



SBGN Competition 2010

**How to represent EGFR pathway?
A Case Study**

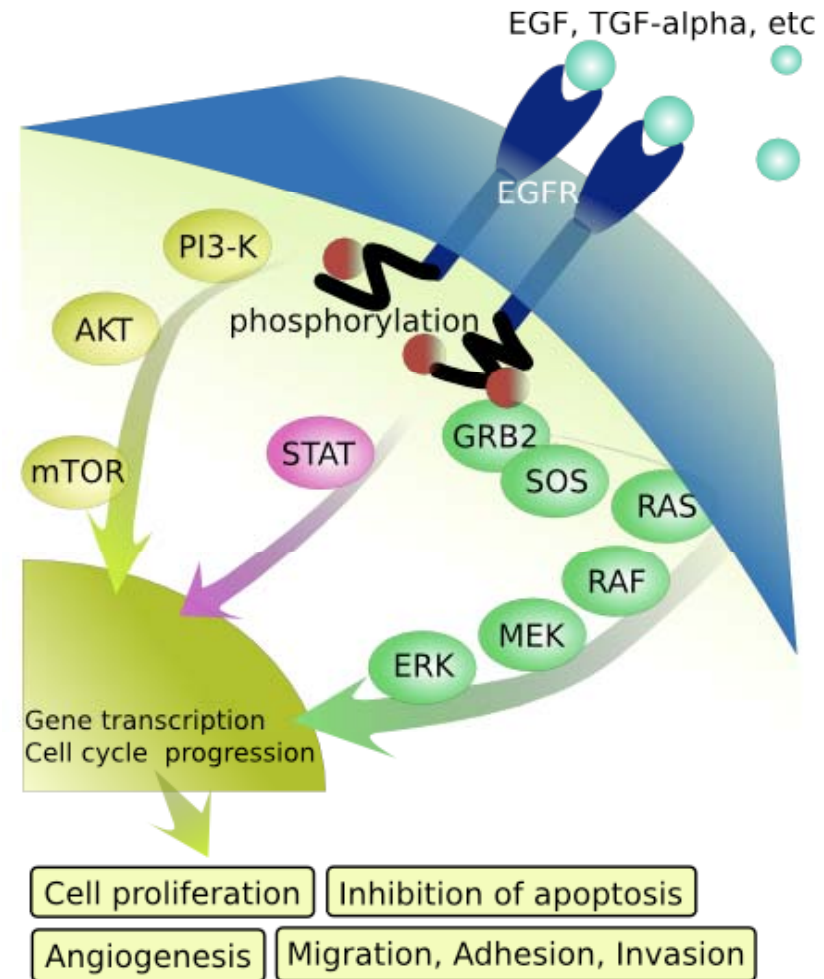
2010/Aug

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JST ERATO Kawaoka Infection-Induced Host Response Network Project

SBGN Process Description Diagram EGFR Signaling Pathway

*What would be
the best representation of
SBGN Process Description
Diagram notation,
using EGFR signaling pathway
as an example?*

