

**Proteins identified from the proteomic analysis of
Burkholderia vietnamiensis G4**

Locus	% coverage	Description
gj 115350111	81.5	histone family protein DNA-binding protein [<i>B.ambifaria</i> AMMD]
gj 115352352	79.9	ketol-acid reductoisomerase [<i>B.ambifaria</i> AMMD]
gj 115350437	79	protein of unknown function DUF796 [<i>B.ambifaria</i> AMMD]
gj 115351862	79	30S ribosomal protein S6 [<i>B.ambifaria</i> AMMD]
gj 115351796	78.8	RNA-binding protein Hfq [<i>B.ambifaria</i> AMMD]
gj 115351817	77.8	thioredoxin [<i>B.ambifaria</i> AMMD]
gj 115350333	77.5	50S ribosomal protein L24 [<i>B.ambifaria</i> AMMD]
gj 115352972	76.8	nitrogen regulatory protein P-II [<i>B.ambifaria</i> AMMD]
gj 115351979	75.8	alkyl hydroperoxide reductase/ Thiol specific antioxidant/ Mal allergen [<i>B.ambifaria</i> AMMD]
gj 115350871	75.4	riboflavin synthase subunit beta [<i>B.ambifaria</i> AMMD]
gj 115350462	74.6	cold-shock DNA-binding domain protein [<i>B.ambifaria</i> AMMD]
gj 3916735	74.2	10 kDa heat shock protein GroES [<i>B.cepacia</i>]
gj 115350323	73.8	50S ribosomal protein L4 [<i>B.ambifaria</i> AMMD]
gj 115350324	73.1	50S ribosomal protein L23 [<i>B.ambifaria</i> AMMD]
gj 115350339	72.7	30S ribosomal protein S5 [<i>B.ambifaria</i> AMMD]
gj 115352600	72.7	50S ribosomal protein L28 [<i>B.ambifaria</i> AMMD]
gj 115358060	69.8	malate dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 115350334	69.3	50S ribosomal protein L5 [<i>B.ambifaria</i> AMMD]
gj 134295913	68.7	hypothetical protein Bcep1808_1809 [<i>B.vietnamiensis</i> G4]
gj 115350326	68.1	30S ribosomal protein S19 [<i>B.ambifaria</i> AMMD]
gj 115352903	68.1	sigma 54 modulation protein/ribosomal protein S30EA [<i>B.ambifaria</i> AMMD]
gj 3916736	67.2	57 kDa heat shock protein GroEL [<i>B.cepacia</i>]
gj 115352195	66.7	phosphopyruvate hydratase [<i>B.ambifaria</i> AMMD]
gj 115352222	66	phasin family protein [<i>B.ambifaria</i> AMMD]
gj 115352747	65.7	succinyl-CoA synthetase subunit beta [<i>B.ambifaria</i> AMMD]
gj 115350593	64.9	glyceraldehyde-3-phosphate dehydrogenase, type I [<i>B.ambifaria</i> AMMD]
gj 587502	64.6	ATPase beta-subunit [<i>B.cepacia</i>]
gj 115350322	64.1	50S ribosomal protein L3 [<i>B.ambifaria</i> AMMD]
gj 115350312	63.3	50S ribosomal protein L10 [<i>B.ambifaria</i> AMMD]
gj 134293952	63	porin, Gram-negative type [<i>B.vietnamiensis</i> G4]
gj 134296389	63	hypothetical protein Bcep1808_2290 [<i>B.vietnamiensis</i> G4]
gj 115350318	62.8	30S ribosomal protein S7 [<i>B.ambifaria</i> AMMD]
gj 115350348	62.5	DNA-directed RNA polymerase subunit alpha [<i>B.ambifaria</i> AMMD]
gj 115351981	62.1	2C-methyl-D-erythritol 2,4-cyclodiphosphate synthase [<i>B.ambifaria</i> AMMD]
gj 115352285	62	bacterioferritin [<i>B.ambifaria</i> AMMD]
gj 115350327	61.5	50S ribosomal protein L22 [<i>B.ambifaria</i> AMMD]
gj 115360798	60.9	Extracellular ligand-binding receptor [<i>B.ambifaria</i> AMMD]
gj 134292382	60.8	hypothetical protein Bcep1808_3668 [<i>B.vietnamiensis</i> G4]

gj 115350341	60.4	50S ribosomal protein L15 [<i>B.ambifaria</i> AMMD]
gj 115351563	60.3	putative serine protein kinase, PrkA [<i>B.ambifaria</i> AMMD]
gj 115350332	59.8	50S ribosomal protein L14 [<i>B.ambifaria</i> AMMD]
gj 115350977	59.2	30S ribosomal protein S1 [<i>B.ambifaria</i> AMMD]
gj 115350149	59	F0F1 ATP synthase subunit B [<i>B.ambifaria</i> AMMD]
gj 115350328	59	30S ribosomal protein S3 [<i>B.ambifaria</i> AMMD]
gj 115350345	58.7	30S ribosomal protein S13 [<i>B.ambifaria</i> AMMD]
gj 115350417	58.6	Glutathione S-transferase, N-terminal domain [<i>B.ambifaria</i> AMMD]
gj 115351963	58	trigger factor [<i>B.ambifaria</i> AMMD]
gj 115350338	57.9	50S ribosomal protein L18 [<i>B.ambifaria</i> AMMD]
gj 115350618	57.9	extracellular solute-binding protein, family 3 [<i>B.ambifaria</i> AMMD]
gj 115350330	57.8	50S ribosomal protein L29 [<i>B.ambifaria</i> AMMD]
gj 115352918	57	phosphopantetheine adenyltransferase [<i>B.ambifaria</i> AMMD]
gj 115350337	56.8	50S ribosomal protein L6 [<i>B.ambifaria</i> AMMD]
gj 15705849	56.8	RecA [<i>B.cepacia</i>]
gj 115353086	56.4	hypothetical protein Bamb_3035 [<i>B.ambifaria</i> AMMD]
gj 115350539	56.3	50S ribosomal protein L27 [<i>B.ambifaria</i> AMMD]
gj 115351727	56.3	extracellular solute-binding protein, family 1 [<i>B.ambifaria</i> AMMD]
gj 115350151	55.8	F0F1 ATP synthase subunit alpha [<i>B.ambifaria</i> AMMD]
gj 134291078	55.7	UspA domain protein [<i>B.vietnamiensis</i> G4]
gj 115350882	55.4	extracellular solute-binding protein, family 1 [<i>B.ambifaria</i> AMMD]
gj 134295800	55.4	OmpW family protein [<i>B.vietnamiensis</i> G4]
gj 115352619	55.2	cold-shock DNA-binding domain protein [<i>B.ambifaria</i> AMMD]
gj 115351011	54.8	peptidase U62, modulator of DNA gyrase [<i>B.ambifaria</i> AMMD]
gj 115350969	54.5	OmpA/MotB domain protein [<i>B.ambifaria</i> AMMD]
gj 115350321	54.4	30S ribosomal protein S10 [<i>B.ambifaria</i> AMMD]
gj 115351860	53.8	30S ribosomal protein S18 [<i>B.ambifaria</i> AMMD]
gj 115350999	53.6	30S ribosomal protein S16 [<i>B.ambifaria</i> AMMD]
gj 115350336	53.4	30S ribosomal protein S8 [<i>B.ambifaria</i> AMMD]
gj 115350349	53.4	50S ribosomal protein L17 [<i>B.ambifaria</i> AMMD]
gj 115350538	53.4	50S ribosomal protein L21 [<i>B.ambifaria</i> AMMD]
gj 115352103	53.3	30S ribosomal protein S2 [<i>B.ambifaria</i> AMMD]
gj 30314430	53.3	outer membrane lipoprotein precursor [<i>B.cepacia</i>]
gj 115350746	53	protein of unknown function DUF883, ElaB [<i>B.ambifaria</i> AMMD]
gj 160222542	53	60 kDa chaperonin [<i>B.cepacia</i>]
gj 115350154	52.5	F0F1 ATP synthase subunit epsilon [<i>B.ambifaria</i> AMMD]
gj 115351778	52.4	acetyl-CoA reductase [<i>B.ambifaria</i> AMMD]
gj 115350971	52	hypothetical protein Bamb_0917 [<i>B.ambifaria</i> AMMD]
gj 115359754	51.9	UspA domain protein [<i>B.ambifaria</i> AMMD]
gj 134291906	51.9	GDP-mannose 4,6-dehydratase [<i>B.vietnamiensis</i> G4]
gj 115350122	51.7	two component transcriptional regulator, winged helix family [<i>B.ambifaria</i> AMMD]
gj 115352642	51.6	Superoxide dismutase [<i>B.ambifaria</i> AMMD]
gj 134291737	51.6	periplasmic protein of unknown function, DUF1488 [<i>B.vietnamiensis</i> G4]
gj 115350375	51.5	protein of unknown function DUF1568 [<i>B.ambifaria</i> AMMD]
gj 115350610	51.4	2-dehydro-3-deoxyphosphogluconate aldolase/4-hydroxy-2-oxoglutarate aldolase [<i>B.ambifaria</i> AMMD]

gij115350627	51.4	50S ribosomal protein L13 [<i>B.ambifaria</i> AMMD]
gij17863800	51.4	ribosomal subunit S21 [<i>B.cepacia</i>]
gij189348233	51.4	putative molecular chaperone [<i>B.multivorans</i> ATCC 17616]
gij115352960	50.9	preprotein translocase subunit SecB [<i>B.ambifaria</i> AMMD]
gij134291736	50.7	heat shock protein Hsp20 [<i>B.vietnamiensis</i> G4]
gij115351822	50.6	50S ribosomal protein L31 type B [<i>B.ambifaria</i> AMMD]
gij115358067	50.6	type II citrate synthase [<i>B.ambifaria</i> AMMD]
gij115350346	50.4	30S ribosomal protein S11 [<i>B.ambifaria</i> AMMD]
gij115350315	50.1	DNA-directed RNA polymerase subunit beta' [<i>B.ambifaria</i> AMMD]
gij134290950	50	histone family protein DNA-binding protein [<i>B.vietnamiensis</i> G4]
gij115350992	49.8	electron transfer flavoprotein beta-subunit [<i>B.ambifaria</i> AMMD]
gij115352343	49.4	30S ribosomal protein S15 [<i>B.ambifaria</i> AMMD]
gij134291889	49.4	putative flagellar transcriptional activator FlhD [<i>B.vietnamiensis</i> G4]
gij134292901	49.2	protein of unknown function DUF883, ElaB [<i>B.vietnamiensis</i> G4]
gij115350331	48.9	30S ribosomal protein S17 [<i>B.ambifaria</i> AMMD]
gij115352237	48.8	glutamine synthetase, type I [<i>B.ambifaria</i> AMMD]
gij115350409	48.7	twin argininte translocase protein A [<i>B.ambifaria</i> AMMD]
gij115358065	48.5	succinate dehydrogenase iron-sulfur subunit [<i>B.ambifaria</i> AMMD]
gij134291916	48.4	NUDIX hydrolase [<i>B.vietnamiensis</i> G4]
gij134292837	48.4	hypothetical protein Bcep1808_4128 [<i>B.vietnamiensis</i> G4]
gij115358036	47.9	6-phosphogluconate dehydrogenase [<i>B.ambifaria</i> AMMD]
gij115350391	47.6	toluene tolerance family protein [<i>B.ambifaria</i> AMMD]
gij115350335	47.5	30S ribosomal protein S14 [<i>B.ambifaria</i> AMMD]
gij115352601	47.3	50S ribosomal protein L33 [<i>B.ambifaria</i> AMMD]
gij115351783	47.1	competence lipoprotein ComL [<i>B.ambifaria</i> AMMD]
gij115351414	46.9	50S ribosomal protein L35 [<i>B.ambifaria</i> AMMD]
gij115351804	46.8	nucleoside diphosphate kinase [<i>B.ambifaria</i> AMMD]
gij115350311	46.6	50S ribosomal protein L1 [<i>B.ambifaria</i> AMMD]
gij115353188	46.4	transcriptional regulators, TraR/DksA family [<i>B.ambifaria</i> AMMD]
gij115350347	46	30S ribosomal protein S4 [<i>B.ambifaria</i> AMMD]
gij115352298	45.9	uncharacterized protein probably involved in high-affinity Fe ²⁺ transport [<i>B.ambifaria</i> AMMD]
gij115352627	45.7	translation elongation factor G [<i>B.ambifaria</i> AMMD]
gij115353180	45.6	S-adenosylmethionine synthetase [<i>B.ambifaria</i> AMMD]
gij115350310	45.5	50S ribosomal protein L11 [<i>B.ambifaria</i> AMMD]
gij115352746	45.4	succinyl-CoA synthetase subunit alpha [<i>B.ambifaria</i> AMMD]
gij115350791	45.3	hypothetical protein Bamb_0737 [<i>B.ambifaria</i> AMMD]
gij115350421	45.1	extracellular solute-binding protein, family 3 [<i>B.ambifaria</i> AMMD]
gij134294323	44.9	histone family protein nucleoid-structuring protein H-NS [<i>B.vietnamiensis</i> G4]
gij171318607	44.9	histone family protein DNA-binding protein [<i>B.ambifaria</i> MEX-5]
gij134291716	44.8	acetoacetyl-CoA reductase [<i>B.vietnamiensis</i> G4]
gij115359110	44.6	hypothetical protein Bamb_4362 [<i>B.ambifaria</i> AMMD]
gij115351836	44.2	ATPase AAA-2 domain protein [<i>B.ambifaria</i> AMMD]
gij115352166	44	Peptidylprolyl isomerase [<i>B.ambifaria</i> AMMD]
gij115352101	43.9	uridylate kinase [<i>B.ambifaria</i> AMMD]
gij115351818	43.8	transcription termination factor Rho [<i>B.ambifaria</i> AMMD]
gij154362145	43.6	RNA polymerase beta subunit [<i>B.cepacia</i>]

gj 115351815	43.5	hypothetical protein Bamb_1764 [<i>B.ambifaria</i> AMMD]
gj 115358620	43.5	Alcohol dehydrogenase GroES domain protein [<i>B.ambifaria</i> AMMD]
gj 115358813	43.3	cold-shock DNA-binding domain protein [<i>B.ambifaria</i> AMMD]
gj 115350309	43.2	transcription antitermination protein NusG [<i>B.ambifaria</i> AMMD]
gj 115351054	43	acyl carrier protein [<i>B.ambifaria</i> AMMD]
gj 115351223	43	transcription elongation factor GreA [<i>B.ambifaria</i> AMMD]
gj 134294822	42.5	GrpE protein [<i>B.vietnamiensis</i> G4]
gj 115352167	42.3	peptidyl-prolyl cis-trans isomerase, cyclophilin type [<i>B.ambifaria</i> AMMD]
gj 115352225	42.3	pyruvate dehydrogenase subunit E1 [<i>B.ambifaria</i> AMMD]
gj 115350242	42.2	S-adenosyl-L-homocysteine hydrolase [<i>B.ambifaria</i> AMMD]
gj 115352973	42.2	protein of unknown function DUF526 [<i>B.ambifaria</i> AMMD]
gj 86370994	42.2	ribosomal S12 protein [<i>B.cepacia</i>]
gj 115352623	42.1	isocitrate dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 115350248	41.8	Extracellular ligand-binding receptor [<i>B.ambifaria</i> AMMD]
gj 134292338	41.5	cationic amino acid ABC transporter, periplasmic binding protein [<i>B.vietnamiensis</i> G4]
gj 115350160	41.4	Extracellular ligand-binding receptor [<i>B.ambifaria</i> AMMD]
gj 115352725	41.4	pyruvate kinase [<i>B.ambifaria</i> AMMD]
gj 115352647	40.9	adenylate kinase [<i>B.ambifaria</i> AMMD]
gj 115350628	40.8	30S ribosomal protein S9 [<i>B.ambifaria</i> AMMD]
gj 115350826	40.5	Phosphomannomutase [<i>B.ambifaria</i> AMMD]
gj 115350747	40.1	Peroxidase [<i>B.ambifaria</i> AMMD]
gj 115351961	40	ATP-dependent protease ATP-binding subunit [<i>B.ambifaria</i> AMMD]
gj 115352102	39.9	elongation factor Ts [<i>B.ambifaria</i> AMMD]
gj 115352900	39.8	OstA family protein [<i>B.ambifaria</i> AMMD]
gj 134291717	39.8	hypothetical protein Bcep1808_6331 [<i>B.vietnamiensis</i> G4]
gj 115352171	39.7	inositol monophosphatase [<i>B.ambifaria</i> AMMD]
gj 115350557	39.5	Redoxin domain protein [<i>B.ambifaria</i> AMMD]
gj 115352342	39.2	polynucleotide phosphorylase/polyadenylase [<i>B.ambifaria</i> AMMD]
gj 115350314	39.1	DNA-directed RNA polymerase subunit beta [<i>B.ambifaria</i> AMMD]
gj 115359727	39.1	3-hydroxybutyrate dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 115350709	39	phosphoribosylaminoimidazole synthetase [<i>B.ambifaria</i> AMMD]
gj 115350408	38.8	histidine triad (HIT) protein [<i>B.ambifaria</i> AMMD]
gj 115350503	38.8	enoyl-CoA hydratase [<i>B.ambifaria</i> AMMD]
gj 134291721	38.6	UspA domain protein [<i>B.vietnamiensis</i> G4]
gj 115350885	38.5	ABC transporter related [<i>B.ambifaria</i> AMMD]
gj 115352113	38.5	2,3,4,5-tetrahydropyridine-2-carboxylate N-succinyltransferase [<i>B.ambifaria</i> AMMD]
gj 115351392	38.4	putative transcriptional regulator, XRE family [<i>B.ambifaria</i> AMMD]
gj 115359179	38.4	Extracellular ligand-binding receptor [<i>B.ambifaria</i> AMMD]
gj 115351980	38.3	Carboxymuconolactone decarboxylase [<i>B.ambifaria</i> AMMD]
gj 134295827	38	protein of unknown function DUF465 [<i>B.vietnamiensis</i> G4]
gj 115358580	37.8	Succinate-semialdehyde dehydrogenase (NAD(P)(+)) [<i>B.ambifaria</i> AMMD]
gj 134295935	37.8	hypothetical protein Bcep1808_1832 [<i>B.vietnamiensis</i> G4]
gj 134296969	37.8	phasin family protein [<i>B.vietnamiensis</i> G4]
gj 115351002	37.7	50S ribosomal protein L19 [<i>B.ambifaria</i> AMMD]

