	D D D D D icd cydC CydC CydD D D D D D CydD D D D D D fepB D D D D D D fepC D D D D D D fepD D D D D D fepG D D D D D D ftsE
atsikanlaranpoinedbiakajūstinas D D D D D D D D D D D D D	D D D D D D IolD D D D D D IptB D D D D D D pstB D D D D D D pstC D D D D D D rffH glmS glnA
	D D D D purB aspS cysS fmt lysS trpS tyrS D D D D ydiF
	D D D D D galU glmS.1 pgm ugd D D D D D D ugd.1 D D D D D D ugd.1
actanBaeseakscrateriem a d a d a d a d a d a d a d a d a d a	secB secF secG secM tolC D D D D D D acrA acrB D D D D D D mrdA tolC.1
	aroD cysE D D D D D D dapA dapB dapD dapE dapF eno fbaA
Biosynthouse Biosy	D D D D D gapA glnA.2 glyA icd.1 rpe rpiA tktA D D D D D D D tktA accB aceE aceF
	adk aroD.1 cysM.1 dapA.1 dapB.1 dapF.1 eno.1 entD fbaA.1 D D D D D D gapA.1
Biosynthes Biosynthes D D D D D D D D D D D D D D D D D D	glmS.2 glyA.1 icd.2 D D D D D ispA ispD ispE ispF ispH D D D D D D lpd pgm.1 D D D D D D rffH.1
D D D D D D D D D D D D D D D D D D D	D D D D Trpe.1 rpiA.1 sucA sucB tktA.1 accB.1 aceE.1 aceF.1 adk.1 aroD.2
	D D D D D D dapA.2 dapB.2 dapF.2 eno.2 entD.1 fbaA.2 glyA.2 glyA.2 gpsA
0 D D D D D D D D D D D D D D D D D D D	hemB hemD hemE hemG hemH D D D D D D icd.3 D D D D D D ispA.1 ispD.1 ispE.1 ispF.1
Biosy D D D D D D D D D D D D D D D D D D D	Split ispH.1 ispH.1 lpd.1 pgm.2 purB.2 ribE ribF ribF ppi.2 purB.2 ribE ribF ppi.2 purB.2 purB.3
	ubiA ubiE D D D D D ubiF D D D D D ubiG D D D D D D ubiH ubiX ynbB entD.2
	fabZ D D D D ydiF.1 accB.2 aceE.2 aceF.2 cysE.2 cysM.3 eno.3 fbaA.3
CAMP resistancemetable CAMP C	fold gap A.3 gly A.3 gly A.3 licd.4 lpd.2 lpd.2 rpe.3 rpi A.3 suc A.2 suc B.2 lbd. A.3 lpd. A
Catigorial (a) Experimental (a) Experime	D D D D D acrA.1 acrB.1 lpxA tolC.2 D D D D D D acrA.1 acrB.1 lpxA tolC.2 D D D D D D location icd.5 lpd.3 sucA.3 sucB.3
	glyA.4 CysE.3 CysM.4 murD murl dnaG dnaQ
roadishbiosyn Dhasiisteep Organia	dnaX holB holC holD ssb accB.3 fabA fabH fabZ.1
	accB.4 fabA.1 fabH.1 fabZ.2 folA folB D D D D D folC folE folK folP
	D D D D D alsK fbaA.4 D D D D D galU.2 pgm.3 D D D D D icd.6 gpsA.1
	D D D D D glyA.5 D D D D D D lpd.4 aceE.4 aceF.4 chbF crr.1 eno.4 fbaA.5
entaimatutiocarbolyydates@leyta O O O O O O O O O O O O O O O O O O O	D D D D D D gapA.4 D D D D D D D pgm.4 GlnA.3 glyA.6 lpd.6 dnaQ.1 dnaT dnaX.1
	holB.1 holC.1 holD.1 priA priB ssb.1 D D D D D suhB
	D D D D D gmhA hldD hldE kdsA kdsB lpxA.1 D D D D D D IpxL D D D D IpxM waaC
etyædætelærosynthæræpolys	waaF waaG waaU D D D D D dapA.3 dapB.3 dapD.1 dapE.1 dapF.3