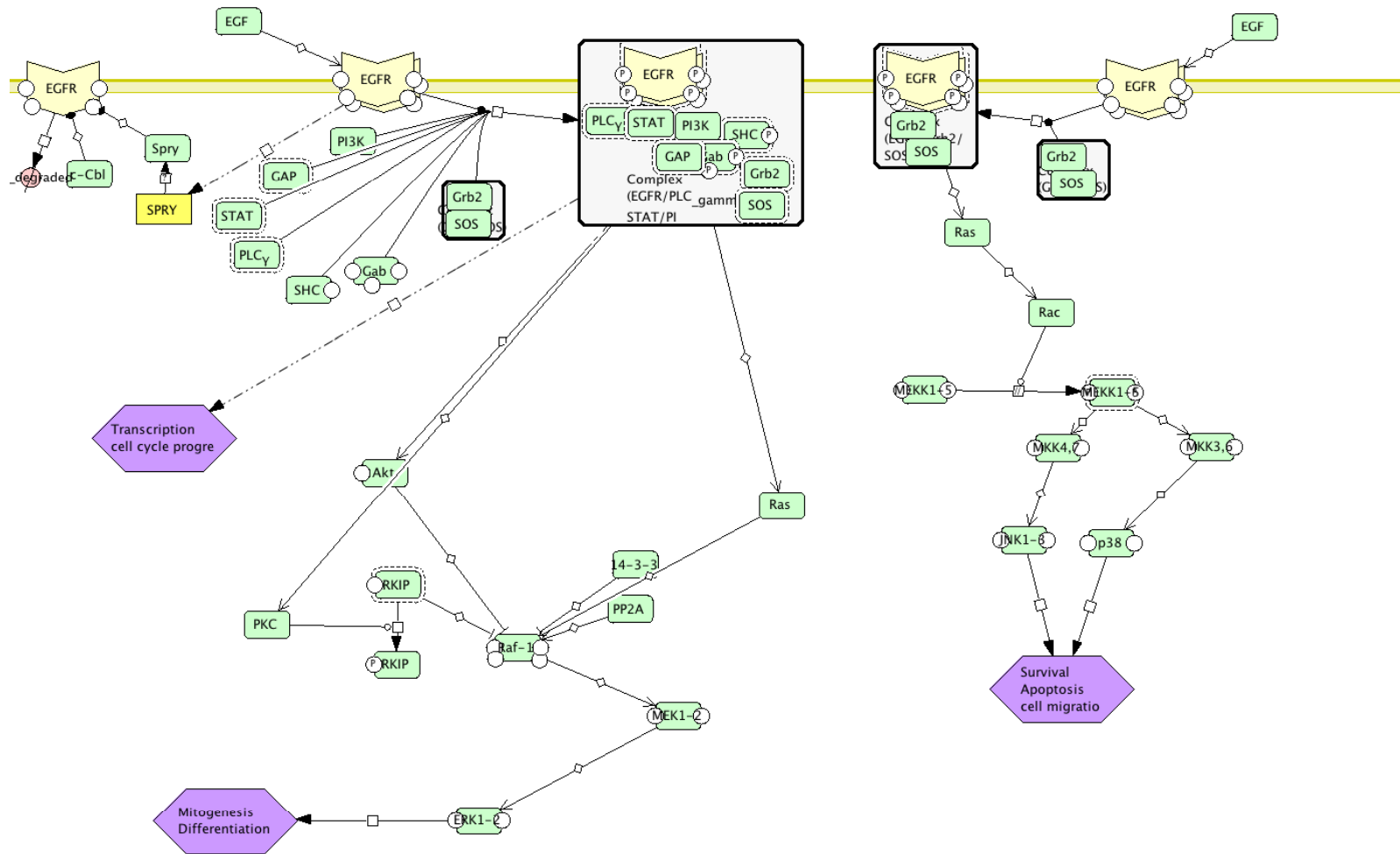


From Wikipedia

Protocol

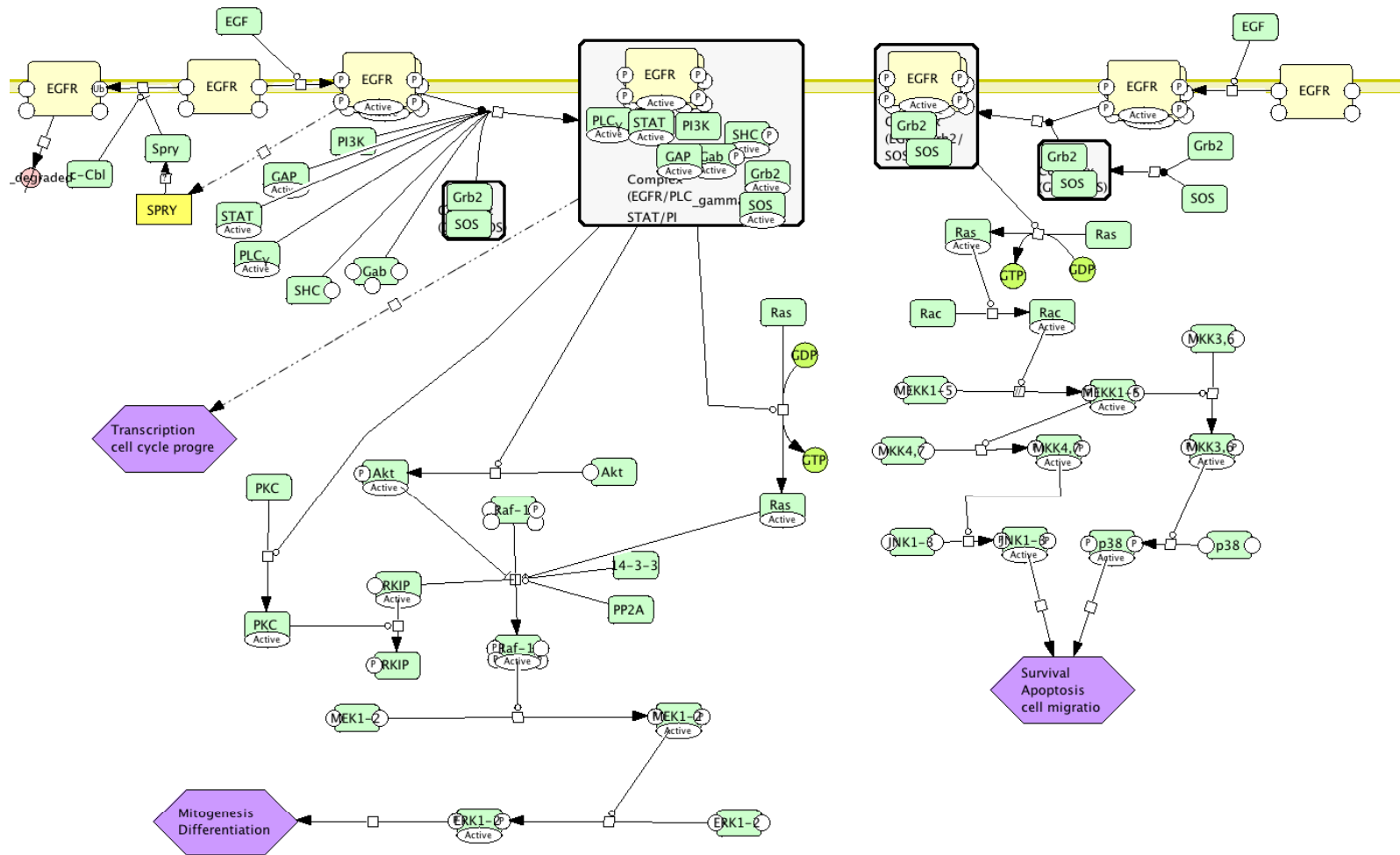
1. Take a model from Panther Pathways Database:
<http://www.pantherdb.org/pathway/>
2. Import the model to CellDesigner Ver.4.1.
<http://celldesigner.org>
3. Re-layout the model.
4. Convert it using SBGN Viewer function of CellDesigner.
5. Export the model image in .SVG or .PDF format.
6. Adjust the detail settings of the model using Illustrator.

