaX	rpIL		Υ
dnaX	rpmC		Υ
dnaX	rpsF		Υ
dnaX	ugpB		Υ
dnaX	yfiF		Υ
•	•		

dnaX	yjgD		Υ
dps	dps	Y	Y
dps	deaD	Υ	
dps	hfq		Υ
dps	hlpA		Y
dps	hrpA	Υ	-
dps	lon	Ϋ́	
dps	ompG	Ϋ́	
dps	rfaD	Ϋ́	Υ
dps	rplW	•	Ϋ́
dps	rplX		Ϋ́
dps	rpsE		Ϋ́
dps	rpsF		Ϋ́
dps	rpsH		Ϋ́
dps	rpsJ		Ϋ́
-	-		Ϋ́
dps	slyD		Y
dps	yibL	V	Y
dsbA	dsbA	Υ	Ϋ́
dsbA	acnA		
dsbA	agaY		Y
dsbA	b2088		Y
dsbA	dctA		Y
dsbA	dsbC		Y
dsbA	fliA		Y
dsbA	hybC		Y
dsbA	katE		Y
dsbA	mltB		Y
dsbA	tap		Υ
dsbA	ydhU		Υ
dsbA	ymcB		Υ
dsbA	ymfN		Υ
dsbC	dsbC	Υ	
dut	dut	Y	
dxs	dxs	Y	
dxs	aceE	Υ	
dxs	adhE	Υ	
dxs	clpA	Υ	
dxs	lon	Υ	
dxs	mreB	Υ	
dxs	pstB	Υ	
dxs	recA	Υ	
dxs	rho	Υ	
dxs	tufA	Υ	
dxs	tufB	Υ	

dxs	yjgL	Υ	
dxs	yqel	Υ	
eno	eno	Υ	Y
eno	b1374		Υ
eno	dnaK	Υ	Υ
eno	fabZ	Υ	
eno	gltX	Υ	
eno	pnp	Υ	Υ
eno	rne	Υ	Υ
eno	rpsE		Υ
eno	yniC	Υ	
era	era	Υ	
era	lon	Υ	
era	rpsB	Υ	
era	rpsE	Υ	
era	tufA	Υ	
era	tufB	Υ	
exo	exo	Υ	Y
exo	dnaK		Υ
exo	rpsB		Υ
fabA	fabA	Υ	Y
fabA	accA	Υ	
fabA	b1742	Υ	
fabA	deaD	Υ	
fabA	ffh	Υ	
fabA	hupA		Υ
fabA	metK	Υ	
fabA	rplA	Υ	
fabA	rplB	Υ	
fabA	rpIS	Υ	
fabA	rpsA	Υ	
fabA	rpsB	Υ	
fabA	rpsE	Υ	
fabA	rpsG	Υ	
fabA	rpsl	Υ	
fabA	rpsM	Υ	
fabA	rpsR		Υ
fabA	tig	Υ	
fabA	tufA	Υ	
fabA	tufB	Υ	
fabB	fabB	Υ	
fabF	fabF	Υ	Υ
fabF	асрР		Υ
fabF	dnaK	Υ	Υ

fabG	fabG	Y	
fabH	fabH	Y	Y
fabl	fabl	Υ	Υ
fabl	b2506	Y	-
fabl	dnaN	Ϋ́	
fabl	rho	Ϋ́	
fabl	rplJ	Ϋ́	
fabl	rpsA	Ý	
fabl	•	Y	
	rpsJ	Y	
fabl	ycbY	ĭ	V
fabl	yjgD		Y
fabZ	fabZ	Y	
fabZ	clpA	Y	
fabZ	ffh	Υ	
fabZ	glnS	Υ	
fabZ	lon	Y	
fabZ	rplA	Y	
fabZ	rpIB	Y	
fabZ	rplC	Υ	
fabZ	rpID	Υ	
fabZ	rpIE	Υ	
fabZ	rpIF	Υ	
fabZ	rpll	Υ	
fabZ	rplJ	Υ	
fabZ	rplM	Υ	
fabZ	rpIS	Υ	
fabZ	rpIV	Υ	
fabZ	rpsA	Υ	
fabZ	rpsB	Υ	
fabZ	rpsC	Υ	
fabZ	rpsD	Υ	
fabZ	rpsE	Υ	
fabZ	rpsG	Υ	
fabZ	tufA	Υ	
fabZ	tufB	Υ	
fadR	fadR	Y	Y
fba	fba	Y	Y
fba	dnaK	Υ	Υ
fba	fldA	Υ	
fba	hsdR	Υ	
fba	rpsB	Υ	
fba	yfeR	Υ	
fdhD	fdhD	Y	Y
fdhD	aceE	Υ	

fdhD fdhD fdhD fdhD fdhD fdnG fdnG	aceF iscS lpdA lysU rpsB fdnG aceE	Y Y Y Y Y	Υ
fdnG	fdhE	Y	Y
fdnG	tufA	Y	Υ
fdnG <i>ffh</i>	tufB <i>ffh</i>	Y Y	
ffh	adhE	Ϋ́	
ffh	clpA	Ϋ́	
ffh	lon	Ý	
ffh	metK	Ϋ́	
ffh	rplA	Υ	
ffh	rpID	Υ	
ffh	rplJ	Υ	
ffh	rpsB	Y	
ffh	rpsD	Y	
ffh	rpsE	Y	
ffh ffh	rpsK tufA	Y Y	
ffh	tufB	Y	
fimB	fimB	Y	Y
fimB	aceF	Ý	•
fimB	rplB	Υ	
fimB	rpIC		Υ
fimB	rplM		Υ
fimB	rpIS		Υ
fimB	rplU		Y
fimB	rplV		Y
fimB fimB	rpIX		Y Y
fimB	rpmB rpsC	Y	Y
fimB	rpsD	Ϋ́	•
fimB	rpsE	-	Υ
fimB	rpsF		Υ
fimB	rpsG	Υ	Υ
fimB	rpsH		Υ
fimB	rpsN		Y
fimB	rpsP		Y
fimB	rpsR		Y Y
fimB	rpsT		Υ

fis	fis dnaK hupA rpIC rpID rpIM rpIS rpIT rpIV rpIW rpmG rpoC rpsB rpsE rpsJ rpsM thiG fkIB	Y	Y Y Y Y Y Y Y Y Y Y
fkIB	entB		Y
fkIB	glpF		Y
fkIB	proY		Y
fkIB	trkH	Υ	Y Y
<i>fkpA</i> fkpA	<i>fkpA</i> b2146	ĭ	Ϋ́
fkpA	carA		Y
fkpA	entE		Ϋ́
fkpA	ydeV		Ϋ́
fldA	fldA	Υ	•
fldA	aidB	Ý	
fldA	clpA	Ϋ́	
fldA	dnaN	Y	
fldA	gcpE	Υ	
fldA	lon	Υ	
fldA	ubiX	Υ	
fliY	fliY	Y	Y
fliY	mopA		Υ
fliY	yaiL		Υ
fliY	ybhK		Y Y
fliY	ygiF		
fliY	yqjD		Υ
fnr	fnr	Y	
folA	folA	Υ	Y
folA	mdoB		Υ
folA	proS		Υ

folA folA	rplM ybhC		Y Y
folA	ycbN		Υ
folA	yieO		Υ
folC	folC	Y	Y
folC	b1588		Υ
frdA	frdA	Y	Y
frdA	b1447		Υ
frdA	yi81_3		Υ
frr	frr	Y	
fruR	fruR	Y	
fruR	rfaD	Υ	
fruR	rpsC	Υ	
fruR	tufA	Υ	
fruR	tufB	Υ	
ftsA	ftsA	Y	
ftsA	accD		Υ
ftsA	aceE	Υ	
ftsA	cutA		Υ
ftsA	dnaJ	Υ	
ftsA	dnaK	Υ	
ftsA	mreB	Υ	
ftsA	nusG		Υ
ftsA	rho		Υ
ftsA	rplK		Υ
ftsA	rplL		Υ
ftsA	rplT		Υ
ftsA	rpIU		Υ
ftsA	rpsB		Υ
ftsA	slyD		Y
ftsA	tdcE		Υ
ftsA	tufA	Y	Y
ftsA	tufB	Υ	
ftsA	yfiF		Y
ftsA	yhhA		Y
ftsE	ftsE	Y	
ftsE	b1368		Y
ftsE	b2639		Y Y
ftsE	cdd		Y
ftsE	ibpA	V	Y
ftsE	mopA	Y Y	Y
ftsE	rfaD	Ť	V
ftsE	rpIB rpIK		Y Y
ftsE	rplK		ľ

ftsE	rplP		Υ
ftsE	rplR		Υ
ftsE	rpIU		Ϋ́
ftsE	rpIV		Ϋ́
ftsE	rpmC		Ϋ́
ftsE	rpsB	Υ	'
ftsE	•	ı	Υ
ftsE	rpsE rpsG		Ϋ́
	rpsG		Ϋ́
ftsE	rpsl		Ϋ́
ftsE	rpsJ		Ϋ́
ftsE	rpsP		Ϋ́
ftsE	sbcC	V	Ť
ftsE	tufA	Y	
ftsE	tufB	Υ	V
ftsE	ushA	\ <u>/</u>	Υ
ftsJ	ftsJ	Y	
ftsJ	aceF	Y	
ftsJ	clpA	Y	
ftsJ	ffh	Y	
ftsJ	lon	Y	
ftsJ	rplA	Y	
ftsJ	rplC	Y	
ftsJ	rpID	Y	
ftsJ	rplJ	Y	
ftsJ	rpIV	Y	
ftsJ	rpsB	Y	
ftsJ	rpsD	Y	
ftsJ	rpsE	Y	
ftsJ	rpsG	Y	
ftsJ	tufA	Y	
ftsJ	tufB	Y	
ftsK	ftsK _		Y
ftsK	aceE	Y	
ftsZ	ftsZ	Y	
ftsZ	add	Y	
ftsZ	dnaK	Y	
ftsZ	ftsK	Y	
ftsZ	fusA	Y	
ftsZ	gatY	Y	
ftsZ	malK	Y	
ftsZ	mreB	Y	
ftsZ	pstB	Y	
ftsZ	recA	Y	
ftsZ	rplE	Υ	

ftsZ	rpsB	Υ	
ftsZ	rpsE	Ϋ́	
ftsZ	rpsJ	Ϋ́	
ftsZ	secA	Ϋ́	
ftsZ	tufA	Ϋ́	
ftsZ	tufB	Ϋ́	
fucU	fucU	ı	Y
fucU	lpdA		Y
fucU	rplL		Y
fucU	rpIM		Y
fucU	rplV		Y
fusA	fusA	Y	Y
fusA	apt	Ϋ́	I
fusA	apt aroB	Ϋ́	
fusA	b1707	1	Υ
fusA	b1707		Y
fusA	citE		Y
fusA	htrE		Y
fusA	nadE	Υ	ı
fusA	rffH	ı	Υ
fusA	tufA	Υ	ı
fusA	tufB	Ϋ́	
fusA	yfcC	ı	Υ
gadA	gadA	Y	•
gadA	dnaK	Ϋ́	Υ
gadA	gadB	Ý	Ϋ́
gadA	ssb	•	Ϋ́
gad/K gadB	gadB	Y	•
gadB	dnaK	Ý	
gadB	gadA	Ϋ́	
gadB	yadG	•	Υ
gapA	gapA	Υ	Ϋ́
gapA	cpxR	•	Ϋ́
gapA	dnaK	Υ	Ϋ́
gapA	gatZ	Ϋ́	•
gapA	rpoA	-	Υ
gapA	rpoB		Y
gapA	rpoC		Υ
gapA	rpoD		Υ
gatB	gatB	Υ	
gatB	b1410	Y	
gatB	cca	Υ	
gatB	tufA	Υ	
gatY	gatY	Y	Y
-	_		

gatY	dnaK	Υ	Υ
			ı
gatY	malT	Y	
gatY	truA	Y	
gatY	tufA		Υ
gatY	tufB	Υ	
gatZ	gatZ	Y	Υ
gcpE	gcpE	Y	
gcpE	ftsZ	Y	
gcpE	rho	Ϋ́	
		Ý	V
gidA	gidA		Υ
gidA	thdF	Y	
gidB	gidB	Υ	Υ
gidB	aceE	Υ	Υ
gidB	aceF	Υ	
gidB	acrB		Υ
gidB	infB		Υ
gidB	IpdA		Υ
gidB	nadC		Ϋ́
gidB	rplM		Ϋ́
	-		
gidB	rpsB		Y
glmS	glmS	Y	Υ
glmS	alaS		Υ
glmS	dnaJ		Υ
glmS	eutH		Υ
glmS	rplV		Υ
glmS	trkH		Υ
glmS	ynbD	Υ	
glmS	yohD	•	Υ
glmU	glmU	Y	'
	-		
glmU	rpsB	Y	
glnB	glnB	Y	
glnB	rpoB	Υ	
glnB	tig	Y	
glnB	yjgD		Υ
glnS	glnS	Y	
gloA	gloA	Y	Υ
gloA	creC		Υ
gloA	dnaK	Y	Υ
gloA	mopA	•	Ϋ́
gloB	gloB	Υ	
		Y	
gltD	gltD		
gltD	gltB	Y	
gltX	gltX	Y	
gltX	b1555	Υ	

glyQ	glyQ	Υ	Υ
glyQ	dnaJ	Ý	Ϋ́
		Ϋ́	'
glyQ	prc	ı	Υ
glyQ	rpID		
glyQ	rpIF		Y
glyQ	rplM		Y
glyQ	rpIP		Y
glyQ	rpIS		Y
glyQ	rplU		Y
glyQ	rpIV		Υ
glyQ	rpmB		Υ
glyQ	rpsB		Υ
glyQ	rpsC	Υ	
glyQ	rpsE		Υ
glyQ	rpsM		Υ
glyQ	rpsP		Υ
glyQ	tufB	Υ	
glyS	glyS	Y	
gmk	gmk	Υ	
greA	greA	Υ	
greA	aceE	Υ	
greA	adhE	Υ	
greA	aptD	Υ	
greA	carB	Υ	
greA	clpX	Υ	
greA	dnaK	Υ	
greA	gapA	Υ	
greA	gatZ	Υ	
greA	guaB	Υ	
greA	metE	Υ	
greA	pstl	Υ	
greA	rho	Υ	
greA	rplB	Υ	
greA	rplD	Υ	
greA	rpIE	Υ	
greA	rplJ	Υ	
greA	rplM	Υ	
greA	rpoA	Υ	
greA	rpoB	Υ	
greA	rpoC	Υ	
greA	rpsB	Υ	
greA	rpsE	Υ	
greA	rpsG	Υ	
greA	rpsJ	Υ	

greA secA Y greA thrS Y greA tufB Y greA yeaG Y greB yeaB Y greB greB Y greB rpoC Y greB rpoC Y greB rpsE Y greB rpsE Y greB rpsE Y grpE dnaK Y grpE dsdX Y grpE entB Y grpE entB Y grpE gadA Y grpE gadA Y grpE gadA Y grpE rplD Y grpE rplD Y grpE rplC Y grpE rpsB Y grpE rplC Y grpE rpsB Y grpE gadA Y grpE gadA Y grpE grpE y grpE rplC Y grpE rplC Y grpE grpE y grpE yy Y grpE grpE Y grpE grpE Y grpE entB Y grpE grpE y grpE yy grpE grpE y grpE grpE y grpE grpE y grpE yy grpE grpE y grpE yy grpE rplD Y grpE rplC Y grpE rplC Y grpE rpsB Y grpE rpsG Y grpE rpsG Y grpE rpsG Y grpE rpsG Y grpE rpsC Y grpE yy grpE thiG Y grpE yy grpA ace grpA y gy				
greA tufB Y greA yeaG Y greA yfiF Y greB greB Y greB rplC Y greB rpoA Y greB rpoC Y greB rpsB Y greB rpsE Y greB rpsE Y greE grpE Y grpE grpE Y grpE dsdX grpE emrY grpE entB Y grpE etaB Y grpE gadA Y grpE gadA Y grpE grpE rplD Y grpE rplD Y grpE rplD Y grpE rplS Y grpE rpsB Y grpE rpsB Y grpE gadA Y grpE gadA Y grpE gadA Y grpE grpE y grpE rplD Y grpE rplD Y grpE rplD Y grpE rplS Y grpE rpsB Y grpE rpsB Y grpE rpsG Y grpE rpsG Y grpE gadA Y grpE grpE Y grpE rpsG Y grpE rpsG Y grpE rpsG Y grpE rpsG Y grpE thiG Y grpE ydbA_2 Y gyrA gyrA Y gyrA dnaX Y gyrA dnaX Y gyrA ffh Y	areA	secA	Υ	
greA tufB Y greA yeaG Y greB yeaB Y greB rplC Y greB rpoA Y greB rpoC Y greB rpoC Y greB rpsB Y greB rpsE Y grpE grpE y grpE dadX Y grpE emrY Y grpE entB Y grpE gadA Y grpE gadA Y grpE gadA Y grpE rplD Y grpE rplD Y grpE rplC Y grpE rpsB Y grpE rpsB Y grpE rpsB Y grpE grpE y grpE ydA Y grpE gadA Y grpE rplD Y grpE rplC Y grpE rplC Y grpE rplC Y grpE rpsB Y grpE rpsG Y grpE rpsG Y grpE rpsG Y grpE thiG Y grpE ydbA_2 Y grpA gyrA Y gyrA aceF Y gyrA add Y gyrA dnaX Y gyrA dnaX Y gyrA ffh Y				
greA yfiF greB greB greB rplC y greB rpoA y greB rpoA y greB rpoB y greB rpoC y greB rpoE y greB rpsE y grpE daaK y grpE daaK y grpE emrY grpE entB grpE evgA grpE gadA y grpE gadB y grpE rplD grpE rplD grpE rplL grpE rpsB y grpE rpsB y grpE rpsB y grpE rpsB y grpE ydbA_2 guaC guaC gyrA gyrA gyrA dnaJ gyrA dnaK y gryB greB daaK y gryA gryA gryA gryA gryA gryA gryA gr				
greA yfiF greB greB yr greB rplC Y greB rpoA Y greB rpoB Y greB rpoC Y greB rpsE y greB rpsE y grpE grpE y y grpE daaK Y grpE daaK Y grpE emrY grpE entB y grpE ettB y grpE gadA Y grpE gadA Y grpE gadA Y grpE grpE y grpE rplD y grpE rplD y grpE rplC y grpE rpsB y grpE rpsB y grpE rpsB y grpE ydbA_2 y graA gyrA y gyrA dnaA y gyrA dnaA y gyrA ffh y				
greB greB y greB rplC y greB rpoA y greB rpoB Y greB rpoC y greB rpsE y greB rpsE y greB rpsE y grpE grpE y y grpE daaK y grpE daaK y grpE emrY y grpE entB y grpE ettB y grpE gadA y grpE gadA y grpE gadB y grpE rplD y grpE rplD y grpE rplS y grpE rpsG y grpE rpsG y grpE rpsG y grpE ttfA y grpE ydbA_2 guaC guaC guaC guaC guaC gyrA dnaX y gyrA dnaX y gyrA dnaX y gyrA ffh y		-		
greB rplC Y greB rpoA Y greB rpoB Y greB rpoC Y greB rpsE Y greB rpsE Y greB rpsE Y grpE grpE Y Y grpE b2392 Y grpE daaK Y grpE daaK Y grpE emrY Y grpE entB Y grpE entB Y grpE gadA Y grpE gadA Y grpE gadB Y grpE rplD Y grpE rplD Y grpE rplS Y grpE rpsG Y grpE rpsG Y grpE rpsC Y grpE tufA Y grpE ydbA_2 guaC guaC Y gyrA gyrA dadd Y gyrA dnaK Y gyrA dnaK Y gyrA ffh Y		=		
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greB rpoB Y greB rpoC Y greB rpsB Y greB rpsE Y greB rpsE Y grpE grpE Y grpE b2392 Y grpE ccmB Y grpE dadK Y grpE dsdX Y grpE emrY Y grpE entB Y grpE evgA Y grpE gadB Y grpE gadB Y grpE rplD Y grpE rplL Y grpE rplS Y grpE rpsB Y grpE rpsB Y grpE rpsG Y grpE rpsG Y grpE tufA Y grpE ydbA_2 Y gyrA gyrA dnaJ Y gyrA dnaJ Y gyrA dnaJ Y gyrA ffh Y				
greB rpoC y greB rpsB Y greB rpsE Y grpE grpE Y grpE b2392 grpE ccmB Y grpE dnaK Y grpE dsdX Y grpE emrY Y grpE entB Y grpE etgA Y grpE gadA Y grpE gadA Y grpE gadB Y grpE rplD Y grpE rplD Y grpE rplC Y grpE rplS Y grpE rpsB Y grpE rpsG Y grpE rpsG Y grpE thiG Y grpE ydbA_2 Y gyrA gyrA dnaJ Y gyrA dnaJ Y gyrA ffh Y				
greB rpsB y greB rpsE y grpE grpE y y grpE b2392 y grpE ccmB y grpE dnaK y grpE dsdX y grpE emrY y grpE entB y grpE evgA y grpE gadA y grpE gadB y grpE rplD y grpE rplD y grpE rplD y grpE rplS y grpE rpsB y grpE rpsB y grpE rpsG y grpE rpsG y grpE thiG y grpE tufA y grpE ydbA_2 y gyrA gyrA dnaJ y gyrA dnaJ y gyrA dnaK y gyrA ffh y				
greB rpsE y y y grpE grpE b2392 y y grpE ccmB y y grpE dnaK y y grpE dsdX y grpE emrY y grpE entB y grpE evgA y grpE gadA y grpE gadA y grpE gadB grpE metK y grpE rplD y grpE rplD y grpE rplD y grpE rplS y grpE rpsB y grpE rpsG y grpE rpsG y grpE rpsG y grpE thiG y grpE tufA y grpE ydbA_2 guaC guaC y y gyrA aceF y gyrA add y gyrA dnaJ y gyrA dnaJ y gyrA dnaK y gyrA ffh y				
grpE grpE y y y y grpE b2392 y y grpE ccmB y y grpE dnaK y y grpE dsdX y grpE emrY y y grpE entB y y grpE evgA y grpE gadA y grpE gadA y grpE gadB y grpE rplD y grpE rplD y grpE rplL y y grpE rplS y grpE rpsS y grpE rpsS y grpE rpsS y grpE rpsG y grpE rpsG y grpE rpsV y grpE thiG y grpE thiG y grpE thiG grpE tufA y grpE ydbA_2 guaC guaC y y gyrA gyrA add y gyrA dnaK y gyrA dnaK y gyrA dnaK y gyrA ffh y		-		
grpE b2392 grpE ccmB grpE dnaK grpE dsdX grpE emrY grpE entB grpE entB grpE evgA grpE gadA y grpE gadB grpE metK grpE rplD grpE rplL grpE rplS grpE rpsS grpE rpsS grpE rpsS grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA dnaK gyrA dnaK gyrA ffh Y Y Y Y Y Y Y Y Y Y Y Y Y	greB	rpsE	Υ	
grpE ccmB grpE dnaK grpE dsdX grpE emrY grpE emrY grpE entB grpE evgA grpE evgA grpE gadA grpE gadB grpE rplD grpE rplD grpE rplS grpE rplS grpE rpsB grpE rpsG grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA gyrA aceF gyrA dnaJ gyrA dnaK gyrA ffh Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	grpE	grpE	Y	Y
grpE dnaK Y grpE dsdX grpE emrY grpE entB grpE evgA grpE evgA grpE gadA Y grpE gadB grpE metK Y grpE rplD grpE rplL grpE rplS grpE rpsB grpE rpsG grpE rpsG grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA gyrA aceF gyrA add gyrA dnaJ gyrA dnaJ gyrA ffh Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y		b2392		Υ
grpE dsdX grpE emrY grpE emrY grpE entB grpE evgA grpE evgA grpE gadA Y grpE gadB grpE rplD grpE rplD grpE rplL grpE rplS grpE rpsS grpE rpsS grpE rpsS grpE rpsS grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA gyrA aceF gyrA add gyrA dnaJ gyrA dnaJ gyrA ffh Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	grpE	ccmB		Υ
grpE dsdX grpE emrY grpE entB grpE evgA grpE evgA grpE flgK grpE gadA Y grpE gadB grpE metK Y grpE rpID grpE rpID grpE rpIS grpE rpSB grpE rpsB grpE rpsG grpE rpsV grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA gyrA aceF gyrA add y gyrA dnaJ gyrA dnaJ gyrA ffh Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y		dnaK	Υ	Υ
grpE emrY grpE entB grpE evgA grpE evgA grpE flgK grpE gadA Y grpE gadB grpE rplD grpE rplD grpE rplL grpE rplS grpE rpsS grpE rpsG grpE rpsG grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA gyrA add gyrA dnaJ gyrA dnaK gyrA ffh Y Y Y Y Y Y Y Y Y Y Y Y Y		dsdX		Υ
grpE entB grpE evgA grpE flgK grpE gadA y grpE gadB grpE metK grpE rplD grpE rplL grpE rplS grpE rpsS grpE rpsS grpE rpsG grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA gyrA add gyrA dnaJ gyrA dnaK gyrA ffh y y y grpE rysG grpC y gyrA dnaK gyrA ffh y				Υ
grpE evgA Y grpE flgK grpE gadA Y grpE gadB Y grpE metK Y grpE rplD Y grpE rplL Y grpE rplS Y grpE rpsB Y grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y gyrA gyrA Y gyrA add Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y				
grpE flgK grpE gadA grpE gadB grpE metK grpE rplD grpE rplD grpE rplS grpE rpsS grpE rpsB grpE rpsG grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA gyrA aceF gyrA add gyrA dnaJ gyrA dnaJ gyrA ffh Y grpE gadB Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y				
grpE gadB Y grpE gadB Y grpE metK Y grpE rpID Y grpE rpIL Y grpE rpIS Y grpE rpsB Y grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y grpE ydbA_2 Y grpA gyrA Y gyrA aceF Y gyrA add Y gyrA dnaJ Y gyrA dnaJ Y gyrA ffh Y				
grpE gadB grpE metK grpE rpID grpE rpIL grpE rpIS grpE rpSB grpE rpsB grpE rpsG grpE rpsV grpE thiG grpE tufA grpE ydbA_2 guaC guaC gyrA gyrA gyrA aceF gyrA add gyrA clpA gyrA dnaJ gyrA dnaK gyrA ffh Y Y Y Y Y Y Y Y Y Y Y Y Y			Υ	
grpE rpID Y grpE rpIL Y grpE rpIS Y grpE rpmG Y grpE rpsB Y grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y grpE ydbA_2 Y grpA gyrA Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA ffh Y			-	Υ
grpE rpID Y grpE rpIL Y grpE rpIS Y grpE rpmG Y grpE rpsB Y grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y guaC guaC Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y			Υ	•
grpE rplS Y grpE rplS Y grpE rpmG Y grpE rpsB Y grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y grpE ydbA_2 Y grpA gyrA Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y			•	Υ
grpE rplS Y grpE rpmG Y grpE rpsB Y grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y guaC guaC Y Y gyrA gyrA Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA ffh Y				
grpE rpmG Y grpE rpsB Y grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y guaC guaC Y Y gyrA gyrA Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA ffh Y				
grpE rpsB Y grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y guaC guaC Y Y gyrA gyrA Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA ffh Y				
grpE rpsG Y grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y guaC guaC Y Y gyrA gyrA Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y		-		
grpE rpsV Y grpE thiG Y grpE tufA Y grpE ydbA_2 Y guaC guaC Y Y gyrA gyrA Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y		•		
grpE thiG Y grpE tufA Y grpE ydbA_2 Y guaC guaC Y Y gyrA gyrA Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y				
grpE tufA Y grpE ydbA_2 Y guaC guaC Y Y gyrA gyrA Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y				
grpE ydbA_2 Y guaC guaC Y Y gyrA gyrA Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y				
guaCguaCYYgyrAgyrAYYgyrAaceFYgyrAaddYgyrAclpAYgyrAdnaJYgyrAdnaKYgyrAffhY				
gyrA gyrA Y Y gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y		_		
gyrA aceF Y gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y		-		
gyrA add Y gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y				Y
gyrA clpA Y gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y				
gyrA dnaJ Y gyrA dnaK Y gyrA ffh Y				
gyrA dnaK Y gyrA ffh Y		-		
gyrA dnaK Y gyrA ffh Y	gyrA	dnaJ		
gyrA ffh Y	gyrA			
	gyrA	ffh	Υ	
gyrA ftsZ Y	gyrA	ftsZ	Υ	
gyrA glyS Y	gyrA	glyS	Υ	

İ			
gyrA	gyrB	Y	Υ
gyrA	hydH	Y	
gyrA	IpdA	Y	
gyrA	malK	Y	
gyrA	metK	Υ	
gyrA	mreB	Υ	
gyrA	pstB	Y	
gyrA	recA	Y	
gyrA	rho	Υ	
gyrA	rpsB	Y	
gyrA	tufA	Υ	Υ
gyrA	tufB	Υ	
gyrA	ycgC	Υ	
gyrA	yjfZ	Υ	
gyrB	gyrB	Y	Y
gyrB	bglJ	Υ	
gyrB	deaD	Υ	
gyrB	glyS	Υ	
gyrB	gyrA	Υ	Y
gyrB	metK	Υ	
gyrB	parE		Υ
gyrB	prsA	Υ	
gyrB	rplJ	Υ	
gyrB	rplW		Υ
gyrB	rpsB	Υ	
gyrB	tufA	Υ	Υ
gyrB	tufB	Υ	
gyrB	yacG	Υ	Υ
helD	helD	Y	
helD	accB		Υ
helD	atpD		Υ
helD	csrA		Υ
helD	dnaK		Υ
helD	gadB		Υ
helD	gapA		Υ
helD	gcvP		Υ
helD	infB		Υ
helD	lipA		Υ
helD	mopA		Υ
helD	pflB		Υ
helD	pgk		Υ
helD	ptsl		Υ
helD	rplC		Υ
helD	rpID		Υ

helD	rplL		Y
helD	rplU		Ϋ́
helD	rpsA		Υ
helD	rpsG		Υ
helD	tpiA		Υ
helD	tufA	Υ	Υ
helD	tufB	Υ	
hepA	hepA	Y	Y
hepA	b2228	Υ	
hepA	fhiA		Y
hepA	nusA	Y	
hepA	rplC	Υ	
hepA	rpoA	Υ	
hepA	rpoB	Y	
hepA	rpoC	Y	Υ
hepA	rpsE	Υ	
hepA	yfcU		Y
hflB	hflB	Υ	Y
hflB	hflC		Y
hflB	nfi		Y
hflB	rpID		Y Y
hflB hflB	rpIL		Ϋ́
hflB	rpIU rpsG		Ϋ́
hfq	hfq	Y	ī
hfq	aidB	Ϋ́	
hfq	cafA	Ý	
hfq	cspC	•	Υ
hfq	csrA		Ϋ́
hfq	deaD	Y	•
hfq	dnaJ	Y	
hfq	dnaK	Υ	
hfq	eno	Υ	
hfq	hlpA	Υ	Υ
hfq	hrpA	Υ	
hfq	hupA	Υ	Υ
hfq	hupB		Υ
hfq	lon	Υ	
hfq	lpxD	Υ	
hfq	nfi		Y
hfq	pnp	Y	
hfq	rho	Y	Υ
hfq	rne	Y	
hfq	rplA	Υ	

hfq	rplB	Υ	
hfq	rpIC	Υ	Υ
hfq	rpID	Ϋ́	Y
hfq	rpll	•	Ϋ́
hfq	rplL		Ϋ́
hfq	rplM		Ϋ́
hfq	rplO		Ϋ́
•	•		Ϋ́
hfq	rpIS		
hfq	rplT		Y
hfq	rpIU		Y
hfq	rplV		Y
hfq	rplX		Y
hfq	rpmB	3.7	Y
hfq	rpoA	Y	
hfq	rpoB	Y	
hfq	rpoC	Y	
hfq	rpsA	Υ	Υ
hfq	rpsB	Υ	Υ
hfq	rpsC	Υ	
hfq	rpsD	Υ	
hfq	rpsE		Υ
hfq	rpsF		Υ
hfq	rpsG		Y
hfq	rpsM	Υ	Y
hfq	rpsP		Υ
hfq	rpsR		Υ
hfq	secA	Υ	
hfq	selB	Υ	
hfq	spoT	Υ	
hfq	tgt	Υ	
hfq	tig	Υ	
hfq	tufA	Υ	
hfq	tufB	Υ	
hfq	vacB	Υ	Y
hfq	ycbY	Υ	
hfq	yciL	Υ	
hfq	yfiF	Υ	
hfq	ygiF	Υ	
himA	himA	Υ	
himA	himD	Υ	
himA	rplA	Y	
himA	rplC	Y	
himA	rplE	Ϋ́	
himA	rpsB	Ϋ́	
	0 -	•	

hlpA rpIV Y hlpA rpIX Y hlpA rpSM Y hlpA spoT Y hlpA vacB Y hns hns Y hns hns Y hns hns Y hns b1410 Y hns b1410 Y hns b1625 Y hns hupA Y hns hupA Y hns malP Y	1			
hIpA rpsM Y hIpA spoT Y hIpA vacB Y hns hns Y hns hns Y hns b1410 Y hns b1625 Y hns hupA Y hns malP Y	-	-		
hIpA spoT Y hIpA vacB Y hns hns Y hns bns Y hns b1410 Y hns b1410 Y hns b1625 Y hns hupA Y hns hupA Y hns malP Y		-		
hIPA vacB Y hns hns Y hns b1s Y hns b1410 Y hns b1625 Y hns hupA Y hns hupA Y hns hupA Y hns malP Y hns rpoB Y hns rpoB Y hns rpoB Y hns secA Y <	-	-		
hns hns y y hns aceE Y hns b1410 Y hns b1625 Y hns hupA Y hns malP Y hns malQ Y hns rpIV Y hns rpOB Y hns rpoB Y hns rpsB Y Y hns secA Y hns ssb Y hns stpA Y hns tig Y holA holA Y holA accB Y holA dacCC Y holA b0878 Y holA dnaK Y holA dnaK Y holA grpE Y holA infB Y holA rpIL Y holA rpIC Y holA rpSC Y holB dnaE Y holB dnaE Y holB dnaE Y holB dnaE Y holB dnaE Y holB y holB dnaE	-	-		
hns b1410 Y hns b1410 Y hns b1625 Y hns hupA Y hns hupA Y hns hupA Y hns malQ Y hns rpIV Y hns rpIX Y	hlpA	vacB		
hns b1410 Y hns b1625 Y hns hupA Y hns hupA Y hns malP Y hns malQ Y hns rpIV Y hns rpOC Y hns rpoB Y hns rpoC Y hns rpoB Y hns rpoC Y hns rpoC Y hns rpoC Y hns secA Y hns stpA Y	hns	hns		Y
hns b1625 hns hupA hns malP hns malP hns malQ hns rpIV hns rpIV hns rpSB Y hns rpoC Y hns rpsB Y hns secA Y hns ssb Y hns stpA hns stpA hns tig holA holA Y holA accB HolA accC Y holA b0878 holA b1428 holA dnaK holA fdnH holA gcvP holA infB holA ppx holA rpIL holA rpIC holA rpSC holB holB Y holB aceE Y holB cysJ holB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y	hns			
hns hupA hns malP hns malQ hns rplV hns rplV hns rplX hns rpoB hns rpoC hns rpsB hns rpsB hns secA hns secA hns ssb hns stpA hns tig holA holA holA holA holA holA holA holA holA	hns	b1410	Υ	
hns malP Y hns malQ Y hns rplV Y hns rplX Y hns rpoB Y hns rpoC Y hns rpsB Y Y hns secA Y hns secA Y hns stpA Y hns tig Y holA holA Y holA accB Y holA bo878 holA b1428 holA dnaK Y holA fdnH Y holA gcvP Y holA grpE Y holA infB Y holA ppx holA rplL Y holA rplC Y holA rplC Y holA rplC Y holA rplC Y holA rplC Y holA rplC Y holA rplC Y holA rplC Y holA rplC Y holA rplC Y holA rpsC Y holB holB Y holB dnaE Y	hns	b1625		
hns malQ Y hns rplV Y hns rplX Y hns rpoB Y hns rpoC Y hns rpsB Y Y hns secA Y hns ssb Y hns stpA Y hns tig Y holA holA Y holA accB Y holA bol878 Y holA dnaK Y holA fdnH Y holA gcvP Y holA infB Y holA ppx Y holA rplL Y holA rplC Y holA rpsC Y holB holB Y holB dnaE Y holB dnaE Y	hns	hupA		Υ
hns rpIV hns rpOB hns rpOB hns rpOC hns rpSB Y hns secA hns secA hns ssb Y hns stpA hns tig holA holA holA y holA accB holA accC holA b0878 holA b1428 holA dnaK holA fdnH holA gcvP holA grpE holA infB holA ppx holA rpIL holA rpIP holA rpIP holA rpSC holA yehB holB holB holB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y	hns	malP		
hns rpoB Y hns rpoC Y hns rpsB Y hns secA Y hns secA Y hns stpA Y hns tig Y holA holA Y holA accB holA accC Y holA b0878 holA dnaK holA fdnH holA gcvP holA grpE holA infB Y holA pepB holA ppx holA rplC holA rplC holA rpsC holB holB Y holB dnaE Y holB dnaE	hns	malQ	Υ	
hns rpoB Y hns rpoC Y hns rpsB Y Y hns secA Y hns ssb Y hns stpA Y hns tig Y holA holA Y holA accB holA accC holA b0878 holA b1428 holA dnaK holA fdnH holA gcvP holA grpE holA infB holA pepB holA ppx holA rplL holA rplC holA rpsC holA yehB holB holB Y holB dnaE Y	hns	rpIV		Υ
hns rpoC Y hns rpsB Y Y hns secA Y hns ssb Y hns stpA Y hns tig Y holA holA Y holA accB Y holA accC Y holA b0878 Y holA dnaK holA dnaK holA gcvP holA grpE holA infB holA ppx holA rplL holA rplP holA rpsC holB dnaE Y holB dnaE Y holB dnaE	hns	rplX		Υ
hns rpsB Y Y hns secA Y hns ssb Y hns stpA Y hns tig Y holA holA Y holA accB Y holA accC Y holA b0878 Y holA dnaK Y holA fdnH Y holA gcvP holA infB Y holA ppx holA rplL Y holA rplV holA rpsC holA yehB holB holB Y holB cysJ holB dnaE Y	hns	rpoB		
hns secA Y hns ssb Y hns stpA Y hns tig Y holA holA Y holA accB holA accC Y holA b0878 Y holA dnaK Y holA fdnH Y holA gcvP holA grpE Y holA infB Y holA pepB Y holA rplL Y holA rplV holA rpsC holA yehB Y holB aceE Y holB cysJ Y holB cysJ Y holB dnaE Y	hns	rpoC		
hns stpA Y hns stpA Y hns tig Y holA holA Y holA accB Y holA accC Y holA b0878 Y holA dnaK Y holA fdnH Y holA gcvP Y holA infB Y holA ppx Y holA rplL Y holA rplC Y holA rpsC Y holB dnaE Y holB dnaE Y	hns	rpsB	Υ	Υ
hns stpA hns tig Y holA holA Y holA accB holA accC Y holA b0878 holA b1428 holA dnaK holA fdnH holA gcvP holA grpE holA infB holA ppx holA rplL holA rplP holA rpsC holB dnaE Y holB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y	hns	secA	Υ	
hns tig Y holA holA Y holA accB Y holA accC Y holA b0878 Y holA b1428 Y holA dnaK Y holA fdnH Y holA gcvP Y holA infB Y holA infB Y holA pepB Y holA rplL Y holA rplP Y holA rplP Y holA rpsC Y holB aceE Y holB dnaE Y	hns	ssb	Υ	
holA holA Y holA accB Y holA accC Y holA b0878 Y holA b1428 Y holA dnaK Y holA fdnH Y holA gcvP Y holA infB Y holA infB Y holA pepB Y holA rplL Y holA rplP Y holA rplV Y holA rpsC Y holB dnaE Y	hns	stpA		Υ
holA accB holA accC holA b0878 holA b1428 holA dnaK holA fdnH holA gcvP holA grpE holA infB holA pepB holA ppx holA rplL holA rplP holA rplV holA rpsC holA yehB holB holB holB cysJ holB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y	hns	tig	Υ	
holA accC holA b0878 Y holA b1428 Y holA dnaK Y holA fdnH Y holA gcvP HolA grpE Y holA infB Y holA pepB Y holA ppx HolA rplL Y HolA rplP Y HolA rplP Y HolA rpsC Y HolB dnaE Y HolB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	holA	holA	Y	
holA b0878 holA b1428 Y holA dnaK holA fdnH Y holA gcvP holA grpE Y holA infB HolA pepB Y holA ppx HolA rplL Y HolA rplP Y HolA rplV HolA rpsC HolA yehB HolB holB Y HolB dnaE Y Y HolB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	holA	accB		Υ
holA b1428 holA dnaK holA fdnH holA gcvP holA grpE holA infB holA pepB holA ppx holA rplL holA rplP holA rplP holA rpsC holA yehB holB holB cysJ holB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y	holA	accC		Υ
holA dnaK holA fdnH holA gcvP holA grpE holA infB holA pepB holA ppx holA rplL holA rplP holA rplV holA rpsC holA yehB holB holB holB cysJ holB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y	holA	b0878		
holA fdnH Y holA gcvP Y holA grpE Y holA infB Y holA pepB Y holA ppx Y holA rplL Y holA rplP Y holA rplV Y holA rpmG Y holA rpsC Y holA yehB Y holB aceE Y holB dnaE Y	holA	b1428		
holA gcvP holA grpE Y holA infB Y holA pepB Y holA ppx HolA rplL Y holA rplP Y holA rplV Y HolA rpmG Y HolA rpsC Y HolA yehB Y HolB aceE Y HolB dnaE Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	holA	dnaK		
holA grpE holA infB holA pepB holA pepB holA ppx holA rplL holA rplP holA rplV holA rpmG holA rpsC holA yehB holB holB holB cysJ holB dnaE Y	holA	fdnH		Υ
holA infB holA pepB holA pepB Y holA ppx holA rplL Y holA rplP Y holA rplV Y holA rpmG Y holA rpsC Y holA yehB Y holB holB Y holB aceE Y holB dnaE Y	holA	gcvP		Υ
holA pepB Y holA ppx Y holA rplL Y holA rplP Y holA rplV Y holA rpmG Y holA rpsC Y holA yehB Y holB holB Y holB aceE Y holB dnaE Y	holA	grpE		Υ
holA ppx holA rplL holA rplL holA rplP holA rplV holA rpmG holA rpsC holA pehB holB y holB aceE holB cysJ Y holB dnaE Y	holA	infB		
holA rplL Y holA rplP Y holA rplV Y holA rpmG Y holA rpsC Y holA yehB Y holB holB Y holB aceE Y holB cysJ Y holB dnaE Y	holA	pepB		
holA rplP Y holA rplV Y holA rpmG Y holA rpsC Y holA yehB Y holB holB Y holB aceE Y holB cysJ Y holB dnaE Y	holA	ppx		Υ
holA rplV Y holA rpmG Y holA rpsC Y holA yehB Y holB holB Y holB aceE Y holB cysJ Y holB dnaE Y	holA	rpIL		
holA rpmG Y holA rpsC Y holA yehB Y holB holB Y holB aceE Y holB cysJ Y holB dnaE Y	holA	rpIP		
holA rpsC Y holA yehB Y holB holB Y holB aceE Y holB cysJ Y holB dnaE Y	holA	rpIV		
holA yehB Y holB holB Y holB aceE Y holB cysJ Y holB dnaE Y	holA	rpmG		Υ
holB holB Y holB aceE Y holB cysJ Y holB dnaE Y	holA	rpsC		
holB aceE Y holB cysJ Y holB dnaE Y	holA	yehB		Υ
holB cysJ Y holB dnaE Y	holB	holB	Υ	
holB dnaE Y	holB	aceE	Υ	
	holB	cysJ		Υ
holB dnaX Y Y	holB	dnaE	Υ	
	holB	dnaX	Υ	Υ