gj 115350843	37.6	protein-L-isoaspartate(D-aspartate) O-methyltransferase [<i>B.ambifaria</i> AMMD]
gj 115350231	37.4	response regulator receiver protein [<i>B.ambifaria</i> AMMD]
gj 115352827	37.3	aspartyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gj 116689141	37.3	50S ribosomal protein L32 [<i>B.cenocepacia</i> HI2424]
gj 115350325	37.1	50S ribosomal protein L2 [<i>B.ambifaria</i> AMMD]
gj 115351507	37.1	extracellular solute-binding protein, family 3 [<i>B.ambifaria</i> AMMD]
gj 115352100	37.1	ribosome recycling factor [<i>B.ambifaria</i> AMMD]
gj 115350369	36.9	glycerol-3-phosphate transporter periplasmic binding protein [<i>B.ambifaria</i> AMMD]
gj 115350436	36.7	protein of unknown function DUF877 [<i>B.ambifaria</i> AMMD]
gj 115351732	36.7	OmpW family protein [<i>B.ambifaria</i> AMMD]
gj 134291907	36.6	exopolysaccharide transport protein family [<i>B.vietnamiensis</i> G4]
gj 115350698	36.5	GrpE protein [<i>B.ambifaria</i> AMMD]
gj 134294235	36.3	F0F1 ATP synthase subunit delta [<i>B.vietnamiensis</i> G4]
gj 134291169	36.1	histone family protein nucleoid-structuring protein H-NS [<i>B.vietnamiensis</i> G4]
gj 115351230	35.8	phosphate ABC transporter, periplasmic phosphate-binding protein [<i>B.ambifaria</i> AMMD]
gj 115352723	35.8	phosphoribosylaminoimidazole-succinocarboxamide synthase [<i>B.ambifaria</i> AMMD]
gj 115350115	35.6	uncharacterized enzyme [<i>B.ambifaria</i> AMMD]
gj 115351436	35.6	ribosome-binding factor A [<i>B.ambifaria</i> AMMD]
gj 115352336	35.5	NADH dehydrogenase subunit C [<i>B.ambifaria</i> AMMD]
gj 134296638	35.3	hypothetical protein Bcep1808_2540 [<i>B.vietnamiensis</i> G4]
gj 115353149	35.2	Ferredoxin--NADP(+) reductase [<i>B.ambifaria</i> AMMD]
gj 115351550	35.1	sulfate ABC transporter, periplasmic sulfate-binding protein [<i>B.ambifaria</i> AMMD]
gj 115350340	35	50S ribosomal protein L30 [<i>B.ambifaria</i> AMMD]
gj 115350434	34.9	Tetratricopeptide TPR_2 repeat protein [<i>B.ambifaria</i> AMMD]
gj 115353194	34.9	acetylglutamate kinase [<i>B.ambifaria</i> AMMD]
gj 115352513	34.8	inorganic pyrophosphatase [<i>B.ambifaria</i> AMMD]
gj 134292692	34.7	LemA family protein [<i>B.vietnamiensis</i> G4]
gj 115353018	34.6	hypothetical protein Bamb_2967 [<i>B.ambifaria</i> AMMD]
gj 115353216	34.6	cell shape determining protein, MreB/Mrl family [<i>B.ambifaria</i> AMMD]
gj 134291891	34.4	UTP-glucose-1-phosphate uridylyltransferase GalU [<i>B.vietnamiensis</i> G4]
gj 134294783	34.4	protein of unknown function DUF330 [<i>B.vietnamiensis</i> G4]
gj 115352094	34.3	outer membrane chaperone Skp (OmpH) [<i>B.ambifaria</i> AMMD]
gj 115350983	34.2	ADP-L-glycero-D-manno-heptose-6-epimerase [<i>B.ambifaria</i> AMMD]
gj 115351435	34.2	translation initiation factor IF-2 [<i>B.ambifaria</i> AMMD]
gj 115350835	34	urease subunit gamma [<i>B.ambifaria</i> AMMD]
gj 115350313	33.9	50S ribosomal protein L7/L12 [<i>B.ambifaria</i> AMMD]
gj 134296812	33.9	RND efflux system, outer membrane lipoprotein, NodT family [<i>B.vietnamiensis</i> G4]
gj 115352888	33.7	single-strand DNA-binding protein [<i>B.ambifaria</i> AMMD]
gj 115351443	33.6	alpha-ketoglutarate decarboxylase [<i>B.ambifaria</i> AMMD]
gj 134292631	33.6	hypothetical protein Bcep1808_3921 [<i>B.vietnamiensis</i> G4]
gj 6683465	33.5	DsbA [<i>B.cepacia</i>]

gj 115350058	33.2	DNA polymerase III subunit beta [<i>B.ambifaria</i> AMMD]
gj 115358095	33.2	aspartate-semialdehyde dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 115358056	33	aconitate hydratase [<i>B.ambifaria</i> AMMD]
gj 115350291	32.9	orotate phosphoribosyltransferase [<i>B.ambifaria</i> AMMD]
gj 115359093	32.9	Lysine decarboxylase [<i>B.ambifaria</i> AMMD]
gj 115350919	32.8	cold-shock DNA-binding domain protein [<i>B.ambifaria</i> AMMD]
gj 115351415	32.8	50S ribosomal protein L20 [<i>B.ambifaria</i> AMMD]
gj 115358650	32.8	cold-shock DNA-binding domain protein [<i>B.ambifaria</i> AMMD]
gj 134293121	32.8	Glutathione S-transferase, C-terminal domain [<i>B.vietnamiensis</i> G4]
gj 115350879	32.7	glucokinase [<i>B.ambifaria</i> AMMD]
gj 115351413	32.7	translation initiation factor IF-3 [<i>B.ambifaria</i> AMMD]
gj 115351962	32.7	ATP-dependent Clp protease proteolytic subunit [<i>B.ambifaria</i> AMMD]
gj 115360688	32.7	cyanate hydratase [<i>B.ambifaria</i> AMMD]
gj 134293177	32.6	porin, Gram-negative type [<i>B.vietnamiensis</i> G4]
gj 115352084	32.5	band 7 protein [<i>B.ambifaria</i> AMMD]
gj 134291603	32.5	transport-associated [<i>B.vietnamiensis</i> G4]
gj 115352964	32.4	carboxyl-terminal protease [<i>B.ambifaria</i> AMMD]
gj 115352992	32.4	hypothetical protein Bamb_2941 [<i>B.ambifaria</i> AMMD]
gj 134294034	32.3	phosphate acetyltransferase [<i>B.vietnamiensis</i> G4]
gj 115352914	32.2	50S ribosomal protein L25/general stress protein Ctc [<i>B.ambifaria</i> AMMD]
gj 134293484	32.2	hypothetical protein Bcep1808_4798 [<i>B.vietnamiensis</i> G4]
gj 134295918	32.2	Ankyrin [<i>B.vietnamiensis</i> G4]
gj 115359728	32.1	acetoacetate decarboxylase [<i>B.ambifaria</i> AMMD]
gj 134293320	32.1	TonB-dependent siderophore receptor [<i>B.vietnamiensis</i> G4]
gj 134296609	32	Methyltransferase type 11 [<i>B.vietnamiensis</i> G4]
gj 115351851	31.7	hypothetical protein Bamb_1800 [<i>B.ambifaria</i> AMMD]
gj 134294434	31.7	ribosomal protein L7/L12 [<i>B.vietnamiensis</i> G4]
gj 134291734	31.5	transport-associated [<i>B.vietnamiensis</i> G4]
gj 134291909	31.5	polysaccharide export protein [<i>B.vietnamiensis</i> G4]
gj 134297154	31.5	hypothetical protein Bcep1808_3063 [<i>B.vietnamiensis</i> G4]
gj 10121186	31.4	RecA [<i>B.cepacia</i>]
gj 115350890	31.4	outer-membrane lipoprotein carrier protein [<i>B.ambifaria</i> AMMD]
gj 115352086	31.3	phosphoenolpyruvate synthase [<i>B.ambifaria</i> AMMD]
gj 134292620	31.3	hypothetical protein Bcep1808_3910 [<i>B.vietnamiensis</i> G4]
gj 115350281	31.1	6-phosphogluconate dehydrogenase, NAD-binding [<i>B.ambifaria</i> AMMD]
gj 115352227	31.1	two component transcriptional regulator, LuxR family [<i>B.ambifaria</i> AMMD]
gj 115352334	31.1	NADH dehydrogenase subunit E [<i>B.ambifaria</i> AMMD]
gj 115352910	31	Tetratricopeptide TPR_2 repeat protein [<i>B.ambifaria</i> AMMD]
gj 116693782	31	NADPH-dependent FMN reductase [<i>B.cenocepacia</i> HI2424]
gj 115350913	30.8	guanylate kinase [<i>B.ambifaria</i> AMMD]
gj 115352521	30.8	Methyltransferase type 11 [<i>B.ambifaria</i> AMMD]
gj 115351859	30.7	50S ribosomal protein L9 [<i>B.ambifaria</i> AMMD]
gj 115350888	30.6	thioredoxin reductase [<i>B.ambifaria</i> AMMD]
gj 115351373	30.6	UTP-glucose-1-phosphate uridylyltransferase [<i>B.ambifaria</i> AMMD]

gi 134294857	30.6	hypothetical protein Bcep1808_0745 [<i>B.vietnamiensis</i> G4]
gi 115352507	30.4	nitrogen regulatory protein P-II [<i>B.ambifaria</i> AMMD]
gi 115351052	30.3	malonyl CoA-acyl carrier protein transacylase [<i>B.ambifaria</i> AMMD]
gi 115352425	30.3	transaldolase B [<i>B.ambifaria</i> AMMD]
gi 134291714	30.3	Alcohol dehydrogenase GroES domain protein [<i>B.vietnamiensis</i> G4]
gi 134293948	30.1	putative signal-transduction protein with CBS domains [<i>B.vietnamiensis</i> G4]
gi 115352875	30	acetyl-CoA acetyltransferase [<i>B.ambifaria</i> AMMD]
gi 115350731	29.9	translocation protein TolB [<i>B.ambifaria</i> AMMD]
gi 115358101	29.8	tryptophan synthase subunit alpha [<i>B.ambifaria</i> AMMD]
gi 115351987	29.6	phage shock protein A, PspA [<i>B.ambifaria</i> AMMD]
gi 115352813	29.6	PpiC-type peptidyl-prolyl cis-trans isomerase [<i>B.ambifaria</i> AMMD]
gi 115352941	29.5	lipoprotein, YaeC family [<i>B.ambifaria</i> AMMD]
gi 115352412	29.4	3-carboxymuconate cyclase-like protein [<i>B.ambifaria</i> AMMD]
gi 134291894	29.4	UDP-glucose 6-dehydrogenase [<i>B.vietnamiensis</i> G4]
gi 134291974	29.4	histone family protein nucleoid-structuring protein H-NS [<i>B.vietnamiensis</i> G4]
gi 115350729	29.3	hypothetical protein Bamb_0675 [<i>B.ambifaria</i> AMMD]
gi 115351053	29.3	3-ketoacyl-(acyl-carrier-protein) reductase [<i>B.ambifaria</i> AMMD]
gi 134291872	29.3	extracellular solute-binding protein, family 3 [<i>B.vietnamiensis</i> G4]
gi 115350435	29.2	Uncharacterised conserved protein UCP028301 [<i>B.ambifaria</i> AMMD]
gi 134296814	29.2	efflux transporter, RND family, MFP subunit [<i>B.vietnamiensis</i> G4]
gi 189352142	29.2	conserved hypothetical protein [<i>B.multivorans</i> ATCC 17616]
gi 115350581	29.1	orotidine 5'-phosphate decarboxylase [<i>B.ambifaria</i> AMMD]
gi 134294993	28.8	6-phosphogluconolactonase [<i>B.vietnamiensis</i> G4]
gi 115350317	28.6	30S ribosomal protein S12 [<i>B.ambifaria</i> AMMD]
gi 115351857	28.4	protein of unknown function DUF47 [<i>B.ambifaria</i> AMMD]
gi 115351878	28.4	Ankyrin [<i>B.ambifaria</i> AMMD]
gi 134294955	28.4	protein-L-isoaspartate(D-aspartate) O-methyltransferase [<i>B.vietnamiensis</i> G4]
gi 134297078	28.4	hypothetical protein Bcep1808_2987 [<i>B.vietnamiensis</i> G4]
gi 115350521	28.3	cell division protein FtsA [<i>B.ambifaria</i> AMMD]
gi 115350551	28.3	ribonucleotide-diphosphate reductase subunit alpha [<i>B.ambifaria</i> AMMD]
gi 115353191	28.3	response regulator receiver protein [<i>B.ambifaria</i> AMMD]
gi 134291328	28.3	Carbonate dehydratase [<i>B.vietnamiensis</i> G4]
gi 115350466	28.2	glutaredoxin-like protein [<i>B.ambifaria</i> AMMD]
gi 115350881	28.2	glucose-6-phosphate 1-dehydrogenase [<i>B.ambifaria</i> AMMD]
gi 115351445	28.2	dihydrolipoamide dehydrogenase [<i>B.ambifaria</i> AMMD]
gi 115353176	28.2	aldo/keto reductase [<i>B.ambifaria</i> AMMD]
gi 115352726	28.1	phosphoglycerate kinase [<i>B.ambifaria</i> AMMD]
gi 134294572	28.1	Uncharacterised conserved protein UCP028301 [<i>B.vietnamiensis</i> G4]
gi 134295564	28.1	hypothetical protein Bcep1808_1456 [<i>B.vietnamiensis</i> G4]
gi 115351842	28	threonine synthase [<i>B.ambifaria</i> AMMD]
gi 115352724	28	fructose-1,6-bisphosphate aldolase [<i>B.ambifaria</i> AMMD]
gi 134291688	28	hypothetical protein Bcep1808_6302 [<i>B.vietnamiensis</i> G4]

gj 134296939	28	intracellular protease, Pfpl family [<i>B.vietnamiensis</i> G4]
gj 115350150	27.9	F0F1 ATP synthase subunit delta [<i>B.ambifaria</i> AMMD]
gj 115350592	27.8	transketolase [<i>B.ambifaria</i> AMMD]
gj 115352300	27.8	aromatic amino acid aminotransferase [<i>B.ambifaria</i> AMMD]
gj 116689828	27.8	RNA-binding protein Hfq [<i>B.cenocepacia</i> HI2424]
gj 115352197	27.7	CTP synthetase [<i>B.ambifaria</i> AMMD]
gj 115352574	27.7	Uncharacterised conserved protein UCP030820 [<i>B.ambifaria</i> AMMD]
gj 115359720	27.7	protein of unknown function DUF1348 [<i>B.ambifaria</i> AMMD]
gj 83721283	27.5	thiol peroxidase [<i>B.thailandensis</i> E264]
gj 115352284	27.4	hydro-lyases, Fe-S type, tartrate/fumarate subfamily, alpha subunit [<i>B.ambifaria</i> AMMD]
gj 134294352	27.4	Carboxymethylenebutenolidase [<i>B.vietnamiensis</i> G4]
gj 134296286	27.4	hypothetical protein Bcep1808_2184 [<i>B.vietnamiensis</i> G4]
gj 5924062	27.4	unknown [<i>B.cepacia</i>]
gj 115351960	27.3	ATP-dependent protease La [<i>B.ambifaria</i> AMMD]
gj 134292900	27.3	hypothetical protein Bcep1808_4191 [<i>B.vietnamiensis</i> G4]
gj 134296935	27.3	Endoribonuclease L-PSP [<i>B.vietnamiensis</i> G4]
gj 115350152	27.1	F0F1 ATP synthase subunit gamma [<i>B.ambifaria</i> AMMD]
gj 115350912	27	hypothetical protein Bamb_0858 [<i>B.ambifaria</i> AMMD]
gj 115353213	26.9	aspartyl/glutamyl-tRNA amidotransferase subunit B [<i>B.ambifaria</i> AMMD]
gj 115351367	26.7	SirA family protein [<i>B.ambifaria</i> AMMD]
gj 115352651	26.7	30S ribosomal protein S20 [<i>B.ambifaria</i> AMMD]
gj 134296410	26.6	glutathione S-transferase-like protein [<i>B.vietnamiensis</i> G4]
gj 115352384	26.5	phosphoribosylamine--glycine ligase [<i>B.ambifaria</i> AMMD]
gj 134291888	26.5	hypothetical protein Bcep1808_6505 [<i>B.vietnamiensis</i> G4]
gj 134293081	26.5	peptidase M23B [<i>B.vietnamiensis</i> G4]
gj 115350599	26.4	SmpA/OmlA domain protein [<i>B.ambifaria</i> AMMD]
gj 115352705	26.4	RND efflux system, outer membrane lipoprotein, NodT family [<i>B.ambifaria</i> AMMD]
gj 116688835	26.4	Peroxidase [<i>B.cenocepacia</i> HI2424]
gj 115350686	26	peptidase U62, modulator of DNA gyrase [<i>B.ambifaria</i> AMMD]
gj 115350522	25.9	cell division protein FtsZ [<i>B.ambifaria</i> AMMD]
gj 134296575	25.9	ecotin precursor [<i>B.vietnamiensis</i> G4]
gj 115352335	25.7	NADH dehydrogenase subunit D [<i>B.ambifaria</i> AMMD]
gj 134291738	25.6	UspA domain protein [<i>B.vietnamiensis</i> G4]
gj 134291105	25.5	Linocin_M18 bacteriocin protein [<i>B.vietnamiensis</i> G4]
gj 134293786	25.5	hypothetical protein Bcep1808_5106 [<i>B.vietnamiensis</i> G4]
gj 115350993	25.4	electron transfer flavoprotein, alpha subunit [<i>B.ambifaria</i> AMMD]
gj 115352729	25.4	branched-chain amino acid aminotransferase [<i>B.ambifaria</i> AMMD]
gj 116690382	25.4	hypothetical protein Bcen2424_2362 [<i>B.cenocepacia</i> HI2424]
gj 134295952	25.4	Glucose-6-phosphate isomerase [<i>B.vietnamiensis</i> G4]
gj 134297253	25.4	aldo/keto reductase [<i>B.vietnamiensis</i> G4]
gj 115352075	25.3	inositol-5-monophosphate dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 134292463	25.2	UspA domain protein [<i>B.vietnamiensis</i> G4]
gj 134294333	25.2	methyl-accepting chemotaxis sensory transducer [<i>B.vietnamiensis</i> G4]
gj 115350385	25.1	thiazole synthase [<i>B.ambifaria</i> AMMD]