Panther Pathways Database Lite View



Panther Pathways Database

SBGN view

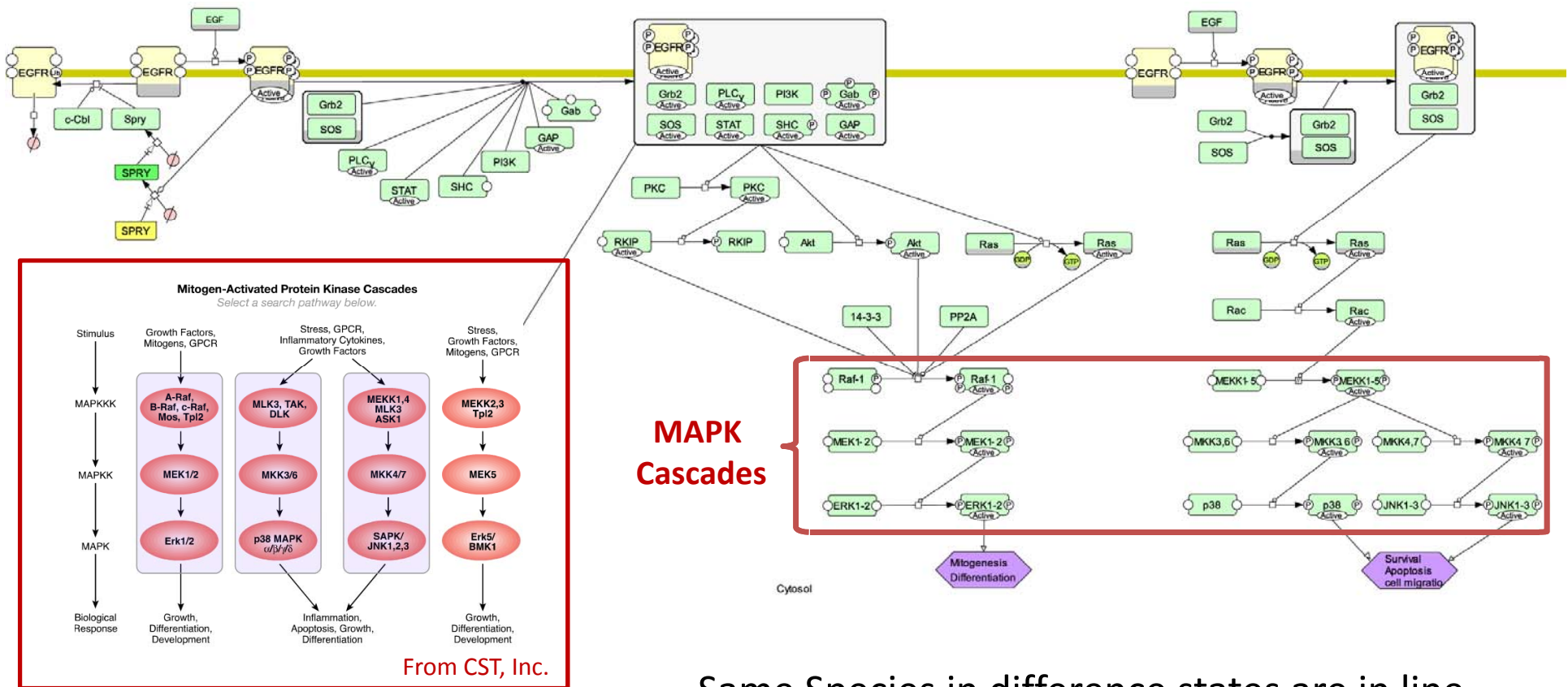


Layout a Signaling Pathway

- **Align:** to illustrate Flow of Information
 - Cascade
 - Input / Output (Top-to-Bottom or Left-to-Right)
- **Merge :** Duplicate Species or Merge them?
- **Connect:** Lines Straight or Orthogonal bend?
- **Modifications:** Catalyse or Inhibit?
- **Complex:** Position of the Components?
- **Dimer :** Dimer? Two Components in parallel?

