SBGN-ED – working with the Systems Biology Graphical Notation

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Overview

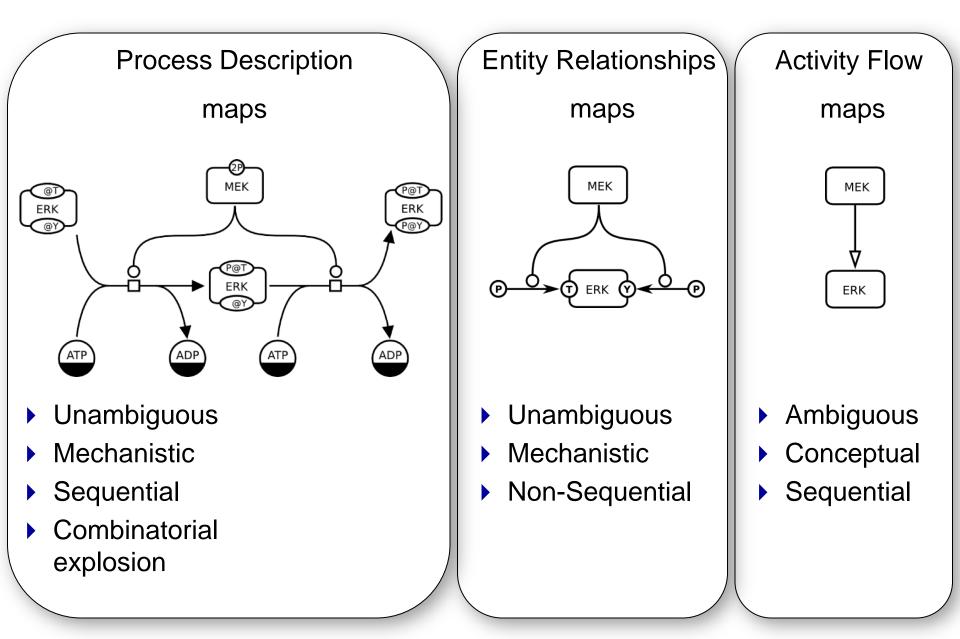
- SBGN
- SBGN-ED and VANTED
- Working with SBGN-ED
 - Installation
 - Editing
 - Validation
 - Import / Export
 - Translation
 - Additional features from VANTED
- Outlook
- Hands on

What is SBGN

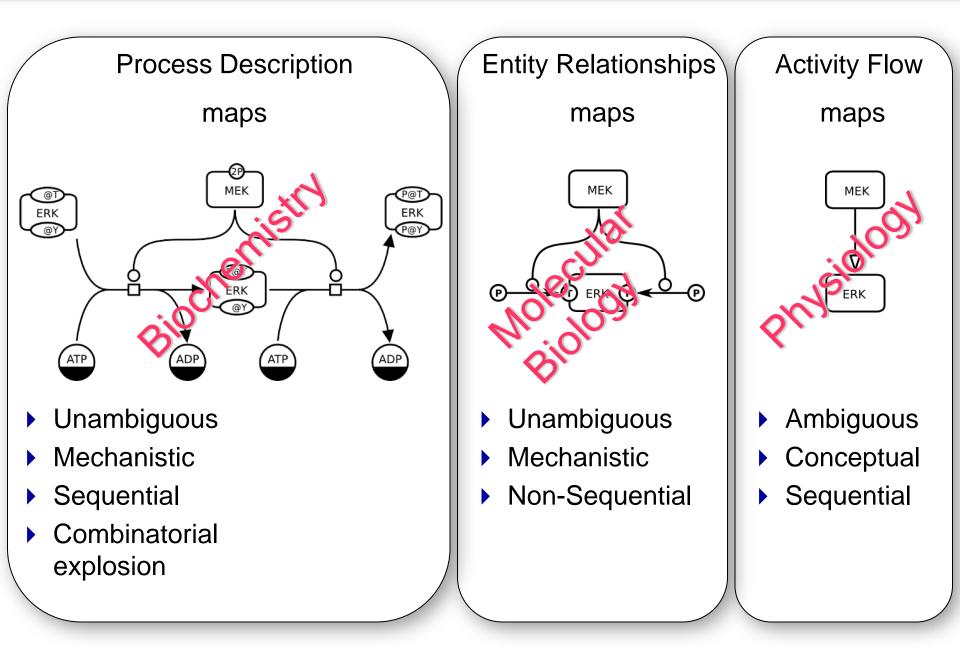
- Visualisation of biological information
- A way to unambiguously describe biochemical and cellular events in graphs
- Can graphically represent quantitative models, biochemical pathways, at different levels of granularity
- Developed over six years by a growing community
- Limited amount of symbols (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