1			
holB	holA	Υ	
holB	holD		Y
holB	rplA	Y	Y
holB	rplC	Υ	V
holB	rplJ		Y Y
holB holB	rpmB	Y	Ť
holB	rpsJ rpsP	Ī	Υ
holB	secA	Y	ı
holB	ugpB	1	Υ
holB	ybbN		Ϋ́
holC	holC	Y	Y
holC	b1374	•	Ϋ́
holC	b1808	Y	•
holC	clpA	Ϋ́	
holC	dnaE	Ý	Υ
holC	dnaK	Ϋ́	-
holC	dnaQ	Y	Υ
holC	dnaX	Y	Y
holC	glnD	Υ	
holC	holA	Υ	Υ
holC	holB	Υ	Υ
holC	holD	Υ	Υ
holC	holE		Y
holC	intE	Υ	
holC	lon	Υ	
holC	priA	Υ	
holC	recQ	Υ	
holC	rplB	Υ	Υ
holC	rplC	Υ	Υ
holC	rplD	Υ	Υ
holC	rpIL		Υ
holC	rpIS		Υ
holC	rpIV _		Y
holC	rpmB		Y
holC	rpmC		Y
holC	rpmG		Y
holC	rpsB	Y	V
holC	rpsE	V	Υ
holC	rpsJ	Υ	V
holC	rpsN	V	Y
holC	sbcB	Y Y	
holC	secA	Ϋ́Υ	Υ
holC	ssb	Ť	Ţ

halo	4	V	V
holC	topA	Y	Y
holC	topB	Y Y	
holC holC	uvrA yhiR	Ϋ́	
holD	holD	Y	Υ
holD	b1410	Y	1
holD	dnaE	Ϋ́	
holD	dnaQ	Ϋ́	
holD	dnaX	Ý	Y
holD	holA	Ϋ́	•
holD	holB	Ϋ́	
holD	holC	Ϋ́	Υ
holD	holE	-	Y
holD	recQ	Υ	
holD	rplC	Υ	
holD	rplL		Υ
holD	rplM	Υ	
holD	rpIU		Υ
holD	rpmB		Υ
holD	rpoC	Υ	
holD	secA	Υ	
holD	ssb	Υ	Υ
holD	topB	Υ	
holE	holE	Y	Y
holE	dnaE	Υ	Υ
holE	dnaK		Υ
holE	dnaQ	Υ	
holE	dnaX	Υ	Y
holE	hns		Υ
holE	holA	Υ	
holE	holB	Υ	
holE	holD	Υ	
holE	katG		Υ
holE	rpIA	Υ	
holE	rplL		Y
holE	rplW		Y
holE	rpsB	Υ	
holE	rpsF	V	Y
holE	rpsJ	Υ	V
holE	rpsP	Y	Y
holE	ssb top A	Y	V
holE	topA	Υ	Y
holE	tufA	ĭ	Y
holE	ugpB		Y

holE	yfiF		Υ
hrpA	hrpA	Υ	-
hrpA	aceE	Ϋ́	
hrpA	aceF	Y	
hrpA	lpdA	Ϋ́	
hrpA	rpll	Ϋ́	
hrpA	rpsA	Ϋ́	
hrpA	rpsB	Ϋ́	
hrpA	rpsC	Ϋ́	
hrpA	rpsE	Ϋ́	
-	rpsG	Ϋ́	
hrpA	•	Ϋ́	
hrpA	rpsM	Ϋ́	
hrpA	rpsS	Ϋ́	
hrpA	tufA		
hrpA	yfiF	Y	
hrpB	hrpB	Y	
hrpB	aceF	Y	
hrpB	lpdA	Y	Y
hrpB	rplL		Y
hscA	hscA	Y	Y
hscA	b2863		Υ
hscA	iscU	Υ	
hscA	pepB		Υ
hscA	rplJ	Υ	Υ
hscA	rpIL		Υ
hscA	rplT		Υ
hscA	secA	Υ	
hscA	ugpB		Υ
hsdM	hsdM	Y	Y
hsdM	elaB		Υ
hsdM	gloB	Υ	
hsdM	hsdR	Υ	Y
hsdM	nfi		Υ
hsdM	uvrB		Υ
hsdM	yigC	Υ	
hsdR	hsdR	Y	Y
hsdR	aceF	Υ	Υ
hsdR	hsdM	Υ	Y
hsdR	IpdA	Υ	Υ
hsdR	rpIL		Υ
hsdS	hsdS	Υ	
hsdS	aceF	Υ	Υ
hsdS	hsdM	Υ	Υ
hsdS	hsdR		Υ
i .			

hsdS	hupA		Y
hsdS	hupB		Y
hsdS	lpdA		Y
hsdS	malP		Y
hsdS	rpID		Υ
hsdS	rplU		Υ
hsdS	rpsG		Υ
hslU	hsIU	Υ	Υ
hslU	accB		Y
hslU	accC		Υ
hslU	accD		Υ
hslU	b1498		Υ
hslU	cca	Υ	
hslU	ccmB		Y
hslU	cysW	Y	
hsIU	dnaJ	Y	Y
hsIU	dnaK	Υ	Υ
hsIU	ftsZ	Y	
hslU	fusA	Y	
hslU	gatZ	Y	
hslU	gpmB	Υ	
hslU	hslV		Y
hslU	iciA		Υ
hslU	lon	Y	
hslU	metE	Y	
hsIU	metK	Υ	
hslU	mglB	V	Y
hsIU	mreB	Y	
hslU	narG	Y	
hslU	nuoC	Υ	
hslU	ompX	V	Y
hsIU	pstB	Y	
hslU	recA	Υ	V
hslU	ribD		Y Y
hslU	rplC		
hslU	rplD	V	Υ
hslU	rplJ	Y	Y
hslU hslU	rpIL		Ϋ́
hslU	rpIM		Y
hslU	rpIP rpsB		Ϋ́
hslU	rpsB rpsE		Ϋ́
	rpsE		Ϋ́
hslU	rpsG		Ϋ́Υ
hslU	rpsJ		Ť

hslU	secA	Υ	
hslU	thrA		Υ
hslU	tufA	Υ	Ϋ́
hslU	tufB	Ϋ́	•
hslU		1	Υ
	ydgA	Υ	ı
hslU	ygeG	Y	
hslV	hslV	Y	V
hslV	accA		Y
hslV	accB		Y
hslV	accC		Υ
hslV	accD		Υ
hslV	ccmB		Υ
hslV	deaD	Υ	
hslV	gabP		Υ
hsIV	hsIU	Υ	
hslV	rplB	Υ	
hslV	rplC	Υ	Υ
hslV	rplD	Υ	Υ
hslV	rpll	Υ	
hslV	rplJ	Υ	
hslV	rplL	•	Υ
hslV	rpIS	Υ	-
hslV	rplU	·	Υ
hslV	rpIV	Υ	Ϋ́
hslV	rplX	•	Ϋ́
hslV	rpmC		Ϋ́
hslV	rpmG		Ϋ́
hslV	rpsG	V	ı
	-	Y	
hslV	rpsM	Υ	V
hslV	rpsP	V	Y
hslV	tufA	Y	
hslV	tufB	Y	
hslV	yciL	Υ	
hslV	yhjD		Y
hslV	yihl		Υ
htpG	htpG	Y	Υ
htpG	hyfG		Υ
htpG	mcrB		Υ
htpG	pepB		Υ
htpG	rplB		Υ
htpG	rpmC		Υ
htpG	rpmG		Υ
htpG	rpoC	Υ	
htpG	rpsF		Υ
•	•		

htpG	rpsG		Υ
htpG	ugpB		Υ
htpG	ycbW		Υ
htpG	yeal		Υ
htpG	yggR		Υ
hupA	hupA	Y	Y
hupA	dnaK	Υ	
hupA	hupB		Υ
hupA	malP	Υ	
hupA	malQ	Υ	
hupA	rpoA	Υ	
hupA	yciL	Υ	
hupB	hupB	Y	Y
hupB	fis		Υ
hupB	himA		Υ
hupB	himD		Υ
hupB	hlpA		Υ
hupB	hns		Υ
hupB	hsdM	Υ	
hupB	hupA		Υ
hupB	malP	Υ	Υ
hupB	malQ	Υ	Υ
hupB	polA	Υ	Υ
hupB	rplC		Υ
hupB	rpoA		Υ
hupB	rpsG		Υ
hupB	seqA		Υ
hupB	yaiD		Υ
hybC	hybC	Υ	Υ
hybC	pepQ	Υ	
hybC	rplL		Y
hybC	rplV		Y
hybC	rpmB		Y
hybC	rpsB		Y
hybC	rpsJ		Y
hybC	rpsN	V	Y
hybC	tufA	Y	Y
hybC	tufB	Y Y	
<i>hybE</i>	<i>hybE</i> b2997	Ϋ́	
hybE <i>hycE</i>	hycE	Ϋ́	
hycl	hycL	Y	V
hycl	aceE	Y	ı
hycl	IpdA	ı	Υ
HyCi	ipuA		ı

1			
hycl	yafQ		Y
hyfG	hyfG	Y	
hyfG	mopA	Y	Y
hypC	hypC	Y	Y
hypC	hycE	Υ	Υ
hypC	hyfG		Υ
hypC	hypD	Υ	Υ
hypC	lysS		Υ
hypC	rpsG		Y
hypC	tig		Υ
hypD	hypD	Υ	Υ
hypD	hypC	Υ	Y
hypD	hypE		Υ
hypD	mopA	Υ	
hypD	tufA	Υ	Υ
hypD	tufB	Υ	
hypE	hypE	Y	
hypF	hypF	Y	
ibpA	ibpA	Υ	Y
ibpA	add	Υ	
ibpA	atpD	Υ	
ibpA	clpA	Υ	Υ
ibpA	clpB		Y Y
ibpA	dnaJ		Υ
ibpA	eno		Υ
ibpA	ftsZ	Υ	
ibpA	gadA	Υ	
ibpA	gatA		Υ
ibpA	gatB		Υ
ibpA	gatY	Υ	
ibpA	gpmA		Υ
ibpA	hrpB		Υ
ibpA	ibpB	Υ	Υ
ibpA	infB		Υ
ibpA	IpdA		Υ
ibpA	malK	Υ	Υ
ibpA	malT	Υ	
ibpA	malZ		Υ
ibpA	metK	Υ	Υ
ibpA	mreB	Υ	
ibpA	msbB		Υ
ibpA	ompC	Υ	
ibpA	pflB	Υ	Υ
ibpA	pnp	Υ	Υ

۸ میرا:			V
ibpA	prc		Y
ibpA	proS	V	Y
ibpA	pstB	Υ	Y Y
ibpA	radC	Υ	Ĭ
ibpA ibpA	rpoB rpoC	Υ	
ibpA	rpsA		Υ
ibpA	rpsB		Ϋ́
ibpA	rpsF		Ϋ́
ibpA	rpsG		Ϋ́
ibpA	tufA	Υ	Ϋ́
ibpA	tufB	Ϋ́	·
ibpA	uspA	-	Υ
ibpA	ybdQ		Y
ibpA	ybiY		Υ
ibpA	yccD		Υ
ibpA	yjjL	Υ	
ibpB	ibpB	Y	
ibpB	add	Υ	
ibpB	gadB	Υ	
ibpB	gatY	Υ	
ibpB	ibpA		Υ
ibpB	IpdA	Υ	
ibpB	malK	Υ	
ibpB	malT	Υ	
ibpB	mreB	Υ	
ibpB	pstB	Υ	
ibpB	tufA	Υ	
iciA	iciA	Υ	
iciA	mopA	Y	
iciA	tufA	Y	
icIR	icIR _	Υ	Y
iclR	aceF		Υ
iclR	rfaD	Υ	V
iclR	rpIA		Y
iclR	rpID		Y
iclR	rplM		Y Y
iclR iclR	rpIT rpILI		Υ Υ
iclR	rpIU rpIV		Ϋ́
icIR	rplX		Ϋ́
iclR	rpsB		Y
iclR	rpsE		Ϋ́
icIR	rpsE		Ϋ́
1011 (ipoi		•

iclR	rpsG		Υ
iclR	rpsH		Y
iclR	rpsM		Ϋ́
	-		Ϋ́
iclR	rpsN		
idnD	idnD	Y	Y
idnD	lpdA	Υ	
idnD	lysS		Υ
idnD	yecC		Υ
idnO	idnO	Y	
ileS	ileS	Υ	Υ
ileS	hyaD		Υ
ileS	ppk		Υ
ileS	rbsB		Υ
ileS	tufA	Υ	-
ileS	tufB	Ý	
ileS	ybbU	•	Υ
ileS	-		Ϋ́
	yjgL <i>i</i> uD	V	Y
ilvB	ilvB	Y	
ilvB	aceE	Υ	Y
ilvB	fumA		Y
ilvB	IpdA	Υ	Υ
ilvB	rplL		Υ
ilvB	rplM		Υ
ilvB	rpsB		Υ
ilvB	ycaH		Υ
infB	infB	Υ	Υ
infB	abc		Υ
infB	dnaK		Υ
infB	grpE		Υ
infB	ilvG_1		Y
infB	phoA		Ϋ́
infB	ribD	Y	•
infB	rplC	Y	Υ
infB	rplM	Y	ı
infB	•	ī	Y
	rpmG	V	Ϋ́
infC	infC	Y	Y
infC	b1410	Y	
infC	cafA	Υ	
infC	ccmB		Υ
infC	dacA		Υ
infC	hupB		Υ
infC	rpIB	Υ	Υ
infC	rpIC	Υ	Y
infC	rpID	Υ	Y

infC infC infC infC infC infC	rpIE rpIJ rpIK rpIL rpIM	Y Y Y	Y Y Y
infC infC	rplO rplP		Y Y
infC infC	rpIS rpIT		Y Y
infC	rplU		Υ
infC	rpIV		Y
infC infC	rpIX rpmG		Y Y
infC	rpsA	Υ	Ϋ́
infC	rpsB	Υ	Υ
infC	rpsC	Y	Y
infC infC	rpsD rpsE	Y Y	Y Y
infC	rpsF	Ϋ́	Ϋ́
infC	rpsG	Υ	Υ
infC	rpsH		Y
infC	rpsl	V	Y
infC infC	rpsJ rpsK	Y Y	Y
infC	rpsM	Ý	Υ
infC	rpsN		Υ
infC	rpsO		Υ
infC	rpsP		Y
infC infC	rpsR rpsS		Y Y
infC	rpsT		Ϋ́
infC	rpsU		Υ
infC	yfiF		Y
infC infC	yhbY ynhD		Y Y
intA	intA	Υ	ı
intA	aceF	Y	Υ
intA	yihl	_	Υ
intC	intC	Y	V
intC intC	aceF cspG	Y	Y Y
intC	lpdA	Υ	•
iscS	iscS	Υ	Y

iscS	cobB	Υ	
iscS	dnaK	Υ	
iscS	dniR		Υ
iscS	fdhD	Υ	
iscS	iscU	Υ	Υ
iscS	yhhP		Υ
iscU	iscU	Υ	Υ
iscU	b1121		Υ
iscU	cfa		Υ
iscU	damX		Υ
iscU	dedA		Υ
iscU	dnaK	Υ	-
iscU	edd		Υ
iscU	hscA	Υ	Y
iscU	iscS	Y	Y
iscU	pheP	-	Y
iscU	proA		Y
iscU	sucD		Y
ispA	ispA	Y	Ϋ́
ispA	b2460	•	Ϋ́
ispA	dnaK		Ϋ́
ispA	flxA		Ϋ́
ispA	purT		Ϋ́
ispA	rplC		Y
ispA	rplK		Ϋ́
ispA	rplL		Ϋ́
ispA	rplM		Ϋ́
ispA	rpmG		Ϋ́
ispA	rpsB		Y
ispA	rpsJ		Ϋ́
katE	katE	Y	Ϋ́
kdsA	kdsA	Ϋ́	Ϋ́
kdsA	b2434	•	Ϋ́
kdsA	gyrA	Υ	•
kdsA	hupA	•	Υ
kdsA	kdsB	Υ	•
kdsA	metK	Ý	
kdsA	mopA	Ϋ́	
kdsA	murE	•	Υ
kdsA	rcsC	Υ	•
kdsA	rfaD	Ý	
kdsA	rplA	Ý	
kdsA	rpIC	•	Υ
kdsA	rpID		Ϋ́
Nusa	יאים		•

1			
kdsA	rplE		Υ
kdsA	rpIF		Υ
kdsA	rpll		Y
kdsA	rpIL		Υ
kdsA	rplM		Y
kdsA	rplN		Υ
kdsA	rpIS	Υ	Υ
kdsA	rplU		Υ
kdsA	rplV	Υ	Υ
kdsA	rplX	Υ	Υ
kdsA	rplY		Υ
kdsA	rpmB		Υ
kdsA	rpmG		Υ
kdsA	rpsB		Υ
kdsA	rpsC	Υ	Υ
kdsA	rpsD		Υ
kdsA	rpsE		Υ
kdsA	rpsG		Υ
kdsA	rpsl		Υ
kdsA	rpsJ		Υ
kdsA	rpsK		Υ
kdsA	rpsM	Υ	
kdsA	rpsN		Υ
kdsA	rpsT		Υ
kdsA	tufA	Υ	Υ
kdsA	tufB	Υ	
kdsA	yagG		Υ
kdsA	ybdN	Υ	
kdsA	yfiF		Υ
kdsA	yihR		Y
kdsB	kdsB	Y	Υ
kdsB	dnaJ	•	Ϋ́
kdsB	dnaK		Y
kdsB	hupB		Ϋ́
kdsB	рерВ		Ϋ́
kdsB	rpID		Ϋ́
kdsB	rplK		Ϋ́
kdsB	rpIL		Ϋ́
kdsB	rpsA		Ϋ́
kdsB	tufA	Υ	•
kdsB	tufB	Ý	
kdsB	yfbS	•	Υ
kdsB	yjaH		Y
kdsB	yjdG		Ϋ́
KU3D	yjuG		ı

kdtA	kdtA	Y	
kdtA	entB		Υ
kdtA	murE		Υ
kdtA	rplA		Υ
kdtA	rplE		Υ
kdtA	rplL		Y
kdtA	rplM		Y
kdtA	rpIS		Υ
kdtA	rplT		Υ
kdtA	rplU		Υ
kdtA	rplV		Υ
kdtA	rpmG		Υ
kdtA	rpsB		Υ
kdtA	rpsE		Y
kdtA	rpsJ		Υ
kdtB	kdtB	Y	-
kdtB	murE		Υ
kdtB	rplC		Y
kdtB	rplD	Y	
kdtB	rplL		Υ
kdtB	rpsE		Y
kdtB	yciA		Υ
ksgA	ksgA	Υ	
IctD	IctD		Y
IctD	glyQ		Υ
IctD	radC		Υ
IctD	rfaD		Υ
IctD	rplV		Υ
IctD	rpmB		Υ
IctD	rpsB		Υ
IdcC	IdcC	Υ	Y
ldcC	cadA	Υ	Υ
<i>lepA</i>	<i>lepA</i>	Υ	Y
lepA	argA		Υ
lepA	b0878		Υ
lepA	b1342		Υ
lepA	hyfE		Υ
lepA	recE		Υ
lepA	rplD		Υ
lepA	rpll		Υ
lepA	rplM		Υ
lepA	rplV		Υ
lepA	rplX		Υ
lepA	rpsB		Υ
•	-		

	_		
lepA	rpsE		Y
lepA	rpsF		Y
lepA	rpsG		Y
lepA	rpsJ		Y
lepA	ycjJ		Y
lepA	yeiN		Y
lepA	yidX		Y
leuS	leuS	Y	Y
leuS	htrE		Y
leuS	map		Y
leuS	rplC		Y
leuS	rpID		Υ
leuS	rplK		Υ
leuS	rplL		Υ
leuS	rplW		Υ
leuS	rpsB		Υ
leuS	rpsJ		Υ
leuS	rpsP		Υ
leuS	yeaP		Υ
<i>lexA</i>	<i>lexA</i>	Y	
lexA	rpsE	Y	
lig 	lig_	Y	Υ
lig	aceE	Y	
lig 	leuS	Υ	
lig 	lpxD		Y
lig 	mreC	3.4	Υ
lig	recE	Υ	
lig	rplL		Y
lig .:	rpsB		Υ
lig 	yaiD	Y	
lig	ykgC		Y
IoIA	IoIA	Y	Y
IoIA	aceE	Y	V
IoIA	b2809	V	Υ
IoIA	lon	Υ	V
IoIA	rplL		Y
IoIA	rplM		Y Y
IoIA	rpIU		Y Y
IoIA	rpIV	V	Y
lolA	rpsJ	Y Y	Υ
<i>lon</i> lon	lon minB	r	Ϋ́Υ
	mipB	Υ	Ĭ
lon	rpIC	Ϋ́Υ	Y
lon	rpll	Ť	Y

lon	rplM	Υ	Υ
	-	Ϋ́	•
lon	rplT	ĭ	V
lon	rplU		Y
lon	rplV		Υ
lon	rpsB	Υ	
lon	rpsC	Υ	
lon	rpsG		Υ
lon	rpsH		Υ
lon	ssb		Υ
lon	tufA	Υ	
lon	ydbA_1	-	Υ
lon	ydb/ (_ \ yjgL		Ϋ́
IpIA	lpIA	Υ	Y
	entC	,	Ϋ́
lplA			
lplA	ybdH		Y
lpxA	lpxA	Y	Y
lpxA	b1686		Y
lpxA	dniR		Υ
lpxA	fhuB		Υ
lpxA	hcaB		Υ
lpxA	hflB		Υ
lpxA	hfq		Υ
lpxA	ribB		Υ
lpxA	rplC		Υ
lpxA	rplF		Υ
lpxA	rplL		Υ
lpxA	rplM		Y
lpxA	rplN		Ϋ́
lpxA	rpIT		Ϋ́
lpxA	rplU		Ϋ́
lpxA	rplV		Y
lpxA	rplX		Y
lpxA	rpmB		Y
lpxA	rpmG		Y
lpxA	rpsB		Υ
lpxA	rpsl		Υ
lpxA	rpsJ		Υ
lpxA	yqcD		Υ
lpxB	lpxB	Υ	
lpxB	bacA		Υ
lpxB	rplM		Υ
lpxB	rpIS		Υ
lpxB	rpsB		Υ
lpxB	torC		Y
			•

lpxB	tufA	Υ	Υ
lpxB	tufB	Υ	
lpxD	lpxD	Y	Υ
lpxD	aspA	,	Ϋ́
-	b1371		Ϋ́
lpxD		V	ı
lpxD	b1555	Υ	V
lpxD	b1605		Y
lpxD	b2810		Y
lpxD	cbpA	Υ	Υ
lpxD	ccmB		Υ
lpxD	dnaJ		Υ
lpxD	glnS	Υ	
lpxD	pssA	Υ	
lpxD	rfaD	Υ	
lpxD	rho	Υ	
lpxD	rnk	•	Υ
lpxD	rplB	Υ	•
lpxD	rplC	•	Υ
lpxD	rpID	Υ	Ϋ́
-	-	Y	
lpxD	rpll	Y	Y
lpxD	rplK		Y
lpxD	rplL		Y
lpxD	rplM		Υ
lpxD	rplO	Υ	Υ
lpxD	rplP		Υ
lpxD	rpIS		Υ
lpxD	rplT		Υ
lpxD	rplU	Υ	Υ
lpxD	rpIV		Υ
lpxD	rplX		Υ
lpxD	rpmG		Y
lpxD	rpsB	Υ	Ϋ́
lpxD	rpsC	Ϋ́	Ϋ́
•	•	Y	Ϋ́
lpxD	rpsD		
lpxD	rpsE	Y	Y
lpxD	rpsG	Υ	Y
lpxD	rpsH		Y
lpxD	rpsJ		Y
lpxD	rpsP		Υ
lpxD	rpsR		Υ
lpxD	rpsT		Υ
lpxD	tyrA	Υ	
lpxD	yciL	Υ	
lpxD	yfiF		Υ

lpxD	yhbY		Y
lpxD	yjfl		Y
lysA	lysA	Y	Υ
lysA	cpxR		Υ
lysS	lysS	Υ	Υ
lysS	b2341	Υ	
lysS	bioA		Υ
lysS	gcd		Υ
lysS	lysU	Υ	Y
lysS	pdxJ		Υ
lysS	rhaS		Υ
lysS	rhsA	Υ	
lysS	tdcD	Υ	
lysS	ygeV		Υ
lysS	ytfG		Υ
lysU	İysU	Y	Y
lysU	aceE	Υ	
lysU	adhE	Υ	
lysU	b1644		Υ
lysU	b2343		Υ
lysU	clpB	Υ	-
lysU	cspC	•	Υ
lysU	cysK		Ϋ́
lysU	dnaN	Υ	•
lysU	dps	Y Y	Υ
lysU	eno	Ý	Ϋ́
lysU	focB	•	Ϋ́
lysU	fusA	Υ	•
lysU	gadA	Ϋ́	
lysU	gadA gadB	Y	
lysU		Y	
_	gapA katE	Y	
lysU		Y	
lysU	lysA	Ĭ	V
lysU	lysS	V	Υ
lysU	mopA	Y	V
lysU	pflB	Y	Υ
lysU	pgk	Y	
lysU	rplB	Y	V
lysU	rpll		Y
lysU	rplX		Y
lysU	rplY		Υ
lysU	rpoB	Y	
lysU	rpoC	Y	
lysU	rpsB	Υ	Υ

lysU	rpsC		Υ
lysU	rpsG		Υ
lysU	rpsJ		Υ
lysU	rpsK	Υ	
lysU	serS		Υ
lysU	tnaA	Υ	
lysU	tufA	Υ	Υ
lysU	tufB	Υ	
lysU	wcal		Υ
lysU	wrbA		Υ
lysU	yagJ		Υ
lysU	yfiD		Υ
lysU	ygiW		Υ
lysU	yhbM		Υ
lysU	yhjX		Υ
lytB	lytB	Υ	Y
lytB	mopA		Υ
lytB	rpID		Υ
lytB	rpIL		Υ
malK	malK	Y	
malP	malP	Y	Υ
malP	aceF	Υ	Υ
malP	hupA		Υ
malP	IpdA	Υ	Υ
malP	pnp	Υ	
malP	rplA	Υ	
malP	rplC	Υ	Υ
malP	rplD	Υ	
malP	rpll		Υ
malP	rplM		Υ
malP	rpIS		Υ
malP	rplU		Υ
malP	rplV		Υ
malP	yhbY		Υ
malT	malT	Y	
malT	dnaJ	Y	
malT	fixB	Y	
malT	fliJ	Y	
malT	metK	Y	
malT	mopA	Y	
malT	mreB	Y	
malT	pstB	Y	
malT	rho	Y	
malT	rplS	Υ	

malT	tufA	V	
malT	tufB	Y Y	
		Y	V
manA	manA	1	Y Y
manA	b1559		
manA	minC		Y
manA	ycbY		Y
manA	yrdC	V	Υ
manX	manX	Υ	
manX	rplV		Y
manX	rpmB		Y
map	map	Y	Y
map	deaD		Y
map	recE		Y
map	rpID		Υ
map	rpll		Υ
map	rplL		Υ
map	rplM		Υ
map	rpIS		Υ
map	rplV		Υ
map	rplW		Υ
map	rpmB		Υ
map	rpmG		Υ
map	rpsB		Υ
map	rpsE		Υ
map	rpsJ		Υ
map	rpsP		Υ
map	rpsT		Υ
map	uidR		Υ
map	yeaP		Υ
map	yegN		Υ
map	yhbY		Υ
map	ylaD		Υ
mazG	mazG	Υ	Υ
mazG	elaB		Y
mazG	mopA	Υ	
mcrB	mcrB	Y	
mcrB	yjeF	-	Υ
menB	menB	Υ	•
menB	rfaD	Ϋ́	Υ
menB	rpsN	•	Ϋ́
menC	menC	Υ	Ϋ́
menD	menD	Ϋ́	Ϋ́
menF	menF	Ϋ́	•
menG	menG	Ϋ́	Υ
,,,,,,,,		ı	,

metK	metK	Y	Y
metK	dnaJ	Y	Υ
metK	dnaK	Y	
metK	glyQ	Y	
metK	mreB	Y	
metK metK	pstB	Y Y	
metK	recA	Ϋ́	
metK	secA tufA	Ϋ́	Υ
mfd	mfd	Y	Y
mfd	tdcC	1	Y
mfd	yffB		Y
moaA	упь тоаА	Υ	ī
moaA	gyrB	1	Υ
moaA	mopA	Υ	Ϋ́
moaA	rplU	'	Ϋ́
moaA	rpsG		Ϋ́
moaA	smpB		Ϋ́
moaA	tufA	Υ	•
moaA	tufB	Ϋ́	
moaB	moaB	Ϋ́	
moaC	moaC	Y	Υ
moaC	rplA	Ϋ́	,
moaC	rpIS	•	Υ
moaC	rplV		Ϋ́
moaC	rpmB		Ϋ́
moaD	moaD	Y	Ϋ́
moaD	rplL	,	Ϋ́
moaE	moaE	Y	Ϋ́
moaE	moaD	Ϋ́	Ϋ́
mobA	mobA	Y	•
mobA	rplV	•	Υ
mobB	mobB	Υ	Y
mobB	moaE	•	Ϋ́
moeA	moeA	Y	Y
moeA	pfkA	Υ	
moeB	moeB	Y	
moeB	dnaK	Y	
mog	mog	Y	Υ
mog	dnaK	Υ	
mog	gyrA	Y	
mopA	mopA	Υ	Υ
mopA	aceE	Υ	
mopA	clpB	Υ	
	-		

mopA	dnaK		Υ
mopA	grpE		Υ
mopA	lon	Υ	
mopA	narG	Υ	
mopA	pyrD		Υ
mopA	rpsB	Υ	
mopA	ybbN	Υ	
mopA	ycgR		Υ
mopA	yhdJ		Υ
mopA	yibA	Υ	
торВ	торВ	Y	Y
mopB	dnaK	Υ	Υ
mopB	dnaN	Υ	Υ
mopB	fepB		Υ
mopB	fusA	Υ	
mopB	htpG	Υ	
mopB	mreB	Υ	
mopB	rfaD	Υ	Υ
mopB	rpsB		Υ
mopB	slyD		Υ
mopB	tig	Υ	Υ
mopB	ybbP		Υ
mopB	ydbA 1		Υ
mreB	mreB	Y	Y
mreB	accA		
mreB	add	Y Y	
mreB	atpD	Υ	
mreB	b1685		Υ
mreB	clpB	Υ	
mreB	dnaJ	Υ	Υ
mreB	dnaK	Υ	Υ
mreB	fusA	Υ	
mreB	gatB		Υ
mreB	gatY	Υ	
mreB	glyQ	Υ	
mreB	gyrA	Υ	
mreB	ibpA		Υ
mreB	kefC	Υ	
mreB	lysU	Υ	
mreB	malK	Υ	
mreB	malT	Υ	
mreB	minD	Υ	
mreB	mopA	Υ	Υ
mreB	narG	Y	
I .			

mreB nuoC mreB nusG mreB ompA mreB ompC mreB ompC mreB pnp mreB pstB mreB recA mreB recA mreB rho mreB rplC mreB rplE mreB rpoB mreB rpoC mreB rpsF mreB secA mreB secA mreB tufA mreB tufA mreB ybdQ mreB ybeD mreB yjgD mreB yjgD mreB yleA mrr mrr mrr mrr mrr mrr mrsA mrsA mrsA rplC mrsA				
mreB ompA Y mreB ompC Y mreB pnp Y mreB pstB Y mreB recA Y mreB rho Y mreB rplC Y mreB rplE Y mreB rpoB Y mreB rpoC Y mreB secA Y mreB secA Y mreB sws Y mreB tufA Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB ybeD Y mreB yleA Y mrsA mrsA Y mrsA mrsA Y mrsA rplK Y mrsA rplK Y mrsA rplK Y mrsA rplK Y mrsA rpsB	mreB	nuoC	Υ	
mreB ompC y mreB pnp y mreB pstB Y mreB recA Y mreB rho Y mreB rplC Y mreB rplC Y mreB rplE Y mreB rpoC Y mreB rpoC Y mreB secA Y Y mreB secA Y Y mreB stufA Y Y mreB tufB Y mreB ybdQ Y mreB ybdQ Y mreB ybeD Y mreB yleA Y mrsA mrsA Y Y mrsA mrsA Y Y mrsA rplC Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y mtlD b1458 mtlD dppA Y mtlD entB mtlD ybbP mukB mukB Y mukB mukB Y mukB mukB Y mukB mukB Y mukB mukB Y	mreB	nusG		Υ
mreB ompC Y mreB pnp Y mreB pstB Y mreB recA Y mreB rho Y mreB rplC Y mreB rplE Y mreB rpoC Y mreB rpoC Y mreB secA Y Y mreB secA Y Y mreB stufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB ybeD Y mreB yleA Y mrsA mrsA Y Y mrsA mrsA Y Y mrsA rplK Y mrsA rplK Y mrsA rplK Y mrsA rplK Y mrsA rpsB	mreB	ompA	Υ	
mreB pstB Y mreB pstB Y mreB recA Y mreB rho Y mreB rplC Y mreB rplE Y mreB rpoB Y mreB rpoC Y mreB rpsF Y mreB secA Y Y mreB sms Y mreB tufA Y Y mreB ybdQ Y mreB ybdQ Y mreB ybeD Y mreB yjgD Y mreB yleA Y mrsA mrsA mrsA Y Y mrsA rplK Y mrsA rplK Y mrsA rplK Y mrsA rpsB	mreB	-	Υ	
mreB pstB Y mreB recA Y mreB rho y mreB rplC Y mreB rplE Y mreB rpoB Y mreB rpoC Y mreB rpsF Y mreB secA Y Y mreB stufA Y Y mreB tybdQ Y mreB ybdQ Y mreB ybeD Y mreB yjgD Y mreB yleA Y mrsA mrsA mrsA Y Y mrsA rplK Y mrsA rplK Y mrsA rplK Y mrsA rpsB Y mrsA r	mreB	-		
mreB recA Y mreB rho y mreB rplC Y mreB rplE Y mreB rpoB Y mreB rpoC Y mreB rpsF Y mreB secA Y Y mreB stufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB yjgD Y mreB yjgD Y mreB yleA Y mrsA mrsA Y Y mrsA rplK Y mrsA rplK Y mrsA rplK Y mrsA rplK Y mrsA rpsB Y mrsA rpsB Y mrsA rpsE Y mrsA rpsE Y mrsA tufB Y mrsA tufB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtiD mtiD Y mtiD b1458 mtiD dppA Y mukB mukB Y mukB acpP	mreB			
mreB rho mreB rpIC mreB rpIE Y mreB rpoB Y mreB rpoC Y mreB rpsF Y mreB secA Y Y mreB sms Y mreB tufA Y Y mreB tybdQ Y mreB ybdQ Y mreB ybeD Y mreB yjgD Y mreB yjgD Y mreB yleA Y mrsA mrsA Y Y mrsA rpIK Y mrsA rpIK Y mrsA rpIK Y mrsA rpIK Y mrsA rpSB Y mrsA rpSB Y mrsA rpSB Y mrsA rpSB Y mrsA tufB Y mrsA tufB Y mrsA tufB Y mrsA tufB Y mrsA tufB Y mrsA yggB Y mriD mtID Y mtID b1458 mtID dppA Y mukB mukB Y mukB mukB Y mukB acpP		-		
mreB rplE Y mreB rpoB Y mreB rpoC Y mreB rpsF Y mreB secA Y Y mreB sms Y mreB tufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB yleA Y mrr mrr Y Y mrsA mrsA Y Y mrsA rplK Y mrsA rplK Y mrsA rplL Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufB Y mrsA tufB Y mrsA yggB Y mrtD mtlD Y Y mtlD b1458 mtlD dppA Y mukB mukB Y mukB mukB Y mukB acpP	mreB	rho		
mreB rplE Y mreB rpoB Y mreB rpoC Y mreB rpsF Y mreB secA Y Y mreB sms Y mreB tufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB yleA Y mrr mrr Y Y mrsA mrsA Y Y mrsA rplK Y mrsA rplK Y mrsA rplL Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufB Y mrsA tufB Y mrsA yggB Y mrtD mtlD Y Y mtlD b1458 mtlD dppA Y mukB mukB Y mukB mukB Y mukB acpP	mreB	rplC		Υ
mreB rpoB Y mreB rpoC Y mreB rpsF Y mreB secA Y Y mreB sms Y mreB tufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB ylgD Y mreB ylgD Y mreB yleA Y mrr mrr Y Y mrsA mrsA Y Y mrsA rplK Y mrsA rplK Y mrsA rplL Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufA Y mrsA yggB Y mrsA yggB Y mtlD mtlD Y mtlD b1458 mtlD dppA mtlD dppA mtlD dppA mtlD entB mtlD ybbP mukB mukB Y mrsA ry mrsB ry mrsB ry mrsB mukB Y mrsB ry mrsB ry mrsB ry mrsA	mreB	•	Υ	
mreBrpoCYmreBrpsFYmreBsecAYYmreBsmsYmreBtufAYYmreBtufBYmreBybdQYmreBybeDYmreByjgDYmreByleAYmrrmrrYYmrsAmrsAYYmrsAmurCYmrsArplKYmrsArplKYmrsArplLYmrsArplWYmrsArpsBYmrsArpsBYmrsArpsBYmrsAtufAYmrsAtufBYmrsAyggBYmtlDmtlDYmtlDdppAYmtlDdppAYmtlDpohAYmukBmukBYmukBmukBYmukBacpPY	mreB	-	Υ	
mreB rpsF mreB secA Y mreB sms Y mreB tufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB yjgD Y mreB yjgD Y mreB yleA Y mrr mrr mrr Y Y mrsA murC Y mrsA rplK Y mrsA rplK Y mrsA rplL Y mrsA rplW Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufB Y mrsA yggB Y mrsA yggB Y mrsA yggB Y mrlD mtlD Y mtlD b1458 mtlD dppA Y mtlD entB Y mukB mukB Y mukB acpP	mreB	_	Υ	
mreB sms Y mreB tufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB yjgD Y mreB yleA Y mrr mrr Y Y mrsA mrsA Y Y mrsA nusG Y mrsA rplK Y mrsA rplK Y mrsA rplW Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufA Y mrsA yggB Y mrsA yggB Y mtlD mtlD Y mtlD b1458 mtlD dppA	mreB	_		Υ
mreB sms Y mreB tufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB yjgD Y mreB yleA Y mrr mrr Y Y mrsA mrsA Y Y mrsA nusG Y mrsA rplK Y mrsA rplK Y mrsA rplW Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufA Y mrsA yggB Y mrsA yggB Y mtlD mtlD Y mtlD b1458 mtlD dppA		•	Υ	
mreB tufA Y Y mreB tufB Y mreB ybdQ Y mreB ybeD Y mreB yjgD Y mreB yjeA Y mrr mrr Y Y mrsA mrsA Y Y mrsA nusG Y mrsA rplK Y mrsA rplL Y mrsA rplV Y mrsA rpsB Y mrsA rpsE Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD dppA mtlD d	mreB	sms	Υ	
mreB ybdQ mreB ybeD mreB yjgD y mreB yleA y mrr mrr Y y mrsA mrsA Y y mrsA nusG y mrsA recA y mrsA rplK y mrsA rplM y mrsA rplW y mrsA rpsB y mrsA rpsE y mrsA tufA y mrsA yggB y mtlD mtlD y mtlD dppA mtlD d	mreB			Υ
mreB yjgD Y mreB yjgD Y mreB yleA Y mrr mrr Y Y mrsA mrsA Y mrsA murC Y mrsA recA Y mrsA rplK Y mrsA rplW Y mrsA rplW Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y mtlD b1458 mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD ybbP mukB mukB Y mukB acpP	mreB	tufB	Υ	
mreB yjgD Y mreB yjgA Y mreB yleA Y mrr mrr Y Y mrsA mrsA Y Y mrsA nusG Y mrsA recA Y mrsA rplK Y mrsA rplW Y mrsA rplV Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA rpsB Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y mtlD b1458 mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD ybbP mukB mukB Y mukB acpP	mreB	ybdQ		Υ
mreB yleA Y mrr mrr Y Y mrsA mrsA Y mrsA murC Y mrsA nusG Y mrsA recA Y mrsA rplK Y mrsA rplL Y mrsA rplW Y mrsA rpsB Y mrsA rpsE Y mrsA rpsJ Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD dppA Y mukB mukB Y mukB acpP	mreB	-		Υ
mrr mrr Y Y mrsA mrsA Y mrsA murC Y mrsA nusG Y mrsA recA Y mrsA rplK Y mrsA rplL Y mrsA rplW Y mrsA rpsB Y mrsA rpsB Y mrsA rpsE Y mrsA rpsJ Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dpbP Y mukB mukB Y mukB acpP Y	mreB	yjgD		Υ
mrsAmrsAYYmrsAmurCYmrsArusGYmrsArecAYmrsArplKYmrsArplLYmrsArplMYmrsArpsBYmrsArpsEYmrsArpsJYmrsAtufAYmrsAtufBYmrsAyggBYmtlDmtlDYmtlDdppAYmtlDdppAYmtlDybbPYmukBmukBYYmukBmukBYYmukBacpPY	mreB		Υ	
mrsA murC mrsA nusG mrsA recA y mrsA rplK y mrsA rplL y mrsA rplW mrsA rplV y mrsA rpsB y mrsA rpsE y mrsA rpsJ y mrsA tufA y mrsA tufB y mrsA yggB y mtlD mtlD y mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD ybbP mukB mukB y mukB acpP	mrr	mrr	Y	Y
mrsA nusG mrsA recA mrsA rplK mrsA rplL mrsA rplL mrsA rplW mrsA rplV mrsA rpsB mrsA rpsE mrsA rpsE mrsA tufA mrsA tufB mrsA yggB mrsA yggB mtlD mtlD y mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD dppA mtlD ybbP mukB mukB y mukB acpP	mrsA	mrsA	Υ	Υ
mrsA recA Y mrsA rplK Y mrsA rplL Y mrsA rplM Y mrsA rplV Y mrsA rpsB Y mrsA rpsE Y mrsA rpsJ Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD b1458 Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD dppA Y mtlD ybbP Y mukB mukB Y mukB acpP Y	mrsA	murC	Υ	
mrsA rplK mrsA rplL mrsA rplM mrsA rplW mrsA rplV mrsA rpsB mrsA rpsE mrsA rpsJ mrsA tufA mrsA tufB mrsA yggB mrsA yggB mtlD mtlD mtlD ybbP mukB mukB y mrsA y mrsB y mrsA ygbP y mrtD ybbP y mukB acpP	mrsA	nusG		Υ
mrsA rplL mrsA rplM mrsA rplV mrsA rplV mrsA rpsB mrsA rpsE mrsA rpsJ mrsA tufA mrsA tufB mrsA yggB mrtlD mtlD mtlD y mtlD dppA mtlD dppA mtlD entB mtlD ybbP mukB mukB y mrsA rpsJ Y mrsA yggB Y mtlD dppA Y mtlD dppA T mtlD dppA T mtlD ybbP T mukB acpP Y T T T T T T T T T T T T T T T T T T	mrsA	recA	Υ	
mrsA rpIM mrsA rpIV mrsA rpsB y mrsA rpsE y mrsA rpsJ y mrsA tufA y mrsA tufB y mrsA yggB y mtID mtID y mtID b1458 y mtID dppA y mtID entB y mtID ybbP mukB mukB y mukB acpP	mrsA	rplK		Υ
mrsA rplV Y mrsA rpsB Y mrsA rpsE Y mrsA rpsJ Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD b1458 Y mtlD dppA Y mtlD entB Y mtlD ybbP Y mukB mukB Y y mukB acpP Y	mrsA	rplL		Υ
mrsA rpsB Y mrsA rpsE Y mrsA rpsJ Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD b1458 Y mtlD dppA Y mtlD entB Y mtlD ybbP Y mukB mukB Y Y mukB acpP Y	mrsA	rplM		Υ
mrsA rpsJ Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD b1458 Y mtlD dppA Y mtlD entB Y mtlD ybbP Y mukB mukB Y Y mukB acpP Y		rplV		
mrsA rpsJ Y mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD b1458 Y mtlD dppA Y mtlD entB Y mtlD ybbP Y mukB mukB Y Y mukB acpP Y	mrsA	rpsB		Υ
mrsA tufA Y mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD b1458 Y mtlD dppA Y mtlD entB Y mtlD ybbP Y mukB mukB Y Y mukB acpP Y	mrsA	-		
mrsA tufB Y mrsA yggB Y mtlD mtlD Y Y mtlD b1458 Y mtlD dppA Y mtlD entB Y mtlD ybbP Y mukB mukB Y Y mukB acpP Y		•		Υ
mrsA yggB Y mtlD mtlD Y Y mtlD b1458 Y mtlD dppA Y mtlD entB Y mtlD ybbP Y mukB mukB Y Y mukB acpP Y	mrsA			
mtIDmtIDYYmtIDb1458YmtIDdppAYmtIDentBYmtIDybbPYmukBmukBYYmukBacpPY	mrsA	tufB	Υ	
mtIDb1458YmtIDdppAYmtIDentBYmtIDybbPYmukBmukBYYmukBacpPY	mrsA	yggB		
mtID dppA Y mtID entB Y mtID ybbP Y mukB mukB Y Y mukB acpP Y			Y	
mtID entB Y mtID ybbP Y mukB mukB Y Y mukB acpP Y				
mtlDybbPYmukBmukBYYmukBacpPY				
mukBmukBYYmukBacpPY				
mukB acpP Y		•		
-			Υ	
mukB mukE Y Y		-		
	mukB	mukE	Υ	Υ