	D D sucB.4 accB.5 aceE.5 aceF.5 adk.2 aroD.3 birA.1 cmk coaE cydA
	D D D D D cydB cysE.4 D D D D D D dapA.4 dapB.4 dapD.2 dapE.2 dapF.4 dcd dnaQ.2 dnaX.2
	dut eno.5 fabA.2 D D fabH.2 fabZ.3 fbaA.6 folA.1 D D D D D D folB.1 D D D D D D folC.1 folD.1 NPEQ folE.1 8 folK 1
	folK.1 folP.1 galU.3 gapA.5 glmS.3 glnA.4 glyA.7 gmhA.1 guaA hemB.1 hemD.1 hemE.1
tabolic pathways The state of	hemG.1 hemH.1 hemL.1 hldD.1 hldE.1 holB.2 holC.2 holC.2 holD.2 icd.7 iscS D D D D D D ispA.2
	ispD.2 ispE.2 ispF.2 ispH.2 kdsA.1 kdsB.1 lipA.1 lipB.1 lipB.1 lpd.7 lpxA.2 D D D D D D lpxL.1
	D D D D D IpxM.1 murD.1 murl.1 nadD nadE D D D D D D nrdA D D D D D D nrdB pgm.5 pgsA.1 D D D D D D purB.3 rffH.2
	ribE.1 ribF.1 D D D D D rpe.4 rpiA.4 sucA.5 sucB.5 D D D D D SuhB.1 thiL thyA tktA.4 tmk
D D D D D D D D D D D D D D D D D D D	UbiA.1
Methane metabolis	eno.6 fbaA.7 glyA.8 accB.6 aceE.6 aceF.6 alsK.1 cysE.5
in diverse environme D D D D D D D D D D D D D	D D D D D CysM.6 dapA.5 dapB.5 dapD.3 dapE.3 dapF.5 eno.7 fbaA.8 foID.2 gapA.6 glnA.5
D D D D D D D D D D D	D D D D D icd.8 D D D D D Ipd.8 paaK pgm.6 D D D D D Trpe.5 rpiA.5 sucA.6 sucB.6 D D D D D tktA.5 ydiF.3
	D D D dam dnaQ.3 dnaX.3 holB.3 holC.3 holD.3 ssb.2
Comparing the control of the co	nadD.1 nadE.1 D D D can glnA.6 fmt.1 folA.2 folD.3 glyA.10
	thyA.1 D D D D D CydA.1 cydB.1 ppa acpS coaE.1 D D D D D galU.4 rpe.6
	D D D D ugd.3 fbaA.9 pgm.7 pgm.7 rpe.7 rpiA.6 tktA.6 D D D D D mrdA.1 murD.2 murJ
Protein example and the substitution of the su	paaK.1 aroD.4 crr.2 ptsl D D D rffH.3 hemB.2
	hemD.2 hemE.2 hemG.2 hemH.2 hemL.2 D D D D D D accB.7 D D D D D D ydiF.4
Protein expans	lepB lspA secA.1 secB.1 secF.1 secG.1 secM.1
ES CONTRACTOR OF THE PROPERTY	guaA.1 holB.4 holC.4 holD.4 D D D D D D nrdA.1 D D D D D D pgm.8 pnp D D D D D D purB.4
D D D D D D D D D D D D D D D D D D D	D D D D spoT cmk.1
D D D D D D D D D D D D D D D D D D D	cmk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holC.5 holD.5 D D D D D nrdA.2 pnp.1 psuK
metabolism Pyrimidine metabolism O O O O O O O O O O O O O O O O O O O	cmk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holC.5 holD.5 nrdA.2 nrdB.2 pnp.1 psuK thyA.2 tmk.1 accB.8 aceE.7 aceF.7 gloB lpd.10 lepB.1 rseP
e metabolism Pyrimidine metabolism O O O O O O O O O O O O O O O O O O O	cmk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holC.5 holD.5 nrdA.2 nrdB.2 pnp.1 psuK thyA.2 tmk.1 accB.8 aceE.7 aceF.7 gloB lpd.10 lepB.1 rseP secA.2 secB.2 secG.2 ribF.2 rplA rplC rplD
