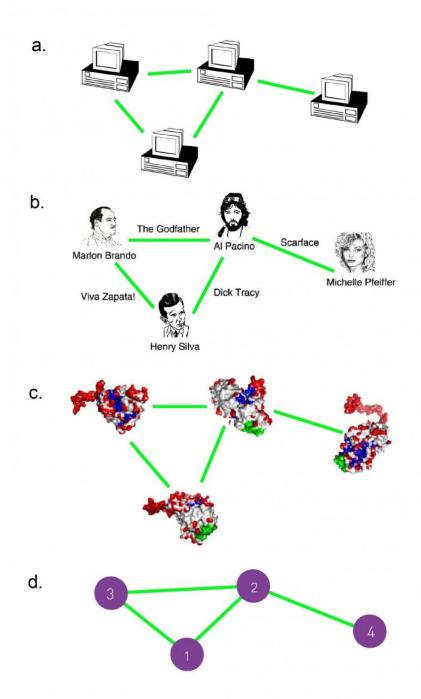
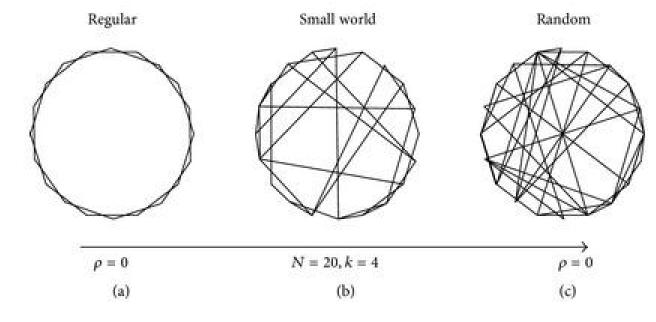


Networks

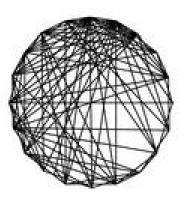
- → Graph:
 - abstractmathematicalrepresentation, datastructure
- → Network
 - phenomenon modelled by graph



Network topology

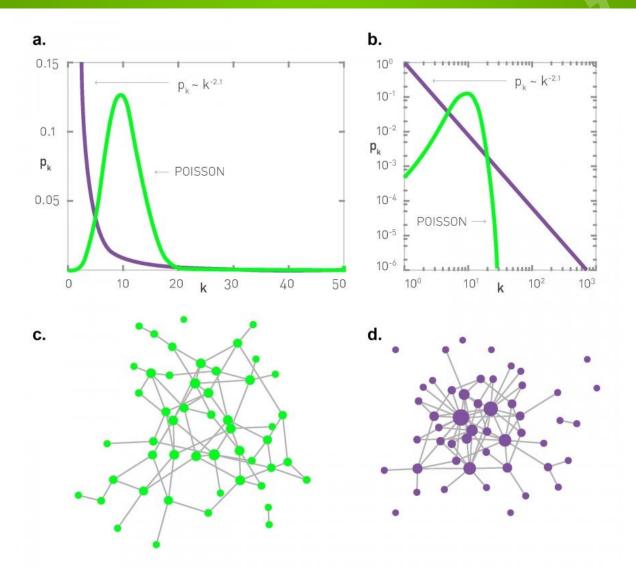


Free-scale

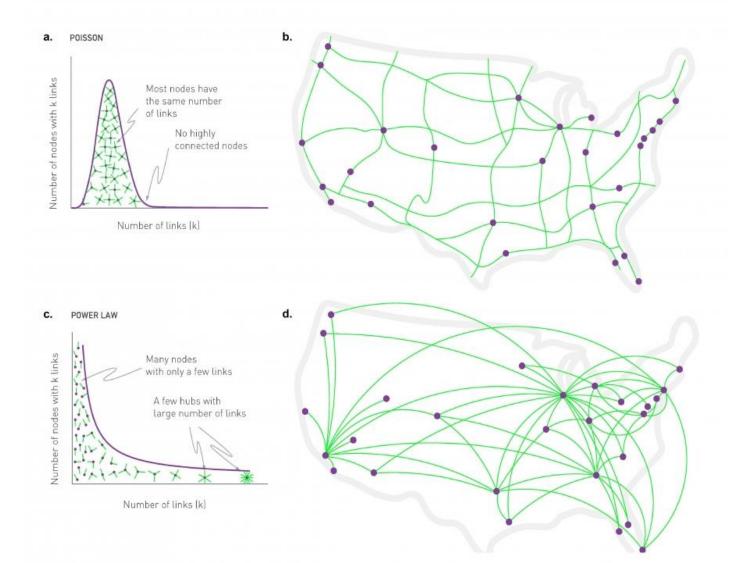


$$N = 30, m_0 = 6, k = 4$$
(d)

