

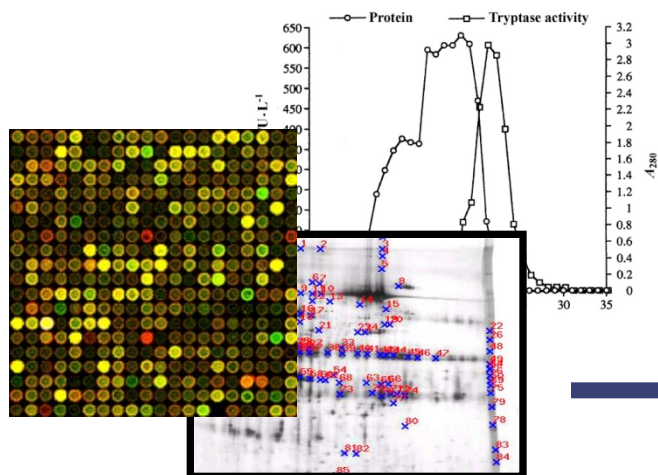
Access to Biochemical Reaction Kinetics Data

Martin Golebiewski

Scientific Databases and Visualization Group

Heidelberg Institute for Theoretical Studies

HITS gGmbH, Germany



Experimental Data

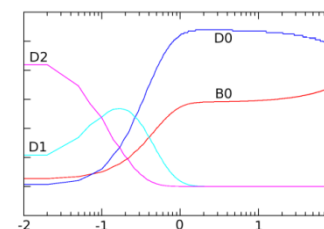
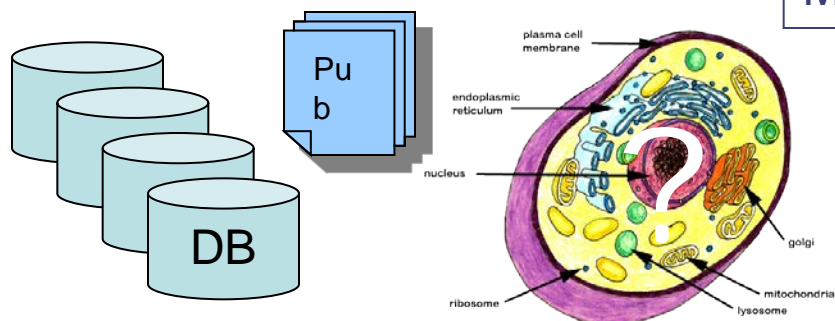
Calculations

$$\begin{aligned}
 [G\alpha]' &= k_1 + (k_2[G\alpha]) - k_3 \frac{[G\alpha][PLC]}{([G\alpha] + K_4)} - k_5 \frac{[G\alpha][Ca_{cyt}]}{([G\alpha] + K_6)} \\
 [PLC]' &= k_7[G\alpha] - k_8 \frac{[PLC]}{([PLC] + K_9)} \\
 [Ca_{cyt}]' &= (Ca_{ER} - Ca_{cyt}) * \frac{k_{10} * Ca_{cyt} * PLC^4}{PLC^4 + K_{11}^4} + k_{12} * PLC + k_{13} * [G\alpha] \\
 &\quad - k_{14} \frac{[Ca_{cyt}]}{([Ca_{cyt}] + K_{15})} - k_{16} \frac{[Ca_{cyt}]}{([Ca_{cyt}] + K_{17})} - k_{18} \frac{[Ca_{cyt}]^n}{([Ca_{cyt}]^n + K_{19}^n)} \\
 &\quad + (Ca_{mit} - Ca_{cyt}) * k_{20} \frac{[Ca_{cyt}]}{([Ca_{cyt}] + K_{21})} \\
 [Ca_{ER}]' &= -(Ca_{ER} - Ca_{cyt}) * \frac{k_{10} * Ca_{cyt} * PLC^4}{PLC^4 + K_{11}^4} + k_{16} \frac{[Ca_{cyt}]}{([Ca_{cyt}] + K_{17})} \\
 [Ca_{Mito}]' &= k_{18} \frac{[Ca_{cyt}]^n}{([Ca_{cyt}]^n + K_{19}^n)} - (Ca_{mit} - Ca_{cyt}) * k_{20} \frac{[Ca_{cyt}]}{([Ca_{cyt}] + K_{21})}
 \end{aligned}$$

