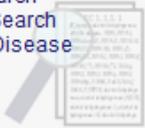
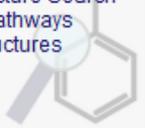
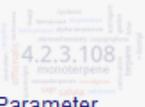


Please enter a search term

Enzyme, Ligand contains

add search field delete search field **start search**

<p><u>Text-based queries</u></p> <ul style="list-style-type: none">Full-text SearchAdvanced SearchEnzyme & Disease 	<p><u>Structure-based queries</u></p> <ul style="list-style-type: none">Ligand Structure SearchMetabolic PathwaysEnzyme Structures 	<p><u>Explorer</u></p> <ul style="list-style-type: none">Enzyme ClassificationTaxTreeProtein folding: CATH / SCOPeOntologies 
<p><u>Visualization</u></p> <ul style="list-style-type: none">Word Maps 4.2.3.108GenomesFunctional Parameter StatisticsMetabolic Pathways 	<p><u>Prediction</u></p> <ul style="list-style-type: none">Membrane HelicesLocalization PredictionEnzymeDetector 	<p><u>Supporting</u></p> <ul style="list-style-type: none">BRENDA Tissue OntologyBiochemical Reactions 

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BRENDA Tutorial

Introduction to the Enzyme Information System

Facts about BRENDA (BRaunschweig ENzyme DAtabase)

- one of the most comprehensive enzyme information repositories
- BRENDA is a member of de.NBI (German Network for Bioinformatics Structure, since 2015)
- BRENDA is an ELIXIR Core Data Resource (since 2018)
- all enzymes, classified by the Enzyme Nomenclature (IUBMB)
- data of molecular biology, biochemistry, medical research, and biotechnology
- furthermore BRENDA includes data from interconnected databases containing results from text mining methods and bioinformatic approaches
- BRENDA is freely available to the scientific community
- more than 80,000 visits of the BRENDA website each month
- major updates of the data in BRENDA are performed twice a year

History and major developments of BRENDA

- BRENDA was created at the former German National Research Center for Biotechnology (GBF, now HZI, Helmholtz Zentrum für Infektionsforschung, Braunschweig, Germany) in 1987
- BRENDA was originally published as a series of book
 - 1st Edition 1990-1997 (Enzyme Handbook)
 - 2nd Edition 2001-2013 (Handbook of Enzymes)
- BRENDA moved to the University of Cologne, Germany
- First online version in 1998 via the SRS system at the EBI
- First website of BRENDA in Cologne
- Transfer of BRENDA into a fully relational database system
- BRENDA moved back to Braunschweig in 2007
- BRENDA is now maintained and further developed at the BRICS - TU Braunschweig

Facts about BRENDA

The main categories are based on the **Enzymes** and the **Metabolites / Ligands**

Enzyme-related data encompasses information on:

- Enzyme and ligand nomenclature
- Organism
- Reaction and specificity
- Kinetic properties
- Structure and role of the ligands
- Stability information
- Ligand-enzyme information
- Enzyme sequence and structure
- Mutants and disease
- Occurrence, isolation, and preparation
- Pathways



BRENDA is the most comprehensive information system on:

- 7862 EC Numbers (July 2019)
- more than 2 Mill. different enzymes
- more than 3 Mill. enzyme data, manually annotated from more than 150,000 literature references

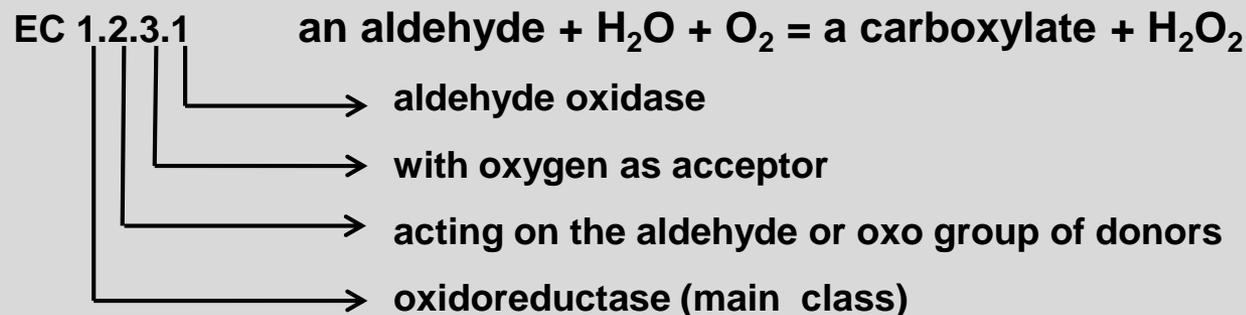
Enzyme Commission numbers (EC Numbers) are defined according to the catalyzed reaction by the IUBMB (International Union of Biochemistry and Molecular Biology)