gij134297084	25.1	ABC-type uncharacterized transport system auxiliary component-like protein [<i>B.vietnamiensis</i> G4]
gij189349329	25.1	peroxiredoxin [<i>B.multivorans</i> ATCC 17616]
gij91784510	25	hypothetical protein Bxe_A1289 [<i>B.xenovorans</i> LB400]
gij115351227	24.9	ATP-dependent metalloprotease FtsH [<i>B.ambifaria</i> AMMD]
gij115351791	24.9	ATP phosphoribosyltransferase regulatory subunit [<i>B.ambifaria</i> AMMD]
gij115350555	24.8	17 kDa surface antigen [<i>B.ambifaria</i> AMMD]
gij115351854	24.8	PhoH family protein [<i>B.ambifaria</i> AMMD]
gij134294984	24.8	transcription antitermination protein NusB [<i>B.vietnamiensis</i> G4]
gij115352530	24.6	Lysine decarboxylase [<i>B.ambifaria</i> AMMD]
gij115351055	24.5	3-oxoacyl-(acyl carrier protein) synthase II [<i>B.ambifaria</i> AMMD]
gij116691095	24.5	aldo/keto reductase [<i>B.cenocepacia</i> HI2424]
gij115350609	24.4	phosphogluconate dehydratase [<i>B.ambifaria</i> AMMD]
gij115351799	24.4	hypothetical protein Bamb_1748 [<i>B.ambifaria</i> AMMD]
gij115352177	24.4	FKBP-type peptidyl-prolyl cis-trans isomerases 2-like protein [<i>B.ambifaria</i> AMMD]
gij134291678	24.4	UspA domain protein [<i>B.vietnamiensis</i> G4]
gij115352707	24.3	efflux transporter, RND family, MFP subunit [<i>B.ambifaria</i> AMMD]
gij134291905	24.1	NAD-dependent epimerase/dehydratase [<i>B.vietnamiensis</i> G4]
gij115351210	23.9	glutathione S-transferase-like protein [<i>B.ambifaria</i> AMMD]
gij115351476	23.9	periplasmic binding protein/LacI transcriptional regulator [<i>B.ambifaria</i> AMMD]
gij115360302	23.9	Peptidylprolyl isomerase [<i>B.ambifaria</i> AMMD]
gij134292090	23.9	Extracellular ligand-binding receptor [<i>B.vietnamiensis</i> G4]
gij134294458	23.9	50S ribosomal protein L6 [<i>B.vietnamiensis</i> G4]
gij115350638	23.8	bifunctional phosphoribosylaminoimidazolecarboxamide formyltransferase/IMP cyclohydrolase [<i>B.ambifaria</i> AMMD]
gij134297229	23.8	Ferredoxin--NADP(+) reductase [<i>B.vietnamiensis</i> G4]
gij115360206	23.7	cationic amino acid ABC transporter, periplasmic binding protein [<i>B.ambifaria</i> AMMD]
gij134292525	23.7	toluene tolerance family protein [<i>B.vietnamiensis</i> G4]
gij115351951	23.5	PpiC-type peptidyl-prolyl cis-trans isomerase [<i>B.ambifaria</i> AMMD]
gij115352497	23.5	two component transcriptional regulator, winged helix family [<i>B.ambifaria</i> AMMD]
gij134295694	23.5	lysine/ornithine N-monooxygenase [<i>B.vietnamiensis</i> G4]
gij134295697	23.5	hypothetical protein Bcep1808_1590 [<i>B.vietnamiensis</i> G4]
gij115350292	23.4	malic enzyme [<i>B.ambifaria</i> AMMD]
gij115350615	23.4	adenylosuccinate lyase [<i>B.ambifaria</i> AMMD]
gij115353023	23.2	extracellular solute-binding protein, family 5 [<i>B.ambifaria</i> AMMD]
gij134296537	23.2	putative transcriptional regulator, GntR family [<i>B.vietnamiensis</i> G4]
gij158702462	23.2	ferric uptake regulator [<i>B.cepacia</i>]
gij4128212	23.2	principal sigma factor [<i>B.cepacia</i>]
gij134290916	23.1	hypothetical protein Bcep1808_5486 [<i>B.vietnamiensis</i> G4]
gij134294994	23.1	glucose-6-phosphate 1-dehydrogenase [<i>B.vietnamiensis</i> G4]
gij115351977	23	osmolarity response regulator [<i>B.ambifaria</i> AMMD]
gij115352229	23	Oligopeptidase A [<i>B.ambifaria</i> AMMD]
gij115352732	23	hypothetical protein Bamb_2681 [<i>B.ambifaria</i> AMMD]
gij115358716	23	transcriptional regulator, MarR family [<i>B.ambifaria</i> AMMD]

gj 115352858	22.9	peptidase M48, Ste24p [<i>B.ambifaria</i> AMMD]
gj 134294595	22.9	nitroreductase [<i>B.vietnamiensis</i> G4]
gj 115351801	22.8	4-hydroxy-3-methylbut-2-en-1-yl diphosphate synthase [<i>B.ambifaria</i> AMMD]
gj 134294574	22.8	protein of unknown function DUF796 [<i>B.vietnamiensis</i> G4]
gj 134294801	22.8	Ferritin, Dps family protein [<i>B.vietnamiensis</i> G4]
gj 115352840	22.7	Endoribonuclease L-PSP [<i>B.ambifaria</i> AMMD]
gj 134294246	22.7	Extracellular ligand-binding receptor [<i>B.vietnamiensis</i> G4]
gj 134295774	22.7	hypothetical protein Bcep1808_1670 [<i>B.vietnamiensis</i> G4]
gj 134297166	22.7	Heavy metal transport/detoxification protein [<i>B.vietnamiensis</i> G4]
gj 115359854	22.6	hypothetical protein Bamb_5109 [<i>B.ambifaria</i> AMMD]
gj 115361014	22.5	extracellular solute-binding protein, family 3 [<i>B.ambifaria</i> AMMD]
gj 115361244	22.5	dihydrolipoamide dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 134296385	22.5	Biopolymer transport protein ExbD/TolR [<i>B.vietnamiensis</i> G4]
gj 115352520	22.4	phosphoenolpyruvate carboxylase [<i>B.ambifaria</i> AMMD]
gj 115352654	22.4	nucleotide-binding protein [<i>B.ambifaria</i> AMMD]
gj 115359687	22.4	malate synthase G [<i>B.ambifaria</i> AMMD]
gj 134295686	22.4	Taurine catabolism dioxygenase TauD/TfdA [<i>B.vietnamiensis</i> G4]
gj 115352160	22.3	aspartate kinase [<i>B.ambifaria</i> AMMD]
gj 115358623	22.3	alpha,alpha-trehalose-phosphate synthase (UDP-forming) [<i>B.ambifaria</i> AMMD]
gj 116688622	22.3	17 kDa surface antigen [<i>B.cenocepacia</i> HI2424]
gj 115352353	22.1	acetolactate synthase 3 regulatory subunit [<i>B.ambifaria</i> AMMD]
gj 134295693	22.1	amino acid adenylation domain [<i>B.vietnamiensis</i> G4]
gj 115352617	21.9	ATP-dependent Clp protease, ATP-binding subunit clpA [<i>B.ambifaria</i> AMMD]
gj 115353103	21.9	porin, Gram-negative type [<i>B.ambifaria</i> AMMD]
gj 115350841	21.8	ABC branched chain amino acid family transporter, periplasmic ligand binding protein [<i>B.ambifaria</i> AMMD]
gj 115350876	21.8	glutamate-1-semialdehyde 2,1-aminomutase [<i>B.ambifaria</i> AMMD]
gj 116688363	21.8	50S ribosomal protein L7/L12 [<i>B.cenocepacia</i> HI2424]
gj 189349441	21.8	putative exported protein [<i>B.multivorans</i> ATCC 17616]
gj 134291690	21.7	hypothetical protein Bcep1808_6304 [<i>B.vietnamiensis</i> G4]
gj 115350612	21.6	carbohydrate kinase, thermoresistant glucokinase family [<i>B.ambifaria</i> AMMD]
gj 115351121	21.6	extracellular solute-binding protein, family 3 [<i>B.ambifaria</i> AMMD]
gj 2738611	21.6	L-ornithine N(5)-oxygenase [<i>B.cepacia</i>]
gj 83719093	21.6	co-chaperonin GroES [<i>B.thailandensis</i> E264]
gj 115351594	21.5	TonB-dependent siderophore receptor [<i>B.ambifaria</i> AMMD]
gj 134292402	21.4	histone family protein nucleoid-structuring protein H-NS [<i>B.vietnamiensis</i> G4]
gj 134295219	21.3	alkyl hydroperoxide reductase/ Thiol specific antioxidant/ Mal allergen [<i>B.vietnamiensis</i> G4]
gj 115352622	21.2	isocitrate dehydrogenase, NADP-dependent [<i>B.ambifaria</i> AMMD]
gj 134295947	21.2	PpiC-type peptidyl-prolyl cis-trans isomerase [<i>B.vietnamiensis</i> G4]
gj 115352506	21.1	outer membrane protein (porin)-like protein [<i>B.ambifaria</i> AMMD]
gj 134294032	21.1	PHB de-polymerase domain protein [<i>B.vietnamiensis</i> G4]
gj 115351434	21	transcription elongation factor NusA [<i>B.ambifaria</i> AMMD]
gj 134292468	21	aldo/keto reductase [<i>B.vietnamiensis</i> G4]

gij115352438	20.9	aminotransferase, class I and II [<i>B.ambifaria</i> AMMD]
gij115357984	20.9	OsmC family protein [<i>B.ambifaria</i> AMMD]
gij116688369	20.9	elongation factor G [<i>B.cenocepacia</i> HI2424]
gij134294706	20.9	2-dehydro-3-deoxy-6-phosphogalactonate aldolase [<i>B.vietnamiensis</i> G4]
gij134295692	20.9	amino acid adenylation domain [<i>B.vietnamiensis</i> G4]
gij115350527	20.8	bifunctional ornithine acetyltransferase/N-acetylglutamate synthase protein [<i>B.ambifaria</i> AMMD]
gij115351222	20.8	carbamoyl phosphate synthase large subunit [<i>B.ambifaria</i> AMMD]
gij115352517	20.8	HemY domain protein [<i>B.ambifaria</i> AMMD]
gij115360458	20.8	UspA domain protein [<i>B.ambifaria</i> AMMD]
gij134296696	20.8	isocitrate dehydrogenase, NADP-dependent [<i>B.vietnamiensis</i> G4]
gij115352206	20.7	lysyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gij116688459	20.7	histidine triad (HIT) protein [<i>B.cenocepacia</i> HI2424]
gij115351798	20.5	Pyrrolo-quinoline quinone [<i>B.ambifaria</i> AMMD]
gij115352566	20.5	leucyl aminopeptidase [<i>B.ambifaria</i> AMMD]
gij115353269	20.5	50S ribosomal protein L34 [<i>B.ambifaria</i> AMMD]
gij116692027	20.5	hypothetical protein Bcen2424_3930 [<i>B.cenocepacia</i> HI2424]
gij134296027	20.5	elongation factor Ts [<i>B.vietnamiensis</i> G4]
gij115352892	20.4	formyltetrahydrofolate deformylase [<i>B.ambifaria</i> AMMD]
gij171319779	20.4	Ferritin Dps family protein [<i>B.ambifaria</i> MEX-5]
gij115350329	20.3	50S ribosomal protein L16 [<i>B.ambifaria</i> AMMD]
gij134292354	20.3	Glutathione S-transferase, N-terminal domain [<i>B.vietnamiensis</i> G4]
gij134296510	20.2	3-carboxymuconate cyclase-like protein [<i>B.vietnamiensis</i> G4]
gij134297105	20.2	arginyl-tRNA synthetase [<i>B.vietnamiensis</i> G4]
gij134292511	20.1	NADPH-dependent FMN reductase [<i>B.vietnamiensis</i> G4]
gij134294657	20.1	bifunctional ornithine acetyltransferase/N-acetylglutamate synthase protein [<i>B.vietnamiensis</i> G4]
gij115351956	20	Glucose-6-phosphate isomerase [<i>B.ambifaria</i> AMMD]
gij115352559	20	protein of unknown function DUF541 [<i>B.ambifaria</i> AMMD]
gij115358064	20	succinate dehydrogenase flavoprotein subunit [<i>B.ambifaria</i> AMMD]
gij134291980	20	flavoprotein WrbA [<i>B.vietnamiensis</i> G4]
gij116688678	19.9	Rare lipoprotein B [<i>B.cenocepacia</i> HI2424]
gij134294025	19.9	Carbohydrate-selective porin OprB [<i>B.vietnamiensis</i> G4]
gij115351097	19.8	GntR domain protein [<i>B.ambifaria</i> AMMD]
gij134294649	19.8	D-alanine--D-alanine ligase [<i>B.vietnamiensis</i> G4]
gij134296968	19.8	acetyl-CoA acetyltransferase [<i>B.vietnamiensis</i> G4]
gij11230854	19.7	ornibactin synthetase F [<i>B.cepacia</i>]
gij115351785	19.7	protein of unknown function DUF465 [<i>B.ambifaria</i> AMMD]
gij115352772	19.7	cytochrome c, class I [<i>B.ambifaria</i> AMMD]
gij134292531	19.7	2-nitropropane dioxygenase, NPD [<i>B.vietnamiensis</i> G4]
gij115350665	19.6	preprotein translocase subunit YajC [<i>B.ambifaria</i> AMMD]
gij115351589	19.6	cyclic peptide transporter [<i>B.ambifaria</i> AMMD]
gij115352953	19.6	cytochrome c oxidase, subunit II [<i>B.ambifaria</i> AMMD]
gij134291881	19.6	histone family protein nucleoid-structuring protein H-NS [<i>B.vietnamiensis</i> G4]
gij134296928	19.6	import inner membrane translocase, subunit Tim44 [<i>B.vietnamiensis</i> G4]

gj 91779140	19.6	DNA-binding protein HU-alpha [<i>B.xenovorans</i> LB400]
gj 115350661	19.5	Ycel family protein [<i>B.ambifaria</i> AMMD]
gj 115360693	19.5	protein of unknown function DUF1348 [<i>B.ambifaria</i> AMMD]
gj 134293175	19.5	histone family protein DNA-binding protein [<i>B.vietnamiensis</i> G4]
gj 134294890	19.5	transport-associated [<i>B.vietnamiensis</i> G4]
gj 115350660	19.4	Ycel family protein [<i>B.ambifaria</i> AMMD]
gj 115359595	19.4	histone family protein nucleoid-structuring protein H-NS [<i>B.ambifaria</i> AMMD]
gj 162389966	19.4	phenylalanyl-tRNA synthetase subunit alpha [<i>B.ambifaria</i> AMMD]
gj 134294673	19.3	hypothetical protein Bcep1808_0561 [<i>B.vietnamiensis</i> G4]
gj 115353141	19.2	extracellular solute-binding protein, family 5 [<i>B.ambifaria</i> AMMD]
gj 115360327	19.2	HAD-superfamily hydrolase, subfamily IA, variant 3 [<i>B.ambifaria</i> AMMD]
gj 115359494	19.1	glucose-6-phosphate 1-dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 115350736	19	short-chain dehydrogenase/reductase SDR [<i>B.ambifaria</i> AMMD]
gj 115350794	19	aspartate carbamoyltransferase catalytic subunit [<i>B.ambifaria</i> AMMD]
gj 115351225	18.9	protein of unknown function UPF0044 [<i>B.ambifaria</i> AMMD]
gj 115361080	18.9	Cupin 2, conserved barrel domain protein [<i>B.ambifaria</i> AMMD]
gj 134295111	18.9	hypothetical protein Bcep1808_1000 [<i>B.vietnamiensis</i> G4]
gj 115352913	18.8	ribose-phosphate pyrophosphokinase [<i>B.ambifaria</i> AMMD]
gj 134293152	18.8	aspartate-semialdehyde dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 30314432	18.8	outer membrane lipoprotein precursor [<i>B.cepacia</i>]
gj 115350562	18.7	3-dehydroquinone dehydratase [<i>B.ambifaria</i> AMMD]
gj 115350251	18.6	histone family protein nucleoid-structuring protein H-NS [<i>B.ambifaria</i> AMMD]
gj 115350873	18.6	hypothetical protein Bamb_0819 [<i>B.ambifaria</i> AMMD]
gj 115352333	18.6	NADH-quinone oxidoreductase, F subunit [<i>B.ambifaria</i> AMMD]
gj 134297208	18.6	Glutathione S-transferase, N-terminal domain [<i>B.vietnamiensis</i> G4]
gj 189349671	18.6	multiple sugar transport system substrate-binding protein [<i>B.multivorans</i> ATCC 17616]
gj 115352706	18.5	transporter, hydrophobe/amphiphile efflux-1 (HAE1) family [<i>B.ambifaria</i> AMMD]
gj 115352963	18.5	phosphoglycerate mutase 1 family [<i>B.ambifaria</i> AMMD]
gj 115357988	18.5	hypothetical protein Bamb_3236 [<i>B.ambifaria</i> AMMD]
gj 115361015	18.5	Antibiotic biosynthesis monooxygenase [<i>B.ambifaria</i> AMMD]
gj 134295544	18.5	alpha/beta hydrolase fold [<i>B.vietnamiensis</i> G4]
gj 134292157	18.4	hypothetical protein Bcep1808_3440 [<i>B.vietnamiensis</i> G4]
gj 115351394	18.3	chorismate synthase [<i>B.ambifaria</i> AMMD]
gj 134294402	18.3	6-phosphogluconate dehydrogenase, NAD-binding [<i>B.vietnamiensis</i> G4]
gj 115351400	18.2	short chain dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 134296331	18.2	Rhodanese domain protein [<i>B.vietnamiensis</i> G4]
gj 115350127	18.1	adenylate cyclase [<i>B.ambifaria</i> AMMD]
gj 116688997	18.1	thioredoxin reductase [<i>B.cenocepacia</i> HI2424]
gj 115350376	18	Glutamate synthase (ferredoxin) [<i>B.ambifaria</i> AMMD]
gj 115350908	18	hypothetical protein Bamb_0854 [<i>B.ambifaria</i> AMMD]
gj 115351008	18	oligoribonuclease [<i>B.ambifaria</i> AMMD]
gj 115350672	17.9	Ferritin, Dps family protein [<i>B.ambifaria</i> AMMD]