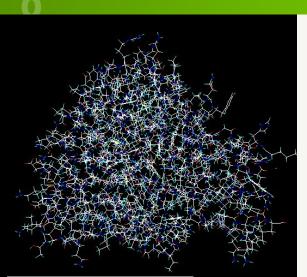


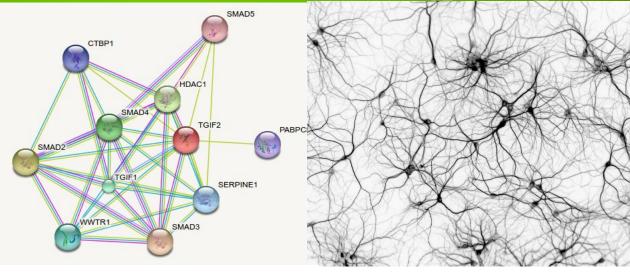




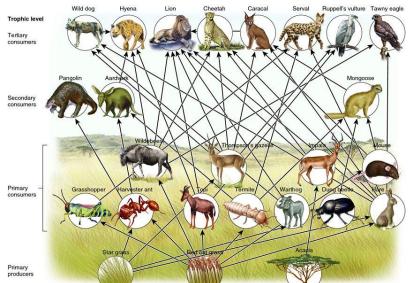


Nested networks



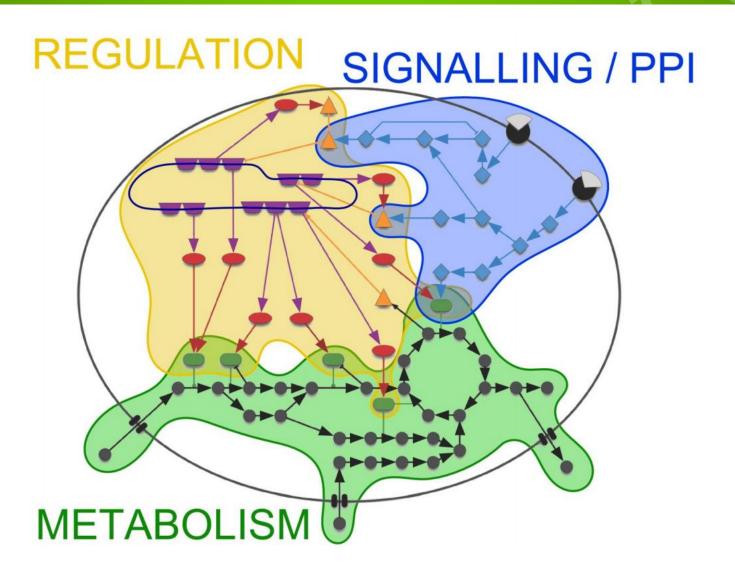




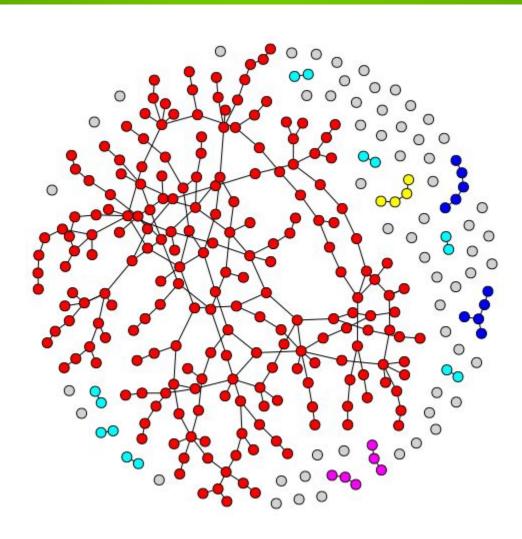




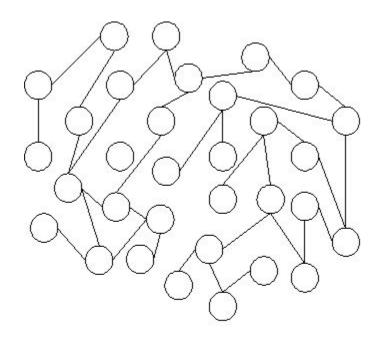
Molecular networks

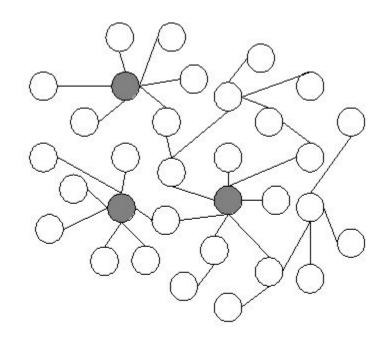


Giant component (sub-graphs)



