

MycoCosm KEGG Browser

KEGG stands for Kyoto Encyclopedia of Genes and Genomes at <http://www.genome.jp/kegg/>, which maintains a curated set of EC-annotated enzymes and their pathways. Each portal's KEGG Browser facilitates display and discovery of MycoCosm's KEGG-annotated genes.

Scenario: You have plated a variety of yeasts on a variety of carbon sources, and discovered that some members of the Pichiaceae grow on galactose (e.g. *Dekkera bruxellensis*) and some do not (e.g. *Pichia membranifaciens*). Use MycoCosm to find genes that could explain this metabolic difference.

- 1) Go to the MycoCosm Pichiaceae PhyloGroup at mycocosm.jgi.doe.gov/Pichiaceae :

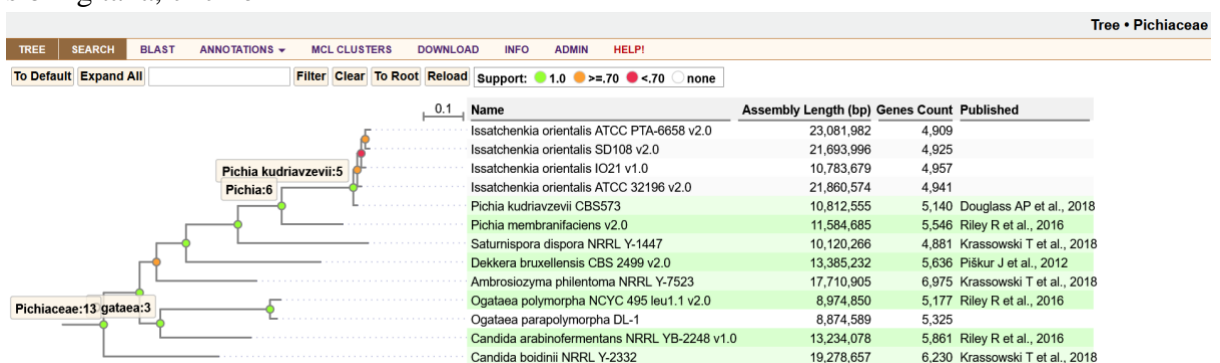
Info • Pichiaceae

TREE SEARCH BLAST ANNOTATIONS MCL CLUSTERS DOWNLOAD INFO **HELP!**

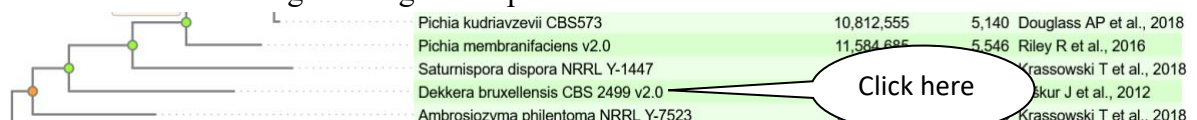
Group Name: Pichiaceae

##	Name	Assembly Length	# Genes	Published
1	Candida arabinofementans NRRL YB-2248 v1.0	13,234,078	5,861	Riley R et al., 2016
2	Candida boidinii NRRL Y-2332	19,278,657	6,230	Krassowski T et al., 2018
3	Dekkera bruxellensis CBS 2499 v2.0	13,385,232	5,636	Piskur J et al., 2012
4	Issatchenkia orientalis ATCC 32196 v2.0	21,860,574	4,941	
5	Issatchenkia orientalis ATCC PTA-6658 v2.0	23,081,982	4,909	
6	Issatchenkia orientalis SD108 v2.0	21,693,996	4,925	
7	Nakazawaea wickerhamii NRRL Y-2563	10,857,285	5,490	Krassowski T et al., 2018
8	Ogataea parapolyomorpha DL-1	8,874,589	5,325	
9	Ogataea polymorpha NCYC 495 leu1.1 v2.0	8,974,850	5,177	Riley R et al., 2016
10	Pichia kudriavzevii CBS573	10,812,555	5,140	Douglass AP et al., 2018
11	Pichia membranifaciens v2.0	11,584,685	5,546	Riley R et al., 2016
12	Saturnispora dispora NRRL Y-1447	10,120,266	4,881	Krassowski T et al., 2018

- 2) To verify that *Dekkera* (which grows on galactose) and *Pichia* (which does not) are sibling taxa, click on 'TREE':



- 3) Click on 'Dekkera' to go to its genome portal:



- 4) Click on “ANNOTATIONS => KEGG” to go to the portal’s KEGG browser:

KEGG • Dekkera bruxellensis CBS 2499 v2.0

SEARCH BLAST BROWSE ANNOTATIONS MCL CLUSTERS SYNTENY DOWNLOAD INFO HOME STATUS HELP!

