



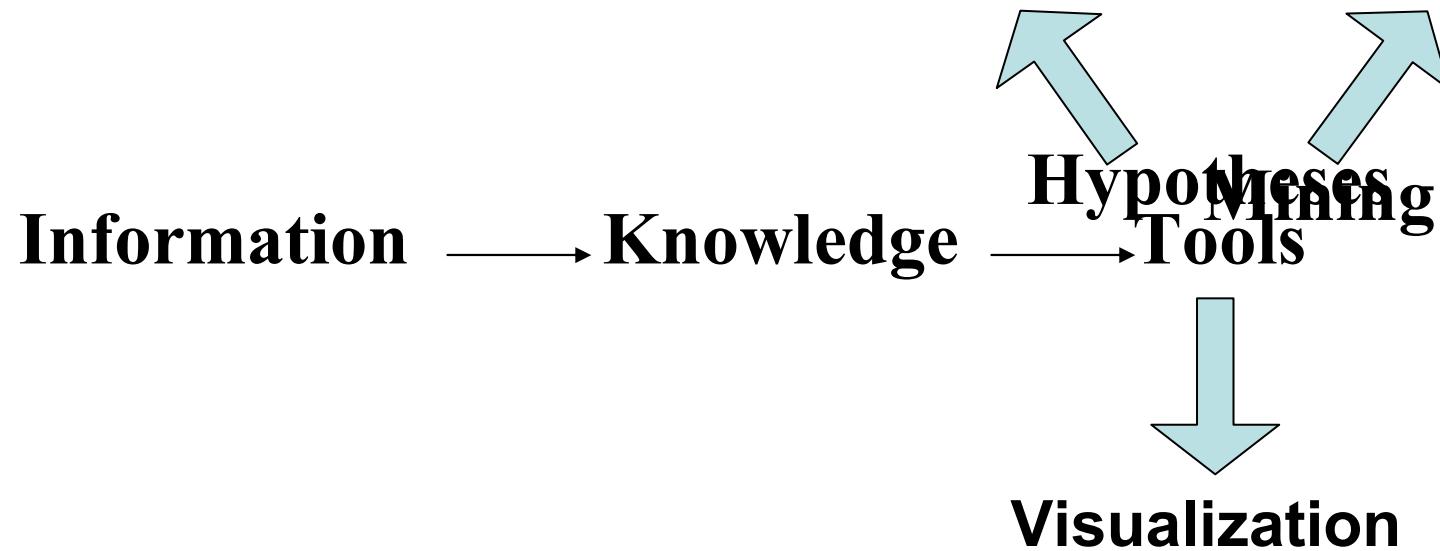
Strategies for Mining the NCI's Screening Databases: *Data (NCI60, Xenograft) Informatics (Bio and Chemo)*

Laboratory of Computational Technologies
*Anders Wallqvist, Ruili Huang, Narmada Thanki,
Xiang-Jun Lu, Alfred Rabow*