Special nodes in scale-free networks

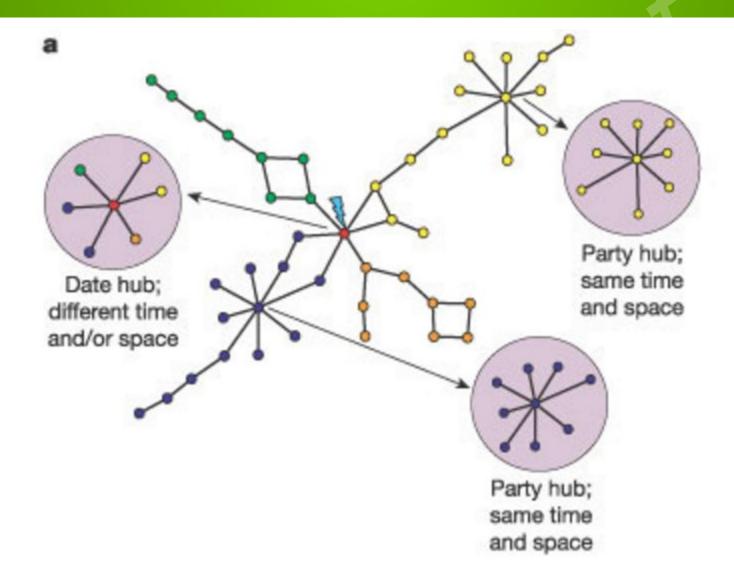




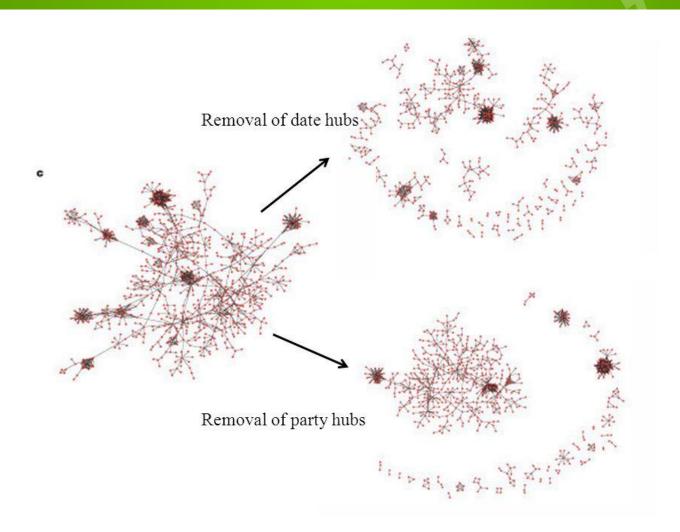
(a) Random network

(b) Scale-free network

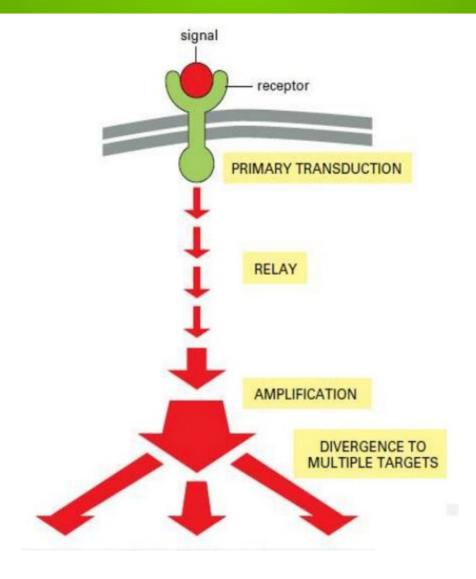




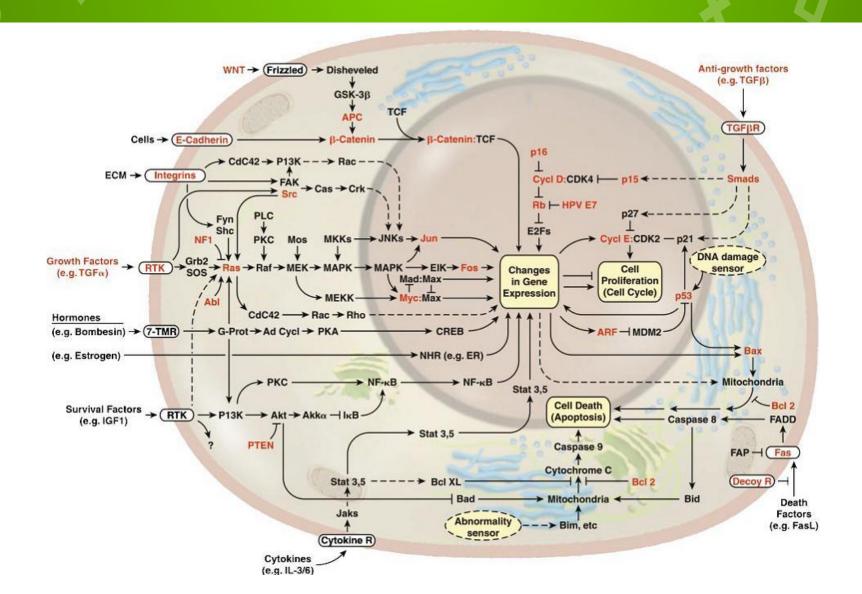
Party hub, date hub





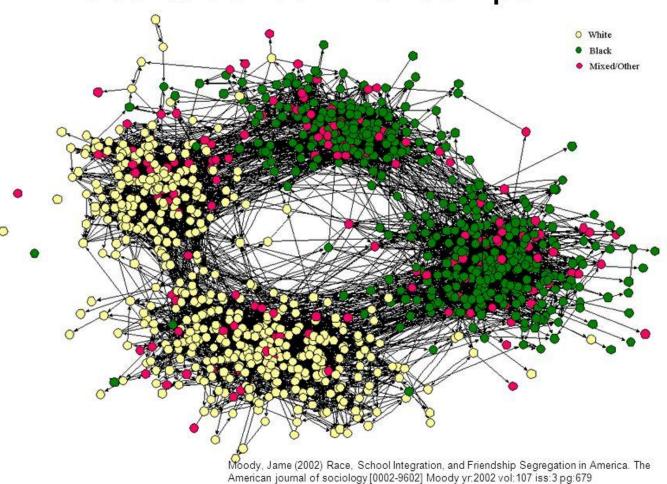


Singnal flow



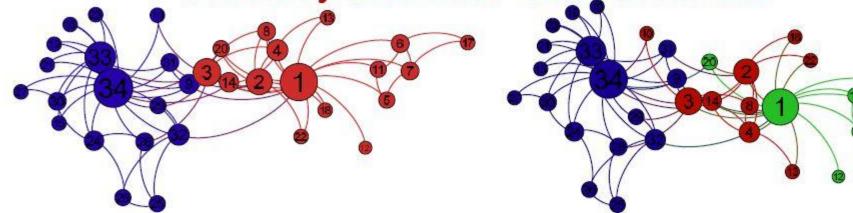
Modules, communities, clusters