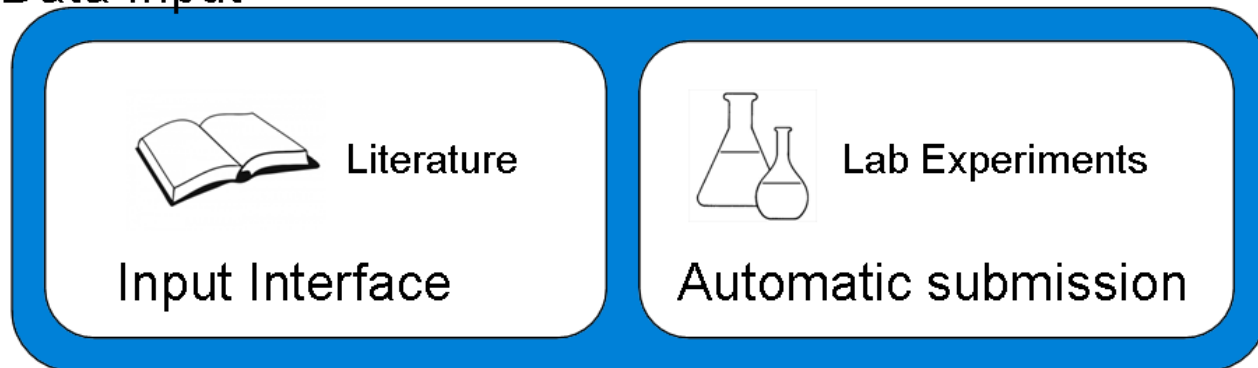
Model

Simulation

Prediction



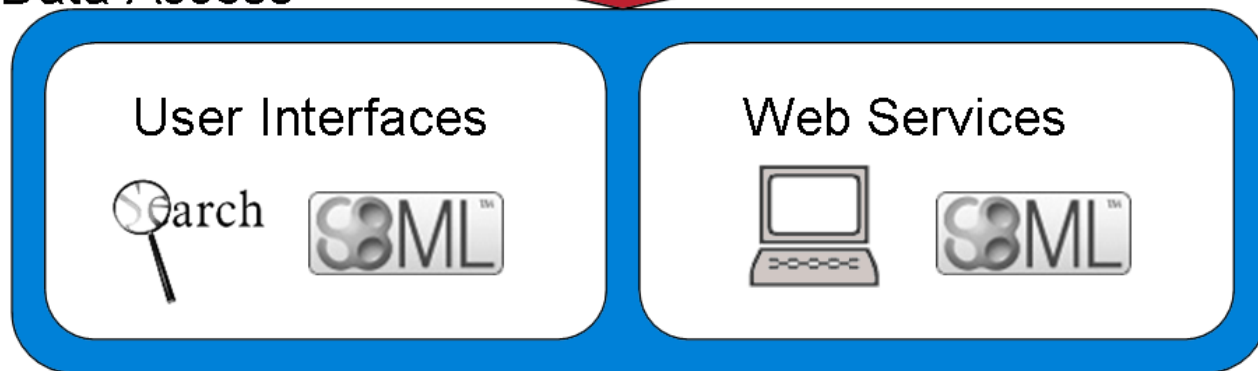
Data Input

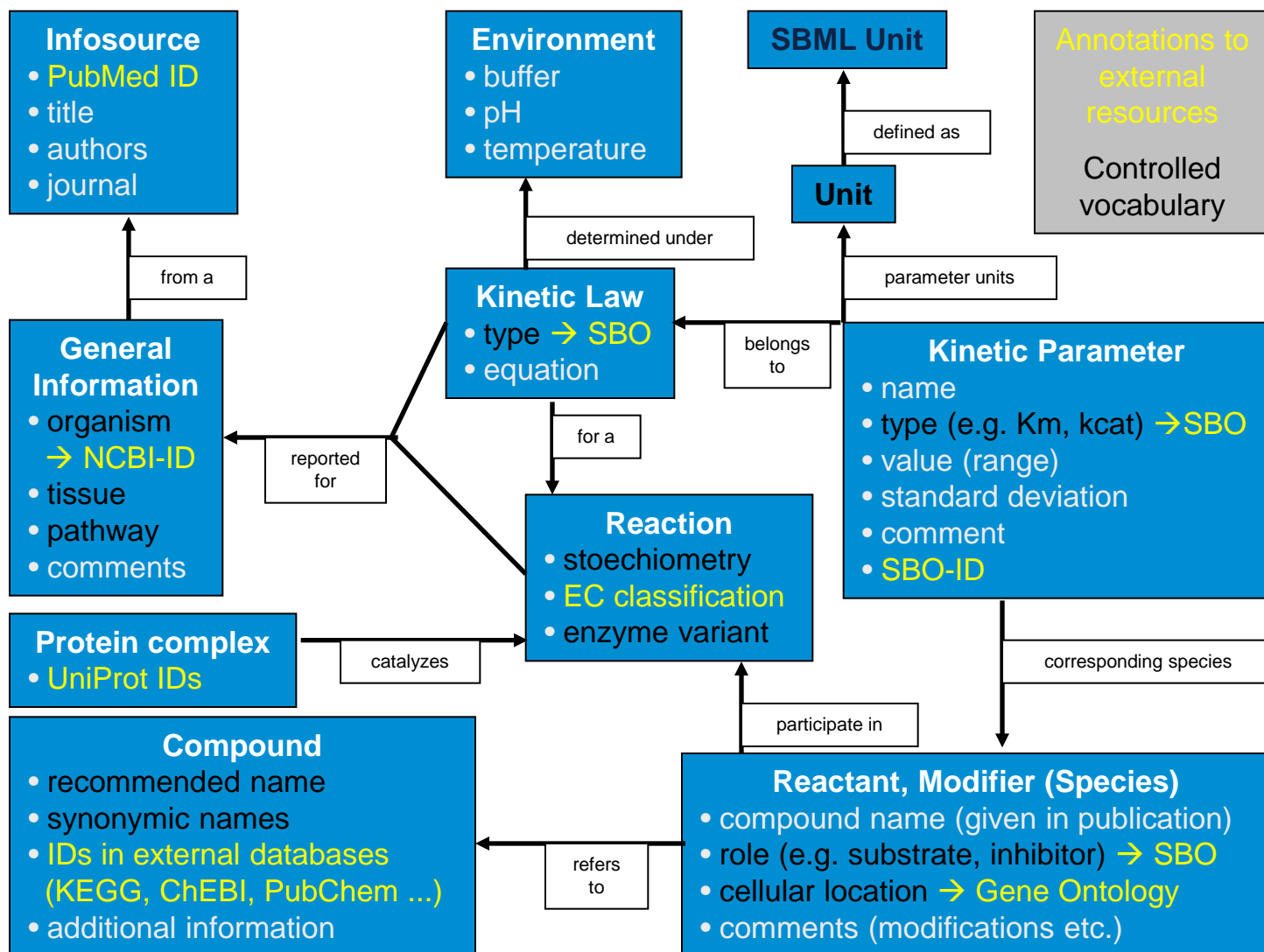


Curation /Annotation



Data Access






SABIO-RK News

- **SABIO-RK: New user interface**, including new search features (e.g. Inhibitors/activators/cofactors as ligands)
- **Ontology/Taxonomy based search**, e.g. for organism (NCBI taxonomy) or tissue (Brenda tissue ontology)
- **New (RESTful) Web Services** for programmatic access to SABIO-RK (currently implemented by platforms like JWS online or Virtual Cell)
- **Access restriction** for sensitive data
- **Extended information** in exported SBML

New user interface

Hi anonymousUser! [Log Out](#)

Heidelberg Institute for Theoretical Studies 

Reaction Search

HOME | CONTACT | HELP | IMPRINT | ADMINISTRATION | LOGIN

Using

☒ Within Current Result?