gj 115352182	17.9	NlpBDapX family lipoprotein [<i>B.ambifaria</i> AMMD]
gj 115352330	17.9	NADH dehydrogenase subunit I [<i>B.ambifaria</i> AMMD]
gj 134297107	17.9	DSBA oxidoreductase [<i>B.vietnamiensis</i> G4]
gj 134297180	17.9	protein of unknown function DUF799 [<i>B.vietnamiensis</i> G4]
gj 115350556	17.8	PfkB domain protein [<i>B.ambifaria</i> AMMD]
gj 115358058	17.8	hypothetical protein Bamb_3307 [<i>B.ambifaria</i> AMMD]
gj 134293528	17.8	short-chain dehydrogenase/reductase SDR [<i>B.vietnamiensis</i> G4]
gj 115352095	17.7	surface antigen (D15) [<i>B.ambifaria</i> AMMD]
gj 115353043	17.7	sulfate ABC transporter, periplasmic sulfate-binding protein [<i>B.ambifaria</i> AMMD]
gj 115353214	17.7	aspartyl/glutamyl-tRNA amidotransferase subunit A [<i>B.ambifaria</i> AMMD]
gj 116688821	17.7	hypothetical protein Bcen2424_0798 [<i>B.cenocepacia</i> HI2424]
gj 134297024	17.7	lipoprotein, YaeC family [<i>B.vietnamiensis</i> G4]
gj 189351726	17.7	putative ABC transport system auxiliary component [<i>B.multivorans</i> ATCC 17616]
gj 115359945	17.6	Glycine hydroxymethyltransferase [<i>B.ambifaria</i> AMMD]
gj 134292961	17.6	3-mercaptopyruvate sulfurtransferase [<i>B.vietnamiensis</i> G4]
gj 134294799	17.6	catalase/peroxidase HPI [<i>B.vietnamiensis</i> G4]
gj 134295186	17.6	HAD-superfamily hydrolase, subfamily IIB [<i>B.vietnamiensis</i> G4]
gj 115352354	17.5	acetolactate synthase 3 catalytic subunit [<i>B.ambifaria</i> AMMD]
gj 134295009	17.5	GCN5-related N-acetyltransferase [<i>B.vietnamiensis</i> G4]
gj 116688659	17.4	Glyoxalase/bleomycin resistance protein/dioxygenase [<i>B.cenocepacia</i> HI2424]
gj 116690768	17.4	CHRD domain containing protein [<i>B.cenocepacia</i> HI2424]
gj 134290920	17.4	Carboxymethylenebutenolidase [<i>B.vietnamiensis</i> G4]
gj 134291741	17.4	putative signal-transduction protein with CBS domains [<i>B.vietnamiensis</i> G4]
gj 134296606	17.4	bifunctional uroporphyrinogen-III synthetase/uroporphyrin-III C-methyltransferase [<i>B.vietnamiensis</i> G4]
gj 134296900	17.4	dimethyladenosine transferase [<i>B.vietnamiensis</i> G4]
gj 189349974	17.4	polar amino acid transport system substrate-binding protein [<i>B.multivorans</i> ATCC 17616]
gj 115350526	17.3	preprotein translocase subunit SecA [<i>B.ambifaria</i> AMMD]
gj 115358648	17.3	electron transfer flavoprotein, alpha subunit [<i>B.ambifaria</i> AMMD]
gj 134294116	17.3	Rhodanese domain protein [<i>B.vietnamiensis</i> G4]
gj 134294637	17.3	ubiquinone biosynthesis protein COQ7-like protein [<i>B.vietnamiensis</i> G4]
gj 115351789	17.2	phosphoribosyltransferase [<i>B.ambifaria</i> AMMD]
gj 115353016	17.2	Methionine synthase [<i>B.ambifaria</i> AMMD]
gj 134295046	17.2	aminopeptidase N [<i>B.vietnamiensis</i> G4]
gj 134295358	17.2	protein of unknown function UPF0044 [<i>B.vietnamiensis</i> G4]
gj 115350288	17.1	N-acetyl-gamma-glutamyl-phosphate reductase [<i>B.ambifaria</i> AMMD]
gj 134297054	17	glutathione synthetase [<i>B.vietnamiensis</i> G4]
gj 115351049	16.9	50S ribosomal protein L32 [<i>B.ambifaria</i> AMMD]
gj 115351059	16.9	sigma E regulatory protein, MucB/RseB [<i>B.ambifaria</i> AMMD]
gj 115351596	16.9	hypothetical protein Bamb_1543 [<i>B.ambifaria</i> AMMD]
gj 115351873	16.9	hypothetical protein Bamb_1822 [<i>B.ambifaria</i> AMMD]
gj 115352289	16.9	MotA/ToIQ/ExbB proton channel [<i>B.ambifaria</i> AMMD]

gj 115358290	16.9	Acetyl-CoA hydrolase [<i>B.ambifaria</i> AMMD]
gj 115358928	16.9	short chain dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 134291069	16.9	3-hydroxybutyryl-CoA dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 134294615	16.9	protein-disulfide isomerase-like protein [<i>B.vietnamiensis</i> G4]
gj 134297114	16.9	short chain dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 115350554	16.8	hypothetical protein Bamb_0500 [<i>B.ambifaria</i> AMMD]
gj 115350822	16.8	UDP-glucose 4-epimerase [<i>B.ambifaria</i> AMMD]
gj 116688684	16.8	2-dehydro-3-deoxyphosphogluconate aldolase/4-hydroxy-2-oxoglutarate aldolase [<i>B.cenocepacia</i> HI2424]
gj 134291087	16.8	hypothetical protein Bcep1808_5663 [<i>B.vietnamiensis</i> G4]
gj 134295840	16.8	Pyrrolo-quinoline quinone [<i>B.vietnamiensis</i> G4]
gj 83721066	16.8	unnamed protein product; Similar to Hcp protein [<i>B.thailandensis</i> E264]
gj 115350495	16.7	ribulose-phosphate 3-epimerase [<i>B.ambifaria</i> AMMD]
gj 115352224	16.7	dihydroliipoamide acetyltransferase [<i>B.ambifaria</i> AMMD]
gj 11559479	16.7	GspG [<i>B.cepacia</i>]
gj 115358099	16.6	tryptophan synthase subunit beta [<i>B.ambifaria</i> AMMD]
gj 116686801	16.6	Linocin_M18 bacteriocin protein [<i>B.cenocepacia</i> HI2424]
gj 134293101	16.6	hypothetical protein Bcep1808_4406 [<i>B.vietnamiensis</i> G4]
gj 115350390	16.5	VacJ family lipoprotein [<i>B.ambifaria</i> AMMD]
gj 115352434	16.5	heat shock protein 90 [<i>B.ambifaria</i> AMMD]
gj 115352519	16.5	porphobilinogen deaminase [<i>B.ambifaria</i> AMMD]
gj 116686888	16.5	HAD-superfamily hydrolase, subfamily IA, variant 3 [<i>B.cenocepacia</i> HI2424]
gj 134291073	16.5	alpha/beta hydrolase fold [<i>B.vietnamiensis</i> G4]
gj 115351422	16.4	uncharacterized lipoprotein-like protein [<i>B.ambifaria</i> AMMD]
gj 116692036	16.4	type II citrate synthase [<i>B.cenocepacia</i> HI2424]
gj 134296618	16.4	OmpA/MotB domain protein [<i>B.vietnamiensis</i> G4]
gj 115350542	16.3	hypothetical protein Bamb_0488 [<i>B.ambifaria</i> AMMD]
gj 115360436	16.3	bifunctional aconitate hydratase 2/2-methylisocitrate dehydratase [<i>B.ambifaria</i> AMMD]
gj 134295690	16.3	periplasmic binding protein [<i>B.vietnamiensis</i> G4]
gj 134297038	16.3	cytochrome c oxidase, subunit II [<i>B.vietnamiensis</i> G4]
gj 134294758	16.2	30S ribosomal protein S9 [<i>B.vietnamiensis</i> G4]
gj 115358569	16.1	acetoacetyl-CoA reductase [<i>B.ambifaria</i> AMMD]
gj 116688829	16.1	serine hydroxymethyltransferase [<i>B.cenocepacia</i> HI2424]
gj 134283475	16.1	cmaB protein [<i>B.pseudomallei</i> 305]
gj 115352571	16	sulfate adenyltransferase, large subunit [<i>B.ambifaria</i> AMMD]
gj 115352822	16	acetyl-CoA acetyltransferase [<i>B.ambifaria</i> AMMD]
gj 115352572	15.9	sulfate adenyltransferase subunit 2 [<i>B.ambifaria</i> AMMD]
gj 115358572	15.9	Alcohol dehydrogenase GroES domain protein [<i>B.ambifaria</i> AMMD]
gj 134293105	15.9	succinate dehydrogenase, cytochrome b subunit [<i>B.vietnamiensis</i> G4]
gj 134294671	15.9	GTPase ObgE [<i>B.vietnamiensis</i> G4]
gj 189350815	15.9	small subunit ribosomal protein S2 [<i>B.multivorans</i> ATCC 17616]
gj 115351403	15.8	3-oxoacid CoA-transferase, A subunit [<i>B.ambifaria</i> AMMD]
gj 115352107	15.8	peptide deformylase [<i>B.ambifaria</i> AMMD]
gj 115352387	15.8	iojap-like protein [<i>B.ambifaria</i> AMMD]
gj 115352905	15.8	HPr kinase/phosphorylase [<i>B.ambifaria</i> AMMD]

gj 115359149	15.8	hypothetical protein Bamb_4401 [<i>B.ambifaria</i> AMMD]
gj 115359511	15.8	cationic amino acid ABC transporter, periplasmic binding protein [<i>B.ambifaria</i> AMMD]
gj 134296393	15.8	uncharacterized protein probably involved in high-affinity Fe ²⁺ transport [<i>B.vietnamiensis</i> G4]
gj 115351503	15.7	ribose-5-phosphate isomerase A [<i>B.ambifaria</i> AMMD]
gj 115352090	15.7	lipid-A-disaccharide synthase [<i>B.ambifaria</i> AMMD]
gj 134292091	15.7	hypothetical protein Bcep1808_3374 [<i>B.vietnamiensis</i> G4]
gj 134294620	15.7	Indole-3-glycerol-phosphate synthase [<i>B.vietnamiensis</i> G4]
gj 134294723	15.7	transketolase [<i>B.vietnamiensis</i> G4]
gj 83720445	15.7	pyridine nucleotide-disulphide oxidoreductase, class II [<i>B.thailandensis</i> E264]
gj 115351590	15.6	amino acid adenylation domain [<i>B.ambifaria</i> AMMD]
gj 115352541	15.6	aspartate alpha-decarboxylase [<i>B.ambifaria</i> AMMD]
gj 116690813	15.6	putative PTS IIA-like nitrogen-regulatory protein PtsN [<i>B.cenocepacia</i> HI2424]
gj 134292523	15.6	VacJ family lipoprotein [<i>B.vietnamiensis</i> G4]
gj 115358006	15.5	GatB/Yqey domain protein [<i>B.ambifaria</i> AMMD]
gj 115358454	15.5	short-chain dehydrogenase/reductase SDR [<i>B.ambifaria</i> AMMD]
gj 116688305	15.5	histone family protein nucleoid-structuring protein H-NS [<i>B.cenocepacia</i> HI2424]
gj 134295160	15.5	protease Do [<i>B.vietnamiensis</i> G4]
gj 134295363	15.5	phosphate ABC transporter, periplasmic phosphate-binding protein [<i>B.vietnamiensis</i> G4]
gj 160699072	15.5	glutamine synthetase, type I [<i>B.mallei</i> ATCC 10399]
gj 115351339	15.4	alanyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gj 115358450	15.4	ThiJ/Pfpl domain protein [<i>B.ambifaria</i> AMMD]
gj 134295561	15.4	uncharacterized lipoprotein-like protein [<i>B.vietnamiensis</i> G4]
gj 38640325	15.4	Bcep22gp17 [B.phage Bcep22]
gj 115352589	15.3	bifunctional N-succinyldiaminopimelate-aminotransferase/acetylornithine transaminase protein [<i>B.ambifaria</i> AMMD]
gj 115358000	15.3	Polyprenyl synthetase [<i>B.ambifaria</i> AMMD]
gj 116690730	15.3	PpiC-type peptidyl-prolyl cis-trans isomerase [<i>B.cenocepacia</i> HI2424]
gj 134293840	15.3	chlorinating enzyme [<i>B.vietnamiensis</i> G4]
gj 189351807	15.3	large subunit ribosomal protein L7/L12 [<i>B.multivorans</i> ATCC 17616]
gj 116688978	15.2	transcription antitermination protein NusB [<i>B.cenocepacia</i> HI2424]
gj 134296596	15.2	outer membrane protein (porin)-like protein [<i>B.vietnamiensis</i> G4]
gj 134291887	15.1	hypothetical protein Bcep1808_6504 [<i>B.vietnamiensis</i> G4]
gj 115352092	15	(3R)-hydroxymyristoyl-(acyl carrier protein) dehydratase [<i>B.ambifaria</i> AMMD]
gj 115352349	15	2-isopropylmalate synthase [<i>B.ambifaria</i> AMMD]
gj 967990	15	outer membrane protein [<i>B.cepacia</i>]
gj 115359090	14.9	peptidase M20 [<i>B.ambifaria</i> AMMD]
gj 134292440	14.9	phospho-2-dehydro-3-deoxyheptonate aldolase [<i>B.vietnamiensis</i> G4]
gj 115350352	14.8	delta-aminolevulinic acid dehydratase [<i>B.ambifaria</i> AMMD]
gj 115351444	14.8	dihydrolipoamide acetyltransferase [<i>B.ambifaria</i> AMMD]

gj 115352213	14.8	scaffold protein [<i>B.ambifaria</i> AMMD]
gj 115352531	14.8	deoxycytidine triphosphate deaminase [<i>B.ambifaria</i> AMMD]
gj 115352652	14.8	ornithine carbamoyltransferase [<i>B.ambifaria</i> AMMD]
gj 134294522	14.8	imidazole glycerol phosphate synthase subunit HisF [<i>B.vietnamiensis</i> G4]
gj 134295476	14.8	hypothetical protein Bcep1808_1367 [<i>B.vietnamiensis</i> G4]
gj 134296332	14.8	molybdopterin binding domain [<i>B.vietnamiensis</i> G4]
gj 134296524	14.8	transaldolase B [<i>B.vietnamiensis</i> G4]
gj 189349244	14.8	probable translation factor [<i>B.multivorans</i> ATCC 17616]
gj 115352073	14.7	bifunctional GMP synthase/glutamine amidotransferase protein [<i>B.ambifaria</i> AMMD]
gj 115359379	14.7	transcriptional regulator, TetR family [<i>B.ambifaria</i> AMMD]
gj 115359705	14.7	NADPH-dependent FMN reductase [<i>B.ambifaria</i> AMMD]
gj 116692461	14.7	UspA domain protein [<i>B.cenocepacia</i> HI2424]
gj 134295695	14.7	TonB-dependent siderophore receptor [<i>B.vietnamiensis</i> G4]
gj 134295696	14.7	folate-dependent phosphoribosylglycinamide formyltransferase PurN-like protein [<i>B.vietnamiensis</i> G4]
gj 115358321	14.6	extracellular solute-binding protein, family 5 [<i>B.ambifaria</i> AMMD]
gj 116689772	14.6	extracellular solute-binding protein, family 1 [<i>B.cenocepacia</i> HI2424]
gj 116690629	14.6	efflux transporter, RND family, MFP subunit [<i>B.cenocepacia</i> HI2424]
gj 134293676	14.6	Alcohol dehydrogenase GroES domain protein [<i>B.vietnamiensis</i> G4]
gj 134295825	14.6	competence lipoprotein ComL [<i>B.vietnamiensis</i> G4]
gj 134296697	14.6	isocitrate dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 134296996	14.6	outer membrane lipoprotein LolB [<i>B.vietnamiensis</i> G4]
gj 115350157	14.5	uroporphyrinogen decarboxylase [<i>B.ambifaria</i> AMMD]
gj 115351790	14.5	adenylosuccinate synthetase [<i>B.ambifaria</i> AMMD]
gj 116693798	14.5	protein of unknown function DUF1348 [<i>B.cenocepacia</i> HI2424]
gj 189348254	14.5	alcohol dehydrogenase [<i>B.multivorans</i> ATCC 17616]
gj 189351773	14.5	small subunit ribosomal protein S4 [<i>B.multivorans</i> ATCC 17616]
gj 3769514	14.5	unknown [<i>B.cepacia</i>]
gj 11230853	14.4	ornibactin receptor precursor [<i>B.cepacia</i>]
gj 116688624	14.4	Redoxin domain protein [<i>B.cenocepacia</i> HI2424]
gj 115351028	14.3	CreA family protein [<i>B.ambifaria</i> AMMD]
gj 115360735	14.3	Carboxymethylenebutenolidase [<i>B.ambifaria</i> AMMD]
gj 134294508	14.3	VacJ family lipoprotein [<i>B.vietnamiensis</i> G4]
gj 115352899	14.2	protein of unknown function DUF1239 [<i>B.ambifaria</i> AMMD]
gj 115353186	14.2	cobalamin synthesis protein, P47K [<i>B.ambifaria</i> AMMD]
gj 134294810	14.2	globin [<i>B.vietnamiensis</i> G4]
gj 134294995	14.2	extracellular solute-binding protein, family 1 [<i>B.vietnamiensis</i> G4]
gj 115350366	14.1	shikimate kinase [<i>B.ambifaria</i> AMMD]
gj 115351585	14.1	ABC transporter related [<i>B.ambifaria</i> AMMD]
gj 116688302	14.1	Extracellular ligand-binding receptor [<i>B.cenocepacia</i> HI2424]
gj 134295171	14.1	elongation factor P [<i>B.vietnamiensis</i> G4]
gj 134296009	14.1	band 7 protein [<i>B.vietnamiensis</i> G4]
gj 160699730	14.1	phasin family protein [<i>B.mallei</i> ATCC 10399]
gj 115351060	14	protease Do [<i>B.ambifaria</i> AMMD]
gj 115351500	14	ribonuclease R [<i>B.ambifaria</i> AMMD]

gij115353128	14	Glutathione S-transferase, N-terminal domain [<i>B.ambifaria</i> AMMD]
gij116689062	14	OmpA/MotB domain protein [<i>B.cenocepacia</i> HI2424]
gij116689575	14	extracellular solute-binding protein, family 3 [<i>B.cenocepacia</i> HI2424]
gij134294332	14	putative CheW protein [<i>B.vietnamiensis</i> G4]
gij115351810	13.9	peptidase M23B [<i>B.ambifaria</i> AMMD]
gij134293587	13.8	UspA domain protein [<i>B.vietnamiensis</i> G4]
gij115351800	13.7	histidyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gij115360171	13.7	O-acetylhomoserine/O-acetylserine sulfhydrylase [<i>B.ambifaria</i> AMMD]
gij116690641	13.7	phospholipase/lecithinase/hemolysin-like protein [<i>B.cenocepacia</i> HI2424]
gij134295970	13.7	alkyl hydroperoxide reductase/ Thiol specific antioxidant/ Mal allergen [<i>B.vietnamiensis</i> G4]
gij134296645	13.7	leucyl aminopeptidase [<i>B.vietnamiensis</i> G4]
gij134297013	13.7	cytochrome bd ubiquinol oxidase, subunit I [<i>B.vietnamiensis</i> G4]
gij115350260	13.6	transport-associated [<i>B.ambifaria</i> AMMD]
gij115352351	13.6	phosphatidylserine decarboxylase [<i>B.ambifaria</i> AMMD]
gij115352607	13.6	Fmu (Sun) domain protein [<i>B.ambifaria</i> AMMD]
gij115352834	13.6	ubiquinone/menaquinone biosynthesis methyltransferase [<i>B.ambifaria</i> AMMD]
gij134292879	13.6	aldehyde dehydrogenase [<i>B.vietnamiensis</i> G4]
gij134296276	13.6	dihydrodipicolinate synthase [<i>B.vietnamiensis</i> G4]
gij115358509	13.5	aromatic amino acid aminotransferase [<i>B.ambifaria</i> AMMD]
gij134294113	13.5	Lysine decarboxylase [<i>B.vietnamiensis</i> G4]
gij91781830	13.5	Putative acyltransferase [<i>B.xenovorans</i> LB400]
gij115350914	13.4	DNA-directed RNA polymerase subunit omega [<i>B.ambifaria</i> AMMD]
gij115352534	13.4	OmpA/MotB domain protein [<i>B.ambifaria</i> AMMD]
gij115360253	13.4	aldehyde dehydrogenase [<i>B.ambifaria</i> AMMD]
gij134294833	13.4	phosphoribosylaminoimidazole synthetase [<i>B.vietnamiensis</i> G4]
gij134295841	13.4	hypothetical protein Bcep1808_1737 [<i>B.vietnamiensis</i> G4]
gij134296641	13.4	protein of unknown function DUF541 [<i>B.vietnamiensis</i> G4]
gij134296951	13.4	peptidase M48, Ste24p [<i>B.vietnamiensis</i> G4]
gij189351857	13.4	putative histone-like nucleoid-structuring protein H-NS [<i>B.multivorans</i> ATCC 17616]
gij115350218	13.3	histone family protein nucleoid-structuring protein H-NS [<i>B.ambifaria</i> AMMD]
gij115351777	13.3	polyhydroxyalkonate synthesis repressor, PhaR [<i>B.ambifaria</i> AMMD]
gij115351989	13.3	large conductance mechanosensitive channel protein [<i>B.ambifaria</i> AMMD]
gij10697312	13.2	aspartate transcarbamoylase [<i>B.cepacia</i>]
gij115352191	13.2	transferase hexapeptide repeat containing protein [<i>B.ambifaria</i> AMMD]
gij189353572	13.1	lysine decarboxylase [<i>B.multivorans</i> ATCC 17616]
gij115352185	13	tryptophanyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gij134294741	13	2-dehydro-3-deoxyphosphogluconate aldolase/4-hydroxy-2-oxoglutarate aldolase [<i>B.vietnamiensis</i> G4]
gij134296016	13	UDP-N-acetylglucosamine acyltransferase [<i>B.vietnamiensis</i> G4]