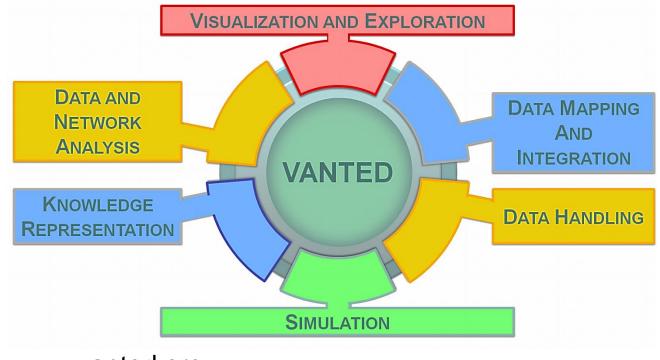


Graph Trinity: Three Languages in One



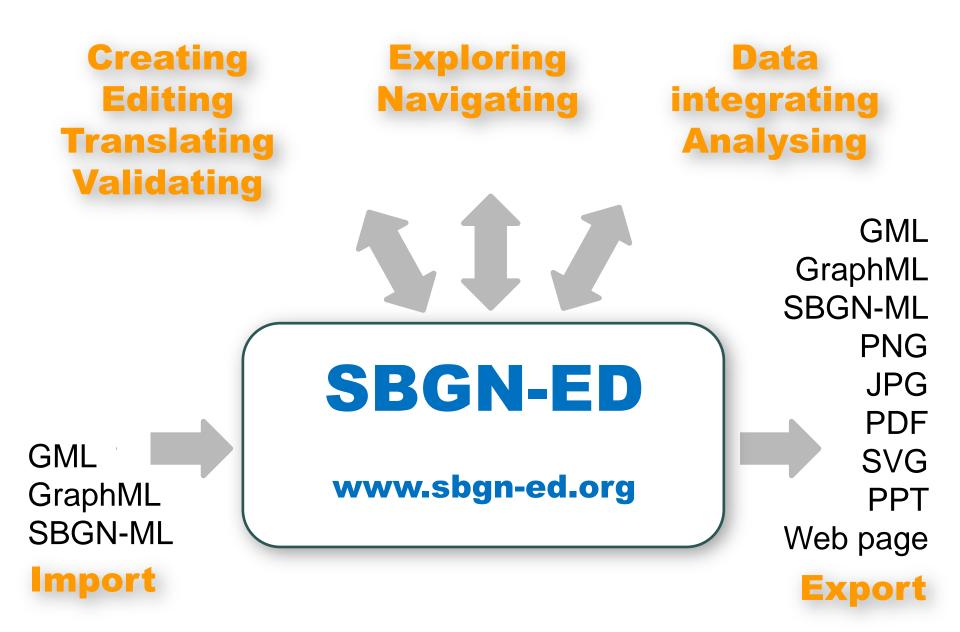
VANTED and SBGN-ED

- 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



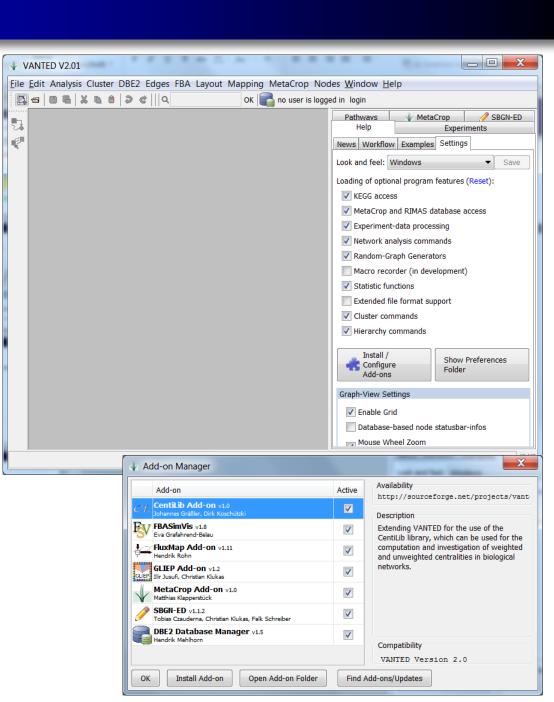
- www.vanted.org
- Open source (sourceforge)