Race & school friendships

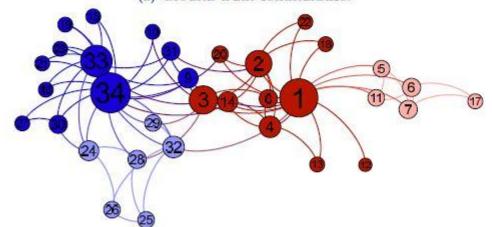


Modules, communities, clusters

Zachary's Karate Club Network

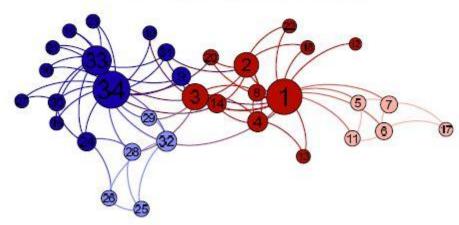


(a) Ground truth communities.



(c) Communities detected with Fine-tuned Q.

(b) Communities detected with Greedy Q.

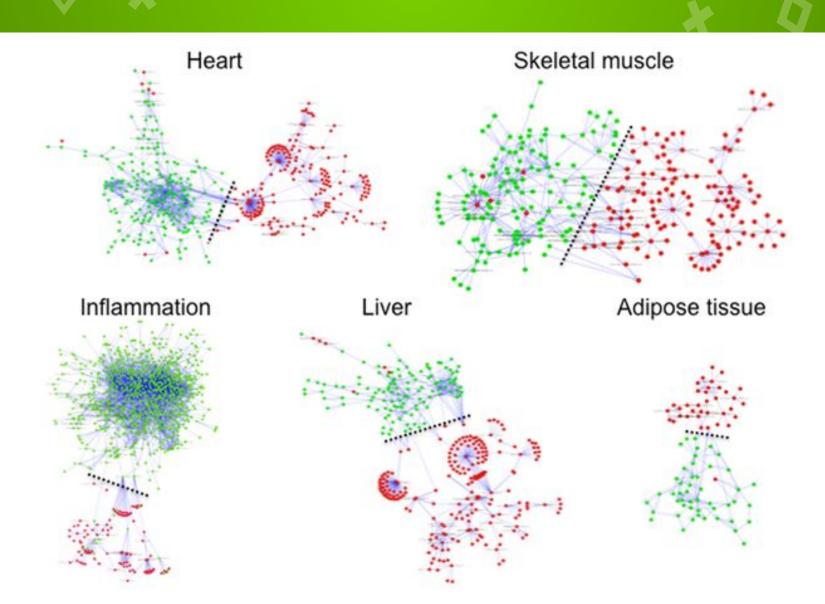


(d) Communities detected with Fine-tuned Q_{ds}.

