			K00656	1	class
			K02035		43 kbp HMO cluster
			K02429	0.8	Sialidase_cluster
			K00048		Fucosidase_cluster
			K15533	0.6	LNT related gene
			K12574		GNB-LNB_cluster
			K02033	0.4	GluNAc-GalNAc_related_gene
			K01714	• .	alpha_mannose_related_gene
			K12308	0.2	urease_related_gene
			K02031	0.2	CRISPR_related_gene
			K02784		ortion it_rolated_game
			K17319	0	subclass
			K02529		Alpha-galactosidase
			K02026		Alpha-L-fucosidase
			K02027		Alpha-mannosidase
			K08483		Beta-galactosidase
			K02034		Beta-lactamase
			K16209 K13059		Binding-protein-dependent transport system Inner membrane protein
			K02025		Dihydrodipicolinate synthase
			K02023		Endo-alpha-N-acetylgalactosaminidase
			K02564		Family 1 extracellular solute-binding protein
			K02032		formate C-acetyltransferase
			K01191		fusion between oligopeptide transport permease oppC and ATP-binding protein oppD
			K17320	ĺ	Galactoside symporter
			K17624	ĺ	Glucosamine–6–phosphate isomerase
			K01443		high-affinity nickel-transporter
			K15342		inner-membrane translocator
			K09951		Integrase
			K02777		L-fuconolactonase
			K02804		L-fucose permease
			K03190		Lacl-type transcriptional regulator
			K07046		lactaldehyde reductase
			K07241		Lacto-N-biose phorylase
			K11960 K11961		LNT beta-1,3-galactosidase
			K11901 K15923		Mandelate racemase/muconate lactonizing domain–containing protein
			K22397		N–Acetylglucosamine–6–phosphate deacetylase
			K19119		N–Acetylmannosamine–6–phosphate 2–epimerase,NanE
			K01206		N–Acetylneuraminate lyase
			K01428		Oligopeptide transport system permease protein
			K01430		Oligopeptide-binding protein oppA
			K03188		Permease protein of ABC transporter system for sugars
			K03189		Phosphocarrier protein HPr
			K18334		Phosphoenolpyruvate-protein phosphotransferase
			K19117		PTS system, glucose–specific IIABC component
			K19118		PTS system, N-acetylglucosamine-specific IIBC component
			K01186		ROK family transcriptional regulator
			K01190		Sialidase A
			K01788		Solute–binding protein of ABC transporter system for sugars
			K03187		Sugar ABC transporter permease
			K07012 K00845		type I–C
			K17318		type I–E
			K00965		urea inner–membrane translocator
<u>C</u> S	<u>र</u> ु	St			Urease accessory protein
subclass class	strain2 strain1	strain3		1	
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