Gene	Description	24h	48h
sll1945	1-deoxy-xylulose 5-phosphate synthase	1.59	
slr2130	3-dehydroquinate synthase	1.78	
ssl3436	50s ribosomal protein 129	1.50	
sll0480	aspartate aminotransferase related protein	1.57	
sll0208	conserved hypothetical protein	1.57	
slr0609	conserved hypothetical protein	1.61	
ssr3451	cytochrome b559 alpha chain	1.64	
smr0006	cytochrome b559 beta chain	1.79	
sll0018	fructose-bisphosphate aldolase (ec 4.1.2.13)	1.82	
slr0151	hypothetical protein	1.77	
sll1618	hypothetical protein	1.52	
slr1472	-identity	1.84	
sll1363	ketol-acid reductoisomerase (ec 1.1.1.86)	1.75	
sll1743	L11 protein	2.36	
sll0947	light repressed protein a	1.52	
sll0258	low potential cytochrome C	1.97	
sll0629	photosystem i reaction centre subunit x (psi-k)	1.77	
sll1471	phycobilisome rod-core linker polypeptide CpcG	2.67	
slr0775	protein-export membrane protein SecF, putative, putative	2.55	
slr1164	ribonucleotide reductase alpha chain	1.53	
slr0653	RNA polymerase sigma factor rpod1	1.59	
sll1931	serine hydroxymethyltransferase	1.57	
sll1688	threonine synthase	1.81	
sll0860	conserved hypothetical protein	1.53	2.14
slr0186	2-isopropylmalate synthase	1.88	1.92
sll0336	acetyl-coenzyme a carboxylase carboxyl transferase subunit beta	1.58	2.17
ssl2084	acyl carrier protein	2.78	3.61
slr0729	conserved hypothetical protein	2.18	3.51
slr2136	conserved hypothetical protein	1.72	1.70
slr1847	conserved hypothetical protein	2.46	3.05
ssl3364	conserved hypothetical protein	1.53	3.24
sll1214	conserved hypothetical protein (orf349)	2.67	2.17
slr0923	conserved hypothetical protein (orf99)	2.78	2.16
sll1185	coproporphyrinogen III oxidase, aerobic	2.00	2.02
sll1663	CpcE-related protein	1.66	2.16
sll0504	diaminopimelate decarboxylase	2.34	1.64
sll1091	geranylgeranyl hydrogenase	2.83	1.80
slr0033	glutamyl-tRNA (Gln) amidotransferase subunit C	1.88	2.48
slr0426	GTP cyclohydrolase i	1.79	2.31
sll0872	hypothetical protein	2.56	2.85
slr1853	PcaC-like protein, putative	1.67	1.89
sll1110	peptide chain release factor RF-1	1.76	1.66
sll1994	porphobilinogen synthase	2.18	2.87
sll1019	probable hydroxyacylglutathione hydrolase	4.39	3.64
sll0807	ribulose-phosphate 3-epimerase (ec 5.1.3.1)	1.67	1.72
		2107	

Table S2 B Up-regulated proteins (fold changes) revealed by quantitative I TRAQ proteomic a

