PC1	negativ	e loadings	PC1	positive	loadings
-1	K03205	type IV secretion system protein VirD4 [EC:7.4.2.8]	1	K01903	succinyl-CoA synthetase beta subunit [EC:6.2.1.5]
-2	K16926	energy-coupling factor transport system substrate-specific component	2	K01092	myo-inositol-1(or 4)-monophosphatase [EC:3.1.3.25]
-3	K02005	HlyD family secretion protein	3	K01902	succinyl-CoA synthetase alpha subunit [EC:6.2.1.5]
-4	K02117	V/A-type H+/Na+-transporting ATPase subunit A [EC:7.1.2.2 7.2.2.1]	4	K01961	acetyl-CoA carboxylase, biotin carboxylase subunit [EC:6.4.1.2 6.3.4.14]
-5	K07133	uncharacterized protein	5	K00526	ribonucleoside-diphosphate reductase beta chain [EC:1.17.4.1]
-6	K16785	energy-coupling factor transport system permease protein	6	K01696	tryptophan synthase beta chain [EC:4.2.1.20]
-7	K02034	peptide/nickel transport system permease protein	7	K07173	S-ribosylhomocysteine lyase [EC:4.4.1.21]
-8	K07052	uncharacterized protein	8	K00674	2,3,4,5-tetrahydropyridine-2,6-dicarboxylate N- succinyltransferase [EC:2.3.1.117]
		branched-chain amino acid transport			2-oxoglutarate dehydrogenase E2 component (dihydrolipoamide succinyltransferase)
-9	K01999	system substrate-binding protein	9	K00658	[EC:2.3.1.61]
-		ATP-binding cassette, subfamily B,			anthranilate synthase component I
10	K06147	bacterial	10	K01657	[EC:4.1.3.27]