SBGN-ED – working with the Systems Biology Graphical Notation

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Overview

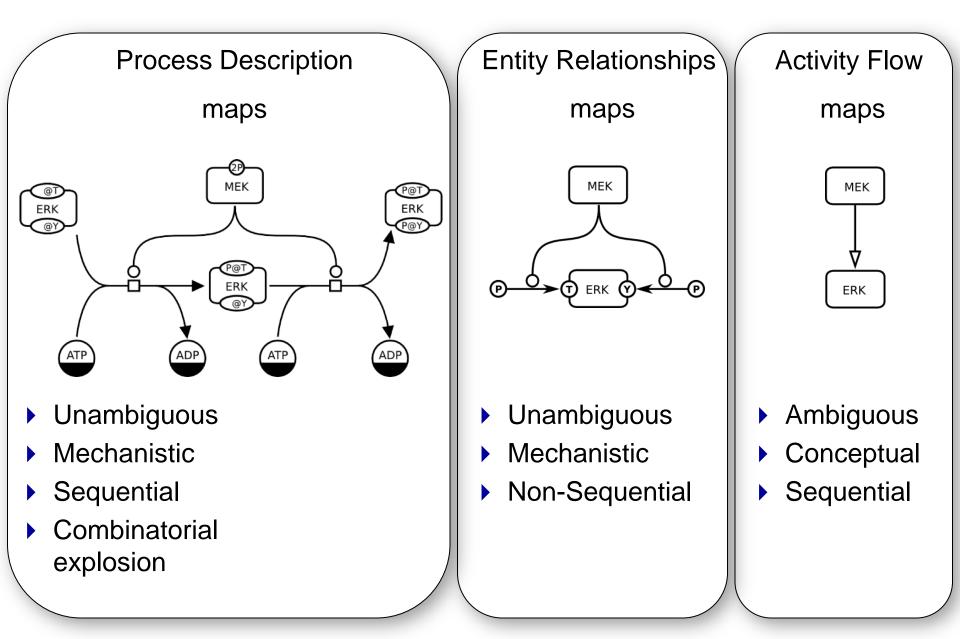
- SBGN
- SBGN-ED and VANTED
- Working with SBGN-ED
 - Installation
 - Editing
 - Validation
 - Import / Export
 - Translation
 - Additional features from VANTED
- Outlook
- Hands on

What is SBGN

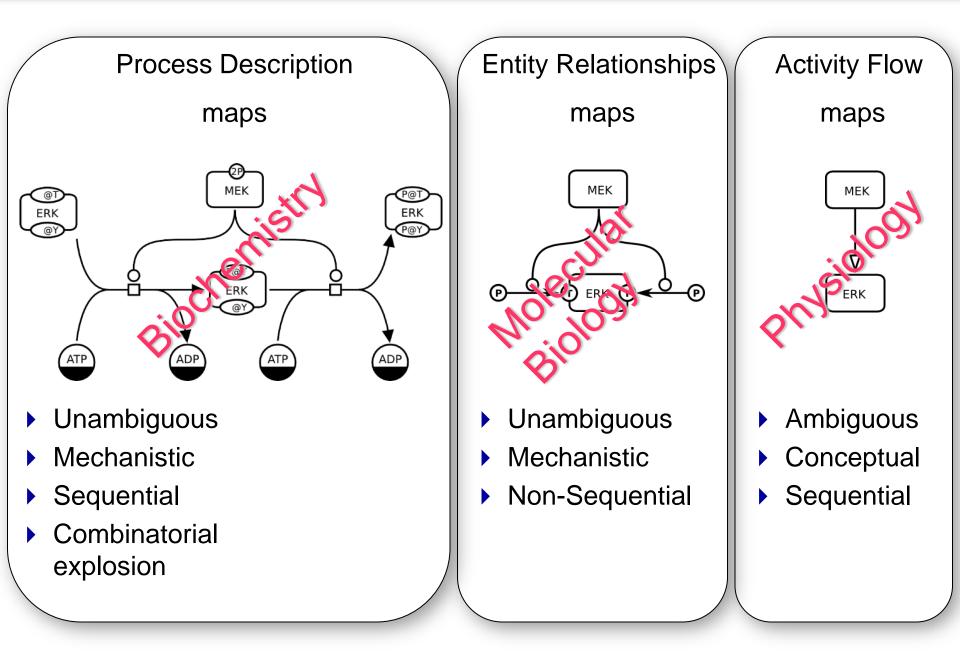
- Visualisation of biological information
- A way to unambiguously describe biochemical and cellular events in graphs
- Can graphically represent quantitative models, biochemical pathways, at different levels of granularity
- Developed over six years by a growing community
- Limited amount of symbols (32) \rightarrow Smooth learning curve
- Three languages
 - Process Descriptions
 - Entity Relationships
 - Activity Flow