		37	V
mukB	mukF	Y	Y
mukB	parC	Y	
mukB	rplB	Y	
mukB	rpIC	Y	
mukB	rpID	Υ	V
mukB	rplL	Υ	Y Y
mukB mukB	rpIM	Ť	Ϋ́
mukB	rpIR rpIV		Ϋ́
mukB	rpIV rpmB		Ϋ́
mukB	•	Υ	Y
mukB	rpsB rpsC	Ϋ́	ĭ
mukB	-	ī	Y
mukB	rpsE rpsG		Y
mukB	rpsJ	Υ	Y
mukB	rpsP	ī	Y
mukB	tufA	Υ	Y
mukB	tufB	Ϋ́	Į.
mukE	mukE	Y	Υ
mukE	mukB	Ϋ́	Ϋ́
mukE	mukF	Ϋ́	Y
mukF	mukF	Ϋ́	Y
murA	murA	Y	Y
murA	b2060	,	Ϋ́
murA	degQ		Ϋ́
murA	ftsl	Υ	•
murA	hepA	•	Υ
murA	рерВ		Ϋ́
murA	prsA	Υ	•
murA	rpID	•	Υ
murA	rplK		Ϋ́
murA	rpIL		Ϋ́
murA	rplM		Ϋ́
murA	rplV		Ϋ́
murA	rpsB		Ϋ́
murA	rpsJ		Ϋ́
murA	tufA	Υ	•
murA	tufB	Ϋ́	
murA	yhgF	-	Υ
murB	murB	Υ	Y
murB	emrY		Y
murB	ydaY		Y
murB	ygjL		Y
murC	murC	Y	Υ
i,			

murC	b1592		Υ
murC	dnaK	Υ	
murC	lipA		Υ
murC	murB	Υ	-
murC	nusG	-	Υ
murC	rplL		Ϋ́
murC	rpsB		Ϋ́
murC	tufA	Υ	Ϋ́
murC	tufB	Ϋ́	•
murC	yfcC	•	Υ
murC	yiaH		Ϋ́
murC	ytfT		Ϋ́
murD	murD	Y	Ϋ́
murD	yffG	,	Ϋ́
murD	yrfB		Ϋ́
murE	murE	Υ	Y
murE	carA	Ϋ́	,
murE	hyfB	•	Υ
murE	katG		Ϋ́
murE			Ϋ́
murE	pepP		Y
	pnp		Υ
murE	rpsB		Ϋ́
murE	yaeL		
murE	ydaY	V	Y
murF	murF	Y	Y
murF	b1085		Y
murF	gltD		Y
murF	mrdB		Y
murF	mukB		Y
murF	yaaF		Y
murF	yedV		Y
murG	murG	Y	
murG	fimD		Y
murG	hupA		Υ
murG	hyfB		Υ
murG	murE		Υ
murG	rplK		Υ
murG	rpIL		Υ
murG	rplM		Υ
murG	rplS		Υ
murG	rplX		Υ
murG	rpmG		Υ
murG	rpsB		Υ
murG	ydbA_1		Υ

murG	yfiF		Υ
murG	yphH		Υ
murl	murl	Y	Y
murl	nusG		Υ
murl	rplK		Υ
murl	rpIL		Υ
murl	rpsB		Υ
murl	tufA	Υ	
murl	tufB	Υ	
mutL	mutL	Υ	Y
mutL	dnaK		Υ
mutL	rpsA		Υ
mutL	tufB	Υ	
mutS	mutS	Y	Y
mutS	tufA	Y	
mutT	mutT	Y	
mutY	mutY	Y	
mutY	dnaK	•	Υ
mutY	fusA		Ϋ́
nadE	nadE	Y	Y
nadE	b2255	Ϋ́	•
nadE	ygfH	Ϋ́	
napA	napA	Y	Υ
napA	aceE	Ϋ́	•
napA	lpdA	Ϋ́	
napA	panC	•	Υ
napA	rpmB		Ϋ́
napA	rpsN		Ϋ́
napA	ushA		Ϋ́
napA	yacL		Ϋ́
napD	napD		Ϋ́
napD	aceF		Ϋ́
napD	mopA		Ϋ́
napD	pta	Υ	Ϋ́
napD	rplW	•	Ϋ́
napD	yajQ		Ϋ́
narG	narG	Υ	Y
narG	dnaJ	Ý	,
narG	mreB	Ý	
narG	narJ	Ý	Υ
narG	narY	Ý	•
narG	recA	Ý	
narG	rfaD	Ϋ́	
narG	tufA	Y	
Haid	luiA	ı	

	4 4 D	V	
narG	tufB	Y	V
narH	narH	Y	Y
narH	narG	Y	Υ
narH	tufA	Y	
narH	tufB	Υ	V
narH	ygjL	V	Y
narJ	narJ	Υ	Y
narJ	aceF		Υ
narJ	b1806	Y	
narJ	lpdA	3.7	Y
narJ	narG	Y	Y
narJ	narH	Υ	Y
narJ	narY		Υ
narJ	ompA	Y	
narJ	pepP	Υ	
narJ	rplL		Υ
narW	narW	Υ	Υ
narW	narG	Υ	Υ
narW	narH	Υ	
narW	narY		Υ
narY	narY	Υ	Υ
narY	narG	Υ	
narY	narH		Υ
narY	pepN	Υ	
narY	ygjL		Υ
narZ	narZ	Υ	Υ
narZ	narH	Υ	Υ
ndk	ndk	Υ	Y
ndk	dnaK	Υ	
ndk	dnaN	Υ	
ndk	ygfA	Υ	
nei	nei	Υ	Y
nei	hupB		Υ
nei	lpdA		Υ
nei	mopA		Υ
nei	nfi		Υ
nei	rpID		Υ
nei	rpIU		Υ
nei	rpmB		Υ
nei	rpsD		Υ
nei	rpsN		Υ
nei	slyD		Υ
nfi	nfi	Υ	Υ
nfi	aceF	Υ	Υ

nfi	himA		Υ
nfi	himD		Υ
nfi	hns		Υ
nfi	hupA		Υ
nfi	hupB		Υ
nfi	lpdA		Υ
nfi	malP		Υ
nfi	mopA		Υ
nfi	nei	Υ	
nfi	rplC		Υ
nfi	rplD		Υ
nfi	rplL		Υ
nfi	rpIS		Υ
nfi	rpmB		Υ
nfi	rpoA		Υ
nfi	rpsE		Υ
nfi	rpsF		Υ
nfi	rpsG		Υ
nfi	rpsM		Υ
nfi	rpsN		Υ
nfi	rpsO		Υ
nfi	rpsR		Υ
nfi	ruvA		Υ
nfi	yaiD		Υ
nfo	nfo	Y	
nfo	aceF	Υ	
nfo	lon	Υ	
nfo	lpdA	Y Y	
nfo	malP	Υ	
nfo	secA	Υ	
nikA	nikA	Y	Y
nikA	tufA		Υ
nikD	nikD	Y	
nikD	aceE	Υ	
nikD	dnaK	Υ	
nikD	mopA	Υ	
nikD	tufA	Υ	
nikE	nikE	Y	
nikE	rplB	Υ	
nikE	rpsC	Υ	
nikE	rpsD	Υ	
nikE	rpsG	Υ	
nikE	secA	Υ	
nikE	srmB	Υ	

nikE	tufB	Y	
nrdA	nrdA	Y	Y
nrdA	mreB	Y	
nrdA	proA	Y	
nrdA	tufA	Y	
nrdA	tufB	Y	
nrdB	nrdB	Y	Y
nrdB	dnaK		Υ
nrdB	dnaN	Υ	
nrdB	mreB	Υ	
nrdB	rhsB	Υ	
nrdB	tufA	Υ	
nrdB	tufB	Υ	
nth	nth	Y	Y
nth	aceE	Υ	Υ
nth	lpdA		Υ
nth	rplW		Υ
nth	tufA		Υ
ntpA	ntpA	Υ	Y
ntpA	aceE		Υ
ntpA	dnaK		Υ
ntpA	lpdA		Υ
nusA	nusA	Υ	Y
nusA	aceE		Υ
nusA	infB	Υ	Υ
nusA	metJ		Υ
nusA	prsA	Υ	
nusA	pstS		Υ
nusA	rplA	Υ	
nusA	rplB	Υ	
nusA	rpIC	Υ	
nusA	rpID	Υ	
nusA	rplO	Υ	
nusA	rpIS		Υ
nusA	rpIV	Υ	Υ
nusA	rplX		Υ
nusA	rpoA	Υ	Υ
nusA	rpoB	Υ	Y
nusA	rpoC	Υ	Υ
nusA	rpoD		Υ
nusA	rpoH	Υ	
nusA	rpoZ		Υ
nusA	rpsC	Υ	
nusA	rpsD	Y	

nusA	rpsE		Υ
nusA	rpsG	Υ	
nusA	tufA	Υ	
nusA	tufB	Υ	
nusA	uspA		Υ
nusA	uvrA		Υ
nusB	nusB	Υ	
nusB	rpIC	Υ	
nusB	rplM	Υ	
nusB	rpsB	Υ	
nusB	rpsD	Υ	
nusB	rpsE	Υ	
nusB	rpsJ	Υ	
nusG	nusG	Υ	Y
nusG	aefA		Υ
nusG	b0878	Υ	
nusG	b2255	Υ	
nusG	deaD	Υ	
nusG	dnaJ	Υ	
nusG	hepA	Υ	
nusG	mreB	Υ	
nusG	pssA	Υ	
nusG	rho	Υ	Υ
nusG	rplB	Υ	
nusG	rpID		Υ
nusG	rplE	Υ	
nusG	rplL		Υ
nusG	rplV		Υ
nusG	rplW		Υ
nusG	rpmB		Υ
nusG	rpoA	Υ	Υ
nusG	rpoB	Υ	
nusG	rpoC	Υ	
nusG	rpsB		Υ
nusG	rpsC	Υ	
nusG	rpsE	Υ	Υ
nusG	rpsF		Υ
nusG	rpsG		Υ
nusG	rpsJ	Υ	Υ
nusG	rpsN		Υ
nusG	rpsP		Υ
nusG	secA	Υ	
nusG	sgbH	Υ	
nusG	yacL	Υ	Υ

	4		W
ogt	ogt		Y
ogt	aceE	V	Y
ogt	aceF	Y	V
ogt	gapA	V	Y
ogt	lpdA	Υ	Y Y
ogt	mopA	Υ	Y
panC panC	panC dnaK	Y	I
panC	panB	Y	
panC	purC	'	Υ
panC	ushA	Υ	Ÿ
parC	parC	Y	Y Y
parC	aceF	Ϋ́	•
parC	gyrA	Ý	
parC	lpdA	Ϋ́	
parC	mreB	Y	
parC	murF	·	Υ
parC	rfaQ		Y
parC	rplL		Y
parC	rplM		Υ
parC	rplV		Υ
parC	rpmA		Υ
parC	rpmG		Υ
parC	rpsB		Υ
parC	rpsJ		Υ
parC	secA	Υ	
parC	slyD		Υ
parE	parE	Υ	Υ
parE	b1200		Υ
parE	b1410	Υ	
parE	b2737		Υ
parE	csgD	Υ	
parE	fucO		Υ
parE	parC		Y
parE	rpID		Y
parE	rpsB		Υ
parE	secA	Y	
parE	ssb	Υ	V
parE	yacE		Y
parE	yebG	V	Υ
parE	yhcC	Y	V
parE	yidK		Y Y
parE	yqcE	V	Ϋ́Υ
pepB	pepB	Υ	Y

pepB pepD pepD pepD pepD pepE pepE pepN pepN pepN pepP pepP pepP pepT pepT	fabH ompA pepD dnaK guaC pepP pepE pepB pepN hyfG mopA pepP guaC pepQ pepT dps fusA	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y
pepT pepT	pepE pfIB	Υ	Y Y
рерТ	proA		Υ
рерТ	tpiA	Υ	
рерТ	tufA	Υ	
рерТ	tufB	Υ	
рерТ	wcaL	Υ	
рерТ	yihl		Υ
pfkA	pfkA	Υ	
pflB	pflB	Υ	Υ
pflB	pykA		Υ
pfIB	tpiA	Y	V
pfIB	yfiD zwf	Υ	Y Y
pflB <i>pgk</i>	pgk	Υ	ľ
pgk	dnaK	,	Υ
pgk	eno		
pgk	hfq		Y Y Y
pgk	pnp		
pgk	rne		Υ
pgk	rpll		Y Y
pgk	rpsF		
pgk	rpsN		Y
pgk	rpsT	17	Y
pheS	pheS	Y	Y Y
pheS	brnQ erfK		Y Y
pheS	CIII		I

h - C	b.a.T	V	
pheS	pheT	Y	V
pheT	pheT	Y	Y
pheT	mfd	V	Y
pheT	pheS	Υ	Y
pheT	secD		Y
pheT	yeaJ		Y
pheT	yghQ		Y
pheT	yjcQ	V	Υ
phoB	phoB	Y	
phrB	phrB :	Υ	
phrB	nfi . D		Υ
plsB	plsB	Y	
plsB	lpdA	Y	
plsB	rho	Υ	
plsB	rplB		Y
plsB	rplC	Y	Υ
plsB	rplD	Υ	
plsB	rplF		Y
plsB	rpll		Y
plsB	rplK		Υ
plsB	rplL		Υ
plsB	rplM		Υ
plsB	rplO		Υ
plsB	rpIS		Υ
plsB	rpIV		Υ
plsB	rplX		Υ
plsB	rpmB		Υ
plsB	rpmG		Υ
plsB	rpsB		Υ
plsB	rpsC	Y	Y
plsB	rpsD	Υ	Υ
plsB	rpsE		Υ
plsB	rpsH		Υ
plsB	rpsl		Υ
plsB	rpsJ		Υ
plsB	srmB	Υ	
plsB	tufA	Υ	
plsB	tufB	Υ	
plsB	yciL		Υ
pncB	pncB	Υ	Y
pncB	rpIS		Υ
pncB	rplV		Υ
pncB	rpsB		Υ
pncB	tufA		Υ

pncB	yeiM		Y
pnp	pnp	Y	Y
pnp	eno	Υ	Y
pnp	hupA		Υ
pnp	metK	Y	
pnp	mreB	Υ	
pnp	pgk		Y
pnp	recA	Y	
pnp	rhlB	Υ	Y
pnp	rne	Y	Y
pnp	rplA		Υ
pnp	rplB	Y	
pnp	rplD	Y	Υ
pnp	rplE	Υ	
pnp	rplF	Y	
pnp	rpll	Y	Y
pnp	rplM	Υ	Y
pnp	rplO		Y
pnp	rplQ		Υ
pnp	rplU		Υ
pnp	rplV		Υ
pnp	rplX		Υ
pnp	rpoA	Υ	
pnp	rpsC	Υ	
pnp	rpsD	Υ	
pnp	rpsE	Y	Y
pnp	rpsG	Υ	Υ
pnp	rpsM	Υ	Υ
pnp	rpsO		Υ
pnp	rpsR		Υ
pnp	rpsS		Υ
pnp	rpsT		Υ
pnp	srmB	Υ	
pnp	yciL	Υ	
polA	polA	Y	Υ
polA	aceF	Υ	Υ
polA	b2324		Υ
polA	fis		Υ
polA	hns		Υ
polA	hrpA		Υ
polA	hsdS		Υ
polA	hupA		Y
polA	hupB		Y
polA	lpdA	Y	Υ

polA	malP	Υ	Υ
polA	metK	Υ	
polA	murF		Υ
polA	rplD		Υ
polA	rpIL		Υ
polA	rplM		Υ
polA	rpmB		Υ
polA	rpoA	Υ	Υ
polA	rpoB	Υ	
polA	rpoC	Υ	
polA	rpsB		Υ
polA	slyD		Υ
polA	tufA	Υ	
polA	tufB	Υ	
polA	ydbA_2		Υ
polA	yidX		
рра	рра	Y	Y Y
ppa	atoD		Υ
ppa	dnaK	Υ	
ppa	dnaN	Υ	
ppa	katG		Υ
ppa	mopA	Υ	
ppa	rfaD	Υ	
ppa	rpIC		Υ
ppa	rplD	Υ	Υ
ppa	rpll		Υ
ppa	rplM		Υ
ppa	rplN		Υ
ppa	rpIS		Υ
ppa	rpIV		Υ
ppa	rpmB		Υ
ppa	rpsB		Υ
ppa	rpsC		Υ
ppa	rpsE		Υ
ppa	rpsG		Υ
ppa	rpsH		Υ
ppa	rpsJ		Υ
ppa	rpsN		Υ
ppa	rpsT		Υ
ppa	secB		Υ
ppa	slyD		Υ
ppa	tig	Υ	Y
ppa	tufA	Υ	
ppa	tufB	Υ	

nna	vha∩		Y
ppa ppa	ybgQ yhhU		Y
ppa	yjcG		Ϋ́
ppa	yjgD		Ϋ́
ppiB	ppiB	Υ	Y
ppiB	fucK		Υ
ppiB	hflB		Υ
ppiB	rhsC		Υ
ppiB	yhjU		Υ
ppiC	ppiC	Y	Y
ppiC	accA		Y
ppiC	accD		Y
ppiC	rpID		Y
ppiC	rplK		Y Y
ppiC ppiC	rplL rplM		Ϋ́
ppiC	rplS		Y
ppiC	rpIV		Ϋ́
ppiC	rplW		Ϋ́
ppiC	rpmC		Y
ppiC	rpsG		Υ
ppiC	yehX		Υ
ppiC	ygaF		Υ
ppk	ppk	Y	
ppk	aceF	Υ	
ppk	lpdA	Y	
prfA	prfA	Υ	Y
prfA	b1327		Y
prfA prfA	basS frdC		Y Y
prfA	hyfR		Y
prfA	pheA		Ϋ́
prfA	rfaK		Ϋ́
prfA	yghE		Y
prfA	yhgA		Υ
prfB	prfB	Y	Y
prfB	b1587		Υ
prfB	b2146		Υ
prfB	ccmB		Y
prfB	hupA		Y
prfB	hupB		Y
prfB	parE		Y Y
prfB prfB	polA rfe		Y Y
prfB	пe		Ť

4D	المسا		V
prfB	rplL		Y
prfB	rpIV		Y
prfB	rpoA		Y Y
prfB	rpsB		Ϋ́
prfB prfB	rpsE		Ϋ́
prfB	rpsJ		Ϋ́
prfB	sgaT ycfX		Ϋ́
prfB	ygjD		Ϋ́
prfB	yjcF		Ϋ́
prfC	prfC	Υ	Y
prfC	gyrB	,	Ϋ́
prfC	pfkA	Υ	•
prfC	smpB	•	Υ
prfC	tgt		Ϋ́
priA	priA	Y	•
priA	tufB	Y Y	
prmA	prmA	Y	Y
prmA	malS	•	Ϋ́
prmA	msbA		Y
prmA	rplK	Υ	Ϋ́
prmA	ydaY		Υ
prmA	yegB		Υ
prmA	yejO		Υ
proA	proA	Y	Y
proA	aceE	Υ	
proA	b2981		Υ
proA	cysA		Υ
proA	dnaN	Υ	
proA	lpxD		Υ
proA	proS		Υ
proA	sfsA		Υ
proA	ugpB		Υ
proA	yehT		Υ
proS	proS	Y	Y
proS	aceE	Υ	
proS	dnaN	Υ	
prsA	prsA	Y	Y
prsA	cobB	Y	Y
prsA	dnaJ	Y	Υ
prsA	dnaK	Y	
prsA	mopA	Y	Υ
prsA	mreB	Υ	
prsA	nusG		Υ

prsA	rho		Υ
prsA	rplB	Υ	Υ
prsA	rplC		Υ
prsA	rpID		Υ
prsA	rpIE	Υ	
prsA	rpll		Υ
prsA	rplJ	Υ	
prsA	rplK		Υ
prsA	rplL		Y Y
prsA	rplM		Υ
prsA	rplN		Υ
prsA	rplO		Υ
prsA	rpIP	Υ	Υ
prsA	rpIS		Υ
prsA	rplU		Υ
prsA	rpIV		Υ
prsA	rplW		Υ
prsA	rplX		Υ
prsA	rpmB		Υ
prsA	rpsB	Υ	Υ
prsA	rpsC	Y Y	Υ
prsA	rpsD	Υ	Υ
prsA	rpsE		Υ
prsA	rpsG	Υ	Υ
prsA	rpsH		Υ
prsA	rpsJ		Υ
prsA	rpsP		Υ
prsA	rpsT		Υ
prsA	tufA	Υ	Υ
prsA	tufB	Υ	
prsA	yciL	Υ	
prsA	yfiF		Υ
prsA	yibA	Υ	
prsA	yrfD		Υ
pssA	pssA	Υ	
pssA	pnp	Υ	
pssA	rpIC	Υ	
pssA	rpIL		Υ
pssA	rpIU		Υ
pssA	rplW		Υ
pssA	rpsA	Υ	
pssA	rpsB	Υ	
pssA	rpsC	Υ	
pssA	rpsD	Υ	

pssA pssA pssA pssA pssA pssB pstB	rpsE rpsO rpsN rpsU ybaU ycbY yfiF pstB adhC	Y Y Y	Y Y Y Y
pstB pstB	cfa dnaJ		Y Y
pstB	gatY	Υ	
pstB	mopA	Y	Y
pstB	mreB	Y	
pstB pstB	narG rhsC	Y Y	
pstB	tufA	Ϋ́	
pstB	tufB	Ϋ́	
pstB	ycfS	'	Υ
pta	pta	Υ	Ϋ́
pta	chpS	,	Ý
pta	dnaJ	Υ	-
pta	tufA	Y	
pta	tufB	Υ	
pth	pth	Υ	Y
pth	ccmB		Υ
pth	lytB		Υ
pth	mopA		Υ
pth	rpID		Υ
pth	rplK		Υ
pth	rplL		Y
pth	rplM		Y
pth	rplV		Y
pth	rpmB		Y Y
pth pth	rpmG rpsB	Υ	Ϋ́
pth	rpsb	ī	Y
pth	tufA	Υ	•
pth	tufB	Ϋ́	
pth	yljA	Y	
purB	purB	Y Y	Υ
purB	aceE	Υ	
purB	rpIV		Υ
purB	rpsB		Υ

purB ydaY purC purC Y y purC cca y purC dnaK Y purC evgA purC guaC purC lhr purC rplL purC tig Y purC yidZ pykA pykA Y pykA dnaK pykA dnaK pykA dnaN Y pykA malX pykA tig Y pyrH pyrH Y pyrH b1834 pyrH cadB pyrH clpA pyrH dnaJ pyrH dnaK pyrH gatA pyrH gatA pyrH gatA pyrH gatZ pyrH hns Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
purC cca purC dnaK purC evgA purC evgA purC guaC purC lhr purC rplL purC tig Y purC yidZ pykA pykA pykA dnaK pykA dnaK pykA dnaN pykA lnt pykA malX pykA tig Y pyrH pyrH pyrH pyrH pyrH cadB pyrH clpA pyrH dnaJ pyrH dnaJ pyrH dnaJ pyrH gatA pyrH gatA pyrH gatZ pyrH gatZ pyrH dnaS pyrH gatZ pyrH dnaS pyrH gatZ pyrH gatZ pyrH hns
purC cca purC dnaK Y purC evgA purC guaC purC lhr purC rplL purC tig Y purC yidZ pykA pykA Y pykA dnaK pykA dnaK pykA lnt pykA malX pykA tig Y pyrH pyrH Y pyrH cadB pyrH clpA pyrH dnaJ pyrH dnaJ pyrH gatA pyrH gatA pyrH gatZ pyrH gatZ pyrH pyrH gatZ pyrH dnaS pyrH gatZ pyrH gatZ pyrH dnaS pyrH gatZ pyrH dnaS pyrH gatZ pyrH dnaS pyrH gatZ pyrH dnaS
purCdnaKYYpurCevgAYpurCguaCYpurClhrYpurCrplLYpurCtigYYpurCyidZYpykApykAYYpykAdnaKYYpykAlntYYpykAtigYYpykAyggRYYpyrHpyrHYYpyrHcadBYYpyrHdnaJYYpyrHdnaJYYpyrHgatAYYpyrHgatBYYpyrHgatBYYpyrHhnsYY
purC evgA Y purC guaC purC lhr Y purC rplL Y purC tig Y Y purC yidZ Y pykA pykA Y Y pykA dnaK Y pykA dnaN Y pykA malX Y pykA tig Y pykA yggR Y pyrH pyrH Y pyrH b1834 pyrH cadB pyrH clpA pyrH dnaJ pyrH dnaJ pyrH dnaK pyrH gatA pyrH gatA pyrH gatZ pyrH hns Y
purC guaC purC lhr purC rplL purC tig Y purC yidZ pykA pykA Y pykA dnaK pykA dnaN y pykA lnt pykA malX ypkA tig Y pykA yggR pyrH pyrH pyrH b1834 pyrH cadB pyrH clpA pyrH dnaJ pyrH dnaJ pyrH dnaK pyrH gatA pyrH gatA pyrH gatZ pyrH hns Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
purC Ihr Y purC rplL Y purC tig Y Y purC yidZ Y pykA pykA Y Y pykA dnaK Y pykA dnaK Y pykA lnt Y pykA malX Y pykA tig Y Y pykA yggR Y pyrH pyrH Y Y pyrH b1834 Y pyrH cadB Y pyrH clpA Y pyrH dnaJ Y pyrH dnaJ Y pyrH dnaK Y pyrH gatA Y pyrH gatB pyrH gatZ Y pyrH hns Y
purC tig Y Y Y PurC yidZ Y PykA pykA pykA Y Y Y PykA dnaK Y PykA dnaN Y PykA Int Y PykA malX Y PykA tig Y Y Y PyrH b1834 Y PyrH cadB PyrH clpA Y PyrH dnaX PyrH dnaX PyrH dnaX PyrH dnaX PyrH dnaX PyrH gatA Y PyrH gatB PyrH gatZ Y PyrH hns Y
purC tig Y Y purC yidZ Y pykA pykA Y Y pykA dnaK Y pykA dnaN Y pykA lnt Y pykA malX Y pykA tig Y Y pykA yggR Y pyrH pyrH Y Y pyrH b1834 Y pyrH cadB Y pyrH clpA Y pyrH dnaJ Y pyrH dnaJ Y pyrH dnaK Y pyrH gatA Y pyrH gatB pyrH gatZ Y pyrH hns Y
purC yidZ Y pykA pykA Y Y pykA dnaK Y Y pykA lnt Y Y pykA malX Y Y pykA tig Y Y pykA yggR Y Y pyrH pyrH Y Y pyrH pyrH Y Y pyrH dnaX Y pyrH gatA Y pyrH gatB Y pyrH gatZ Y pyrH hns Y
pykApykAYYpykAdnaKYpykAlntYpykAmalXYpykAtigYYpykAyggRYpyrHpyrHYYpyrHb1834YpyrHcadBYpyrHclpAYpyrHdnaJYpyrHdnaKYpyrHgatAYpyrHgatBYpyrHgatZYpyrHhnsY
pykA dnaK pykA dnaN pykA lnt pykA malX pykA tig pykA yggR pyrH pyrH pyrH b1834 pyrH cadB pyrH clpA pyrH dnaJ pyrH dnaJ pyrH dnaK pyrH gatA pyrH gatZ pyrH hns Y
pykA dnaN Y pykA Int Y pykA malX Y pykA tig Y pykA yggR Y pyrH pyrH Y pyrH b1834 Y pyrH cadB Y pyrH clpA Y pyrH dnaJ Y pyrH dnaK Y pyrH fusA Y pyrH gatA Y pyrH gatZ Y pyrH hns Y
pykA Int Y pykA tig Y Y pykA yggR Y Y pyrH pyrH Y Y pyrH b1834 Y Y pyrH cadB Y Y pyrH clpA Y Y pyrH dnaJ Y Y pyrH fusA Y Y pyrH gatA Y Y pyrH gatB Y Y pyrH hns Y Y
pykA tig Y Y pykA tig Y Y pykA yggR Y pyrH pyrH Y Y pyrH b1834 Y Y pyrH clpA Y Y pyrH dnaJ Y Y pyrH dnaK Y Y pyrH gatA Y Y pyrH gatB Y Y pyrH gatZ Y Y pyrH hns Y Y
pykA tig Y Y pykA yggR Y pyrH pyrH Y Y pyrH b1834 Y pyrH cadB Y pyrH clpA Y pyrH dnaJ Y pyrH fusA Y pyrH gatA Y pyrH gatB Y pyrH hns Y
pykA yggR Y pyrH pyrH Y pyrH b1834 Y pyrH cadB Y pyrH clpA Y pyrH dnaJ Y pyrH fusA Y pyrH gatA Y pyrH gatB Y pyrH gatZ Y pyrH hns Y
pyrH pyrH Y Y pyrH b1834 Y pyrH cadB Y pyrH clpA Y pyrH dnaJ Y pyrH fusA Y pyrH gatA Y pyrH gatB Y pyrH gatZ Y pyrH hns Y
pyrH b1834 Y pyrH cadB Y pyrH clpA Y pyrH dnaJ Y pyrH dnaK Y pyrH fusA Y pyrH gatA Y pyrH gatB Y pyrH gatZ Y pyrH hns Y
pyrH cadB Y pyrH clpA Y pyrH dnaJ Y pyrH dnaK Y pyrH fusA Y pyrH gatA Y pyrH gatB Y pyrH gatZ Y pyrH hns Y
pyrH clpA Y pyrH dnaJ Y pyrH dnaK Y pyrH fusA Y pyrH gatA Y pyrH gatB Y pyrH gatZ Y pyrH hns Y
pyrHdnaJYpyrHdnaKYpyrHfusAYpyrHgatAYpyrHgatBYpyrHgatZYpyrHhnsY
pyrH dnaK Y pyrH fusA Y pyrH gatA Y pyrH gatB Y pyrH gatZ Y pyrH hns Y
pyrH gatA Y pyrH gatB Y pyrH gatZ Y pyrH hns Y
pyrH gatB Y pyrH gatZ Y pyrH hns Y
pyrH gatZ Y pyrH hns Y
pyrH hns Y
pyrH htrA Y
pyrH hupA Y
pyrH hupB Y
pyrH lpxC Y
pyrH mopA Y
pyrH recA Y
pyrH rho Y
pyrH rplA Y
pyrH rplB Y
pyrH rplC Y
pyrH rplD Y
pyrH rplE Y
pyrH rpll Y
pyrH rplK Y pyrH rplL Y
pyrH rplM Y

pyrH	rplR		Y
pyrH	rpIS		Y
pyrH	rpIU		Y
pyrH	rpIV		Y
pyrH	rplW		Y
pyrH	rpmA		Y Y
pyrH	rpmG		Ϋ́
pyrH	rpsA		Ϋ́
pyrH	rpsB		Ϋ́
pyrH	rpsC		Ϋ́
pyrH	rpsE rpsG		Ϋ́
pyrH	rpsJ		Υ
pyrH pyrH	rpsP		Ϋ́
pyrH	secA		Ϋ́
pyrH	slyD		Ϋ́
pyrH	thiE		Ϋ́
pyrH	tufA		Ϋ́
pyrH	uspA		Ϋ́
pyrH	ybdQ		Ϋ́
pyrH	yfiF		Ϋ́
pyrH	yhbY		Ϋ́
qor	qor	Υ	Y
qor	dnaK	Ϋ́	,
rbfA	rbfA	Ϋ́	Y
rcsA	rcsA	•	Ϋ́
rcsA	add		Ý
rcsA	cadA		Ϋ́
rcsA	dnaK		Y
rcsA	hupA		Y
rcsA	hycC		Υ
rcsA	rplB		Υ
rcsA	rpIC		Υ
rcsA	rpID		Υ
rcsA	rplE		Υ
rcsA	rplM		Υ
rcsA	rplP		Υ
rcsA	rpIS		Υ
rcsA	rplT		Υ
rcsA	rpIU		Υ
rcsA	rpIV		Υ
rcsA	rplW		Υ
rcsA	rpmC		Υ
rcsA	rpmG		Υ

rcsA	rpsB		Υ
rcsA	rpsF		Υ
rcsA	rpsG		Υ
rcsA	rpsM		Υ
rcsA	rpsN		Υ
rcsB	rcsB	Y	Y
rcsB	b2247		Υ
rcsB	b2249		Υ
rcsB	b2878		Υ
rcsB	crp		Υ
rcsB	dnaK	Υ	
rcsB	glnB	Υ	
rcsB	IIdP		Υ
rcsB	mesJ		Υ
rcsB	mgtA		Υ
rcsB	polA		Υ
rcsB	prmA		Υ
rcsB	rplC	Υ	
rcsB	rplE		Υ
rcsB	rpll		Υ
rcsB	rplM	Υ	Υ
rcsB	rpIS		Υ
rcsB	rpIV		Υ
rcsB	rpmB		Υ
rcsB	rpoA		Υ
rcsB	rpsC	Υ	Υ
rcsB	rpsD	Υ	
rcsB	rpsJ	Υ	
rcsB	rpsU	Υ	
rcsB	rspJ	Υ	
rcsB	tufB	Υ	
rcsB	yagA		Υ
rcsB	ycbN		Υ
rcsB	yccC		Y
rcsB	ydbA_2		Y
rcsB	yhiE		Y
rcsB	yhjQ		Y
rcsB	ymcA	\	Υ
rcsB	yojN	Y	
recB	recB	Y	
recB	aceE	Y	
recB	aceF	Y	
recB	lpdA	Y	
recB	recC	Υ	

	_		
recB	recD	Υ	
recB	rpsB	Υ	
recB	tufB	Υ	
recD	recD	Y	
recD	aceE	Υ	
recD	lpdA	Υ	
recD	mopA	Υ	
recD	recB	Υ	
recD	recC	Υ	
recE	recE	Y	Υ
recE	aceE	Υ	
recE	lig	Υ	
recF	recF	Y	
recF	mopA	Υ	Υ
recF	uidC		Υ
recG	recG	Y	
recG	aceF	Υ	
recJ	recJ	Y	Y
recJ	aceE	Υ	
recJ	aceF	Υ	
recJ	dnaJ	Υ	
recJ	dnaK		Υ
recJ	galR		Υ
recJ	lpdA	Υ	Υ
recJ	mtlD		Υ
recJ	rnhA	Υ	Υ
recJ	rplD		Υ
recJ	rpIV		Υ
recJ	rpmB		Υ
recJ	rpsP		Υ
recJ	ruvA		Υ
recJ	sbcB	Υ	
recJ	ssb	Υ	Υ
recJ	topB	Υ	
recJ	tufA	Υ	
recJ	tufB	Υ	
recJ	yhbY		Υ
recN	recN	Υ	
recN	ydcP	Y	
recQ	recQ	Y	
recQ	aceE	Y	
recQ	aceF	Y	Υ
recQ	lpdA	Y	Y
recQ	rpmB		Ϋ́
	•		

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recQ	rpsN	V	Y
recQ	ssb	Y	Υ
recQ	topA	Y	
rep	rep	Y	
rfaD	rfaD	Y	Y
rfaD	b1484		Υ
rfaD	cysM	Y	
rfaD	dnaK	Y	
rfaD	entF	Υ	
rfaD	rplC		Υ
rfaD	rplO	Υ	
rfaD	rpIS		Y
rfaD	rpsG	Υ	Υ
rfaD	secA		Υ
rfaD	torA		Υ
rfaD	ybjX	Υ	
rfaD	ydaY		Υ
rfaD	yhhN		Υ
rfaD	yhjS		Υ
rfbC	rfbC	Υ	
rfbC	mopA	Υ	
rhIB	rhIB	Y	Y
rhlB	aceE	Υ	
rhlB	dnaK	Υ	
rhIB	eno	Υ	Υ
rhIB	pnp	Υ	Υ
rhlB	rne	Υ	Υ
rhlB	rplA		Υ
rhlB	rplC	Υ	Υ
rhlB	rpll	Υ	
rhlB	rplK	Υ	
rhlB	rplM		Υ
rhlB	rplS		Υ
rhlB	rplU		Υ
rhlB	rplV	Υ	Υ
rhlB	rpmB		Υ
rhlB	rpsC	Υ	
rhlB	rpsD	Υ	
rhlB	rpsE		Υ
rhlB	rpsF		Υ
rhlB	rpsG	Υ	Υ
rhlB	rpsJ	Υ	
rhlB	rpsP		Υ
rhlB	rpsS		Υ
l .	•		

rhlB	rpsT		Υ
rhlB	tufB	Υ	•
			V
rho	rho	Υ	Y
rho	gsp		Υ
rho	menF		Υ
rho	nusG		Υ
rho	rplC		Υ
rho	rpID		Υ
rho	rplL		Υ
rho	rpsB		Y
rho	tufA	Υ	•
rho	tufB	Ϋ́	
		ı	V
rho	yhbY		Y
ribB	ribB	Υ	Y
ribB	arcB		Υ
ribB	atpD		Υ
ribB	b2373		Υ
ribB	ybaL		Υ
ribD	ribD	Υ	Υ
ribD	accB		Υ
ribD	accC		Υ
ribD	accD		Υ
ribD	agaR		Y
ribD	b1586		Ϋ́
ribD	cvpA		Ϋ́
ribD	fruB		Ϋ́
ribD			Ϋ́
	hupA		Ϋ́
ribD	mopA		
ribD	pbpC		Y
ribD	ribH		Y
ribD	rplC		Υ
ribD	rpID		Υ
ribD	rplE		Υ
ribD	rpIS		Υ
ribD	rplU		Υ
ribD	rpsB		Υ
ribD	rpsD		Υ
ribD	rpsG		Υ
ribD	ycil		Ϋ́
ribF	ribF	Υ	-
ribF	aceE	Ϋ́	
ribF	acrD	·	Υ
ribF	recD		Ϋ́
	ribH	Υ	Y
ribH	דוטוו	1	1

ribH	feaB		Υ
ribH	gltJ		Υ
ribH	rfaD	Υ	Υ
ribH	rplC	Υ	Υ
ribH	rplD		Υ
ribH	rplM		Υ
ribH	rplT		Υ
ribH	rplU		Υ
ribH	rpIV		Υ
ribH	rpmG		Υ
ribH	tufA	Υ	
ribH	tufB	Υ	
ribH	yehQ		Υ
rna	rna	Y	
rnb	rnb	Y	
rnb	rpsE	Y	
rnc	rnc	Y	
rnc	rnb	Y	
rnd	rnd	Y	
rne	rne	Y	Y
rne	deaD	Ý	•
rne	dnaK	Ϋ́	Υ
rne	eno	Ϋ́	Ý
rne	fabZ	Ϋ́	-
rne	fruR	Ϋ́	
rne	hfq	•	Υ
rne	pnp	Υ	Y
rne	rhIB	Ϋ́	Ϋ́
rne	rplA	Ϋ́	•
rne	rpIB	Ϋ́	Υ
rne	rpIC	Ϋ́	Ϋ́
rne	rpID	Ϋ́	Ϋ́
rne	rpll	Ϋ́	•
rne	rplK	Ϋ́	Υ
rne	rplL	•	Ϋ́
rne	rplM		Ϋ́
rne	rplU		Ϋ́
rne	rplV	Υ	Ϋ́
rne	rplW	•	Ϋ́
rne	rplX		Ϋ́
rne	rpsB	Υ	•
rne	rpsC	Ϋ́	
rne	rpsD	Ϋ́	Υ
rne	rpsE	•	Y
1116	ips⊏		I

1			
rne	rpsF		Υ
rne	rpsG	Υ	Υ
rne	rpsJ		Υ
rne	rpsP		Υ
rne	rpsT		Υ
rne	rpsU	Υ	
rne	srmB	Υ	
rne	tufA	Υ	
rne	tufB	Υ	
rne	vacB	Υ	
rne	ycbY	Υ	
rne	yciL	Υ	
rne	yfiF		Υ
rne	yfiQ	Υ	
rnhA	rnhA	Y	
rnhA	aceE	Υ	
rnhA	aceF	Υ	Υ
rnhA	hupA		Υ
rnhA	lpdA	Υ	
rnhA	malP	Υ	
rnhA	priA	Υ	
rnhA	recJ	Υ	
rnhA	recQ	Υ	
rnhA	rplB	Υ	
rnhA	rpmB		Υ
rnhA	rpsC	Υ	
rnhA	rpsE		Υ
rnhA	rpsl		Υ
rnhA	rpsN		Υ
rnhA	rpsT		Υ
rnhA	sbcB	Υ	
rnhA	ssb	Υ	Υ
rnhA	topA	Υ	Υ
rnhA	topB	Υ	
rnhA	yhbY		Υ
rnpA	rnpA	Y	Y
rnpA	aceF	Υ	
rnpA	clpA	Υ	
rnpA	deaD	Υ	
rnpA	hupA		Υ
rnpA	lpdA	Υ	
rnpA	ppk	Υ	
rnpA	pssA	Υ	Υ
rnpA	rho	Υ	

1			
rnpA	rplA	Υ	Y
rnpA	rplB	Υ	Υ
rnpA	rpIC	Υ	Y
rnpA	rpID		Y
rnpA	rplF		Y
rnpA	rpll		Υ
rnpA	rplK		Υ
rnpA	rpIL		Υ
rnpA	rplM		Υ
rnpA	rplN		Υ
rnpA	rplO		Υ
rnpA	rpIP		Y
rnpA	rplQ		Y
rnpA	rplR		Υ
rnpA	rpIS		Υ
rnpA	rplT		Υ
rnpA	rplU		Υ
rnpA	rplV		Υ
rnpA	rplX		Υ
rnpA	rpmC		Υ
rnpA	rpsA	Υ	
rnpA	rpsB	Υ	Υ
rnpA	rpsC	Υ	Υ
rnpA	rpsD	Υ	
rnpA	rpsE	Υ	Υ
rnpA	rpsF		Υ
rnpA	rpsG	Υ	Υ
rnpA	rpsH	Υ	Υ
rnpA	rpsl		Y
rnpA	rpsJ		Υ
rnpA	rpsK		Υ
rnpA	rpsM	Υ	Υ
rnpA	rpsN		Υ
rnpA	rpsO		Υ
rnpA	rpsP		Υ
rnpA	rpsR		Υ
rnpA	rpsT		Υ
rnpA	rpsU		Υ
rnpA	spoT	Υ	
rnpA	ycbY	Υ	
rnpA	ycfL	Y	
rnpA	yciL	Υ	
rnpA	yfiF	Υ	
rnt	rnt	Y	
I			

rnt	aceF	Υ	
rpe	rpe	Y	Y
rpe	dnaK		Υ
rpe	dnaN	Υ	Υ
rpe	eno		Υ
rpe	hsdM		Υ
rpe	selD		Υ
rpe	trg		Υ
rpe	ydaY		Υ
rpe	yedD		Υ
rpe	yghK		Υ
rpe	yhfQ		Υ
rph	rph	Υ	
rph	dnaJ	Υ	
rph	dnaK	Υ	
rph	mreB	Υ	
rph	narG	Υ	
rph	pstB	Υ	
rph	recA	Υ	
rph	rfaD	Υ	
rph	tufA	Υ	
rph	tufB	Υ	
rpiB	rpiB	Υ	
rpiB	rplJ	Υ	
rpiB	rplL		Y
rpiB	rpmG		Υ
rpID	rpID	Y	
rpID	clpA	Y	
rpID	deaD	Y	
rpID	dnaK	Y	
rpID	dps	Y	
rpID	fusA	Y	
rpID	gatZ	Y	
rpID	pflB	Y	
rpID	rho	Y	
rpID	rpIA	Y	
rpID	rplB	Y	
rpID	rpIC	Y	
rpID	rpIE	Y	
rpID	rpIF	Y	
rpID	rplJ rplM	Y Y	
rpID	rpIM	Ϋ́	
rpID	rpIN rpIO	Ϋ́Υ	
rplD	rplQ	ĭ	

!	IV /	V		
rpID	rplV	Y		
rpID	rplX	Y		
rpID	rpoC	Y		
rpID	rpsB	Y		
rpID	rpsC	Y		
rpID	rpsD	Y		
rpID	rpsE	Y		
rpID	rpsG	Y		
rpID	rpsJ	Y		
rpID	rpsK	Y		
rpID	rpsM	Y		
rpID	secA	Y		
rpID	srmB	Y		
rpID	tufB	Y		
rpID	vacB	Y		
rplD	ygiF	Y		
rpIW	rpIW	Y	Υ	
rplW	rplA	Υ	Υ	
rplW	rplB	Υ	Υ	
rplW	rplC	Υ	Υ	
rplW	rpID	Υ	Υ	
rplW	rplE		Υ	
rplW	rplF		Υ	
rplW	rpll		Υ	
rplW	rplJ	Υ		
rplW	rplK		Υ	
rplW	rpIL		Υ	
rplW	rplM		Υ	
rplW	rplO		Υ	
rplW	rplR		Υ	
rplW	rpIS		Υ	
rplW	rpIU		Υ	
rplW	rpIV		Υ	
rplW	rplX		Υ	
rplW	rplY		Υ	
rplW	rpmB		Υ	
rplW	rpmC		Υ	
rplW	rpmG		Υ	
rplW	rpsC		Υ	
rplW	secA	Υ		
rpoA	rpoA	Υ	Y	
rpoA	aceE	Υ		
rpoA	aceF	Υ		
rpoA	aspS	Υ	Υ	
•	-			

l.			
rpoA	cspC		Y
rpoA	greB	Y	
rpoA	hepA	Υ	Y
rpoA	hupA		Υ
rpoA	mreB	Υ	
rpoA	nusA	Υ	Υ
rpoA	nusG	Υ	Υ
rpoA	pssA	Υ	
rpoA	rfaK	Υ	
rpoA	rho	Υ	
rpoA	rplA	Υ	
rpoA	rpIB	Υ	
rpoA	rpIC	Υ	
rpoA	rpID	Υ	Υ
rpoA	rplE	Υ	
rpoA	rpIL		Υ
rpoA	rplM	Υ	Y
rpoA	rplO	Υ	
rpoA	rplP		Υ
rpoA	rpIS		Υ
rpoA	rpIU		Υ
rpoA	rplW		Υ
rpoA	rplX		Υ
rpoA	rpmA		Υ
rpoA	rpmB		Υ
rpoA	rpmG		Υ
rpoA	rpoB	Υ	Υ
rpoA	rpoC	Υ	Υ
rpoA	rpoD	Υ	Υ
rpoA	rpoZ	Υ	Υ
rpoA	rpsA	Υ	
rpoA	rpsB	Υ	
rpoA	rpsC	Υ	
rpoA	rpsD	Υ	
rpoA	rpsE	Υ	Υ
rpoA	rpsF		Υ
rpoA	rpsG	Υ	
rpoA	rpsJ	Υ	
rpoA	rpsM	Υ	
rpoA	rpsN		Υ
rpoA	rpsO		Υ
rpoA	rpsP		Υ
rpoA	rpsS		Υ
rpoA			Υ

rpoA	rpsU		Υ
rpoA	wecG		Υ
rpoB	rpoB	Y	Υ
rpoB	atpD	Υ	
rpoB	b1192	•	Υ
rpoB	b1629		Ϋ́
rpoB	b2881		Ϋ́
	cca	Υ	
rpoB		Y	
rpoB	clpA	Y	V
rpoB	cycA	.	Y
rpoB	cysB	Y	
rpoB	hepA	Y	
rpoB	hupA	Υ	Υ
rpoB	mutL		Υ
rpoB	nusA	Υ	Υ
rpoB	nusG	Υ	Υ
rpoB	pstB		Υ
rpoB	rho	Υ	
rpoB	rhsE		Υ
rpoB	ribD		Υ
rpoB	rplB	Υ	
rpoB	rpID	-	Υ
rpoB	rplK		Ϋ́
rpoB	rplL		Ϋ́
rpoB	rpIV		Ϋ́
-	rpmG		Ϋ́
rpoB	•	V	
rpoB	rpoA	Y	Y
rpoB	rpoC	Y	Y
rpoB	rpoD	Y	Y
rpoB	rpoH	Y	
rpoB	rpoN	Υ	
rpoB	rpoS		Υ
rpoB	rpoZ	Υ	Υ
rpoB	rpsC	Υ	
rpoB	rpsE		Υ
rpoB	rpsF		Υ
rpoB	rpsJ	Υ	Υ
rpoB	rpsN		Υ
rpoB	rpsT		Υ
rpoB	rpsU		Υ
rpoB	secA	Υ	
rpoB	torR	-	Υ
rpoB	tufA	Υ	•
rpoB	tufB	Ϋ́	
IPOD	luib	I	

