SBGN in Use

Learning and using
Process Description language

Topics

- Teaching/learning SBGN
- SBGN PD in Excel
- Representing incomplete information
- Aesthetics

Teaching/learning SBGN

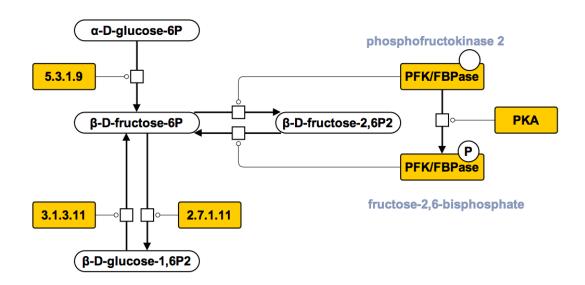
- MSc students, Computational Systems Biology course, School of Informatics, University of Edinburgh
- MSc students, Pathway Reconstruction course, Department of Pathway Medicine, University of Edinburgh

Guidelines

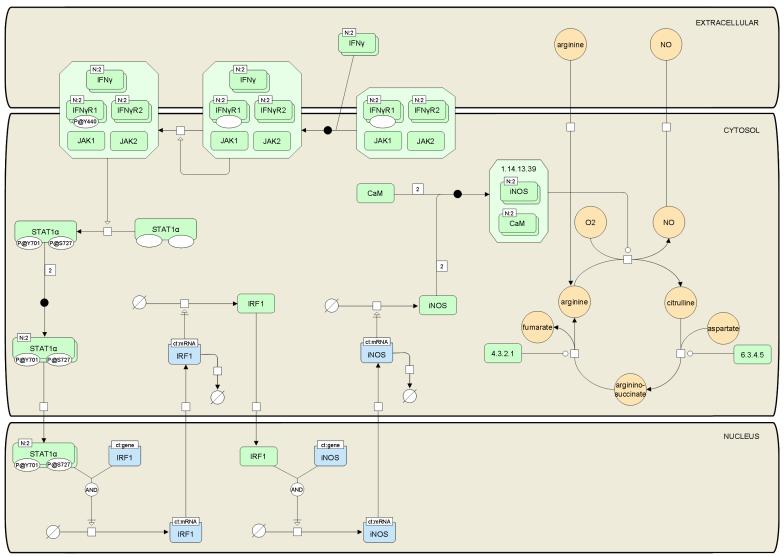
- Good first examples are very important
- Hands-on experience: drawing
- The tool is important (easy-to-create nice-looking diagrams, flexibility)
- Steps
 - 1. Attempt to visualize a textually described system (1st example)
 - 2. Redrawing the main example diagram (2nd example)
 - Pathway reconstruction: visualization in SBGN PD + annotation in Excel

Text to Diagram

Conversion of F6P into F2,6P2 is catalyzed by PFK. Conversion of F2,6P2 into F6P is catalyzed by FBPase. Both PFK and FBPase activities are the functions of one protein. Unphosphorylated protein has PFK activity and phosphorylated protein has FBPase activity. Phosphorylation of the protein is catalyzed by PKA.