- \rightarrow one state = one glyph
- \rightarrow one entity = one glyph
- \rightarrow conceptual level

Graph Trinity: Three Languages in One

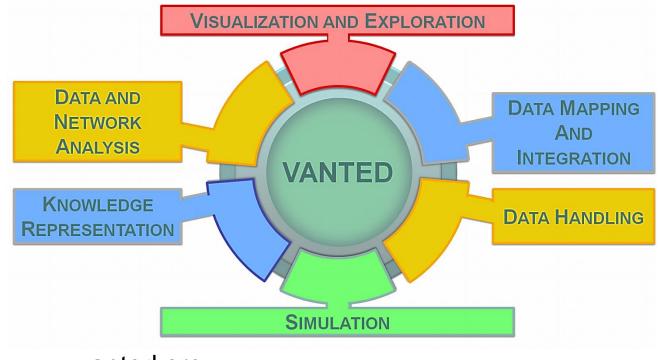


Graph Trinity: Three Languages in One



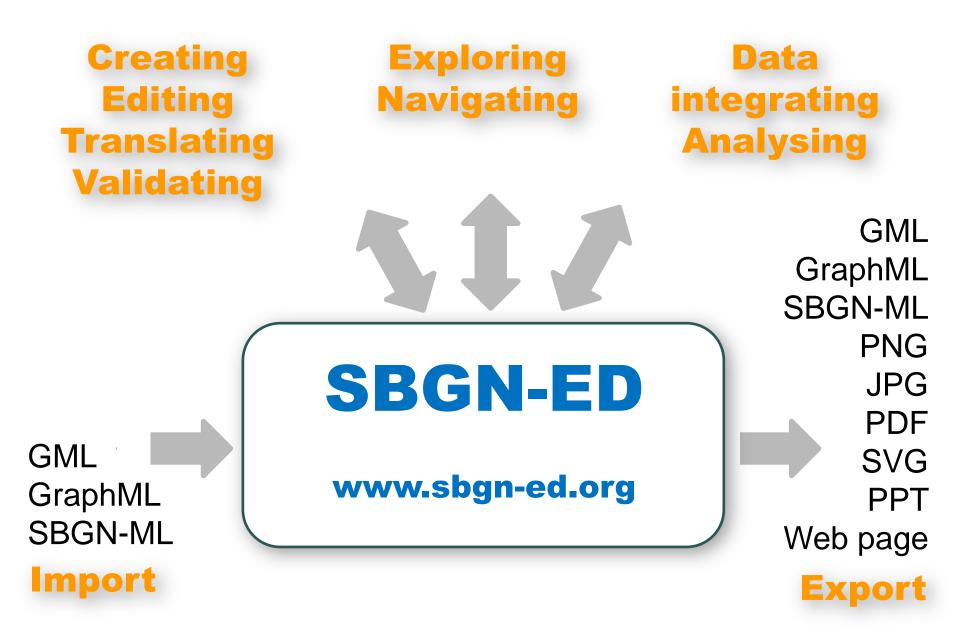
VANTED and SBGN-ED

- SBGN-ED is an add-on (extension) of VANTED
- VANTED framework
 - Integration, analysis and visual exploration of experimental data in the context of biological networks
 - Main functions