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

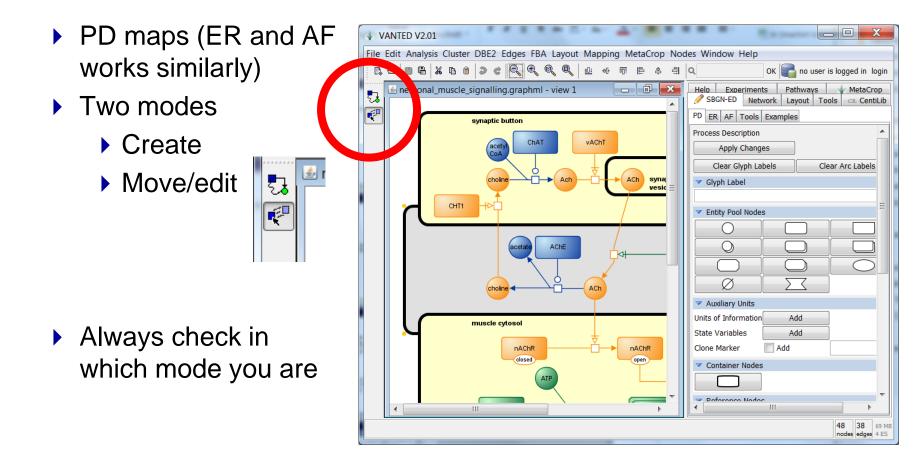
| Help | Ext | periments |
|------------------------------------|--------------|------------------|
| Pathways | MetaCrop | SBGN-ED |
| PD ER AF To | ols Examples | |
| Process Descript | ion | A |
| Apply Ch | anges | |
| Clear Glypt | n Labels | Clear Arc Labels |
| 🔻 Glyph Label | | |
| label | | |
| 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

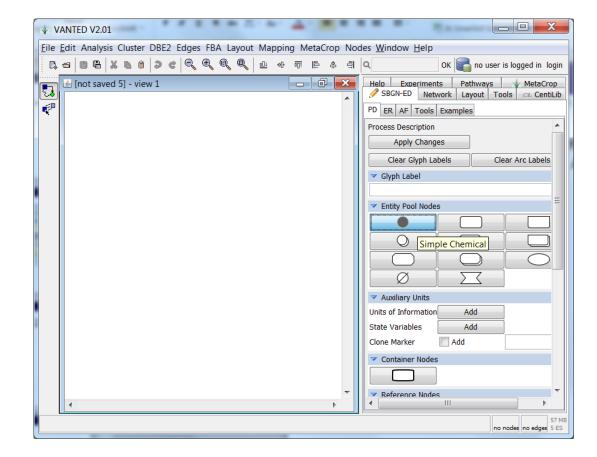
- Files
 - File (menu) → Open
 - ▶ File formats: GML, GraphML, SBGN-ML, KGML
- Pathway database (Pathway tab)
 - MetaCrop
 - Choose pathway
- Examples
 - ▶ SBGN-ED \rightarrow Examples (tab)
 - Choose example

Creating and Editing SBGN Maps

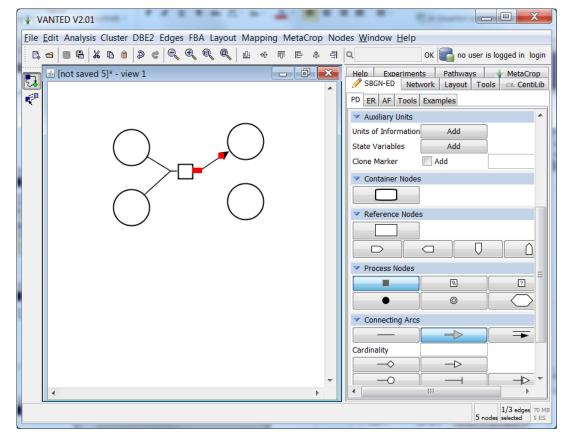
| Star VANTED V2.0 | A VANTED V20 | | |
|---|---------------------------------------|--|--|
| VANTED V2.0 | | | |
| Eile Edit Analysis Cluster Edges Layout Mapping Nodes Window Help | | а | ОК |
| [] | п с ок | | Help Experiments Pathways SBGN-ED Network Layout Tools |
| 💼 🕼 neuronal_muscle_signalling.graphml - 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 | |