gj 189351863	13	carboxymethylenebutenolidase [<i>B.multivorans</i> ATCC 17616]
gj 115350684	12.9	phospho-2-dehydro-3-heoxyheptonate aldolase [<i>B.ambifaria</i> AMMD]
gj 115359889	12.9	protein of unknown function UPF0047 [<i>B.ambifaria</i> AMMD]
gj 115360277	12.9	hypothetical protein Bamb_5534 [<i>B.ambifaria</i> AMMD]
gj 116690511	12.9	bifunctional N-succinyldiaminopimelate-aminotransferase/acetylornithine transaminase protein [<i>B.cenocepacia</i> HI2424]
gj 134295343	12.9	Glutathione S-transferase, N-terminal domain [<i>B.vietnamiensis</i> G4]
gj 189351255	12.9	acetylornithine aminotransferase [<i>B.multivorans</i> ATCC 17616]
gj 189352365	12.9	putative lipoprotein [<i>B.multivorans</i> ATCC 17616]
gj 115353190	12.8	ATP-dependent protease ATP-binding subunit [<i>B.ambifaria</i> AMMD]
gj 116688989	12.8	6-phosphogluconolactonase [<i>B.cenocepacia</i> HI2424]
gj 116690741	12.8	acyl-CoA dehydrogenase domain protein [<i>B.cenocepacia</i> HI2424]
gj 134294340	12.8	hypothetical protein Bcep1808_0224 [<i>B.vietnamiensis</i> G4]
gj 134294589	12.8	Water Stress and Hypersensitive response [<i>B.vietnamiensis</i> G4]
gj 134295605	12.8	RNA chaperone Hfq [<i>B.vietnamiensis</i> G4]
gj 115350319	12.7	elongation factor G [<i>B.ambifaria</i> AMMD]
gj 115351591	12.7	amino acid adenylation domain [<i>B.ambifaria</i> AMMD]
gj 115352196	12.7	2-dehydro-3-deoxyphosphooctonate aldolase [<i>B.ambifaria</i> AMMD]
gj 115358526	12.7	UspA domain protein [<i>B.ambifaria</i> AMMD]
gj 134292438	12.7	PRC-barrel domain protein [<i>B.vietnamiensis</i> G4]
gj 134293649	12.7	Succinate-semialdehyde dehydrogenase (NAD(P)(+)) [<i>B.vietnamiensis</i> G4]
gj 134296733	12.7	adenylate kinase [<i>B.vietnamiensis</i> G4]
gj 134295884	12.6	threonine synthase [<i>B.vietnamiensis</i> G4]
gj 134295896	12.6	PhoH family protein [<i>B.vietnamiensis</i> G4]
gj 134296445	12.6	phosphatidylserine decarboxylase [<i>B.vietnamiensis</i> G4]
gj 134296972	12.6	Cobyrinic acid a,c-diamide synthase [<i>B.vietnamiensis</i> G4]
gj 115352576	12.5	transcriptional regulator CysB-like protein [<i>B.ambifaria</i> AMMD]
gj 115350202	12.4	FAD dependent oxidoreductase [<i>B.ambifaria</i> AMMD]
gj 115351853	12.4	alkyl hydroperoxide reductase/ Thiol specific antioxidant/ Mal allergen [<i>B.ambifaria</i> AMMD]
gj 115352414	12.4	polyhydroxyalkanoate depolymerase, intracellular [<i>B.ambifaria</i> AMMD]
gj 115358872	12.4	putative signal-transduction protein with CBS domains [<i>B.ambifaria</i> AMMD]
gj 134292688	12.4	outer membrane efflux protein [<i>B.vietnamiensis</i> G4]
gj 115351020	12.3	fumarate hydratase [<i>B.ambifaria</i> AMMD]
gj 116688703	12.3	30S ribosomal protein S9 [<i>B.cenocepacia</i> HI2424]
gj 116689322	12.3	phosphate ABC transporter, periplasmic phosphate-binding protein [<i>B.cenocepacia</i> HI2424]
gj 134293450	12.3	methyl-accepting chemotaxis sensory transducer [<i>B.vietnamiensis</i> G4]
gj 134294625	12.3	ribulose-phosphate 3-epimerase [<i>B.vietnamiensis</i> G4]
gj 134297260	12.3	protein of unknown function DUF484 [<i>B.vietnamiensis</i> G4]
gj 134297272	12.3	nucleoid occlusion protein [<i>B.vietnamiensis</i> G4]
gj 115352139	12.2	glutamyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]

gj 115352332	12.2	NADH dehydrogenase subunit G [<i>B.ambifaria</i> AMMD]
gj 134295820	12.2	acetyl-CoA reductase [<i>B.vietnamiensis</i> G4]
gj 189350933	12.2	phasin family protein [<i>B.multivorans</i> ATCC 17616]
gj 116690548	12.1	translation elongation factor G [<i>B.cenocepacia</i> HI2424]
gj 116691123	12.1	exodeoxyribonuclease III Xth [<i>B.cenocepacia</i> HI2424]
gj 134294242	12.1	Prephenate dehydratase [<i>B.vietnamiensis</i> G4]
gj 134294354	12.1	Extracellular ligand-binding receptor [<i>B.vietnamiensis</i> G4]
gj 115350574	12	2-dehydro-3-deoxy-6-phosphogalactonate aldolase [<i>B.ambifaria</i> AMMD]
gj 115350604	12	Rare lipoprotein B [<i>B.ambifaria</i> AMMD]
gj 115351124	12	Redoxin domain protein [<i>B.ambifaria</i> AMMD]
gj 115351959	12	PpiC-type peptidyl-prolyl cis-trans isomerase [<i>B.ambifaria</i> AMMD]
gj 115358093	12	isopropylmalate isomerase small subunit [<i>B.ambifaria</i> AMMD]
gj 134291545	12	transport-associated [<i>B.vietnamiensis</i> G4]
gj 134294488	12	3-dehydroquininate synthase [<i>B.vietnamiensis</i> G4]
gj 134295159	12	sigma E regulatory protein, MucB/RseB [<i>B.vietnamiensis</i> G4]
gj 115351071	11.9	elongation factor P [<i>B.ambifaria</i> AMMD]
gj 134291893	11.9	predicted glycosyl transferase [<i>B.vietnamiensis</i> G4]
gj 134292239	11.9	aldehyde dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 134293103	11.9	malate dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 134294279	11.9	cytochrome c, class I [<i>B.vietnamiensis</i> G4]
gj 134294490	11.9	glycerol-3-phosphate transporter periplasmic binding protein [<i>B.vietnamiensis</i> G4]
gj 115350188	11.8	glycine cleavage system H protein [<i>B.ambifaria</i> AMMD]
gj 116689500	11.8	50S ribosomal protein L20 [<i>B.cenocepacia</i> HI2424]
gj 116689532	11.8	dihydrolipoamide dehydrogenase [<i>B.cenocepacia</i> HI2424]
gj 134294571	11.8	Tetratricopeptide TPR_2 repeat protein [<i>B.vietnamiensis</i> G4]
gj 134295573	11.8	transcription elongation factor NusA [<i>B.vietnamiensis</i> G4]
gj 134295811	11.8	Alcohol dehydrogenase, zinc-binding domain protein [<i>B.vietnamiensis</i> G4]
gj 189349587	11.8	pyrimidine operon attenuation protein / uracil phosphoribosyltransferase [<i>B.multivorans</i> ATCC 17616]
gj 115350795	11.7	dihydroorotase [<i>B.ambifaria</i> AMMD]
gj 116689940	11.7	lipolytic enzyme, G-D-S-L family [<i>B.cenocepacia</i> HI2424]
gj 134295071	11.7	OmpA/MotB domain protein [<i>B.vietnamiensis</i> G4]
gj 171320747	11.7	peptidyl-prolyl cis-trans isomerase cyclophilin type [<i>B.ambifaria</i> MEX-5]
gj 115350541	11.6	gamma-glutamyl kinase [<i>B.ambifaria</i> AMMD]
gj 115350652	11.6	putative RNA methylase [<i>B.ambifaria</i> AMMD]
gj 116688550	11.6	protein-disulfide isomerase-like protein [<i>B.cenocepacia</i> HI2424]
gj 116689064	11.6	hypothetical protein Bcen2424_1041 [<i>B.cenocepacia</i> HI2424]
gj 134294633	11.6	enoyl-CoA hydratase [<i>B.vietnamiensis</i> G4]
gj 134294875	11.6	transcriptional regulator, LysR family [<i>B.vietnamiensis</i> G4]
gj 134296848	11.6	general secretion pathway protein H [<i>B.vietnamiensis</i> G4]
gj 115350931	11.5	fructose-1,6-bisphosphatase [<i>B.ambifaria</i> AMMD]
gj 115358107	11.5	O-succinylhomoserine sulphydrylase [<i>B.ambifaria</i> AMMD]
gj 115359283	11.5	ABC transporter related [<i>B.ambifaria</i> AMMD]
gj 115359332	11.5	enoyl-CoA hydratase [<i>B.ambifaria</i> AMMD]
gj 116691933	11.5	Peptidylprolyl isomerase [<i>B.cenocepacia</i> HI2424]
gj 116692966	11.5	phosphate acetyltransferase [<i>B.cenocepacia</i> HI2424]

gj 134293107	11.5	succinate dehydrogenase flavoprotein subunit [<i>B.vietnamiensis</i> G4]
gj 134296995	11.5	Tetratricopeptide TPR_2 repeat protein [<i>B.vietnamiensis</i> G4]
gj 14572550	11.5	RuvB [<i>B.cepacia</i>]
gj 115350932	11.4	aminopeptidase N [<i>B.ambifaria</i> AMMD]
gj 115359750	11.4	aldo/keto reductase [<i>B.ambifaria</i> AMMD]
gj 116689854	11.4	50S ribosomal protein L31 type B [<i>B.cenocepacia</i> HI2424]
gj 134292441	11.4	protein of unknown function DUF480 [<i>B.vietnamiensis</i> G4]
gj 134293978	11.4	short chain dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 115357980	11.3	molybdenum ABC transporter, periplasmic molybdate-binding protein [<i>B.ambifaria</i> AMMD]
gj 189350468	11.3	pyoverdine biosynthesis gene [<i>B.multivorans</i> ATCC 17616]
gj 115350560	11.2	acetyl-CoA carboxylase biotin carboxylase subunit [<i>B.ambifaria</i> AMMD]
gj 115352901	11.2	ABC transporter related [<i>B.ambifaria</i> AMMD]
gj 115360930	11.2	UTP-glucose-1-phosphate uridylyltransferase [<i>B.ambifaria</i> AMMD]
gj 115350156	11.1	Prephenate dehydratase [<i>B.ambifaria</i> AMMD]
gj 115350880	11.1	6-phosphogluconolactonase [<i>B.ambifaria</i> AMMD]
gj 115351593	11.1	lysine/ornithine N-monooxygenase [<i>B.ambifaria</i> AMMD]
gj 116689821	11.1	phosphoribosyltransferase [<i>B.cenocepacia</i> HI2424]
gj 134295351	11.1	Lytic transglycosylase, catalytic [<i>B.vietnamiensis</i> G4]
gj 134295216	11	extracellular solute-binding protein, family 3 [<i>B.vietnamiensis</i> G4]
gj 189350947	11	glutamine synthetase [<i>B.multivorans</i> ATCC 17616]
gj 115350212	10.9	flagellin [<i>B.ambifaria</i> AMMD]
gj 115350296	10.9	transcriptional regulator, AsnC family [<i>B.ambifaria</i> AMMD]
gj 115350576	10.9	periplasmic binding protein/LacI transcriptional regulator [<i>B.ambifaria</i> AMMD]
gj 115352341	10.9	Alcohol dehydrogenase, zinc-binding domain protein [<i>B.ambifaria</i> AMMD]
gj 115352610	10.9	isoleucyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gj 116691099	10.9	S-adenosylmethionine synthetase [<i>B.cenocepacia</i> HI2424]
gj 134295574	10.9	translation initiation factor IF-2 [<i>B.vietnamiensis</i> G4]
gj 134296988	10.9	sigma 54 modulation protein/ribosomal protein S30EA [<i>B.vietnamiensis</i> G4]
gj 115350475	10.8	putative ABC transporter ATP-binding protein [<i>B.ambifaria</i> AMMD]
gj 115353021	10.8	DSBA oxidoreductase [<i>B.ambifaria</i> AMMD]
gj 115360696	10.8	Alpha/beta hydrolase fold-3 domain protein [<i>B.ambifaria</i> AMMD]
gj 134294938	10.8	Phosphomannomutase [<i>B.vietnamiensis</i> G4]
gj 134297058	10.8	protein of unknown function DUF526 [<i>B.vietnamiensis</i> G4]
gj 115350985	10.7	cysteine synthase B [<i>B.ambifaria</i> AMMD]
gj 115352406	10.7	DEAD/DEAH box helicase domain protein [<i>B.ambifaria</i> AMMD]
gj 116689892	10.7	50S ribosomal protein L9 [<i>B.cenocepacia</i> HI2424]
gj 134295673	10.7	hypothetical protein Bcep1808_1566 [<i>B.vietnamiensis</i> G4]
gj 115361243	10.6	pyruvate dehydrogenase complex dihydrolipoamide acetyltransferase [<i>B.ambifaria</i> AMMD]
gj 116692259	10.6	extracellular solute-binding protein, family 5 [<i>B.cenocepacia</i> HI2424]
gj 134295025	10.6	ribonuclease PH [<i>B.vietnamiensis</i> G4]
gj 189349830	10.6	conserved hypothetical protein [<i>B.multivorans</i> ATCC 17616]

gj 115352699	10.5	periplasmic solute binding protein [<i>B.ambifaria</i> AMMD]
gj 115352720	10.5	Sua5/YciO/YrdC/YwIC family protein [<i>B.ambifaria</i> AMMD]
gj 115353236	10.5	DNA topoisomerase III [<i>B.ambifaria</i> AMMD]
gj 116688692	10.4	extracellular solute-binding protein, family 3 [<i>B.cenocepacia</i> HI2424]
gj 116689933	10.4	PpiC-type peptidyl-prolyl cis-trans isomerase [<i>B.cenocepacia</i> HI2424]
gj 116691351	10.4	pyruvate dehydrogenase complex dihydrolipoamide acetyltransferase [<i>B.cenocepacia</i> HI2424]
gj 134296965	10.4	hypothetical protein Bcep1808_2874 [<i>B.vietnamiensis</i> G4]
gj 160696587	10.4	6,7-dimethyl-8-ribityllumazine synthase [<i>B.mallei</i> ATCC 10399]
gj 171316822	10.4	Substrate-binding region of ABC-type glycine betaine transport system [<i>B.ambifaria</i> MEX-5]
gj 115350540	10.3	GTPase ObgE [<i>B.ambifaria</i> AMMD]
gj 115351584	10.3	Taurine catabolism dioxygenase TauD/TfdA [<i>B.ambifaria</i> AMMD]
gj 115351920	10.3	hypothetical protein Bamb_1869 [<i>B.ambifaria</i> AMMD]
gj 115352091	10.3	UDP-N-acetylglucosamine acyltransferase [<i>B.ambifaria</i> AMMD]
gj 115353226	10.3	response regulator receiver protein [<i>B.ambifaria</i> AMMD]
gj 115359175	10.3	ABC transporter related [<i>B.ambifaria</i> AMMD]
gj 116688661	10.3	transketolase [<i>B.cenocepacia</i> HI2424]
gj 116688778	10.3	3-methyl-2-oxobutanoate hydroxymethyltransferase [<i>B.cenocepacia</i> HI2424]
gj 134292690	10.3	protein of unknown function DUF477 [<i>B.vietnamiensis</i> G4]
gj 134295095	10.3	electron transfer flavoprotein, alpha subunit [<i>B.vietnamiensis</i> G4]
gj 115350921	10.2	chorismate mutase [<i>B.ambifaria</i> AMMD]
gj 115351953	10.2	phosphoribosylformylglycinamide synthase [<i>B.ambifaria</i> AMMD]
gj 134294696	10.2	UDP-N-acetylmuramate [<i>B.vietnamiensis</i> G4]
gj 134297109	10.2	extracellular solute-binding protein, family 5 [<i>B.vietnamiensis</i> G4]
gj 134297182	10.2	Curli production assembly/transport component CsgG [<i>B.vietnamiensis</i> G4]
gj 189349113	10.2	outer membrane porin [<i>B.multivorans</i> ATCC 17616]
gj 115352409	10.1	phosphoribosylglycinamide formyltransferase 2 [<i>B.ambifaria</i> AMMD]
gj 116688507	10.1	OmpA/MotB domain protein [<i>B.cenocepacia</i> HI2424]
gj 134292529	10.1	malate synthase G [<i>B.vietnamiensis</i> G4]
gj 134296656	10.1	Extracellular ligand-binding receptor [<i>B.vietnamiensis</i> G4]
gj 115350700	10	chaperone protein DnaK [<i>B.ambifaria</i> AMMD]
gj 115350898	10	septum site-determining protein MinD [<i>B.ambifaria</i> AMMD]
gj 115350986	10	lytic murein transglycosylase B [<i>B.ambifaria</i> AMMD]
gj 115351125	10	NAD-dependent epimerase/dehydratase [<i>B.ambifaria</i> AMMD]
gj 115351879	10	hydrolase, TatD family [<i>B.ambifaria</i> AMMD]
gj 115352183	10	dihydrodipicolinate synthase [<i>B.ambifaria</i> AMMD]
gj 115352608	10	phosphoribosylglycinamide formyltransferase [<i>B.ambifaria</i> AMMD]
gj 115360410	10	Linocin_M18 bacteriocin protein [<i>B.ambifaria</i> AMMD]
gj 134290885	10	Glycerone kinase [<i>B.vietnamiensis</i> G4]
gj 134295835	10	band 7 protein [<i>B.vietnamiensis</i> G4]
gj 115351834	9.9	hypothetical protein Bamb_1783 [<i>B.ambifaria</i> AMMD]
gj 115351945	9.9	3-hydroxyacyl-CoA dehydrogenase, NAD-binding [<i>B.ambifaria</i> AMMD]