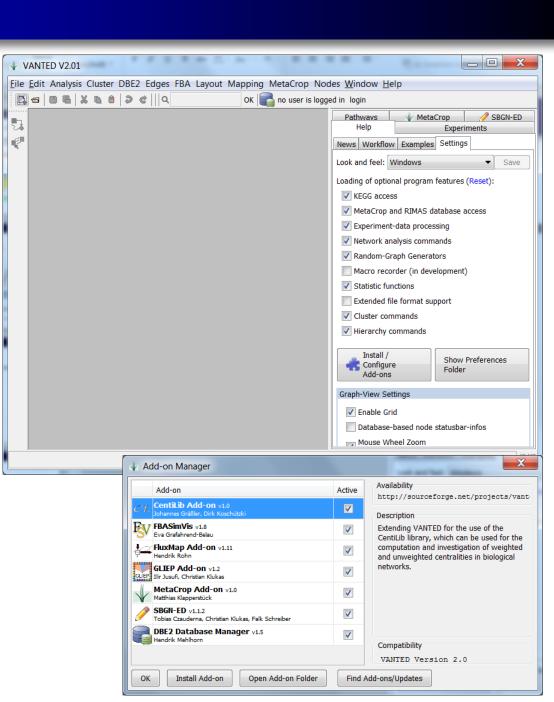
- www.vanted.org
- Open source (sourceforge)

VANTED and SBGN-ED



Installation

- 1. Install VANTED
 - 1. www.vanted.org
 - Webstart version or stand alone version (download)
- 2. Download SBGN-ED
 - 1. www.sbgn-ed.org
 - 2. Put it into arbitrary directory
- Help (tab) → Settings (subtab) → Install / Configure Add-ons
 - 1. Install from directory



SBGN-ED Tabs

- Three languages
 - Process Description (PD)
 - Entity Relationship (ER)
 - Activity Flow (AF)
- Tools
 - Validate
 - Translate
- Examples
- Other functionalities such as Import, Export, Layout, Data Mapping, Network Analysis etc are part of the underlying VANTED system and part of VANTEDs menus

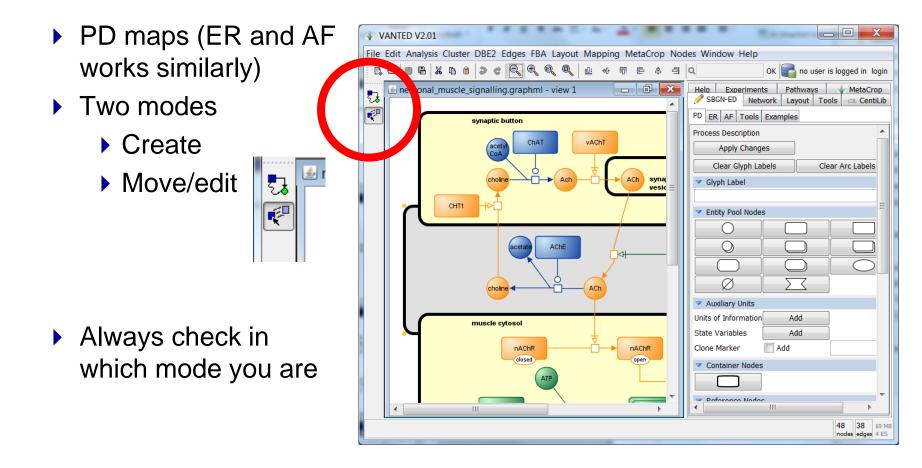
Help	Ext	periments
Pathways	MetaCrop	SBGN-ED
PD ER AF To	ols Examples	
Process Descript	ion	A
Apply Ch	anges	
Clear Glypt	n Labels	Clear Arc Labels
🔻 Glyph Label		
label		
Entity Pool N	odes	=
0		
Ø		
 Auxiliary Unit 	ts	
Units of Informa	tion Add	
State Variables	Add	
Clone Marker	Add	marker
Container No	des	
Reference No	odes	•
•	111	4

Import

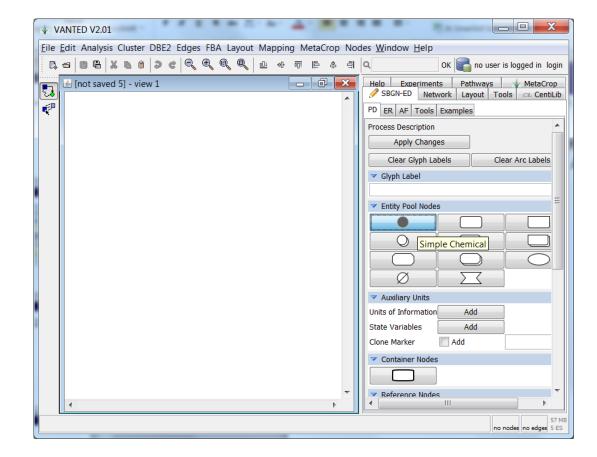
- Files
 - File (menu) → Open
 - ▶ File formats: GML, GraphML, SBGN-ML, KGML
- Pathway database (Pathway tab)
 - MetaCrop
 - Choose pathway
- Examples
 - ▶ SBGN-ED \rightarrow Examples (tab)
 - Choose example