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rpoB	yacL		Y
rpoB	yahO	Υ	
rpoB	ybhF	Υ	
rpoB	yidE		Υ
rpoC	rpoC	Υ	Y
rpoC	b0499	•	Y
rpoC	b2372		Ϋ́
rpoC	cysQ		Ϋ́
rpoC	flxA		Ϋ́
rpoC	focA		Ϋ́
-		V	I
rpoC	greB	Υ	V
rpoC	hemC	3.4	Y
rpoC	hepA	Υ	Y
rpoC	hupA		Υ
rpoC	malP	Y	
rpoC	nusA	Υ	
rpoC	nusG	Υ	Y
rpoC	rpIB	Υ	Υ
rpoC	rpIC	Υ	Υ
rpoC	rpID	Υ	
rpoC	rpll		Υ
rpoC	rplM	Υ	
rpoC	rplO	Υ	
rpoC	rpIP	•	Υ
rpoC	rplQ	Υ	•
rpoC	rpIS	•	Υ
rpoC	rpIV		Ϋ́
rpoC	rpIX	V	Y
rpoC	rpoA	Y	Y
rpoC	rpoB	Y	Y
rpoC	rpoD	Y	Y
rpoC	rpoZ	Υ	Y
rpoC	rpsC		Y
rpoC	rpsD	Υ	Υ
rpoC	rpsE	Υ	Υ
rpoC	rpsG	Υ	Υ
rpoC	rpsJ		Υ
rpoC	rpsM	Υ	
rpoC	rpsP		Υ
rpoC	ygcP		Υ
rpoC	yjbH		Υ
rpoD	rpoD	Y	Y
rpoD	accA	Ý	•
rpoD	infB	Ý	Υ
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rpoD	nusA	Y	Y
rpoD	рерВ		Υ
rpoD	rho		Υ
rpoD	rplA	Y	Υ
rpoD	rplB	Υ	
rpoD	rplD	Υ	Υ
rpoD	rplL		Υ
rpoD	rplO	Υ	
rpoD	rpIS		Υ
rpoD	rpIU		Υ
rpoD	rpIV	Υ	
rpoD	rpmB		Υ
rpoD	rpoA	Υ	Υ
rpoD	rpoB	Y	Y
rpoD	rpoC	Υ	Υ
rpoD	rpoZ	Ϋ́	Ϋ́
rpoD	rpsA	•	Y
rpoD	rpsB	Y	•
rpoD	rpsC	Ϋ́	
rpoD	rpsD	Ϋ́	
rpoD	rpsE	Ϋ́	
rpoD	rpsF	ı	Υ
rpoD	rpsG	Υ	Ϋ́
rpoD	tufA	ı	Ϋ́
rpoD	yeeX		Ϋ́
rpoH	rpoH	Y	Y
rpoH	dnaK	Ϋ́	Ϋ́
rpoH	qor	Y	Ϋ́
rpoH	rplC	Ϋ́	•
rpoH	rplL	ı	Υ
rpoH	rpoA	Y	•
rpoH	rpoB	Ϋ́	
rpoH	rpoC	Ϋ́	
rpoH	rpsD	Ϋ́	
rpoN	rpoN	Y	Υ
rpoN	dnaK	Ϋ́	Ϋ́
rpoN	manX	Ϋ́	1
rpoN	rplD	Ϋ́	
	•	Ϋ́	
rpoN	rpIM	Y	V
rpoN	rpoA	Y	Y Y
rpoN	rpoB	Ϋ́	Ĭ
rpoN	rpoC	Ϋ́Υ	
rpoN	yraM		
rpoS	rpoS	Υ	

rpoS	manX	Υ	
rpoS	rplA	Y	
rpoS	rpoA	Ϋ́	
rpoS	rpoB	Ϋ́	
rpoS	rpoC	Y	
rpoS	rpsB	Y	
rpoS	tufA	Ϋ́	
rpoZ	rpoZ	Y	Υ
rpoZ	b1731	Y	7
_	dnaK	Y	
rpoZ		Ϋ́	
rpoZ	hepA nusA	Y Y	
rpoZ		Y	V
rpoZ	nusG	Ϋ́	Y
rpoZ	rpIB	Y	V
rpoZ	rplL		Y
rpoZ	rplW	V	Υ
rpoZ	rpoA	Y	V
rpoZ	rpoB	Y	Y
rpoZ	rpoC	Y	Y
rpoZ	rpoD	Y	Υ
rpoZ	rpsB	Y	
rpoZ	rpsG	Y	
rpoZ	rpsJ	Y	
rpoZ	rpsM	Υ	
rpoZ	rpsT		Υ
rpoZ	spoT	Υ	
rpoZ	thiG		Υ
rpoZ	tig	Υ	
rpoZ	ygjl		Y
rpoZ	ykgC		Y
rpsE	rpsE	Υ	Y
rpsE	gyrA		Y
rpsE	gyrB		Υ
rpsE	infC		Υ
rpsE	rplC		Υ
rpsE	rplU	_	Y
rpsE	rpsA	Υ	Υ
rpsE	rpsB	Υ	Υ
rpsE	rpsC	Υ	Υ
rpsE	rpsD	Υ	Υ
rpsE	rpsF	Υ	Υ
rpsE	rpsG	Υ	Υ
rpsE	rpsH		Υ
rpsE	rpsl	Υ	

rpsE rpsJ Y Y	
rpsE rpsL Y	
rpsE rpsM Y Y	
rpsE rpsP Y	
rpsE rpsS Y	
rpsE rpsT Y	
rpsE tufA Y	
rpsE tufB Y	
rsuA rsuA Y Y	•
rsuA rpsB Y	
rsuA rpsD Y	
rsuA rpsl Y	
rsuA rpsJ Y Y	
rsuA rpsM Y	
rsuA rpsP Y	
rsuA ycjF Y	
ruvB ruvB Y Y	
ruvB clpB Y	
ruvB dnaJ Y Y	
ruvB dnaK Y Y	
ruvB mopA Y Y	
ruvB rho Y	
ruvB rpsB Y	
ruvB rpsJ Y	
ruvB tufA Y	
sbcC sbcC Y Y	•
sbcC b1410 Y	
secA secA Y Y	•
secA aas Y	
secA b2710 Y	
secA bioB Y	
secA dnaK Y	
secA gapC_1 Y	
secA gatZ Y	
secA infC Y	
secA metK Y	
secA minD Y	
secA motA Y	
secA mreB Y	
secA rho Y	
secA rpID Y	
secA rplK Y	
secA rplL Y	
secA rplV Y	

secA	rpsB		Υ
secA	secB	Υ	Υ
secA	tufA	Υ	Υ
secA	tufB	Υ	
secA	ybiU		Υ
secA	ycaO	Υ	
secA	ycfB		Υ
secB	secB	Υ	Y
secB	b0362		Υ
secB	cpxR		Υ
secB	İon	Υ	Υ
secB	secA	Υ	
selB	selB	Y	Y
selB	aceE	Υ	
selB	aceF	Υ	
selB	dnaJ	Υ	Υ
selB	pta	Υ	
selB	rho		Υ
selB	rplA		Υ
selB	rplB	Υ	
selB	rplC	Υ	Υ
selB	rplD		Υ
selB	rpll		Υ
selB	rpIL		Υ
selB	rplM		Υ
selB	rplO		Υ
selB	rpIS		Υ
selB	rpIT		Υ
selB	rpIU		Υ
selB	rplV		Υ
selB	rplX		Υ
selB	rplY		Υ
selB	rpmB		Υ
selB	rpsA		Υ
selB	rpsC		Υ
selB	rpsD	Υ	Υ
selB	rpsE		Υ
selB	rpsF		Υ
selB	rpsG		Υ
selB	rpsT		Υ
selB	srmB	Υ	Υ
selB	vacB	Υ	
selB	yciL	Υ	
selB	yfiF	Υ	Y
•	•		

serC	serC	Υ	Y
serC	dnaK	Ϋ́	•
serC	selB	'	Υ
		V	Y
serS	serS	Υ	
serS	recF		Y
serS	yeiR		Υ
serS	yfjD		Υ
serS	yhjN		Υ
serS	yjfN		Υ
sfhB	sfhB	Υ	Y
sfhB	pepQ	Υ	
sfhB	ycdF		Υ
slpA	slpA	Υ	Y
slpA	adhC		Υ
slpA	ebgA		Υ
slpA	eda		Ϋ́
slpA	flgl		Ϋ́
slpA	glvG		Y
slpA	hupA		Ϋ́
	nuol		Ϋ́
slpA			
slpA	rpll		Y
slpA	rpsA		Y
slpA	rpsE		Y
slpA	rpsG		Y
slpA	rpsM		Y
slpA	rpsN		Υ
slpA	rpsP		Υ
slpA	ycgX		Υ
slpA	ycjJ		Υ
slpA	ydhU		Υ
slpA	ygiR		Υ
slpA	yhbY		Υ
slyD	slyD	Υ	Y
slyD	gĺtL		Υ
slyD	hupA		Υ
slyD	hupB		Υ
slyD	hypB		Ϋ́
slyD	sapD		Ϋ́
slyD	serS		Ϋ́
slyD			Ϋ́
_	sgaE ydel		Ϋ́
slyD	-		Ϋ́
slyD	yeaB		
slyD	yfiD		Y
slyD	yhjQ		Υ

smpB	smpB	Υ	Y
smpB	aidB	Υ	
smpB	clpA	Υ	
smpB	gyrA		Υ
smpB	gyrB		Υ
smpB	hlpA		Υ
smpB	hupA		Υ
smpB	hupB		Υ
smpB	infB	Υ	
smpB	malP	Υ	
smpB	parC	Υ	
smpB	rplF	Υ	Υ
smpB	rplM		Υ
smpB	rplX		Υ
smpB	rpmB		Υ
smpB	rpsA	Υ	Υ
smpB	rpsB	Υ	Υ
smpB	rpsC	Υ	
smpB	rpsE		Υ
smpB	rpsN		Υ
smpB	secA	Υ	
smpB	selB	Υ	
smpB	ycbB	Υ	
smpB	yeeX		Υ
smpB	yfiF	Υ	
speE	speE	Υ	
spoT	spoT	Υ	Y
spoT	acpP		Υ
spoT	deaD	Υ	
spoT	hrpA	Υ	
spoT	hupA		Υ
spoT	mukB	Υ	
spoT	rplB	Υ	Υ
spoT	rpIC	Υ	Y
spoT	rpID	Υ	Υ
spoT	rpll		Υ
spoT	rpIL		Υ
spoT	rplM		Υ
spoT	rplO		Υ
spoT	rplQ		Υ
spoT	rplR		Υ
spoT	rpIS		Υ
spoT	rplU		Υ
spoT	rplV	Υ	Υ

spoT	rplX		Y
spoT	rpmB		Ϋ́
spoT	rpoC	Υ	•
spoT	rpsA	Ϋ́	
spoT	rpsB		Υ
spoT	rpsC	Υ	•
spoT	rpsD	Ϋ́	
spoT	rpsE		Υ
spoT	rpsF		Y
spoT	rpsG	Υ	Ϋ́
spoT	rpsH	•	Ϋ́
spoT	rpsl		Ϋ́
spoT	rpsJ		Ϋ́
spoT	rpsK	Υ	•
spoT	rpsk	'	Υ
spoT	rpsN		Ϋ́
spoT	rpsN		Y
spoT	rpsR		Y
spoT	rpsT		Ϋ́
spoT	spoU	Υ	•
spoT	srmB	Ϋ́	
spoT	tolR	Ϋ́	
spoT	tufA	Ϋ́	
spoT	yciL	Ϋ́	
spoT	yfjB	•	Υ
spoU	spoU	Υ	•
srIR	srIR	,	Υ
srlR	b1410		Ϋ́
srlR	dnaK		Ϋ́
srlR	entC		Ϋ́
srlR	hupA		Ϋ́
srlR	hupB		Ϋ́
srIR	pflB		Ϋ́
srlR	rplA		Ϋ́
srlR	rplB		Ϋ́
srlR	rplC		Ϋ́
srlR	rpID		Ϋ́
srlR	rplE		Ϋ́
srlR	rplM		Ϋ́
srlR	rplO		Ϋ́
srlR	rplR		Ϋ́
srlR	rplT		Ϋ́
srlR	rplU		Ϋ́
srlR	rpIV		Ϋ́
1	. 1		<u>-</u>

srIR srIR srIR srIR srIR srIR srIR srMB	rpIW rpmC rpmG rpsA rpsB rpsE ydaT yfiF srmB dnaJ	Y Y	Y Y Y Y Y
srmB	rplA	Υ	
srmB	rplB	Υ	
srmB	rpID	Υ	Υ
srmB	rplL		Υ
srmB	rpIV	Y	Υ
srmB	rplW		Υ
srmB	rplX	Y	Υ
srmB	rpsB	Y	
srmB	rpsC	Y	
srmB	tufA	Y	
srmB	tufB	Y	
srmB	ycbY	Y	
srmB	yciL	Y	
ssb	ssb 50050	Y	Υ
ssb	b0359	Y	
ssb	dnaK	Y	
مامم	a A	\/	
ssb	gyrA	Υ	v
ssb	hupA		Y
ssb ssb	hupA malP	Υ	Y
ssb ssb ssb	hupA malP mtgA	Y Y	Y
ssb ssb ssb	hupA malP mtgA nohA	Y Y Y	Y
ssb ssb ssb ssb	hupA malP mtgA nohA parC	Y Y Y Y	Y
ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA	Y Y Y Y	Y
ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG	Y Y Y Y Y	Y
ssb ssb ssb ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG recJ	Y Y Y Y Y	Y
ssb ssb ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG recJ	Y Y Y Y Y Y	Y
ssb ssb ssb ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG recJ	Y Y Y Y Y	Y
ssb ssb ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG recJ recQ rplB	Y Y Y Y Y Y Y	
ssb ssb ssb ssb ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG recJ recQ rplB rplC	Y Y Y Y Y Y Y	Y
ssb ssb ssb ssb ssb ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG recJ recQ rplB rplC rplD	Y Y Y Y Y Y	Y Y
ssb ssb ssb ssb ssb ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG recJ recQ rplB rplC rplD rplL	Y Y Y Y Y Y	Y Y Y Y
ssb ssb ssb ssb ssb ssb ssb ssb ssb ssb	hupA malP mtgA nohA parC priA recG recJ recQ rplB rplC rplD rplL rplS	Y Y Y Y Y Y	Y Y Y Y

ssb	rpmG		Υ
ssb	rpoA		Υ
ssb	rpoB	Υ	
ssb	rpoC	Υ	
ssb	rpsD	Υ	
ssb	rpsJ		Υ
ssb	rpsU		Υ
ssb	sbcB	Υ	Υ
ssb	secA	Υ	
ssb	slyD		Υ
ssb	thdF	Υ	
ssb	topB	Υ	
ssb	tufA	Υ	Υ
ssb	tufB	Υ	
ssb	yfbG	Υ	
ssb	yfiF	Υ	
sspA	sspA	Y	
sspA	dnaK		Υ
sspA	hrsA		Υ
sspA	lysU		Υ
sspA	mopA		Υ
sspA	pflB		Υ
sspA	rpll		Υ
sspA	rpoA		Υ
sspA	rpoB	Υ	
sspA	rpoC	Υ	
sspA	serS		Υ
sspA	thiG		Υ
sspA	tyrP		Υ
sspA	ybaT		Υ
sspB	sspB	Υ	Υ
sspB	clpA		Υ
sspB	dnaE	Υ	
sspB	dnaK		Υ
sspB	elaB		Υ
sspB	rho		Υ
sspB	rpID		Υ
sspB	rpsG		Υ
sspB	slyD		Υ
sspB	tufA	Υ	
sspB	ybdN		Υ
sspB	yffG		Υ
sucA	sucA	Y	Υ
sucA	hflK	Υ	

sucA	lipA	Y	
sucA	lpdA	Υ	Υ
sucA	rplC		Υ
sucA	sucB	Υ	Υ
sucB	sucB	Υ	Y
sucB	lpdA	Y	
sucB	sucA	Υ	
sucB	ybaY		Υ
sucB	yedO	Υ	
surA	surA	Y	Y
surA	celB		Υ
surA	ilvA		Υ
surA	lpdA		Υ
surA	rplL		Υ
surA	rpsN		Υ
surA	ydaY		Υ
tag	tag	Υ	
tag	aceE	Υ	
tag	rpmC		Υ
tag	rpsE		Y Y Y
tag	serS		Υ
tdcD	tdcD	Υ	
tdcD	glf		Υ
tdcD	lysS	Υ	
tdcD	tufA	Υ	
tgt	tgt	Υ	Y
tgt	aceE	Υ	
tgt	aceF	Υ	
tgt	deaD	Υ	
tgt	hupA		Υ
tgt	hupB		Υ
tgt	rplA	Υ	
tgt	rplB	Υ	Υ
tgt	rpIC		Υ
tgt	rpID		Υ
tgt	rpll	Υ	Υ
tgt	rplL		Υ
tgt	rplM		Υ
tgt	rplQ		Υ
tgt	rpIS		Υ
tgt	rpIT		Υ
tgt	rplU		Υ
tgt	rpIV		Υ
tgt	rpIX		Υ
เลเ	· P·X		•

tgt rpoB Y tgt rpoB Y tgt rpsA Y tgt rpsE Y tgt rpsG Y tgt rpsU Y tgt rpsN Y tgt smpB Y tgt ycbY Y tgt yciL Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF db8A Y thdF dbA Y thdF hupA Y thdF mdh Y thdF rplD Y thdF rplD Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsD Y thdF rpsB Y thd				
tgt rpoB Y tgt rpsA Y tgt rpsE Y tgt rpsG Y tgt rpsL Y tgt rpsN Y tgt smpB Y tgt ycbY Y tgt yciL Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF dsbA Y thdF dsbA Y thdF mdh Y thdF rplC Y thdF rplD Y thdF rplW Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsG Y thdF rpsB Y thdF rpsG Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsG Y thdF rpsD Y thdF rpsG Y thdF rpsG Y thdF rpsG Y thdF rpsG Y thdF rpsG Y thdF rpsD Y thdF rpsD Y thdF rpsG Y thdF rpsD	tat	rnmR		V
tgt rpsA Y tgt rpsE Y tgt rpsG Y tgt rpsN Y tgt rpsN Y tgt smpB Y tgt ycbY Y tgt yciL Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF dsbA Y thdF dbA Y thdF mdh Y thdF nplD Y thdF rplD Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsB Y thdF rpsG Y thdF rpsC Y thdF rpsB Y thdF rpsC Y thdF rpsB Y thdF rpsC Y thdF rpsB Y thdF rpsC Y thdF rpsB Y thdF rpsC Y thdF rpsC Y thdF rpsB Y thdF rpsC Y t		•	V	1
tgt rpsE Y tgt rpsG Y tgt rpsL Y tgt rpsN Y tgt smpB Y tgt ycbY Y tgt yciL Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF dcm Y thdF dbA Y thdF lpdA Y thdF rplC Y thdF rplD Y thdF rpsB Y thdF rpsB Y thdF rpsG Y thdF rpsG Y thdF rpsG Y thdF rpsO Y thdF secA Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y thrS dnaK		•		
tgt rpsG Y tgt rpsN Y tgt rpsN Y tgt smpB Y tgt ycbY Y tgt yciL Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF dcm Y thdF dsbA Y thdF hupA Y thdF mdh Y thdF rplC Y thdF rplD Y thdF rpsB Y thdF rpsB Y thdF rpsG Y thdF rpsG Y thdF rpsO Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y thrS dnaK		-	Y	
tgt rpsL Y tgt rpsN Y tgt smpB Y tgt ycbY Y tgt yciL Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF dcm Y thdF dsbA Y thdF lpdA Y thdF lpdA Y thdF rplC Y thdF rplM Y thdF rpsB Y thdF rpsB Y thdF rpsG Y thdF rpsN Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ydA Y thdF secA Y thdF ydA Y thdF spdA Y thdF rpsN Y thdF rpsN Y thdF rpsA Y thdF rpsN Y thdF secA Y thdF ydA Y thdF ydA Y thdF rpsA Y thdF rpsN Y thdF rpsN Y thdF rpsN Y thdF sapF Y thdF ydA Y thdF ydA Y thdF ydA Y thrS thrS Y thrS dnaK	tgt	rpsE		Υ
tgt rpsL Y tgt rpsN Y tgt smpB Y tgt ycbY Y tgt yciL Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF dcm Y thdF dsbA Y thdF lpdA Y thdF lpdA Y thdF rplC Y thdF rplM Y thdF rpsB Y thdF rpsB Y thdF rpsG Y thdF rpsN Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ydA Y thdF secA Y thdF ydA Y thdF spdA Y thdF rpsN Y thdF rpsN Y thdF rpsA Y thdF rpsN Y thdF secA Y thdF ydA Y thdF ydA Y thdF rpsA Y thdF rpsN Y thdF rpsN Y thdF rpsN Y thdF sapF Y thdF ydA Y thdF ydA Y thdF ydA Y thrS thrS Y thrS dnaK	tgt	rpsG		Υ
tgt rpsN Y tgt smpB Y tgt ycbY Y tgt yciL Y tgt yfiF Y Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF dcm Y thdF dsbA Y thdF lpdA Y thdF lpdA Y thdF rplD Y thdF rplD Y thdF rpsB Y Y thdF rpsG Y thdF rpsA Y thdF		rpsL		Υ
tgt ycbY Y tgt yciL Y tgt yfiF Y Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF dcm Y thdF dsbA Y thdF lpdA Y thdF lpdA Y thdF rplC Y thdF rplD Y thdF rplW Y thdF rpsB Y Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF ydA Y thdF sapF Y thdF ydA Y thdF secA Y thdF ydA Y thdF secA Y thdF ydA Y thdF ydA Y thdF ydA Y thdF ydA Y thdF rpsN Y thdF rpsN Y thdF rpsN Y thdF ydA Y thdF ydA Y thdF ydA Y thdF ydA Y thrS thrS Y thrS dnaK		•		
tgt ycbY tgt yciL Y tgt yfiF Y tgt yhjO Y thdF thdF Y thdF aceE Y ThdF b0816 Y thdF dcm Y thdF dsbA Y thdF hupA ThdF hupA ThdF rplC ThdF rplD Y ThdF rplD Y ThdF rpsB Y ThdF rpsB Y ThdF rpsB Y ThdF rpsG ThdF rpsN ThdF rpsN ThdF rpsN ThdF secA ThdF ydA Y ThdF ydA Y ThdF secA ThdF ydA Y ThdF ydA Y ThdF ydA Y ThdF rpsN ThdF rpsN ThdF rpsN ThdF rpsN ThdF rpsN ThdF rpsN ThdF ydA ThrS thrS Y ThrS dnaK		-		
tgt yfiF Y Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF b0816 Y thdF dsbA Y thdF gidA Y thdF lpdA Y thdF rplC Y thdF rplD Y thdF rplW Y thdF rpsB Y Y thdF rpsG Y thdF rpsA Y thdF secA Y thdF ycdP thdF yidA Y thrS thrS Y thrS dnaK			V	•
tgt yfiF Y Y tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF b0816 thdF dsbA Y thdF gidA Y thdF hupA thdF hupA thdF rplC thdF rplD Y thdF rplD Y thdF rplW thdF rpsB Y Y thdF rpsG thdF rpsJ Y thdF rpsJ Y thdF sapF thdF secA thdF ydA Y thdF ydA Y thdF secA thdF ydA Y thdF ydA Y thdF rpsN Y thdF rpsN Y thdF rpsN Y thdF sapF thdF secA thdF ydA Y thrS thrS Y Y thrS dnaK		-		
tgt yhjO Y thdF thdF Y Y thdF aceE Y Y thdF b0816 thdF dcm Y thdF dsbA Y thdF gidA Y thdF hupA Y thdF lpdA Y thdF rplC Y thdF rplD Y thdF rplM Y thdF rplW Y thdF rpsB Y Y thdF rpsG Y thdF rpsJ Y thdF rpsJ Y thdF sapF Y thdF secA Y thdF yidA Y thrS thrS Y thrS dnaK		-		V
thdF thdF Y Y thdF aceE Y thdF b0816 thdF dcm Y thdF dsbA Y thdF gidA Y thdF lpdA Y thdF ndh Y thdF rplC Y thdF rplD Y thdF rplM Y thdF rpsB Y thdF rpsB Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF thdF secA Y thdF ydA Y thdF ydA Y thdF ydA Y thdF ydA Y thdF rpsG Y thdF rpsN Y thdF rpsN Y thdF rpsN Y thdF sapF Y thdF ydA Y thdF ydA Y thrS thrS Y thrS dnaK		-	Y	
thdF b0816 thdF b0816 thdF dcm thdF dsbA thdF gidA Y thdF pidA thdF lpdA thdF mdh thdF rplC thdF rplD thdF rplM thdF rplW thdF rpsB Y thdF rpsB Y thdF rpsG thdF rpsG thdF rpsJ thdF rpsN thdF rpsN thdF sapF thdF secA thdF ycdP thdF yidA thrS thrS Y thrS dnaK				
thdF b0816 thdF dcm Y thdF dsbA Y thdF gidA Y thdF hupA ThdF hupA ThdF lpdA Y ThdF rplC ThdF rplD ThdF rplM ThdF rplW ThdF rpsB ThdF rpsB ThdF rpsG ThdF rpsG ThdF rpsN ThdF rpsN ThdF rpsN ThdF sapF ThdF secA ThdF ydA ThrS thrS Ty ThdF ydA ThrS thrS Ty ThdF TysG ThdF ydA ThrS thrS Ty ThdF ydA ThrS thrS Ty ThdF ydA ThrS thrS Ty ThdF ydA ThdF ydA ThrS thrS Ty ThdF ydA ThrS thrS Ty ThdF ydA ThrS thrS Ty ThdF ydA ThrS thrS Ty ThdF ydA ThrS Ty ThdF ydA ThrS Ty ThdF ydA ThrS Ty ThdF ydA ThrS ThrS Ty ThdF ydA ThrS ThrS Ty ThdF ydA ThrS ThrS Ty ThdF ydA ThrS ThrS ThrS ThrS ThrS ThrS ThrS ThrS		thdF		
thdF dsbA Y thdF gidA Y thdF gidA Y thdF hupA Y thdF lpdA Y thdF mdh Y thdF rplC Y thdF rplD Y thdF rplM Y thdF rplW Y thdF rpsB Y Y thdF rpsG Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ydA Y thrS thrS Y Y thrS dnaK Y	thdF	aceE	Y	Υ
thdF gidA Y thdF gidA Y thdF hupA thdF lpdA Y thdF mdh Y thdF rplC Y thdF rplD Y thdF rplW Y thdF rprW Y thdF rpsB Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF sapF Y thdF secA Y thdF ydA Y thrS thrS Y Y thrS dnaK	thdF	b0816		Υ
thdF gidA Y thdF hupA Y thdF lpdA Y thdF mdh Y thdF rplC Y thdF rplD Y thdF rplM Y thdF rplW Y thdF rpmG Y thdF rpsB Y Y thdF rpsG Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ydA Y thrS thrS Y Y thdF drs dnaK Y	thdF	dcm		Υ
thdF hupA Y thdF lpdA Y thdF mdh Y thdF rplC Y thdF rplD Y thdF rplM Y thdF rplV Y thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ydA Y thrS thrS Y Y thrS dnaK	thdF	dsbA		Υ
thdF hupA Y thdF lpdA Y thdF mdh Y thdF rplC Y thdF rplD Y thdF rplM Y thdF rplV Y thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ydA Y thrS thrS Y Y thrS dnaK	thdF	gidA	Υ	Y
thdF mdh Y thdF rplC Y thdF rplD Y thdF rplM Y thdF rplW Y thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK	thdF			Υ
thdF rpIC Y thdF rpID Y thdF rpIM Y thdF rpIV Y thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF yidA Y thrS thrS Y thrS dnaK Y	thdF	lpdA		Υ
thdF rpID Y thdF rpIM Y thdF rpIV Y thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK	thdF	mdh		Υ
thdF rpIM Y thdF rpIV Y thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thdF dark	thdF	rplC		Υ
thdF rpIV Y thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thdS dnaK	thdF	rplD		Υ
thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thdS dnaK Y	thdF	rplM		Υ
thdF rpmG Y thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thdS dnaK Y	thdF	rpIV		Υ
thdF rpsB Y Y thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK Y	thdF			Υ
thdF rpsF Y thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK Y		-	Υ	
thdF rpsG Y thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK Y		•	•	
thdF rpsJ Y thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK Y				
thdF rpsN Y thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK Y				
thdF sapF Y thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK Y		-		
thdF secA Y thdF ycdP Y thdF yidA Y thrS thrS Y Y thrS dnaK Y		•		
thdF ycdP Y thdF yidA Y thrS thrS Y thrS dnaK Y		-		
thdF yidA Y thrS thrS Y Y thrS dnaK Y				
thrS thrS Y Y thrS dnaK Y		-		
thrS dnaK Y	thdF	-		
	thrS	thrS	Υ	
thrS eno Y	thrS	dnaK		
	thrS	eno		
thrS ibpA Y	thrS	ibpA		
thrS rpe Y	thrS	rpe		Υ
thrS rpIA Y Y	thrS	rplA	Υ	Υ
thrS rplC Y	thrS	rplC		Υ
thrS rpll Y	thrS	rpll		Υ
thrS rplM Y	thrS	-		Υ

tig yicP Y tktA tktA Y tmk tmk Y Y tmk rplL Y Y tmk tufA Y Y tnaA tnaA Y Y tnaA dnaK Y Y topA topA Y Y topA alaS Y topA b0878 Y	thrs thrs ssssssssssssssssssssssssssssss	rpIS rpIV rpsA rpsB rpsF rpsG rpsM rpsN rpsN rpsN rpsN rpsN rpsD rpsN rpsD rpsD rpIV rpmC rpsA rpsB rpsC rpsC rpsC rpsC rpsC rpsC rpsC rpsC	YY	Y
tktA tktA Y tmk tmk Y Y tmk rplL Y tmk tufA Y Y tnaA tnaA Y Y tnaA dnaK Y Y topA topA Y Y topA alaS Y topA b0878 Y		-		
tmk rplL Y tmk tufA Y tnaA tnaA Y Y tnaA dnaK Y Y topA topA Y Y topA aceF Y Y topA alaS Y topA b0878 Y				
tmk tufA Y tnaA tnaA Y Y tnaA dnaK Y Y topA topA Y Y topA aceF Y Y topA alaS Y topA b0878 Y			Υ	
tnaA tnaA Y Y tnaA dnaK Y Y topA topA Y Y topA aceF Y Y topA alaS Y topA b0878 Y		-		
topA topA Y Y topA aceF Y Y topA alaS Y topA b0878 Y	tnaA	tnaA	Υ	
topA aceF Y Y topA alaS Y topA b0878 Y				
topA alaS Y topA b0878 Y				
topA b0878 Y	-			Y
•	-			
	topA	gabP	1	Υ

topA	hns	Υ	
topA	hupA		Υ
topA	hupB		Υ
topA	kdgK		Υ
topA	lpdA	Υ	-
topA	malP	Y	
topA	pssA	Y	
topA	recJ	Y	
topA	recQ	Y	
topA	rfaL	•	Υ
topA	rob	Υ	-
topA	rplB	Y	
topA	rplC	Y	Υ
topA	rplM	•	Ϋ́
topA	rpIS	Υ	-
topA	rplV	Ϋ́	Υ
topA	rpmB	•	Ý
topA	rpmG		Y Y Y
topA	rpoA	Υ	•
topA	rpoB	Ϋ́	
topA	rpoC	Ϋ́	Υ
topA	rpoZ	•	Y
topA	rpsC	Υ	•
topA	rpsE	Υ	
topA	rpsG	Υ	
topA	rpsJ		Υ
topA	rpsM	Υ	
topA	rpsT		Υ
topA	rstA	Υ	
topA	srmB	Υ	
topA	ssb	Υ	
topA	topB	Υ	
topA	ybiT		Υ
topA	yciL	Υ	
topB	topB	Y	Υ
topB	aceE	Υ	
topB	aceF	Υ	Υ
topB	ccmB		Υ
topB	gcl		Y Y Y
topB	hupA		Υ
topB	IpdA	Υ	Υ
topB	recD	Υ	
topB	recQ	Υ	
topB	rplC	Υ	Υ
	-		

topB	rplD		Υ
topB	rplJ		Υ
topB	rpIL		Υ
topB	rplM		Υ
topB	rplS		Υ
topB	rplU		Y
topB	rplV		Υ
topB	rplX		Υ
topB	rpmB		Υ
topB	rpmC		Υ
topB	rpmG		Υ
topB	rpsB	Υ	Υ
topB	rpsC	Υ	
topB	rpsE		Υ
topB	rpsN		Υ
topB	rpsP		Y
topB	ssb	Υ	Y
topB	topA	Υ	
topB	ybeW		Υ
topB	ydaH		Υ
topB	yjcD		Υ
tpiA	tpiA	Y	Y
tpiA	dnaK	Y	
tpiA	entB		Y
tpiA	lysU -		Y
tpiA	pepT	Y	Y
tpiA	pflB	Y	Y
tpiA	rpmB		Y
tpiA	rpoA		Y
tpiA	rpsB		Y
tpiA	yfgB	V	Y
tpiA trmA	yfiD trmA	Y Y	Y Y
trmA	aceE	Ϋ́	1
trmA	rne	1	Υ
trmA	rplL		Ϋ́
trmA	yhbJ	Υ	•
trpS	trpS	Y	Υ
trpS	rplM	•	Ϋ́
trpS	rpIV		Ϋ́
trpS	rpsB		Υ
trpS	rpsE		Y
trxC	trxC	Y	Y Y
trxC	cmr		Υ
1			

trxC	codA		Υ
trxC	dnaK		Υ
trxC	glgP		Υ
trxC	mglB		Υ
trxC	murD	Υ	
trxC	ydbA 2		Υ
tsf	tsf	Υ	Y
tsf	asr	•	Ϋ́
tsf	ecpD		Ϋ́
tsf	hsdM		Ϋ́
tsf	rpsE		Ϋ́
tsf	tufA		Ϋ́
		V	ı
tsf	tufB	Y Y	Υ
tufA	tufA		
tufA	tsf	Y	Υ
tufA	tufB	Y	
tufB	tufB	Y	
tufB	acrF		Y
tufB	afuC		Y
tufB	b1664		Y
tufB	dnaJ		Y
tufB	dnaK		Υ
tufB	metK	Y	
tufB	mreB	Υ	
tufB	rpsB		Υ
tufB	rpsE		Υ
tufB	secA	Υ	Υ
tufB	tsf	Υ	Υ
tufB	tufA	Υ	Y
tufB	valS		Υ
tufB	xylA		Υ
tufB	yaaA		Υ
tufB	ybdQ		Υ
tufB	ybeD		Υ
tyrA	tyrA	Y	
tyrR	tyrR	Y	Υ
tyrR	aceF	Υ	
tyrS	tyrS	Y	Υ
tyrS	dnaK	Υ	Υ
tyrS	secA	Υ	
tyrS	sucB		Υ
tyrS	ybgF		Υ
ubiB	ubiB	Y	Y
ubiB	crp		Υ
ı	'		

ubiB ssb \	<i>(</i>
ubiB yacL	Υ
,	ΥΫ́Υ
	(
ubiC aceF	Y
ubiC rpmB	
	Y
•	1
	1
ubiG ubiG	ΥΥ
ubiG rplV	Y Y Y
ubiH ubiH	ΥΥ
ubiH rfaD \	<i>(</i>
	ΥΥ
ucpA lysS	Υ
ucpA yraQ	Y
_ =	Y Y
	Ϋ́Υ
	Y Y
	Υ
usg nusG	
usg pykA 	Y
usg rplL	Y
usg rpoA	Υ
usg rpoB	Υ
	Y Y
usg rpsC \	<i>(</i>
usg tufA \	/ Y
	<i>(</i>
usg ygfZ	Υ
_	ΥΥ
uup aldA	Υ
uup b2146	Y
·	· (
uup yaiP	Υ
uup ydaY	Ϋ́
	' '
	Y
	ΥΥ
uvrB hns	Y
uvrB hupA	Y
uvrB hupB	Υ
uvrB lpdA \	· · · · · · · · · · · · · · · · · · ·
uvrB nfi	Υ

uvrB	rpmB		Υ
uvrB	yhaD	Υ	
uvrC	uvrC	Υ	Υ
uvrC	aceE	Ϋ́	•
uvrC	b1487	•	Υ
uvrC	b2503		Ϋ́
uvrC	cspC		Ϋ́
uvrC	glnK		Ϋ́
			Ϋ́
uvrC	hupA	V	
uvrC	lpdA	Y	Y
uvrC	pntA		Y
uvrC	rplA	Y	Y
uvrC	rplB	Υ	Y
uvrC	rpIC	Υ	Y
uvrC	rplD		Y
uvrC	rpll		Υ
uvrC	rplM		Υ
uvrC	rplN		Υ
uvrC	rpIS		Y
uvrC	rplT		Υ
uvrC	rplU		Υ
uvrC	rplV		Υ
uvrC	rplY		Υ
uvrC	rpmB		Υ
uvrC	rpsA		Υ
uvrC	rpsB	Υ	
uvrC	rpsC	Υ	Y
uvrC	rpsD	Υ	Υ
uvrC	rpsE		Υ
uvrC	rpsF		Υ
uvrC	rpsG	Υ	Υ
uvrC	rpsH		Υ
uvrC	rpsJ		Υ
uvrC	rpsK	Υ	•
uvrC	rpsL		Υ
uvrC	rpsM	Υ	Ϋ́
uvrC	rpsN	•	Ϋ́
uvrC	rpsO		Ϋ́
uvrC	rpsR		Ϋ́
uvrC	rpsR		Ϋ́
uvrC	rps5		Y
uvrC	sucA		Y
uvrC			Ϋ́
	sucB	V	Ĭ
uvrC	yfiB	Y	

uvrC	yfiF		Υ
uvrC	yhbY		Υ
uvrC	yjeF	Υ	
uvrD	uvrD	Y	Y
uvrD	abc		Υ
uvrD	accA		Υ
uvrD	aceE		Υ
uvrD	aceF		Υ
uvrD	agaR		Υ
uvrD	araG		Υ
uvrD	b2879		Υ
uvrD	clpA		Υ
uvrD	dgt		Υ
uvrD	fis		Υ
uvrD	hns		Υ
uvrD	hupA		Υ
uvrD	hupB		Υ
uvrD	lpdA	Υ	Υ
uvrD	malP		Υ
uvrD	pepN		Υ
uvrD	pqiB		Υ
uvrD	rplC		Υ
uvrD	rplL		Υ
uvrD	rplM		Υ
uvrD	rplV		Υ
uvrD	rplW		Υ
uvrD	rpmC		Υ
uvrD	rpoB		Υ
uvrD	rpsB		Υ
uvrD	rpsE		Υ
uvrD	rpsF		Υ
uvrD	rpsG		Υ
uvrD	rpsN		Υ
uvrD	secA		Υ
uvrD	tgt		Υ
uvrY	uvrY	Y	
uvrY	fimZ		Υ
uvrY	lpdA	Υ	
uvrY	rpIL		Υ
uvrY	tufA	Υ	
uvrY	ybeW		Y
vacB	vacB	Y	Y
vacB	cca	Y	
vacB	deaD	Υ	

vacB	himA		Y
vacB	hupA		Υ
vacB	infC	Υ	
vacB	pgm	Υ	
vacB	pssA	Υ	
vacB	rplA	Υ	Υ
vacB	rpIB	Υ	Υ
vacB	rpIC	Υ	Υ
vacB	rpID	Υ	Υ
vacB	rpIF	Y	
vacB	rpll	Υ	Υ
vacB	rplM	Υ	Υ
vacB	rplN	Υ	
vacB	rplO	Y	Υ
vacB	rpIP		Υ
vacB	rplT		Y
vacB	rplU		Y
vacB	rplV		Y
vacB	rplW		Y
vacB	rplX		Y
vacB	rpsA	Y	
vacB	rpsB	Y	Y
vacB	rpsC	Y	
vacB	rpsD	Y	Y
vacB	rpsE	Y	Y
vacB	rpsF	Y	Y
vacB	rpsG	Y	Y
vacB	rpsH		Y
vacB	rpsl		Y
vacB	rpsK		Y
vacB	rpsM	Y	Y
vacB	rpsN		Y
vacB	rpsO	V	Y Y
vacB	rpsP	Y	Ϋ́Υ
vacB	rpsR		Ϋ́
vacB	rpsT		
vacB	rpsU	V	Y
vacB vacB	slpA	Y Y	
vacB	tgt vebY	Ϋ́	
vacB vacB	ycbY yceC	Ϋ́	
vacB	ycec yfiF	Ϋ́	
vacB	y iir ygiF	Y	
vacb valS	ygır valS	Y	
vais	vais	1	