slr1417	T10O24.11	1.85	3.62
sll1980	thiol:disulfide interchange protein txla	1.62	2.40
sll0381	hypothetical protein		2.01
slr1761	12 kDa FK506-binding protein (fkBP) (peptidyl-prolyl)		2.04
ssl3436	50s ribosomal protein 129		1.57
slr0242	bacterioferritin comigratory protein		3.34
sll1654	conserved hypothetical protein		1.74
slr0823	conserved hypothetical protein		2.56
sll1336	conserved hypothetical protein		1.69
sll1734	conserved hypothetical protein		2.19
ssl0352	conserved hypothetical protein		1.64
slr0581	conserved hypothetical protein		2.08
ssl1707	conserved hypothetical protein		2.92
ssr3122	conserved hypothetical protein		2.61
sll0634	conserved hypothetical protein		1.69
ssr2998	conserved hypothetical protein		1.50
sll0662	conserved hypothetical protein		1.85
ssl2982	conserved hypothetical protein (orf75)		1.92
slr1828	ferredoxin ii		2.43
slr0739	geranylgeranyl diphosphate synthase (SelGGPS)		1.66
sll0057	grpe protein, fragment-related		1.62
slr1600	hypothetical protein		1.92
ssr1853	hypothetical protein		4.10
slr1339	hypothetical protein		1.56
sll0630	hypothetical protein		3.45
slr0455	hypothetical protein		2.17
slr1431	hypothetical protein		2.06
slr0476	hypothetical protein		1.69
sll1620	hypothetical protein		1.75
sll1549	hypothetical protein		2.06
ssl0461	hypothetical protein		1.73
ssl0242	hypothetical protein		10.31
slr1160	hypothetical protein		2.39
slr0111	hypothetical protein		1.90
ssr3402	hypothetical protein		2.11
slr1915	hypothetical protein		2.63
sll0162	hypothetical protein		1.57
slr1970	hypothetical protein		3.77
slr1472	-identity		1.92
sll1074	leucyl-tRNA synthetase		2.73
slr1198	LsfA		2.02
slr1030	Mg-Protoporhyrin IX		1.82
sll1209	M-protein, putative		1.61
sll1732	NADH dehydrogenase subunit 5		2.18
slr0043	nrtC-like transporter protein (cmpC)		1.91
sll0199	ntss01_ORF03920		4.44
slr1887	PBG-D (aa1-344)		1.86
slr0394	phosphoglycerate kinase		2.39

ssr2831	photosystem i reaction centre subunit iv	2.39
ssr3383	phycobilisome 7.8 kd linker polypeptide, allophycocyanin-associated, core	1.57
slr2051	phycobilisome rod-core linker polypeptide cpcg	2.13
slr1783	probable transcriptional regulator ycf29	1.82
ssl0563	product binds the terminal electron accepting Fe-S centers of photosystem I	2.07
slr1846	promoter active fragment E3	1.51
slr0114	regulatory protein, putative	1.77
slr0194	ribose 5-phosphate isomerase, putative	1.64
sll0145	ribosome recycling factor (ribosome releasing factor) (rrf)	1.68
sll1594	rubisco operon transcriptional regulator	1.56
sl10039	similar to sir1042 protein of Synechocystis sp. strain PCC 6803	1.79
slr1271	teichoic acid biosynthesis protein a-related	1.77
slr1139	thioredoxin	2.61
slr1992	vitami B12 transporter protein/glutathione peroxidase	2.16