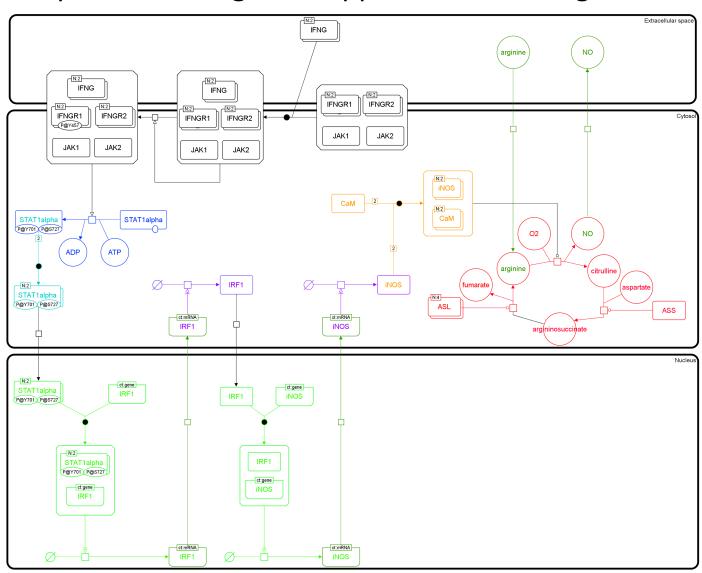


The main example: as many symbols as possible on a small diagram



iNOS pathway: the last version

http://sourceforge.net/apps/mediawiki/sbgnbricks



SBGN PD in Excel

Developing a database with SBGN PD compatibility

SBGN PD symbols -> Text

| Group | Element | Example |
|-------------------|------------------------|--|
| Entity Pool Nodes | unspecified entity | |
| Entity Pool Nodes | simple chemical | Default |
| Entity Pool Nodes | macromolecule | [mt:prot]STAT1alpha <p@y701,s727></p@y701,s727> |
| Entity Pool Nodes | macromolecule multimer | [mt:prot]STAT1alpha <p@y701,s727>[N:2]</p@y701,s727> |
| Entity Pool Nodes | nucleic acid feature | [ct:gene]iNOS |
| Entity Pool Nodes | nucleic acid feature | [ct:mRNA]iNOS |
| Entity Pool Nodes | perturbing agent | |
| Entity Pool Nodes | source/sink | Ø Ø |
| Entity Pool Nodes | complex | { [mt:prot]iNOS[N:2]; [mt:prot]CaM[N:2] } |
| Auxiliary Units | unit of information | [mt:prot] |
| Auxiliary Units | state variable | <p@y701,s727></p@y701,s727> |
| Auxiliary Units | clone marker | Automatically |
| Process Nodes | process | >> process >>, << process >> |
| Process Nodes | omitted process | >>/\>>, < \ > |
| Process Nodes | uncertain process | >>?>>, < > |
| Process Nodes | association | >> association >> |
| Process Nodes | dissociation | >> dissociation >> |
| Connecting Arcs | consumtion | Input column |
| Connecting Arcs | production | Output column |
| Connecting Arcs | modulation | Modulation column |
| Connecting Arcs | stimulation | Stimulation column |
| Connecting Arcs | catalysis | Catalysis column |
| Connecting Arcs | inhibition | Inhibition column |
| Connecting Arcs | necessary stimulation | Necessary stimulation column |

1. Short names for entity pool nodes

```
Entity Marker / Alias / Short Name
```

"STAT1alphaP"

STAT TalphaP

"STAT1alphaP*2"

"IFNGR_complex"

"IFNGR_active"

"iNOS"

"iNOS gene"
"iNOS mRNA"

"iNOS-CaM"

Entity Definition

[mt:prot]STAT1alpha<P@Y701,S727>

[mt:prot]STAT1alpha<P@Y701,S727>[N:2]

{ [mt:prot]INFG[N:2]; [mt:prot]INFGR1[N:2]; [mt:prot]INFGR2[N:2];

[mt:prot]JAK1; [mt:prot]JAK2 }

{ [mt:prot]INFG[N:2]; [mt:prot]INFGR1<P@Y440>[N:2]; [mt:prot]INFGR2*2;

[mt:prot]JAK1; [mt:prot]JAK2 }

[mt:prot]iNOS

[ct:gene]iNOS

[ct:mRNA]iNOS

{ [mt:prot]iNOS[N:2]; [mt:prot]CaM[N:2] }

[LABEL]

<LABEL>

{complex description}

unit of information

state variable

complex

"SHORT_NAME"

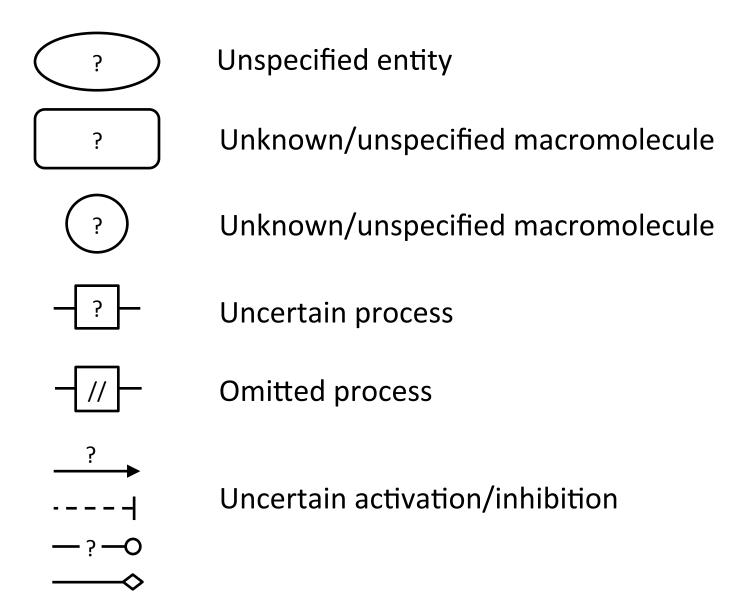
2. Stimulation/inhibition and logic gates

| Input | Process Type and Direction | Output | Stimulation | Catalysis "iNOS- | Necessary Stimulation | Inhibi tion |
|-------------|-----------------------------|----------------------------------|----------------|------------------|----------------------------------|----------------|
| J | >> process >> >> process >> | citruline + NO "STAT1alpha2P" | "IFNGR_active" | CaM" | | |
| "empty set" | >> process >> | "IRF1_mRNA" | - | | "STAT1alpha2P" AND "IRF1_gene | , |

One line – one process plus all the connected links (activation/inhibition etc.)