NIH/NCI/DCTD/DTP/STB
Drs. Doroshow, Collins, Shoemaker

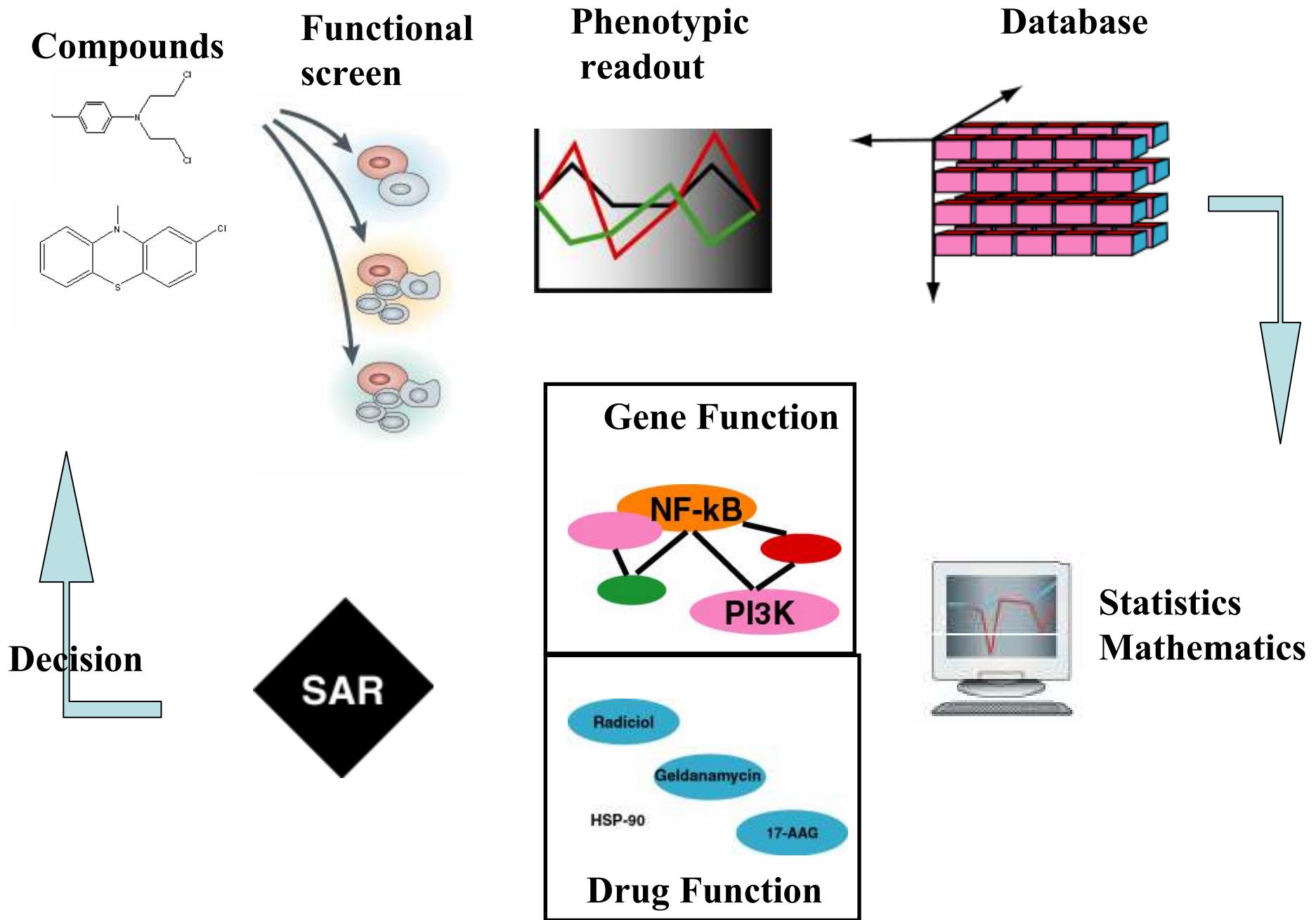


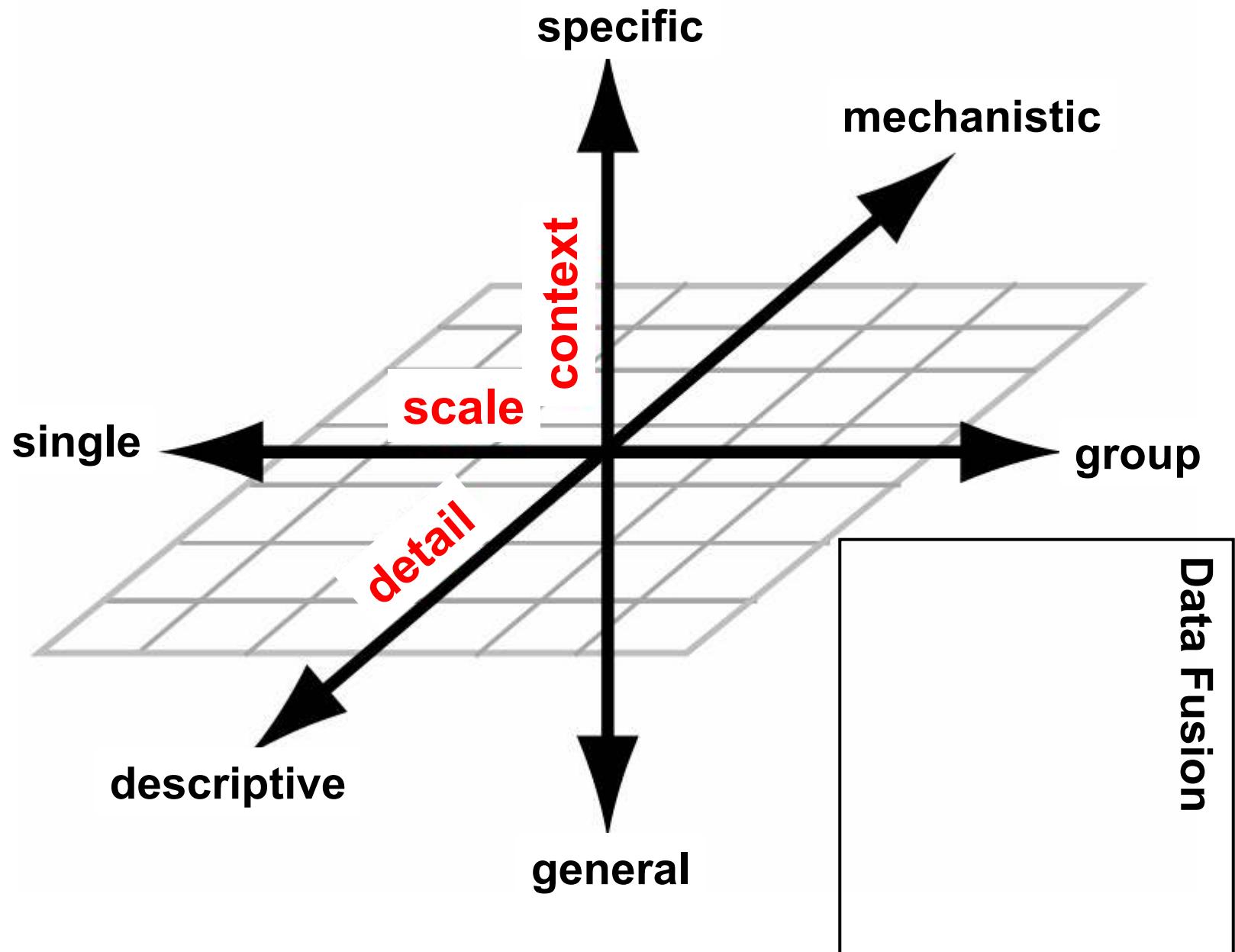
spheroid.ncifcrf.gov

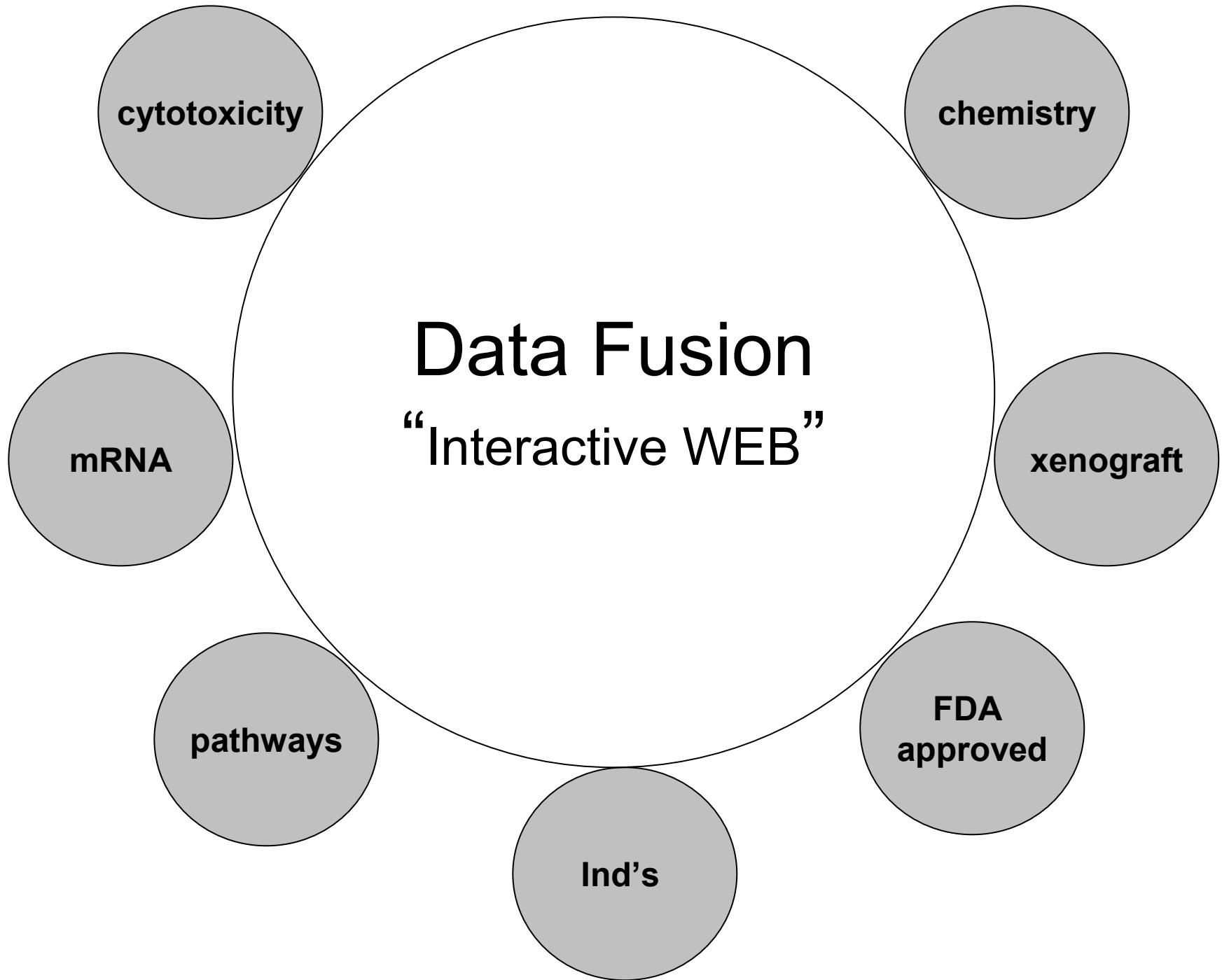


- Successes
- Pitfalls
- Strategies
- Recommendations

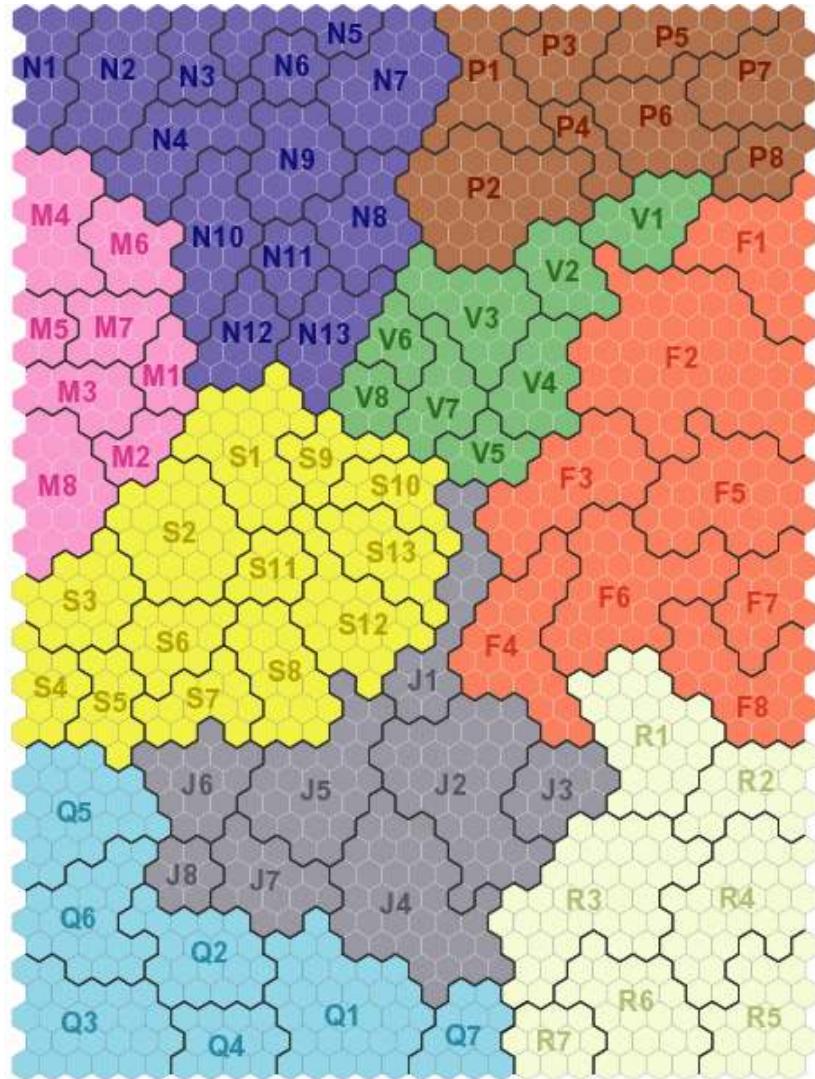
Data Generation / Data Analysis



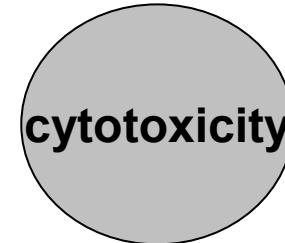




GI₅₀ SOM