gij115352404	9.9	hypothetical protein Bamb_2353 [<i>B.ambifaria</i> AMMD]
gij115360742	9.9	parB-like partition proteins [<i>B.ambifaria</i> AMMD]
gij134292965	9.9	Polyprenyl synthetase [<i>B.vietnamiensis</i> G4]
gij115350143	9.8	parB-like partition proteins [<i>B.ambifaria</i> AMMD]
gij115350586	9.8	malic enzyme [<i>B.ambifaria</i> AMMD]
gij115351780	9.8	poly(R)-hydroxyalkanoic acid synthase, class I [<i>B.ambifaria</i> AMMD]
gij115352575	9.8	nitrite/sulfite reductase, hemoprotein beta-component, ferredoxin domain protein [<i>B.ambifaria</i> AMMD]
gij115352597	9.8	4-hydroxy-3-methylbut-2-enyl diphosphate reductase [<i>B.ambifaria</i> AMMD]
gij115353130	9.8	Lytic transglycosylase, catalytic [<i>B.ambifaria</i> AMMD]
gij115358285	9.8	phosphoenolpyruvate phosphomutase [<i>B.ambifaria</i> AMMD]
gij134296395	9.8	aromatic amino acid aminotransferase [<i>B.vietnamiensis</i> G4]
gij189349395	9.8	polar amino acid transport system substrate-binding protein [<i>B.multivorans</i> ATCC 17616]
gij189350331	9.8	2-oxoglutarate dehydrogenase E2 component [<i>B.multivorans</i> ATCC 17616]
gij115350153	9.7	F0F1 ATP synthase subunit beta [<i>B.ambifaria</i> AMMD]
gij115350631	9.7	tyrosyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gij115353212	9.7	protein of unknown function DUF344 [<i>B.ambifaria</i> AMMD]
gij116689911	9.7	ankyrin repeat protein [<i>B.cenocepacia</i> HI2424]
gij134291718	9.7	UspA domain protein [<i>B.vietnamiensis</i> G4]
gij115350187	9.6	glycine dehydrogenase [<i>B.ambifaria</i> AMMD]
gij115351040	9.6	ribonuclease, Rne/Rng family [<i>B.ambifaria</i> AMMD]
gij134294613	9.6	translation-associated GTPase [<i>B.vietnamiensis</i> G4]
gij160695740	9.6	ATP synthase F0, B subunit [<i>B.mallei</i> ATCC 10399]
gij115351387	9.5	FAD linked oxidase domain protein [<i>B.ambifaria</i> AMMD]
gij115352228	9.5	Methenyltetrahydrofolate cyclohydrolase [<i>B.ambifaria</i> AMMD]
gij115359542	9.5	short-chain alcohol dehydrogenase [<i>B.ambifaria</i> AMMD]
gij115360459	9.5	2-keto-4-pentenoate hydratase-like protein [<i>B.ambifaria</i> AMMD]
gij134295691	9.5	cyclic peptide transporter [<i>B.vietnamiensis</i> G4]
gij134296512	9.5	polyhydroxyalkanoate depolymerase, intracellular [<i>B.vietnamiensis</i> G4]
gij115358575	9.4	Alcohol dehydrogenase, zinc-binding domain protein [<i>B.ambifaria</i> AMMD]
gij115359036	9.4	ABC transporter related [<i>B.ambifaria</i> AMMD]
gij134294033	9.4	acetate kinase [<i>B.vietnamiensis</i> G4]
gij134294740	9.4	phosphogluconate dehydratase [<i>B.vietnamiensis</i> G4]
gij115350692	9.3	inorganic polyphosphate/ATP-NAD kinase [<i>B.ambifaria</i> AMMD]
gij115352162	9.3	acetyl-CoA carboxylase carboxyltransferase subunit alpha [<i>B.ambifaria</i> AMMD]
gij189349358	9.3	NADP-dependent oxaloacetate-decarboxylating malate dehydrogenase [<i>B.multivorans</i> ATCC 17616]
gij115350828	9.2	hypothetical protein Bamb_0774 [<i>B.ambifaria</i> AMMD]
gij115353008	9.2	hypothetical protein Bamb_2957 [<i>B.ambifaria</i> AMMD]
gij115350587	9.1	guanine-specific ribonuclease N1 and T1 [<i>B.ambifaria</i> AMMD]
gij115358001	9.1	1-deoxy-D-xylulose-5-phosphate synthase [<i>B.ambifaria</i> AMMD]
gij116691352	9.1	dihydroliipoamide dehydrogenase [<i>B.cenocepacia</i> HI2424]
gij134292443	9.1	short chain dehydrogenase [<i>B.vietnamiensis</i> G4]

gi 134294446	9.1	50S ribosomal protein L2 [<i>B.vietnamiensis</i> G4]
gi 134295878	9.1	ATPase AAA-2 domain protein [<i>B.vietnamiensis</i> G4]
gi 134296384	9.1	MotA/TolQ/ExbB proton channel [<i>B.vietnamiensis</i> G4]
gi 189349826	9.1	3-demethylubiquinone-9 3-methyltransferase [<i>B.multivorans</i> ATCC 17616]
gi 116688627	9	acetyl-CoA carboxylase biotin carboxylase subunit [<i>B.cenocepacia</i> HI2424]
gi 134283218	9	tetratricopeptide repeat protein [<i>B.pseudomallei</i> 305]
gi 134296427	9	NADH dehydrogenase subunit G [<i>B.vietnamiensis</i> G4]
gi 134296894	9	glycyl-tRNA synthetase subunit beta [<i>B.vietnamiensis</i> G4]
gi 83717742	9	Prokaryotic protein of unknown function (DUF849) family [<i>B.thailandensis</i> E264]
gi 115350670	8.9	catalase/peroxidase HPI [<i>B.ambifaria</i> AMMD]
gi 115351374	8.9	valyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gi 134294789	8.9	Ycel family protein [<i>B.vietnamiensis</i> G4]
gi 134295336	8.9	malonate decarboxylase gamma subunit [<i>B.vietnamiensis</i> G4]
gi 134297267	8.9	two component transcriptional regulator, Fis family [<i>B.vietnamiensis</i> G4]
gi 115351088	8.8	electron transport protein SCO1/SenC [<i>B.ambifaria</i> AMMD]
gi 115352801	8.8	transporter-associated region [<i>B.ambifaria</i> AMMD]
gi 115359692	8.8	toluene tolerance family protein [<i>B.ambifaria</i> AMMD]
gi 134293372	8.8	aminotransferase, class V [<i>B.vietnamiensis</i> G4]
gi 134294243	8.8	uroporphyrinogen decarboxylase [<i>B.vietnamiensis</i> G4]
gi 134297210	8.8	Lytic transglycosylase, catalytic [<i>B.vietnamiensis</i> G4]
gi 189350441	8.8	LysR family Cys regulon transcriptional activator [<i>B.multivorans</i> ATCC 17616]
gi 116688379	8.7	50S ribosomal protein L16 [<i>B.cenocepacia</i> HI2424]
gi 116690575	8.7	nucleotide-binding protein [<i>B.cenocepacia</i> HI2424]
gi 115350519	8.6	D-alanine--D-alanine ligase [<i>B.ambifaria</i> AMMD]
gi 116690037	8.6	ribosome recycling factor [<i>B.cenocepacia</i> HI2424]
gi 134293901	8.6	hydroxylamine reductase [<i>B.vietnamiensis</i> G4]
gi 134296025	8.6	ribosome recycling factor [<i>B.vietnamiensis</i> G4]
gi 162389967	8.6	glutamate synthase subunit beta [<i>B.ambifaria</i> AMMD]
gi 115350483	8.5	translation-associated GTPase [<i>B.ambifaria</i> AMMD]
gi 115351417	8.5	phenylalanyl-tRNA synthetase subunit beta [<i>B.ambifaria</i> AMMD]
gi 115352518	8.5	bifunctional uroporphyrinogen-III synthetase/uroporphyrin-III C-methyltransferase [<i>B.ambifaria</i> AMMD]
gi 115358103	8.5	FoC bifunctional protein [<i>B.ambifaria</i> AMMD]
gi 115360278	8.5	protein of unknown function DUF883, ElaB [<i>B.ambifaria</i> AMMD]
gi 116693234	8.5	Substrate-binding region of ABC-type glycine betaine transport system [<i>B.cenocepacia</i> HI2424]
gi 134291238	8.5	HAD-superfamily hydrolase, subfamily IA, variant 3 [<i>B.vietnamiensis</i> G4]
gi 171321248	8.5	chlorinating enzyme [<i>B.ambifaria</i> MEX-5]
gi 115350484	8.4	hypothetical protein Bamb_0430 [<i>B.ambifaria</i> AMMD]
gi 115351779	8.4	acetyl-CoA acetyltransferase [<i>B.ambifaria</i> AMMD]
gi 116688916	8.4	Phosphomannomutase [<i>B.cenocepacia</i> HI2424]
gi 116688991	8.4	extracellular solute-binding protein, family 1 [<i>B.cenocepacia</i> HI2424]
gi 189350477	8.4	lysine N6-hydroxylase [<i>B.multivorans</i> ATCC 17616]

gj 115351609	8.3	cobalamin biosynthesis protein CobW [<i>B.ambifaria</i> AMMD]
gj 115351760	8.3	ABC transporter related [<i>B.ambifaria</i> AMMD]
gj 115352533	8.3	Cobyrinic acid a,c-diamide synthase [<i>B.ambifaria</i> AMMD]
gj 116690911	8.3	Serine-type D-Ala-D-Ala carboxypeptidase [<i>B.cenocepacia</i> HI2424]
gj 115350970	8.2	DNA gyrase subunit A [<i>B.ambifaria</i> AMMD]
gj 116690230	8.2	uncharacterized protein probably involved in high-affinity Fe ²⁺ transport [<i>B.cenocepacia</i> HI2424]
gj 116691903	8.2	hypothetical protein Bcen2424_3806 [<i>B.cenocepacia</i> HI2424]
gj 116692965	8.2	acetate kinase [<i>B.cenocepacia</i> HI2424]
gj 134294496	8.2	protein of unknown function DUF1568 [<i>B.vietnamiensis</i> G4]
gj 134296335	8.2	hypothetical protein Bcep1808_2233 [<i>B.vietnamiensis</i> G4]
gj 189351543	8.2	ribose-phosphate pyrophosphokinase [<i>B.multivorans</i> ATCC 17616]
gj 116689635	8.1	putative serine protein kinase, PrkA [<i>B.cenocepacia</i> HI2424]
gj 116689842	8.1	peptidase M23B [<i>B.cenocepacia</i> HI2424]
gj 134293680	8.1	phosphoesterase [<i>B.vietnamiensis</i> G4]
gj 134295949	8.1	phosphoribosylformylglycinamide synthase [<i>B.vietnamiensis</i> G4]
gj 161648989	8.1	bifunctional aconitate hydratase 2/2-methylisocitrate dehydratase [<i>B.vietnamiensis</i> G4]
gj 189348076	8.1	branched-chain amino acid transport system substrate-binding protein [<i>B.multivorans</i> ATCC 17616]
gj 115351412	8	threonyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gj 116690124	8	transferase hexapeptide repeat containing protein [<i>B.cenocepacia</i> HI2424]
gj 134293745	8	aldo/keto reductase [<i>B.vietnamiensis</i> G4]
gj 134294440	8	elongation factor G [<i>B.vietnamiensis</i> G4]
gj 134296487	8	hypothetical protein Bcep1808_2388 [<i>B.vietnamiensis</i> G4]
gj 115351442	7.9	GTP-binding protein TypA [<i>B.ambifaria</i> AMMD]
gj 115352668	7.9	Cyclopropane-fatty-acyl-phospholipid synthase [<i>B.ambifaria</i> AMMD]
gj 115357996	7.9	Carboxymethylenebutenolidase [<i>B.ambifaria</i> AMMD]
gj 116690819	7.9	Tetratricopeptide TPR_2 repeat protein [<i>B.cenocepacia</i> HI2424]
gj 83721398	7.9	isocitrate dehydrogenase [<i>B.thailandensis</i> E264]
gj 115352994	7.8	Serine-type D-Ala-D-Ala carboxypeptidase [<i>B.ambifaria</i> AMMD]
gj 116690492	7.8	sulfate adenyltransferase subunit 2 [<i>B.cenocepacia</i> HI2424]
gj 134294441	7.8	elongation factor Tu [<i>B.vietnamiensis</i> G4]
gj 134294998	7.8	ABC transporter related [<i>B.vietnamiensis</i> G4]
gj 160695879	7.8	translation elongation factor Tu [<i>B.mallei</i> ATCC 10399]
gj 115351634	7.7	Glutathione S-transferase, N-terminal domain [<i>B.ambifaria</i> AMMD]
gj 115351852	7.7	polysaccharide deacetylase [<i>B.ambifaria</i> AMMD]
gj 115353079	7.7	UDP-N-acetylglucosamine pyrophosphorylase [<i>B.ambifaria</i> AMMD]
gj 115359630	7.7	PRC-barrel domain protein [<i>B.ambifaria</i> AMMD]
gj 116693806	7.7	3-hydroxybutyrate dehydrogenase [<i>B.cenocepacia</i> HI2424]
gj 134296402	7.7	hypothetical protein Bcep1808_2303 [<i>B.vietnamiensis</i> G4]
gj 171319800	7.7	OmpW family protein [<i>B.ambifaria</i> MEX-5]
gj 189348250	7.7	conserved hypothetical protein [<i>B.multivorans</i> ATCC 17616]
gj 189349365	7.7	glyceraldehyde 3-phosphate dehydrogenase [<i>B.multivorans</i>

		ATCC 17616]
gj 115351220	7.6	carbamoyl phosphate synthase small subunit [<i>B.ambifaria</i> AMMD]
gj 115353019	7.6	arginyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gj 134294686	7.6	17 kDa surface antigen [<i>B.vietnamiensis</i> G4]
gj 171315488	7.6	conserved hypothetical protein [<i>B.ambifaria</i> MEX-5]
gj 83655611	7.6	DNA gyrase subunit B [<i>B.cepacia</i>]
gj 115351375	7.5	UvrD/REP helicase [<i>B.ambifaria</i> AMMD]
gj 115352097	7.5	1-deoxy-D-xylulose 5-phosphate reductoisomerase [<i>B.ambifaria</i> AMMD]
gj 115353020	7.5	Sporulation domain protein [<i>B.ambifaria</i> AMMD]
gj 115358719	7.5	protein of unknown function DUF1255 [<i>B.ambifaria</i> AMMD]
gj 134295026	7.5	hypothetical protein Bcep1808_0915 [<i>B.vietnamiensis</i> G4]
gj 134296323	7.5	Oligopeptidase A [<i>B.vietnamiensis</i> G4]
gj 37654966	7.5	flagellin [<i>B.vietnamiensis</i>]
gj 115360893	7.4	hypothetical protein Bamb_6152 [<i>B.ambifaria</i> AMMD]
gj 116688419	7.4	glycerol-3-phosphate transporter periplasmic binding protein [<i>B.cenocepacia</i> HI2424]
gj 116688912	7.4	UDP-glucose 4-epimerase [<i>B.cenocepacia</i> HI2424]
gj 116689825	7.4	HflC protein [<i>B.cenocepacia</i> HI2424]
gj 134294934	7.4	UDP-glucose 4-epimerase [<i>B.vietnamiensis</i> G4]
gj 134296917	7.4	3-hydroxyacyl-CoA dehydrogenase, NAD-binding [<i>B.vietnamiensis</i> G4]
gj 189349328	7.4	adenosine kinase [<i>B.multivorans</i> ATCC 17616]
gj 115351564	7.3	hypothetical protein Bamb_1511 [<i>B.ambifaria</i> AMMD]
gj 115352122	7.3	alanine racemase [<i>B.ambifaria</i> AMMD]
gj 134291841	7.3	Alcohol dehydrogenase, zinc-binding domain protein [<i>B.vietnamiensis</i> G4]
gj 134292247	7.3	peptidase S10, serine carboxypeptidase [<i>B.vietnamiensis</i> G4]
gj 134294413	7.3	malic enzyme [<i>B.vietnamiensis</i> G4]
gj 134295901	7.3	50S ribosomal protein L9 [<i>B.vietnamiensis</i> G4]
gj 134296406	7.3	peptidase M14, carboxypeptidase A [<i>B.vietnamiensis</i> G4]
gj 134296605	7.3	HemY domain protein [<i>B.vietnamiensis</i> G4]
gj 171318106	7.3	ribosomal protein S2 [<i>B.ambifaria</i> MEX-5]
gj 189349608	7.3	phosphomannomutase [<i>B.multivorans</i> ATCC 17616]
gj 189351043	7.3	polyribonucleotide nucleotidyltransferase [<i>B.multivorans</i> ATCC 17616]
gj 83719268	7.3	acetyl-CoA reductase [<i>B.thailandensis</i> E264]
gj 115350400	7.2	imidazoleglycerol-phosphate dehydratase [<i>B.ambifaria</i> AMMD]
gj 115350681	7.2	globin [<i>B.ambifaria</i> AMMD]
gj 115351858	7.2	replicative DNA helicase [<i>B.ambifaria</i> AMMD]
gj 115352798	7.2	PhoH family protein [<i>B.ambifaria</i> AMMD]
gj 115358567	7.2	UspA domain protein [<i>B.ambifaria</i> AMMD]
gj 115359753	7.2	transcriptional regulator, Crp/Fnr family [<i>B.ambifaria</i> AMMD]
gj 116690135	7.2	lipoprotein releasing system, transmembrane protein, LolC/E family [<i>B.cenocepacia</i> HI2424]
gj 116690544	7.2	isocitrate dehydrogenase [<i>B.cenocepacia</i> HI2424]
gj 134294357	7.2	histone family protein nucleoid-structuring protein H-NS [<i>B.vietnamiensis</i> G4]
gj 134294509	7.2	toluene tolerance family protein [<i>B.vietnamiensis</i> G4]
gj 171321751	7.2	branched-chain amino acid aminotransferase [<i>B.ambifaria</i> MEX-

