

Taxon	RefSeqID	Function
<i>Pseudomonas</i> sp. ATCC13985	WP009049534.1	hypothetical protein
<i>Pseudomonas</i> sp. ATCC13985	WP014335965.1	retention module-containing protein
<i>Pseudomonas</i> sp. ATCC13985	WP009041592.1	DUF4347 domain-containing protein
<i>Pseudomonas</i> sp. ATCC13985	WP016491125.1	cyclic pyranopterin monophosphate synthase MoaC
<i>Pseudomonas</i> sp. ATCC13985	WP016702937.1	1-deoxy-D-xylulose-5-phosphate synthase
<i>Pseudomonas</i> sp. ATCC13985	WP020297948.1	matrixin family metalloprotease
<i>Pseudomonas</i> sp. ATCC13985	WP011063911.1	hypothetical protein
<i>Pseudomonas</i> sp. ATCC13985	WP009049534.1	hypothetical protein
<i>Pseudomonas</i> sp. ATCC13985	WP017737192.1	acetyl-CoA carboxylase biotin carboxylase subunit
<i>Pseudomonas</i> sp. ATCC13985	WP003186930.1	type I secretion C-terminal target domain-containing protein
<i>Pseudomonas</i> sp. ATCC13985	WP008088213.1	acetyl/propionyl/methylcrotonyl-CoA carboxylase subunit alpha
<i>Pseudomonas</i> sp. ATCC13985	WP016704055.1	autotransporter domain-containing protein
<i>Pseudomonas</i> sp. ATCC13985	WP008047384.1	arginine-ornithine antiporter
<i>Pseudomonas</i> sp. ATCC13985	WP009042593.1	type I secretion C-terminal target domain-containing protein
<i>Arthrobacter</i> sp. KBS0702	WP011693971.1	ABC transporter ATP-binding protein
<i>Janthinobacterium</i> sp. KBS0711	WP010399297.1	alpha-hydroxy-acid oxidizing protein
<i>Janthinobacterium</i> sp. KBS0711	WP013236088.1	cytochrome c oxidase subunit II
<i>Janthinobacterium</i> sp. KBS0711	WP010393863.1	mechanosensitive ion channel family protein
<i>Janthinobacterium</i> sp. KBS0711	WP010395693.1	DNA gyrase subunit A
<i>Janthinobacterium</i> sp. KBS0711	WP010394499.1	c-type cytochrome biogenesis protein CcsB
<i>Janthinobacterium</i> sp. KBS0711	WP009239942.1	SufS family cysteine desulfurase
<i>Janthinobacterium</i> sp. KBS0711	WP010394527.1	GTPase
<i>Janthinobacterium</i> sp. KBS0711	WP020701617.1	flagellar type III secretion system pore protein FlpP
<i>Janthinobacterium</i> sp. KBS0711	WP008451236.1	ATP-dependent RNA helicase HrpA
<i>Janthinobacterium</i> sp. KBS0711	WP018059363.1	methionine-tRNA ligase
<i>Janthinobacterium</i> sp. KBS0711	WP008450068.1	chemotaxis response regulator protein-glutamate methylesterase
<i>Janthinobacterium</i> sp. KBS0711	WP020656832.1	valine-tRNA ligase
<i>Janthinobacterium</i> sp. KBS0711	WP018058224.1	acetyl-CoA C-acetyltransferase
<i>Janthinobacterium</i> sp. KBS0711	WP020701147.1	ribonucleotide-diphosphate reductase subunit beta
<i>Janthinobacterium</i> sp. KBS0711	WP017877147.1	DNA polymerase III subunit alpha
<i>Janthinobacterium</i> sp. KBS0711	WP018058636.1	sulfate ABC transporter permease subunit CysW
<i>Janthinobacterium</i> sp. KBS0711	WP020702753.1	30S ribosomal protein S2
<i>Janthinobacterium</i> sp. KBS0711	WP017876556.1	methyl-accepting chemotaxis protein
<i>Janthinobacterium</i> sp. KBS0711	WP010397062.1	aminotransferase class I/II-fold pyridoxal phosphate-dependent enzyme
<i>Janthinobacterium</i> sp. KBS0711	WP020704252.1	methionine adenosyltransferase
<i>Janthinobacterium</i> sp. KBS0711	WP002611629.1	substrate-binding domain-containing protein
<i>Janthinobacterium</i> sp. KBS0711	WP005664034.1	F0F1 ATP synthase subunit alpha