Format: **Four** numbers separated by periods, e.g. 1.2.3.1

Main Enzyme Classes:

- 1 Oxidoreductases
- 2 Transferases
- 3 Hydrolases
- 4 Lyases
- 5 Isomerases
- 6 Ligases
- 7 Translocases

Numbers represent from left to right a progressively finer classification scheme



(A)

Please enter a search term

Enzyme, Ligand contains

[add search field](#) [delete search field](#) [start search](#)

(B)

<p><u>Text-based queries</u></p> <ul style="list-style-type: none">Full-text SearchAdvanced SearchEnzyme & Disease	<p><u>Structure-based queries</u></p> <ul style="list-style-type: none">Ligand Structure SearchMetabolic PathwaysEnzyme Structures	<p><u>Explorer</u></p> <ul style="list-style-type: none">Enzyme ClassificationTaxTreeProtein folding: CATH / SCOPeOntologies
<p><u>Visualization</u></p> <ul style="list-style-type: none">Word MapsGenomesFunctional Parameter StatisticsMetabolic Pathways	<p><u>Prediction</u></p> <ul style="list-style-type: none">Membrane HelicesLocalization PredictionEnzymeDetector	<p><u>Supporting</u></p> <ul style="list-style-type: none">BRENDA Tissue OntologyBiochemical Reactions

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The main search options:

- Quick access (A)
- and more specific queries (B)
- Classic View (C)
- ...further details in the corresponding BRENDA tutorials



EC-Number	Enzyme Name	Organism	Protein	Full text	Ligand	Advanced Search
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/> Search		Display <input type="text" value="10"/> ▾		entries		
Nomenclature		Reaction & Specificity			Functional Parameters	
<ul style="list-style-type: none"> Enzyme Names EC-Number 		<ul style="list-style-type: none"> Pathway Catalysed Reaction Reaction Type Natural Substrates & Products Substrates & Products Inhibitors Cofactors Metals/Ions Activating Compounds Ligands 			<ul style="list-style-type: none"> K_M Value k_{cat}/K_M Value K_i Value IC_{50} Value pI Value Turnover Number Specific Activity pH Optimum pH Range Temperature Optimum Temperature Range Kinetic ENzyme DATA 	
Organism-related information		<ul style="list-style-type: none"> Biochemicals Reactions 			<ul style="list-style-type: none"> <ul style="list-style-type: none"> K_M Value k_{cat}/K_M Value K_i Value IC_{50} Value pI Value Turnover Number Specific Activity pH Optimum pH Range Temperature Optimum Temperature Range Kinetic ENzyme DATA 	
<ul style="list-style-type: none"> Organism Source Tissue Localization 						
Isolation & Preparation		<ul style="list-style-type: none"> Biochemicals Reactions 			<ul style="list-style-type: none"> <ul style="list-style-type: none"> K_M Value k_{cat}/K_M Value K_i Value IC_{50} Value pI Value Turnover Number Specific Activity pH Optimum pH Range Temperature Optimum Temperature Range Kinetic ENzyme DATA 	
<ul style="list-style-type: none"> Purification Cloned Expression Renatured Crystallization 						
Stability		Enzyme Structure			Disease, Engineering & Application	
<ul style="list-style-type: none"> pH Stability Temperature Stability General Stability Organic Solvent Stability Oxidation Stability Storage Stability 		<ul style="list-style-type: none"> Sequence 3D-Structure Molecular Weight Subunits Posttranslational Modification 			<ul style="list-style-type: none"> Disease/ Diagnostics Engineering Application 	
					References	

BRENDA data and information fields „classic view“