[Send Selected Entries to SBML File](#)

Rows/page: 15

Other Reaction View Entry View

Product

- Reaction
- Reaction Participant
- Enzyme
- Tissue
- Organism
- Publication
- Parameter
- Cellular Location
- Pathway
- Signalling
- Kinetic Mechanism
- Type
- Entry Date

Any Role

- Substrate
- Product
- Inhibitor
- Catalyst
- Cofactor
- Activator
- Other Modifier

Reactants

☒ Product

☒ Glycocholate

Enzymes

☒ EC Numbers

2.3.1.65

Tissues

☒ liver (BTO)

Reaction	Enzyme		Tissue	Organism	Parameters (besides concentration)	Environment	
	Protein	Enzyme Variant (w)ildtype (m)utant				°C	pH
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	liver	Homo sapiens	Vmax Km	37	8.4
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q63276	liver	Rattus norvegicus	Vmax Km	30	7.2
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	liver	Homo sapiens	Vmax Km	37	7
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	liver	Homo sapiens	Vmax Km	37	7
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	liver	Homo sapiens	Vmax Km	37	8.25
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	liver	Homo sapiens	Vmax Km	37	8.25
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	liver	Homo sapiens	Vmax Km	37	8
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	liver	Homo sapiens	Vmax Km	37	8
Glycine + Cholesteryl-CoA = Coenzyme A + Glycocholate	2.3.1.65		liver	Bos taurus	Km Ki	30	8

Pages: [1]

[Send Selected Entries to SBML File](#)

[Send Selected Entries to SBML File](#)

Your Current

Criteria:

[RESET](#)
☒ Reactants

☒ Product

Glycocholate ☒
☒ Enzymes

☒ EC Numbers

2.3.1.65 ☒
☒ Tissues

liver (BTO) ☒

Pages:[1]

There are 9 Kinetic Law Entries

	Reaction	Enzyme			Tissue	Organism
		ECNumber	Protein	Enzyme Variant (w)ildtype (m)utant		
<input checked="" type="checkbox"/>	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	wildtype (w)	liver	Homo sapiens

Entry ID: 37536

General information		
Organism	Homo sapiens	
Tissue	liver	
EC Class	2.3.1.65	
SABIO reaction id	3479	
Variant	wildtype	

Substrates		
name	location	comment
Glycine	cytosol	-
Choloyl-CoA	cytosol	-

Products		
name	location	comment
Coenzyme A	cytosol	-
Glycocholate	cytosol	-

Modifiers				
name	location	effect	comment	protein complex
Glycine N-choloyltransferase(Enzyme)	cytosol	Modifier-Catalyst	-	Q14032 ;

Enzyme (protein data)				
	UniProt-ID	name	mol. weight (kDa)	deviation (kDa)
subunit	Q14032	-	-	-
complex	-	-	50.0	2.0

Kinetic Law	
type	formula
Michaelis-Menten	$V_{max} \cdot A / (K_m + A)$

Parameter								
name	type	species	start val.	end val.	deviat.	unit	comment	
A	concentration	Glycine	0.25	10.0	-	mM	-	
B	concentration	Choloyl-CoA	1.15	-	-	mM	saturating	
Vmax	Vmax	-	0.77	-	-	$\mu\text{mol}/(\text{min} \cdot \text{mg})$	purified enzyme	
Km	Km	Glycine	5.8	-	-	mM	-	

Experimental conditions			
	start value	end value	unit
pH	8.4	-	-
temperature	37.0	-	°C
buffer	100 mM Potassium phosphate		


General comment	
-----------------	--

<input checked="" type="checkbox"/>	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q63276	wildtype (w)	liver	Rattus norvegicus
<input checked="" type="checkbox"/>	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	wildtype (w)	liver	Homo sapiens

Please Select Attribute

☐ Within Current Result?Using 

Select: 1 entries

Rows/page: 15 

2

[Send Selected Entries to SBML File](#)















Other

Reaction View

Entry View

Pages:[1]

There are 7 Kinetic Law Entries

	Reaction	Enzyme			Tissue	Organism	Parameters (besides concentration)	Environment		
		ECNumber	Protein	Enzyme Variant (w)ildtype (m)utant				°C	pH	
	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	wildtype (w)	liver	Homo sapiens	Vmax Km	37	8.4	
	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	wildtype (w)	liver	Homo sapiens	Vmax Km	37	7	
	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	wildtype (w)	liver	Homo sapiens	Vmax Km	37	7	
	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	C235S (m) mutant (m)	liver	Homo sapiens	Vmax Km	37	8.25	
	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	wildtype (w)	liver	Homo sapiens	Vmax Km	37	8.25	
	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	wildtype (w)	liver	Homo sapiens	Vmax Km	37	8	
	Glycine + Choloyl-CoA = Coenzyme A + Glycocholate	2.3.1.65	Q14032	wildtype (w)	liver	Homo sapiens	Vmax Km	37	8	

1

SABIO-RK SBML Export

Entry ID: 20147

Substrates							
name	location		comment				
Glycine	cytosol		-				
Choloyl-CoA	cytosol		-				
Products							
name	location		comment				
Coenzyme A	cytosol		-				
Glycocholate	cytosol		-				
Modifiers							
name	location	effect	comment	protein complex			
Glycine N-choloyltransferase(Enzyme)	cytosol	Modifier-Catalyst	-	Q14032 ;			
Enzyme (protein data)							
	UniProt-ID	name	mol. weight (kDa)		deviation (kDa)		
subunit	Q14032	-			-		
complex	-	-	50.0				
Kinetic Law							
type			formula				
Michaelis-Menten			$V_{max} * B / (K_m + B)$				
Parameter							
name	type	species	start val.	end val.	deviat.	unit	comment
B	concentration	Glycine	0.0	20.0	-	mM	-
Km	Km	Glycine	0.02	-	-	M	-
A	concentration	Choloyl-CoA	200.0	300.0	-	μM	-
Vmax	Vmax	-	15.3	-	-	nmol/(min*mg)	cytosolic protein
E	concentration	Enzyme	0.2	-	-	mg/ml	-
Experimental conditions							
	start value		end value		unit		
temperature	37.0				°C		
pH	8.0				-		
buffer	50 mM potassium phosphate						
comment	-						
Reference							
title	author	year	journal	volume	pages	PubMed	
Subcellular organization of bile acid amidation in human liver: a key issue in regulating the biosynthesis of bile salts	Solaas K, Ulvestad A, Soreide O, Kase BF	2000	J Lipid Res	41	1154-62	10884298	

Pages:[1]



2

Send Selected Entries to SBML File

Search



[Save Model](#)

Enter name of model:

Export parameters normalized to SI base units ☒

[Save Model on Disk as SBML](#)

[Back to Results](#)



[Sample File](#)

– Search criteria:

Reactant:

– Pathway:

[Glycolysis classical](#)

Enzyme:

Publication:

Protein:

Sign. modific.:

Sign. event:

– Organism:

[Homo sapiens](#)

– Tissue:

[liver](#)

Cell. loc.:

Exp. cond.:

Kin. data:

SABIO-RK SBML Export

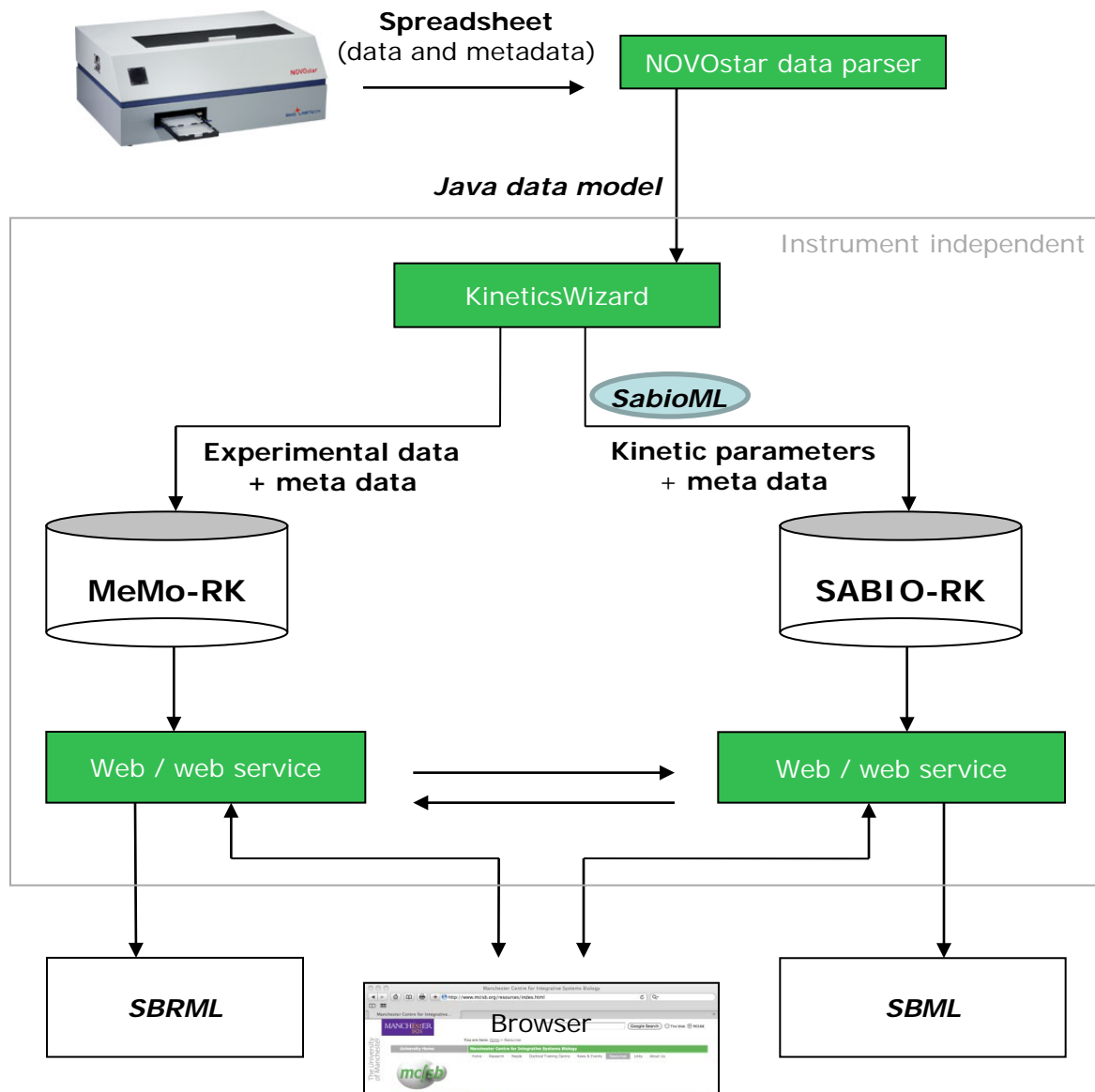
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          (http://sabiork.h-its.org/)
          (c) 2005-2010 HITS gGmbH http://www.h-its.org
        </p><br/>
        To cite SABIO-RK Database, please use
        "http://www.ncbi.nlm.nih.gov/pubmed/17822389"
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        Wittig U., Golebiewski M., Kania R., Krebs O., Mir S., Weidemann A., Anstein S., Sar
        Lecture Notes in Computer Science, 4075: 94-103.
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- Currently up to **SBML Level 2 Version 4**
- **Reaction Kinetics Warehouse:**
Reactions, kinetic equations and parameters (with corresponding units) from different database entries can be exported in one SBML file
- Data is annotated (RDF and SBOterms) according to **MIRIAM**
- Annotations include **SABIO-RK Ids** (reaction and kineticlaw) for tracking
- Optional **normalization of kinetic parameters** to SI base units
- Model can also be exported as human readable PDF → **SBML2LaTeX**





Enzyme kinetics informatics: from instrument to browser

Neil Swainston^{1,†}, Martin Golebiewski^{2,†},
Hanan L. Messiha¹, Naglis Malys¹, Renate
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Ulrike Wittig², Douglas B. Kell¹, Pedro
Mendes^{1,3}, Wolfgang Müller², Norman W.
Paton¹, Isabel Rojas²