Creating and Editing SBGN Maps

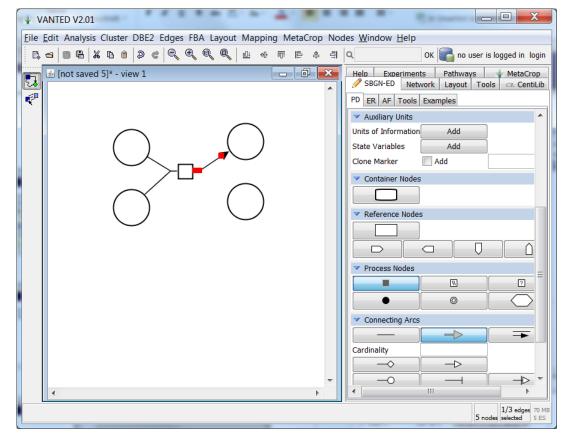
Star VANTED V2.0	A VANTED V20		
VANTED V2.0			
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[]	п с ок		Help Experiments Pathways SBGN-ED Network Layout Tools
💼 🕼 neuronal_muscle_signalling.graphml - view 1 👘 🛋	Help Experiments	Help Experiments Pathways SBGN-ED Network Layout Tools	
2a 🖉	Pathways SBGN-ED Network Layout Tools	PD ER AF Tools Examples	PD ER AF Tools Examples
synaptic button	PD ER AF Tools Examples	Entity Relationship	Activity Flow Apply Changes
ChAT VAChT	Process Description	Apply Changes	
	Apply Changes	Clear Glyph Labels Clear Arc Labels	Clear Glyph Labels
chaine Ach Ach synaptic vesicle	Clear Glyph Labels Clear Arc Labels	✓ Glyph Label	✓ Glyph Label
	✓ Glyph Label		
		✓ Interactors	Activity Nodes
active ACHE	✓ Entity Pool Nodes		Unit of Information
synaptic cleft		✓ Auxiliary Units	
synaput ciert		Units of Information Add	Add Add Add
		State Variables Add	
muscle cytosol		Add	Add Add
		Add	✓ Container Nodes
	✓ Auxiliary Units	~	
	Units of Information Add		Units of Information Add
	Clone Marker Add		▼ Reference Nodes
myosin Ca2+	✓ Container Nodes	Logical Operators	
acin (ATP)	Container Nodes		
	▼ Reference Nodes	✓ Statements	Modulating Arcs
		Cardinality	
		Unit of Information	
(ATP)	✓ Process Nodes		Logical Operators
muscle contraction		0	AND OR NOT
			5
	Connecting Arcs	▼ Influences	
	Cardinality	Unit of Information	
	✓ Logical Operators	✓ Reference Nodes	
			7 7 38 MB
		21 28 40 MB nodes edges 1 ES	nodes edges 1 ES
	48 38 46 MB nodes edges 1 ES	jinoues jeoges 1 ES	



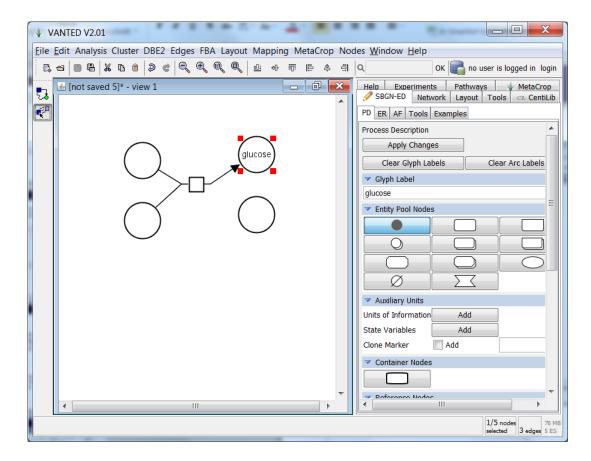
- 1. Select glyph
- 2. Click in canvas



- Step by step drawing of the map
- Information can either be added imidiatly or changed later

