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# eLignin a database on microbial catabolism of aromatic compounds derived from lignin



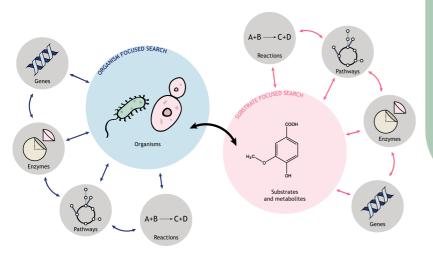
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### BACKGROUND AND AIM

We here present a new database named **eLignin** for collection of data on metabolic pathways and enzymes for the **microbial conversion of lignin and lignin-derived aromatic compounds** to metabolites of the cellular central carbon catabolism.

The aim of eLignin is to bring the existing bibliome together in a single searchable platform. By doing so, we aim to facilitate the overview of the field and generate new connections between different aspects of the molecular biology of lignin- and aromatic catabolism that occurs in nature. Such knowledge will be an important foundation for the creation of new strategies for metabolic engineering of microorganisms to produce value-added products from lignin and/or its derivatives.



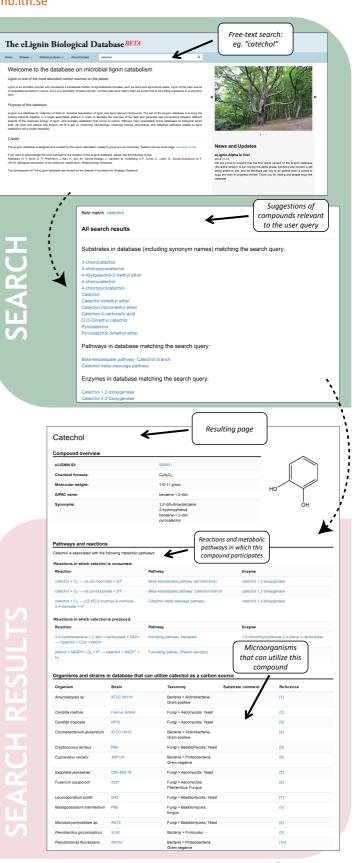
#### www.elignindatabase.com

Figure 1: Schematic representation of an organism and a substrate focused search in eLignin. Please keep in mind that although the database was designed from this point-of-view, it is possible to begin a search in any of the available categories (enzymes, genes, metabolic pathways, organisms, reactions and substrates)

Table 1. The dataset in eLignin, as of March 201		
	Entry	Count
5	Substrates	87
	Organisms	104
Meta	bolic pathways	14
	Reactions	34
	Enzymes	25
2	Genes	31
R	References	131

## -CONCLUSIONS AND OUTLOOK-

- eLignin will fill a gap between the existing biological databases by combining microbiology, molecular biology and metabolism related to lignin catabolism into a single repository.
- By facilitating the overview of the field, we hope that eLignin will be an useful aid in experimental design and metabolic engineering.
- The database is available at <u>www.elignindatabase.com</u>





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