MAP-1: Align

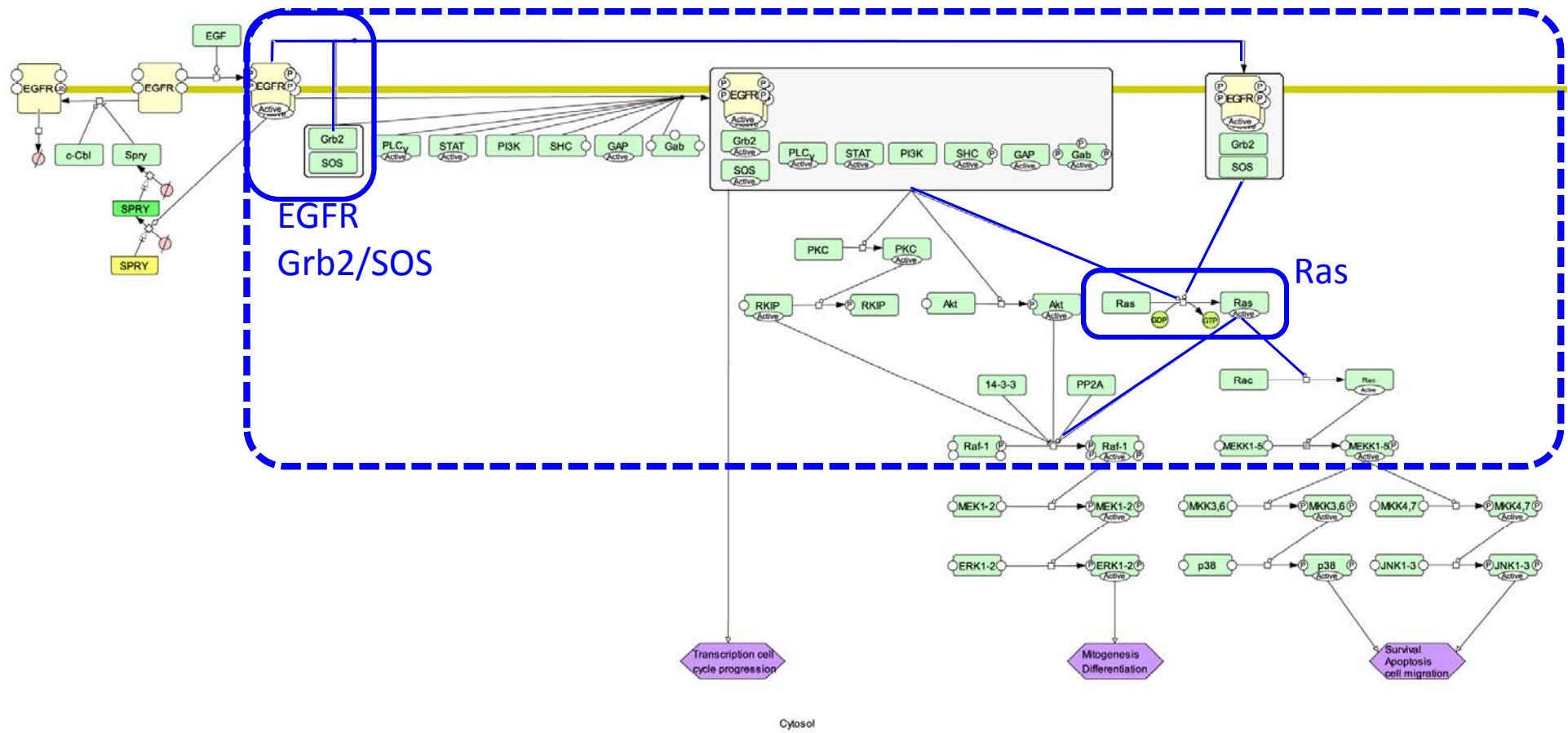
Align Components Vertically and Horizontally