1			
xerD	xerD	Υ	
xerD	acpP		Υ
xerD	mukB		Υ
xseA	xseA	Υ	
xseA	b1579	Υ	
xthA	xthA	Y	
xthA	aceE	Ϋ́	
xthA	aceF	Y	
xthA	lpdA	Ϋ́	
xthA	malP	Ϋ́	
		Υ	V
yabC	yabC	Y	Y
yabC	rpIC		Y
yabC	rpIL		Y
yabC	rpsJ		Υ
yacG	yacG	Y	Υ
yacG	cysN		Υ
yacG	gyrA	Υ	Υ
yacG	gyrB	Υ	Υ
yacL	yacL	Υ	Y
yacL	b2255		Y
yacL	crp		Y
yacL	elaB		Y
yacL	fur		Υ
yacL	gltA		Y
yacL	hepA	Υ	Υ
yacL	hfq		Y
yacL	nuoG		Υ
yacL	nusA	Υ	
yacL	nusG	Υ	Υ
yacL	pnp		Υ
yacL	proV		Υ
yacL	rplB	Υ	Υ
yacL	rplC	Υ	
yacL	rplD	Υ	
yacL	rplM	Υ	Υ
yacL	rplN		Υ
yacL	rplO	Υ	Υ
yacL	rplP		Υ
yacL	rplQ		Y
yacL	rplU		Ϋ́
yacL	rplV		Ϋ́
yacL	rpmB		Ϋ́
yacL	rpoA	Υ	Y
yacL	rpoB	Ϋ́	Ϋ́
, ,	- 1	•	•

wool	***	V	V
yacL yacL	rpoC rpoZ	Y Y	Y Y
yacL yacL	rpsA	I	Y
_	rpsA	Υ	1
yacL yacL	rpsC	ı	Υ
yacL	rpsD	Υ	Y
yacL	rpsE	'	Y
yacL	rps⊏		Ϋ́
yacL	rpsJ		Ϋ́
yacL	rpsK		Ϋ́
yacL	rpsl		Ϋ́
yacL	rpsM	Υ	•
yacL	rpsN		Υ
yacL	rpsT		Ϋ́
yacL	rpsU		Ϋ́
yacL	rsuA		Ϋ́
yacL	slyD		Ϋ́
yacL	ybdQ		Ϋ́
yacL	yqjl		Y
yadB	yadB	Υ	Y
yadB	aceF		Y
yadB	artQ		Υ
yadB	lpdA		Υ
yadB	rpIL		Υ
yadB	yhdP		Υ
yadB	ypjA		Υ
yadF	yadF	Υ	
yadF	tufA	Υ	
yadF	tufB	Υ	
yaeC	yaeC	Y	Y
yaeC	araG		Υ
yaeC	b2249		Υ
yaeC	tufA		Υ
yaeC	yabF		Υ
yaeC	yahK		Υ
yaeC	ymcD		Υ
yaeS	yaeS	Y	
yajQ	yajQ	Y	
yajQ	dnaN	Y	
yajQ	napD		Y
ybaB	ybaB		Y
ybaB	deaD		Y
ybaB	wzxC		Y
ybaB	ychN		Υ

ybaD	<i>ybaD</i> b2520	Y	Y
ybaD			Ϋ́
ybaD ybaD	glyS rplL		Ϋ́
ybaD	rplU		Ϋ́
ybaD	rpsJ		Ϋ́
ybaD	ydbK		Ϋ́
ybab	ybaK	Y	Y
ybaK	b1310	,	Ϋ́
ybaK	cpdB		Ϋ́
ybaK	IpdA	Υ	•
ybaK	rhsC	ı	Υ
ybaK	rplL		Ϋ́
ybaK	rpsB		Ϋ́
ybaK	rpsJ		Y
ybaK	ydjA		Ϋ́
ybaK	yhaD		Ϋ́
ybaX	ybaX	Υ	Y
ybaX	aidB		Y
ybaX	dnaK		Υ
ybaX	napD		Υ
ybaX	rplL		Υ
ybaX	rpsJ		Υ
ybaX	tufA	Υ	Υ
ybaX	yegT		Υ
ybaX	yfiD		Υ
ybaZ	ybaZ		Υ
ybaZ	b1624		Υ
ybaZ	hupA		Υ
ybaZ	lysS		Υ
ybaZ	marA		Υ
ybaZ	rplA		Υ
ybaZ	rplJ		Υ
ybaZ	rplL		Y
ybaZ	rpsB		Y
ybaZ	rpsJ		Y
ybaZ	ydgA		Y
ybaZ	ygcA	V	Υ
ybbA	ybbA	Y	
ybbL ybbl	ybbL mon^	Y	
ybbL	mopA	Y Y	V
<i>ybbN</i> ybbN	<i>ybbN</i> dnaN	Ϋ́Υ	Υ
ybbN	holE	ı	Υ
yoon	HUIL		ı

ybbN	yjjK		Υ
ybbU	ybbU		Y
ybbU	hupA		Υ
ybbU	rplA		Υ
ybbU	rplB		Υ
ybbU	rplR		Υ
ybbU	rpIS		Υ
ybbU	rplU		Υ
ybbU	rplX		Υ
ybbU	rpsE		Υ
ybbU	rpsF		Υ
ybbU	rpsG		Υ
ybbU	rpsL		Υ
ybbU	rpsM		Υ
ybbU	rpsN		Υ
ybbU	rpsO		Υ
ybcJ	ybcJ	Y	Y
ybcJ	aceF	Υ	Υ
ybcJ	aidB	Υ	
ybcJ	deaD	Υ	
ybcJ	dnaJ	Υ	Υ
ybcJ	eno		Υ
ybcJ	hupA		Υ
ybcJ	hupB		Υ
ybcJ	lpdA	Υ	
ybcJ	pnp	Υ	Υ
ybcJ	pssA	Υ	
ybcJ	rhlB	Υ	
ybcJ	rne		Υ
ybcJ	rplA	Υ	Υ
ybcJ	rplC	Υ	Υ
ybcJ	rplD		Υ
ybcJ	rpIF		Υ
ybcJ	rpll	Υ	Υ
ybcJ	rplK		Υ
ybcJ	rpIL		Υ
ybcJ	rplM		Υ
ybcJ	rplQ		Υ
ybcJ	rplR		Υ
ybcJ	rpIS		Υ
ybcJ	rpIT		Υ
ybcJ	rplU		Υ
ybcJ	rpIV		Υ
ybcJ	rplX		Υ

ybcJ	rpmG		Υ
ybcJ	rpoA	Υ	
ybcJ	rpoC	Ϋ́	
ybcJ	rpsA	Ϋ́	Υ
ybcJ	rpsB	Ϋ́	Ϋ́
ybcJ	rpsC	Ý	Ϋ́
ybcJ	rpsE	•	Ϋ́
ybcJ	rpsF		Ϋ́
ybcJ	rpsl		Ϋ́
ybcJ	srmB	Υ	•
ybcJ	vacB	Ϋ́	Υ
ybcJ	yajQ	Y	•
ybcJ	ycbY	Y	
ybcJ	yceC	Ϋ́	
_	-	Y	v
ybcJ	yciL	Ϋ́	Y
ybcJ	yfiF	Ϋ́	Υ
ybcJ	ygiF	Ϋ́	V
ybcJ	yhiR	Ť	Y
ybcJ	yibL		Y
ybcJ	yihl	\ \	Y
ybdQ	ybdQ	Y	Υ
ybdQ	dnaJ	Υ	
ybdQ	dnaK	Y	
ybdQ	mopA	Υ	
ybdQ ybdQ	mopA rfaD		
ybdQ ybdQ ybdQ	mopA rfaD rplL	Υ	Y
ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW	Υ	Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL	Υ	Y
ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW	Y	
ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW secA	Y	Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW secA secB	Y Y	Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW secA secB tufB	Y Y Y	Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW secA secB tufB ycaO	Y Y Y Y	Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW secA secB tufB ycaO ybeA	Y Y Y Y	Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB	Y Y Y Y Y	Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB ybeB	Y Y Y Y Y	Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB ybeB b1200	Y Y Y Y Y Y Y Y Y	Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybeA ybeA ybeB ybeB	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB ybeB b1200 cca rplD	Y Y Y Y Y Y Y Y Y	Y Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybeA ybeA ybeB ybeB ybeB	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB ybeB b1200 cca	Y Y Y Y Y Y Y Y Y	Y Y Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybeA ybeA ybeB ybeB ybeB ybeB	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB ybeB b1200 cca rplD rplL rplN	Y Y Y Y Y	Y Y Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybeA ybeA ybeB ybeB ybeB ybeB	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB b1200 cca rplD rplL rplN rplS	Y Y Y Y Y	Y Y Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybeA ybeB ybeB ybeB ybeB ybeB ybeB ybeB	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB b1200 cca rplD rplL rplN rplS yehL	Y Y Y Y Y	Y Y Y Y Y Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybeA ybeB ybeB ybeB ybeB ybeB ybeB ybeB ybeB	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB b1200 cca rplD rplL rplN rplS yehL yehQ	Y Y Y Y Y	Y Y Y Y Y Y Y Y
ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybdQ ybeA ybeB ybeB ybeB ybeB ybeB ybeB ybeB	mopA rfaD rplL rplW secA secB tufB ycaO ybeA rpsB b1200 cca rplD rplL rplN rplS yehL	Y Y Y Y Y	Y Y Y Y Y Y Y

ybeW yhjK Y ybeY ybeY Y ybeY ybeY Y ybeY ybeZ Y ybeZ ybeZ Y ybeZ aceE Y ybeZ aceE Y ybeZ deaD Y ybeZ fabZ Y ybeZ hupA Y ybeZ pssA Y ybeZ rpIB Y ybeZ rpIB Y ybeZ rpIC Y ybeZ rpIE Y ybeZ rpIE Y ybeZ rpII Y ybeZ r	ybeD ybeW ybeW ybeW ybeW ybeW	ybeD tufA ybeW b2443 clpA rplJ rpsB	Y Y Y Y Y	Υ
ybeY ybeZ Y ybeY yihK Y ybeZ ybeZ Y ybeZ aceE Y ybeZ aceF Y ybeZ b2710 Y ybeZ deaD Y ybeZ fabZ Y ybeZ hupA Y ybeZ pssA Y ybeZ rpIA Y Y ybeZ rpIB Y ybeZ rpIB Y ybeZ rpIE Y Y ybeZ rpIE Y Y ybeZ rpII Y Y ybeZ rpII Y Y ybeZ rpII Y ybeZ rpIX Y ybeZ rpIX Y ybeZ rpIX Y ybeZ rpmB Y ybeZ rpsA Y ybeZ rpsA Y ybeZ rpsB Y	ybeW	yhjK		
ybeY yihK ybeZ ybeZ Y ybeZ aceE Y ybeZ aceF Y ybeZ b2710 Y ybeZ deaD Y ybeZ fabZ Y ybeZ hupA Y ybeZ pssA Y ybeZ rpIA Y ybeZ rpIB Y ybeZ rpIB Y ybeZ rpIE Y ybeZ rpIE Y ybeZ rpII Y ybeZ rpII Y ybeZ rpII Y Y ybeZ rpII Y Y ybeZ rpII Y Y ybeZ rpII Y Y ybeZ rpII Y Y Y ybeZ rpII Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	-	-		Y
ybeZ aceE Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	ybeY	yihK		
ybeZ aceF Y Y ybeZ b2710 Y ybeZ deaD Y ybeZ fabZ Y ybeZ hupA Y ybeZ pssA Y ybeZ rplA Y Y ybeZ rplB Y ybeZ rplC Y Y ybeZ rplE Y Y ybeZ rplE Y Y ybeZ rplF Y Y ybeZ rplJ Y ybeZ rplJ Y ybeZ rplU Y ybeZ rplU Y ybeZ rplU Y ybeZ rplD Y ybeZ rplU Y ybeZ rplU Y ybeZ rplU Y ybeZ rplU Y ybeZ rplO Y ybeZ rplU Y ybeZ rplO Y ybeZ rplU Y ybeZ rplU Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplO Y ybeZ rplY Y ybeZ rplY Y ybeZ rplY Y ybeZ rplX Y ybeZ rpmB Y ybeZ rpsA Y ybeZ rpsA Y ybeZ rpsB Y	-	-		Υ
ybeZ b2710 Y ybeZ deaD Y ybeZ hupA Y ybeZ lpdA Y ybeZ rplA Y Y ybeZ rplA Y Y ybeZ rplB Y Y ybeZ rplD Y Y ybeZ rplE Y Y ybeZ rplI Y Y ybeZ rplM Y Y ybeZ rplN Y Y ybeZ rplN Y Y ybeZ rplV Y Y ybeZ rplX Y Y ybeZ rpmB Y Y ybeZ rpsA Y ybeZ rpsA Y ybeZ rpsB Y	I -			V
ybeZ deaD Y ybeZ hupA Y ybeZ lpdA Y ybeZ pssA Y ybeZ rplA Y Y ybeZ rplB Y ybeZ rplD Y Y ybeZ rplE Y Y ybeZ rplF Y Y ybeZ rplI Y Y ybeZ rplI Y Y ybeZ rplI Y Y ybeZ rplI Y Y ybeZ rplI Y Y ybeZ rplI Y Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplI Y ybeZ rplY ybeZ rplV Y ybeZ rplV Y ybeZ rplY ybeZ rplY ybeZ rpmB ybeZ rpsA Y ybeZ rpsA Y ybeZ rpsB Y	•			ı
ybeZ hupA Y ybeZ pssA Y ybeZ rpIA Y Y ybeZ rpIB Y Y ybeZ rpIC Y Y ybeZ rpID Y Y ybeZ rpIE Y Y ybeZ rpIF Y Y ybeZ rpIJ Y Y ybeZ rpIM Y Y ybeZ rpIS Y Y ybeZ rpIV Y Y ybeZ rpIX Y Y ybeZ rpsA Y Y ybeZ rpsA Y Y ybeZ rpsB Y Y	_			
ybeZ pssA Y ybeZ rpIA Y Y ybeZ rpIB Y ybeZ rpIC Y Y ybeZ rpID Y Y ybeZ rpIE Y Y ybeZ rpIF Y Y ybeZ rpII Y Y ybeZ rpIJ Y ybeZ rpIJ Y ybeZ rpIM Y ybeZ rpIN Y ybeZ rpIN Y ybeZ rpIP Y ybeZ rpIP Y ybeZ rpIP Y ybeZ rpIV Y ybeZ rpIV Y ybeZ rpIX Y	_		Υ	
ybeZ rpsA Y ybeZ rplB Y ybeZ rplC Y Y ybeZ rplD Y Y ybeZ rplE Y Y ybeZ rplF Y Y ybeZ rplJ Y Y ybeZ rplM Y Y ybeZ rplO Y Y ybeZ rplS Y Y ybeZ rplV Y Y ybeZ rplX Y Y ybeZ rpmB Y Y ybeZ rpsA Y Y ybeZ rpsA Y Y	-	_	V	
ybeZ rpIA Y Y ybeZ rpIB Y Y ybeZ rpID Y Y ybeZ rpIE Y Y ybeZ rpIF Y Y ybeZ rpII Y Y ybeZ rpIM Y Y ybeZ rpIO Y Y ybeZ rpIF Y Y ybeZ rpIF Y Y ybeZ rpIV Y Y ybeZ rpIX Y Y ybeZ rpsA Y Y ybeZ rpsA Y Y	_	•		Y
ybeZ rpIB Y ybeZ rpIC Y Y ybeZ rpIE Y Y ybeZ rpIE Y Y ybeZ rpII Y Y ybeZ rpIM Y Y ybeZ rpIM Y Y ybeZ rpIP Y Y ybeZ rpIT Y Y ybeZ rpIV Y Y ybeZ rpIX Y Y ybeZ rpmB Y ybeZ rpsA Y ybeZ rpsB Y	-	•		Υ
ybeZ rpIC Y Y ybeZ rpIE Y Y ybeZ rpIF Y Y ybeZ rpII Y Y ybeZ rpIM Y Y ybeZ rpIM Y Y ybeZ rpIP Y Y ybeZ rpIT Y Y ybeZ rpIV Y Y ybeZ rpIX Y Y ybeZ rpmB Y Y ybeZ rpsA Y Y ybeZ rpsA Y Y	_	•		•
ybeZ rplE Y Y ybeZ rplF Y Y ybeZ rplJ Y Y ybeZ rplM Y Y ybeZ rplO Y Y ybeZ rplP Y Y ybeZ rplT Y Y ybeZ rplV Y Y ybeZ rplX Y Y ybeZ rpmB Y Y ybeZ rpsA Y Y ybeZ rpsA Y Y	-	_	Υ	Υ
ybeZ rpIF Y Y ybeZ rpII Y Y ybeZ rpIL Y Y ybeZ rpIM Y Y ybeZ rpIO Y Y ybeZ rpIF Y Y ybeZ rpIT Y Y ybeZ rpIV Y Y ybeZ rpIX Y Y ybeZ rpmB Y Y ybeZ rpsA Y Y ybeZ rpsB Y Y	_	-		
ybeZ rpll Y Y ybeZ rplJ Y ybeZ rplM Y Y ybeZ rplO Y ybeZ rplP Y ybeZ rplS Y Y ybeZ rplU Y Y ybeZ rplV Y Y ybeZ rplY Y Y ybeZ rpmB Y ybeZ rpsA Y ybeZ rpsB Y ybeZ rpsB Y	_	-		
ybeZ rplJ Y ybeZ rplM Y Y ybeZ rplO Y ybeZ rplP Y ybeZ rplS Y Y ybeZ rplT Y Y ybeZ rplV Y Y ybeZ rplX Y Y ybeZ rpmB Y Y ybeZ rpsA Y Y ybeZ rpsB Y Y	_	-		
ybeZ rplL Y ybeZ rplM Y Y ybeZ rplO Y ybeZ rplP Y ybeZ rplS Y Y ybeZ rplU Y Y ybeZ rplV Y Y ybeZ rplX Y Y ybeZ rpmB Y ybeZ rpsA Y ybeZ rpsB Y ybeZ rpsB Y		-		ĭ
ybeZ rpIM Y Y ybeZ rpIO Y ybeZ rpIF Y ybeZ rpIS Y ybeZ rpIT Y ybeZ rpIV Y ybeZ rpIX Y ybeZ rpIF Y ybeZ rpmB Y ybeZ rpsA Y ybeZ rpsB Y		•	•	Υ
ybeZ rpIP Y ybeZ rpIS Y Y ybeZ rpIT Y ybeZ rpIU Y Y ybeZ rpIX Y Y ybeZ rpIX Y Y ybeZ rpmB Y Y ybeZ rpsA Y Y ybeZ rpsB Y Y	•	-	Υ	
ybeZ rpIS Y Y ybeZ rpIT Y ybeZ rpIV Y Y ybeZ rpIX Y Y ybeZ rpIY Y Y ybeZ rpmB Y Y ybeZ rpsA Y Y ybeZ rpsB Y Y	_	-		Υ
ybeZ rplT Y ybeZ rplU Y ybeZ rplV Y Y ybeZ rplX Y Y ybeZ rpmB Y Y ybeZ rpsA Y Y ybeZ rpsB Y Y	_			
ybeZ rpIU Y ybeZ rpIV Y Y ybeZ rpIX Y Y ybeZ rpmB Y ybeZ rpmG Y ybeZ rpsA Y ybeZ rpsB Y ybeZ rpsB Y	_	-	Y	
ybeZ rpIV Y Y ybeZ rpIX Y Y ybeZ rpIY Y ybeZ rpmB Y ybeZ rpsA Y ybeZ rpsB Y ybeZ rpsB Y	_	-		
ybeZ rplX Y Y ybeZ rplY Y ybeZ rpmB Y ybeZ rpmG Y ybeZ rpsA Y ybeZ rpsB Y		-	Υ	
ybeZ rpmB Y ybeZ rpmG Y ybeZ rpsA Y ybeZ rpsB Y Y		_		
ybeZ rpmG Y ybeZ rpsA Y ybeZ rpsB Y Y	_	-		
ybeZ rpsA Y ybeZ rpsB Y Y	_	•		
ybeZ rpsB Y Y	_	-	V	Y
	-	-		V
, , ~ ~ · POO	ybeZ	rpsC	•	Y

ybeZ	rpsD	Υ	
ybeZ	rpsE	Y	Υ
ybeZ	rps⊑	Y	Y
ybeZ	rpsM	Ý	•
ybeZ	rpsN	•	Υ
ybeZ	rpsP		Ϋ́
ybeZ	rpsT		Y
ybeZ	secA	Υ	•
ybeZ	uvrD	•	Υ
ybeZ	vacB	Υ	
ybeZ	ybcJ		Υ
ybeZ	ycbY	Υ	
ybeZ	ycfB	Υ	
ybeZ	yciL	Υ	
ybeZ	yfcB	Υ	
ybeZ	yfiF	Υ	
ybeZ	yhbY		Υ
ybgC	ybgC	Y	
ybgC	b1410	Υ	
ybgC	rplB	Υ	
ybgC	rplC	Υ	
ybgC	rplW		Υ
ybgC	rpsD	Υ	
ybhF	ybhF	Υ	Y
ybhF	aceE	Υ	Υ
ybhF	aceF	Υ	
ybhK	ybhK	Y	Y
ybhK	aceE		Y
ybhK	b2247		Y
ybhK	basS		Y
ybhK	ccmH		Y
ybhK	ftsK		Y
ybhK	glgC		Y
ybhK	gsk idaT		Y Y
ybhK	idnT	V	Ϋ́
ybhK	lpdA	Y	Ϋ́
ybhK ybhK	narl rplC		Ϋ́
ybhK	rpID		Ϋ́
ybhK	rplM		Ϋ́
ybhK	rplU		Ϋ́
ybhK	rplV		Y
ybhK	rplW		Ϋ́
ybhK	rpoB	Υ	'
your	ipob	ı	

ybhK	rpsB		Υ
ybhK	rpsE		Υ
ybhK	rpsF		Υ
ybhK	rpsG		Υ
ybhK	secA		Υ
ybhK	valS		Υ
ybhK	ybhF		Υ
ybhK	yfdE		Υ
ybjX	ybjX	Y	
ybjX	aceF		Υ
ybjX	dnaJ		Υ
ybjX	hupA		Υ
ybjX	hupB		Y
ybjX	lpdA	Υ	•
ybjX	pssA	Ϋ́	
ybjX	rfaD	Ϋ́	
ybjX	rplA	Ϋ́	Υ
ybjX	rplB	Ϋ́	Ϋ́
ybjX	rplC	Ϋ́	Ϋ́
ybjX	rplD	Ϋ́	Ϋ́
ybjX	rplE	Ϋ́	Ϋ́
ybjX	rplF	Ϋ́	Ϋ́
ybjX	rpll	•	Ý
ybjX	rplJ		Ϋ́
ybjX	rplL		Ϋ́
ybjX	rplM	Υ	Ϋ́
ybjX	rplO	•	Ϋ́
ybjX	rplQ		Ϋ́
ybjX	rpIS		Y
ybjX	rpIT		Ϋ́
	rplU		Y
ybjX ybjX	rpIV	Υ	1
	rplX	1	Υ
ybjX			Y
ybjX	rpmB rpmC		Y
ybjX			Y
ybjX	rpmG	Υ	Ĭ
ybjX	rpoC	Y	V
ybjX	rpsA rpsB	Ϋ́Υ	Y Y
ybjX	rpsB	Υ Υ	
ybjX	rpsC		Υ
ybjX	rpsD	Υ	V
ybjX	rpsE		Y
ybjX	rpsF	V	Y
ybjX	rpsG	Υ	Υ

1			
ybjX	rpsM		Υ
ybjX	rpsS		Υ
ybjX	rpsT		Υ
ybjX	srmB	Υ	
ybjX	tufA	Υ	Υ
ybjX	tufB	Υ	
ybjX	vacB	Υ	
ybjX	yciL	Υ	
ycaJ	ycaJ	Υ	
ycaJ	cadB		Υ
ycaJ	napH		Υ
ycaJ	potC		Υ
ycaJ	rplL		Υ
ycaJ	yihN		Υ
ycaO	ycaO	Υ	Υ
ycaO	rpmC		Υ
ycaO	yjcW		Y
ycbL	ycbL	Υ	Y
ycbL	fliM	•	Ϋ́
ycbL	sucA		Y
ycbL	yahJ		Ý
ycbY	ycbY	Υ	•
ycbY	b1410	Ý	
ycbY	rplA	Y	
ycbY	yfiF	Ý	
yceA	yceA	Y	
yceA	aceF	Ý	
yceA	dnaK	Ý	
yceA	IpdA	Ϋ́	
yceA	rplL	•	Υ
yceA	yfjN		Ϋ́
yceC	yceC	Y	Y
yceC	accA	Ϋ́	,
yceC	aceE	Ϋ́	
yceC	aceF	Ϋ́	
yceC	aidB	Ý	
yceC	b1452	Ϋ́	Υ
yceC	deaD	Ϋ́	I
yceC	hrpA	Y	
yceC	hupA	•	Y
yceC	hupB		Y
yceC	IpdA	Υ	Y
-	-	Ϋ́	Y
yceC	pssA rbl⊏	Ϋ́	f
yceC	rhIE	Y	

1 _			
yceC	rplA	Υ	Y
yceC	rplB	Υ	Y
yceC	rpIC	Y	Y
yceC	rpID	Y	Y
yceC	rplE	Y	
yceC	rplF	Υ	Y
yceC	rpll	Υ	Y
yceC	rplL		Υ
yceC	rplM	Υ	Υ
yceC	rplN		Υ
yceC	rplO	Υ	Υ
yceC	rpIP		Υ
yceC	rplQ		Υ
yceC	rplR		Y
yceC	rpIS	Υ	Υ
yceC	rplT		Υ
yceC	rplU		Υ
yceC	rpIV	Υ	Υ
yceC	rplX	Υ	Υ
yceC	rpmB	Υ	
yceC	rpoB	Υ	
yceC	rpoC	Υ	
yceC	rpsA	Υ	
yceC	rpsB	Y	Y
yceC	rpsC	Y	Y
yceC	rpsD	Υ	
yceC	rpsE	Υ	Υ
yceC	rpsF		Υ
yceC	rpsG	Υ	Y
yceC	rpsM	Υ	Y
yceC	rpsN		Y
yceC	rpsP		Y
yceC	rpsR		Y
yceC	rpsT		Y
yceC	secA	Υ	
yceC	spoT	Υ	
yceC	srmB	Y	
yceC	vacB	Y	Υ
yceC	ybcJ		Υ
yceC	ybjD	Υ	
yceC	ycbY	Υ	Υ
yceC	yciL	Υ	Y
yceC	yfiF	Υ	Y
yceC	ygiF	Y	

	الماد	V	
yceC	yhiR	Y	V
yceC	yibL		Y Y
yceC	ymfC	Υ	I
yceH ycfB	yceH ycfB	Y	Υ
ycfB	dniR	Y	ı
ycfB	melR	Y	
ycfF	ycfF	Ϋ́	Υ
ycfF	b1501	,	Ϋ́
ycfF	dnaK	Y	
ycfF	dnaN	Ý	
ycfF	dniR	-	Υ
ycfF	tufA	Υ	-
ycfF	yjhH		Υ
ycfH	ycfH	Υ	Υ
ycfH	gcd		Υ
ycfH	tufB	Υ	
ycfX	ycfX	Y	Y
ycfX	yhdM	Υ	
ycgC	ycgC	Y	Υ
ycgC	b1372		Υ
ycgC	b2463	Υ	
ycgC	cysP		Υ
ycgC	dnaK	Υ	
ycgC	yjeH		Υ
ycgE	ycgE	Υ	
ycgE	hipA		Υ
ycgE	IpdA	Υ	
ychA	ychA	Y	
ychB	ychB	Y	Y
ychB	glmS		Y
ychJ	ychJ	Y	
ychJ	hyfB		Y
yciH	yciH		Y
ycil	ycil	V	Y
yciL	yciL	Y	Y
yciL	aceE	Y	
yciL	aidB	Υ	V
yciL	alaS bro A	Y	Y Y
yciL	hrpA	Ť	Υ Υ
yciL yciL	hupA	Υ	ſ
yciL	pnp pssA	Y	
yciL	rplA	Y	Υ
y CIL	i þiA	ſ	ſ

yciL yciL yciL yciL yciL yciL yciL yciL	rpIB rpIC rpID rpIE rpIF rpII rpIM rpIS rpIU rpIV	Y Y Y Y Y	Y Y Y Y Y Y
yciL yciL yciL yciL yciL yciL yciL yciL	rpIX rpIY rpmB rpsA rpsB rpsC rpsD rpsE	Y Y Y Y	Y Y Y Y
yciL yciL yciL yciL yciL yciL	rpsF rpsG rpsJ rpsM rpsN rpsP	Y Y Y	Y Y Y
yciL yciL yciL yciL yciL	rpsR rpsT rpsU srmB vacB	Y Y	Y Y Y Y Y
yciL yciL yciL yciL yciL	ybhF ycbY yfiF ygiF yhbP	Y Y	Y Y Y
yciL yciL yciL yciL yciL	yhbY yhiN yhiR yibL ymfC	Y	Y Y Y Y Y
yciO ycjC ydcP ydcP ydcP	yciO ycjC ydcP kbl rho	Y Y Y	Y Y Y

ydcP	rplA	Υ	Υ
ydcP	rpIC		Υ
ydcP	rpll		Υ
ydcP	rplL		Y
ydcP	rpIS		Ϋ́
ydcP	rplT		Ϋ́
ydcP	rplV		Ϋ́
ydcP	rplX		Ϋ́
ydcP	rpmG		Ϋ́
ydcP	rpsA		Ϋ́
ydcP	rpsC		Ϋ́
ydcP	rpsE		Ϋ́
ydcP	rpsG		Ϋ́
ydcP	rpsN		Ϋ́
-	selB		Ϋ́
ydcP ydcP	tufA	Υ	ī
_		Ϋ́	Υ
ydhD	ydhD	7	
ydhD	bolA	V	Y Y
ydiA	ydiA	Y	7
ydiA	aceE	Y	
ydiA	pepB	Y	
ydiA	sfhB	Y	Υ
ydiA	tufA	Y	
ydiA	tufB	Y	
ydiB	ydiB	Y	
ydiB	yggG	2.2	Y
yeaA	yeaA	Υ	Υ
yeaA	yecK		Υ
yeaG	yeaG	Υ	Y
yeaG	dgt	Υ	
yeaG	dnaN	Υ	
yeaU	yeaU	Y	Υ
yeaU	tufA	Υ	
yeaZ	yeaZ	Y	Υ
yeaZ	aroK	Υ	
yeaZ	ygjD	Υ	
yebC	yebC	Y	Y
yebC	argS		Υ
yebC	goaG		Υ
yebC	hdeD		Υ
yebC	sseA		Υ
yebK	yebK	Y	
yebK	b1579	Υ	
yebK	hflB	Υ	
•			

yebM	yebM	Y	
yebM	aceE	Y	
yebM	aceF	Υ	
yebM	b1410	Υ	
yebM	lpdA	Υ	
yebM	mopA	Υ	
yebM	rpID	Υ	
yebM	rplL		Y
yebM	rpIV		Y
yebM	rpmG	V	Υ
yebM	rpsE	Y Y	Υ
yedO	<i>yedO</i> rplA	Ϋ́	Y
yedO yedO	rplC	Ϋ́	
yedO	rpID	Ϋ́	
yedO	rpsC	Ϋ́	
yedO	rpsD	Ϋ́	
yedO	rpsE	Y	
yedO	sucB	Υ	Υ
yedO	ygiP		Υ
yedW	yedW	Y	
yedW	b1410	Υ	
yeiP	yeiP	Υ	Υ
yejF	yejF	Υ	
yejF	dnaE		Y
yejF	lpdA	Υ	V
yejF	oppD		Y Y
yejF	rpIC tufB	Υ	Y
yejF	yfcB	Ĭ	Υ
yejF <i>yfaO</i>	yfaO	Y	Y
yfcB	yfcB	Y	Ϋ́
yfcB	b1372	Ϋ́	•
yfcB	dnaK	Υ	
yfcB	rplC	Υ	Υ
yfcB	rpsJ		Υ
yfcB	tig	Υ	Υ
yfcB	yncC		Υ
yfgB	yfgB	Y	Υ
yfgB	aas	Y	
yfgB	aceE	Y	
yfgB	aceF	Y Y	
yfgB	clpA	Y Y	
yfgB	clpB	Ĭ	

yfgB	dapF		Υ
yfgB	deaD	Υ	
yfgB	dnaJ	Υ	Υ
yfgB	elaB		Y
yfgB	fruR	Υ	•
yfgB	fusA	Ϋ́	
yfgB	gatY	Ϋ́	
	gyrB	Ϋ́	
yfgB vfaB		'	Υ
yfgB	hlpA bro A	Υ	1
yfgB	hrpA	Ţ	V
yfgB	hupA	V	Υ
yfgB	infB	Υ	
yfgB	infC		Υ
yfgB	lon	Υ	
yfgB	lpdA		Y
yfgB	lysS		Y Y Y
yfgB	lysU		Υ
yfgB	malT	Υ	
yfgB	melB		Y
yfgB	metK	Υ	
yfgB	mreB	Υ	
yfgB	nuoC	Υ	
yfgB	ompC	Υ	
yfgB	phoE	Υ	
yfgB	phoR		Υ
yfgB	pnp	Υ	
yfgB	rcsB		Υ
yfgB	recA	Υ	
yfgB	rne	Υ	
yfgB	rplA	Υ	Υ
yfgB	rplB	Υ	Υ
yfgB	rpIC	Υ	Υ
yfgB	rplD	Υ	Υ
yfgB	rplE	Y	Y
yfgB	rplF	-	Y
yfgB	rpll	Υ	Y
yfgB	rplL	-	Y
yfgB	rplM	Υ	Y
yfgB	rplO	Ϋ́	Ϋ́
yfgB	rplR	-	Ϋ́
yfgB	rpIS	Υ	Ϋ́
yfgB	rplT	-	Ϋ́
yfgB	rplU		Ϋ́
yfgB	rplV	Υ	Ϋ́
yigo	ipiv	•	•

yfgB	rplW		Y
yfgB	rpIX	Υ	Y
yfgB	rpmA		Υ
yfgB	rpmB		Y
yfgB	rpmC		Y
yfgB	rpoC	Υ	
yfgB	rpsA	Υ	Y
yfgB	rpsB	Υ	Y
yfgB	rpsC	Υ	Y
yfgB	rpsD	Υ	
yfgB	rpsE	Υ	Υ
yfgB	rpsF		Υ
yfgB	rpsG	Υ	Υ
yfgB	rpsJ	Υ	Υ
yfgB	rpsN		Y
yfgB	rpsP		Υ
yfgB	rpsR		Υ
yfgB	secA	Υ	Υ
yfgB	selB	Υ	Υ
yfgB	spoT	Υ	
yfgB	srmB	Υ	Y
yfgB	tsr		Υ
yfgB	tufA	Υ	Υ
yfgB	tufB	Υ	
yfgB	vacB	Υ	Υ
yfgB	wcaD		Υ
yfgB	yaiU		Υ
yfgB	ycbY	Υ	
yfgB	ycfF		Υ
yfgB	yciL	Υ	Y
yfgB	ydcP	Υ	
yfgB	yeeX		Y
yfgB	yeiJ	3.6	Y
yfgB	yfiF	Υ	Y
yfgB	ygaM . <u>-</u>	3.6	Υ
yfgB	ygiF	Υ	
yfgB	yhbY		Y
yfgB	yhiR	Υ	Y
yfgB	yibL	V	Y
yfhA	yfhA	Y	V
yfhE	yfhE	Y	Y
yfhE	clpB htpC	Υ	V
yfhE	htpG		Y
yfhE	rplA		Υ

yfhE	rplJ		Y
yfhE	rplK		Ϋ́
yfhE	rplL		Ϋ́
yfhE	rpmC		Ϋ́
yfhE	rpmG		Ý
yfhE	rpoC	Υ	•
yfhE	rpsB	·	Υ
yfhE	rpsG		Y
yfhE	rpsN		Υ
yfhE	tufA		Υ
yfhE	tufB	Υ	
yfhE	ugpB		Υ
yfhE	yhcL		Υ
yfiA	yfiA		Y
yfiD	yfiD	Y	Y
yfiD	b1696		Υ
yfiD	lysU	Υ	
yfiD	parE	Υ	
yfiD	pflB	Υ	Y
yfiD	rpsT		Υ
yfiD	tdcE	Υ	
yfiD	tpiA	Υ	
yfiF	yfiF	Υ	Y
yfiF	b2520		Υ
yfiF	dacA		Υ
yfiF	deaD	Υ	
yfiF	holB	Υ	
yfiF	hupA		Y
yfiF	pssA	Y	
yfiF	rplA	Y	Y
yfiF	rplB	Y	Y
yfiF	rpIC	Y	Y
yfiF	rpIF	Y	
yfiF	rpll	Υ	V
yfiF	rpIP		Y
yfiF	rpIR		Y Y
yfiF	rplT rpl\/		Ϋ́
yfiF yfiF	rpIV rpsC	Υ	ĭ
yfiF	rpsC	Ϋ́	Υ
yfiF	rpsE	Ϋ́	Ϋ́
yfiF	rpsE	Ϋ́	Ϋ́
yfiF	rpsG	Ϋ́	Ϋ́
yfiF	rpsH	•	Ϋ́
, <i>,</i>			•

yfiF	rpsK		Υ
yfiF	rpsL	V	Y Y
yfiF yfiF	rpsM rpsN	Υ	Ϋ́Υ
yfiF	rpsO		Ϋ́
yfiF	rpsP		Υ
yfiF	rpsR		Υ
yfiF	rpsS		Y
yfiF	rpsT		Y
yfiF yfiF	rpsU vacB	Υ	Y
yfiF	ycbY	Ϋ́	
yfiF	yceC	Y	
yfiF	yciL	Y	
yfiF	ynhD		Υ
yfiQ	yfiQ	Υ	V
yfiQ yfiQ	b2434 dnaJ		Y Y
yfiQ	hupB		Ϋ́
yfiQ	hypC		Ϋ́
yfiQ	lpdA		Υ
yfiQ	mopA		Υ
yfiQ	rplC		Y
yfiQ	rplM		Y
yfiQ yfiQ	rplO rplP		Y Y
yfiQ	rplQ		Ϋ́
yfiQ	rplV		Y
yfiQ	rpmB		Υ
yfiQ	rpsB		Y
yfiQ	rpsE		Y
yfiQ yfiQ	rpsJ rpsM		Y Y
yfiQ	rpsN		Ϋ́
yfiQ	rpsS		Υ
yfiQ	rpsT		Υ
yfiQ	tufA		Y
yfiQ	yjfQ vfiB	V	Υ
yfjB yfjK	yfjB yfjK	Y Y	
yijiK yfjK	dnaK	,	Υ
yfjK	gapA		Ϋ́
yfjK	mopA		Υ
yfjK	rplC		Υ

£:17	II		V
yfjK	rplL		Y
yfjK	rplW		Y
yfjK	tufA	V	Y Y
ygaG	<i>ygaG</i> dnaK	Y Y	Y
ygaG ygbB	ygbB	Y	
<i>ygbB</i> ygbB	flgD	1	Υ
ygbB	rfaD	Υ	'
ygbB	rpsJ	•	Υ
ygbM	ygbM	Y	•
ygcM	удсМ	Ϋ́	
ygcP	ygcP	,	Υ
ygcP	ccmH		Ϋ́
ygcP	dnaK		Y
ygcP	rplC		Y
ygcP	rplD		Ϋ́
ygcP	rplU		Υ
ygcP	rpIV		Υ
ygcP	rpmG		Υ
ygcP	rpoA		Υ
ygcP	rpoC		Υ
ygcP	rpsD		Υ
ygcP	ybjO		Υ
ygcW	ygcW	Y	
ygdP	ygdP	Y	Y
ygdP	dapF	Υ	Υ
ygdP	deaD	Υ	
ygdP	hupA		Υ
ygdP	rplA	Υ	Υ
ygdP	rplB	Υ	Y
ygdP	rplC	Y	Y
ygdP	rpID	Y	Υ
ygdP	rplE	Y	V
ygdP	rplF	Y	Y
ygdP	rpll	Y Y	Y
ygdP	rplJ rplM	Ϋ́	Υ
ygdP	rpIN	Y	I
ygdP ygdP	rplO	Ϋ́	Υ
ygdP	rpIP	•	Ϋ́
ygdP	rplQ		Ϋ́
ygdP	rplR	Y	Ϋ́
ygdP	rpIS	Ý	Ϋ́
ygdP	rplT		Y
ygur	ipii		ı

ygdP	rplU		Y
ygdP	rplV	Υ	Y
ygdP	rpIX	•	Ϋ́
ygdP	rplY		Ϋ́
ygdP	rpmB		Ϋ́
ygdP	rpsA	Υ	Ϋ́
ygdP	rpsB	Ý	Ϋ́
ygdP	rpsC	•	Ϋ́
ygdP	rpsD	Υ	•
ygdP	rpsE	Ý	Υ
ygdP	rpsF	•	Ϋ́
ygdP	rpsG	Υ	Ϋ́
ygdP	rpsM	Ϋ́	Ϋ́
ygdP	rpsR	'	Ϋ́
ygdP	rpsS		Ϋ́
ygdP	rpsT		Ϋ́
ygdP	spoT	Υ	•
ygdP	srmB	Ý	
ygdP	vacB	Ϋ́	
ygdP	yajQ	Ϋ́	
ygdP	ycbY	Ϋ́	
ygdP	yceC	Ϋ́	
ygdP	yciL	-	Υ
ygdP	yfiF	Υ	Y
ygeV	ygeV	Υ	
ygeV	lon	Υ	
ygeV	lpdA	Υ	
ygeV	tufA	Υ	
ygeV	tufB	Υ	
ygfA	ygfA	Υ	Y
ygfA	b2146		Υ
ygfA	galM		Υ
ygfA	iscU		Υ
ygfA	mopA		Υ
ygfA	ndk	Y	Υ
ygfA	pqiB		Υ
ygfA	rpsB		Υ
ygfA	ybiN		Υ
ygfF	ygfF	Υ	
ygfF	lpdA	Υ	
yggH	yggH	Υ	Υ
yggH	aceE	Υ	
yggH	b1410	Y	
yggH	b1451	Y	

1			
yggH	deaD	Y	
yggH	dnaK	Υ	
yggH	glnS	Υ	
yggH	hrpA	Υ	
yggH	lpdA	Υ	
yggH	mglB	Υ	
yggH	pssA	Υ	
yggH	rplB	Υ	
yggH	rpIC	Υ	
yggH	rpID	Υ	
yggH	rpIE	Y	
yggH	rplF	Y	
yggH	rpll	Υ	
yggH	rplJ	Υ	
yggH	rplM	Υ	
yggH	rpIS	Υ	
yggH	rpIV	Υ	Υ
yggH	rpIX	Υ	
yggH	rpsA	Υ	
yggH	rpsB	Υ	
yggH	rpsC	Υ	
yggH	rpsD	Υ	
yggH	rpsE	Υ	
yggH	rpsF		Υ
yggH	rpsG	Υ	Y
yggH	rpsM	Y	-
yggH	rpsP	Y	
yggH	rpsT	-	Υ
yggH	rpsU	Υ	•
yggH	srmB	Ý	
yggH	ybeZ	Ϋ́	
yggH	ycbY	Ϋ́	
yggH	yfiF	Ý	
yggH	ygiF	Ý	
yggS	yggS	Y	
yggS	fdnG	Ϋ́	
yggS	rpsJ	Ϋ́	
yggV	yggV	Y	Υ
yggW	ygg v yggW	Y	•
		Y	Υ
yggX vah∆	yggX yghA	Y	Y
yghA	<i>ygriA</i> dnaK	Ϋ́	I
yghA		Ϋ́Υ	
yghA	dnaN		
ygiC	ygiC	Y	

ygiF	ygiF	Υ	Υ
ygiF	aceE	Υ	
ygiF	aceF	Y	.
ygiF	ompA	Y	Υ
ygiF	ompC	Y Y	
ygiF	phoE rpmB	Y	Y
ygiF	ygjD	Υ	Y
<i>ygjD</i> ygjD	aceE	Ϋ́	I
ygjD	hybG	ı	Y
ygjD	rplL		Ϋ́
ygjD	rpoA		Ϋ́
ygjD	rpoB		Ϋ́
ygjD	rpsJ		Ϋ́
ygjD	tufA	Υ	-
ygjD	tufB	Y	
ygjD	yeaZ	Υ	
ygjH	ygjH	Y	
yhaD	yhaD	Υ	Y
yhaR	yhaR	Υ	
yhbC	yhbC	Υ	Y
yhbC	rpsE	Υ	
yhbC	rpsG	Υ	
yhbC	yihl		Υ
yhbH	yhbH	Y	Υ
yhbH	dnaN	Υ	
yhbJ	yhbJ	Υ	Y
yhbJ	acpP		Υ
yhbJ	cspC		Υ
yhbJ	mopA	Υ	
yhbJ	rplV		Υ
yhbJ	rpsN		Υ
yhbJ	tufA	Y	
yhbU	yhbU	Y	
yhbU	aceE	Y	
yhbU	aceF	Y	
yhbU	lon	Y	
yhbU	lpdA	Y	V
yhbU	mopA	Y	Y
yhbU	tufA	Y Y	
yhbU	tufB	Ϋ́Υ	
yhbU yhbV	yhbV yhbV	Y	
yhbY	yhbY	ı	Υ
yiio i	yiioi		I

yhbY	rplB	Υ	
yhbY	rplC	Υ	
yhbY	rpID	Υ	
yhbY	rplK	Υ	
yhbY	rplM	Υ	
yhbY	rplT		Υ
yhbY	rplU		Y
yhbY	rpsB	Υ	-
yhbY	rpsG	Ϋ́	
yhbY	yceC	Y	
yhbY	yciL	Ϋ́	
yhbY	yfiF	Ϋ́	
yhbY	zntA	•	Υ
yhbZ	yhbZ	Y	Y Y
yhbZ	b1685	,	Ϋ́
yhbZ	b2097	Υ	•
yhbZ	pepE	•	Υ
yhbZ	prIC		Ϋ́
yhbZ	rplL		Ϋ́
yhbZ	rpoA		Ϋ́
_	sfhB	Υ	ı
yhbZ		Ϋ́	
yhcC	yhcC	Ϋ́	
yhcC	alkB		
yhcC	phoE	Y	
yhcC	yhcQ	Y	V
yhcJ	yhcJ	Y	Y
yhcJ	yedO	V	Y
yheS	yheS	Υ	Y
yheS	sucA	V	Y
yheS	tufA	Y	Y
yhfR	yhfR	Υ	V
yhfR	topB	V	Y
yhhF	yhhF	Υ	V
yhhF	dnaK	V	Y
yhhF	tufA	Y	
yhhG	yhhG	Υ	Y
yhhG	rplL		Υ
yhhG	tufA	Y	\ <u>/</u>
yhhP	yhhP	Y	Υ
yhhP	aceE	Υ	3.7
yhhP	gapA		Y
yhhP	hlpA	3.4	Y
yhhP	iscS	Υ	3.7
yhhP	rplL		Y

yhiF rpsD Y yhiR yhiR Y yhiR aceE Y yhiR mopA Y yhiR yihI Y yiaE yiaE Y yiaE pgm Y yiaJ yiaJ Y yibA yibA Y yibK yibK Y yibL yibL Y yibL rpIA Y yibL rpIC Y yibL rpIM Y
yhiR aceE Y yhiR mopA Y yhiR yihl Y yiaE yiaE Y Y yiaE pgm Y yiaJ yiaJ Y yibA yibA Y yibK yibK Y yibL yibL Y Y yibL aceF yibL rplA Y yibL rplC Y
yhiR yihI Y yiaE yiaE Y Y yiaE pgm Y yiaJ yiaJ Y yibA yibA Y yibK yibK Y yibL yibL Y yibL rplA Y yibL rplC Y
yiaE yiaE Y Y yiaE pgm Y yiaJ yiaJ Y yibA yibA Y yibK yibK Y yibL yibL Y yibL rplA Y yibL rplC Y
yiaE pgm Y yiaJ yibA Y yibA yibA Y yibK yibK Y yibL yibL Y yibL rplA Y yibL rplC Y
yiaJ yiaJ Y yibA yibA Y yibK yibK Y yibL yibL Y yibL rpIA Y yibL rpIC Y
yibAyibAYyibKyibKYyibLyibLYyibLrpIAYyibLrpICY
yibKyibKYyibLyibLYyibLrpIAYyibLrpICY
yibLyibLYyibLrplAYyibLrplCY
yibL aceF Y yibL rpIA Y yibL rpIC Y
yibL rpIA Y yibL rpIC Y
yibL rplC Y
vibl rnIM V
, ,
yibL rpIR Y
yibL rplS Y
yibL rplT Y
yibL rplV Y yibL rpmB Y
yibL rpmB Y yibL rpsB Y
yibL rpsE Y
yibL rpsG Y
yibL rpsN Y
yibL ybcJ Y
yibL yfiF Y
yibL yhbY Y
yibL ymfC Y
yicC yicC Y Y
yicC dnaJ Y yicC dnaK Y
yicC dnaK Y yicC lpdA Y
yicC ipuA i yicC rplW Y

yicC	rpoB	Υ	
-	-	•	V
yicC	rpsB		Y
yicC	rpsP		Υ
yicC	tufA	Υ	
yicC	tufB	Υ	
yicC	yidY		Υ
yidA	yidA	Υ	Υ
yidA	dnaK	Υ	
yidA	mopA	Ϋ́	
yidA	tufB	Ϋ́	
•		Y	
yieM	yieM		
yieM	tufA	Y	
yieM	yniC	Y	
yieN	yieN	Υ	Y
yieN	accA	Υ	
yieN	accB		Υ
yieN	accC		Υ
yieN	b1579	Υ	
yieN	cadA	Y	Υ
yieN	dnaK	Ϋ́	Y
yieN	frdA	Ϋ́	•
		Ϋ́	
yieN	fucU	Y	V
yieN	ggt_		Y
yieN	grpE		Y
yieN	infB		Υ
yieN	IdcC	Υ	
yieN	lpdA		Υ
yieN	rplJ	Υ	
yieN	rpIL		Υ
yieN	rpsB	Υ	
yieN	rpsP		Υ
yieN	tufB	Υ	-
yigW_2		Ϋ́	
yigZ	yigZ	Ϋ́	
		,	Υ
yigZ	topA	V	ı
yigZ	yihK	Y	
yihA	yihA	Y	
yihl	yihl	Y	
yihl	rplB	Υ	
yihl	rpIC	Υ	
yihl	rplD	Υ	
yihl	secA	Υ	
yihl	spoT	Υ	
yihl	ycbY	Ϋ́	
, ,	,	•	