Rabow et al. J Med Chem (2002)
Wallqvist et al. JCIM (2006)

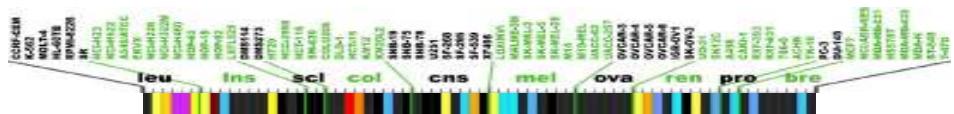


Successes

NCI₆₀: lung, renal, colorectal, ovary, breast, prostate, central nervous system, melanoma, leukemia

~100,000 compounds

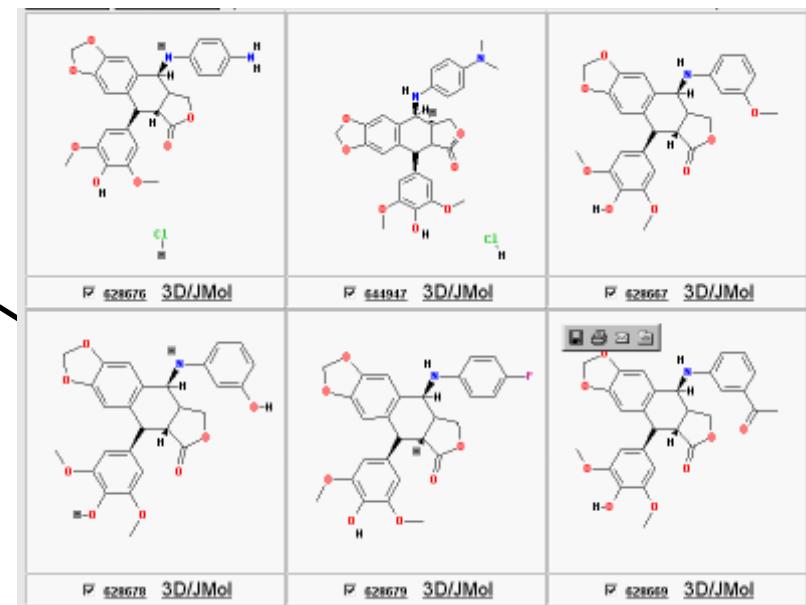
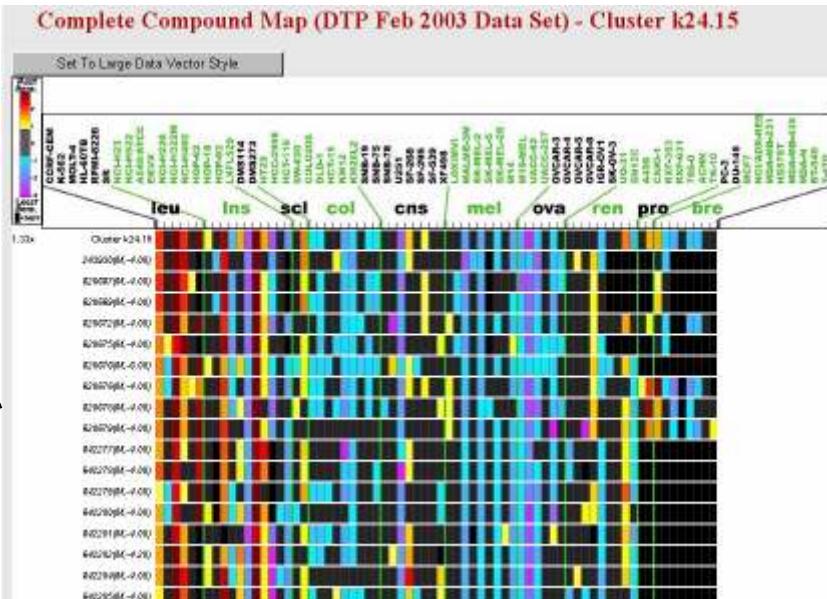
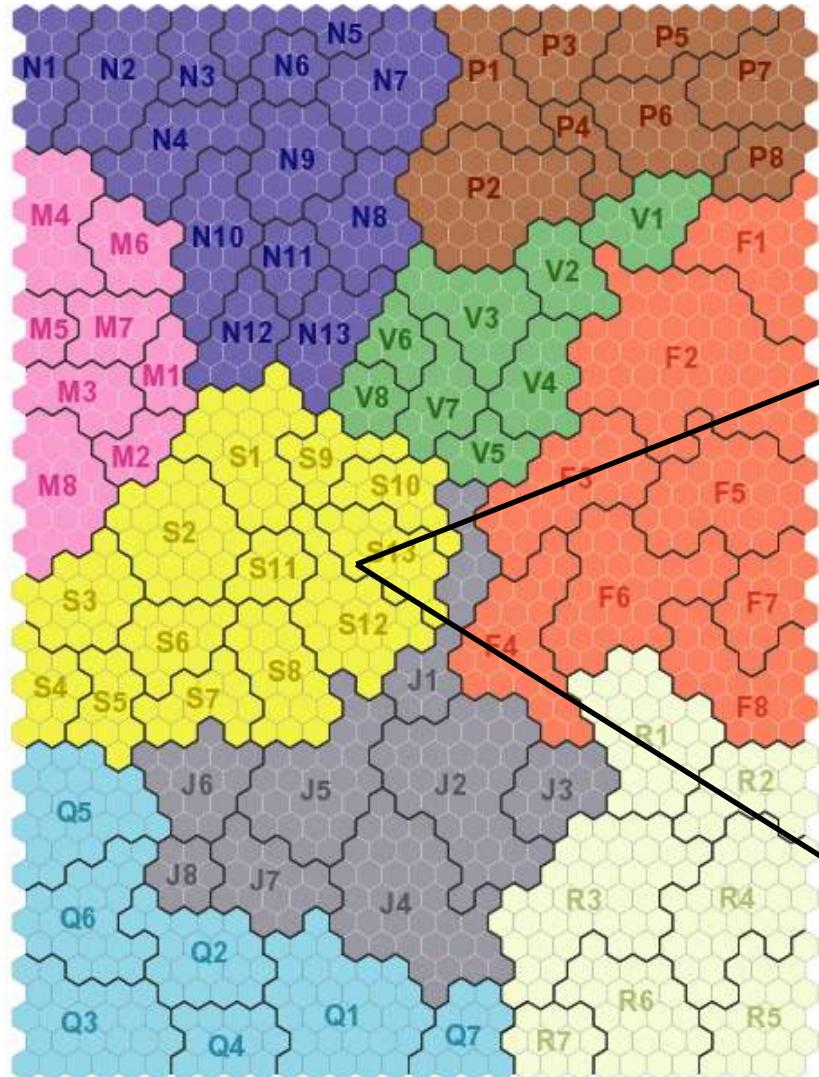
GI₅₀: 50% cancer cell growth inhibition concentration