Same Species in difference states are in line
 Highlight **Cascade** pattern

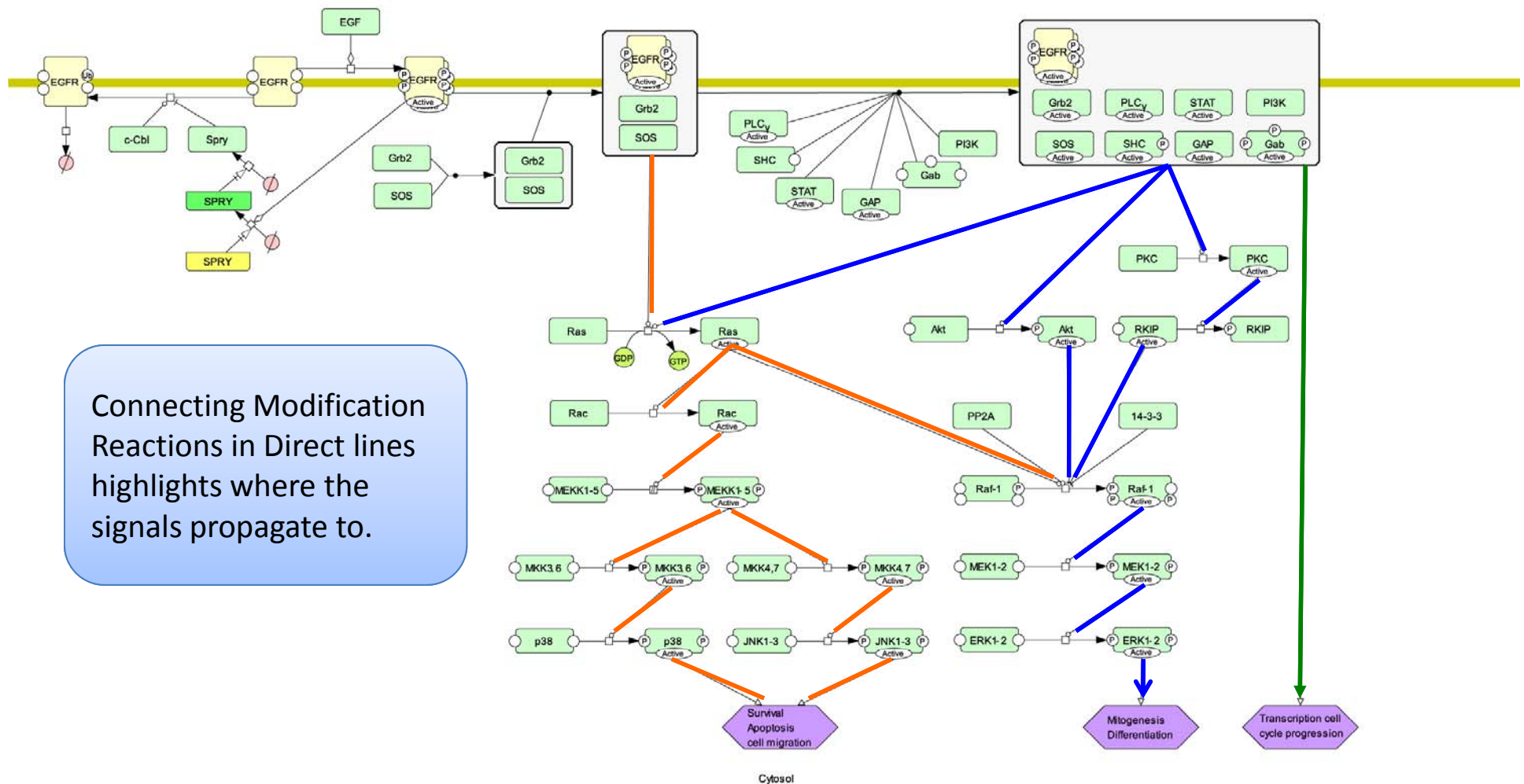
MAP-2 : Merge

Merge the common components



MAP-3-1: Connect - Straight

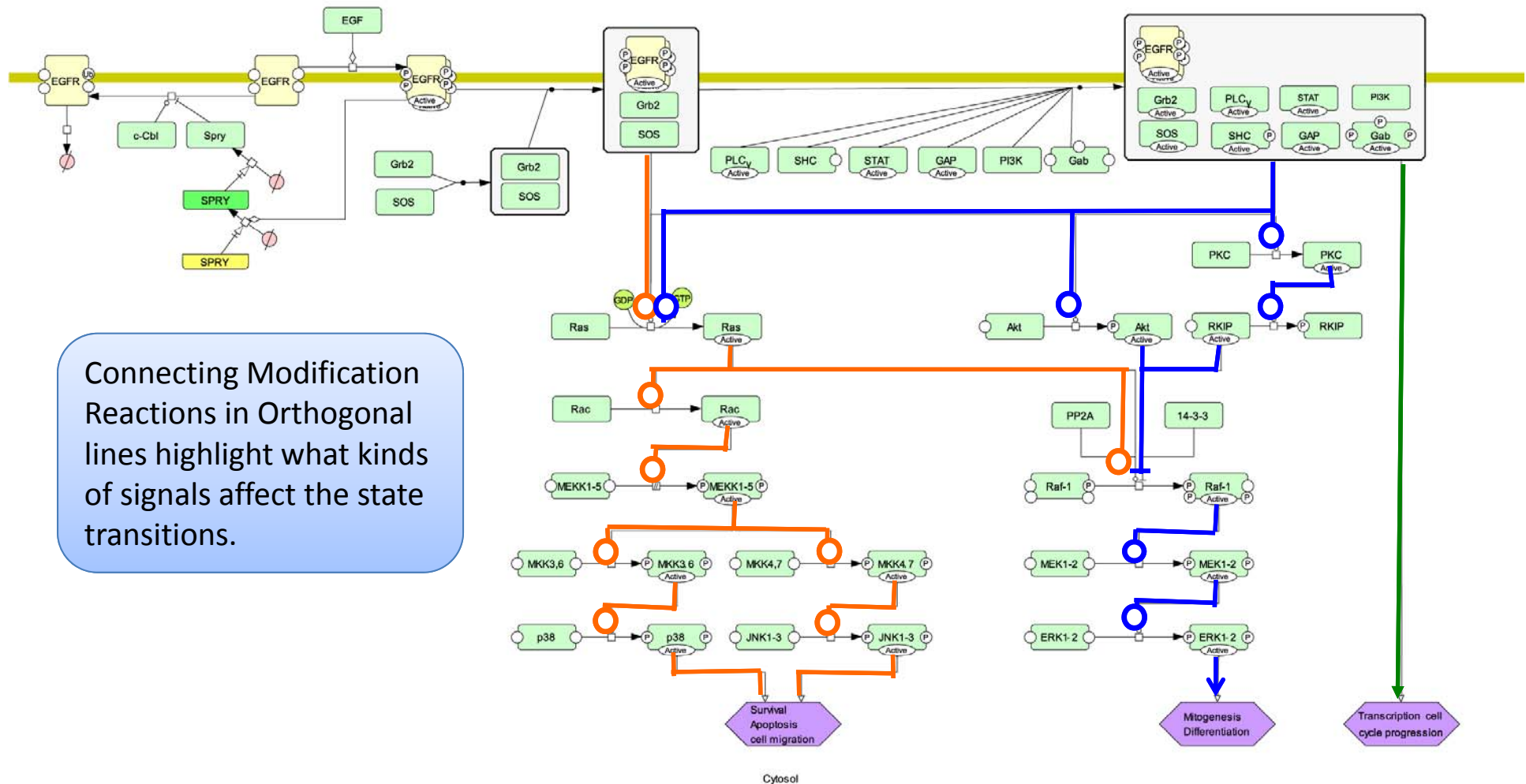
Connecting Modification Reactions in Direct lines