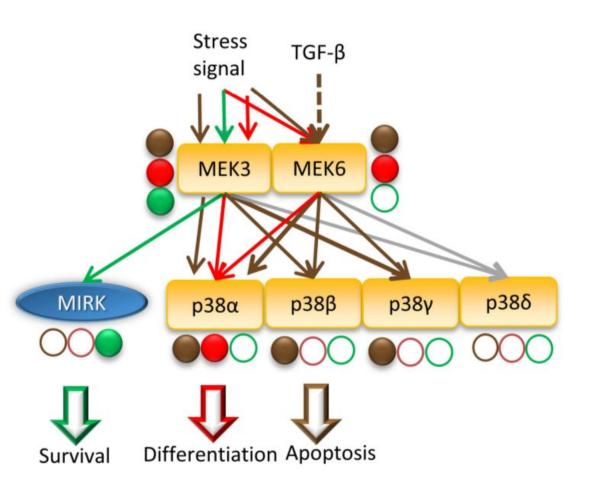
Probabilistic networks

- → By the present methods, we can study all possible molecular interactions
- Many interactions
- → High false-positive rate
- → All connections can not be concurrent at the same time:
 - common binding surface
 - different expression
 - different localization

Context specific via different expression

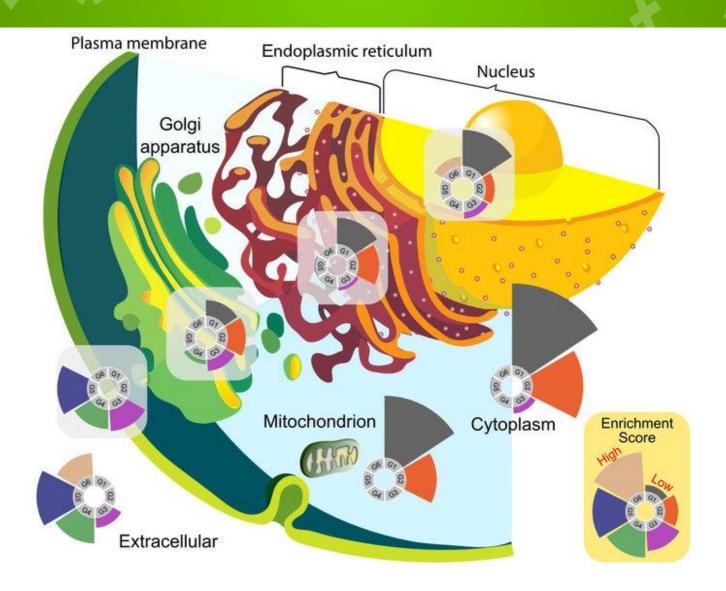


Context specific via different expression

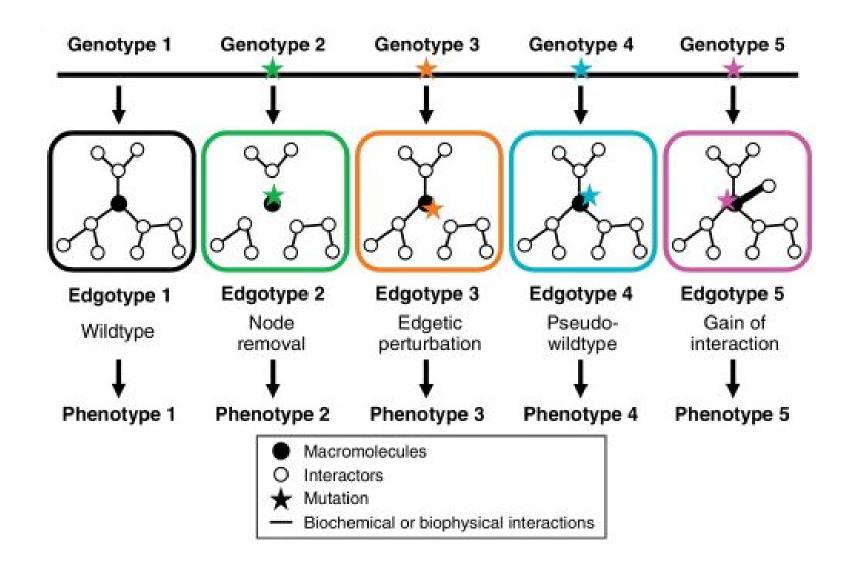


- Liver and gall bladder
- Cardiovascular
- Peripheral Nervous System

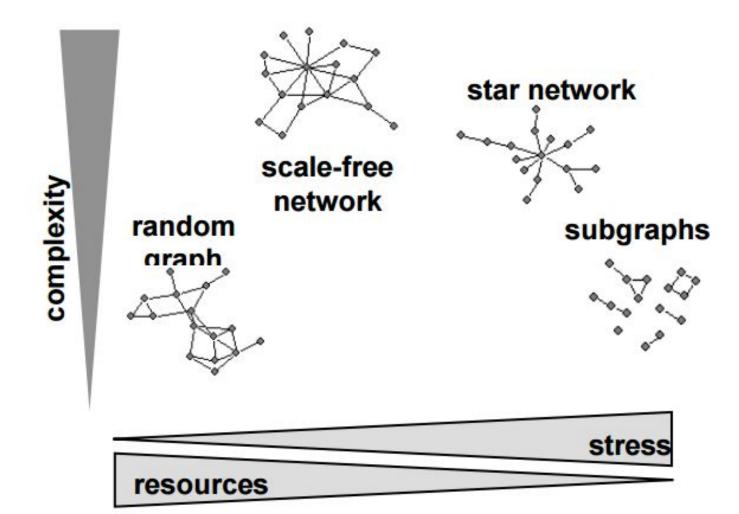
Context specific via subcellular localization



Edgetics - ???