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nalysis

slr0769 co slr0957 hy sll1080 hy sll1757 co sll0783 hy sll0172 hy slr1735 glu slr1071 Bs slr0058 hy slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	TP phosphoribosyltransferase onserved hypothetical protein ypothetical protein ypothetical 36.3 kd lipoprotein precursor, putative onserved hypothetical protein ypothetical protein ypothetical protein ypothetical protein lutamine transport ATP-binding protein glnq saA ypothetical protein pha-l-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein tilizing regulatory protein tutC	0.003 0.210 0.295 0.583 0.352 0.354 0.369 0.373 0.566 0.567 0.433 0.459 0.582 0.481 0.490 0.561 0.517	
slr0957 hy sll1080 hy sll1757 co sll0783 hy ssl1918 hy sll0172 hy slr1735 glu slr1735 glu slr1735 hy slr0518 alp slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	ypothetical protein ypothetical 36.3 kd lipoprotein precursor, putative onserved hypothetical protein ypothetical protein ypothetical protein lutamine transport ATP-binding protein glnq saA ypothetical protein pha-1-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.219 0.295 0.583 0.352 0.354 0.369 0.373 0.566 0.567 0.433 0.433 0.459 0.582 0.481 0.490 0.561	
sll1080 hy sll1757 co sll0783 hy ssl1918 hy sll0172 hy sll0172 hy slr1735 glu slr1717 Bs slr0058 hy slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	ypothetical 36.3 kd lipoprotein precursor, putative onserved hypothetical protein ypothetical protein ypothetical protein utamine transport ATP-binding protein glnq saA ypothetical protein pha-1-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.295 0.583 0.352 0.354 0.369 0.373 0.566 0.567 0.433 0.459 0.582 0.481 0.490 0.561	
sll1757 co sll0783 hy ssl1918 hy sll0172 hy slr1735 glu slr1735 glu slr171 Bs slr0058 hy slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	onserved hypothetical protein ypothetical protein ypothetical protein ypothetical protein lutamine transport ATP-binding protein glnq saA ypothetical protein pha-1-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.583 0.352 0.354 0.369 0.373 0.566 0.567 0.433 0.459 0.582 0.481 0.490 0.561	
sll0783 hy ssl1918 hy sll0172 hy slr1735 glu slr1735 glu slr171 Bs slr0058 hy slr0518 alp slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	ypothetical protein ypothetical protein ypothetical protein lutamine transport ATP-binding protein glnq saA ypothetical protein pha-1-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.352 0.354 0.369 0.373 0.566 0.567 0.433 0.459 0.582 0.481 0.490 0.561	
ssl1918 hy sll0172 hy slr1735 glu slr1735 glu slr058 hy slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	ypothetical protein ypothetical protein lutamine transport ATP-binding protein glnq saA ypothetical protein pha-1-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.354 0.369 0.373 0.566 0.567 0.433 0.459 0.582 0.481 0.490 0.561	
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slr1735 glu slr171 Bs slr0058 hy slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	lutamine transport ATP-binding protein glnq saA ypothetical protein pha-l-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.373 0.566 0.567 0.433 0.459 0.582 0.481 0.490 0.561	
slr1171 Bs slr0058 hy slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	saA ypothetical protein pha-l-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.566 0.567 0.433 0.459 0.582 0.481 0.490 0.561	
slr1171 Bs slr0058 hy slr0518 alp slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	saA ypothetical protein pha-l-arabinofuranosidase b precursor b)-related onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.567 0.433 0.459 0.582 0.481 0.490 0.561	
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slr0374 co sll1426 hy slr2101 co sll1483 tra slr1773 hy	onserved hypothetical protein ypothetical protein onserved hypothetical protein ansforming growth factor induced protein ypothetical protein	0.459 0.582 0.481 0.490 0.561	
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sll1483 tra slr1773 hy	ansforming growth factor induced protein ypothetical protein	0.490 0.561	
sll1483 tra slr1773 hy	ansforming growth factor induced protein ypothetical protein	0.561	
s110779 11ti	tilizing regulatory protein tutC	0 517	
1 July au		0.517	
sll1054 co	onserved hypothetical protein	0.527	
	DP-3-0-acyl N-acetylglucosamine deacetylase	0.537	
slr1678 50	OS ribosomal protein L21	0.528	
sll0051 co	onserved hypothetical protein	0.533	
sll1933 Dr	naJ1 protein	0.534	
slr1693 pil	llg protein, putative	0.603	
sll0103 co	onserved hypothetical protein	0.613	
sll1089 hy	ypothetical protein	0.617	
slr1958 hy	ypothetical protein	0.623	
slr1410 pu	utative WD-repeat protein, putative	0.623	
slr0201 su	accinate dehydrogenase subunit C, putative	0.628	
ssr1600 pu	utative anti-sigma factor antagonist, putative	0.631	
slr2060 hy	ypothetical protein	0.641	
sll1981 ac	cetolactate synthase	0.643	
sll0753 me	ethylenetetrahydrofolate dehydrogenase	0.646	
	ypothetical protein	0.647	
sll0923 pu	utative transmembrane protein Wzc, putative	0.651	
sll0180 hy	ypothetical 41.2 kd lipoprotein in hded-gada intergenic region pre	0.653	
	onserved hypothetical protein	0.654	
slr1924 Al	lkaline D-peptidase	0.657	
sll0272 hy	ypothetical protein	0.663	
ssl2501 hy	ypothetical protein	0.136	0.081
	esponse regulator SrrB, putative	0.295	0.253
sll1498 ()		0.376	0.584
slr1830 po	oly-beta-hydroxybutyrate polymerase	0.398	0.348
sll1223 ho		0.399	0.362