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FEBS

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Author Information

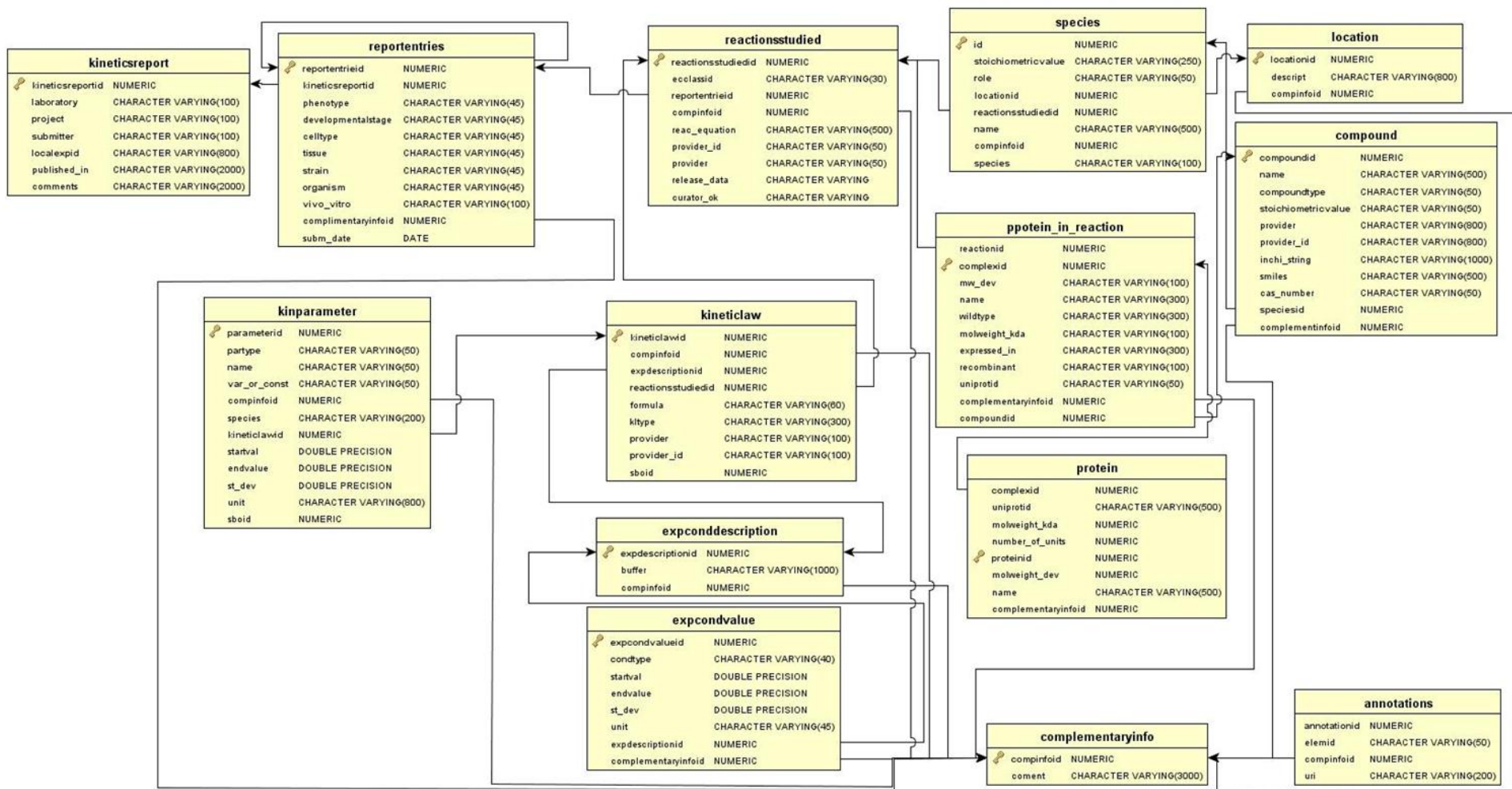
- ¹ Manchester Centre for Integrative Systems Biology, University of Manchester, UK
- ² Heidelberg Institute for Theoretical Studies, Germany
- ³ Virginia Bioinformatics Institute, Virginia Tech, Blacksburg, VA, USA

*Correspondence: N. Swainston, Manchester Centre for Integrative Systems Biology, University of Manchester, Manchester M1 7DN, UK Fax: +44 161 306 8918 Tel:  +44 161 306 5146  E-mail: neil.swainston@manchester.ac.uk Website: <http://www.mcisb.org>