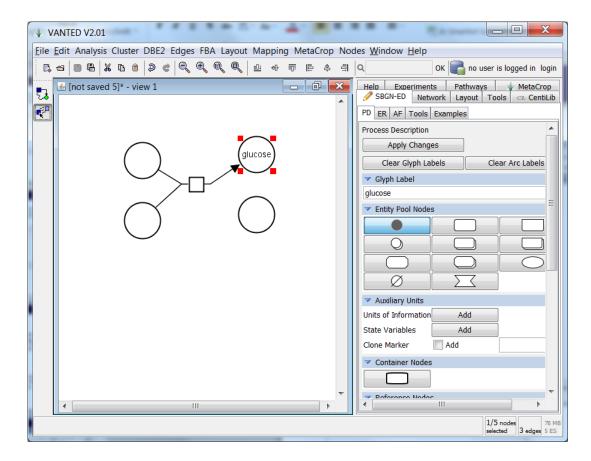
- 1. Select glyph
- 2. Click in canvas



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

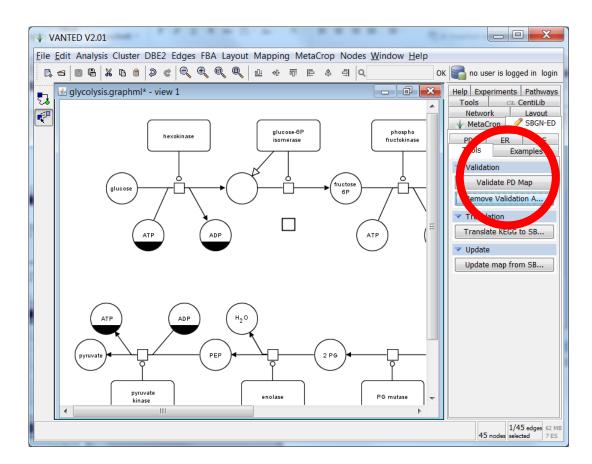


- Change elements later
 - 1. Change modus
 - 2. Add/change information within tab
 - 3. Click apply changes



Validation

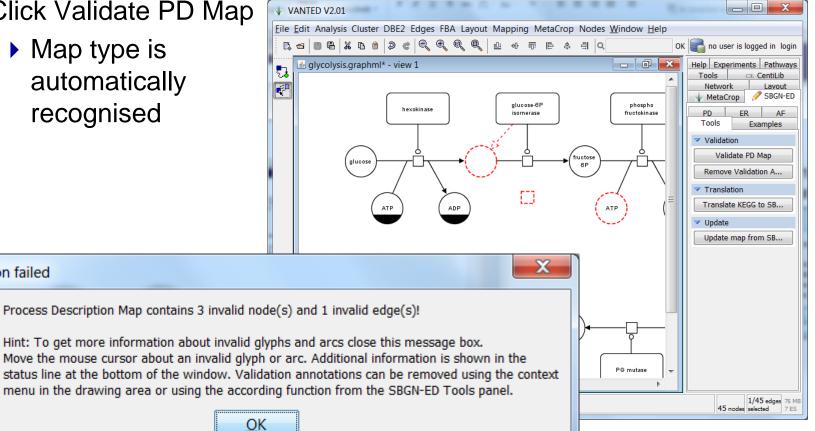
- In this example
 - Wrong edge
 - Missing clone marker
 - Unconnected node
 - Missing label



