

KEGG Orthology (KO)

- Integrar a informação genômica às redes protéicas através dos números EC.
- Utiliza identificadores comuns nos genomas e nas vias metabólicas.
- Grupos eram formados por análise manual de um banco de similaridades combinada com características evolutivas.
- Informação propagada utilizando ferramentas computacionais (KOALA)
- Peça chave para o sistema hierárquico do KEGG.
- 18949 KOs com 5320103 proteínas.



Exemplo: Kegg Orthology



ORTHOLOGY K00134

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Entry	K00134	KO
Name	GAPDH, gapA	
Definition	glyceraldehyde 3-phosphate dehydrogenase [EC:1.2.1.12]	
Pathway	ko00010 Glycolysis / Gluconeogenesis ko00710 Carbon fixation in photosynthetic organisms ko01200 Carbon metabolism ko01230 Biosynthesis of amino acids ko04066 HIF-1 signaling pathway ko05010 Alzheimer's disease	
Module	M00001 Glycolysis (Embden-Meyerhof pathway), glucose => pyruvate M00002 Glycolysis, core module involving three-carbon compounds M00003 Gluconeogenesis, oxaloacetate => fructose-6P M00165 Reductive pentose phosphate cycle (Calvin cycle) M00166 Reductive pentose phosphate cycle, ribulose-5P => glyceraldehyde-3P M00308 Semi-phosphorylative Entner-Doudoroff pathway, gluconate => glycerate-3P M00552 D-galactonate degradation, De Ley-Doudoroff pathway, D-galactonate => glycerate-3P	
Brite	KEGG Orthology (KO) [BR: ko00001] Metabolism Overview 01200 Carbon metabolism K00134 GAPDH, gapA; glyceraldehyde 3-phosphate dehydrogenase 01230 Biosynthesis of amino acids K00134 GAPDH, gapA; glyceraldehyde 3-phosphate dehydrogenase Carbohydrate metabolism 00010 Glycolysis / Gluconeogenesis K00134 GAPDH, gapA; glyceraldehyde 3-phosphate dehydrogenase Energy metabolism 00710 Carbon fixation in photosynthetic organisms K00134 GAPDH, gapA; glyceraldehyde 3-phosphate dehydrogenase	

All links

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