Table S2 A Down-regulated proteins (fold changes) revealed by quantitative I TRAQ p

	conserved hypothetical protein	0.404	0.323
	ChpA, putative	0.427	0.496
	cyanelle ribosomal protein S4	0.476	0.497
	hypothetical protein	0.512	0.623
sll1821	chloroplast ribosomal protein L13	0.527	0.617
	ribosomal protein L19	0.536	0.446
slr1276	hypothetical protein	0.539	0.509
sll0374	urea/short-chain amide ABC transporter, ATP-binding protein, puta	0.542	0.322
slr1694	appA protein	0.548	0.476
slr1289	isocitrate dehydrogenase [nadp] (ec 1.1.1.42)	0.554	0.380
sll1811	50s ribosomal protein 118	0.558	0.468
	conserved hypothetical protein	0.563	0.492
slr0369	mexF	0.568	0.466
sll1380	conserved hypothetical protein (orf238)	0.571	0.577
sl10420	urease beta subunit	0.582	0.653
sll0368	pyrimidine operon regulatory protein PyrR	0.585	0.589
sll1388	conserved hypothetical protein	0.597	0.659
ssl3437	30s ribosomal protein s17	0.608	0.444
sll1296	ChpA, putative	0.616	0.489
slr1363	conserved hypothetical protein	0.625	0.518
sll1358	conserved hypothetical protein	0.625	0.438
slr1793	transaldolase	0.625	0.621
sll1423	NtcA	0.632	0.607
slr1406	hypothetical protein	0.634	0.624
slr1275	hypothetical protein	0.641	0.502
	NADH dehydrogenase	0.647	0.449
slr1753	hypothetical protein	0.652	0.513
sll1118	conserved hypothetical protein	0.653	0.601
	30S ribosomal protein S7	0.654	0.575
sll0781	conserved hypothetical protein	0.656	0.371
slr1881	0		0.528
sll0528	(), putative		0.524
slr0250	17.3 kd protein in mura-rpon intergenic region precursor, putative		0.575
	30s ribosomal protein s11		0.520
slr0628	30S ribosomal protein S14		0.518
sll1804	30S ribosomal protein S3		0.513
	30s ribosomal protein s8		0.335
slr1994	3-oxoacyl-(acyl-carrier protein) reductase		0.235
	4-alpha-glucanotransferase (ec 2.4.1.25) (amylomaltase)		0.594
	50s ribosomal protein 117		0.377
	50s ribosomal protein 12		0.647
sll1803	50s ribosomal protein 122		0.607
-	50s ribosomal protein 123		0.463
	50S ribosomal protein L27		0.552
	50s ribosomal protein 14		0.563
-	50s ribosomal protein 16		0.521
	ABC transporter, ATP-binding protein		0.275
	ABC transporter, putative		0.525
	- · · · · · · · · · · · · · · · · · · ·		