Validation failed

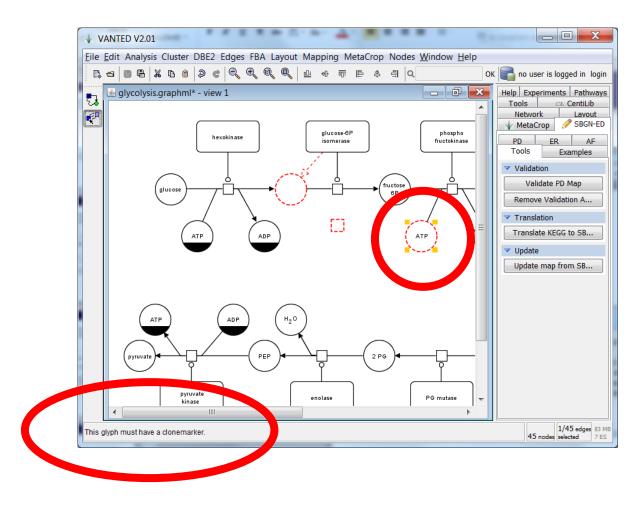


Map type is automatically recognised

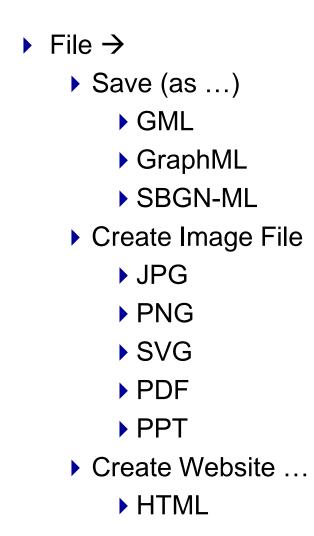


Validation

 Additional information



Export



| * 1 | VANTED V2.01 | R & lowerse | |
|----------|--|-----------------|--|
| File | Edit Analysis Cluster DBE2 Edges FBA Layout Mapping MetaCrop Nodes Window | / Help | |
| I | New | Strg+N | iser is logged in login |
| 9 | New View Open | Strg+O | periments Pathways |
| | Save | Strg+S | ork Lavout |
| | Save as | Strg+Umschalt+S | ER AF |
| | Create Image File | | Examples |
| 0 | Create Website | c | ation |
| | 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

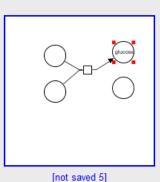
| of the currently opened network files. Use the menu command Edit/Add Link to o | nmand to create a simple website, containing links to graphical representations create links assigned to network nodes, which point to other networks or to nd publish your own library of interlinked network files, possibly 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 |

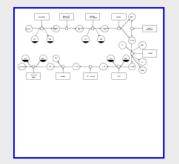
Overview

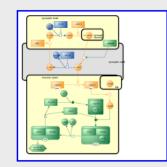
VANTED graph to website export. VANTED is available at http://vanted.ipk-gatersleben.de.