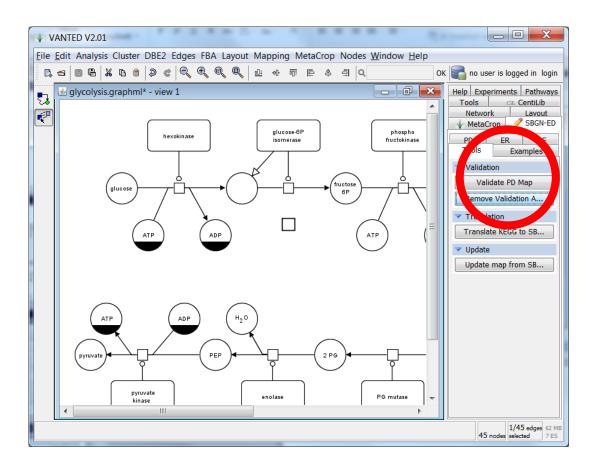


- Change elements later
 - 1. Change modus
 - 2. Add/change information within tab
 - 3. Click apply changes



Validation

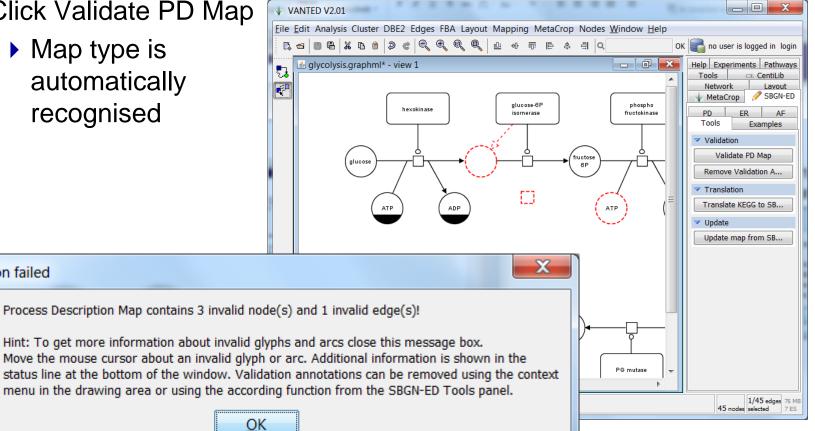
- In this example
 - Wrong edge
 - Missing clone marker
 - Unconnected node
 - Missing label