		5]
gj 189351140	7.2	YjgF family putative translation initiation inhibitor [<i>B.multivorans</i> ATCC 17616]
gj 189351606	7.2	carboxy-terminal protease [<i>B.multivorans</i> ATCC 17616]
gj 115352093	7.1	UDP-3-O-[3-hydroxymyristoyl] glucosamine N-acyltransferase [<i>B.ambifaria</i> AMMD]
gj 116688482	7.1	extracellular solute-binding protein, family 3 [<i>B.cenocepacia</i> HI2424]
gj 116688695	7.1	ABC transporter related [<i>B.cenocepacia</i> HI2424]
gj 116689202	7.1	bifunctional N-succinyldiaminopimelate-aminotransferase/acetylornithine transaminase protein [<i>B.cenocepacia</i> HI2424]
gj 134293099	7.1	aconitate hydratase [<i>B.vietnamiensis</i> G4]
gj 134294561	7.1	extracellular solute-binding protein, family 3 [<i>B.vietnamiensis</i> G4]
gj 134295581	7.1	GTP-binding protein TypA [<i>B.vietnamiensis</i> G4]
gj 53718021	7.1	thiol:disulfide interchange protein [<i>B.pseudomallei</i> K96243]
gj 91779712	7.1	thiol:disulfide interchange protein [<i>B.pseudomallei</i> K96243]
gj 91779712	7.1	3-hydroxybutyryl-CoA dehydrogenase [<i>B.xenovorans</i> LB400]
gj 115353232	7	methionyl-tRNA formyltransferase [<i>B.ambifaria</i> AMMD]
gj 116690430	7	outer membrane protein (porin)-like protein [<i>B.cenocepacia</i> HI2424]
gj 116691071	7	Ferredoxin--NADP(+) reductase [<i>B.cenocepacia</i> HI2424]
gj 134296431	7	NADH dehydrogenase subunit C [<i>B.vietnamiensis</i> G4]
gj 134297221	7	extracellular solute-binding protein, family 5 [<i>B.vietnamiensis</i> G4]
gj 115350968	6.9	3-demethylubiquinone-9 3-methyltransferase [<i>B.ambifaria</i> AMMD]
gj 134293406	6.9	extracellular solute-binding protein, family 5 [<i>B.vietnamiensis</i> G4]
gj 134295074	6.9	phosphoserine aminotransferase [<i>B.vietnamiensis</i> G4]
gj 134296619	6.9	methionyl-tRNA synthetase [<i>B.vietnamiensis</i> G4]
gj 134296902	6.9	PpiC-type peptidyl-prolyl cis-trans isomerase [<i>B.vietnamiensis</i> G4]
gj 134297080	6.9	Serine-type D-Ala-D-Ala carboxypeptidase [<i>B.vietnamiensis</i> G4]
gj 171322437	6.9	extracellular solute-binding protein family 5 [<i>B.ambifaria</i> MEX-5]
gj 189350585	6.9	acetoacetyl-CoA reductase [<i>B.multivorans</i> ATCC 17616]
gj 189352638	6.9	citrate synthase [<i>B.multivorans</i> ATCC 17616]
gj 83720868	6.9	Phage integrase [<i>B.thailandensis</i> E264]
gj 115350493	6.8	anthranilate synthase component I [<i>B.ambifaria</i> AMMD]
gj 115351843	6.8	homoserine dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 115360899	6.8	acetyl-CoA synthetase [<i>B.ambifaria</i> AMMD]
gj 116688387	6.8	50S ribosomal protein L6 [<i>B.cenocepacia</i> HI2424]
gj 116692004	6.8	6-phosphogluconate dehydrogenase [<i>B.cenocepacia</i> HI2424]
gj 134295094	6.8	electron transfer flavoprotein beta-subunit [<i>B.vietnamiensis</i> G4]
gj 83716265	6.8	thiotemplate mechanism natural product synthetase [<i>B.thailandensis</i> E264]
gj 134294455	6.7	50S ribosomal protein L5 [<i>B.vietnamiensis</i> G4]
gj 134294850	6.7	serine hydroxymethyltransferase [<i>B.vietnamiensis</i> G4]
gj 134296376	6.7	acetyl-CoA synthetase [<i>B.vietnamiensis</i> G4]
gj 115351565	6.6	hypothetical protein Bamb_1512 [<i>B.ambifaria</i> AMMD]
gj 134294675	6.6	prolyl-tRNA synthetase [<i>B.vietnamiensis</i> G4]
gj 134295936	6.6	polymorphic outer membrane protein [<i>B.vietnamiensis</i> G4]
gj 134296437	6.6	polynucleotide phosphorylase/polyadenylase [<i>B.vietnamiensis</i> G4]

gij160696872	6.6	GTP-binding protein YchF [<i>B.mallei</i> ATCC 10399]
gij115352527	6.5	argininosuccinate lyase [<i>B.ambifaria</i> AMMD]
gij115358985	6.5	Carbohydrate-selective porin OprB [<i>B.ambifaria</i> AMMD]
gij116690646	6.5	fructose-1,6-bisphosphate aldolase [<i>B.cenocepacia</i> HI2424]
gij134295557	6.5	hypothetical protein Bcep1808_1449 [<i>B.vietnamiensis</i> G4]
gij134295958	6.5	ATP-dependent Clp protease proteolytic subunit [<i>B.vietnamiensis</i> G4]
gij83718546	6.5	3-hydroxyacyl-CoA dehydrogenase/enoyl-CoA hydratase/isomerase family protein [<i>B.thailandensis</i> E264]
gij115350547	6.4	signal recognition particle protein [<i>B.ambifaria</i> AMMD]
gij115351062	6.4	GTP-binding protein LepA [<i>B.ambifaria</i> AMMD]
gij115351485	6.4	type II secretion system protein [<i>B.ambifaria</i> AMMD]
gij115358766	6.4	extracellular solute-binding protein, family 5 [<i>B.ambifaria</i> AMMD]
gij115359376	6.4	aldehyde dehydrogenase [<i>B.ambifaria</i> AMMD]
gij116688128	6.4	F0F1 ATP synthase subunit alpha [<i>B.cenocepacia</i> HI2424]
gij134296316	6.4	phasin family protein [<i>B.vietnamiensis</i> G4]
gij134296330	6.4	glutamine synthetase, type I [<i>B.vietnamiensis</i> G4]
gij91781310	6.4	ABC branched chain amino acid family transporter, periplasmic ligand binding protein [<i>B.xenovorans</i> LB400]
gij91784774	6.4	Microcin-processing peptidase [<i>B.xenovorans</i> LB400]
gij115350701	6.3	chaperone protein DnaJ [<i>B.ambifaria</i> AMMD]
gij115350737	6.3	serine hydroxymethyltransferase [<i>B.ambifaria</i> AMMD]
gij116690345	6.3	polyhydroxyalkanoate depolymerase, intracellular [<i>B.cenocepacia</i> HI2424]
gij167838193	6.3	50S ribosomal protein L15 [<i>B.thailandensis</i> MSMB43]
gij171320789	6.3	trigger factor [<i>B.ambifaria</i> MEX-5]
gij189350986	6.3	bacterioferritin [<i>B.multivorans</i> ATCC 17616]
gij189351783	6.3	large subunit ribosomal protein L6 [<i>B.multivorans</i> ATCC 17616]
gij115350342	6.2	preprotein translocase subunit SecY [<i>B.ambifaria</i> AMMD]
gij116686681	6.2	hypothetical protein Bcen2424_6303 [<i>B.cenocepacia</i> HI2424]
gij116693836	6.2	GTP cyclohydrolase I [<i>B.cenocepacia</i> HI2424]
gij134291711	6.2	beta-lactamase domain protein [<i>B.vietnamiensis</i> G4]
gij134296319	6.2	pyruvate dehydrogenase subunit E1 [<i>B.vietnamiensis</i> G4]
gij83716594	6.2	nonribosomal peptide synthetase, putative [<i>B.thailandensis</i> E264]
gij91781882	6.2	Putative heat shock protein [<i>B.xenovorans</i> LB400]
gij100917445	6.1	hypothetical protein Bmal10_03001194 [<i>B.mallei</i> 10399]
gij115350367	6.1	3-dehydroquinate synthase [<i>B.ambifaria</i> AMMD]
gij115352588	6.1	CDP-6-deoxy-delta-3,4-glucoseen reductase [<i>B.ambifaria</i> AMMD]
gij115359631	6.1	phospho-2-dehydro-3-deoxyheptonate aldolase [<i>B.ambifaria</i> AMMD]
gij116690050	6.1	hypothetical protein Bcen2424_2029 [<i>B.cenocepacia</i> HI2424]
gij134293374	6.1	phosphoenolpyruvate phosphomutase [<i>B.vietnamiensis</i> G4]
gij134295708	6.1	cobalamin biosynthesis protein CobW [<i>B.vietnamiensis</i> G4]
gij134296028	6.1	30S ribosomal protein S2 [<i>B.vietnamiensis</i> G4]
gij134296086	6.1	peptidyl-prolyl cis-trans isomerase, cyclophilin type [<i>B.vietnamiensis</i> G4]
gij160698114	6.1	argininosuccinate synthase [<i>B.mallei</i> ATCC 10399]
gij189349421	6.1	X-Pro aminopeptidase [<i>B.multivorans</i> ATCC 17616]
gij91778430	6.1	Methylpurine-DNA glycosylase (MPG) [<i>B.xenovorans</i> LB400]
gij116688683	6	phosphogluconate dehydratase [<i>B.cenocepacia</i> HI2424]

gij116690356	6	transaldolase B [<i>B.cenocepacia</i> HI2424]
gij134292355	6	short-chain alcohol dehydrogenase [<i>B.vietnamiensis</i> G4]
gij134294652	6	cell division protein FtsZ [<i>B.vietnamiensis</i> G4]
gij134294842	6	hypothetical protein Bcep1808_0730 [<i>B.vietnamiensis</i> G4]
gij115350606	5.9	gamma-glutamyl phosphate reductase [<i>B.ambifaria</i> AMMD]
gij115350624	5.9	dihydroorotase [<i>B.ambifaria</i> AMMD]
gij116690628	5.9	transporter, hydrophobe/amphiphile efflux-1 (HAE1) family [<i>B.cenocepacia</i> HI2424]
gij134295442	5.9	helix-turn-helix- domain containing protein, AraC type [<i>B.vietnamiensis</i> G4]
gij134296193	5.9	hypothetical protein Bcep1808_2091 [<i>B.vietnamiensis</i> G4]
gij115360195	5.8	argininosuccinate synthase [<i>B.ambifaria</i> AMMD]
gij116688126	5.8	F0F1 ATP synthase subunit B [<i>B.cenocepacia</i> HI2424]
gij116689113	5.8	fumarate hydratase [<i>B.cenocepacia</i> HI2424]
gij116690115	5.8	uncharacterized lipoprotein [<i>B.cenocepacia</i> HI2424]
gij134296317	5.8	dihydroliipoamide dehydrogenase [<i>B.vietnamiensis</i> G4]
gij189351795	5.8	large subunit ribosomal protein L2 [<i>B.multivorans</i> ATCC 17616]
gij189353570	5.8	deacylase [<i>B.multivorans</i> ATCC 17616]
gij115352928	5.7	cytochrome bd ubiquinol oxidase, subunit I [<i>B.ambifaria</i> AMMD]
gij115358106	5.7	amidophosphoribosyltransferase [<i>B.ambifaria</i> AMMD]
gij115358899	5.7	NADH:flavin oxidoreductase/NADH oxidase [<i>B.ambifaria</i> AMMD]
gij115360051	5.7	molecular chaperone-like protein [<i>B.ambifaria</i> AMMD]
gij116692007	5.7	peptidase M23B [<i>B.cenocepacia</i> HI2424]
gij134292973	5.7	RNA polymerase sigma factor RpoD [<i>B.vietnamiensis</i> G4]
gij134294473	5.7	delta-aminolevulinic acid dehydratase [<i>B.vietnamiensis</i> G4]
gij134294747	5.7	extracellular solute-binding protein, family 3 [<i>B.vietnamiensis</i> G4]
gij134295582	5.7	alpha-ketoglutarate decarboxylase [<i>B.vietnamiensis</i> G4]
gij189349477	5.7	molecular chaperone [<i>B.multivorans</i> ATCC 17616]
gij91780922	5.7	hypothetical protein Bxe_C0904 [<i>B.xenovorans</i> LB400]
gij115350875	5.6	riboflavin biosynthesis protein RibD [<i>B.ambifaria</i> AMMD]
gij115351466	5.6	RNA chaperone Hfq [<i>B.ambifaria</i> AMMD]
gij115352535	5.6	methionyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gij115360740	5.6	hypothetical protein Bamb_5999 [<i>B.ambifaria</i> AMMD]
gij116688994	5.6	ABC transporter related [<i>B.cenocepacia</i> HI2424]
gij134292217	5.6	TonB-dependent siderophore receptor [<i>B.vietnamiensis</i> G4]
gij134296443	5.6	2-isopropylmalate synthase [<i>B.vietnamiensis</i> G4]
gij134296805	5.6	periplasmic solute binding protein [<i>B.vietnamiensis</i> G4]
gij189348710	5.6	putative universal stress protein [<i>B.multivorans</i> ATCC 17616]
gij189351944	5.6	F-type H ⁺ -transporting ATPase delta chain [<i>B.multivorans</i> ATCC 17616]
gij91781739	5.6	single-strand DNA-binding protein [<i>B.xenovorans</i> LB400]
gij115359710	5.5	Cobyrinic acid a,c-diamide synthase [<i>B.ambifaria</i> AMMD]
gij116688362	5.5	50S ribosomal protein L10 [<i>B.cenocepacia</i> HI2424]
gij134294891	5.5	hypothetical protein Bcep1808_0779 [<i>B.vietnamiensis</i> G4]
gij134295459	5.5	cellulose synthase regulator protein [<i>B.vietnamiensis</i> G4]
gij134296504	5.5	DEAD/DEAH box helicase domain protein [<i>B.vietnamiensis</i> G4]
gij171318173	5.5	protein of unknown function DUF306 Meta and HslJ [<i>B.ambifaria</i> MEX-5]
gij189351808	5.5	large subunit ribosomal protein L10 [<i>B.multivorans</i> ATCC 17616]

gj 189352627	5.5	aconitate hydratase 1 [<i>B.multivorans</i> ATCC 17616]
gj 115359586	5.4	methylcitrate synthase [<i>B.ambifaria</i> AMMD]
gj 115359622	5.4	Radical SAM domain protein [<i>B.ambifaria</i> AMMD]
gj 116690722	5.4	glycyl-tRNA synthetase subunit beta [<i>B.cenocepacia</i> HI2424]
gj 160696708	5.4	aspartate carbamoyltransferase, 44 kDa non-catalytic chain [<i>B.mallei</i> ATCC 10399]
gj 70779411	5.4	TraL [<i>B.cepacia</i>]
gj 83717138	5.4	JamP [<i>B.thailandensis</i> E264]
gj 83720213	5.4	polyketide synthase [<i>B.thailandensis</i> E264]
gj 91777827	5.4	methylcitrate synthase [<i>B.xenovorans</i> LB400]
gj 115351370	5.3	Peptidoglycan-binding LysM [<i>B.ambifaria</i> AMMD]
gj 115352823	5.3	3-hydroxyacyl-CoA dehydrogenase, NAD-binding [<i>B.ambifaria</i> AMMD]
gj 134291080	5.3	nitrate reductase, alpha subunit [<i>B.vietnamiensis</i> G4]
gj 189349528	5.3	glycine hydroxymethyltransferase [<i>B.multivorans</i> ATCC 17616]
gj 115359463	5.2	beta-ketoadipyl CoA thiolase [<i>B.ambifaria</i> AMMD]
gj 116688714	5.2	bifunctional phosphoribosylaminoimidazolecarboxamide formyltransferase/IMP cyclohydrolase [<i>B.cenocepacia</i> HI2424]
gj 116690158	5.2	pyruvate dehydrogenase subunit E1 [<i>B.cenocepacia</i> HI2424]
gj 115352932	5.1	PTS system, N-acetylglucosamine-specific IIBC subunit [<i>B.ambifaria</i> AMMD]
gj 115358564	5.1	UspA domain protein [<i>B.ambifaria</i> AMMD]
gj 116690456	5.1	OmpA/MotB domain protein [<i>B.cenocepacia</i> HI2424]
gj 134292248	5.1	peptidase S8 and S53, subtilisin, kexin, sedolisin [<i>B.vietnamiensis</i> G4]
gj 134296608	5.1	phosphoenolpyruvate carboxylase [<i>B.vietnamiensis</i> G4]
gj 160698040	5.1	conserved hypothetical protein [<i>B.mallei</i> ATCC 10399]
gj 171320670	5.1	MltA-interacting MipA family protein [<i>B.ambifaria</i> MEX-5]
gj 171321799	5.1	peptidase M48 Ste24p [<i>B.ambifaria</i> MEX-5]
gj 115350368	5	deoxyguanosinetriphosphate triphosphohydrolase-like protein [<i>B.ambifaria</i> AMMD]
gj 115351699	5	hypothetical protein Bamb_1647 [<i>B.ambifaria</i> AMMD]
gj 115352805	5	glycyl-tRNA synthetase subunit beta [<i>B.ambifaria</i> AMMD]
gj 116688138	5	Extracellular ligand-binding receptor [<i>B.cenocepacia</i> HI2424]
gj 171318238	5	carbamoyl-phosphate synthase, small subunit [<i>B.ambifaria</i> MEX-5]
gj 189350717	5	peptidyl-prolyl cis-trans isomerase D [<i>B.multivorans</i> ATCC 17616]
gj 189351544	5	large subunit ribosomal protein L25 [<i>B.multivorans</i> ATCC 17616]
gj 83720370	5	pmbA protein [<i>B.thailandensis</i> E264]
gj 115360290	4.9	UDP-glucose 6-dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 115360454	4.9	nitrate reductase, beta subunit [<i>B.ambifaria</i> AMMD]
gj 116690740	4.9	3-hydroxyacyl-CoA dehydrogenase, NAD-binding [<i>B.cenocepacia</i> HI2424]
gj 116690337	4.8	DEAD/DEAH box helicase domain protein [<i>B.cenocepacia</i> HI2424]
gj 160697011	4.8	hippurate hydrolase [<i>B.mallei</i> ATCC 10399]
gj 171320062	4.8	transketolase [<i>B.ambifaria</i> MEX-5]
gj 2935157	4.8	type I flagellin [<i>B.cepacia</i>]
gj 115352311	4.7	peptidase M14, carboxypeptidase A [<i>B.ambifaria</i> AMMD]
gj 116690367	4.7	heat shock protein 90 [<i>B.cenocepacia</i> HI2424]