yihK	yihK	Y	Y
yihQ	yihQ	Υ	
yihW	yihW	Υ	
yihZ	yihZ	Υ	
yihZ	cca	Υ	
yihZ	rplC		Υ
yihZ	rplV		Υ
yihZ	yihl		Υ
yiiD	yiiD	Y	Υ
yiiD	acpP		Υ
yiiD	dnaK		Υ
yiiD	gadB		Υ
yiiD	gapA		Υ
yiiD	prkB	Υ	
yiiD	rpIL		Υ
yiiD	yfjW		Υ
yiiD	yleB	Υ	
yjaD	yjaD		Υ
yjbJ	yjbJ	Y	Υ
yjdG	yjdG	Υ	
yjeA	yjeA	Y	Υ
yjeA	aceE		Υ
yjeE	yjeE	Υ	Y
yjeE	dnaK		Y
yjeE	gadB		Y
yjeE	gapA		Y
yjeE	metE		Υ
yjeE	mopA		Y
yjeE	pflB		Y
yjeE	purA		Ϋ́
yjeE	rplL		Ϋ́
yjeE	rpsA		Ϋ́
yjeE	yiiD	Υ	·
yjeE	yjeF	Ϋ́	
yjeF	yjeF	Y	Y
yjeF	aceF	Ϋ́	Ϋ́
yjeF	lpdA	Ϋ́	Ϋ́
yjeF	yabB	Ϋ́	Ý
yjeF	yjeE	Ϋ́	-
yjeQ	yjeQ yjeQ	Y	Υ
yjeQ	aceE	•	Ϋ́
yjeQ	b2451		Ϋ́
yjeQ	dnaK		Ϋ́
yjeQ	fusA		Ϋ́
7,100	.40/ (1

yjeQ	gapA		Υ
yjeQ	mopA		Υ
yjeQ	pabB		Υ
yjeQ	rplC		Υ
yjeQ	rpID		Υ
yjeQ	rplL		Υ
yjeQ	rplM		Υ
yjeQ	rpIV		Υ
yjeQ	rplW		Υ
yjeQ	rpmG		Υ
yjeQ	rpsA		Υ
yjeQ	rpsB		Υ
yjeQ	rpsC		Υ
yjeQ	rpsG		Υ
yjeQ	rpsJ		Υ
yjeQ	rpsM		Y
yjeQ	rpsN		Y
yjeQ	rpsP		Y
yjeQ	tufA		Y
yjeQ	wcaJ		Y
yjeQ	yaiW		Y
yjeQ	ybaL		Y
yjeQ	ybeR		Y
yjeQ	ydaY		Y
yjeQ	ydiA		Y
yjfH	yjfH	Y	Υ
yjfH	b0703	Υ	
yjfH	rplC		Y
yjfH	rplF		Y
yjfH	rpll		Y
yjfH	rplM		Y
yjfH	rpIR		Y
yjfH	rpIS		Y
yjfH	rpIV		Y
yjfH	rpmB		Y
yjfH	rpmG		Y
yjfH	rpsB		Y
yjfH	rpsE		Y
yjfH	rpsG		Y
yjfH	rpsJ		Y Y
yjfH	rpsU	V	Y
yjfH	yjeE vioE	Y	V
yjfH vifO	yjeF vifO	Υ	Y Y
yjfQ	yjfQ	γ	γ

vif○	aceF	Y	
yjfQ yjfQ	b1410	Y	
yjfQ	b2494		Υ
yjfQ	b2520		Ϋ́
yjfQ	dnaJ		Y
yjfQ	dnaK	Υ	
yjfQ	hupB		Υ
yjfQ	IpdA		Υ
yjfQ	lysR		Υ
yjfQ	mopA	Υ	Υ
yjfQ	rplA	Y	
yjfQ	rpIC	Υ	Y
yjfQ	rpID		Y
yjfQ	rplL		Y
yjfQ	rplM		Y
yjfQ vifQ	rpIN rpIS		Y Y
yjfQ yjfQ	rpIS rpIU		Y
yjfQ	rpIV		Y
yjfQ	rpmB		Ϋ́
yjfQ	rpsA	Υ	•
yjfQ	rpsB	•	Υ
yjfQ	rpsC		Y
yjfQ	rpsD	Υ	Υ
yjfQ	rpsE	Υ	Υ
yjfQ	rpsG		Υ
yjfQ	rpsH		Υ
yjfQ	rpsl		Υ
yjfQ	rpsJ		Υ
yjfQ	rpsL		Y
yjfQ	rpsM	Υ	Y
yjfQ	rpsN		Y
yjfQ	rpsS		Y
yjfQ	rpsT	V	Y Y
yjfQ yjfQ	tufA tufB	Y Y	Y
yjfQ yjfQ	yagX	ĭ	Y
yjfQ	ybgH		Y
yjfQ	ybgri yhaJ		Ϋ́
yjgD	yjgD		Y
yjgD	aceF	Υ	•
yjgD	gshA	·	Υ
yjgD	guaC		Υ
yjgD	lpdA	Υ	

r			
yjgD	yhfM		Υ
yjgD	yjfG		Υ
yjgD	yniC	Υ	
yjgH	yjgH	Y	Y
yjgH	pta	Υ	
yjgL	yjgL	Υ	
yjgL	rplW	-	Υ
yjhG	yjhG	Y	Y
yjhH	yjn:C yjhH	Ý	,
ујјі	ујін і УјјІ	Y	Y
	יעל dnaK	,	Ϋ́
yjjl vii\/		Υ	1
yjjV viiv	yjj∨		
yjjV	tufA	Y	
yleA	yleA	Y	Υ
yleA	dnaJ	Υ	
yleA	dnaK		Υ
yleA	mreB	Υ	
yleA	pstB	Υ	
yleA	tufA		Υ
yleA	tufB	Υ	
yleB	yleB	Y	Y
yleB	ydaC		Υ
yncC	yncC	Y	
yncC	rplC	Υ	
ynhC	ynhC	Y	
ynhC	ynhE	Υ	
ynhD	ynhD	Y	
ynhD	rplA	Υ	
ynhD	ynhC	Y	
ynhD	ynhE	Ϋ́	
ynhE	ynhE	Y	Y
ynnE ynhE	ynhC	Ý	Ϋ́
ynhE	ynhD	Ý	•
yohl	yohl	Y	Y
yohl	aceE	1	Y
_			Y
yohl	ampE		
yohl	cynT		Y
yohl	entE		Y
yohl	fumC		Y
yohl	hupA		Y
yohl	intA		Y
yohl	lpdA		Y
yohl	nfo		Y
yohl	recG		Υ

yohl yohl yohl yohl yohl yohl yohl yohl	rpIM rpIN rpIR rpIV rpsB rpsE rpsG rpsJ secA ydaY ydiB yfjW yqaB dnaK	Y Y Y	Y Y Y Y Y Y Y
yqaB yqaB	mopA tufA	Υ	
yqeA	yqeA	Y	
yqgF	yqgF	Y	
yqiE	yqiE	Y	
yqiE	sgbH	Y Y	Υ
<i>yraL</i> yraL	<i>yraL</i> aceF	Ϋ́	Y
yraL yraM	yraM	Y	
yraM	hycG	,	Υ
yraN	yraN		Y
yraN	aceE		Ϋ́
yraN	aceF		Y
yraN	hupB		Υ
yraN	IpdA		Υ
yraN	rplM		Υ
yraN	rpmB		Υ
yrbF	yrbF	Υ	Y
yrbF	dnaK	Υ	
yrbF	hybC		Y
yrbF	mopA	Υ	Y
yrbF	rplL	V	Y
yrbF	tufA	Y Y	Y
<i>yrdC</i> yrdC	<i>yrdC</i> rplK	7	Y Y
yrdC	rpIV		Ϋ́
yrdC	rplW		Ϋ́
yrdC	rpsB		Ϋ́
yrdC	rpsE		Ϋ́
yrdC	rpsJ		Υ

yrdC	rpsP		Υ
yrdD	yrdD	Υ	
yrfE	yrfE	Y Y	
yrfH	yrfH	Υ	Y
yrfH	rpll		Υ
yrfH	rplM		Υ
yrfH	rplN		Υ
yrfH	rpIS		Υ
yrfH	rplV		Υ
yrfH	rpmB		Υ
yrfH	rpsB	Υ	Υ
yrfH	rpsC		Υ
yrfH	rpsE		Υ
yrfH	rpsG		Υ
yrfH	rpsJ		Υ
yrfH	sbcC		Υ
yrfH	yagG		Υ
yrfH	ycfS		Υ
yrfH	ydhA		Υ
yrfl	yrfl	Υ	Y
yrfl	arcB		Υ
yrfl	nikD		Υ
yrfl	rplC		Υ
yrfl	rpsB		Υ
yrfl	rpsC		Υ
yrfl	rpsE		Υ
yrfl	rpsG		Υ
yrfl	rpsJ		Υ
yrfl	rpsL		Y
yrfl	tufA		Υ
yrfl	tufB	Y	
zwf	zwf	Y	Υ
zwf	dnaN	Υ	

Supplementary Table 2. Functional annotation and experimental information for all bait genes reported in this study

<i>Bait</i> aas	Annotation bifunctional: 2-acylglycerophospho-ethanolamine acyl transferase (N-terminal); acyl-acyl carrier protein	Essei	ntial Tagg	ed Purified
	synthetase (C-terminal)			
abgB	putative peptidase, p-aminobenzoyl-glutamate utilization, with Zn-dependent exopeptidase domain and carboxypeptidase G2, dimerisation domain		Y	Υ
iccA	acetylCoA carboxylase, carboxytransferase subunit alpha	Υ	Υ	Υ
ассВ	acetyl-CoA carboxylase, biotin carboxyl carrier protein subunit	Y		
accC	acetyl CoA carboxylase, biotin carboxylase subunit	Y	Y	Y
accD acpD	acetylCoA carboxylase, carboxyltranferase subunit beta NADH-azoreductase, FMN-dependent	Y	Y Y	Y
асрВ	acyl carrier protein	Υ	Ϋ́	Υ
acpS	holo-[acyl-carrier-protein] synthase (CoA:apo-[acyl-carrier-protein] pantetheinephosphotransferase)	Y	Y	-
ada add	bifunctional: transcriptional regulator of DNA repair (N-terminal); O6-methylguanine-DNA methyltransferase adenosine deaminase		Υ	Υ
adhC	alcohol dehydrogenase class III		Υ	Υ
adhE	multifunctional: acetaldehyde-CoA dehydrogenase (N-terminal); iron-dependent alcohol dehydrogenase (C-terminal); pyruvate-formate lyase deactivase		Y	Y
adiA	arginine decarboxylase, inducible by acid, catabolic		Υ	Υ
adk	adenylate kinase	Υ	Y	Y
agaA	putative N-acetylglucosamine-6-phosphate deacetylase			
agaY	tagatose 6-phosphate aldolase 1, subunit together with AgaZ		Υ	Υ
ahpC	alkyl hydroperoxide reductase, C22 subunit, thioredoxin-like, detoxification of hydroperoxides		Y	Y
aidB	putative acyl-CoA dehydrogenase (flavoprotein) , adaptive response (transcription activated by Ada)	v	Y Y	Y
alaS alkA	alanyl-tRNA synthetase 3-methyl-adenine DNA glycosylase II, inducible	Υ	Y Y	Y Y
alkB	oxidative demethylase of N1-methyladenine or N3-methylcytosine DNA lesions, repair of alkylated DNA		Ϋ́	•
allB	allantoinase	Υ	Ϋ́	
allR	transcriptional repressor of allantoin metabolism (IcIR family)		Υ	Υ
alsA	allose transport protein (ABC superfamily, atp_bind)		Υ	
alsE	putative hexose phosphate epimerase with ribulose-phoshate binding barrel		Y	
apt	adenine phosphoribosyltransferase	Υ	Y	Y
araC argB	transcriptional regulator of arabinose catabolism (AraC/XylS family) acetylglutamate kinase		Y Y	Υ
argR	transcriptional repressor of arginine synthesis (ArgR familiy)		Ϋ́	Ϋ́
argS	arginine tRNA synthetase	Υ	•	•
arnA	UDP-D-glucuronate dehydrogenase		Υ	Υ
aroB	dehydroquinate synthase	Υ	Υ	
aroE	dehydroshikimate reductase, NAD(P)-binding		Υ	Υ
artP	arginine transport protein (ABC superfamily, atp_bind)	v	Y	Y
asnS aspA	asparagine tRNA synthetase	Y	Y Y	Y Y
aspA aspP	aspartate ammonia-lyase (aspartase) adenosine diphosphate sugar pyrophosphatase (ADP-ribose pyrophosphatase)		Ϋ́	Ϋ́
aspS	aspartate tRNA synthetase	Υ	Y	Ϋ́
astD	succinylglutamic semialdehyde dehydrogenase		Υ	Υ
atpA	membrane-bound ATP synthase, F1 sector, alpha-subunit		Υ	Υ
atpC	membrane-bound ATP synthase, F1 sector, epsilon-subunit		Y	
atpD	membrane-bound ATP synthase, F1 sector, beta-subunit		Y Y	Υ
atpG atpH	membrane-bound ATP synthase, F1 sector, gamma-subunit membrane-bound ATP synthase, F1 sector, delta-subunit		Ϋ́	T
b0100			•	
	unknown CDS			
0165	unknown CDS			
	unknown CDS			
	CP4-6 prophage; IS911, putative transposase unknown CDS		Y	
00302	unknown CDS unknown CDS			
	unknown CDS			
00359	putative acyl transferase with trimeric LpxA-like domain		Υ	
	unknown CDS			
	unknown CDS			
	conserved protein unknown CDS			
	putative regulator with homeodomain-like DNA binding domain		Υ	
	DLP12 prophage		Ϋ́	
	unknown CDS		-	
0703	putative rhs protein		Υ	
			Y	
00947	putative 2Fe-2S protein with ferredoxin-like NADP-linked domain and 2Fe-2S ferredoxin-like domain		Υ	Y
	unknown CDS			
1016				
01016 01028	putative malonyl-CoA:Acyl carrier protein transacylase			
1016				

·				
	e14 prophage; putative exisionase			
b1146	e14 prophage; putative regulator			
	conserved hypothetical protein			
	unknown CDS			
	unknown CDS		Υ	
	unknown CDS		Υ	
	conserved protein with PYP-like sensor domain		Υ	Υ
	Rac prophage		Υ	
	Rac prophage; putative DNA replication protein		Υ	
	Rac prophage			
b1369	Rac prophage; putative outer membrane protein		Y	
b1371	Rac prophage; putative outer membrane protein		Y	
b1410	putative methylase with S-adenosyl-L-methionine-dependent methyltransferase domain and alpha/beta-		Υ	
h1420	hydrolase domain			
b1420	unknown CDS unknown CDS			
	putative LpxA-like enzyme			
	conserved hypothetical protein		Υ	
	putative transposase		Ϋ́	
	unknown CDS			
	conserved protein		Υ	
	conserved protein			
	conserved hypothetical protein with homeodomain-like domain			
	conserved protein			
	Qin prophage; putative tail fiber protein		Υ	
b1560	Qin prophage		•	
b1588	putative oxidoreductase subunit with Formate dehydrogenase/DMSO reductase, domains 1-3 and ADC-like			
b1589	putative 4Fe-4S ferredoxin-type protein			
b1598	putative enzyme with serine protease-like domain		Υ	Υ
b1668	putative oxidoreductase with NAD(P)/FAD-binding domain		Y	Ý
	conserved protein with DEATH domain		•	-
	conserved hypothetical protein		Υ	
	conserved hypothetical protein		Υ	Υ
b1773	putative fructose-bisphosphate aldolase		Υ	Υ
b1936	unknown CDS			
b2228	putative membrane protein			
b2299	putative enzyme (Nudix hydrolase)		Υ	Υ
b2384	putative endoglucanase with Zn-dependent exopeptidase domain		Υ	
b2385	putative peptidase with creatinase/prolidase N-terminal domain and creatinase/aminopeptidase		Υ	
b2443	CPZ-55 prophage		Υ	
b2506	conserved hypothetical protein			
b2865	putative lipoprotein, outer membrane		Υ	
b4285	KpLE2 phage-like element; putative transposase			
barA	hybrid sensory histidine kinase in two-component regulatory system with UvrY		Υ	
bglG	transcriptional antiterminator of bgl operon, phosphorylated by BgIF	Υ	Υ	Υ
bglJ	transcriptional activator for the transport and utilization of arbutin and salicin (LuxR/UhpA family)			
bioH	carboxylesterase in pimeloyl-CoA (biotin precursor) synthesis with alpha/beta-hydrolase domain		Υ	Υ
birA	bifunctional: biotin-[acetylCoA carboxylase] holoenzyme synthetase; transcriptional repressor of biotin	Υ	Υ	Υ
	synthesis (BirA family)			
bolA	transcriptional activator of morphogenic pathway (BolA family), important in general stress response		Υ	Υ
btuF	cyano-cobalamin transport protein (ABC superfamily, peri_bind)		Υ	
cadA	lysine decarboxylase 1		Υ	Υ
carA	carbamoyl phosphate synthetase, glutamine amidotransferase small subunit		Υ	Υ
cbpA	curved DNA-binding protein, co-chaperone of DnaK (Hsp40 family)		Υ	Y
cca	tRNA nucleotidyl transferase		Y	Y
cedA	cell division activator		Υ	Y
cgtA	putative GTP-binding protein with nucleoside triP hydrolase domain	Y	Y	Y
cheA	chemotactic sensory histidine kinase (soluble) in two-component regulatory system with CheB and CheY,		Υ	Υ
-114/	senses chemotactic signal			V
cheW	purine-binding chemotaxis protein; regulation chemotactic response regulator in two-component regulatory system with CheA, transmits signals to FliM		Y Y	Y Y
cheY				Ť
			ī	
che7	flagelllar motor component			Y
cheZ	flagelllar motor component chemotactic response, CheY protein phophatase		Y	Y Y
cho	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair		Y Y	Y Y
cho cirA	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB		Y Y Y	Υ
cho	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair		Y Y	
cho cirA clpA	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB ATP-dependent specificity subunit of clpA-clpP serine protease		Y Y Y	Y Y
cho cirA clpA clpB	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB ATP-dependent specificity subunit of clpA-clpP serine protease ATP-dependent protease, Hsp 100, part of multi-chaperone system with DnaK, DnaJ, and GrpE		Y Y Y Y	Y Y Y
cho cirA clpA clpB clpP	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB ATP-dependent specificity subunit of clpA-clpP serine protease ATP-dependent protease, Hsp 100, part of multi-chaperone system with DnaK, DnaJ, and GrpE proteolytic subunit of clpA-clpP ATP-dependent serine protease		Y Y Y Y Y	Y Y Y
cho cirA clpA clpB clpP clpS	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB ATP-dependent specificity subunit of clpA-clpP serine protease ATP-dependent protease, Hsp 100, part of multi-chaperone system with DnaK, DnaJ, and GrpE proteolytic subunit of clpA-clpP ATP-dependent serine protease modulator of ClpA substrate specificity	Y	Y Y Y Y Y	Y Y Y
cho cirA clpA clpB clpP clpS clpX	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB ATP-dependent specificity subunit of clpA-clpP serine protease ATP-dependent protease, Hsp 100, part of multi-chaperone system with DnaK, DnaJ, and GrpE proteolytic subunit of clpA-clpP ATP-dependent serine protease modulator of ClpA substrate specificity ATPase, chaperone subunit of serine protease	Y	Y Y Y Y Y Y	Y Y Y Y
cho cirA clpA clpB clpP clpS clpX coaD	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB ATP-dependent specificity subunit of clpA-clpP serine protease ATP-dependent protease, Hsp 100, part of multi-chaperone system with DnaK, DnaJ, and GrpE proteolytic subunit of clpA-clpP ATP-dependent serine protease modulator of ClpA substrate specificity ATPase, chaperone subunit of serine protease CMP-deoxy-D-manno-octulosonate-lipid A transferase (phosphopantetheine adenylyltransferase)	Y	Y Y Y Y Y Y	Y Y Y Y Y
cho cirA clpA clpB clpP clpS clpX coaD cobB	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB ATP-dependent specificity subunit of clpA-clpP serine protease ATP-dependent protease, Hsp 100, part of multi-chaperone system with DnaK, DnaJ, and GrpE proteolytic subunit of clpA-clpP ATP-dependent serine protease modulator of ClpA substrate specificity ATPase, chaperone subunit of serine protease CMP-deoxy-D-manno-octulosonate-lipid A transferase (phosphopantetheine adenylyltransferase) putative enzyme with DHS-like NAD/FAD-binding domain transcriptional regulator, cyclic AMP receptor protein (CAMP-binding family), interacts with RNAP cysteine sulfinate desulfinase	Y	Y Y Y Y Y Y Y	Y Y Y Y Y Y Y
cho cirA clpA clpB clpP clpS clpX coaD cobB	flagelllar motor component chemotactic response, CheY protein phophatase endonuclease in nucleotide excision repair outer membrane pore protein, receptor for colicin I, requires TonB ATP-dependent specificity subunit of clpA-clpP serine protease ATP-dependent protease, Hsp 100, part of multi-chaperone system with DnaK, DnaJ, and GrpE proteolytic subunit of clpA-clpP ATP-dependent serine protease modulator of ClpA substrate specificity ATPase, chaperone subunit of serine protease CMP-deoxy-D-manno-octulosonate-lipid A transferase (phosphopantetheine adenylyltransferase) putative enzyme with DHS-like NAD/FAD-binding domain transcriptional regulator, cyclic AMP receptor protein (CAMP-binding family), interacts with RNAP	Y	Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y

1 .				
cspA	major cold shock protein 7.4, transcription antiterminator of hns, ssDNA-binding property		Υ	Υ
cspB	Qin prophage; cold shock protein		Υ	Υ
cspC	cold shock protein, transcription antiterminator, affects expression of rpoS and uspA		Υ	Υ
cspD	DNA replication inhibitor, nucleic acid-binding domain		Υ	Υ
cspE	RNA chaperone, transcription antiterminator, affects expression of rpoS and uspA		Υ	Υ
cspF	Qin prophage; cold shock protein			
cspG	low-temperature-responsive gene, nucleic acid-binding domain		Υ	Υ
cspH	cold shock-like protein, nucleic acid-binding domain		Y	
cspl	Qin prophage; cold shock-like protein		Ϋ́	
cutA	periplasmic divalent cation tolerance protein; cytochrome c biogenesis		Ϋ́	Υ
cynT	carbonic anhydrase		Ϋ́	•
cysB	transcriptional regulator of biosynthesis of L-cysteine and regulator of sulfur assimilation (LysR familiy)		Ϋ́	Υ
cysS	cysteine tRNA synthetase	Υ	Ý	Ý
cytR	transcriptional repressor for genes of nucleoside catabolism and recycling (GalR/Lacl family)	•	•	•
dam	DNA adenine methylase		Υ	Υ
dbpA	ATP-dependent RNA helicase, stimulated by 23S rRNA		Ϋ́	'
dcm	DNA cytosine methylase		Ϋ́	Υ
			ı	ī
dctA	citrate and C4-dicarboxylic acids transport protein (DAACS family)		V	V
dcuR	response regulator in two-component regulatory system with DcuS, regulates anaerobic fumarate respiration		Y	Y
ddIA	D-alanine-D-alanine ligase A	Υ	Υ	Υ
ddpX	D-Ala-D-Ala dipeptidase, Zn-dependent			
deaD	cold-shock DeaD box ATP-dependent RNA helicase	Y	Y	Y
def	peptide deformylase	Υ	Υ	Υ
deoC	2-deoxyribose-5-phosphate aldolase, NAD(P)-linked		Υ	Υ
der	GTP-binding protein, essential for cell growth	Υ	Υ	Υ
dfp	bifunctional: 4'-phosphopantothenoylcysteine decarboxylase; phosphopantothenoylcysteine	Υ	Υ	Υ
	synthetase, FMN-binding			
dhaH	putative PTS family enzyme I and HPr components		Υ	Υ
dhaK1	putative dihydroxyacetone kinase		Υ	Υ
dinG	LexA-regulated (SOS) repair enzyme		Υ	
dinJ	damage-inducible protein J		Υ	
dinP	DNA polymerase IV, devoid of proofreading, damage-inducible protein P		Υ	
dmsD	twin-arginine leader-binding protein			
dnaA	DNA replication initiator protein, transcriptional regulator of replication and housekeeping genes	Υ	Υ	Υ
dnaB	replicative DNA helicase; chromosome replication; chain elongation	Ϋ́	Ý	Ϋ́
dnaC	chromosome replication (initiation and chain elongation) with nucleoside triP hydrolase domain	Ÿ	Ϋ́	•
		Ϋ́	Ϋ́	v
dnaE	DNA polymerase III, alpha subunit			Y
dnaG	DNA biosynthesis; DNA primase	Y	Y	Y
dnaJ	heat shock protein (Hsp40), co-chaperone with DnaK	Υ		Υ
			Υ	
dnaK	chaperone Hsp70 in DNA biosynthesis/cell division	Ϋ́	Υ	Ϋ́
dnaK dnaN	DNA polymerase III, beta-subunit			
		Υ	Υ	
dnaN	DNA polymerase III, beta-subunit	Υ	Y Y	Υ
dnaN dnaQ dnaT	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I	Y Y	Y Y Y	Y Y Y
dnaN dnaQ dnaT dnaX	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III	Y Y	Y Y Y Y	Y Y Y
dnaN dnaQ dnaT	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of	Y Y	Y Y Y	Y Y Y
dnaN dnaQ dnaT dnaX dpiA	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family)	Y Y	Y Y Y Y Y	Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain	Y Y	Y Y Y Y Y	Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation	Y Y	Y Y Y Y Y Y	Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD	Y Y	Y Y Y Y Y Y	Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B	Y Y Y	Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C	Y Y Y Y	Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC dut	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase	Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC dut dxs	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding	Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P)	Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp endA	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I	Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp endA eno	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase	Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC dut dxs efp endA eno	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division	Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp endA eno	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization	Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC dut dxs efp endA eno	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division	Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC dut dxs efp endA eno era eutD	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization	Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC efp endA eno era eutD exbB	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase III, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins	Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC efp endA eno era eutD exbB exbD	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase III, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins	Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp endA eno era eutD exbB exbD exoX	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase III, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity	Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp endA eno era eutD exbB exbD fabA fabB	DNA polymerase III; beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase I	Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC dut dxs efp endA eno era eutD exbB exbD fabB fabB	DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III: tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase I malonyl-CoA-[acyl-carrier-protein] transacylase	Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC dut dxs efp endA eno era eutD exbB exbD fabB fabB fabB	DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase II malonyl-CoA-[acyl-carrier-protein] synthase II	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp endA eno exa eutD exbB exbD fabB fabB fabB fabB fabB fabB	DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase II malonyl-CoA-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase II	Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp endA eno exa eutD exbB exbD fabB fabB fabB fabB fabB fabB fabB fabB	DNA polymerase III. beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III. tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase II malonyl-CoA-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase III; acetylCoA ACP transacylase	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC dut dxs efp endA eno era eutD exbB fabA fabB fabB fabB fabB fabB fabB fabB fabB	DNA polymerase III. beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III. tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase I malonyl-CoA-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase III; acetylCoA ACP transacylase enoyl-[acyl-carrier-protein] reductase 8-oxoacyl-[acyl-carrier-protein] reductase 8-oxoacyl-[acyl-carrier-protein] reductase 8-oxoacyl-[acyl-carrier-protein] reductase 8-oxoacyl-[acyl-carrier-protein] reductase 8-oxoacyl-[acyl-carrier-protein] reductase 8-oxoacyl-[acyl-carrier-protein] reductase 9-oxoacyl-[acyl-carrier-protein] reductase 9-oxoacyl-[acyl-carrier-protein] reductase 9-oxoacyl-[acyl-carrier-protein] reductase (NADH)	Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC erp endA eno exaB exbD exoX fabA fabB fabB fabB fabB fabB fabB fabB fabB	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase I-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase I malonyl-CoA-[acyl-carrier-protein] synthase I 3-oxoacyl-[acyl-carrier-protein] reductase 3-oxoacyl-[acyl-carrier-protein] reductase 0NADH) (3R)-hydroxymyristol acyl carrier protein dehydratase	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dps dsbA dsbC dusB dusC dut dxs efp endA eno era eutD exbB exbD fabA fabB fabD fabF fabG fabH fabI fabZ fadA	DNA polymerase III, beta-subunit DNA polymerase III: epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase I malonyl-CoA-[acyl-carrier-protein] synthase I nalonyl-CoA-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase III; acetyl-CoA ACP transacylase enoyl-[acyl-carrier-protein] reductase 3-oxoacyl-[acyl-carrier-protein] reductase 3-oxoacyl-[acyl-carrier-protein] reductase 3-oxoacyl-[acyl-carrier-protein] reductase (NADH) (3R)-hydroxymyristol acyl carrier protein dehydratase) 3-ketoacyl-CoA thiolase; (thiolase I, acetyl-CoA transferase), in complex with FadB catalyzes EC 2.3.1.16	Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbA dsbC dusB dusC erp endA eno exaB exbD exoX fabA fabB fabB fabB fabB fabB fabB fabB fabB	DNA polymerase III, beta-subunit DNA polymerase III, epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III, tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxyuridinetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase I malonyl-CoA-[acyl-carrier-protein] synthase I malonyl-CoA-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase III; acetylCoA ACP transacylase 3-oxoacyl-[acyl-carrier-protein] reductase 3-oxoacyl-[acyl-carrier-protein] reductase (NADH) (3R)-hydroxymyristol acyl carrier protein dehydratase 3-ketoacyl-CoA thiolase; (thiolase I, acetyl-CoA transferase), in complex with FadB catalyzes EC 2.3.1.16 multifunctional: 3-hydroxybutyryl-CoA epimerase, delta(3)-cis-delta(2)-trans-enoyl-CoA isomerase, enoyl-CoA	Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbC dusB dusC dut dxs efp endA eno era eutD exbB fabA fabB fabB fabB fabB fabB fabB fabB fabB	DNA polymerase III. epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III. tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxy-didnetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] reductase (NADH) (3R)-hydroxymyristol acyl carrier protein dehydratase 3-ketoacyl-CoA thiolase; (thiolase I, acetyl-CoA transferase), in complex with FadB catalyzes EC 2.3.1.16 multifunctional: 3-hydroxybutryl-CoA epimerase, delta(3)-cis-delta(2)-trans-enoyl-CoA isomerase, enoyl-CoA hydratase (N-terminal); 3-hydroxyacyl-CoA dehydrogenase (C-terminal)	Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbC dusB dusC dut dxs efp endA eno exa eutD exbB exbD fabF fabG fabH fabI fabI fabI fabI fabI fabI fadB	DNA polymerase III. epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III. tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase I, activated by N-terminal of DsbD (tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase I malonyl-CoA-[acyl-carrier-protein] transacylase 3-oxoacyl-[acyl-carrier-protein] reductase 3-oxoacyl-[acyl-carrier-protein] synthase III; acetylCoA ACP transacylase enoyl-[acyl-carrier-protein] synthase III; acetylCoA ACP transacylase 3-oxoacyl-[acyl-carrier-protein] synthase III; acetylCoA ACP transacylase enoyl-[acyl-carrier-protein] synthase II, acetyl-CoA transferase), in complex with FadB catalyzes EC 2.3.1.16 multifunctional: 3-hydroxybutryl-CoA epimerase, delta(3)-cis-delta(2)-trans-enoyl-CoA isomerase, enoyl-CoA hydratase (N-terminal); 3-hydroxyacyl-CoA dehydrogenase (C-terminal)	Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
dnaN dnaQ dnaT dnaX dpiA dsbC dusB dusC dut dxs efp endA eno era eutD exbB fabA fabB fabB fabB fabB fabB fabB fabB fabB	DNA polymerase III. epsilon subunit, 3-5 exonucleolytic proofreading function primosomal protein I DNA polymerase III. tau and gamma subunits; DNA elongation factor III response regulator in two-component regulatory system with DpiB, regulation of citrate fermentation and of plasmid inheritance (OmpR family) stress response DNA-binding protein with ferritin-like domain periplasmic protein disulfide isomerase I, disulfide bond formation protein disulfide isomerase II, activated by N-terminal of DsbD tRNA-dihydrouridine synthase B tRNA-dihydrouridine synthase C deoxy-didnetriphosphatase 1-deoxy-D-xylulose 5-phosphate synthase; flavoprotein, thiamin-binding elongation factor P (EF-P) DNA-specific endonuclease I enolase GTPase believed to be involved in coordination of cell cycle, energy metabolism, cell division putative phosphate acetyltransferase in ethanolamine utilization uptake of enterobactin; tonB-dependent uptake of B colicins uptake of enterobactin; tonB-dependent uptake of B colicins DNA exonuclease X, degrades ss and ds DNA with 3'-5' polarity beta-hydroxydecanoyl thioester dehydrase (trans-2-decenoyl-ACP isomerase) 3-oxoacyl-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] synthase II 3-oxoacyl-[acyl-carrier-protein] reductase (NADH) (3R)-hydroxymyristol acyl carrier protein dehydratase 3-ketoacyl-CoA thiolase; (thiolase I, acetyl-CoA transferase), in complex with FadB catalyzes EC 2.3.1.16 multifunctional: 3-hydroxybutryl-CoA epimerase, delta(3)-cis-delta(2)-trans-enoyl-CoA isomerase, enoyl-CoA hydratase (N-terminal); 3-hydroxyacyl-CoA dehydrogenase (C-terminal)	Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y

fadR	transcriptional regulator (positive regulator of fabB and fabA, negative regulator of fad), with acyl-CoA (GntR family)		Υ	Υ
arR	transcriptional repressor for TCA cycle, fatty acyl-responsive transcriptional regulator (GntR family)		Υ	
baA	fructose-bisphosphate aldolase, class II	Υ	Υ	Υ
baB	fructose-bisphosphate aldolase class I		Υ	Υ
dhD	formate dehydrogenase formation protein	Υ	Υ	Υ
dhF	formate dehydrogenase H, selenopolypeptide subunit			
dnG	formate dehydrogenase-N, alpha subunit, nitrate-inducible		Υ	Υ
dnH	formate dehydrogenase-N, Fe-S beta subunit, nitrate-inducible			
dnl	formate dehydrogenase-N, cytochrome B556(Fdn) gamma subunit, nitrate-inducible			
doG	formate dehydrogenase-O, major subunit			
doH	formate dehydrogenase-O, Fe-S subunit			
ecA	KpLE2 phage-like element; outer membrane porin, receptor for ferric citrate, in multi-component regulatory system with cytoplasmic Fecl (sigma factor) and membrane-bound FecR			
ecB	KpLE2 phage-like element; citrate-dependent iron (III) transport protein (ABC superfamily, peri bind)		Υ	
ерВ	ferric enterobactin tranport protein (ABC superfamily, peri_bind)		Ý	
h	4.5S-RNP protein, GTP-binding export factor, part of signal recognition particle with 4.5 RNA	Υ	Ý	Υ
nuD	hydroxamate-dependent iron transport protein (ABC superfamily, peri bind)	•	•	•
mB	tyrosine recombinase, regulator of fimA		Υ	Υ
mE	tyrosine recombinase, regulator of fimA		Ϋ́	'
	DNA-binding protein for site-specific recombination and inversion, transcription of rRNA and tRNA operons, and		Ϋ́	Υ
S	DNA replication			ĭ
хВ	putative electron transfer flavoprotein, NAD/FAD-binding domain and ETFP adenine nucleotide-binding domain-like		Y	
klΒ	FKBP-type 22KD peptidyl-prolyl cis-trans isomerase (rotamase)		Y	Y
крΑ	FKBP-type peptidyl-prolyl cis-trans isomerase (rotamase)		Y	Y
κрВ	FKBP-type peptidyl-prolyl cis-trans isomerase (rotamase)		Υ	Υ
dA	flavodoxin 1	Υ	Υ	Υ
gE	flagellar biosynthesis; hook protein			
gK	flagellar biosynthesis; hook-filament junction protein 1		Υ	
hC	transcriptional regulator of flagellar class II biosynthesis, anaerobic respiration and the Entner-Doudoroff pathway, tetramer with FlhD		Y	
hD	transcriptional regulator of flagellar class II biosynthesis, anaerobic respiration and the Entner-Doudoroff pathway, tetramer with FlhC		Υ	
iC	flagellar biosynthesis; flagellin, filament structural protein			
iD	flagellar biosynthesis; filament capping protein, enables filament assembly			
iE	flagellar biosynthesis; basal-body component			
iG	flagellar biosynthesis; component of motor switching and energizing			
iH	flagellar biosynthesis; putative export of flagellar proteins			
il	flagellum-specific ATP synthase		Υ	
iJ	flagellar fliJ protein			
iM	flagellar biosynthesis; component of motor switch and energizing			
iN	flagellar biosynthesis; component of motor switch and energizing		Υ	
iO	flagellar biosynthesis			
iS	flagellar biosynthesis; repressor of class 3a and 3b operons (RflA activity)			
liΥ	cysteine transport protein (ABC superfamily, peri_bind)	Υ	Υ	Υ
nr	transcriptional regulator of aerobic, anaerobic respiration, osmotic balance (CAMP-binding family)		Υ	Υ
Αlc	dihydrofolate reductase type I, trimethoprim resistance	Υ	Υ	Υ
olC	bifunctional: folylpolyglutamate synthase; dihydrofolate synthase	Υ	Υ	Υ
dΑ	fumarate reductase, anaerobic, catalytic and NAD/flavoprotein subunit	-	Ϋ́	Ϋ́
e	flavin reductase, FAD = preferred substrate		Ϋ́	Ϋ́
	·	Υ	Ϋ́	Y
rr D	ribosome releasing factor	ī		
uR	transcriptional regulator of the control of carbon and energy metabolism (GalR/Lacl family)	.,	Y	Y
sA	cell division protein with ATPase domain, involved in recruitment of FtsK to Z ring	Υ	Υ	Υ
sE	putative transport protein (ABC superfamily, atp_bind)	Υ	Υ	Υ
sl	division-specific transpeptidase, penicillin-binding protein 3			
sK	cell division protein required for chromosome partitioning with P-loop containing nucleoside triphosphate hydrolase domain		Υ	Υ
sX	integral membrane cell division protein		Υ	
sΥ	cell division protein: membrane binding (N-terminal); GTPase domain (C-terminal)	Υ	Ý	
sZ	tubulin-like GTP-binding protein and GTPase, forms circumferential ring in cell division	Ϋ́	Ϋ́	Υ
ucU	conserved protein of fucose operon	•	Ϋ́	Y
	·	Υ		
usA	protein chain elongation factor EF-G, GTP-binding	T	Y	Y
adA	glutamate decarboxylase A, isozyme, PLP-dependent		Y	Y
adB	glutamate decarboxylase, PLP-dependent, isozyme beta		Y	Υ
alR	transcriptional repressor for galactose utilization (GalR/Lacl family)		Y	
арА	glyceraldehyde-3-phosphate dehydrogenase A		Υ	Υ
arK	glycerate kinase I		Υ	Υ
atB	PTS family enzyme IIB, galactitol-specific		Υ	Υ
atY	tagatose 6-phosphate aldolase 2, subunit with GatZ		Υ	Υ
atZ	tagatose 6-phosphate aldolase 2, subunit with GatY		Y	Y
	putative O-sialoglycoprotein endopeptidase, with actin-like ATPase domain	Υ	Ý	Ý
			Ϋ́	•
ср	alutamate dehydrogenase NADP-specific			
cp dhA	glutamate dehydrogenase, NADP-specific	Υ		V
	glutamate dehydrogenase, NADP-specific lucose-inhibited division protein, oxidoreductase-like with FAD/NAD(P)-binding domain	Y Y	Ý Y	Y Y

glmS glmU	L-glutamine:D-fructose-6-phosphate aminotransferase bifunctional: N-acetyl glucosamine-1-phosphate uridyltransferase (N-terminal); glucosamine-1-phosphate acetyl transferase (C-terminal)	Y Y	Y Y	Y
glnB	regulatory protein (P-II) for nitrogen assimilation by glutamine synthetase (ATase)		Υ	Υ
ginS	glutamine tRNA synthetase	Υ	Ý	Ý
-		•	Ϋ́	Ϋ́
gloA	glyoxalase I, nickel isomerase			
gloB	putative hydroxyacylglutathione hydrolase with metallo-hydrolase/oxidoreductase domain		Y	Y
gltD	glutamate synthase, small subunit, nucleotide-binding, 4Fe-4S protein		Υ	Υ
gltF	transcriptional regulator of glutamate synthase, induction of Ntr enzymes		Υ	
gltX	glutamate tRNA synthetase, catalytic subunit	Υ	Υ	Υ
glyQ	glycine tRNA synthetase, alpha subunit	Υ	Υ	Υ
glyS	glycine tRNA synthetase, beta subunit	Υ	Υ	Υ
gmk	guanylate kinase	Ÿ	Ý	Ý
gnsB	Qin prophage; suppressor of cold/temperature-sensitive mutants, affects levels of unsaturated fatty acids, similar to GnsA	-	Ϋ́	•
aro A	transcription elongation factor, cleaves 3' nucleotide of paused mRNA		Υ	Υ
greA	· · · · · · · · · · · · · · · · · · ·			
greB	transcription elongation factor and transcript cleavage	.,	Y	Y
groL	chaperone Hsp60 (GroEL), part of GroE chaperone system	Υ	Υ	Υ
groS	chaperone Hsp10 (GroES), part of GroE chaperone system	Υ	Υ	Υ
grpE	Hsp 24 nucleotide exchange factor	Υ	Υ	Υ
gspE	putative protein exporter, transport across outer membrane (General Secretory Pathway) with P-loop containing nucleoside triphosphate hydrolase domain)		
guaC	GMP reductase		Υ	Υ
gyrA	DNA gyrase, subunit A, type II topoisomerase	Υ	Υ	Υ
gyrB	DNA gyrase, subunit B (type II topoisomerase)	Y	Y	Y
hda	regulatory factor involved in inactivation of DnaA	Ý	Ý	Ý
		•	Ϋ́	Y
nelD	DNA helicase IV			
nexR	putative transcriptional regulator with phosphosugar-binding domain		Y	Y
nflB	ATP-dependent zinc-metallo protease		Υ	Υ
nfq	host factor I for bacteriophage Q beta replication, plays a role in degradation of RNA transcripts		Υ	Υ
nimA	integration host factor (IHF), alpha subunit, DNA-bending protein, DNA replication		Υ	Υ
nimD	integration host factor (IHF), beta subunit, site-specific recombination		Υ	Υ
hisS	histidine tRNA synthetase	Υ	Υ	Υ
nldD	ADP-L-glycero-D-mannoheptose-6-epimerase, NAD(P)-binding		Ϋ́	Y
	periplasmic molecular chaperone for outer membrane proteins	Υ	Ϋ́	Ý
hlpA		1		T
nmsR	putative transport protein (VGP family)		Y	
nns nofB	transcriptional regulator, DNA-binding protein HLP-II (HU, BH2, HD, NS), increases DNA thermal stability putative integral membrane protein involved in biogenesis of finition from transport, DNA with D lean containing pu		Y Y	Y
	uptake with P-loop containing nucleoside triphosphate hydrolysis domain		v	.,
holA	DNA polymerase III, delta subunit, probably ATP hydrolase	Υ	Υ	Υ
holB	DNA polymerase III, delta prime subunit	Υ	Υ	Υ
noIC	DNA polymerase III, chi subunit		Υ	Υ
noID	DNA polymerase III, psi subunit		Υ	Υ
holE	DNA polymerase III, theta subunit		Υ	Υ
nrpA	helicase, ATP-dependent		Υ	Υ
nrpB	helicase, ATP-dependent		Ϋ́	Ý
•				
nscA	chaperone (Hsp70 family), believed to be involved in assembly of Fe-S clusters		Y	Y
nscB	co-chaperone protein Hsc20, believed to be involved in assembly of Fe-S clusters		Y	Y
nscC	putative heatshock protein (Hsp70 family), with actin-like ATPase domain and C-terminal substrate-binding domain		Y	Y
hsdM	DNA methylase M, host modification		Y	Υ
nsdR	endonuclease R, host restriction		Υ	Υ
nsdS			\/	Υ
	specificity determinant for hsdM and hsdR		Υ	
nslJ	specificity determinant for hsdM and hsdR heat shock protein hslJ		Υ Υ	
	heat shock protein hslJ			Υ
nsIO	heat shock protein hslJ heat shock protein 33, redox regulated chaperone		Y Y	
nsIO nsIR	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding		Y Y Y	Υ
nsIO nsIR nsIU	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone		Y Y Y	Y Y
nsIO nsIR nsIU nsIV	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease		Y Y Y Y	Y Y Y
nsIO nsIR nsIU nsIV ntpG	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5		Y Y Y	Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC		Y Y Y Y Y	Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation		Y Y Y Y Y	Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation		Y Y Y Y Y	Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit		Y Y Y Y Y Y	Y Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation		Y Y Y Y Y	Υ
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit		Y Y Y Y Y Y	Y Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC nybE nybO	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase-2, small subunit		Y Y Y Y Y Y Y Y	Y Y Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC nybE nybO nycE	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase-2, small subunit hydrogenase 3, large subunit (part of FHL complex)		Y Y Y Y Y Y Y	Y Y Y Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC nybE nybO nycE	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase-2, small subunit hydrogenase 3, large subunit (part of FHL complex) protease involved in processing C-terminal end of HycE		Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC nybE nybC nycE nycI nyfG	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase 3, large subunit (part of FHL complex) protease involved in processing C-terminal end of HycE hydrogenase 4 subunit		Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y
hsIJ hsIO hsIR hsIU hsIV htpG htrC hupA hupB hybC hybE hybO hycE hydG hyfG hyfR hypA	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase 2, small subunit hydrogenase 3, large subunit (part of FHL complex) protease involved in processing C-terminal end of HycE hydrogenase 4 subunit transcriptional activator for expression of hydrogenase 4 genes, interacts with sigma 54 (EBP family) guanine-nucleotide-binding protein in formate-hydrogenlyase system, functions as nickel donor for HycE of		Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC nybE nybC nycE nycI nyfG nyfR	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase 3, large subunit (part of FHL complex) protease involved in processing C-terminal end of HycE hydrogenase 4 subunit transcriptional activator for expression of hydrogenase 4 genes, interacts with sigma 54 (EBP family) guanine-nucleotide-binding protein in formate-hydrogenlyase system, functions as nickel donor for HycE of hydrogenlyase 3		Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC nybE nybC nycE nycI nyfG nyfR nypA	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase 2, small subunit hydrogenase 3, large subunit (part of FHL complex) protease involved in processing C-terminal end of HycE hydrogenase 4 subunit transcriptional activator for expression of hydrogenase 4 genes, interacts with sigma 54 (EBP family) guanine-nucleotide-binding protein in formate-hydrogenlyase system, functions as nickel donor for HycE of		Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
hsIO hsIR hsIU hsIV htpG htrC hupA hupB hybC hybE hybO hycE hycI hyfG hyfR	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase 3, large subunit (part of FHL complex) protease involved in processing C-terminal end of HycE hydrogenase 4 subunit transcriptional activator for expression of hydrogenase 4 genes, interacts with sigma 54 (EBP family) guanine-nucleotide-binding protein in formate-hydrogenlyase system, functions as nickel donor for HycE of hydrogenlyase 3		Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y
nsIO nsIR nsIU nsIV ntpG ntrC nupA nupB nybC nybE nybC nycE nycI nyfG nyfR nypA	heat shock protein hslJ heat shock protein 33, redox regulated chaperone heat shock protein 15, DNA/RNA-binding ATPase component of the HslUV protease, also functions as molecular chaperone peptidase component of the HslUV protease chaperone Hsp90, heat shock protein C 62.5 heat shock protein htrC DNA-binding protein HU-alpha (HU-2), plays a role in DNA replication and in rpo translation DNA-binding protein HU-beta, NS1 (HU-1), plays a role in DNA replication and in rpo translation hydrogenase-2, large subunit putative hydrogenase hydrogenase-2, small subunit hydrogenase 3, large subunit (part of FHL complex) protease involved in processing C-terminal end of HycE hydrogenase 4 subunit transcriptional activator for expression of hydrogenase 4 genes, interacts with sigma 54 (EBP family) guanine-nucleotide-binding protein in formate-hydrogenlyase system, functions as nickel donor for HycE of hydrogenlyase 3 hydrogenase expression/formation protein		Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y