slr1993 acetyl-coa acetyltransferase	0.325
slr1898 acetylglutamate kinase (n-acetylglutama)	0.306
slr1022 acetylornithine aminotransferase	0.575
sll1085 aerobic glycerol-3-phosphate dehydrogenase, putative	0.629
sll1270 amino acid ABC transporter, permease protein	0.277
sll0224 amino-acid abc transporter binding protein	0.610
sll0108 ammonium transporter	0.234
slr0585 argininosuccinate synthase	0.433
sll1327 ATP synthase gamma chain	0.416
slr1768 B-cell receptor associated protein-related protein, putative	0.593
sll1314 c4-dicarboxylate-binding periplasmic protein precursor, putative	0.397
slr0257 carboxyl terminal protease	0.564
slr1751 carboxyl terminal protease	0.552
slr0798 cation-transporting ATPase, P-type (pacS)	0.584
sll1822 chloroplast 30s ribosomal protein s9	0.344
slr0929 chromosome partitioning protein, ParA family	0.585
sll0401 citrate synthase	0.420
sll0185 conserved hypothetical protein	0.361
sll0335 conserved hypothetical protein	0.254
sll0596 conserved hypothetical protein	0.413
sll0749 conserved hypothetical protein	0.666
sll0772 conserved hypothetical protein	0.608
sll0788 conserved hypothetical protein	0.264
sll0997 conserved hypothetical protein	0.573
sll1039 conserved hypothetical protein	0.400
sll1218 conserved hypothetical protein	0.591
sll1252 conserved hypothetical protein	0.556
sll1271 conserved hypothetical protein	0.392
sll1306 conserved hypothetical protein	0.487
sll1367 conserved hypothetical protein	0.656
sll1825 conserved hypothetical protein	0.429
slr0083 conserved hypothetical protein	0.626
slr0551 conserved hypothetical protein	0.555
slr0809 conserved hypothetical protein	0.613
slr1104 conserved hypothetical protein	0.664
slr1179 conserved hypothetical protein	0.666
slr1203 conserved hypothetical protein	0.662
slr1260 conserved hypothetical protein	0.354
slr1815 conserved hypothetical protein	0.403
slr1908 conserved hypothetical protein	0.636
slr1926 conserved hypothetical protein protein	0.595
sll1744 cyanelle ribosomal protein L1	0.606
sll1096 cyanelle ribosomal protein S12	0.456
ssr0482 cyanelle ribosomal protein S16	0.607
slr2002 cyanophycin synthetase	0.612
sll0813 cytochrome c oxidase subunit II	0.511
slr1379 cytochrome d ubiquinol oxidase subunit I	0.593
sll1258 Deoxycytidine triphosphate deaminase family protein	0.612

slr2058 DNA topoisomerase i	0.615
slr1265 DNA-directed RNA polymerase gamma chain	0.519
sll1427 DO serine protease	0.588
slr0839 ferrochelatase	0.583
slr1233 fumarate reductase flavoprotein subunit	0.484
slr1143 GGDEF family protein	0.589
slr1238 glutathione synthetase	0.563
slr0884 glyceraldehyde 3-phosphate dehydrogenase 1	0.298
sll0301 HglK	0.426
slr1259 homology to a plant EST:RICS2753A	0.569
sll1226 hoxH	0.664
sll1514 HspA	0.522
sll1221 hydrogenase subunit (hoxF)	0.465
sll1080 hypothetical 36.3 kd lipoprotein precursor, putative	0.297
sll0069 hypothetical protein	0.485
sll0172 hypothetical protein	0.397
sll0446 hypothetical protein	0.602
sll0606 hypothetical protein	0.534
sll0625 hypothetical protein	0.594
sll0854 hypothetical protein	0.598
sll0877 hypothetical protein	0.612
sll0911 hypothetical protein	0.630
sll1049 hypothetical protein	0.313
sll1217 hypothetical protein	0.451
sll1545 hypothetical protein	0.439
sll1571 hypothetical protein	0.597
sll1862 hypothetical protein	0.571
slr0226 hypothetical protein	0.595
slr0439 hypothetical protein	0.491
slr0575 hypothetical protein	0.629
slr0637 hypothetical protein	0.519
slr0645 hypothetical protein	0.565
slr0658 hypothetical protein	0.649
slr0708 hypothetical protein	0.534
slr0818 hypothetical protein	0.460
slr1178 hypothetical protein	0.644
slr1385 hypothetical protein	0.533
slr1612 hypothetical protein	0.629
slr1616 hypothetical protein	0.507
slr1681 hypothetical protein	0.318
slr1931 hypothetical protein	0.411
slr1944 hypothetical protein	0.388
slr1951 hypothetical protein	0.588
slr2004 hypothetical protein	0.577
slr2018 hypothetical protein	0.516
slr2144 hypothetical protein	0.467
ssl1918 hypothetical protein	0.310
slr0721 malate oxidoreductase	0.657