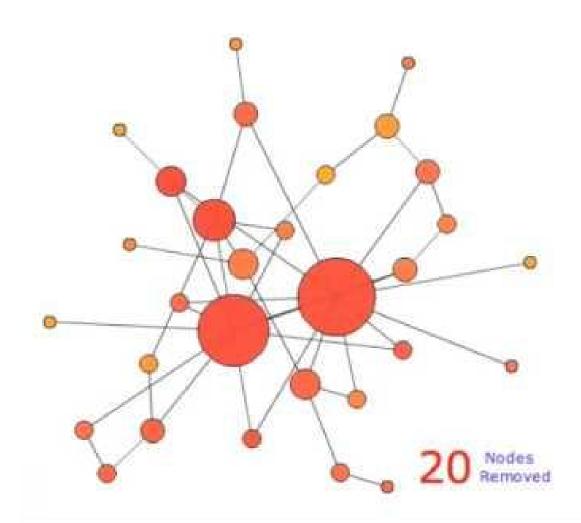
Topological phase transformations



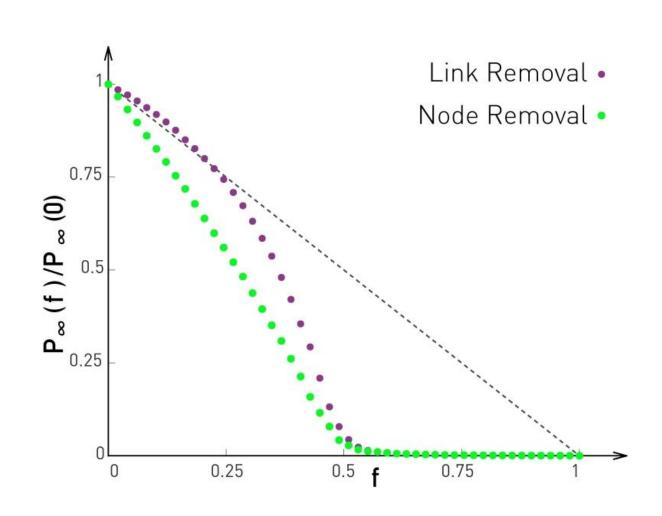
Network stability

- Network stability indicates that a given network responds to external influences
- External impact: Termination of point or connection
- → Attack strategy:
 - random
 - planned

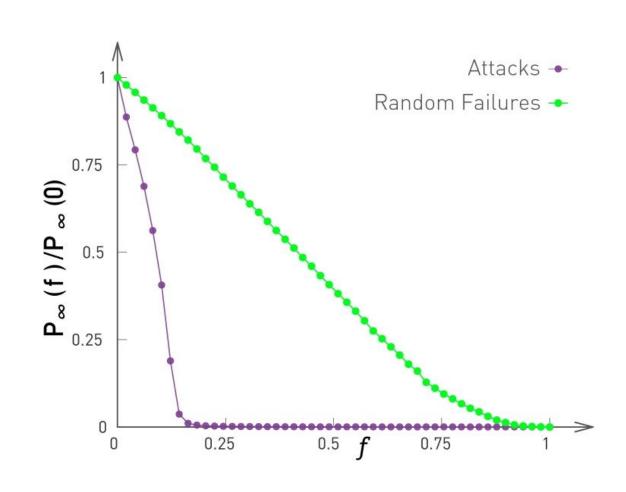
Network stability



Network stability - scale-free networks



Network stability - scale-free networks



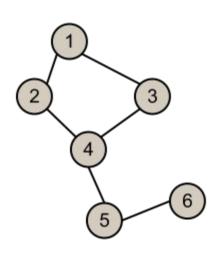
Storing network data - datastructure

Adjacency list

→ Adjacency matrix

Adjacency matrix

Undirected Graph & Adjacency Matrix



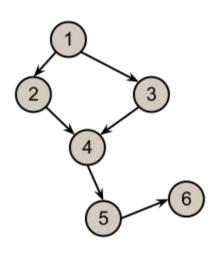
Undirected Graph

	1	2	3	4	(5)	6
1	0_	1	1	0	0	0
2	1	0	0	1	0	0
3	1	0	0	1	0	0
4	0	1	1	0	1	0
(5)	0	0	0	1	0	1
6	0	0	0	0	1	0