ibpA	small heat shock protein		Y	Υ
ibpB	small heat shock protein		Y	Y
iciA	inhibitor of replication initiation; transcriptional regulator of dnaA and argK (LysR family)		Y	Y
iclR	transcriptional repressor for glyoxylate bypass (IcIR family)		Y	Y
dnD	L-idonate 5-dehydrogenase, NAD-binding		Y	Y
dnO	5-keto-D-gluconate-5-reductase		Y	Y
leS	isoleucine tRNA synthetase	Υ	Υ	Υ
IvB	acetolactate synthase I, large subunit, valine-sensitive	Υ	Υ	Υ
lvH	acetolactate synthase III, valine-sensitive, small subunit		Υ	
infA	protein chain initiation factor IF-1	Υ	Υ	
nfB	protein chain initiation factor IF-2		Υ	Υ
infC	protein chain initiation factor IF-3	Υ	Υ	Υ
ntA	CP4-57 prophage; integrase		Υ	Υ
ntB	KpLE2 phage-like element; P4-like integrase			
ntD	DLP12 prophage; integrase		Υ	
ntQ	Qin prophage; putative transposase			
ntR	Rac prophage; putative transposase/integrase			
ntS	CPS-53 (KpLE1) prophage; Sf6-like integrase		Υ	Υ
ntZ	CPZ-55 prophage; putative integrase		Υ	
scS	cysteine desulfurase (tRNA sulfurtransferase), PLP-dependent	Υ	Υ	Υ
scU	putative Fe-S assembly protein	Υ	Υ	Υ
spA	geranyltranstransferase (=farnesyldiphosphate synthase)	Υ	Υ	Υ
spE	4-diphosphocytidyl-2C-methyl-D-erythritol kinase	Υ	Υ	Υ
spF	2C-methyl-D-erythritol 2,4-cyclodiphosphate synthase	Y	Y	Y
spG	1-hydroxy-2-methyl-2-(E)-butenyl 4-diphosphate synthase	Ÿ	Ý	Ý
spH	1-hydroxy-2-methyl-2-(E)-butenyl 4-diphosphate reductase, 4Fe-4S protein	Ÿ	Ý	Ÿ
katE	catalase; hydroperoxidase HPII (III), RpoS-dependent	•	Ϋ́	Ý
kdsA	3-deoxy-D-manno-octulosonic acid 8-P synthetase	Υ	Ϋ́	Ý
		Ϋ́		Ý
kdsB	CTP:CMP-3-deoxy-D-manno-octulosonate transferase		Y	Y
kdtA	3-deoxy-D-manno-octulosonic-acid transferase (KDO transferase)	Y	Y	Y
kefC	K+ efflux antiporter, glutathione-regulated, NAD(P)-binding (CPA2 family)		Y	.,
(sgA	S-adenosylmethionine-6-N',N'-adenosyl (rRNA) dimethyltransferase, kasugamycin resistance		Υ	Υ
acl	transcriptional repressor of lactose catabolism (GalR/Lacl family)			
acY	galactoside permease (lactose permease, M protein) (MFS family)		.,	
amB	maltoporin, high-affinity receptor for maltose and maltoseoligosaccharides; phage lambda receptor		Υ	
dcA	L, D-carboxypeptidase A (in murein recycling)		Y	Y
dcC	lysine decarboxylase 2, constitutive		Υ	Υ
ерА	GTP-binding elongation factor	Υ	Υ	Υ
leuS	leucine tRNA synthetase	Υ	Υ	Υ
exA	transcriptional repressor for SOS response (signal peptidase of LexA family)		Υ	Υ
ligA	DNA ligase	Υ	Υ	Υ
igB	DNA ligase			
ldD	L-lactate dehydrogenase, FMN-linked	Υ	Υ	Υ
olA	periplasmic chaperone effects translocation of lipoproteins from inner membrane to outer membrane	Υ	Υ	Υ
lolB	outer membrane component involved in lipoprotein localization	Υ	Υ	
oID	transport protein of outer membrane lipoproteins (ABC superfamily, atp_bind)	Υ	Υ	
on	DNA-binding ATP-dependent protease La; heat shock K-protein		Υ	Υ
pΙΑ	lipoate-protein ligase A	Υ	Ý	Ý
рхА	UDP-N-acetylglucosamine acetyltransferase	Ϋ́	Ϋ́	Ÿ
	tetraacyldisaccharide-1-P synthase	Ÿ	Ý	Ÿ
рхВ				
рхС	UDP-3-O-acyl N-acetylglucosamine deacetylase	Y Y	Y Y	Υ
rp	UDP-3-O-(3-hydroxymyristoyl)-glucosamine N-acyltransferase transcriptional regulator of lrp regulon and for high-affinity branched-chain amino acid transport system (AsnC family)	•	Y	T
uxS	quorum-sensing protein, produces autoinducer - acyl-homoserine lactone-signaling molecules		Υ	Υ
ysA	diaminopimelate decarboxylase, PLP-binding		Ϋ́	Ϋ́
ysA ysS	lysine tRNA synthetase, constitutive		Ϋ́	Ϋ́
ysU ysU	lysine tRNA synthetase, inducible; heat shock protein	Υ	Ϋ́	Ý
	accessory protein to ABC-type macrolide transport protein MacB	•	Ϋ́	1
macA	bifunctional: putative malic oxidoreductase (N-terminal); putative phosphotransacetylase (C-terminal)		Ϋ́Υ	Υ
naeB	maltose transport protein, chemotaxis (ABC superfamily, peri bind)		Y Y	Y
nalE			Ϋ́Υ	
nalF malG	maltose transport protein (ABC superfamily, membrane)			
nalG	maltose transport protein (ABC superfamily, membrane)		Y	V
nalK	bifunctional: maltose transport protein (ABC superfamily, atp_bind) (N-terminal); phenotypic repressor of mal		Υ	Y
nall/	operon (C-terminal)		~	
nalM	periplasmic protein of mal regulon		Y	V
malP	maltodextrin phosphorylase		Y	Y
malT	transcriptional activator of maltose utilization, binds maltotriose (inducer) and ATP (LysR familiy)		Y	Y
manA	mannose-6-phosphate isomerase	Y	Y	Y
manX	PTS family enzyme IIA (N-terminal); enzyme IIB (C-terminal), mannose-specific		Y	Y
map_	methionine aminopeptidase	Υ	Y	Υ
marR	transcriptional repressor for antibiotic resistance and oxidative stress		Υ	
ma=C	conserved protein		Υ	Υ
nazG ncrA	e14 prophage; restriction of DNA at 5-methylcytosine residues			

mcrB	component of 5-methylcytosine-specific restriction enzyme McrBC		Υ	Υ
melR	transcriptional activator of melibiose catabolism (AraC/XyIS family)		Υ	
menB	dihydroxynaphthoic acid synthetase		Y	Υ
	o-succinylbenzoyl-CoA synthase		Ý	Ϋ́
menC				
menD	bifunctional: 2-oxoglutarate decarboxylase; SHCHC synthase		Υ	Υ
menE	o-succinylbenzoate-CoA ligase		Υ	
menF	isochorismate synthase (isochorismate hydroxymutase 2), menaquinone biosynthesis		Υ	Υ
metK	methionine adenosyltransferase 1 (AdoMet synthetase)		Υ	Υ
metN			Ý	Ϋ́
	D- and L-methionine transport protein (ABC superfamily, atp_bind)			
metQ	D-methionine transport protein (ABC superfamily, peri_bind)		Υ	Υ
mfd	transcription-repair ATP-dependent coupling factor	Υ	Υ	Υ
mglB	galactose transport protein (ABC superfamily, peri_bind)		Υ	
mhpR	transcriptional activator for 3-hydroxyphenylpropionate degradation (IcIR family)		Υ	
	1 31 31 1 0 (3)			V
miaB	involved in methylthiolation of isopentenylated A37 derivatives in tRNA, Fe-S protein		Y	Υ
mipA	scaffolding protein for murein-synthesizing holoenzyme, outer membrane protein		Υ	
mltD	lytic murein transglycosylase C, membrane-bound		Υ	
moaA	molybdopterin biosynthesis protein A		Υ	Υ
moaB	molybdopterin biosynthesis protein B		Ϋ́	Ϋ́
moaC	molybdenum cofactor biosynthesis protein C		Υ	Υ
moaD	molybdenum cofactor biosynthesis protein D		Υ	Υ
moaE	molybdopterin converting factor, subunit 2		Υ	Υ
mobA	molybdopterin-guanine dinucleotide synthase		Υ	Υ
mobB	molybdopterin-guanine dinucleotide biosynthesis protein B, GTP-binding		Ý	Ϋ́
				T
modE	transcriptional repressor for molybdate uptake		Υ	
modF	molybdenum transport protein (ABC superfamily, atp_bind)	Υ		
moeA	molybdopterin biosynthesis protein, molybdenum incorporation step		Υ	Υ
moeB	ATP-dependent adenylate transferase, modifies MoaD		Ý	Y
mogA	putative molybdochetalase in molybdopterine biosynthesis, metal incorporation step		Υ	Υ
mraW	S-adenosyl-dependent methyl transferase		Υ	Υ
mraZ	conserved hypothetical protein		Υ	
mreB	split gene in K-12 with ATPase domain, associated with mecillinam resistance, cell shape		Υ	Υ
mrr	restriction of methylated adenine		Ϋ́	Ϋ́
mrsA	phosphoglucosamine mutase	Υ	Υ	Υ
msrB	methionine sulfoxide reductase		Υ	Υ
mtID	mannitol-1-phosphate dehydrogenase, NAD(P)-binding	Υ	Υ	Υ
mukB	kinesin-like cell division protein involved in sister chromosome partitioning		Υ	Υ
	· · · · · · · · · · · · · · · · · · ·		Ý	Ϋ́
mukE	putative killing protein suppressor			
mukF	mukF protein (killing factor KICB)		Υ	Υ
murA	UDP-N-acetylglucosamine 1-carboxyvinyltransferase	Υ	Υ	Υ
murB	UDP-N-acetylenolpyruvoylglucosamine reductase, FAD-binding	Υ	Υ	Υ
murC		Ý	Ý	Ϋ́
	UDP-N-acetyl-muramate:alanine ligase, L-alanine adding enzyme			
murD	UDP-N-acetylmuramoylalanine-D-glutamate ligase	Υ	Υ	Υ
murE	UDP-N-acetylmuramoylalanyl-D-glutamate 2,6-diaminopimelate ligase	Υ	Υ	Υ
murF	D-alanine:D-alanine-adding enzyme	Υ	Υ	Υ
murG		Ý	Ý	Ϋ́
illulG	UDP-N-acetylglucosamine:N-acetylmuramyl-(pentapeptide) pyrophosphoryl-undecaprenol N-	ī	ı	ī
	acetylglucosamine transferase			
murl	glutamate racemase	Υ	Υ	Υ
mutH	putative methyl-directed mismatch repair protein with restriction endonuclease-like domain		Υ	
mutL	enzyme in methyl-directed mismatch repair, stimulates binding of Vsr and MutS to heteroduplex DNA		Υ	Υ
	formamidopyrimidine DNA glycosylase, also acts on 5-formyluracil and 5-hydroxymethyluracil		Ϋ́	•
mutM				
mutS	methyl-directed mismatch repair, recognizes exocyclic adducts of guanosine		Υ	Υ
mutT	7,8-dihydro-8-oxoguanine-triphosphatase, prefers dGTP		Υ	Υ
mutY	adapina DNA aluanadana			
	adenine DNA glycosylase		Υ	Υ
nadE	0, ,	Υ		
nadE	NAD synthetase, prefers NH3 over glutamine	Y	Υ	Υ
nanE	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain	Y Y	Y Y	Y Y
nanE napA	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB		Υ	Υ
nanE	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain		Y Y	Y Y
nanE napA napB	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA		Y Y	Y Y Y
nanE napA napB napD	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein		Y Y Y	Y Y
nanE napA napB napD napF	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer		Y Y Y	Y Y Y
nanE napA napB napD napF napG	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer		Y Y Y	Y Y Y
nanE napA napB napD napF napG narG	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit		Y Y Y Y	Y Y Y Y
nanE napA napB napD napF napG	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer		Y Y Y Y	Y Y Y
nanE napA napB napD napF napG narG	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit		Y Y Y Y	Y Y Y Y
nanE napA napB napD napF napG narG narH	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly		Y Y Y Y Y	Y Y Y Y
nanE napA napB napD napF napG narG narH narJ	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function		Y Y Y Y Y Y	Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit		Y Y Y Y Y Y Y	Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW narY	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit nitrate reductase 2, alpha subunit		Y Y Y Y Y Y Y	Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit		Y Y Y Y Y Y Y	Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW narY narZ ndk	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase		Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narG narH narJ narW narY narZ ndk nei	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit nitrate reductase 2, alpha subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease VIII, DNA glycosylase activity for 5-formyluracil and 5-hydroxymethyluracil		Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narG narH narJ narW narY narZ ndk nei nfi	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 2, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, alpha subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease VIII, DNA glycosylase activity for 5-formyluracil and 5-hydroxymethyluracil endonuclease V (deoxyinosine 3'endoduclease)		Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW narY narY nard ndk nei nfi nfo	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit nitrate reductase 2, alpha subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease V (deoxyinosine 3'endoduclease) endonuclease IV, with intrinsic 3'-5' exonuclease activity		Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narG narH narJ narW narY narZ ndk nei nfi	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 2, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, alpha subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease VIII, DNA glycosylase activity for 5-formyluracil and 5-hydroxymethyluracil endonuclease V (deoxyinosine 3'endoduclease)		Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW narY narY nard ndk nei nfi nfo	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit nitrate reductase 2, alpha subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease V (deoxyinosine 3'endoduclease) endonuclease IV, with intrinsic 3'-5' exonuclease activity		Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW narY narZ ndk nei nfi nfo nikA nikB	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, alpha subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease VIII, DNA glycosylase activity for 5-formyluracil and 5-hydroxymethyluracil endonuclease V (deoxyinosine 3'endoduclease) endonuclease IV, with intrinsic 3'-5' exonuclease activity nickel transport protein (ABC superfamily, peri_bind) nickel transport protein (ABC superfamily, membrane)		Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW narY narZ ndk nei nfi nfo nikA nikB nikC	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit nitrate reductase 2, alpha subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease VIII, DNA glycosylase activity for 5-formyluracil and 5-hydroxymethyluracil endonuclease V, with intrinsic 3'-5' exonuclease activity nickel transport protein (ABC superfamily, peri_bind) nickel transport protein (ABC superfamily, membrane) nickel transport proein (ABC superfamily, membrane)		Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW narY narZ ndk nei nfi nfo nikA nikB nikC nikD	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, Fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease VIII, DNA glycosylase activity for 5-formyluracil and 5-hydroxymethyluracil endonuclease V (deoxyinosine 3'endoduclease) endonuclease V, with intrinsic 3'-5' exonuclease activity nickel transport protein (ABC superfamily, peri_bind) nickel transport protein (ABC superfamily, membrane) nickel transport protein (ABC superfamily, membrane) nickel transport protein (ABC superfamily, papi_bind)		Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y
nanE napA napB napD napF napG narG narH narJ narW narY narZ ndk nei nfi nfo nikA nikB nikC	NAD synthetase, prefers NH3 over glutamine putative ManNAc-6P epimerase, NAD(P)-linked, with ribulose-phoshate binding barrel domain periplasmic nitrate reductase, large subunit, in complex with NapB periplasmic nitrate reductase, small subunit, cytochrome C550, in complex with NapA periplasmic nitrate reductase assembly protein Fe-S ferredoxin-type protein, electron transfer Fe-S ferredoxin-type protein, electron transfer nitrate reductase 1, alpha subunit nitrate reductase 1, fe-S (beta) subunit nitrate reductase 1, delta subunit, chaperone required for molybdenum cofactor assembly nitrate reductase 2, delta subunit, assembly function nitrate reductase 2, beta subunit nitrate reductase 2, alpha subunit nitrate reductase 2, alpha subunit nucleoside diphosphate kinase endonuclease VIII, DNA glycosylase activity for 5-formyluracil and 5-hydroxymethyluracil endonuclease V, with intrinsic 3'-5' exonuclease activity nickel transport protein (ABC superfamily, peri_bind) nickel transport protein (ABC superfamily, membrane) nickel transport proein (ABC superfamily, membrane)		Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y

nikR	transcriptional repressor of nickel transport, nickel-responsive		Υ	Υ
nohA	Qin prophage; packaging protein NU1		.,	
norV	flavorubredoxin (FIRd), bifunctional NO and O2 reductase		Υ	Υ
nrdA	ribonucleoside diphosphate reductase 1, alpha subunit	Υ	Υ	Υ
ırdB	ribonucleoside-diphosphate reductase 1, beta subunit	Υ	Υ	Υ
nth	endonuclease III; DNA glycosylase/apyrimidinic (AP) lyase, acts on 5-formyluracil and 5-hydroxymethyluracil		Υ	Υ
ntpA	dATP pyrophosphohydrolase, MutT-like		Υ	Υ
nudE	Nudix hydrolase, active on adenosine(5')triphospho(5')adenosine, adenosine(5')diphospho(5')adenosine, ADP-ribose and NADH		Υ	Υ
nudG	CTP pyrophosphohydrolase		Υ	Υ
nusA	transcription pausing; L factor	Υ	Ý	Ý
nusB	transcription termination; L factor	Y	Ý	Ÿ
านรG	component in transcription antitermination	Ÿ	Ÿ	Ϋ́
	·	•	Ϋ́	Ϋ́
ogt	O-6-alkylguanine-DNA/cysteine-protein methyltransferase		ī	ı
ompG	outer membrane pore protein		Υ	
oaaD	putative subunit of multicomponent oxygenase, phenylacetic acid degradation			
oaal	putative phenylacetic acid degradation protein with thioesterase/thiol ester dehydrase-isomerase domain		Y	
oaaK	phenylacetyl-CoA ligase, phenylacetic acid degradation		Y	Y
oaaX	transcriptional repressor for phenylacetic acid degradation		Y	Y
oanC	pantothenate synthetase		Y	Υ
oarC	DNA topoisomerase IV, subunit A	Υ	Υ	Υ
oarE	DNA topoisomerase IV, subunit B	Υ	Υ	Υ
рерА	aminopeptidase A, a cyteinylglycinase		Y	
ерВ	aminopeptidase B, a cysteinylglycinase		Υ	Υ
pepD	aminopeptidase D (aminoacyl-histidine dipeptidase)		Υ	Υ
pepE	(alpha)-aspartyl dipeptidase		Υ	Υ
epN	aminopeptidase N, a cysteinylglycinase		Υ	Υ
рерР	proline aminopeptidase P II		Υ	Υ
pepQ	proline dipeptidase		Υ	Υ
рерТ	putative aminotripeptidase with Zn-dependent exopeptidase domain		Υ	Υ
ofkA	6-phosphofructokinase I		Υ	Υ
oflB	pyruvate formate lyase I, induced anaerobically		Υ	Υ
ogk	phosphoglycerate kinase		Υ	Υ
oheS	phenylalanine tRNA synthetase, alpha-subunit	Υ	Υ	Υ
oheT	phenylalanine tRNA synthetase, beta-subunit	Υ	Υ	Υ
ohoB	response regulator in two-component regulatory system with PhoR (or CreC), regulation of Pi uptake (OmpR family)		Υ	Υ
phoL	putative phosphate starvation-inducible protein (ATP-binding) with P-loop containing nucleoside triphosphate hydrolase domain		Y	Υ
phrB pin	deoxyribodipyrimidine photolyase (photoreactivation), FAD-binding e14 prophage; inversion of adjacent DNA		Y	Y
pinQ	Qin prophage; putative resolvase (recombinase)		Υ	
pinR	Rac prophage; putative transposon resolvase		Υ	Υ
pldA	outer membrane phospholipase A		Υ	
plsB	glycerolphosphate acyltransferase	Υ	Υ	Υ
plsC	1-acyl-sn-glycerol-3-phosphate acyltransferase	Υ	Υ	
plsX	fatty acid/phospholipid synthesis protein, methyltransferase domain		Υ	
pncB	nicotinate phosphoribosyltransferase	Υ	Υ	Υ
pnp	polynucleotide phosphorylase, has polyadenylase activity		Υ	Υ
polA	multifunctional DNA polymerase I: 5'->3' exonuclease (N-terminal); 3'->5' polymerase; 3'->5' exonuclease (C-terminal)	Υ	Y	Υ
polB	DNA polymerase II and 3' -> 5' exonuclease			
рохА	putative lysyl-tRNA synthetase with Class II aaRS and biotin synthetase domains	Υ	Υ	Υ
рра	inorganic pyrophosphatase	Υ	Υ	Υ
opiA	peptidyl-prolyl cis-trans isomerase A (rotamase A)		Υ	
ppiB	peptidyl-prolyl cis-trans isomerase B (rotamase B)		Υ	Υ
ppiC	peptidyl-prolyl cis-trans isomerase C (rotamase C)		Υ	Υ
opk	polyphosphate kinase, component of RNA degradosome		Υ	Υ
prc	carboxy-terminal protease for penicillin-binding protein 3			
prfA	peptide chain release factor RF-1	Υ	Υ	Υ
prfB	peptide chain release factor RF-2	Υ	Υ	Υ
prfC	peptide chain release factor RF-3; possible GTP-binding factor		Υ	Υ
oriA	primosomal protein N' (factor Y) directs replication fork assembly at D-loops, ATP-dependent		Y	Y
oriB	primosomal replication protein N		Y	
orkB	putative phosphoribulokinase with nucleoside triP hydrolase domain		Ϋ́	
ormA	methylation of 50S ribosomal subunit protein L11	Υ	Ý	Υ
oroA	gamma-glutamylphosphate reductase	Y	Ϋ́	Y
oroS	proline tRNA synthetase	Ϋ́	Ÿ	Ý
	phosphoribosylpyrophosphate synthetase	Ϋ́	Ϋ́	Ϋ́
are A	phosphatidylserine synthase (CDP-diacylglycerol-serine O-phosphatidyltransferase)	•	Y	Y
	high-affinity phosphate transport protein (ABC superfamily, atp_bind)	Υ	Ϋ́	Ϋ́
pssA		•	Ϋ́	Y
pssA pstB	phosphotransacetylase (phosphate acetyltransferase)			
pssA pstB pta	phosphotransacetylase (phosphate acetyltransferase)	Y	V	V
pssA pstB pta pth	peptidyl-tRNA hydrolase	Υ	Y	Y Y
prsA pssA pstB pta pth ptpS purB		Y Y	Y Y Y	

purC	phosphoribosylaminoimidazole-succinocarboxamide synthetase (SAICAR synthetase)	Υ	Υ	Y
pykA	pyruvate kinase II, glucose-stimulated	Y	Y	Y
pyrH	uridylate kinase	Y	Y	Y
gor	quinone oxidoreductase, NADPH-dependent	•	Ϋ́	Ϋ́
radC	associated with replication fork, possible DNA repair protein		•	•
rapA	ATPase associated with RNA polymerase and transcriptional activator		Υ	Υ
rbbA	ribosome-associated ATPase, ATP-binding domain (N-terminal)		Ϋ́	•
rbfA	30S ribosome-binding factor, role in processing of 16S rRNA		Ϋ́	Υ
rbsB	D-ribose transport protein (ABC superfamily, peri bind)		Ϋ́	•
rcsA	transcriptional activator of capsular/exo- polysaccharide synthesis (LuxR/UhpA family)		Ϋ́	Υ
rcsB	response regulator (positive) in two-component regulatory system with RcsC and YojN, regulates capsule		Ý	Ϋ́
	biosynthesis, cell division genes, OsmC expression, repressor of flhDC operon (LuxR/UhpA familiy)		·	•
rdgC	putative ribonuclease involved in removal of stalled replication fork with Rh-like domain		.,	
recA	DNA strand exchange and recombination protein with protease and nuclease activity		Y	
recB	exonuclease V, beta chain with recC and recD: 5' and 3' nuclease, ATPase, recombinase, helicase		Y	Υ
recC	exonuclease V, gamma chain with recB and recD: 5' and 3' nuclease, ATPase, recombinase, helicase		Y	
recD	exonuclease V, alpha chain with recC and recD: 5' and 3' nuclease, ATPase, recombinase, helicase		Y	Y
recE	Rac prophage; exonuclease VIII, ds DNA exonuclease, 5' -> 3'-specific		Y	Y
recF	gap repair protein with nucleoside triP hydrolase domain, part of RecFOR complex that targets RecA to ssDNA-dsDNA junction		Y	Υ
recG	DNA helicase, ATP-dependent resolution of Holliday junctions, branch migration		Υ	Υ
recJ	ssDNA exonuclease, 5'> 3'-specific, Mg-dependent	Υ	Υ	Υ
recN	protein used in recombination and DNA repair with nucleoside triphosphate hydrolase domain		Υ	Υ
recO	gap repair protein, part of RecFOR complex that targets RecA to ssDNA-dsDNA junction		Υ	
recQ	ATP-dependent DNA helicase		Υ	Υ
recR	gap repair protein with type I DNA topoisomerase domain, part of RecFOR complex that targets RecA to ssDNA-		Υ	
	dsDNA junction			
relB			Υ	
relE	Qin prophage; part of two-component toxin-antitoxin system with RelE, transcriptional repressor of relBE operon Qin prophage; part of two-component toxin-antitoxin system with RelB, transcriptional corepressor of relBE operon		Υ	
rep	Rep helicase, a single-stranded DNA-dependent ATPase		Υ	Υ
rfbC	dTDP-4,deoxyrhamnose 3,5 epimerase		Ϋ́	Ϋ́
rhlB	putative ATP-dependent helicase with nucleoside triP hydrolase domain		Ϋ́	Ϋ́
rhIE	putative ATP-dependent RNA helicase with P-loop hydrolase domain		Ý	
rho	transcription termination factor Rho; polarity suppressor	Υ	Ý	Υ
rhsC	RhsC protein in RhsC element	•	•	•
ribB	3,4 dihydroxy-2-butanone-4-phosphate synthase	Υ	Υ	Υ
ribD	bifunctional: diaminohydroxyphosphoribosylaminopyrimidine deaminase (N-terminal); 5-amino-6-(5-	•	Ϋ́	Ϋ́
with E	phosphoribosylamino) uracil reductase (C-terminal)	Υ	Υ	v
ribF	bifunctional: flavokinase; FAD synthetase	T	Ϋ́	Y Y
ribH	riboflavin synthase, beta chain		Ϋ́	Ϋ́
rimB rluB	Gm2251 methyltransferase of 23S rRNA		Ϋ́	Ϋ́
rluC	pseudouridine synthase (makes pseudouridine 2605 in 23 S RNA)		Ϋ́	Ϋ́
	23S rRNA pseudouridylate synthase pseudouridine synthase (pseudouridines 1911, 1915, 1917 in 23S RNA)		Ϋ́	Ϋ́
rluD	pseudouridine synthase (pseudouridines 1911, 1913, 1917 ili 233 KNA)		Ϋ́	1
rluE	RNase I, cleaves phosphodiester bond between any two nucleotides		Ϋ́	Υ
rna			Ϋ́	Ϋ́
rnb	RNase II, mRNA degradation		Ϋ́	Ϋ́
rnc	RNase III, ds RNA		-	
rnd	RNase D, processes tRNA precursor	Υ	Y Y	Y Y
rne	RNase E: endoribonuclease for rRNA processing and mRNA degradation	T	Y	
rng	RNase G (ribonuclease G)		Ϋ́	Y Y
rnhA	RNase HI, degrades RNA of DNA-RNA hybrids		Ϋ́	T
rnhB	RNAse HII		Ϋ́	
rnk	regulator of nucleoside diphosphate kinase	Υ	Ϋ́	v
rnpA	RNase P, protein C5 component, processes tRNA, 4.5S RNA	T	Y	Y Y
rnr	RNase R, 3'-5' exoribonuclease		Ϋ́	
rnt	RNase T, degrades tRNA, has exonuclease and ssDNAse activity	v		Y
rpe	D-ribulose-5-phosphate 3-epimerase RNase PH	Υ	Y	Y
rph			Y	Y
rpiB	ribose 5-phosphate isomerase B, also acts as allose 6-phosphate isomerase		Y	Y
rpID	50S ribosomal subunit protein L4, regulates expression of S10 operon		Y Y	Y Y
rplW	50S ribosomal subunit protein L23	v		
rpoA	RNA polymerase, alpha subunit	Y	Y	Y Y
rpoB	RNA polymerase, beta subunit	Y	Y	
rpoC	RNA polymerase, beta prime subunit	Y	Y	Y
rpoD	sigma D (sigma 70) factor of RNA polymerase	Υ	Y	Υ
rpoE	sigma E (sigma 24) factor of RNA polymerase, response to periplasmic stress (TetR/ArcR family)		V	V
rpoH	sigma H (sigma 32) factor of RNA polymerase; transcription of heat shock and stress proteins		Y	Y
rpoN	sigma N (sigma 54) factor of RNA polymerase		Y	Y
rpoS	sigma S (sigma 38) factor of RNA polymerase, major sigma factor during stationary phase		Y	Y
rpoZ	RNA polymerase, omega subunit		Y Y	Y Y
rpsE	30S ribosomal subunit protein S5		ı	ī
rpsJ	30S ribosomal subunit protein S10			

rpsM	30S ribosomal subunit protein S13		V	V
rraA	regulator of RNaseE	V	Y	Y
rrmJ	23 S rRNA methyltransferase	Υ	Y Y	Y Y
rsuA rtcR	16S rRNA pseudouridylate 516 synthase sigma N (sigma 54)-dependent transcriptional activator of RNA 3'-terminal phosphate cyclase (EBP familiy)		Ţ	T
ruvA	Holliday junction helicase, subunit A		Υ	
uvB	Holliday junction helicase, subunit B		Ϋ́	Υ
uvC	Holliday junction nuclease		Ϋ́	
sapA	peptide transport protein (ABC superfamily, peri_bind)		Ϋ́	
sbcB	exonuclease I, 3'> 5'-specific; deoxyribophosphodiesterase		Ϋ́	
sbcC	ATP-dependent dsDNA exonuclease		Υ	Υ
sbcD	ATP-dependent dsDNA exonuclease		Υ	
secA	preprotein translocase, ATPase secretion component (General Secretory Pathway)	Υ	Υ	Υ
secB	molecular chaperone in protein export, enhances activity of SecA (General Secretory Pathway)		Υ	Υ
selB	selenocysteinyl-tRNA-specific translation factor		Υ	Υ
seqA	negative modulator of replication initiation		Υ	
serC	3-phosphoserine/phosphohydroxythreonine aminotransferase		Υ	Υ
serS	serine tRNA synthetase; also charges selenocystein tRNA with serine	Υ	Υ	Υ
sgbH	3-keto-L-gulonate 6-phosphate decarboxylase		Υ	
sgcE	KpLE2 phage-like element; putative ribulose-phosphate 3-epimerase		Υ	
slyD	FKBP-type peptidyl prolyl cis-trans isomerase (rotamase)		Υ	Υ
smf_2	split CDS, fragment 1		Υ	
smpB	trans-translation protein, binds tmRNA and tRNA		Υ	Υ
speE -	spermidine synthase (putrescine aminopropyltransferase)	.,	Y	Y
spoT	bifunctional: (p)ppGpp synthetase II; guanosine-3',5'-bis pyrophosphate 3'-pyrophosphohydrolase	Υ	Y	Y
spoU	putative tRNA/rRNA methyltransferase		Y	Y
srIR	transcriptional repressor for glucitol utilization (DeoR family)		Y	Y
srmB	ATP-dependent RNA helicase	V	Y	Y
ssb	ssDNA-binding protein controls activity of RecBCD nuclease	Υ	Y	Y
sspA	stringent starvation protein A, activator of transcription for bacteriophage P1 late genes		Y	Y Y
sspB	stringent starvation protein B, specificity factor for ClpXP protease		Y Y	ī
ssuB stfR	alkanesulfonate transport protein (ABC superfamily, atp_bind)		ĭ	
sucA	Rac prophage; putative tail fiber protein		Υ	Υ
sucA	2-oxoglutarate decarboxylase, component of the 2-oxoglutarate dehydrogenase complex, thiamin-binding	Υ	Ϋ́	Ý
sufB	dihydrolipoyltranssuccinate transferase, component of the 2-oxoglutarate dehydrogenase complex putative transport protein associated with Fe-S cluster assembly	•	Ϋ́	Ý
sufC	putative transport protein associated with Fe-S cluster assembly (ABC superfamily, atp. bind)		Ϋ́	Ϋ́
sufD	required for stability of Fe-S component of FhuF		Ϋ́	Ϋ́
surA	peptidyl-prolyl cis-trans isomerase (PPlase), involved in maturation of outer membrane proteins		Ϋ́	Ϋ́
tadA	tRNA-specific adenosine deaminase	Υ	Ϋ́	
tag	3-methyl-adenine DNA glycosylase I, constitutive	•	Ý	Υ
tatA	twin-arginine translocase subunit, sec-independent protein export		Ý	•
tatB	twin-arginine translocase subunit, sec-independent protein export		Ϋ́	
tatC	twin-arginine translocase subunit, sec-independent protein export		Y	
tatD	DNase, cytoplasmic (possibly with b3841)		Y	
tatE	component of sec-independent translocase		Υ	
tdcD	propionate kinase/acetate kinase C. anaerobic		Υ	Υ
dcF	conserved protein with YjgF-like domain		Υ	Υ
gt	tRNA-guanine transglycosylase		Υ	Υ
hrS	threonine tRNA synthetase	Υ	Υ	Υ
thyA	thymidylate synthetase	Υ	Υ	Υ
tig	peptidyl-prolyl cis/trans isomerase (trigger factor), molecular chaperone involved in cell division		Υ	Υ
tkrA	2-keto-D-gluconate reductase (2-ketoaldonate reductase)		Υ	Υ
tktA	transketolase 1 thiamin-binding, isozyme	Υ	Υ	Υ
tmk	thymidylate kinase	Υ	Υ	Υ
naA	tryptophan deaminase, PLP-dependent		Υ	Υ
olB	required for outer membrane integrity, uptake of group A colicins, and translocation of phage DNA		Υ	
tonB	energy transducer; uptake of iron, cyanocobalimin; sensitivity to phages, colicins		Υ	
topA	DNA topoisomerase type I, omega protein	Υ	Υ	Υ
орВ	DNA topoisomerase III, type 1		Υ	Υ
orA	trimethylamine N-oxide reductase system I, with TorC		Υ	
orC	cytochrome c-type protein in TMAO respiration; with TorA, also negative regulator of tor operon		Y	
orD	cytoplasmic chaperone involved in maturation of TorA		Y	
piA	triosephosphate isomerase	Υ	Υ	Υ
ra8_3	KpLE2 phage-like element; transposase for IS30			_
rmA	tRNA (uracil-5-)-methyltransferase	Y	Y	Υ
trmD	tRNA (guanine-7-)-methyltransferase	Y	Y	
trmE	GTPase involved in tRNA modification and in thiophene and furan oxidation	Y	Υ	Υ
trmU	tRNA (5-methylaminomethyl-2-thiouridylate)-methyltransferase	Υ	Y	Υ
rpR	transcriptional repressor for tryptophan biosynthesis (TrpR family)		Υ	
trpS	tryptophan tRNA synthetase	Υ	Y	Y
rxC	thioredoxin 2, redox factor		Υ	Υ
tsf ttuC	protein chain elongation factor EF-Ts	Υ	Υ	Υ
	putative tartrate dehydrogenase		Υ	Υ

tufB	protein chain elongation factor EF-Tu (duplicate of tufB)	Υ	Υ	Υ
	protein chain elongation factor EF-Tu; possible GTP-binding factor (duplicate of tufA)	Υ	Υ	Υ
typA	GTP-binding elongation factor family protein with P-loop containing nucleoside triphosphate hydrolase domain		Υ	Υ
tyrA	bifunctional: chorismate mutase T (N-terminal); prephenate dehydrogenase (C-terminal)		Υ	Υ
tyrR	transcriptional regulator of aromatic amino acid biosynthesis and aromatic amino acid transport (EBP family)		Υ	Υ
tyrS	tyrosine tRNA synthetase	Υ	Υ	Υ
ubiC	chorismate pyruvate lyase		Υ	Υ
ubiE	bifunctional: 2-octaprenyl-6-methoxy-1,4-benzoquinone methylase; S-adenosylmethionine:2-DMK methyltransferase		Υ	Y
ubiF	2-octoprenyl-3-methyl-6-methoxy-1,4-benzoquinone hydroxylase		Υ	Υ
ubiG	bifunctional: 3-demethylubiquinone-9 3-methyltransferase; 2-octaprenyl-6-hydroxy phenol methylase		Υ	Υ
ubiH	2-octaprenyl-6-methoxyphenol hydroxylase, FAD/NAD(P)-binding		Υ	Υ
ubiX	3-octaprenyl-4-hydroxybenzoate carboxy-lyase		Υ	
ucpA	putative oxidoreductase, NAD(P)-binding domain		Υ	Υ
umuC	component of DNA polymerase V with UmuD		Υ	
ung	uracil-DNA-glycosylase		Υ	Υ
uppS	undecaprenyl pyrophosphate synthetase (di-trans,poly-cis-decaprenylcistransferase)	Υ	Υ	Υ
usg	putative dehydrogenase with NAD(P)-binding domain and Glyceraldehyde-3-phosphate dehydrogenase-like, C-terminal domain	Υ	Υ	Υ
uup	putative transport protein (ABC superfamily, atp_bind)		Υ	Υ
uvrA	DNA excision repair enzyme subunit, with UvrBC		Υ	Υ
uvrB	ATP-dependent DNA excision repair enzyme UvrAC		Υ	Υ
uvrC	DNA exisiton repair enzyme together with UvrAB	Υ	Υ	Υ
uvrD	DNA-dependent ATPase I and helicase II		Ϋ́	Ϋ́
ıvrY	putative regulator in two-component regulatory system with BarA (LuxR/UhpA familiy)		Ý	Ϋ́
/alS	valine tRNA synthetase	Υ	Ý	Ý
ves	cold-shock induced protein	-	Ý	•
vcaH	GDP-mannose mannosyl hydrolase, colanic acid synthesis		•	
xerC	site-specific tyrosine recombinase		Υ	
kerD	site-specific tyrosine recombinase		Ϋ́	Υ
kni	exonuclease IX, 5'-3' exonuclease		Ϋ́	Ϋ́
kseA	exonuclease VII, large subunit		Ϋ́	Ϋ́
kseB	exonuclease VII, small subunit		Ϋ́	'
kthA			Ϋ́	Υ
	exonuclease III		ī	ī
yaaX	conserved hypothetical protein		Υ	Υ
yacG	conserved hypothetical protein			
yacL	conserved hypothetical protein		Y	Y
yadB	putative glutamyl t-RNA synthetase with nucleotidylyl transferase domain		Y	Y
yadF	carbonic anhydrase		Y	Υ
yaeJ	conserved protein with RF2 (polypeptide chain release factor 2) domain		Y	
yafM	conserved hypothetical protein		Υ	
yagA	CP4-6 prophage; putative transposase			
/agl	CP4-6 prophage; putative transcriptional regulator (IcIR family)			
yahA	putative transcriptional repressor (LuxR/UhpA family)			
/ahO	conserved hypothetical protein			
/ajQ	conserved protein			
/baB	conserved hypothetical protein		Υ	Υ
	annon and protoin		Υ	Υ
ybaD	conserved protein		Y Y	Y Y
/baD /baK	conserved hypothetical protein with YbaK-like domain		Υ	Υ
/baD /baK /baV	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2		Y Y Y	Y Y Y
/baD /baK /baV	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain		Y Y Y	Y Y Y
/baD /baK /baV /baX	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain		Y Y Y	Y Y Y
/baD /baK /baV /baX /baZ /bbA	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind)		Y Y Y Y Y	Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind)		Y Y Y Y Y Y	Y Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL ybbN	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind)		Y Y Y Y Y Y	Y Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL ybbN	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind)		Y Y Y Y Y Y	Y Y Y Y Y Y
/baD /baK /baV /baX /baZ /bbA /bbL /bbN /bcJ	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain		Y Y Y Y Y Y	Y Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL ybbN ybcJ	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein		Y Y Y Y Y Y	Y Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL ybbN ybcJ ybcK	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase		Y Y Y Y Y Y Y	Y Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL ybbN ybcJ ybcK ybdN	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain		Y Y Y Y Y Y Y	Y Y Y Y Y Y
/baD /baK /baV /baX /baZ /bbA /bbL /bbN /bcJ /bcK /bdN /bdO	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy)	Y	Y Y Y Y Y Y Y	Y Y Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL ybbN ybcJ ybcK ybdN ybdO ybdQ	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain	Y	Y Y Y Y Y Y Y Y	Y Y Y Y Y Y
rbaD rbaK rbaV rbaX rbaZ rbbA rbbL rbbN rbcJ rbcK rbdN rbdQ rbdQ rbeA rbeB	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein	Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y
ybaD ybaK ybaV ybaZ ybbA ybbL ybbN ybcJ ybcK ybdO ybdQ ybeA ybeB ybeD	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein	Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
/baD /baK /baV /baX /baZ /bbA /bbL /bbN /bcJ /bcK /bdN /bdQ /beA /beB /beD	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein		Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
rbaD rbaK rbaV rbaX rbaZ rbbA rbbL rbbN rbcJ rbcK rbdN rbdQ rbeA rbeB rbeD rbeY rbgC	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain		Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
/baD /baK /baV /baX /baZ /bbA /bbL /bbD /bcJ /bcJ /bdQ /bdQ /beB /beD /beD /beY /bgC	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain putative carboxylase		Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
/baD /baK /baV /baX /baZ /bbA /bbD /bbN /bcJ /bcK /bdN /bdO /bdQ /beB /beB /beB /bgJ /bgJ	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain putative carboxylase putative transport protein (ABC superfamily, atp_bind)	Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
/baD /baK /baV /baX /baZ /bbA /bbL /bbN /bcJ /bcK /bdO /bdQ /beA /beB /beD /bgJ /bgC	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain putative carboxylase putative transport protein (ABC superfamily, atp_bind) putative phosphatase/sulfatase with NAD(P)-binding domain		Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL ybbN ybcJ ybcK ybdO ybdQ ybeA ybeB ybeB ybeP ybeY ybgJ ybbF	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain putative carboxylase putative transport protein (ABC superfamily, atp_bind) putative phosphatase/sulfatase with NAD(P)-binding domain conserved protein, with phosphatase-like domain	Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
ybaD ybaK ybaX ybaZ ybbA ybbL ybbN ybcJ ybcK ybdO ybdQ ybeA ybeB ybeD ybeY ybgC ybgF ybgF ybhK ybiV	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein monserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain putative carboxylase putative transport protein (ABC superfamily, atp_bind) putative phosphatase/sulfatase with NAD(P)-binding domain conserved protein, with phosphatase-like domain conserved protein, phosphatase-like domain	Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
ybaD ybaK ybaY ybaX ybaZ ybbA ybbL ybbN ybcJ ybcK ybdN ybdQ ybeB ybeB ybeD ybbeY ybgJ ybhF ybjU ybjU ybjU	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain putative carboxylase putative transport protein (ABC superfamily, atp_bind) putative phosphatase/sulfatase with NAD(P)-binding domain conserved protein, with phosphatase-like domain conserved protein, phosphatase-like domain conserved hypothetical protein	Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
ybaD ybaK ybaK ybaV ybaX ybaZ ybbA ybbL ybbN ybcJ ybcK ybdQ ybeA ybeB ybeD ybeY ybgC ybgJ ybhF ybiV ybiV	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain putative carboxylase putative transport protein (ABC superfamily, atp_bind) putative phosphatase/sulfatase with NAD(P)-binding domain conserved protein, with phosphatase-like domain conserved protein, phosphatase-like domain conserved hypothetical protein conserved hypothetical protein	Υ	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y
ybaD ybaK ybaV ybaX ybaZ ybbA ybbL ybbC ybbC ybbC ybbQ ybeA ybeB ybeD ybbF ybbI ybjQ ybiV ybjQ ybjQ ybjQ ybjQ ybjQ ybjQ ybjQ ybjQ	conserved hypothetical protein with YbaK-like domain conserved hypothetical protein with domain like RuvA domain 2 putative (aluminum) resistance protein with adenine nucleotide alpha hydrolase domain putative methylated DNA-protein cysteine methyltransferase, C-terminal domain putative transport protein (ABC superfamily, atp_bind) putative putrescine/spermidine transport protein (ABC superfamily, atp_bind) putative protein prenylyltransferase domain putative RNA-binding protein DLP12 prophage; putative recombinase conserved protein with adenine nucleotide alpha hydrolase domain putative transcriptional regulator with periplasmic binding protein domain (LysR familiy) universal stress protein UP12, flavoprotein, ETFP adenine nucleotide-binding domain conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein conserved hypothetical protein with thioesterase/thiol ester dehydrase-isomerase domain putative carboxylase putative transport protein (ABC superfamily, atp_bind) putative phosphatase/sulfatase with NAD(P)-binding domain conserved protein, with phosphatase-like domain conserved protein, phosphatase-like domain conserved hypothetical protein	Υ	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y	Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y Y