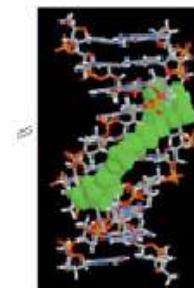
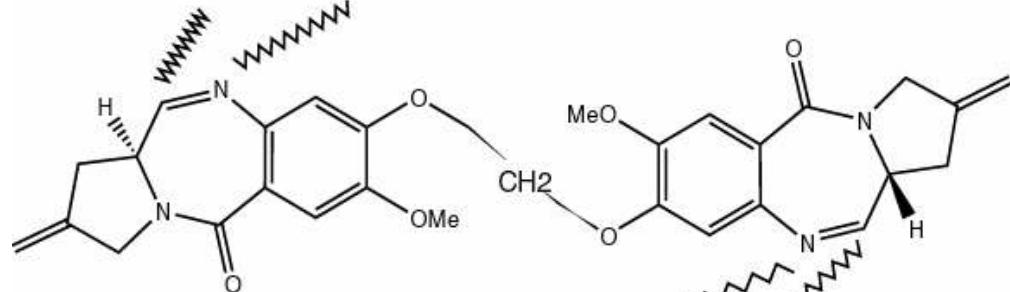
Legend to Cellular Growth Inhibition SOM

- Nucleic Acid Biosynthesis
- Xenobiotic Metabolism
- Membrane Transport/Integrity
- Mitosis
- Oxidative Metabolism
- Kinase/Phosphatase
- Work in Progress
-

GI₅₀ SOM



X ----- G ----- A ----- T ----- C ----- X



X

C

T

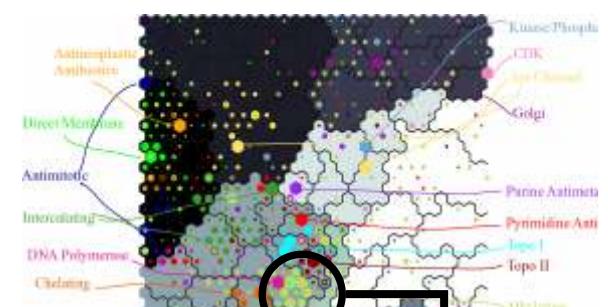
A

G

X

SJG-136 NSC694501

Neidle & Thurston Nat. Rev. 2005



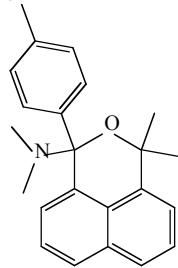
Alkylating

chemistry

success

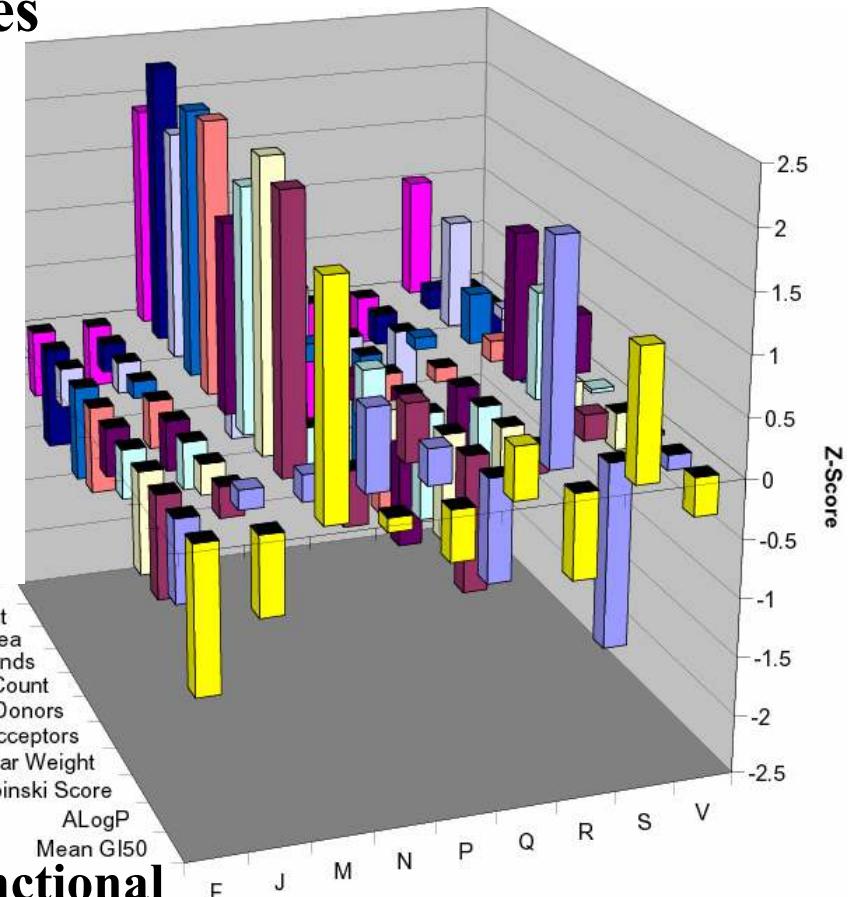
Huang et al. J Med Chem 49:1964-1979 (2006)

Atoms and bonds Physical properties



- Mean GI₅₀
- ALogP
- Lipinski Score
- Molecular Weight
- Hydrogen Bond Acceptors
- Hydrogen Bond Donors
- Parent Atom Count
- Rotatable Bonds
- Polar Surface Area
- Parent Molecular Weight
- LSUFC

LSUFC
Parent Molecular Weight
Polar Surface Area
Rotatable Bonds
Parent Atom Count
Hydrogen Bond Donors
Hydrogen Bond Acceptors
Molecular Weight
Lipinski Score
ALogP
Mean GI₅₀



Mechanism of Action Categories:

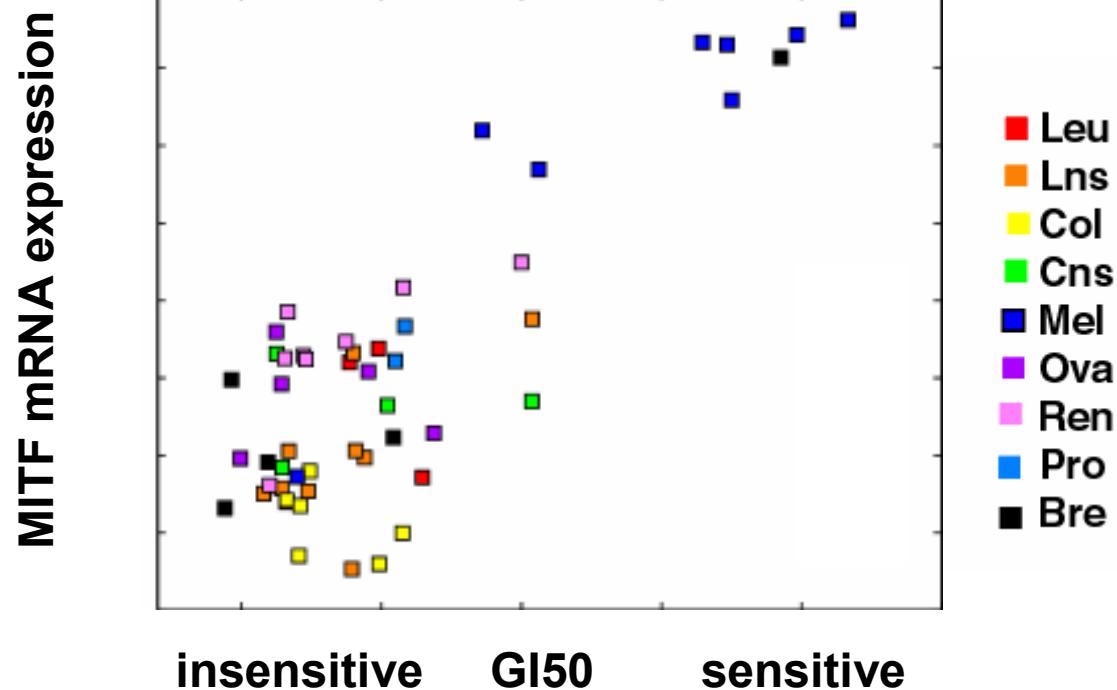
- [M] Anti-mitotic.....large and functional
- [S] DNA synthesis.....low lipophilicity
- [P] Phosphatase/kinase.....most diverse signal
- [R] Membrane active.....high lipophilicity
- [Q] Xenobiotic metabolism...reactive groups