Connecting Modification Reactions in Direct lines highlights where the signals propagate to.

MAP-3-2: Connect - Orthogonal

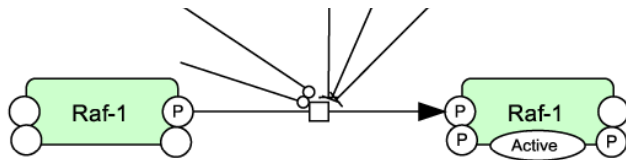
Connecting Modification in Orthogonal



Pattern: Modifications at Process Node

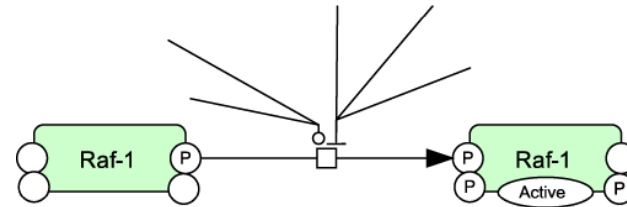
A

Easy to identify the number of modifications but the connections are overlapped around the process node.



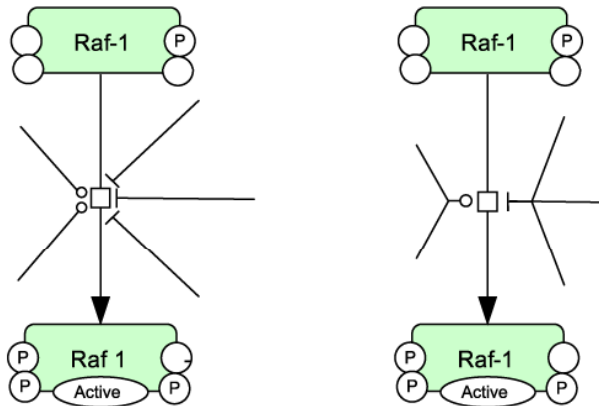
B

Easy to identify the number of modifications and connection types can easily detect.



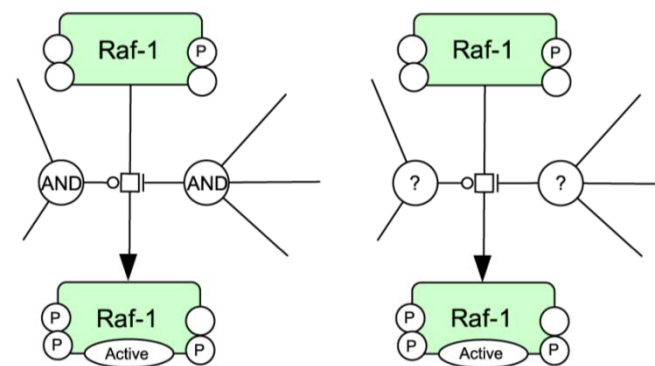
C

Drawing the same type of modifications lines from the same direction.