1			.,	.,
ycbL	putative enzyme with metallo-hydrolase/oxidoreductase domain		Y	Y
ycbY	putative methyltransferase with S-adenosyl-L-methionine-dependent methyltransferase domain		Y	Υ
yccD	conserved protein		Y	.,
yccR	conserved protein		Y	Υ
yccT	conserved hypothetical protein		Y	
yccU	putative NAD(P)-binding enzyme		Y	Y
yccX	putative acylphosphatase		Υ	Y
ycdH	putative oxidoreductase component with FMN-binding split barrel domain	Υ	Υ	
ycdK	conserved protein with YjgF-like domain	ı	Ϋ́	
ycdM	putative enzyme with luciferase-like ATPase domain		Ϋ́	Υ
yceA	conserved hypothetical protein with Rhodanese/Cell cycle control phosphatase domain		Ϋ́	ī
yceB yceH	conserved hypothetical protein		Ϋ́	Υ
ycfC	conserved hypothetical protein membrane-associated protein of unknown function	Υ	'	'
ycfF	putative inhibitor of protein kinase C, contains a transferase domain	•	Υ	Υ
ycfH	putative metallo-dependent hydrolase (with domain)		Ϋ́	Ϋ́
ycfL	conserved hypothetical protein		Ý	•
ycfQ	putative regulator with homeodomain-like DNA binding domain (TetR/AcrR family)		•	
ycfX	putative transcriptional regulator with ATPase domain (NagC/XyIR (ROK) family		Υ	Υ
ycgE	putative transcriptional repressor with DNA-binding domain (MerR family)		Ϋ́	Ϋ́
ycgF	putative FMN-linked oxidoreductase		Ϋ́	Ϋ́
ychA	putative transcriptional regulator with tetratricopeptide repeats (TPR) domain		Y	Y
ychF	putative GTP-binding protein with nucleoside triP hydrolase domain			
ychJ	conserved hypothetical protein		Υ	Υ
ychN	putative phosphatase		Υ	
yciH	conserved hypothetical protein with eIF1-like domain		Υ	Υ
ycil	conserved hypothetical protein		Υ	Υ
yciO	conserved protein with YrdC/RibB domain		Υ	Υ
yciU	conserved hypothetical protein		Υ	Υ
ycjC	putative oxidoreductase/putative regulator with RmlC-like domain and DNA-binding domain		Υ	Υ
ycjV	putative sugar transport protein (ABC superfamily, atp_bind)		Υ	
ycjY	conserved hypothetical protein, alpha/beta-hydrolase domain		Υ	Υ
ydaG	Rac prophage		Υ	
ydaY	Rac prophage			
ydcJ	conserved hypothetical protein		Υ	Υ
ydcP	putative collagenase		Υ	Υ
ydcQ	putative regulator with DNA-binding domain		Υ	
ydcY	unknown CDS		Υ	
yddH	putative enzyme with FMN-binding split barrel domain			
yddJ	unknown CDS			
yddV	conserved protein			
yddW	putative (trans)glycosidase		Υ	
yddX	unknown CDS			
ydeP	putative formate dehydrogenase, related to acid resistance with formate dehydrogenase/DMSO reductase, domains 1-3 and ADC-like domain		Υ	
vdfD	Qin prophage		Υ	
ydfR ydfT	Qin prophage; putative antitermination potein Q		'	
ydfV	Qin prophage			
ydfW	Qin prophage			
ydfX	Qin prophage			
ydfZ	unknown CDS			
ydgH	conserved hypothetical protein		Υ	Υ
ydgJ	putative NAD(P)-binding dehydrogenase, with glyceraldehyde-3-P dehydrogenase-like and NAD(P)-binding		Ϋ́	Ϋ́
, -3-	domains			
ydgM	putative 4Fe-4S ferredoxin-type protein		Υ	
ydgP	conserved protein		Υ	
ydgT	putative hemolysin expression modulating protein HHA domain		Υ	
ydhD	conserved protein with thioredoxin-like domain		Υ	Υ
ydhF	putative oxidoreductase, NAD(P)-linked domain		Υ	Υ
ydhH	conserved hypothetical protein with actin-like ATPase domain		Υ	Υ
ydhL	unknown CDS		Y	.,
ydhM	putative transcriptional regulator with homeodomain-like DNA binding domain (TetR/AcrR family)		Y	Y
ydhR	conserved hypothetical protein		Y	Y
ydhV	putative aldehyde ferridoxin oxidoreductase		Y	Y
ydiA	conserved protein with periplasmic binding protein II-like domain	v	Y	Y Y
ydiB	quinate/shikimate 5-dehydrogenase, NAD(P)-binding	Υ	Y Y	Y Y
ydiH ydiF	unknown CDS nutative transcriptional regulator with DNA-hinding Winged helix domain (DeoP family)		Ϋ́Υ	Ϋ́Υ
ydjF yeaB	putative transcriptional regulator with DNA-binding Winged helix domain (DeoR family) conserved hypothetical protein, MutT-like		Ϋ́Υ	ı
уеаБ yeaG	conserved hypothetical protein, muti-like conserved protein, nucleotide triphosphate hydrolase domain		Ϋ́	Υ
yeaG yeaM	putative transcriptional regulator (AraC/XylS family)		Ϋ́	'
yeaZ	putative glycoprotein endopeptidase, actin-like ATPase domain		Ϋ́	Υ
yebC	conserved protein with YebC-like domain	Υ	Ϋ́	Ϋ́
yecC	putative amino acid transport protein (ABC superfamily, atp_bind)	•	Ϋ́	•
yedO	D-cysteine desulfhydrase, PLP-dependent enzyme		Ϋ́	Υ

yedW yeeN	putative response regulator in two-component regulatory system (OmpR family) conserved protein with YebC-like domain	Υ	Y Y	Y Y
yeeS	CP4-44 prophage; putative DNA repair protein (RadC family)	•	•	•
egL	conserved protein with Integrin A (or I) domain		Υ	
/eiP	putative elongation factor		Υ	Υ
/ejF	putative oligopeptide transport protein (ABC superfamily, atp_bind)		Υ	Υ
/ejH	putative ATP-dependent helicase with nucleoside triP hydrolase domain		Y	
yfaO faV	putative enzyme (Nudix hydrolase)		Y	Υ
yfaX yfbQ	putative transcriptional regulator with DNA-binding Winged helix domain (IcIR family) putative PLP-dependent aminotransferase		Y Y	Υ
/fcB	N5-glutamine methyltransferase, modifies ribosomal protein L3		Ϋ́	Ϋ́
yfcK	conserved protein, FAD/NAD(P)-binding domain		Ϋ́	Ϋ́
yfeR	putative transcriptional regulator with periplasmic binding protein domain (LysR family)		Υ	
yffH	conserved protein with MutT-like domain		Υ	
yfgB	putative pyruvate formate lyase activating enzyme 2		Υ	Υ
/fhA	putative response regulator in two-component regulatory system (EBP family)		Y	Y
yfhM .fi.^	conserved protein with prenyltransferase domain		Y Y	Y Y
yfiA yfiB	ribosome associated factor, stabilizes ribosomes against dissociation putative outer membrane protein		Ϋ́	ī
yfiD	putative outer memorane protein putative formate acetyltransferase withPFL-like glycyl radical domain		Ϋ́	Υ
,∙∠ √fiF	putative tRNA/rRNA methyltransferase		Ϋ́	Ϋ́
yfiQ	putative acyl-CoA synthetase, NAD(P)-binding, ATP-binding		Υ	Υ
yfjB	NAD kinase	Υ	Υ	Υ
yfjK	CP4-57 prophage		Υ	Υ
/fjW	CP4-57 prophage			
/fjY	CP4-57 prophage; putative DNA repair protein putative epimerase/isomerase with Xylose isomerase-like domain		V	V
ygbM ygcP	putative anti-terminator regulatory protein with FMN-linked oxidoreductase domain		Y Y	Y Y
/gcW	putative deoxygluconate dehydrogenase with NAD(P)-binding domain		Ϋ́	Ϋ́
/gdP	nucleotide hydrolase, acts on adenosine(5')-pentaphospho-(5')-adenosine (Nudix family)		Ý	Ý
ygeG	conserved protein with tetratricopeptide repeats (TPR) domain			
ygeV	putative transcriptional regulator protein with GAF, PYP-like sensor, NTP hydrolysis, and FIS-like domains		Υ	Υ
/gfA	putative ligase	Υ	Y	Y
/gfF	putative oxidoreductase, NAD(P)-binding domain		Y	Υ
/gfH /gfT	propionyl-CoA:succinate-CoA transferase putative oxidoreductase: Fe-S subunit (N-terminal); nucleotide-binding domain (C-terminal)		Y Y	
yggH	tRNA (m7G46) methyltransferase, SAM-dependent		Ý	Υ
yggS	putative enzyme with PLP-binding domain		Ϋ́	Ϋ́
yggV	dITP/dXTP pyrophosphatase		Y	Y
yggW	putative oxidase		Υ	Υ
yggX	conserved hypothetical protein		Υ	Υ
yghA	putative oxidoreductase, NAD(P)-binding domain		Y	Y
ygiC	putative glutathione-like synthetase		Y	Y
ygiF ygjH	conserved hypothetical protein putative tRNA synthetase		Y Y	Y Y
yhbC	conserved hypothetical protein with YhbC-like domain	Υ	Ϋ́	Ϋ́
yhbG	putative transport protein (ABC superfamily, atp_bind)	Ϋ́	Ϋ́	-
yhbH	putative sigma N (sigma 54) modulator		Y	Υ
yhbJ	conserved protein, nucleotide triphosphate hydrolase domain		Υ	Υ
/hbU	putative protease		Υ	Υ
/hbV	putative protease		Y	Y
yhbY	putative RNA binding protein		Y Y	Y Y
yhcC yhcF	putative enzyme putative transcriptional regulator		Y	Y
/hcQ	putative transcriptional regulator putative membrane located multidrug resistance protein			
/hdJ	putative methyltransferase with SAM-dependent methyltransferase domain			
/hdZ	putative amino acid transport protein (ABC superfamily, atp_bind)		Υ	
/heS	putative transport protein (ABC superfamily, atp_bind)		Υ	Υ
/hfN	fructoselysine-6-P deglycase		Y	
/hfR /haC	putative transcriptional repressor with DNA-binding Winged helix domain (GntR family)		Y	Y
/hgG /hhF	putative transcriptional regulator with DNA-binding Winged helix domain	Υ	Υ	Υ
/hhF /hhP	putative methyltransferase with SAM-dependent methyltransferase domain small ubiquitous RNA-binding protein required for normal growth, cytoplasmic	ı	Ϋ́Υ	Ϋ́
hiF	putative transcriptional regulator (LuxR/UhpA family)		Ϋ́	Ϋ́
hiR	putative methyltransferase with S-adenosyl-L-methionine-dependent methyltransferase domain		Ϋ́	Ϋ́
riaJ	transcriptional repressor (IcIR family)		Y	Y
ribΑ	putative lyase with ARM repeat domain		Υ	Υ
/ibK	putative tRNA/rRNA methyltransferase		Υ	Υ
/ibL	conserved protein		Y	Y
/icC	conserved protein	Υ	Y	Y
/idA	conserved protein, phosphatase-like domain		Y	Y
/idC	preprotein translocase, substrates includes membrane components of ATP synthase and the SecYEG translocase		Y	
/idP	putative transcriptional repressor with DNA-binding Winged helix domain (GntR family)	Υ		

yieM	conserved protein with Integrin A (or I) domain		Y	Y
yieN	putative transcriptional regulator		Y	Y
yigW_2	conserved hypothetical protein; possible second part of tatD		Υ	Υ
yigZ	putative elongation factor, with GTP-binding EF-G domain		Υ	Υ
yihA	putative GTPase with nucleoside triP hydrolase domain, involved in coordination of cell cycle	Υ	Υ	Υ
yihl	conserved protein		Υ	Υ
yihQ	putative alpha-xylosidase		Υ	Υ
yihU	putative oxidoreductase with NAD(P)-binding domain		Υ	
yihW	putative glycerol-3-phosphate regulon repressor with DNA-binding Winged helix domain (DeoR family)		Y	Y
yihZ	D-Tyr-tRNA(Tyr) deacylase		Y	Y
yiiD	putative acyltransferase		Y	Y
yjaD	conserved hypothetical protein, MutT-like protein		Y	Y
yjbJ 	unknown CDS with YmbJ domain	Υ	Y Y	Y Y
yjeE	putative enzyme with nucleoside triP hydrolase domain	T	Ϋ́	Ϋ́
yjeF yjeQ	putative kinase with ribokinase-like domain putative enzyme with 2 nucleoside triP hydrolase domains	Υ	Ϋ́	Ϋ́
yjeQ yjfQ	putative transcriptional repressor with DNA-binding Winged helix domain (DeoR family)	Ϋ́	Ý	Ϋ́
yjfZ	unknown CDS	•	•	•
yjgD	conserved hypothetical protein		Υ	Υ
yjgH	putative translation factor		Ϋ́	Ϋ́
yjgl yjgl	putative oxidoreductase, NAD(P)-binding domain		Ϋ́	•
yjgL	conserved protein		Ϋ́	Υ
yjhG	putative dehydratase		Ϋ́	Ϋ́
yjhH	KpLE2 phage-like element; putative synthase		Y	Y
yjhl	KpLE2 phage-like element; putative transcriptional repressor (IcIR family)		Υ	
yjhR	KpLE2 phage-like element; frameshift suppressor contains phospolipase D/nuclease			
yjjl	conserved hypothetical protein with PFL-like glycyl radical domain		Υ	Υ
yjjV	putative hydrolase with metallo-dependent hydrolase domain		Υ	Υ
ykfG	CP4-6 prophage; putative DNA repair protein			
ykfl	CP4-6 prophage			
ykiA	unknown CDS			
ymdA	conserved hypothetical protein			
ymfB	bifunctional: thiamin pyrimidine pyrophosphate hydrolase; thiamin pyrophosphate hydrolase		Υ	Υ
ymfP	e14 prophage			
ymfQ	e14 prophage		.,	
ymfS	e14 prophage		Y	
ymgD	unknown CDS		Υ	
ynaK	Rac prophage			
ynbD	putative phosphatase, phosphotyrosine protein with phosphatase II domain putative transcriptional regulator with DNA-binding Winged helix domain (GntR familiy)		Υ	Υ
yncC yncD	putative outer membrane porin protein		Ϋ́	ı
yncH	unknown CDS		Ϋ́	
yncJ	unknown CDS		'	
yneC	conserved hypothetical protein		Υ	
yneG	conserved hypothetical protein		Ϋ́	
ynfB	conserved hypothetical protein		Ý	Υ
ynfE	putative oxidoreductase subunit with Formate dehydrogenase/DMSO reductase, domains 1-3 and ADC-like		•	•
ynfK	putative dethiobiotin synthetase with nucleotide triphosphate hydrolase domain		Υ	Υ
ynfL	putative transcriptional regulator with periplasmic binding protein domain (LysR family)			
ynfN	Qin prophage			
yniC	putative enzyme, with a phosphatase-like domain		Υ	
yoaA	putative ATP-dependent helicase with nucleotide triphosphate hydrolase domain, SOS repair		Υ	Υ
yojN	putative sensory histidine kinase (phosphotransfer intermediate) associated with the RcsBC two-component		Υ	
	regulatory system			
ypdD	putative PTS family Hpr component (N-terminal); enzyme I component (middle); enzyme IIA component (C-		Υ	Υ
	terminal)			
yphE	putative sugar transport protein (ABC superfamily, atp_bind)		Y	V
yqaB	putative phosphoglucomutase, contains a phosphatase-like domain		Y	Y Y
yqeA	putative carbamate kinase with carbamate kinase-like domain putative acetyl-CoA acetyltransferase with thiolase domain		Y Y	Ţ
yqeF	putative acetyl-CoA acetyltransferase with thiolase domain putative transcriptional regulator with C-terminal, effector domain of the bipartite response regulator		ſ	
yqel yqgF	conserved hypothetical protein with ribonuclease H-like domain	Υ	Υ	Υ
yqgr yraL	putative enzyme with Cobalt precorrin-4 methyltransferase domain	•	Ϋ́	Ϋ́
yra∟ yraM	putative enzyme with 3 periplasmic binding protein-like domains		Ϋ́	Ϋ́
yraN yraN	conserved hypothetical protein with restriction endonuclease-like domain		Ϋ́	Ϋ́
yrbF	putative transport protein (ABC superfamily, atp_bind)		Ϋ́	Ϋ́
, . ~ .	putative RNA-binding protein with unique protein fold, with YrdC/RibB domain	Υ	Ý	Ϋ́
	putative DNA topoisomerase	-	Ý	Ϋ́
yrdC			•	
yrdC yrdD	putative sugar transport protein (ABC superfamily, atp_bind)		Υ	
yrdC yrdD ytfR	·	Y	Y Y	
yrdC yrdD	putative sugar transport protein (ABC superfamily, atp_bind)	Y		
yrdC yrdD ytfR ytfS	putative sugar transport protein (ABC superfamily, atp_bind) putative sugar transport protein with P-loop containing nucleoside triphosphate hydrolase domain	Y	Υ	Y
yrdC yrdD ytfR ytfS znuA	putative sugar transport protein (ABC superfamily, atp_bind) putative sugar transport protein with P-loop containing nucleoside triphosphate hydrolase domain high-affinity Zn transport protein (ABC superfamily, peri_bind)	Y	Y Y	Y

Supplementary Table 3. Protein-protein interactions reported here that have previously reported by DIP, STRING, BIND and PROLINKS databases

Protect	BUTLAND	-	DIP DAT	ABASE		OTHER DATABAS	SE .
Decc Decc	ProtA	ProtB		Exp method	STRING		PROLINKS
Sept							
Sept	acpP						
bassa	acpP						
dnaE				N+O			
dnakd dnakd t+3 E dnakd Habel B Habel Habel <td></td> <td></td> <td>_</td> <td>G+I+K</td> <td>S</td> <td></td> <td></td>			_	G+I+K	S		
dnak gpE 1+3 P B eno ppp 1+2+3 N nusA rpoA A S nusA rpoA A S nusA rpoA A S nusA rpoA A S nusB rpoB A S nusB rpoB A S pbB 11-2 D+L pbB me L-M-O S me dnak L S me dnak L S poA rpoZ B S poA rpoZ B S poA rpoZ B S poC rpoZ F S poC rpoZ F S poC rpoZ F S poC rpoZ P S sech pac D F S				_			
Pop Pop				P		В	
Nussa					S		
Nussa			1+2+3		•		
Nuss							
pheS			1	**	Ü		
DPL			4:0	Α			
pstB			1+2	D+I			
me enc							
me							
POPA							
POD							н
POB POC	rpoA	rpoZ					
POC POD A + H+J S POC POC POZ F S S S S S POC POC POZ F S S S S S S S S S			112		S		
POC POZ F S S S S S S S S S			172		S		
Sec	rpoC	rpoZ		F	S		
Success			1				
### ### ### ### ### ### ### ### ### ##			1+2	D+E+L	5		
tef tufA P acpP acpS bolA ydnD S capA dacK cbpA dnakk S clpA rpoB clpA spoT H dlcC cbpA dnakk S clpA rpoB clpA spoT H dlpP dnakk S clpX clpB S cspB S	thdF	gidA			s		
bold ydhD S cadA IdCC S cbpA dnaK S clpA rpoB H clpA spoT H clpA cspB S cspB cspB S cspB cspC S cspB cspE S dnaC ds B </td <td></td> <td>tufA</td> <td></td> <td>Р</td> <td></td> <td></td> <td></td>		tufA		Р			
CadA							
cbpA dnak S clpA spoT H clpA spoT H clpA spoT H clpA spoT S clpA spoT S clpX clpB S cspA cspC S cspB cspA S cspB cspA S cspB cspB S cspB cspA S dnaC dnaX S blq cspA B cspA blpA H <					S		
Color	cbpA	dnaK					
cipP							
clpX					s		п
cspB cspB cspE S cspB cspE S dnaE dnaX S dnaC holE S dnaA hscA S dnaQ dnaX S dnaQ holE S gadA gadB S hfq rpB S holD S B hfyC hyB S hfyC hyB S <	clpX	clpB			S		
cspB cspE S dnaE holE S dnaE holE S dnaC dnaX S dnaQ holE S gadA gadB S hfq rpB S holD s B hard hblD S hsolD hsolD S hsolD s B hypC hybC hybC h							
daaE dnaX S daaE holE S daaK hscA S daaC dnaX S daaC dnaX S daaC holE S gadA gadB S hfq rpB S hfq rpB S himA himD B holB holA S B holB holA S B H holC							
dalAK hscA S daAQ dnaX S dnaQ holE S gadA gadB S hfq rpB S hfq rpB S hfq rpBG S himA himD B holB dnaX S B holB dnaX S B holB dnaX S B holC holD S B hysC hysE S B hypC hysE S B hypC hysE S B hypC hysD S B infC rpsB S S lipA S B <t< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td></t<>							
dnaQ							
daaQ							
gadA gadB S	dnaQ						
hfq	gadA				S		
htig							
himA himD B holB dnaX S B holC holA S B H holC holD S B H holC holD S B B hsdM hsdR S B H hsdM hsdR S B B hypC hybE S B B hypC hybE S B B hypC hybE S B B infC rpsB S S Interpretable S infC rpsR S S Interpretable S Interpretable S Interpretable S Interpretable S Interpretable S Interpretable S							
holB holA S B H H holC dnaX S B H H holC dnaX S B B H holC holD S B B B H holC holD S B B B H holD S B B B H holM hsdR S S B B H holD S S B B H holD S S B B H holD S S B B H holD S S B B H holD S S B B H holD S S B B H holD S S B B H holD S S B B H holD S S B B H holD S S B B H holD S S B B B H holD S S B B B B B B B B B B B B B B B B B						В	
holC dnaX S B holD holD S B B hsdA nifU S B hsdM hsdR S B hybC hybE S B hypC hypE S S B hypC hypD S S ibpA ibpB S S inIC rpsB S S S inIC rpsR S S S mukB mukE S S S mukB mukE S S S mukB mukE S S S S mukB mukE S S S S S S S S S S S S S S S S S S S							
holC holD S B hscA nifU hscA nifU S hscA nifU hscA S hstV hslU S B hypC hypE S B hypC hypE S S hypC hypD S S infC rpSB S S infC rpSB S S infC rpSB S S infC rpSB S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC rpSC S infC						В	н
hsdM hsdR S B hslV hslU S B hypC hypE S B hypC hypD S Is lipA lipB S Is infC rpsB S Is infC rpsB S Is infC rpsC S Is infC rpsC S Is infC rpsC S Is infC rpsC S In infC rpsC S In infC rpsC S In mukB mukE S B marG narG narG narG narG narG In S narG narG S B narG narG S B narG narG S B nusG rpoA S B <tr< td=""><td></td><td></td><td></td><td></td><td></td><td>В</td><td></td></tr<>						В	
hslV hslU S B hypC hyeE S B hypC hyeE S B hypC hypD S B hypA hypB S B infC rpsB S InfC infC rpsC S InfC infC rpsR S <td< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td></td<>							
hypC hypE S hypC hypD S lbpA ibpB S inIC rpsB S inIC rpsC S inIC rpsR S lyS lysU S mukB mukE S mukB mukF S mukB mukF S narG narJ S narG narJ S narH narG S narJ narH S narG rpo S nusG rpoA S nusG rpoA S nusG rpoA S prib yfiD S prib ys F <td></td> <td></td> <td></td> <td></td> <td></td> <td>В</td> <td></td>						В	
hypC hypD S ibpA ibpB S infC rpsB S infC rpsB S infC rpsB S infC rpsB S lysS lysU S mukB mukB S mukB mukF S mukB mukF S mukB mukF S marG narG S nard narG S nusG rpoC S nusG rpoC S nusG rpoC S nusG rpoA S pBB S P pBB S P						В	
infC rpsB S infC rpsC S infC rpsR S lysS lysU S mukB mukE S mukB mukF S narG narJ S narG narY S narH narG S nusG rpoB S nusG rpoA S parC gyrA S piB yfiB S piB yfiB S piB S P ppA rpS P recJ sbcB S P rpoA rpsB S P rpoA rpsB S P rpoA </td <td></td> <td></td> <td></td> <td></td> <td>S</td> <td></td> <td></td>					S		
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infC rpsR S lysS lysU S mukB mukF S mukB mukF S marG narJ S narG narJ S narH narG S narJ narH S nusG rpoB S nusG rpoB S nusG rpoA S nusG rpoA S parC gyrA S prD milB S pstB yfD S ppp milB S pstB S R ppA ps S recD recB S recD recB S recD rpsB S rpoA rpsB S rpoA rpsB S rpoA rpsB S rpoA rpsD S							
mukB mukE S mukB mukF S narG narJ S narG narY S narH narG S narJ narH S nusG rpoB S nusG rpoA S nusG rpoA S nusG rpoA S parC gyrA S parB yfiD S ppB yfiD S ppA rpB S recU recU sc recD recB S recD rpSG S rpA rpB S <							
mukB mukF S narG narJ S narG narJ S narH narG S narJ narH S narJ narH S nusG rpoB S nusG rpoB S nusG rpoA S nusG rpoA S parC gyrA S pill yfiD S pp pflB S pp prhB S ppA psB S rpA rpsB S rpA rpsB S rpA rpsB S rpA rpsB S rpA rpB S <t< td=""><td></td><td>lysU</td><td></td><td></td><td>S</td><td></td><td></td></t<>		lysU			S		
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narG narY S B narH narG S B nusG rpoB S B nusG rpoB S NanusG nusG rpoA S NanusG secA psiC gyrA S Page Page pill yfiD S Page Pagee Pagee Pagee Pagee Pagee Pagee Pagee Pagee Pagee							
narH narG S B narJ narH S N nusG rpoB S N nusG rpoC S N nusG rpoA S N parC gyrA S P pflB yflD S P pflB yflD S P pplB yflD S P pplB yflD S P ppt pstB mopA S P pecU recB S P recU recB S P recU recB S P rpoA rpsC S P rpoA rpsB S P rpoA rpsB S P rpoA rpsE S P rpoA rpsB S P rpoA rplD S P <td< td=""><td>narG</td><td>narY</td><td></td><td></td><td>S</td><td></td><td></td></td<>	narG	narY			S		
nusG rpoB S nusG rpoC S nusG rpoA S parC gyrA S pflB yflD S pflB yflD S pptB mopA S pstB mopA S recD recB S recJ sbcB S rmhA ssb H rpoA rpsC S rpoA rpsC S rpoA rpsE S rpoA rpsB S rpoA rpsE S rpoA rpsE S rpoA rpsE S rpoA rplD S rpoA rplD S rpoA rplB S rpoA rplB S rpoB rpoC rpsE rpoC rpsE S rpoC rpsE S	narH	narG				В	
nusG rpoC S nusG rpoA S nusG secA S parC gyrA S pflB yfiD S pflB yfiD S pplB S P pplB S P pplB S P pplB S P recU sbcB S recU sbcB S rmA rsbC S rpoA rpsC S rpoA rpsC S rpoA rpsG S rpoA rpsG S rpoA rpsG S rpoA rplD S rpoA rplB S rpoA rplB S rpoA rplB S rpoB S H rpoC rpsE S rpoC rpsE S							
nusG rpoA S nusG secA S parC gyrA S pfIB yfID S ppp rhIB S pstB mopA S recD recB S recJ sbcB S rhA ssb H rpoA rpsC S rpoA rpsB S rpoA rpsB S rpoA rpsG S rpoA rpsE S rpoA rpsE S rpoA rpID S rpoA rpID S rpoA rpID S rpoA rpIB S rpoB S H rpoB FR FR rpoC rpsE S rpoC rpsE S rpoC rpsE S rpoH rpoB S	nusG				S		
parC gyrA S pflB yfiD S prp rhiB S pstB mopA S recD recB S recLJ sbcB S recLJ sbcB S recLJ sbcB S recLJ sbcB S rpoA rpsC S rpoA rpsB S rpoA rpsB S rpoA rpsG S rpoA rpsG S rpoA rpsD S rpoA rpiD S rpoB S H rpoC rpsE S rpoC rpsE S	nusG	rpoA			S		
pfIB yfID S ppp rhIB S pstB mopA S recD recB S recJ sbcB S rmhA ssb H rpoA rpsC S rpoA rpsB S rpoA rpoB S rpoA rpsE S rpoA rpsE S rpoA rpsD S rpoA rplD S rpoA rplD S rpoA rplD S rpoA rplB S rpoA rplB S rpoA rplB S rpoA rplB S rpoB rpoB S rpoC rpsE S rpoC rpsE S rpoC rpsE S rpoD S H rpoH rpoB S							
ppp rhIB S pstB mopA S recU recB S recU sbcB S redJ sbcB S rmA ssb H rpoA rpsC S rpoA rpsB S rpoA rpsD S rpoA rpsB S rpoA rpID S rpoA rpID S rpoA rpIB S rpoB rpoB S rpoC rpsE S rpoH rpoB S rpoH rpoB S rpoH rpoB S rpoH rpoH rpoH					S		
recD	pnp	rhIB			S		
recJ sbcB S mhA ssb S mhA ssb rpoA rpsC S rpoA rpsB S rpoB rpoB S rpoB rpoB S rpoB rpoB S rpoB rpoB S rpoC rpsB S rpoC rpsB S rpoC rpsB S rpoD nusA S rpoH rpoB S H rpoH rpoB S H rpoH rpoB S B H ssb recG H ssb recG S s							
mhA ssb H rpoA rpsC S rpoA rpsB S rpoA rpsB S rpoA rpsC S rpoA rpsE S rpoA rpsD S rpoA rpID S rpoA rpIC S rpoA rpIB S rpoB rpoB S rpoC rpsE S rpoC rpsE S rpoC rpsB S rpoC rpsD S rpoD nusA S rpoH rpoB S H rpoH rpoB S B sb recG H B usg rpoC S							
TpoA	rnhA	ssb					Н
TPOA							
rpoA rpsG S rpoA rpsE S rpoA rpsD S rpoA rplD S rpoA rplD S rpoA rplB S rpoB rpoB S rpoB rpoS H rpoC rpsE S rpoC rpsB S rpoC rpsB S rpoH rpoB S rpoH dnaK S ssb recG H tsf tufB B usg rpoC S							
TPOA							
TpoA TpID S FpoA TpoB TpoB TpoB TpoC TpsE S FpoC TpsG S FpoD TpoD S FpoD TpoB S FpoH TpoB S H TpoH TpoB S TpoH TpoB S S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB S TpoH TpoB	rpoA	rpsE			S		
rpoA rpIC S rpoA rpIB S rpoB rpoD S H rpoB rpoS H H rpoC rpsE S F rpoC rpsG S F rpoC rpsD S F rpoH rpoB S H rpoH rea S B ssb rea H B usg rpoC S S							
rpoA rplB S rpoB rpoD S H rpoB rpoS H H rpoC rpsE S F rpoC rpsG S F rpoD nusA S F rpoH rpoB S H rpoH dnaK S B ssb recG H H tsf tufB B B usg rpoC S S							
rpoB rpoD S H rpoB rpoS H rpoC rpsE S rpoC rpsG S rpoD nusA S rpoH rpoB S H rpoH dnaK S B ssb recG H H tsf tufB B B usg rpoC S S		rpIB			S		
rpoC rpsE S rpoC rpsG S rpoC rpsD S rpoD nusA S rpoH rpoB S H rpoH dnaK S B ssb recG H H tsf tufB B B usg rpoC S	rpoB	rpoD					
rpoC rpsG S rpoC rpsD S rpoD rpsG S rpoH rpoB S H rpoH dnaK S B ssb recG H B tsf tufB B B usg rpoC S					S		Н
rpoC rpsD S rpoD nusA S rpoH rpoB S H rpoH dnaK S B ssb recG H H tsf tufB B B usg rpoC S S					S		
rpoH rpoB S H rpoH dnaK S B ssb recG H H tsf tufB B B usg rpoC S	rpoC	rpsD			S		
rpoH dnaK S B ssb recG H H tsf tufB B B usg rpoC S							ш
ssb recG H tsf tufB B usg rpoC S						В	п
usg rpoC S	ssb	recG			-		
							В

Experimental	
Affinity chromat	Α
Alanine scanning	В
Biochemical	С
Copurification	D
Cosedimentation	Ε
Cross-linking	F
Electron microscopy	G
Filter overlay assay	Н
Gel filtration chormat	1
Genetic	J
Immunoblotting	Κ
Immunoprec	L
Native gel electrophoresis	М
Surface plasmon resonance	Ν
Two hybrid	0
x-ray	Ρ

Computational 1 Chromo 1 Fusion 2 Phylo 3

Supplementary Table 4. Proteins removed for network attack analysis

Bait	Connections
deaD	36
hfq	34
yfgB	33
yciL	32
dnaK	30
rpoC	28
yceC	28
rpoA	27
ybeZ	23
yfiF	23
rnpA	22
rpID	22
rplC	20
rplB	19
rpsG	19
vacB	19
mreB	18
rpsE	18
dnaJ	17
rpsD	16

Supplementary Table 5. Protein-protein interactions of bacterial proteins which are conserved in >= 125 genomes

Intorpotion	Pactorial Hamalagues (495)	Archaal Hamalagues (45)	Fukanyatia Hamalanya (7)	Total (4.40)
Interaction alaS-dnaK [R]	Bacterial Homologues (125) 125	Archeal Homologues (16)	Eukaryotic Homologues (7)	Total (148) 139
cbpA-dnaK	125	6	6	137
cbpA-rpIB	124	6	6	136
cbpA-rpID	124	0	5	129
cbpA-rpIV	124	0	1	125
cbpA-rpoC	125	6	6	137
cbpA-rpsB	125	0	5	130
cbpA-rpsC	125	6	4	135
cbpA-rpsC cbpA-rpsG	125	3	6	134
cbpA-rpsG cbpA-srmB	115	6	6	127
clpA-rpoB [R]	121	3	6	130
clpA-rpoB [K]	121	3	6	130
clpP-dnaK	119	0	5	124
clpP-mopA	119	0	5	124
clpP-mopA	119	0	5	124
deaD-rpIA	112	11	3	124
deaD-rplB	111	15	7	133
	112		6	125
deaD-rpIC	112	7 9	5	125
deaD-rpIM	112	9 15	5	132
deaD-rpsC				
deaD-rpsE	112	11	7	130
deaD-rpsG	112	11	6 2	129
deaD-rpsH	112 112	11		125
deaD-rpsI		14	6	132
deaD-rpsJ	112	15	4	131
deaD-rpsM	112	14	5	131
dnaE-dnaX	125	0	0	125
dnaJ-deaD	112	5	7	124
dnaJ-dnaK	125	6	7	138
dnaJ-mreB [R]		6	7	137
dnaJ-pstB	125 121	6	7	138
dnaJ-recA		1	2 7	124
dnaJ-srmB [R]	115	6		128
dnaK-grpE [R]	115	6	6	127
dnaK-hscA	125	7	7	139
eno-dnaK	123	7	7	137
eno-pnp [R]	117	15	5	137
ftsE-mopA	123	16	7	146
ftsZ-fusA	115	12	1	128
gyrA-gyrB [R]	125	6	7	138
gyrA-mreB [R]	124	6	7	137
gyrA-pstB hslU-dnaJ	125 119	6 0	7 6	138 125
hslU-dnaK	119	0	6	125
hslU-mreB	118	1	6	125
	119		6	125
hsIU-pstB	124	3 0		125
infC-rpIB	125	0	1 1	125
infC-rpsB	125	0		126
infC-rpsC	125		1	126
infC-rpsD		0	1	
infC-rpsE	125	0	1	126
infC-rpsG	125	0	1	126
infC-rpsH	125	0	1	126
infC-rpsI	125	0	1	126
infC-rpsJ	125	0	1	126
infC-rpsM	125	0	1	126
infC-rpsR	124	0	1	125
lysS-lysU [R]	123	16	7	146
mopA-clpB	120	2	6	128
mreB-dnaK	124	7	7	138

mreB-pnp [R]	118	7	5	130
	124	8	7	139
mreB-pstB [R]				
mreB-rpoB	124	8	7	139
mreB-rpoC	124	8	7	139
-	124	0	1	125
mreB-secA [R]				
mreB-tufB [R]	124	8	7	139
parC-gyrA	125	6	7	138
pheS-pheT [R]	123	12	5	140
pnp-rhlB [R]	106	15	5	126
polA-rpoA	125	0	2	127
polA-rpoC	125	0	5	130
prsA-mopA	116	16	6	138
pstB-mopA	123	16	7	146
rhlB-eno	110	16	7	133
	120			
rpID-clpA		0	4	124
rpID-dnaK	124	0	5	129
rpID-rpoC	124	0	5	129
				125
rpID-secA	124	0	1	
rpoA-rpIB	124	0	2	126
rpoA-rpIC	125	0	2	127
	124	0	2	126
rpoA-rpID				
rpoA-rpIL	125	0	1	126
rpoA-rpIM	125	0	2	127
	125		2	127
rpoA-rpoB [R]		0		
rpoA-rpoC [R]	125	0	2	127
rpoA-rpoD [R]	125	0	1	126
	125	0	1	126
rpoA-rpsB				
rpoA-rpsC	125	0	1	126
rpoA-rpsD	125	0	1	126
rpoA-rpsE	125	0	2	127
rpoA-rpsG	125	0	2	127
rpoB-rpoC [R]	125	16	7	148
rpoB-rpoD [R]	125	0	1	126
rpoB-rpoH [R]	125	0	1	126
rpoC-rpIB	124	16	7	147
rpoC-rplC	125	7	6	138
rpoC-rpID	124	0	5	129
rpoC-rpoD [R]	125	0	1	126
rpoC-rpsD	125	0	1	126
rpoC-rpsE	125	12	7	144
rpoC-rpsG	125	11	6	142
rpoH-dnaK	125	0	1	126
	·=-	-	· ·	
rpoS-rpoB [R]	125	0	1	126
rpoS-rpoC	125	0	1	126
secA-tufB [R]	125	0	1	126
selB-dnaJ	125	6	7	138
selB-srmB	115	16	7	138
tgt-rplB	109	16	5	130
tgt-rpsG	110	11	4	125
topA-rpoC	122	16	7	145
topB-rpIM	119	9	5	133
tpiA-dnaK	123	2	7	132
tsf-tufB [R]	125	0	5	130
tufA-tsf	125	0	5	130
tufA-tufB [R]	125	16	7	148
uvrC-rpsC	120	4	0	124
uvrC-rpsE	120	4	0	124
uvrC-rpsH	120	4	0	124
vacB-rplB	113	8	7	128
vacB-rpsE	114	6	7	127
yceC-rpIA	125	0	3	128
yceC-rpIB	124	Ö	7	131
yceC-rpIC	125	0	6	131

yceC-rpID	124	0	5	129
yceC-rpIE	125	0	6	131
yceC-rpIF	125	0	2	127
yceC-rplM	125	0	5	130
yceC-rpIV	124	0	1	125
yceC-rpsB	125	0	5	130
yceC-rpsC	125	0	5	130
yceC-rpsD	125	0	1	126
yceC-rpsE	125	0	7	132

[R] denotes that the reciprocal interaction was also observed

() indicates the number of genomes used in the analysis

Supplementary Table 6. Gene neighborhood predicted interactions reported in this study

Bait	Prey	Species	REGULONDB Operon (E. coli)
accC	accB	Escherichia coli K12	accBC
accD	accA	Bacillus subtilis	accbo
accD	aas	Mycobacterium tuberculosis H37Rv	
acpP	fabB	Mesorhizobium loti	
acpP	fabF	Escherichia coli K12	fabHDG-acpP-fabF
acpP		Escherichia coli K12	fabHDG-acpP-fabF
acpl	fabZ	Sinorhizobium meliloti	таыты б-асрт -таы
		Escherichia coli K12	b2341 b2342
	dnaN		02041 02042
cbpA	dnaK	Synechocystis PCC6803	
clpP	rfaD	Agrobacterium tumefaciens C58	
cspC		Pseudomonas aeruginosa	
-	rpsO	Mycoplasma pneumoniae	
dnaJ	•	Escherichia coli K12	dnaKJ
dnaK		Mycobacterium leprae	anarto
dnaK	sapA	•	
gidA	thdF	Borrelia burgdorferi	
hsdM		Escherichia coli K12	hsdMS
hslV	hslU	Escherichia coli K12	hsIVU
		Escherichia coli K12	hypABCDE
kdsA	rfaD	Caulobacter crescentus	
mukE	mukB	Escherichia coli K12	smtA-mukFEB
narH	narG	Escherichia coli K12	narGHJI
narJ	narH	Escherichia coli K12	narGHJI
narY	narG	Bacillus subtilis	
nusA	infB	Escherichia coli K12	nusA-infB
pheS	pheT	Escherichia coli K12	pheST-himA
recD	recB	Escherichia coli K12	recD recB ptr
rnpA	rplT	Rickettsia conorii	
rpoA	rpsD	Escherichia coli K12	rpsMKD-rpoA-rpIQ
rpoB	rpoC	Escherichia coli K12	rpoBC
rpoC	rpsG	Borrelia burgdorferi	
sucB	sucA	Escherichia coli K12	sucABCD
topB	ssb	Xylella fastidiosa	
ybdQ	rfaD	Nostoc sp	
yceC	yfiF	Sinorhizobium meliloti	
yciL	rpsJ	Campylobacter jejuni	
yfiF	rpsF	Pseudomonas aeruginosa	
ygdP	srmB	Listeria innocua	
ygdP	yfiF	Fusobacterium nucleatum	
ygjD	yeaZ	Mycoplasma pulmonis	
ynhD	ynhE	Escherichia coli K12	ynhA b1680 ynhC ynhD ynhE

Supplementary Table 7. List of the 49 COGs bacterial genomes used in this study

Bacteria (10):

Aquifex aeolicus

Thermotoga maritima

Chlamydia trachomatis

Chlamydophila pneumoniae CWL029

Treponema pallidum

Borrelia burgdorferi

Synechocystis

Nostoc sp. PCC 7120

Fusobacterium nucleatum

Deinococcus radiodurans

Actinobacteria (4):

Corynebacterium glutamicum

Mycobacterium tuberculosis H37Rv

Mycobacterium tuberculosis CDC1551

Mycobacterium leprae

Gramplus (12):

Clostridium acetobutylicum

Lactococcus lactis

Streptococcus pyogenes M1 GAS

Streptococcus pneumoniae TIGR4

Staphylococcus aureus N315

Listeria innocua

Bacillus subtilis

Bacillus halodurans

Ureaplasma urealyticum

Mycoplasma pulmonis

Mycoplasma pneumoniae

Mycoplasma genitalium

Gamma (10):

Escherichia coli K12

Escherichia coli O157:H7 EDL933

Escherichia coli O157:H7

Yersinia pestis

Salmonella typhimurium LT2

Buchnera sp. APS

Pseudomonas aeruginosa

Haemophilus influenzae

Pasteurella multocida

Xylella fastidiosa 9a5c

Proteobacteria (6):

Neisseria meningitidis MC58

Neisseria meningitidis Z2491

Ralstonia solanacearum

Helicobacter pylori 26695

Helicobacter pylori J99

Campylobacter jejuni

Alpha (7):

Agrobacterium tumefaciens strain C58 (Cereon)

Sinorhizobium meliloti

Brucella melitensis

Mesorhizobium loti

Caulobacter crescentus CB15

Rickettsia prowazekii

Rickettsia conorii