Chemistries: Modeling GI50

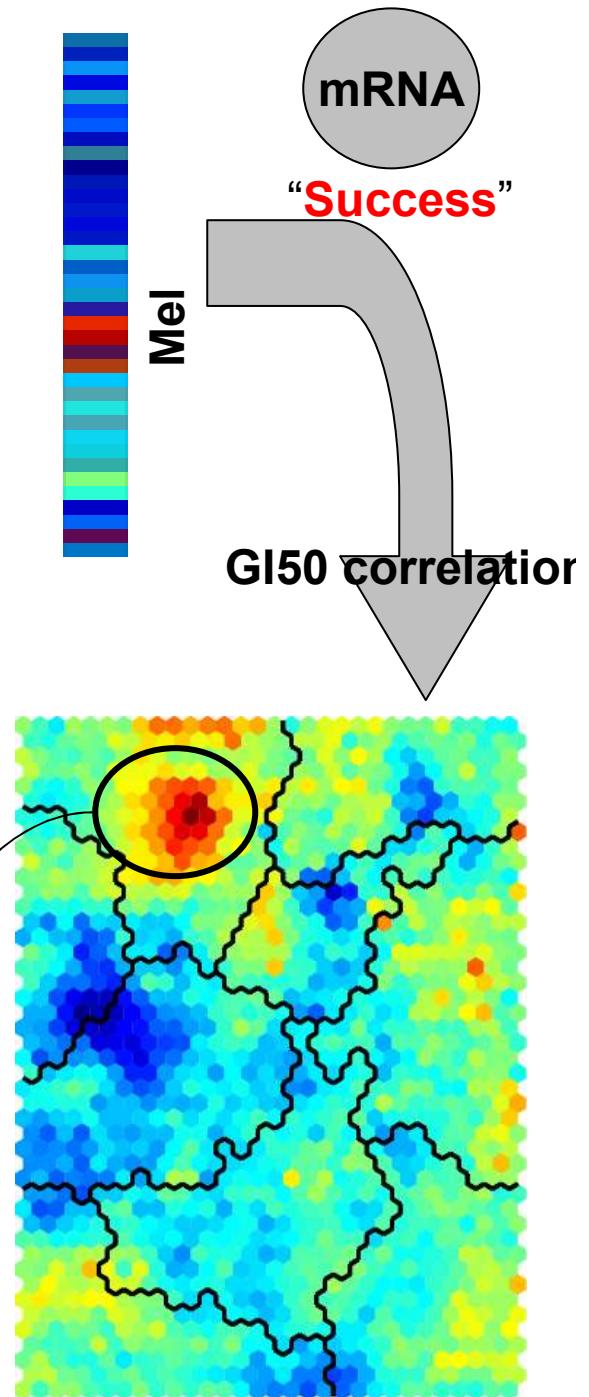
Selecting potent compounds

- large (many previous studies)**
- complex: many features
(Oprea, Blake, Veber, Veith)**
- local effects (SOM regionalization)
➤ potency scales with selectivity
(Huang et al.)**