sll1053membrane fusion protein mtre precursor, putative0.488sll124methyl-accepting chemotaxis protein, putative0.372slr0904Mg(2+) chelatase family protein0.602sll0080N-acctyl-glutamate semialdehyde dehydrogenase ArgL0.595slr0891N-acctyl-glutamate semialdehyde dehydrogenase ArgL0.595sll1453nitrate transporter protein (mtD) homolog (mpD)0.291sll1452ntrC-like transporter protein (cmpC)0.451sll0002penicillin-binding protein0.519sll1679periplasmic serine protease0.538sll10920phosphoenolpyruvate carboxylase0.530sll0395phosphoglycerate mutase, putative0.576slr1254phytoene dehydrogenase0.650sll1649pilin0.514slr0162pilin biogenesis protein-related0.500sll2009processing protease, putative0.637sll2009processing protease, putative0.654slr0655putative acontate hydratase0.544slr0655putative acontate hydragenase0.518sll2009processing protein enlated0.577sll1404putative acontate hydragenase0.518sll1641putative alcohol dehydrogenase0.518sll1642putative alcohol dehydrogenase0.518sll1641putative alcohol dehydrogenase0.518sll1641putative alcohol dehydrogenase0.518sll1641putative alcohol dehydrogenase0.518sll1641putative al		
$ slr0904 Mg(2+) chelatase family protein 0.602 \\ sll0080 N-acetyl-glutamate semialdehyde dehydrogenase ArgL 0.595 \\ slr0891 N-acetyl-muramoyl-L-alanine amidase, putative, putative 0.658 \\ sll1453 nirate transporter protein (mrD) homolog (cmpD) 0.291 \\ sll1452 nrtC-like transporter protein (cmpC) 0.451 \\ sll0002 penicillin-binding protein 1 0.519 \\ sll1079 periplasmic serine protease 0.588 \\ sll1196 Pfk1 0.661 \\ sll0920 phosphoenolpyruvate carboxylase 0.530 \\ sll0920 phosphoenolpyruvate carboxylase 0.530 \\ sll0395 phosphoglycerate mutase, putative 0.576 \\ slr1254 phytoene dehydrogenase 0.650 \\ sll0162 pilin biogenesis protein-related 0.500 \\ sll1064 pilin 0.574 \\ slr0162 pilin biogenesis protein-related 0.500 \\ sll2009 processing protease, putative 0.637 \\ sll2009 processing protease, putative 0.654 \\ slr065 putative acohate hydragenase 0.544 \\ slr065 putative acohate hydrogenase 0.518 \\ sll2009 processing protease, putative 0.654 \\ slr0651 putative acohate hydragenase 0.518 \\ sll0742 putative alcohol dehydrogenase 0.518 \\ sll0641 putative carboxylase 0.518 \\ sll0553 putative ortotin -related 0.577 \\ sll1641 putative glutamate decarboxylase 0.518 \\ sll0553 putative protein -related 0.577 \\ sll1641 putative glutamate decarboxylase 0.518 \\ sll0715 putative protein-related 0.577 \\ sll041 putative alcohol dehydrogenase 0.518 \\ sll0541 pyruvate-flavodoxin oxidoreductase 0.588 \\ sll0741 pyruvate-flavodoxin oxidoreductase 0.588 \\ sll0741 pyruvate-flavodoxin oxidoreductase 0.589 \\ sll0743 treprotein-related 0.514 \\ sll0773 streptogramin B lactonase, putative 0.639 \\ sll0773 treptogramin B lactonase, putative 0.639 \\ sll0743 trenestore $	sll1053 membrane fusion protein mtrc precursor, putative	0.488
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sll1750 urease alpha subunit 0.657	slr1031 tyrosyl-tRNA synthetase	0.636
	sll0764 urea/short-chain amide ABC transporter, ATP-binding protein, putative	0.337
sll0915 zinc protease, putative 0.595	sll1750 urease alpha subunit	0.657
	sll0915 zinc protease, putative	0.595

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