†*

These authors contributed equally to this work

SABIOML: Exchange Format for Experimental Kinetic Data



SABIOML: Exchange Format for Experimental Kinetic Data

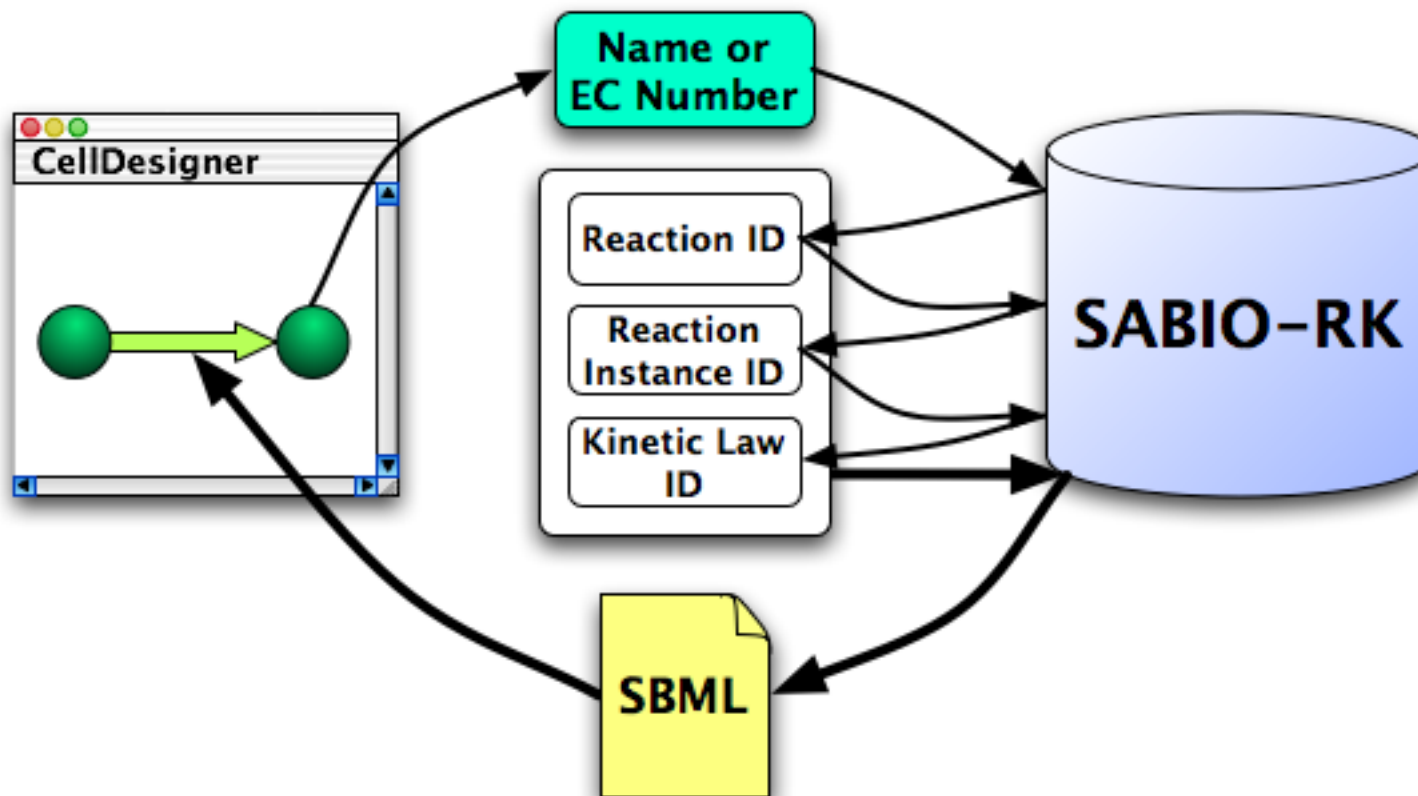
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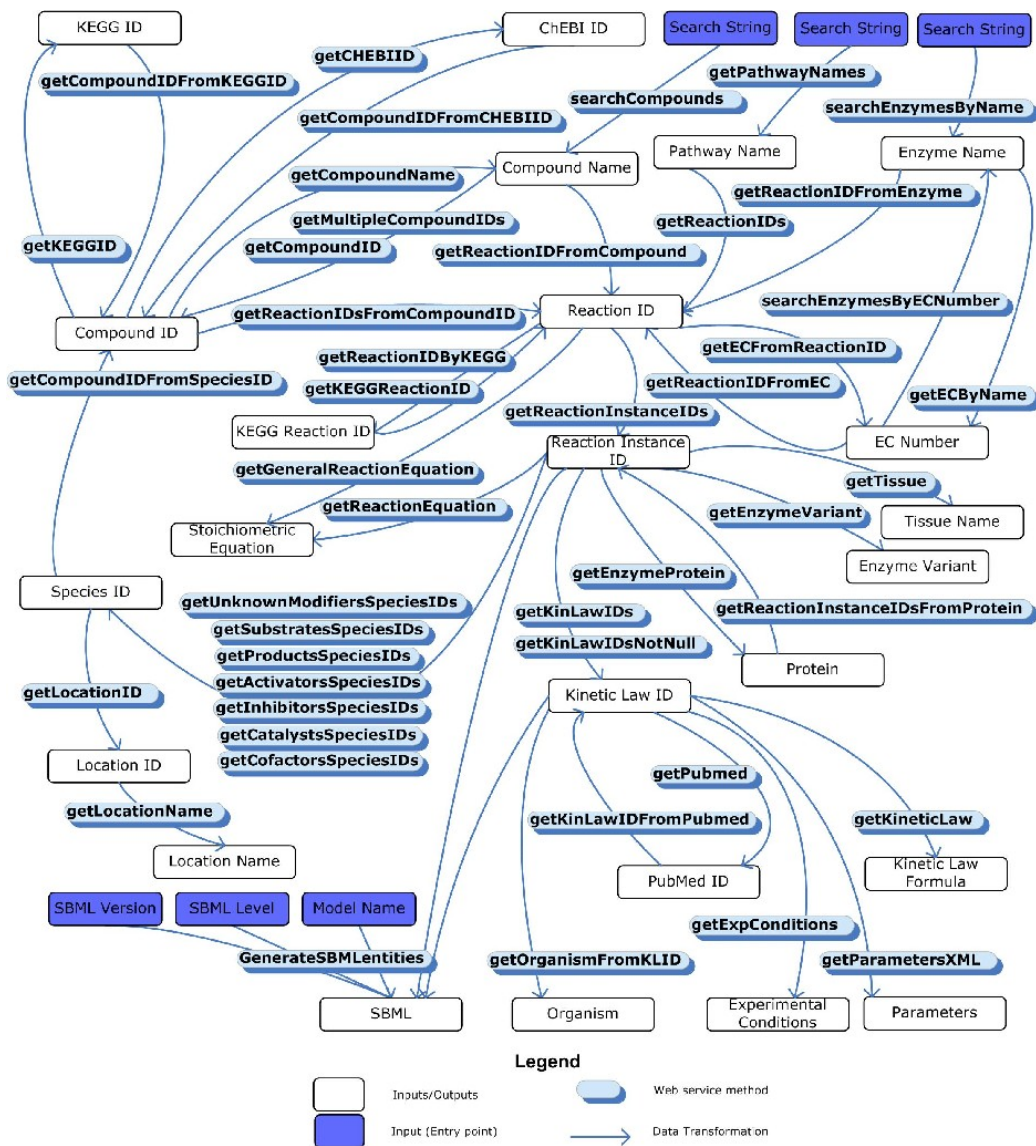