Gene Expression vs GI50

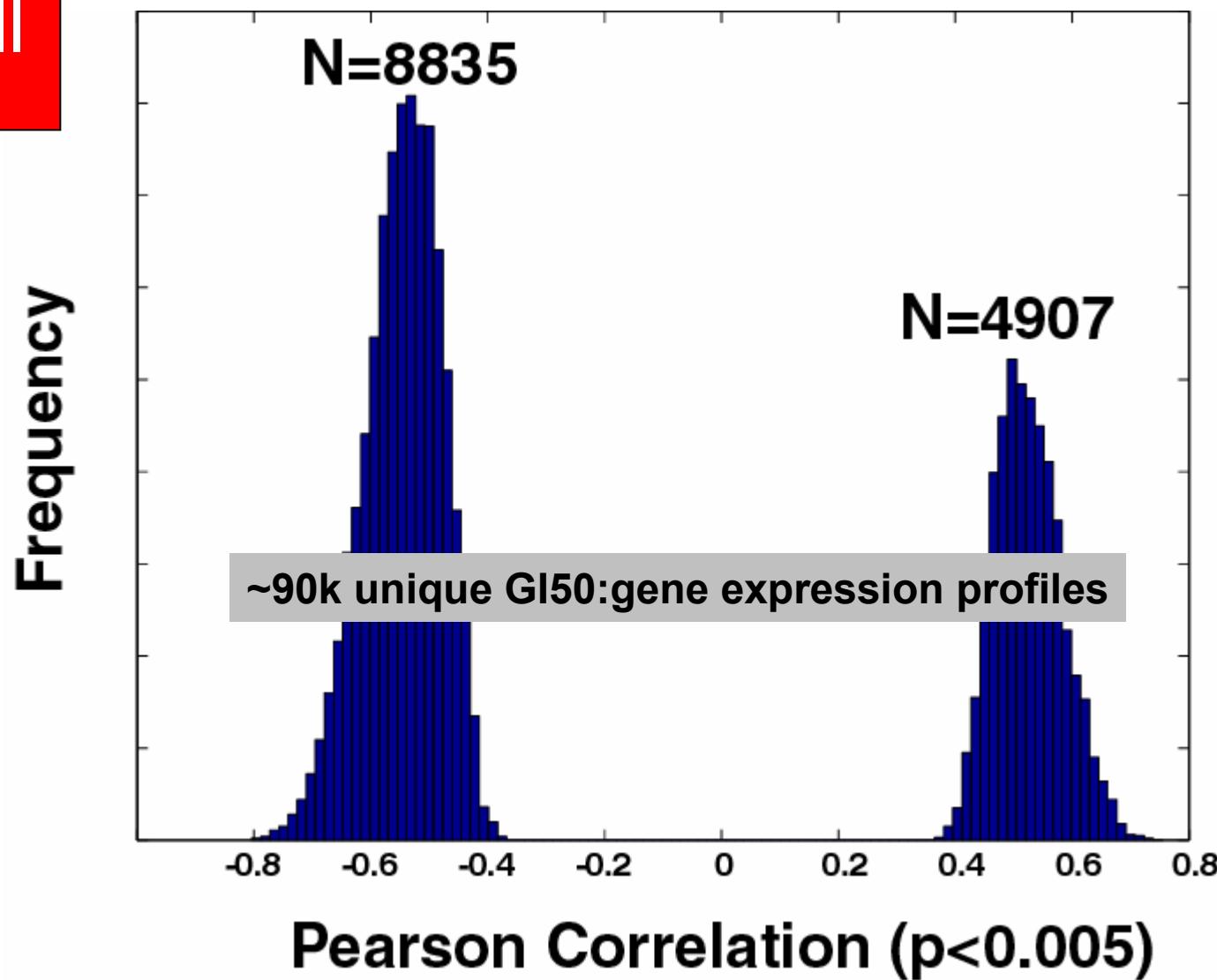


6 NSCs selected from highest + correlations
Hypothemycin, LF, PD98059
Rosen et al.
Sellers et al.

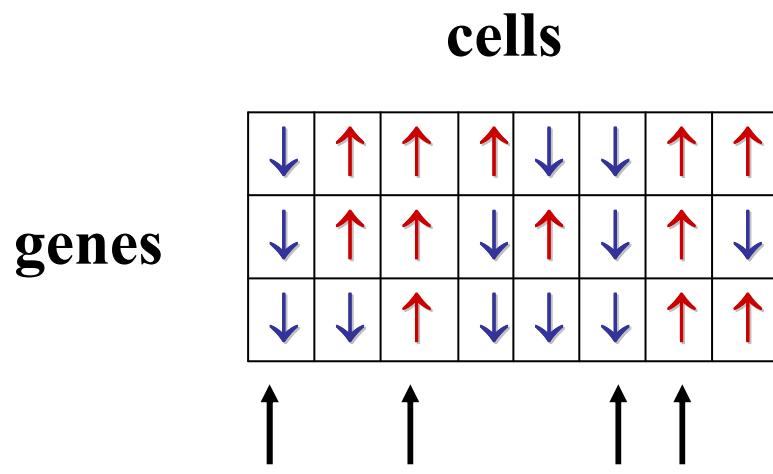


Pitfall

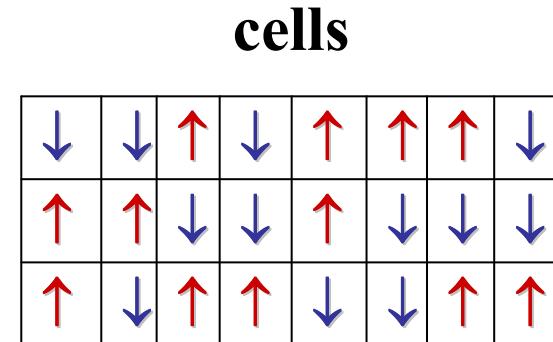
GI50:Gene Expression Correlations



Linking Pathway Gene Expression to GI₅₀



coherent pathway



non-coherent pathway

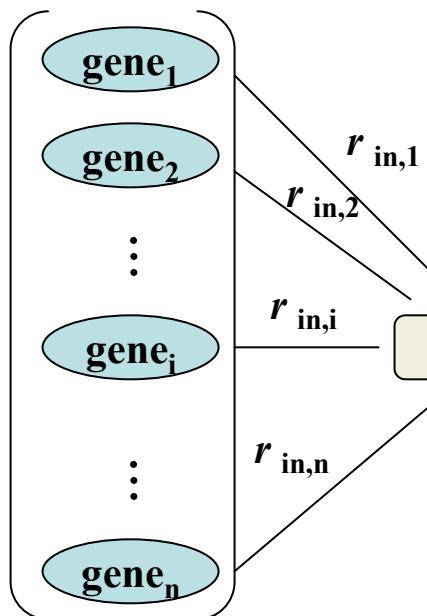
Huang et al. Genomics 87:315-328 (2006)

Linking Pathway Gene Expression to GI₅₀

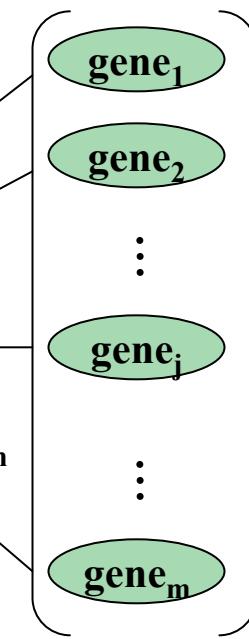
Huang et al. Genomics 87:315-328 (2006)

For pathway P :

Genes *in P*



Genes *not in P*



Pearson Correlation: r

$$R_{in} = \{r_{in,1}, r_{in,2}, \dots, r_{in,i}, \dots, r_{in,n}\}$$

↔ Kruskal-Wallis → H, p

$$R_{out} = \{r_{out,1}, r_{out,2}, \dots, r_{out,j}, \dots, r_{out,m}\}$$

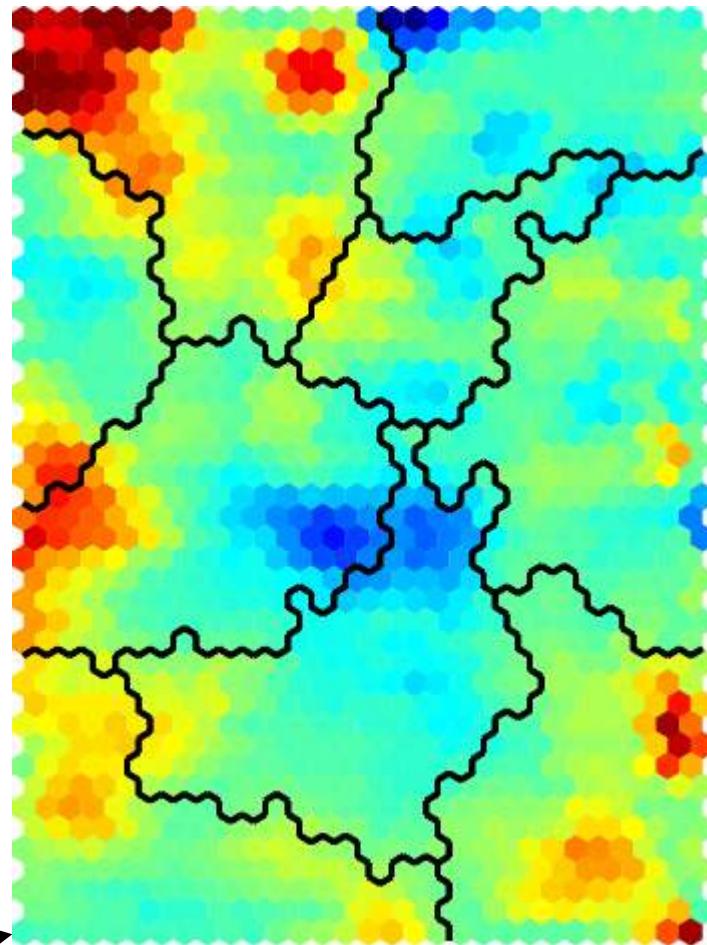
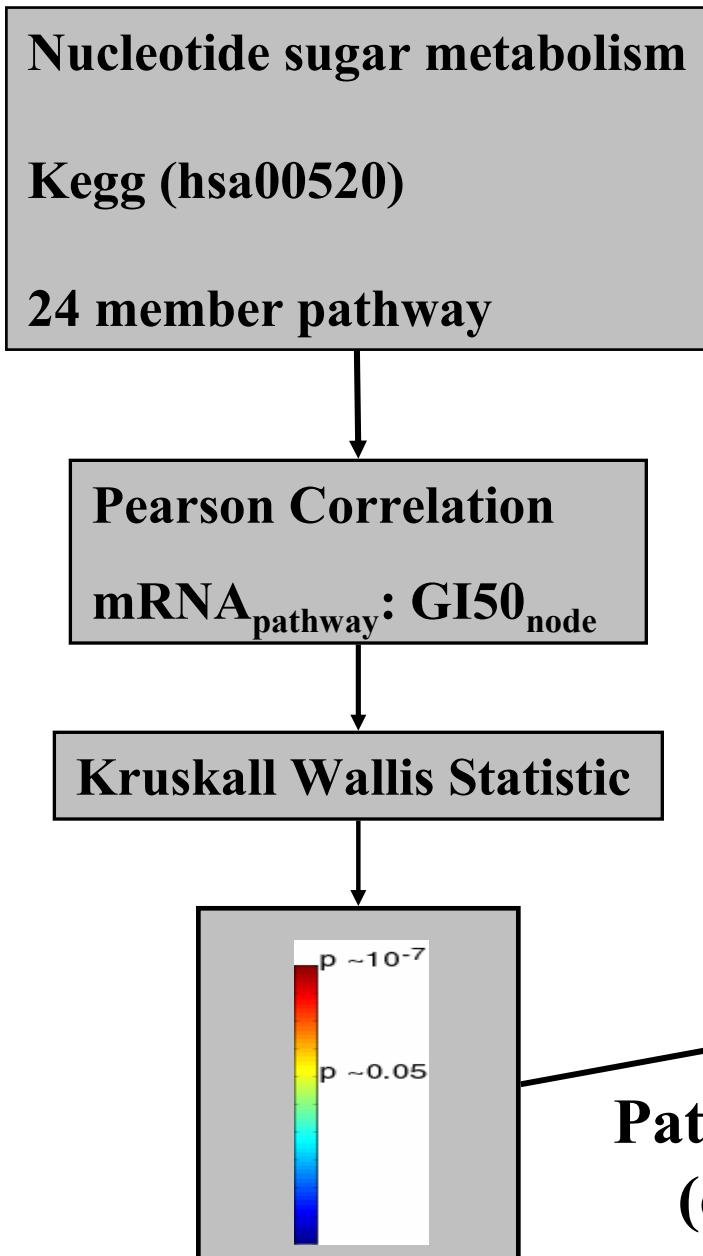
$$R_{in} > R_{out} \Rightarrow H > 0$$