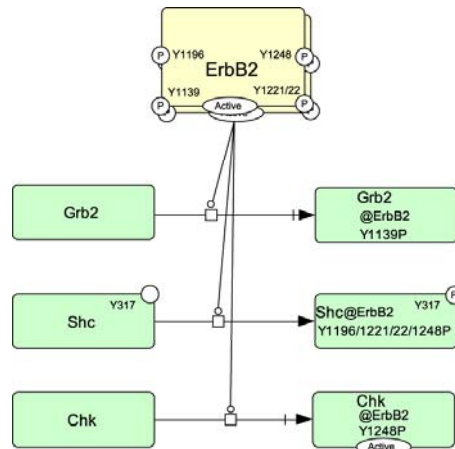
D

Use a Boolean logic gates



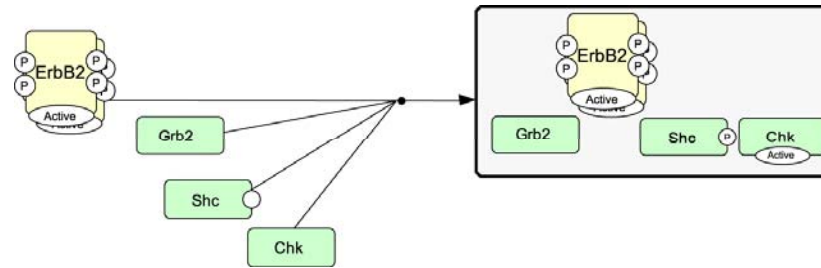
Pattern: Receptors 1

A Show details of modifications

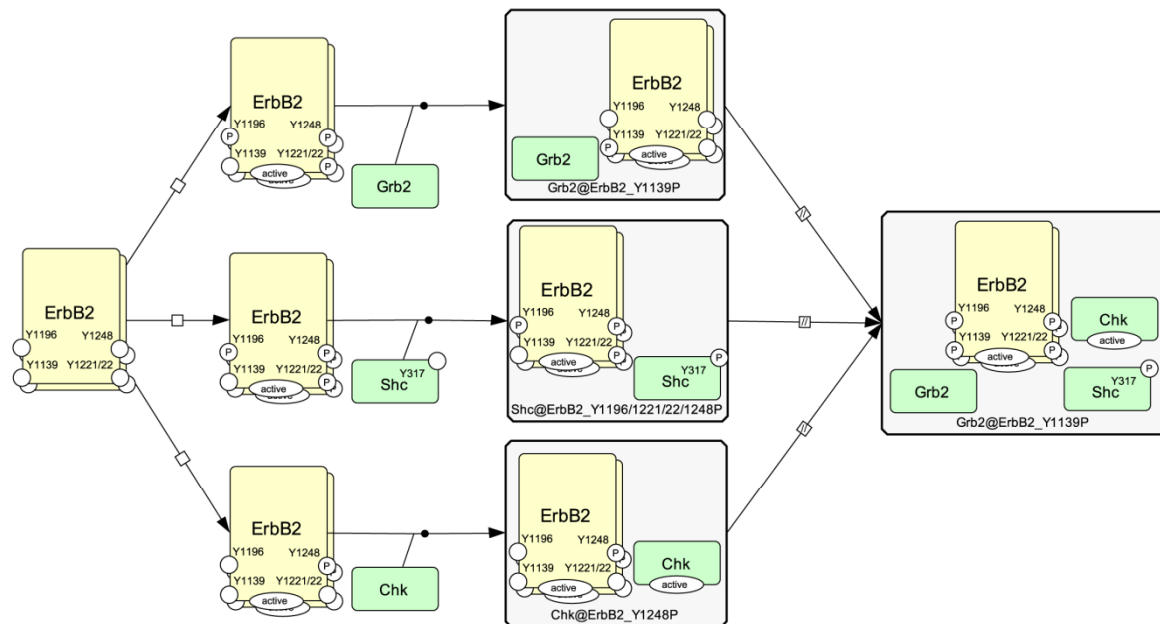


Oda et al. 2005

B Show flow of state transition (simplified)