SABIO-RK API Access

Integration into Modeling Platforms

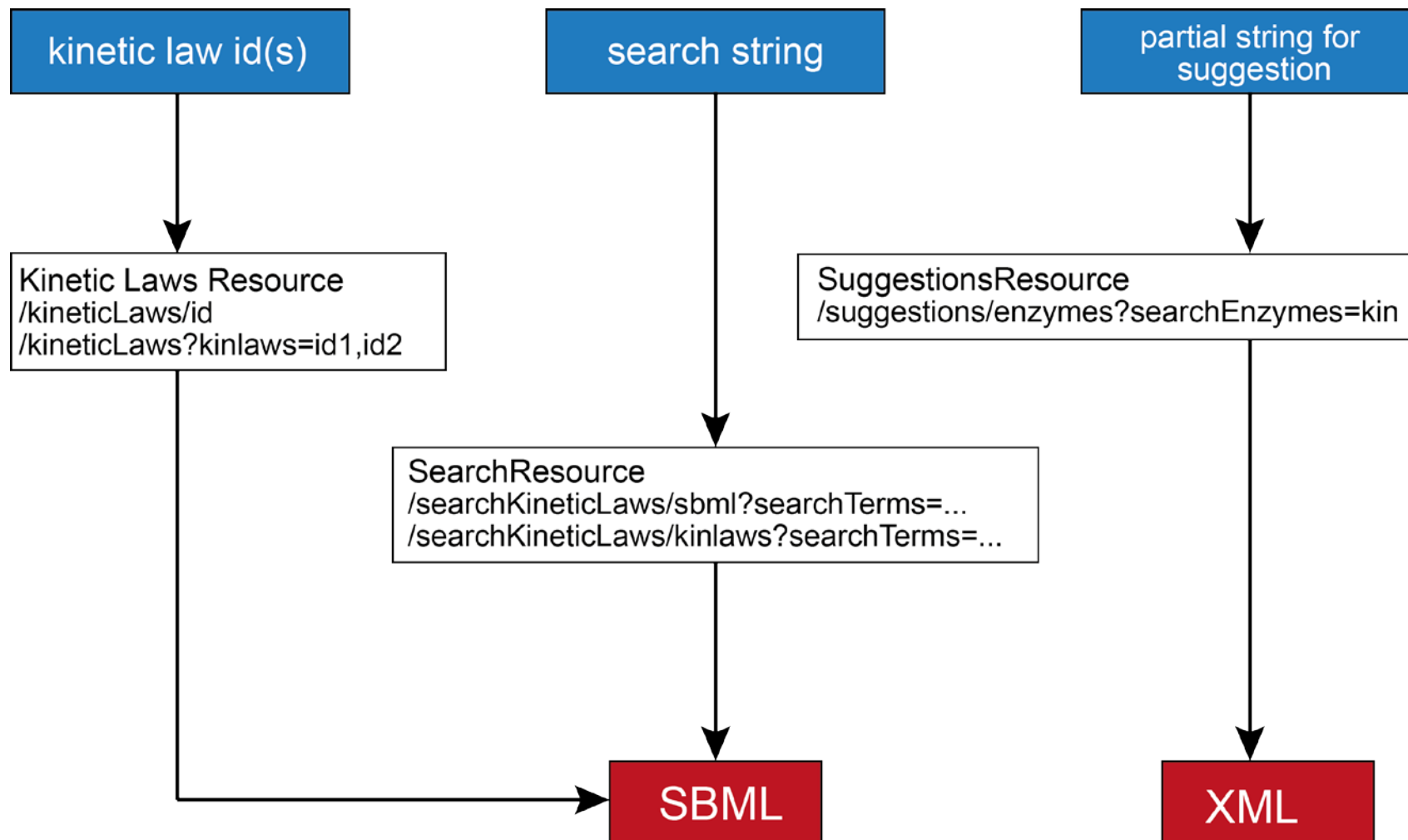


<http://www.celldesigner.org>

SABIO-RK API Access (Web Services)



- JAW-WS based
- Integration possible in modeling platforms or simulation tools (e.g. CellDesigner)
- Cross-linking with other databases (e.g. ChEBI)
- Data export in **SBML** supported



Example requests:

Entries may be requested directly if the database entry ID is known

<http://sabio.h-its.org/sabioRestWebServices/kineticLaws/20147>

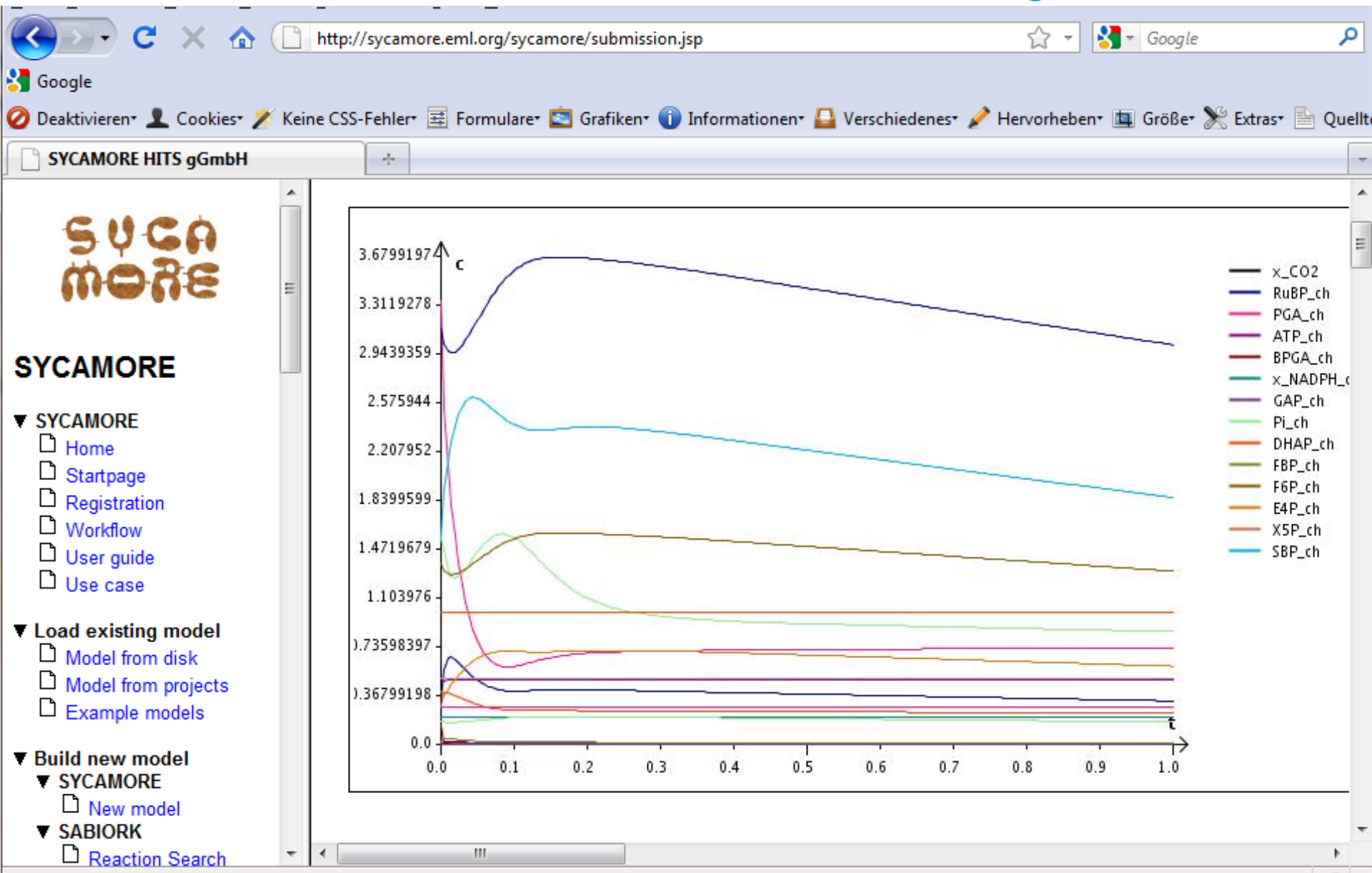
Entries may be searched for using the same search options available in the browser search interface