$$R_{in} < R_{out} \Rightarrow H < 0$$

Drug is significantly associated with P if: $H > 0$ and $p < 0.05$

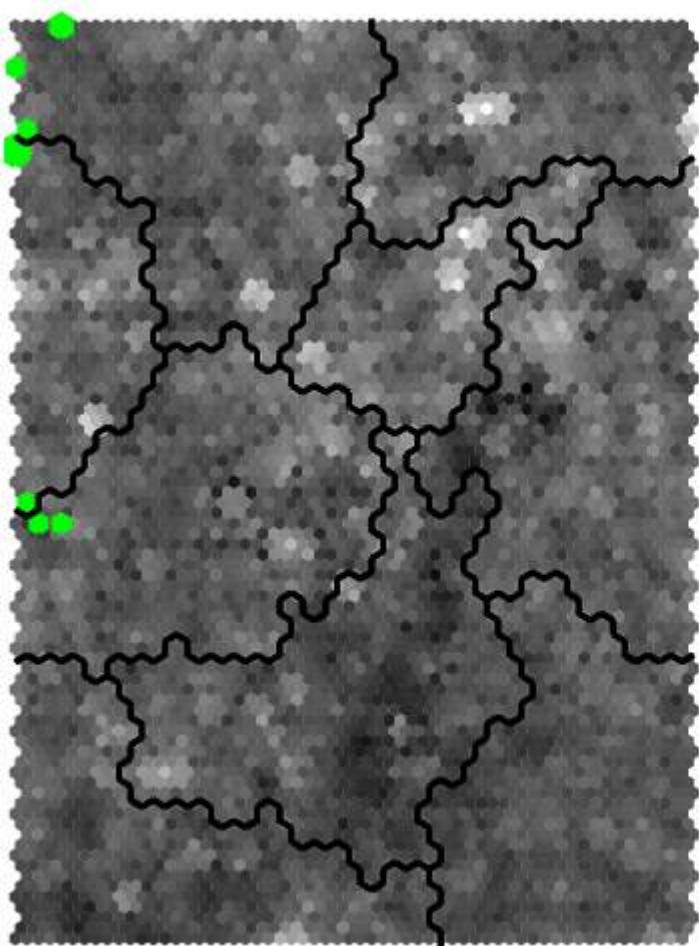
H defines a Fitness Score for pathways against GI₅₀

Relating Fitness Scores to Drug Response

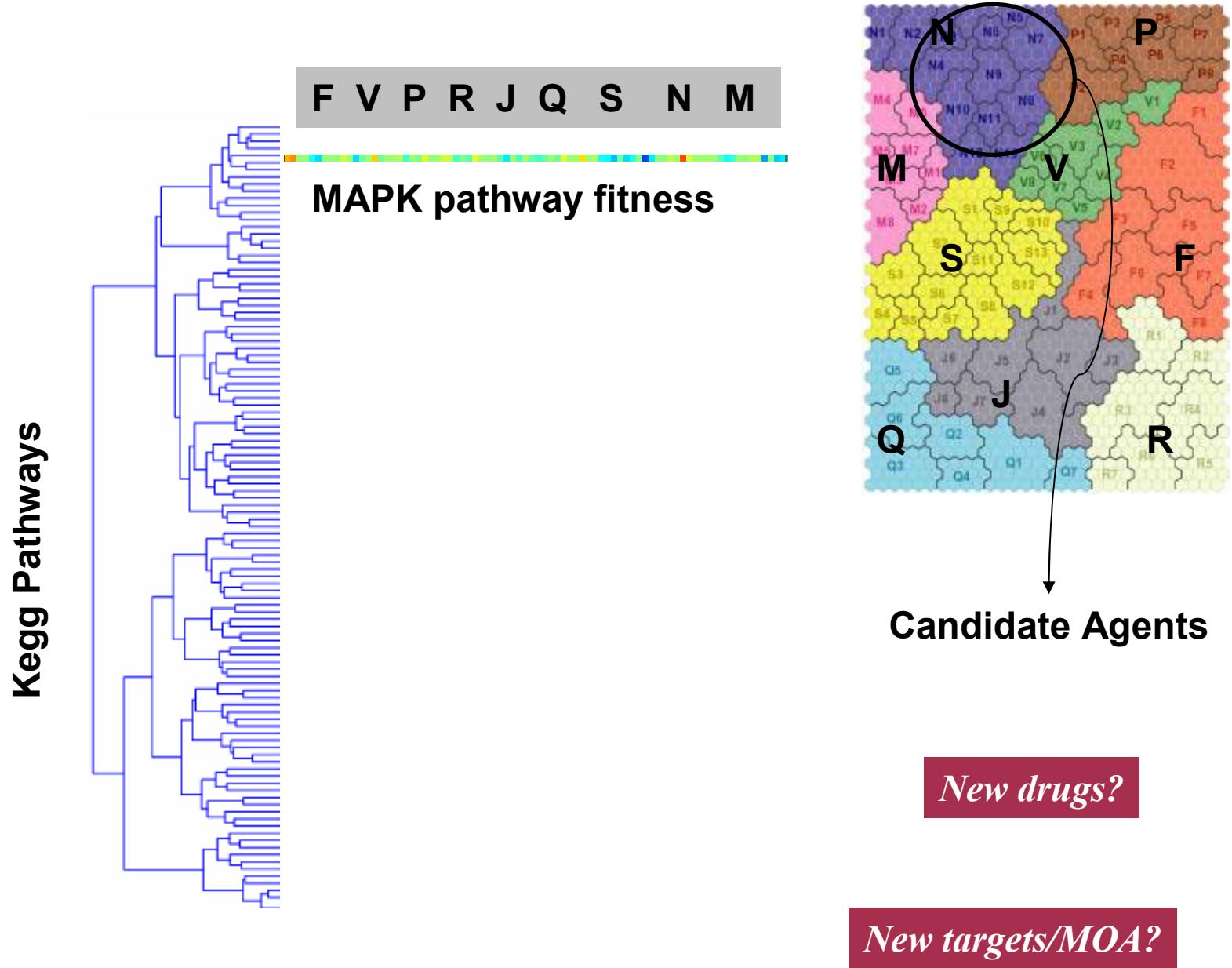


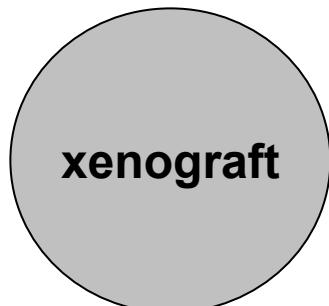
Pathway Fitness
(coherence)

Potential inhibitors of L-asparaginase
biosynthesis: Mokotoff JMC, 1981,
Richards, Ann Rev, 2006