Incomplete knowledge representation

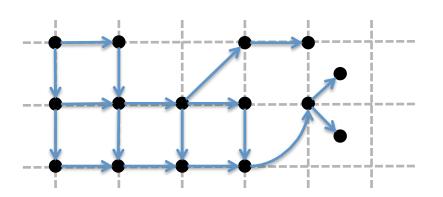
Uncertain connections

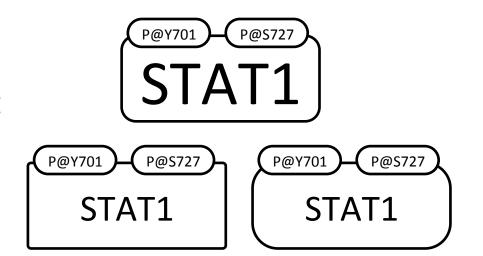


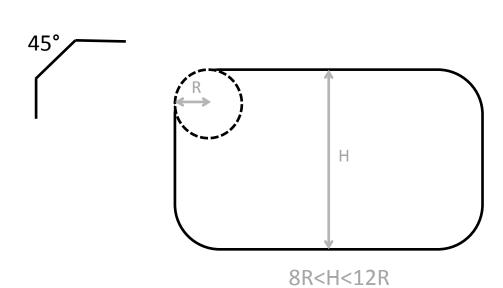
Aestetics

Readability

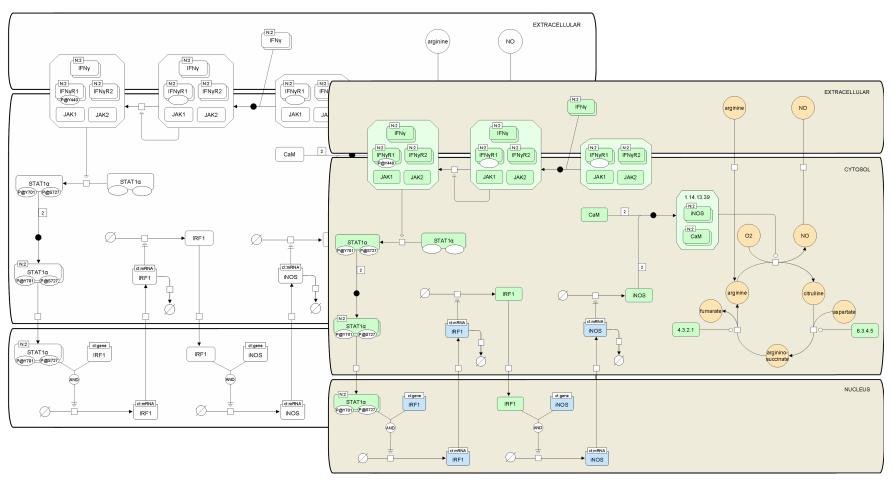
- The smallest font size
- Default shape size/font size
- Layout
- Patterns
- Irregularities







Colours?



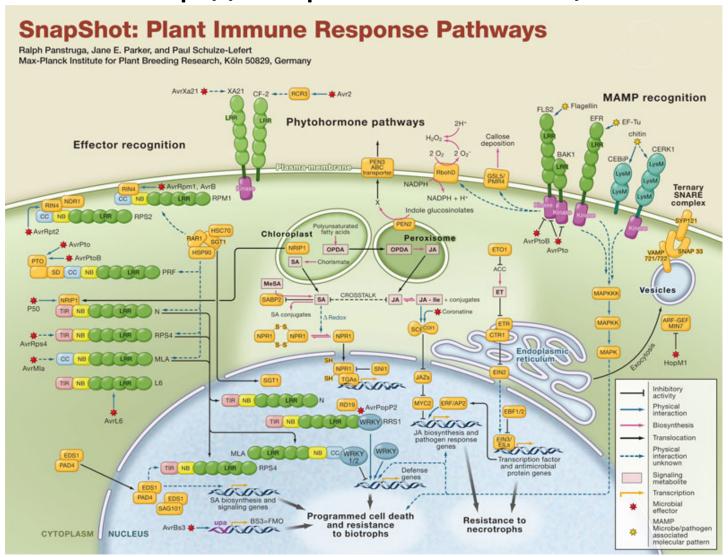
Leave it up to users without offering a default colour scheme? Will a nice default colour scheme help to promote SBGN?

Default colour scheme?

http://colorschemedesigner.com/



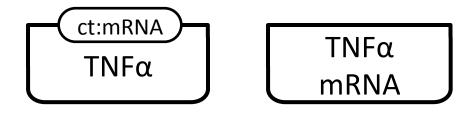
Adopting any existing colour scheme? http://snapshots.cell.com/



Example diagrams

What kind of examples?

Can use and have to use



The same information can be represented in different ways and the set of the examples should reflect that

Thank you

SBGN-9

Edinburgh, 29 Apr – 2 May, 2013