Adjacency Matrix

Adjacency matrix

Directed Graph & Adjacency Matrix



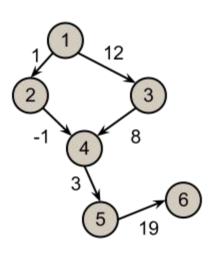
Undirected Graph

	1	2	3	4	(5)	6
1	0_	1	1	0	0	0
2	-1	0	0	1	0	0
3	-1	0	0	1	0	0
4	0	-1	-1	0	1	0
(5)	0	0	0	-1	0	1
6	0	0	0	0	-1	0

Adjacency Matrix

Adjacency matrix

Weighted Directed Graph & Adjacency Matrix



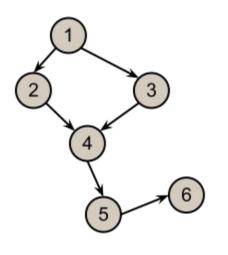
Weighted Directed Graph

	1	2	3	4	5	6
1	0	1	12	0	0	0
2	-1	0	0	-1	0	0
3	-12	0	0	8	0	0
4	0	1	-8	0	3	0
(5)	0	0	0	-3	0	19
6	0	0	0	0	-19	0

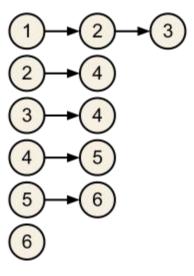
Adjacency Matrix

Adjacency list

Directed Graph & Adjacency List



Undirected Graph



Adjacency Matrix

Adjacency list - file formats

- → CSV, TSV
 - table
- → SQL
 - relational database
- → XML
 - table like
- → SBML
 - special XML
- → BioPax
 - special XML
- → PSI-MI
 - complex format syntax and ontology

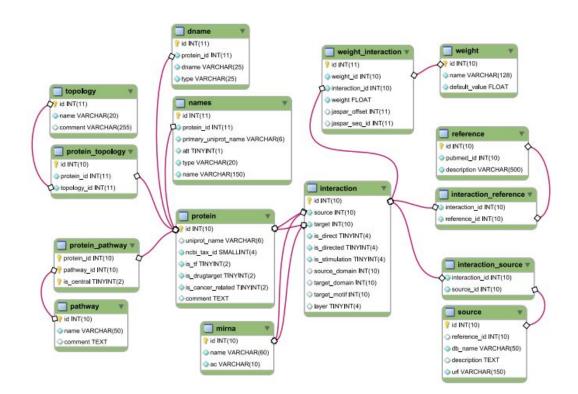
Storing network data - file formats - CSV, TSV

- Semicolon or Tabbed Text Format
- → Can be edited with MS Excel
- It is easy to read with the program and manually

```
source name; source uniprotAC; source speciesID; source species; source topology; source pathways; target name; ta
JAK2;060674;ENSG00000096968;H. sapiens;Mediator;JAK/STAT(core);PTPN11;Q06124;ENSG00000179295;H. sapiens;Co-
PTPN11;Q06124;ENSG00000179295;H. sapiens;Co-factor,Scaffold;RTK(non-core),JAK/STAT(non-core);JAK2;060674;ENS
IRS1; P35568; ENSG00000169047; H. sapiens; Mediator, Scaffold; RTK(core), JAK/STAT(core); JAK1; P23458; ENSG000001624
JAK1; P23458; ENSG00000162434; H. sapiens; Mediator; RTK(core), JAK/STAT(core); IRS1; P35568; ENSG00000169047; H. sapiens
GSK3B; P49841; ENSG00000082701; H. sapiens; Mediator, Co-factor; RTK(non-core), RTK(core), Hedgehog(core), TGF(core)
AXIN1;015169;ENSG00000103126;H. sapiens;Mediator,Scaffold;RTK(non-core),TGF(non-core),TGF(core),WNT/Wingles
MAP2K1;Q02750;ENSG00000169032;H. sapiens;;RTK(core),Hedgehog(core);MAPK3;P27361;ENSG00000102882;H. sapiens;
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SMAD3; P84022; ENSG00000166949; H. sapiens; Mediator, Transcription factor; RTK(core), NHR(core), TGF(core), WNT/Wind
ESR1; P03372; ENSG00000091831; H. sapiens; Receptor, Transcription factor; NHR(core), TGF(non-core); SMAD3; P84022; EN
PEA15;Q15121;ENSG00000162734;H. sapiens;Co-factor;RTK(non-core);MAPK3;P27361;ENSG00000102882;H. sapiens;Med
MAPK3; P27361; ENSG00000102882; H. sapiens; Mediator; RTK(core), JAK/STAT(core), TGF(core); PEA15; Q15121; ENSG0000010
IL10RB;Q08334;ENSG00000243646;H. sapiens;Ligand ;JAK/STAT(core);IL28RA;Q8IU57;ENSG00000185436;H. sapiens;Rec
IL10RB;Q08334;ENSG00000243646;H. sapiens;Ligand ;JAK/STAT(core);IL28RA;Q8IU57;ENSG00000185436;H. sapiens;Rec
SMURF2;Q9HAU4;ENSG00000108854;H. sapiens;Co-factor,Endocytosis related ,Scaffold;TGF(non-core),WNT/Wingless
SMAD2;Q15796;ENSG00000175387;H. sapiens;Mediator,Transcription factor;RTK(core),TGF(core),WNT/Wingless(core
SMAD3; P84022; ENSG00000166949; H. sapiens; Mediator, Transcription factor; RTK(core), NHR(core), TGF(core), WNT/Wind
NOTCH1;P46531;ENSG00000148400;H. sapiens;Receptor,Endocytosis related;Notch(core),WNT/Wingless(core);SMAD3
MAP2K2;P36507;ENSG00000126934;H. sapiens;Mediator;RTK(core);MAPK1;P28482;ENSG00000100030;H. sapiens;Mediato
```