<i>Janthinobacterium</i> sp. KBS0711	WP010462720.1	ABC transporter permease
<i>Janthinobacterium</i> sp. KBS0711	WP008452953.1	ethanolamine ammonia-lyase subunit EutB
<i>Janthinobacterium</i> sp. KBS0711	WP017878982.1	amino-acid N-acetyltransferase
<i>Janthinobacterium</i> sp. KBS0711	WP011979236.1	filamentous hemagglutinin N-terminal domain-containing protein
<i>Janthinobacterium</i> sp. KBS0711	WP008446499.1	cation acetate symporter
<i>Janthinobacterium</i> sp. KBS0711	WP008120030.1	urea ABC transporter substrate-binding protein
<i>Janthinobacterium</i> sp. KBS0711	WP010399469.1	HAMP domain-containing protein
<i>Janthinobacterium</i> sp. KBS0711	WP010399755.1	acetolactate synthase 3 catalytic subunit
<i>Janthinobacterium</i> sp. KBS0711	WP020201433.1	iron-sulfur cluster assembly protein IscA
<i>Variovorax</i> sp. KBS0712	WP007851335.1	amidohydrolase
<i>Variovorax</i> sp. KBS0712	WP017331249.1	FAD-binding protein
<i>Variovorax</i> sp. KBS0712	WP012203717.1	SDR family oxidoreductase
<i>Variovorax</i> sp. KBS0712	WP003053566.1	isocitrate lyase
<i>Variovorax</i> sp. KBS0712	WP019655940.1	ATP-dependent chaperone ClpB
<i>Variovorax</i> sp. KBS0712	WP009124777.1	LamB/YcsF family protein
<i>Variovorax</i> sp. KBS0712	WP015014884.1	ATP-dependent Clp protease ATP-binding subunit ClpX
<i>Variovorax</i> sp. KBS0712	WP013541265.1	ABC transporter ATP-binding protein
<i>Variovorax</i> sp. KBS0712	WP013543193.1	proline-tRNA ligase
<i>Variovorax</i> sp. KBS0712	WP019651994.1	aldehyde dehydrogenase family protein
<i>Variovorax</i> sp. KBS0712	WP007837263.1	DUF802 domain-containing protein
<i>Variovorax</i> sp. KBS0712	WP007828806.1	PAS domain S-box protein
<i>Variovorax</i> sp. KBS0712	WP012746430.1	thioredoxin-disulfide reductase
<i>Variovorax</i> sp. KBS0712	WP013800395.1	methylcrotonoyl-CoA carboxylase
<i>Variovorax</i> sp. KBS0712	WP015866795.1	polyphosphate kinase 2
<i>Variovorax</i> sp. KBS0712	WP019658558.1	ATP-binding cassette domain-containing protein
<i>Variovorax</i> sp. KBS0712	WP018903541.1	sn-glycerol-3-phosphate ABC transporter ATP-binding protein UgpC
<i>Variovorax</i> sp. KBS0712	WP003164005.1	glutathione S-transferase family protein
<i>Variovorax</i> sp. KBS0712	WP013538653.1	urocanate hydratase
<i>Variovorax</i> sp. KBS0712	WP013538795.1	thiamine pyrophosphate-binding protein
<i>Variovorax</i> sp. KBS0712	WP007835149.1	transaldolase
<i>Variovorax</i> sp. KBS0712	WP013543213.1	preprotein translocase subunit SecA
<i>Variovorax</i> sp. KBS0712	WP013540041.1	acyl-CoA dehydrogenase
<i>Variovorax</i> sp. KBS0712	WP013539022.1	hypothetical protein
<i>Variovorax</i> sp. KBS0712	WP006297699.1	elongation factor G
<i>Variovorax</i> sp. KBS0712	WP017328757.1	cytochrome P450
<i>Variovorax</i> sp. KBS0712	WP007828445.1	LLM class flavin-dependent oxidoreductase
<i>Variovorax</i> sp. KBS0712	WP009051660.1	phosphatase PAP2/dual specificity phosphatase family protein
<i>Variovorax</i> sp. KBS0712	WP018904684.1	amino acid adenylation domain-containing protein
<i>Variovorax</i> sp. KBS0712	WP013540507.1	oxidoreductase
<i>Variovorax</i> sp. KBS0712	WP007830805.1	pyridoxal phosphate-dependent aminotransferase
<i>Variovorax</i> sp. KBS0712	WP018906716.1	DUF962 domain-containing protein