Validation failed

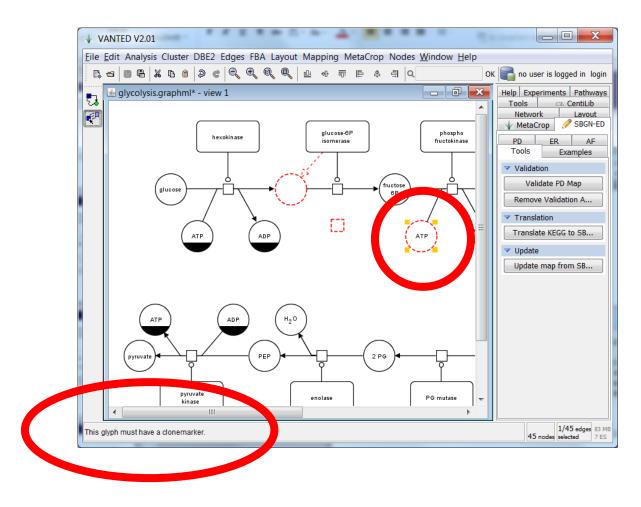


Map type is automatically recognised

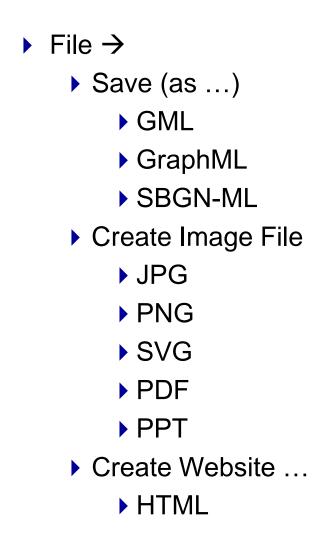


Validation

 Additional information



Export



* 1	VANTED V2.01	R & lowerse	
File	Edit Analysis Cluster DBE2 Edges FBA Layout Mapping MetaCrop Nodes Window	/ Help	
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9	New View Open	Strg+O	periments Pathways
	Save	Strg+S	ork Lavout
	Save as	Strg+Umschalt+S	ER AF
	Create Image File		Examples
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	Print Export Data	Strg+P	alidate PD Map
	Create Random Network		ove Validation A
6	SchemahiddenmappingLayoutGreyTransporterLabel.gml		slation
8	glycolysis_sbgn.gml		slate KEGG to SB
	glycolysis.gml		te
6	sample_foodweb.graphml		te map from SB
	Glycolysis_Gluconeogenesis_Pyruvate_metabolism_regulogs_arab_yeast_reduced.gml		
	Close	Strg+W	
	Exit	Alt+F4	
	PEP 2P6	_	
	PG muta	se 🔪 🗸	
			1/45 edges 65 MB 45 nodes selected 7 ES

Website Export

- Steps with several options for website design
- Clickable overview and clickable SBGN maps

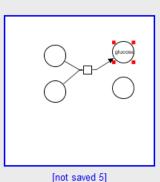
of the currently opened network files. Use the menu command Edit/Add Link to o	nmand to create a simple website, containing links to graphical representations create links assigned to network nodes, which point to other networks or to nd publish your own library of interlinked network files, possibly with
Please specify the title for the whole website	e and for each loaded network:
	character, the networks will be put into categories. etwork into the group 'A' sown with network name 'B'.
	<h1>My Pathway Website</h1>
Website-Title	
Graph [not saved 5]	[not saved 5]
Graph glycolysis.graphml	glycolysis
neuronal_muscle_signalling.graphml	neuronal_muscle_signalling
	OK

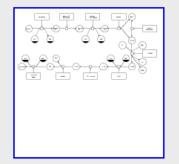
Overview

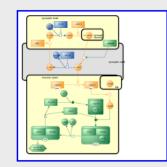
VANTED graph to website export. VANTED is available at http://vanted.ipk-gatersleben.de.

[not saved 5] glycolysis neuronal_muscle_signalling

Feedback





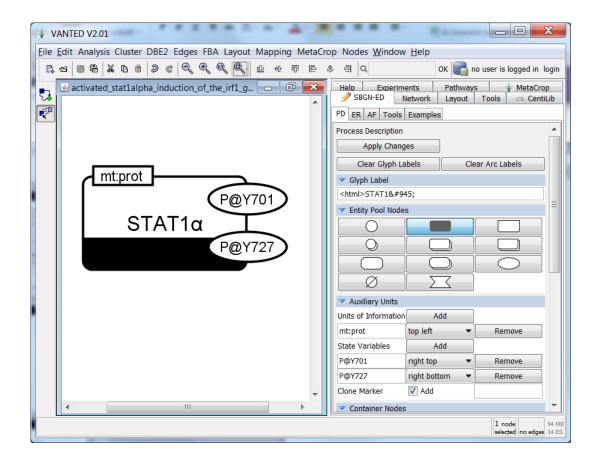


glycolysis

neuronal_muscle_signalling

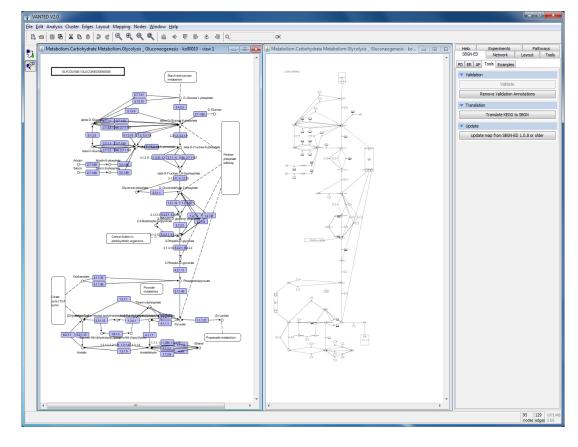
More Complicated Glyphs

- With unit of information (mt:prot)
- With state variable (P@Y701)
- With Clone markers
- HTML tags for text
- And so on