Storing network data - file formats - SQL

- → Database management language
- → It can be used remotely for a database server
- → Can be edited manually and programmatically



Storing network data - file formats - XML

- → "Incomplete" tables are practical
- → "Tag" format
- → It is difficult to read, hand-handy

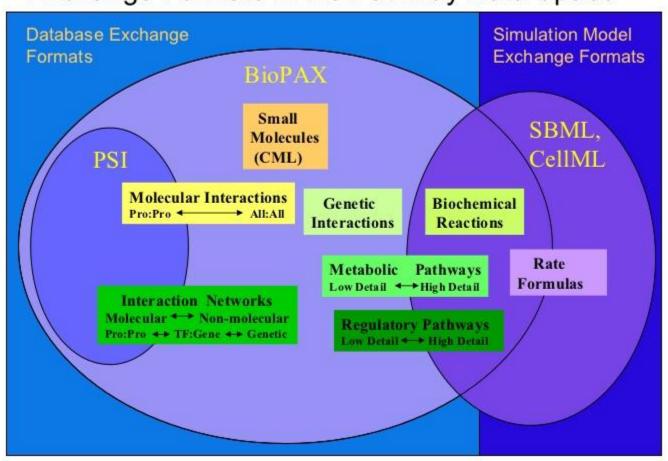
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         <unit kind="gram" exponent="-1"/>
         <unit kind="second" exponent="-1" multiplier="0.000277777777777778"/>
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     </unitDefinition>
   </listofUnitDefinitions>
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     <compartment id="c" name="cytosol" size="1"/>
     <compartment id="e" name="extracellular" size="1"/>
   </listOfCompartments>
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     <species id="M 12dgr120 c" name="1-2-Diacyl-sn-glycerol-didodecanoyl-n-C120" compartment="c" charge="/</pre>
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           KEGG ID: C00641
           PubChem ID: 3914
           ChEBI ID: 17815
```

Storing network data - file formats - PSI-MI

- Proteomics Standards Initiative Molecular Interaction
- → Standard format for expressing molecular contact data
- → Two format:
 - ♦ XML
 - ♦ TSV
- → MI ontology

Storing network data - file formats

Exchange Formats in the Pathway Data Space

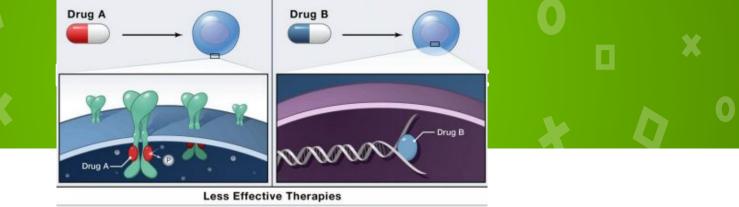


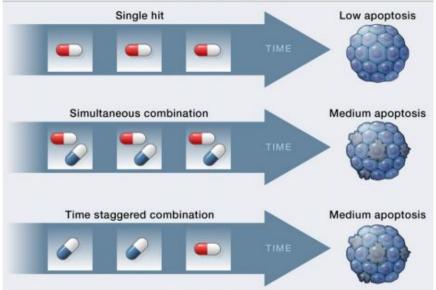


Papers (optional):

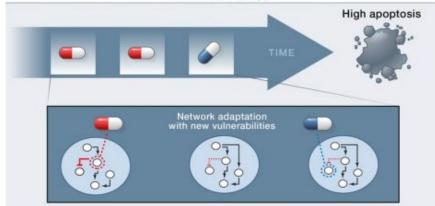
Michael J. Lee, Albert S. Ye, Alexandra K. Gardino, Anne Margriet Heijink, Peter K. Sorger, Gavin MacBeath, Michael B. Yaffe Sequential Application of Anticancer Drugs Enhances Cell Death by Rewiring Apoptotic Signaling Networks Cell, Volume 149, Issue 4, 11 May 2012, Pages 780-794

Preview: Janine T. Erler, Rune Linding Network Medicine Strikes a Blow against Breast Cancer Cell, Volume 149, Issue 4, 11 May 2012, Pages 731–733





Effective Therapy



Aplication of network biology

Combination Erl-Dox Nanoparticles in vivo