<http://sabio.h-its.org/sabioRestWebServices/searchKineticLaws/sbml?searchTerms=ORGANISM=Homo sapiens;TISSUE=liver>

Suggestions for search terms can be done

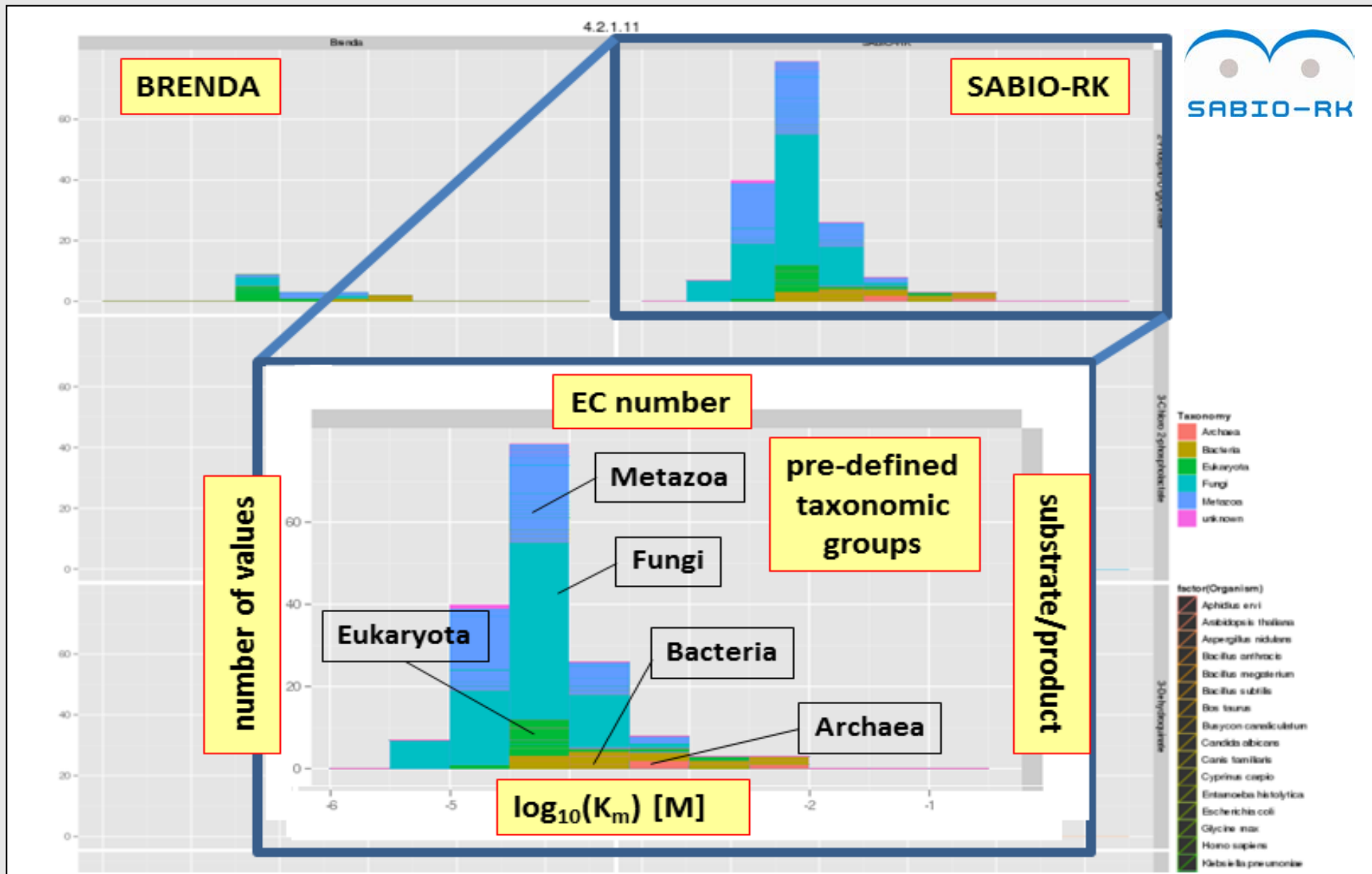
<http://sabio.h-its.org/sabioRestWebServices/suggestions/compounds?searchCompounds=glycoch>

Seamless Data Transfer to SYCAMORE for Modelling



SABIO-RK Content

<i>Organism</i>	Entries (total)	Mutants	Reactions (distinct)	EC number (distinct)	Reaction rates	Km	Kinetic laws
Total 660	37343	10616	4951	1070	27305	29852	17673
Homo sapiens	7028	2156	1233	348	5353	5757	3738
Rattus norvegicus	4286	759	821	285	2723	3188	2178
Escherichia coli	3828	1912	577	188	2976	3155	1988
Saccharomyces cerevisiae	1518	451	220	76	1044	1300	805
Bos taurus	982	59	276	86	579	722	471
Oryctolagus cuniculus	849	125	180	50	450	626	322
Sus scrofa	842	295	158	68	669	434	750
Mus musculus	809	92	236	92	456	290	603
Enterococcus faecalis	538	42	155	51	320	157	478
Gallus gallus	487	167	78	35	405	521	217







<http://sabio.h-its.org>