Select Model Set(s) to View:

Dekkera bruxellensis CBS 2499 v2.0/FilteredModels1 (ver 1)
Pichia membranifaciens v2.0/FilteredModels1 (ver 1)
Ogataea polymorpha NCYC 495 leu1.1 v2.0/FilteredModels2 (ver 1)
Candida arabinoferrmentans NRRL YB-2248 v1.0/FilteredModels1 (ver 1)

apply

Other Functions
View KEGG Metabolic Pathways
View KEGG Regulatory Pathways
Search KEGG
Enzyme Commission Numbers

KEGG Metabolic Pathway

Amino Acid Metabolism

Pathway	models in Dekkera bruxellensis CBS 2499 v2.0/FilteredModels1 (ver 1)
Amino sugar and nucleotide sugar metabolism	206
Alanine, aspartate and glutamate metabolism	27
Arginine and proline metabolism	45

- 5) Scroll down to the ‘Carbohydrate Metabolism’ section, and find the subsection ‘Galactose metabolism’. Dekkera has 24 genes annotated to this metabolic pathway:

Carbohydrate Metabolism	332
Amino sugar and nucleotide sugar metabolism	68
Ascorbate and aldarate metabolism	21
Butanoate metabolism	34
C5-Branched dibasic acid metabolism	2
Citrate cycle (TCA cycle)	28
Fructose and mannose metabolism	46
Galactose metabolism	24
Glycolysis / Gluconeogenesis	47
Glyoxylate and dicarboxylate metabolism	10
Inositol phosphate metabolism	27

- 6) Click on ‘Galactose metabolism’ to drill down into the KEGG hierarchy and list the EC numbers of that pathway.
- 7) Go to the ‘Select Model Set(s) to View’ list box and select *Dekkera* and *Pichia* and click the ‘apply’ button. The *Dekkera* and *Pichia* galactose metabolism gene counts are side-by-side and may be directly compared. Galactokinase (EC = 2.7.1.6) and UDPglucose--hexose-1-phosphate uridylyltransferase (2.7.7.12) are each present in *Dekkera* but not in *Pichia*:

KEGG • Dekkera bruxellensis CBS 2499 v2.0

SEARCH BLAST BROWSE ANNOTATIONS MCL CLUSTERS SYNTENY DOWNLOAD INFO HOME STATUS HELP!

Select Model Set(s) to View:

Dekkera bruxellensis CBS 2499 v2.0/FilteredModels1 (ver 1)
Pichia membranifaciens v2.0/FilteredModels1 (ver 1)
Ogataea polymorpha NCYC 495 leu1.1 v2.0/FilteredModels2 (ver 1)
Candida arabinoferrmentans NRRL YB-2248 v1.0/FilteredModels1 (ver 1)

apply

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Search KEGG

MAP00052: Galactose metabolism

[Summary View | Model View | View KEGG Map]

EC Number	Description	models in Dekkera bruxellensis CBS 2499 v2.0/FilteredModels1 (ver 1)	models in Pichia membranifaciens v2.0/FilteredModels1 (ver 1)	models in all selected model sets
1.1.1.120	galactose 1-dehydrogenase (NADP+)	0	0	0
1.1.1.16	galactitol 2-dehydrogenase	0	0	0

Dekkera bruxellensis CBS 2499 v2.0/FilteredModels1 (ver 1)
Pichia membranifaciens v2.0/FilteredModels1 (ver 1)
Ogataea polymorpha NCYC 495 leu1.1 v2.0/FilteredModels2 (ver 1)
Candida arabinofementans NRRL YB-2248 v1.0/FilteredModels1 (ver 1)

[View KEGG Metabolic Pathways](#)
[View KEGG Regulatory Pathways](#)
[Search KEGG](#)

[[Summary View](#) | [Model View](#) | [View KEGG Map](#)]

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- Riley R, Haridas S, Wolfe KH, Lopes MR, Hittinger CT, Göker M, Salamov AA, Wisecaver JH, Long TM, Calvey CH, Aerts AL, Barry KW, Choi C, Clum A, Coughlan AY, Deshpande S, Douglass AP, Hanson SJ, Klenk HP, LaButti KM, Lapidus A, Lindquist EA, Lipzen AM, Meier-Kolthoff JP, Ohm RA, Otiillar RP, Pangilinan JL, Peng Y, Rokas A, Rosa CA, Scheuner C, Sibirny AA, Slot JC, Stielow JB, Sun H, Kurtzman CP, Blackwell M, Grigoriev IV, Jeffries TW. Comparative genomics of biotechnologically important yeasts. *Proc Natl Acad Sci U S A*. 2016 Aug

30;113(35):9882-7. doi: 10.1073/pnas.1603941113. Epub 2016 Aug 17. PubMed PMID: 27535936; PubMed Central PMCID: PMC5024638.