gj 171319798	4.7	extracellular solute-binding protein family 1 [<i>B.ambifaria</i> MEX-5]
gj 189349662	4.7	conserved hypothetical protein [<i>B.multivorans</i> ATCC 17616]
gj 189351516	4.7	phasin family protein [<i>B.multivorans</i> ATCC 17616]
gj 91778564	4.7	Alkanesulfonate monooxygenase [<i>B.xenovorans</i> LB400]
gj 115358074	4.6	Glutathione S-transferase, C-terminal domain [<i>B.ambifaria</i> AMMD]
gj 115360514	4.6	3-hydroxybutyryl-CoA dehydrogenase [<i>B.ambifaria</i> AMMD]
gj 116686519	4.6	porin, Gram-negative type [<i>B.cenocepacia</i> HI2424]
gj 116690937	4.6	arginyl-tRNA synthetase [<i>B.cenocepacia</i> HI2424]
gj 134291715	4.6	Poly-beta-hydroxybutyrate polymerase domain protein [<i>B.vietnamiensis</i> G4]
gj 134295552	4.6	phenylalanyl-tRNA synthetase subunit beta [<i>B.vietnamiensis</i> G4]
gj 134295886	4.6	aminotransferase AlaT [<i>B.vietnamiensis</i> G4]
gj 134296379	4.6	hydro-lyases, Fe-S type, tartrate/fumarate subfamily, alpha subunit [<i>B.vietnamiensis</i> G4]
gj 189351992	4.6	3-hydroxybutyrate dehydrogenase [<i>B.multivorans</i> ATCC 17616]
gj 189353469	4.6	3-oxoacyl-[acyl-carrier protein] reductase [<i>B.multivorans</i> ATCC 17616]
gj 115350320	4.5	elongation factor Tu [<i>B.ambifaria</i> AMMD]
gj 115350432	4.5	protein of unknown function DUF876 [<i>B.ambifaria</i> AMMD]
gj 115351218	4.5	Lytic transglycosylase, catalytic [<i>B.ambifaria</i> AMMD]
gj 116689850	4.5	transcription termination factor Rho [<i>B.cenocepacia</i> HI2424]
gj 134294256	4.5	Conserved TM helix repeat-containing protein [<i>B.vietnamiensis</i> G4]
gj 134296813	4.5	transporter, hydrophobe/amphiphile efflux-1 (HAE1) family [<i>B.vietnamiensis</i> G4]
gj 115350603	4.4	leucyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gj 115351149	4.4	amino acid permease-associated region [<i>B.ambifaria</i> AMMD]
gj 134290981	4.4	methyl-accepting chemotaxis sensory transducer [<i>B.vietnamiensis</i> G4]
gj 134294992	4.4	glucokinase [<i>B.vietnamiensis</i> G4]
gj 134295571	4.4	pseudouridine synthase [<i>B.vietnamiensis</i> G4]
gj 37654970	4.4	flagellin [<i>B.cepacia</i>]
gj 83720413	4.4	lipoprotein, putative [<i>B.thailandensis</i> E264]
gj 116690130	4.3	CTP synthetase [<i>B.cenocepacia</i> HI2424]
gj 134291725	4.3	Alcohol dehydrogenase GroES domain protein [<i>B.vietnamiensis</i> G4]
gj 189352477	4.3	1-deoxy-D-xylulose-5-phosphate synthase [<i>B.multivorans</i> ATCC 17616]
gj 115350664	4.2	preprotein translocase subunit SecD [<i>B.ambifaria</i> AMMD]
gj 115351432	4.2	Pseudouridine synthase, Rsu [<i>B.ambifaria</i> AMMD]
gj 116689522	4.2	translation initiation factor IF-2 [<i>B.cenocepacia</i> HI2424]
gj 116690446	4.2	phosphoenolpyruvate carboxylase [<i>B.cenocepacia</i> HI2424]
gj 116690486	4.2	leucyl aminopeptidase [<i>B.cenocepacia</i> HI2424]
gj 134295141	4.2	ribonuclease, Rne/Rng family [<i>B.vietnamiensis</i> G4]
gj 134297014	4.2	cytochrome d ubiquinol oxidase, subunit II [<i>B.vietnamiensis</i> G4]
gj 160697735	4.2	pmbA protein [<i>B.mallei</i> ATCC 10399]
gj 83720849	4.2	polyketide synthase [<i>B.thailandensis</i> E264]
gj 91778311	4.2	Putative senescence marker protein-30 (SMP-30) [<i>B.xenovorans</i> LB400]
gj 91781641	4.2	Na ⁺ /phosphate symporter [<i>B.xenovorans</i> LB400]

gij115350981	4.1	UDP-glucose 6-dehydrogenase [<i>B.ambifaria</i> AMMD]
gij115358008	4.1	RNA polymerase sigma factor RpoD [<i>B.ambifaria</i> AMMD]
gij115358570	4.1	Poly-beta-hydroxybutyrate polymerase domain protein [<i>B.ambifaria</i> AMMD]
gij134295511	4.1	valyl-tRNA synthetase [<i>B.vietnamiensis</i> G4]
gij171320557	4.1	lipoprotein, YaeC family [<i>B.ambifaria</i> MEX-5]
gij83716219	4.1	JamP [<i>B.thailandensis</i> E264]
gij83717801	4.1	hypothetical protein BTH_I12167 [<i>B.thailandensis</i> E264]
gij91785036	4.1	hypothetical protein Bxe_A0744 [<i>B.xenovorans</i> LB400]
gij115358875	4	porin, Gram-negative type [<i>B.ambifaria</i> AMMD]
gij116689945	4	trigger factor [<i>B.cenocepacia</i> HI2424]
gij160698338	4	tryptophanyl-tRNA synthetase [<i>B.mallei</i> ATCC 10399]
gij189350945	4	two-component system nitrogen regulation response regulator [<i>B.multivorans</i> ATCC 17616]
gij189351008	4	conserved hypothetical protein [<i>B.multivorans</i> ATCC 17616]
gij115352235	3.9	nitrogen metabolism transcriptional regulator, NtrC, Fis Family [<i>B.ambifaria</i> AMMD]
gij115352911	3.9	outer membrane lipoprotein LolB [<i>B.ambifaria</i> AMMD]
gij116690941	3.9	extracellular solute-binding protein, family 5 [<i>B.cenocepacia</i> HI2424]
gij116692829	3.9	porin, Gram-negative type [<i>B.cenocepacia</i> HI2424]
gij134292209	3.9	Carbamoyl-phosphate synthase L chain, ATP-binding [<i>B.vietnamiensis</i> G4]
gij134294432	3.9	50S ribosomal protein L1 [<i>B.vietnamiensis</i> G4]
gij134294682	3.9	ribonucleotide-diphosphate reductase subunit alpha [<i>B.vietnamiensis</i> G4]
gij134297017	3.9	phosphoenolpyruvate-protein phosphotransferase [<i>B.vietnamiensis</i> G4]
gij189350645	3.9	ATP-dependent Clp protease ATP-binding subunit [<i>B.multivorans</i> ATCC 17616]
gij115352620	3.8	multicopper oxidase, type 3 [<i>B.ambifaria</i> AMMD]
gij134291911	3.8	mannose-1-phosphate guanylyltransferase/mannose-6-phosphate isomerase [<i>B.vietnamiensis</i> G4]
gij134295671	3.8	putative serine protein kinase, PrkA [<i>B.vietnamiensis</i> G4]
gij134296834	3.8	phosphoglycerate kinase [<i>B.vietnamiensis</i> G4]
gij161621317	3.8	23S rRNA 5-methyluridine methyltransferase [<i>B.xenovorans</i> LB400]
gij115351335	3.7	glutaminyl-tRNA synthetase [<i>B.ambifaria</i> AMMD]
gij115359594	3.7	GCN5-related N-acetyltransferase [<i>B.ambifaria</i> AMMD]
gij116686584	3.7	RND efflux system, outer membrane lipoprotein, NodT family [<i>B.cenocepacia</i> HI2424]
gij116690040	3.7	30S ribosomal protein S2 [<i>B.cenocepacia</i> HI2424]
gij134294435	3.7	DNA-directed RNA polymerase subunit beta [<i>B.vietnamiensis</i> G4]
gij134295121	3.7	fumarate hydratase [<i>B.vietnamiensis</i> G4]
gij23820938	3.7	glucose dehydrogenase [<i>B.cepacia</i>]
gij115351586	3.6	iron-hydroxamate transporter permease subunit [<i>B.ambifaria</i> AMMD]
gij115358296	3.6	dihydroxy-acid dehydratase [<i>B.ambifaria</i> AMMD]
gij115358839	3.6	hydroxylamine reductase [<i>B.ambifaria</i> AMMD]
gij116688364	3.6	DNA-directed RNA polymerase subunit beta [<i>B.cenocepacia</i> HI2424]

gj 116690212	3.6	acetyl-CoA synthetase [<i>B.cenocepacia</i> HI2424]
gj 116690792	3.6	acetyl-CoA acetyltransferase [<i>B.cenocepacia</i> HI2424]
gj 134295959	3.6	trigger factor [<i>B.vietnamiensis</i> G4]
gj 160697439	3.6	acetate--CoA ligase [<i>B.mallei</i> ATCC 10399]
gj 189351369	3.6	RND efflux system outer membrane lipoprotein [<i>B.multivorans</i> ATCC 17616]
gj 189352150	3.6	aconitate hydratase 2 [<i>B.multivorans</i> ATCC 17616]
gj 116692055	3.5	tryptophan synthase subunit beta [<i>B.cenocepacia</i> HI2424]
gj 189348990	3.5	DNA topoisomerase type I [<i>B.multivorans</i> ATCC 17616]
gj 115360455	3.4	nitrate reductase, alpha subunit [<i>B.ambifaria</i> AMMD]
gj 134293725	3.4	nitrite reductase (NAD(P)H), large subunit [<i>B.vietnamiensis</i> G4]
gj 134294238	3.4	F0F1 ATP synthase subunit beta [<i>B.vietnamiensis</i> G4]
gj 189351801	3.4	elongation factor EF-G [<i>B.multivorans</i> ATCC 17616]
gj 91783628	3.4	Putative alcohol dehydrogenase cytochrome c subunit precursor, cytochrome c553 [<i>B.xenovorans</i> LB400]
gj 171317669	3.3	Lysine decarboxylase [<i>B.ambifaria</i> MEX-5]
gj 100914669	3.2	hypothetical protein Bmal10_03004030 [<i>B.mallei</i> 10399]
gj 115357994	3.2	DNA polymerase I [<i>B.ambifaria</i> AMMD]
gj 171319466	3.2	3-oxoacyl-(acyl-carrier-protein) synthase 2 [<i>B.ambifaria</i> MEX-5]
gj 171319605	3.2	extracellular solute-binding protein family 5 [<i>B.ambifaria</i> MEX-5]
gj 189351669	3.2	peptide/nickel transport system substrate-binding protein [<i>B.multivorans</i> ATCC 17616]
gj 53722877	3.2	LysR-family transcriptional regulator [<i>B.pseudomallei</i> K96243] LysR-family transcriptional regulator [<i>B.pseudomallei</i> K96243]
gj 115350225	3.1	CheA signal transduction histidine kinases [<i>B.ambifaria</i> AMMD]
gj 115351159	3.1	thiamine biosynthesis protein ThiC [<i>B.ambifaria</i> AMMD]
gj 134295353	3.1	carbamoyl phosphate synthase small subunit [<i>B.vietnamiensis</i> G4]
gj 134295836	3.1	HflK protein [<i>B.vietnamiensis</i> G4]
gj 171317285	3.1	extracellular solute-binding protein family 5 [<i>B.ambifaria</i> MEX-5]
gj 189351515	3.1	acetyl-CoA C-acetyltransferase [<i>B.multivorans</i> ATCC 17616]
gj 91784259	3.1	dihydrolipoamide acetyltransferase [<i>B.xenovorans</i> LB400]
gj 91784987	3.1	Translation elongation factor G [<i>B.xenovorans</i> LB400]
gj 116688760	3	globin [<i>B.cenocepacia</i> HI2424]
gj 116689038	3	aminopeptidase N [<i>B.cenocepacia</i> HI2424]
gj 116689431	3	alanyl-tRNA synthetase [<i>B.cenocepacia</i> HI2424]
gj 116689668	3	hypothetical protein Bcen2424_1646 [<i>B.cenocepacia</i> HI2424]
gj 116690264	3	NADH dehydrogenase subunit G [<i>B.cenocepacia</i> HI2424]
gj 134283477	3	metallo-beta-lactamase family protein [<i>B.pseudomallei</i> 305]
gj 134294734	3	leucyl-tRNA synthetase [<i>B.vietnamiensis</i> G4]
gj 33570923	3	elongation factor Tu [Bordetella pertussis Tohama I]
gj 116693052	2.9	Enoyl-CoA hydratase/isomerase [<i>B.cenocepacia</i> HI2424]
gj 134294272	2.9	glycine dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 134296685	2.9	isoleucyl-tRNA synthetase [<i>B.vietnamiensis</i> G4]
gj 171321878	2.9	transporter, hydrophobe/amphiphile efflux-1 (HAE1) family [<i>B.ambifaria</i> MEX-5]
gj 115351787	2.8	RNA binding S1 domain protein [<i>B.ambifaria</i> AMMD]
gj 116689531	2.8	dihydrolipoamide acetyltransferase [<i>B.cenocepacia</i> HI2424]
gj 167839999	2.8	hypothetical protein Bpse38_25219 [<i>B.thailandensis</i> MSMB43]
gj 171321879	2.8	RND efflux system, outer membrane lipoprotein, NodT family

		[<i>B.ambifaria</i> MEX-5]
gj 53722078	2.8	hypothetical bacteriophage replication protein [<i>B.pseudomallei</i> K96243]
gj 134294497	2.7	Glutamate synthase (ferredoxin) [<i>B.vietnamiensis</i> G4]
gj 134296700	2.7	translation elongation factor G [<i>B.vietnamiensis</i> G4]
gj 115352110	2.6	chromosome segregation protein SMC [<i>B.ambifaria</i> AMMD]
gj 115359514	2.6	porin, Gram-negative type [<i>B.ambifaria</i> AMMD]
gj 116690620	2.6	periplasmic solute binding protein [<i>B.cenocepacia</i> HI2424]
gj 134295001	2.6	thioredoxin reductase [<i>B.vietnamiensis</i> G4]
gj 171320857	2.6	translation elongation factor G [<i>B.ambifaria</i> MEX-5]
gj 189349581	2.6	chaperonin GroEL [<i>B.multivorans</i> ATCC 17616]
gj 189351613	2.6	Amt family ammonium transporter [<i>B.multivorans</i> ATCC 17616]
gj 83719255	2.6	elongation factor G [<i>B.thailandensis</i> E264]
gj 161525461	2.5	ATPase AAA-2 domain protein [<i>B.multivorans</i> ATCC 17616]
gj 115360252	2.4	acetolactate synthase [<i>B.ambifaria</i> AMMD]
gj 115360412	2.4	ATPase AAA-2 domain protein [<i>B.ambifaria</i> AMMD]
gj 134295072	2.4	DNA gyrase subunit A [<i>B.vietnamiensis</i> G4]
gj 189350910	2.4	competence protein [<i>B.multivorans</i> ATCC 17616]
gj 100916983	2.3	hypothetical protein Bmal10_03001635 [<i>B.mallei</i> 10399]
gj 134291992	2.3	Extracellular ligand-binding receptor [<i>B.vietnamiensis</i> G4]
gj 83718787	2.3	pyruvate dehydrogenase subunit E1 [<i>B.thailandensis</i> E264]
gj 83721530	2.3	dihydroaeruginic acid synthetase [<i>B.thailandensis</i> E264]
gj 91777514	2.3	ABC xylose transporter, periplasmic ligand binding protein [<i>B.xenovorans</i> LB400]
gj 116689403	2.2	cellulose synthase operon C domain protein [<i>B.cenocepacia</i> HI2424]
gj 134294300	2.2	hypothetical protein Bcep1808_0177 [<i>B.vietnamiensis</i> G4]
gj 134295584	2.1	dihydroliipoamide dehydrogenase [<i>B.vietnamiensis</i> G4]
gj 162228938	2.1	acetyl-CoA synthetase [<i>B.cenocepacia</i> HI2424]
gj 189351861	2.1	branched-chain amino acid transport system substrate-binding protein [<i>B.multivorans</i> ATCC 17616]
gj 116688365	2	DNA-directed RNA polymerase subunit beta' [<i>B.cenocepacia</i> HI2424]
gj 116693372	2	peptidase S10, serine carboxypeptidase [<i>B.cenocepacia</i> HI2424]
gj 4128219	1.9	transposase [<i>B.cepacia</i>]
gj 83716416	1.9	peptide synthetase, putative [<i>B.thailandensis</i> E264]
gj 91777244	1.9	succinate dehydrogenase flavoprotein subunit [<i>B.xenovorans</i> LB400]
gj 91778087	1.9	Cellulose synthase (UDP-forming) [<i>B.xenovorans</i> LB400]
gj 91782419	1.9	phosphoenolpyruvate carboxylase [<i>B.xenovorans</i> LB400]
gj 115352837	1.8	FAD linked oxidase domain protein [<i>B.ambifaria</i> AMMD]
gj 116693762	1.8	malate synthase G [<i>B.cenocepacia</i> HI2424]
gj 171318018	1.8	peptidase M61 domain protein [<i>B.ambifaria</i> MEX-5]
gj 189350329	1.8	GTP-binding protein [<i>B.multivorans</i> ATCC 17616]
gj 115358682	1.7	protein of unknown function DUF470 [<i>B.ambifaria</i> AMMD]
gj 115359737	1.7	hypothetical protein Bamb_4992 [<i>B.ambifaria</i> AMMD]
gj 116693816	1.7	uncharacterized domain [<i>B.cenocepacia</i> HI2424]
gj 134291880	1.7	excinuclease ABC, A subunit [<i>B.vietnamiensis</i> G4]
gj 189350322	1.7	translation initiation factor IF-2 [<i>B.multivorans</i> ATCC 17616]
gj 83716389	1.7	Rhs element Vgr protein [<i>B.thailandensis</i> E264]

gij83721240	1.7	dipeptide ABC transporter, periplasmic dipeptide-binding protein [<i>B.thailandensis</i> E264]
gij160697538	1.6	conserved hypothetical protein [<i>B.mallei</i> ATCC 10399]
gij100917414	1.5	hypothetical protein Bmal10_03001163 [<i>B.mallei</i> 10399]
gij115361133	1.5	chaperonin GroEL [<i>B.ambifaria</i> AMMD]
gij167837458	1.5	formate dehydrogenase, beta subunit [<i>B.thailandensis</i> MSMB43]
gij167839356	1.5	TonB-dependent siderophore receptor [<i>B.thailandensis</i> MSMB43]
gij189351806	1.5	DNA-directed RNA polymerase, beta subunit [<i>B.multivorans</i> ATCC 17616]
gij52210850	1.5	putative DnaK chaperone protein [<i>B.pseudomallei</i> K96243]
gij115351986	1.4	putative DnaK chaperone protein [<i>B.pseudomallei</i> K96243]
gij189349722	1.4	penicillin-binding protein, 1A family [<i>B.ambifaria</i> AMMD]
gij189349722	1.4	membrane alanyl aminopeptidase [<i>B.multivorans</i> ATCC 17616]
gij189351805	1.4	DNA-directed RNA polymerase, beta' subunit [<i>B.multivorans</i> ATCC 17616]
gij134294436	1.3	DNA-directed RNA polymerase subunit beta' [<i>B.vietnamiensis</i> G4]
gij189348496	1.3	type IV secretion system protein [<i>B.multivorans</i> ATCC 17616]
gij53720952	1.3	putative bacteriophage protein [<i>B.pseudomallei</i> K96243]
gij116689662	1.2	amino acid adenylation domain [<i>B.cenocepacia</i> HI2424]
gij91784463	1.2	Putative Transcriptional regulator, Fis family [<i>B.xenovorans</i> LB400]
gij171320877	1.1	methionyl-tRNA synthetase [<i>B.ambifaria</i> MEX-5]
gij134280841	1	Chaperone clpB [<i>B.pseudomallei</i> 305]
gij100917296	0.8	hypothetical protein Bmal10_03001521 [<i>B.mallei</i> 10399]
gij171320382	0.8	Beta-ketoacyl synthase [<i>B.ambifaria</i> MEX-5]
gij83718897	0.8	peptide synthetase, putative [<i>B.thailandensis</i> E264]
gij160698787	0.7	valyl-tRNA synthetase [<i>B.mallei</i> ATCC 10399]
gij189350474	0.7	D-alanine-poly(phosphoribitol) ligase [<i>B.multivorans</i> ATCC 17616]
gij189352577	0.7	superfamily II helicase [<i>B.multivorans</i> ATCC 17616]