metabolism Pyrimidine metabolism O O O O O O O O O O O O O O O O O O O	cmk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holC.5 holD.5 nrdA.2 nrdB.2 pnp.1 psuK thyA.2 tmk.1 accB.8 aceE.7 aceF.7 gloB lpd.10 lepB.1 rseP secA.2 secB.2 secG.2 ribE.2 ribF.2 rplA rplC rplD rplK rplM rplN rplO rplP rplQ rplR rplS rplT rplY rpmA
metabolism Pyrimidine metabolism O O O O O O O O O O O O O O O O O O O	cmk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holC.5 holD.5 nrdA.2 nrdB.2 pnp.1 psuK thyA.2 tmk.1 accB.8 aceE.7 aceF.7 gloB lpd.10 lepB.1 rseP secA.2 secB.2 secG.2 ribE.2 ribF.2 rplA rplC rplD rplK rplM rplN rplO rplP rplQ rplR rplS rplT rplY rpmA rpmB rpmC rpmD rpmG rpmH rpml rpmJ rpsF rpsl rpsJ rpsK
Ribosome Riboslavin m@abouismseRsinggate metabolism Pyrimidine metabolism of the state of the st	cmk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holC.5 holD.5 nrdA.2 pnp.1 psuK thyA.2 tmk.1 accB.8 aceE.7 aceF.7 gloB lpd.10 lepB.1 rseP secA.2 secB.2 secG.2 ribE.2 ribF.2 rplA rplC rplD rplK rplM rplN rplO rplP rplQ rplR rplS rplT rplY rpmA rpmB rpmC rpmD rpmG rpmH rpmJ rpsF rpsI rpsJ
Ribosome Riboslavin m@abouismseRsinggate metabolism Pyrimidine metabolism of the state of the st	mk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holD.5 holD.5 holD.5 ppp.1 psuK thyA.2 tmk.1 accB.8 aceE.7 aceF.7 gloB pd.10 lepB.1 rseP secA.2 secB.2 secG.2 ribE.2 ribF.2 rplA rplC rplD rplK rplN rplN rplN rplN rplN rplN rplN rplN
Ribosome Riboslavin m@abouismseRsinggate metabolism Pyrimidine metabolism of the state of the st	cmk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holC.5 holD.5 holC.5 holD.5 mrdA.2 pnp.1 psuK ttyA.2 tmk.1 accB.8 aceE.7 aceF.7 gloB pld.10 lepB.1 rseP secA.2 secB.2 secG.2 ribE.2 ribF.2 ribF.2 rplA rplC rplD rplK rplM rplN rplN rplO rplP rplR rplS rps
Ribosome Riboslavin m@abouismseRsinggate metabolism Pyrimidine metabolism of the state of the st	mk.1 dcd.1 dnaX.5 dut.1 holB.5 holC.5 holD.5 mrdB.2 prp.1 psuK thyA.2 tmk.1 accB.8 aceE.7 aceF.7 gloB pd.10 lepB.1 rseP secA.2 secB.2 secG.2 ribE.2 ribE.2 ribE.2 ribF.2 rplA rplC rplC rplD rplK rplB rpmC rpmB rpmC
Ribosome Riboslavin m@abouismseRsinggate metabolism Pyrimidine metabolism of the state of the st	
Column 1 Column 2 Column 3	mk.1 dcd.1 dnaQ.5 dnaX.5 dut.1 holB.5 holC.5 holC.5 holC.5 holC.5 rrdA.2 rrdB.2 ppp.1 psuK thyA.2 tmk.1 accB.8 accE.7 accF.7 gloB pl.10 lepB.1 rseP secA.2 secB.2 secG.2 ribE.2 ribE.2 ribF.2 ribF.2 rplA rplC rplD rplK rplM rplN rplO rplP rplQ rplR rplS rplT rplY rpmA rpmB rpmC rpmD rpmG rpmH rpmJ rpsF rpsI rpsI rpsI rpsI rpsI rpsI rpsI rpsI