Table 2 | Continued

Gene ID	Predicted function	Gene	Replicate 1	Replicate
METTOv1_370002	NAD-linked formate dehydrogenase, subunit D	fdsD	368	312
METTOv1_220028	NAD-linked formate dehydrogenase, subunit A	fdhA2	9	7
C1-ASSIMILATION:SE	RINE CYCLE			
METTOv1_130002	Phosphoenolpyruvate carboxylase	ррс2	104	89
METTOv1_400011	Glycerate kinase	gckA	229	211
METTOv1_400012	Conserved protein of unknown function	orf1	626	708
METTOv1_400013	Malyl-CoA lyase/beta-methylmalyl-CoA lyase	mclA	1713	1615
METTOv1_400014	Phosphoenolpyruvate carboxylase	ppc1	141	139
METTOv1_400015	Malate thiokinase, small subunit	mtkB	516	485
METTOv1_400016	Malate thiokinase, large subunit	mtkA	534	455
METTOv1_400017	Methenyltetrahydrofolate cyclohydrolase	fch	355	281
METTOv1_400018	NADP-dependent methylenetetrahydrofolate dehydrogenase	mtdA	281	243
METTOv1_400019	2-Hydroxyacid dehydrogenase NAD-binding	hprA	375	348
METTOv1_400020	Serine-glyoxylate transaminase	sga	1840	1969
METTOv1_400021	Formate-tetrahydrofolate ligase	ftfL	448	412
METTOv1_670019	Serine hydroxymethyltransferase	glyA	1342	1197
METTOv1_20135	Enolase	eno	432	408
	P PATHWAY AND PHB CYCLE			
METTOv1_100079	Acetyl-CoA acetyltransferase	phaA	597	561
METTOv1_100080	Acetoacetyl-CoA reductase	phaB	1160	1060
METTOv1_50006	Crotonase	croR	235	275
METTOv1 110068	Crotonyl-CoA reductase	ccr	577	523
METTOv1_60013	Ethylmalonyl-CoA mutase	ecm	187	162
METTOv1_510010	Methylsuccinyl-CoA dehydrogenase	ibd	309	295
METTOv1_110043	Mesaconyl-CoA hydratase	meaC	341	317
METTOv1_30129	Methylmalonyl-CoA epimerase	epm	428	394
METTOv1_220010	Malyl-CoA lyase/beta-Methylmalyl-CoA lyase	mclA2	137	135
METTOv1_200020	Acetyl/propionyl-CoA carboxylase	ррсА	353	310
METTOv1_220035	Propionyl-CoA carboxylase	ррсВ	472	455
METTOv1_50067	Methylmalonyl-CoA mutase, large subunit	mcmA	201	188
METTOv1_10062	Methylmalonyl-CoA mutase small subunit B	mcmB	144	144
METTOv1_270063	3-Hydroxybutyrate dehydrogenase	bdhA	235	232
METTOv1_130047	Poly-beta-hydroxybutyrate polymerase	phaC	30	30
METTOv1_200042	Acetoacetate decarboxylase	aad	123	114
METTOv1_200022	Acetoacetyl-coenzyme A synthetase	aas	116	114
METTOv1 630008	Polyhydroxyalkanoate depolymerase	phaZ	237	263
C1-ASSIMILATION: TC	. , , , , , , , , , , , , , , , , , , ,	J		
METTOv1_360040	Malate dehydrogenase	mdh	539	473
METTOv1_360041	Succinyl-CoA synthetase, beta subunit	sucC	660	631
METTOv1_510003	Succinyl-CoA synthetase, alpha subunit	sucD	1198	1135
METTOv1_510002	2-Oxoglutarate dehydrogenase E1	sucA	236	237
METTOv1_370050	2-Oxoglutarate dehydrogenase E2	sucB	191	181
METTOv1_80046	Succinate:ubiquinone oxidoreductase	sdhB	327	348
METTOv1_80046	Succinate:ubiquinone oxidoreductase	sdhA	311	299
METTOV1_80051	Succinate:ubiquinone oxidoreductase, cytochrome b556 subunit	sdhC	318	329
METTOV1_40061	Fumarate hydratase	fum	196	185
METTOV1_40001 METTOV1 1080004	2-Oxoacid ferredoxin oxidoreductase	ofr	79	77
	BOLISM AND ANAPLEROTIC CO ₂ -FIXATION	UII	7 3	11
METTOv1_70038	Phosphoenolpyruvate synthase	nne	28	26
_	Pyruvate carboxylase	pps	28 145	140
METTOv1_120036	Acetyl-coenzyme A carboxylase subunit beta	pcx	246	228
METTOv1_830002	ACETYI-COEHZYITIE A CAIDOXYIASE SUDUIIIL DETA	accD	240	220

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