C Show state transition and the details of modifications

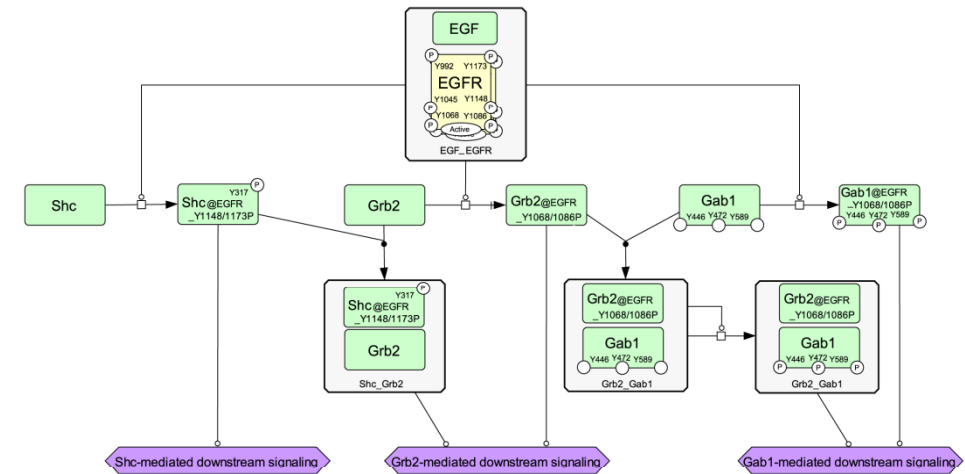
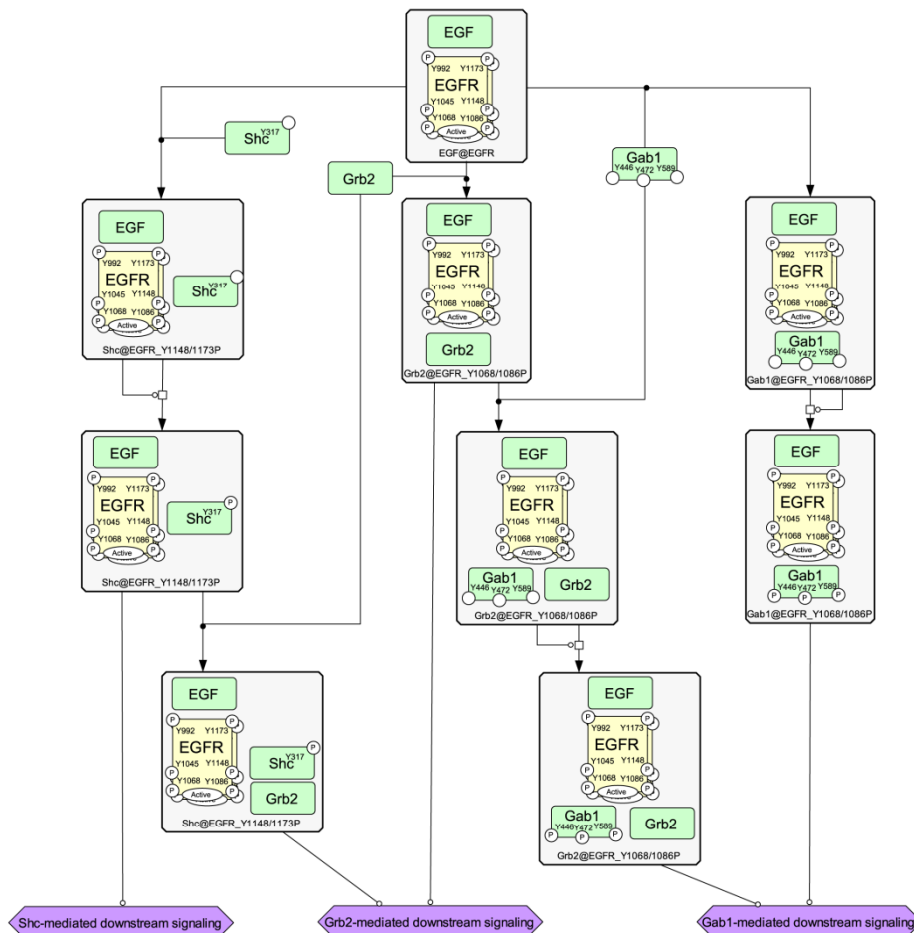


Pattern: Receptors 2

Oda et al. 2005

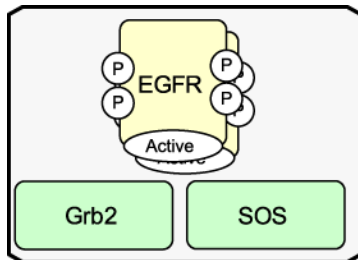
A Show everything (details of modifications of all the component in the complexes.)

B Highlight modifications only

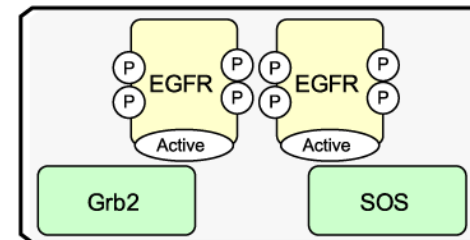


Pattern: Dimerization

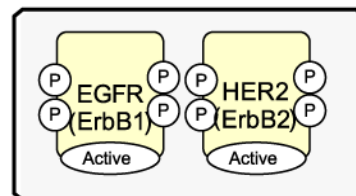
A Receptor as a Dimer



B Two receptors in a complex



C Different receptors form a complex



Supplementary files

- Files:
 - EGF_receptor_signaling_pathway(.xml/.png)*[Original Panther Pathways Models](#)
 - MAP1_Align(.xml /.ai /.png)
 - MAP2_Merge(.xml /.ai /.png)
 - MAP3-1_Connect-Straight(.xml /.ai /.png)
 - MAP3-2_Connect-Orthogonal(.xml /.ai /.png)
 - Modifications (.xml /.ai /.png)
 - Receptors1_ErbB2 (.xml)
 - Receptors1 (.ai /.png)
 - Receptors2_Oda-et al2005_A (.xml /.ai /.png)
 - Receptors2_Oda-et al2005_B (.xml /.ai /.png)
 - Dimerization (.ai /.png)