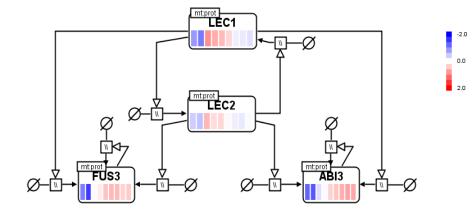
Translating SBGN Maps from KEGG

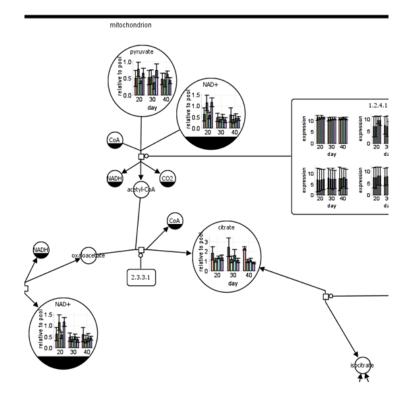
 From KGML files due to KEGG restrictions



Data Integrating and Analysing in SBGN Maps

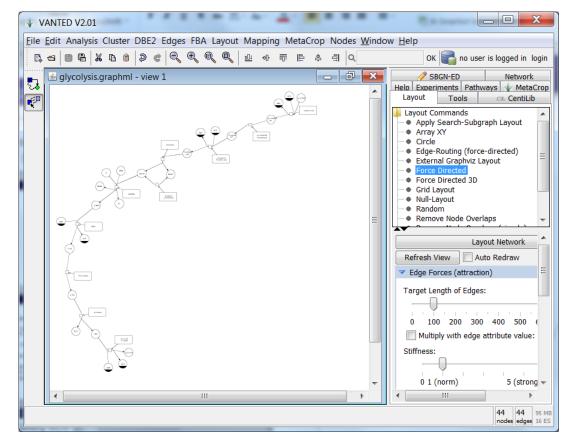
- VANTED provides many methods for
 - Data mapping and integration
 - Data and network analysis
- Also available in SBGN-ED





Layout

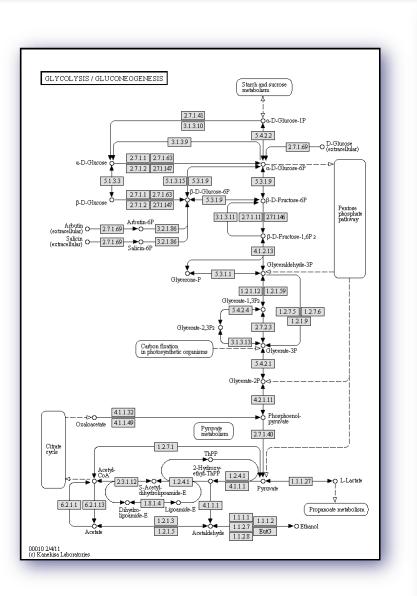
 VANTED provides many standard layout algorithm

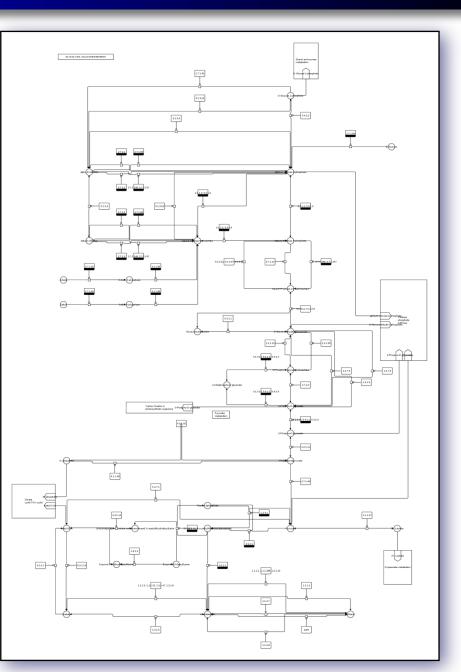


Outlook

- Next version (under development)
 - More importers (in VANTED)
 - ► SBML (all up to L3V1) (JSBML)
 - BioPAX (L2,L3) (Paxtools V4)
 - More export possibilities (in VANTED) (SBML and BioPAX) planed, but needs validation!
 - Constraint-based automatic layout
 - More interaction possibilities
 - Map translations

Outlook Screenshots





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