

Taxon	RefSeqID	Function
<i>Pseudomonas</i> sp. ATCC13985	WP009049534.1	hypothetical protein
<i>Pseudomonas</i> sp. ATCC13985	WP009048342.1	BapA prefix-like domain-containing protein
<i>Pseudomonas</i> sp. ATCC13985	WP009049534.1	hypothetical protein
<i>Pseudomonas</i> sp. ATCC13985	WP007920721.1	DUF2931 family protein
<i>Pseudomonas</i> sp. ATCC13985	WP009041592.1	DUF4347 domain-containing protein
<i>Pseudomonas</i> sp. ATCC13985	WP009042593.1	type I secretion C-terminal target domain-containing protein
<i>Pseudomonas</i> sp. ATCC13985	WP007920721.1	DUF2931 family protein
<i>Pseudomonas</i> sp. ATCC13985	WP008047384.1	arginine-ornithine antiporter
<i>Pseudomonas</i> sp. ATCC13985	WP016059839.1	hypothetical protein
<i>Janthinobacterium</i> sp. KBS0711	WP020656832.1	valine-tRNA ligase
<i>Janthinobacterium</i> sp. KBS0711	WP010399297.1	alpha-hydroxy-acid oxidizing protein
<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	WP010394527.1	GTPase
<i>Janthinobacterium</i> sp. KBS0711	WP009239942.1	SufS family cysteine desulfurase
<i>Janthinobacterium</i> sp. KBS0711	WP013236088.1	cytochrome c oxidase subunit II
<i>Janthinobacterium</i> sp. KBS0711	WP017875079.1	electron transfer flavoprotein-ubiquinone oxidoreductase
<i>Janthinobacterium</i> sp. KBS0711	WP020701617.1	flagellar type III secretion system pore protein FlpP
<i>Janthinobacterium</i> sp. KBS0711	WP020701147.1	ribonucleotide-diphosphate reductase subunit beta
<i>Janthinobacterium</i> sp. KBS0711	WP008450068.1	chemotaxis response regulator protein-glutamate methylesterase
<i>Janthinobacterium</i> sp. KBS0711	WP018059363.1	methionine-tRNA ligase
<i>Janthinobacterium</i> sp. KBS0711	WP020702753.1	30S ribosomal protein S2
<i>Janthinobacterium</i> sp. KBS0711	WP018058636.1	sulfate ABC transporter permease subunit CysW
<i>Janthinobacterium</i> sp. KBS0711	WP019923632.1	bacterioferritin
<i>Janthinobacterium</i> sp. KBS0711	WP019139956.1	ABC transporter ATP-binding protein
<i>Janthinobacterium</i> sp. KBS0711	WP005664034.1	F0F1 ATP synthase subunit alpha
<i>Janthinobacterium</i> sp. KBS0711	WP020704252.1	methionine adenosyltransferase
<i>Janthinobacterium</i> sp. KBS0711	WP010394499.1	c-type cytochrome biogenesis protein CcsB
<i>Janthinobacterium</i> sp. KBS0711	WP010397062.1	aminotransferase class I/II-fold pyridoxal phosphate-dependent enzyme
<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>Janthinobacterium</i> sp. KBS0711	WP017878982.1	amino-acid N-acetyltransferase
<i>Janthinobacterium</i> sp. KBS0711	WP008452953.1	ethanolamine ammonia-lyase subunit EutB
<i>Janthinobacterium</i> sp. KBS0711	WP010462720.1	ABC transporter permease
<i>Janthinobacterium</i> sp. KBS0711	WP010401750.1	leucine-tRNA ligase
<i>Variovorax</i> sp. KBS0712	WP014006191.1	hypothetical protein
<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	WP013541265.1	dipeptide ABC transporter ATP-binding protein
<i>Variovorax</i> sp. KBS0712	WP015014884.1	ATP-dependent Clp protease ATP-binding subunit ClpX
<i>Variovorax</i> sp. KBS0712	WP019651994.1	aldehyde dehydrogenase family protein
<i>Variovorax</i> sp. KBS0712	WP009124777.1	5-oxoprolinase subunit PxpA
<i>Variovorax</i> sp. KBS0712	WP007827819.1	AMP-binding 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	WP013540385.1	RNA polymerase sigma factor RpoD
<i>Variovorax</i> sp. KBS0712	WP018903541.1	sn-glycerol-3-phosphate ABC transporter ATP-binding protein UgpC
<i>Variovorax</i> sp. KBS0712	WP019658558.1	ATP-binding cassette domain-containing protein
<i>Variovorax</i> sp. KBS0712	WP007830496.1	tryptophan synthase subunit beta
<i>Variovorax</i> sp. KBS0712	WP015867550.1	formyltetrahydrofolate deformylase
<i>Variovorax</i> sp. KBS0712	WP013538653.1	urocanate hydratase
<i>Variovorax</i> sp. KBS0712	WP007830805.1	pyridoxal phosphate-dependent aminotransferase
<i>Variovorax</i> sp. KBS0712	WP007828445.1	MsnO8 family LLM class oxidoreductase
<i>Variovorax</i> sp. KBS0712	WP006297699.1	elongation factor G
<i>Curtobacterium</i> sp. KBS0715	WP017886346.1	Asp23/Gls24 family envelope stress response protein
<i>Curtobacterium</i> sp. KBS0715	WP017886981.1	AMP-binding protein
<i>Curtobacterium</i> sp. KBS0715	WP017888661.1	response regulator
<i>Curtobacterium</i> sp. KBS0715	WP018170758.1	glutamate synthase large subunit
<i>Curtobacterium</i> sp. KBS0715	WP017887928.1	glycoside hydrolase family 92 protein
<i>Curtobacterium</i> sp. KBS0715	WP017886488.1	sigma-70 family RNA polymerase sigma factor
<i>Curtobacterium</i> sp. KBS0715	WP012039163.1	response regulator transcription factor
<i>Curtobacterium</i> sp. KBS0715	WP014049992.1	helix-turn-helix domain-containing protein
<i>Curtobacterium</i> sp. KBS0715	WP017886283.1	organic hydroperoxide resistance protein
<i>Curtobacterium</i> sp. KBS0715	WP017886367.1	AAA domain-containing protein
<i>Curtobacterium</i> sp. KBS0715	WP015889560.1	acyl-CoA dehydrogenase
<i>Curtobacterium</i> sp. KBS0715	WP009740872.1	HAMP domain-containing protein