[not saved 5] glycolysis neuronal_muscle_signalling

Feedback





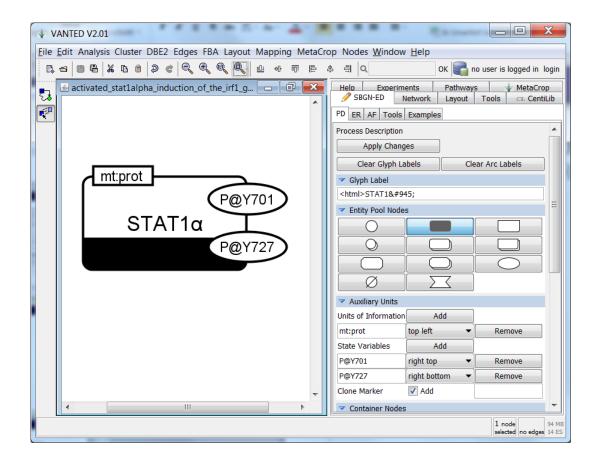


glycolysis

neuronal_muscle_signalling

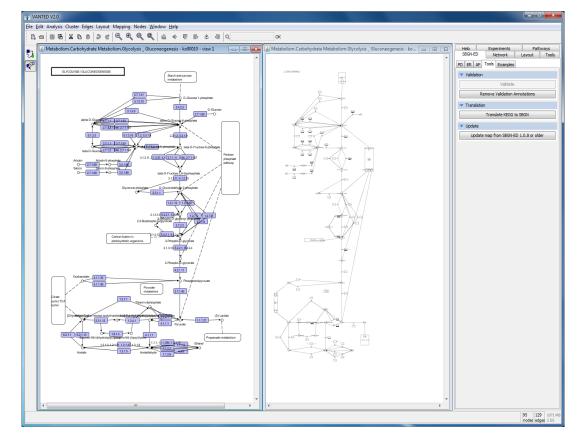
More Complicated Glyphs

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



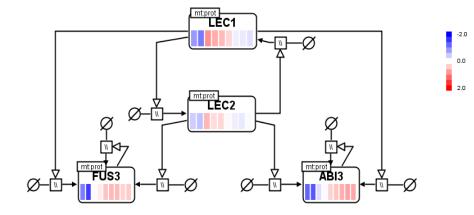
Translating SBGN Maps from KEGG

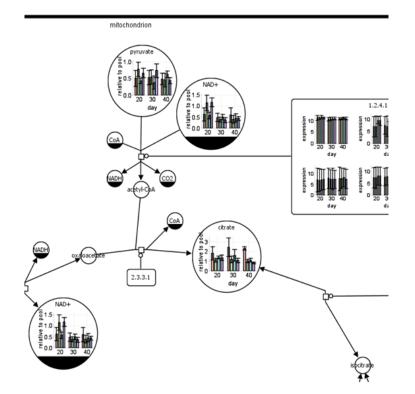
 From KGML files due to KEGG restrictions



Data Integrating and Analysing in SBGN Maps

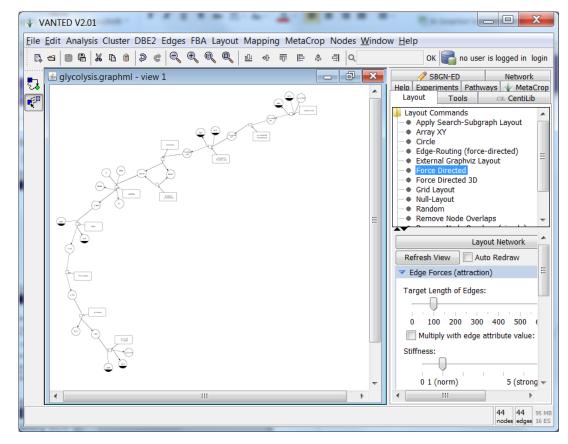
- VANTED provides many methods for
 - Data mapping and integration
 - Data and network analysis
- Also available in SBGN-ED





Layout

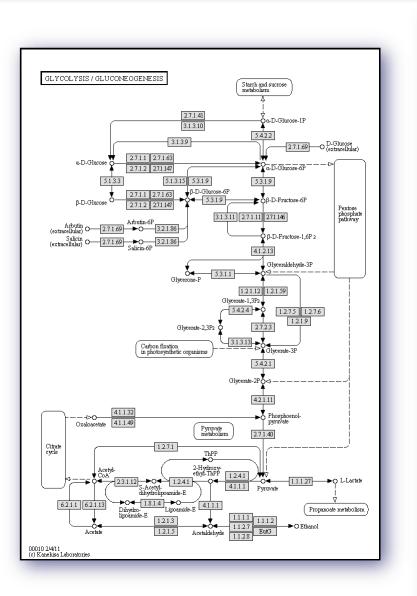
 VANTED provides many standard layout algorithm

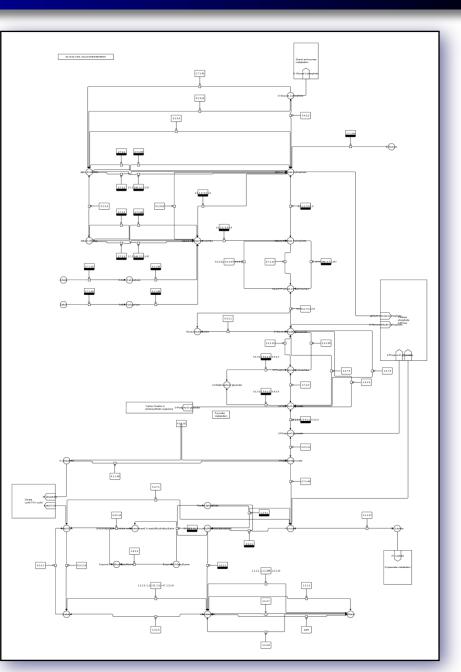


Outlook

- 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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