● Inhibitors and structural analogs





success

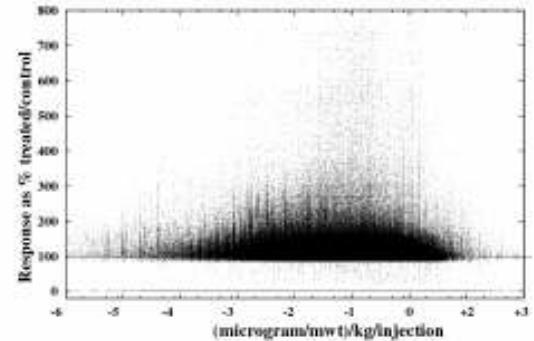
Xenograft data

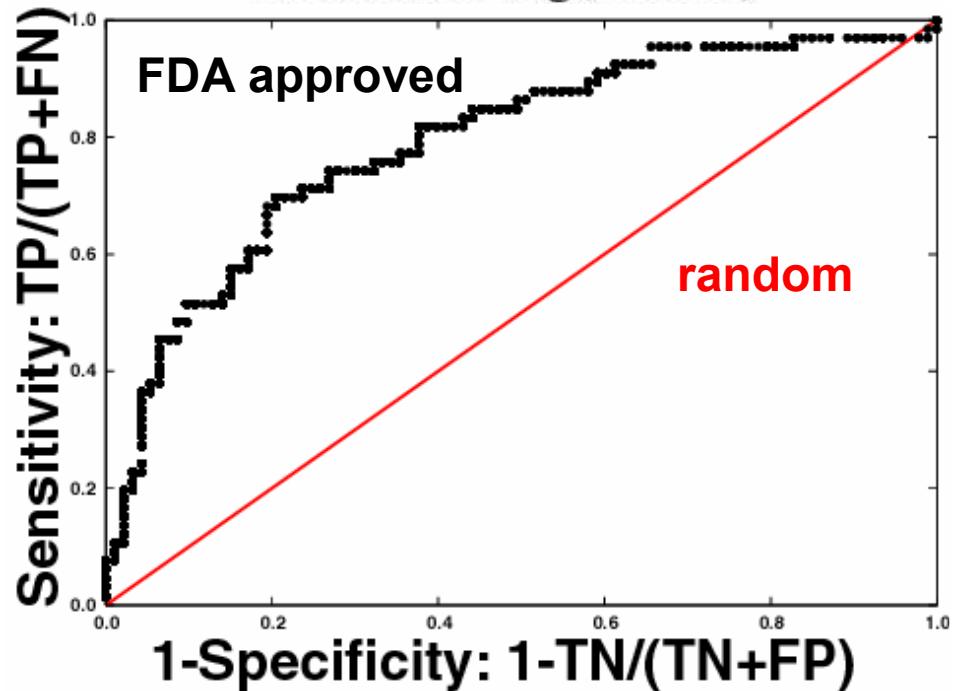
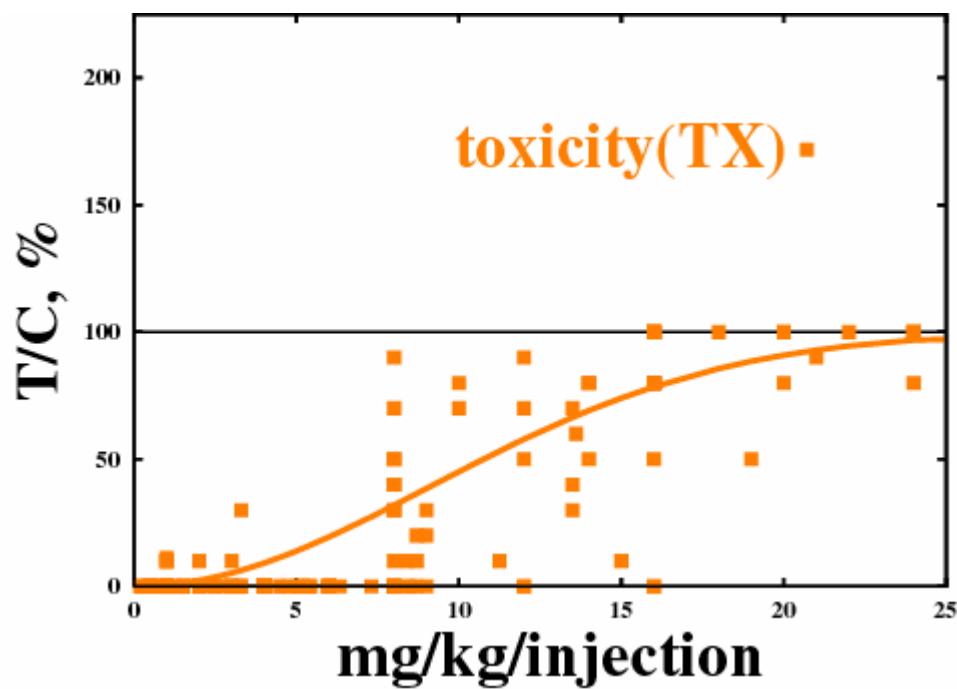
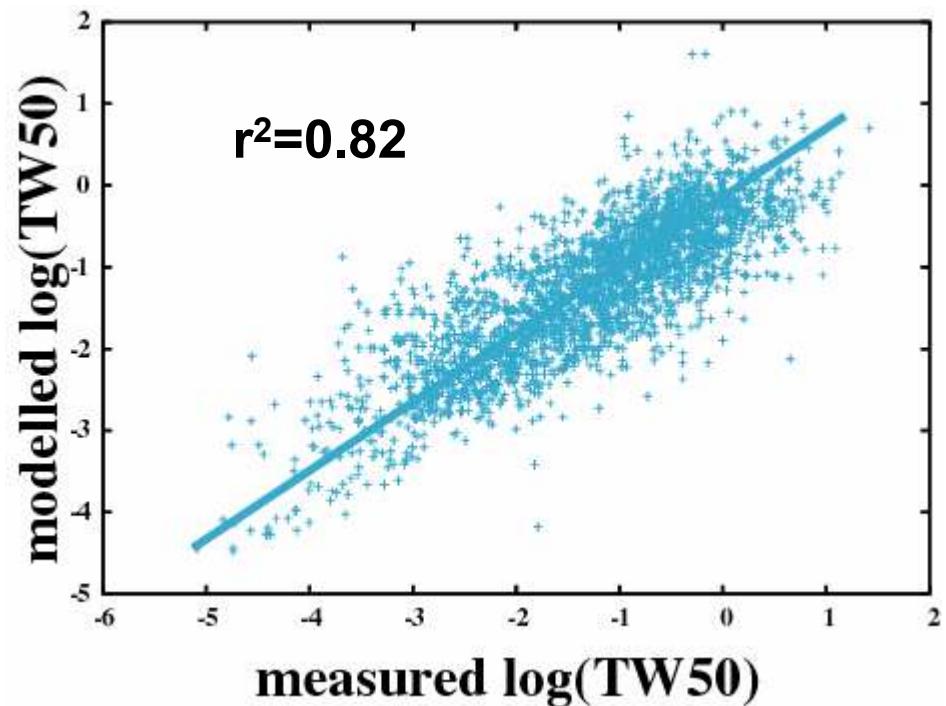
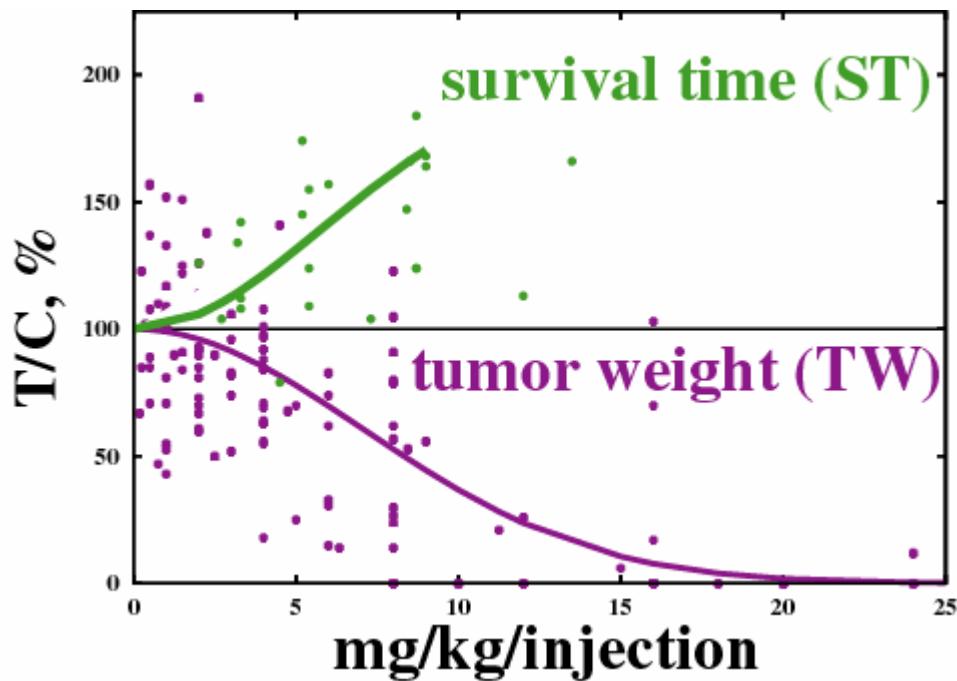
Experimental Design

- 1363 NSC tested
- 31 formulations
- 187 treatment schedules
- 50 tumor models
- 6 implantation sites
- 15 mice strains
- >5,000 combinations of experiments

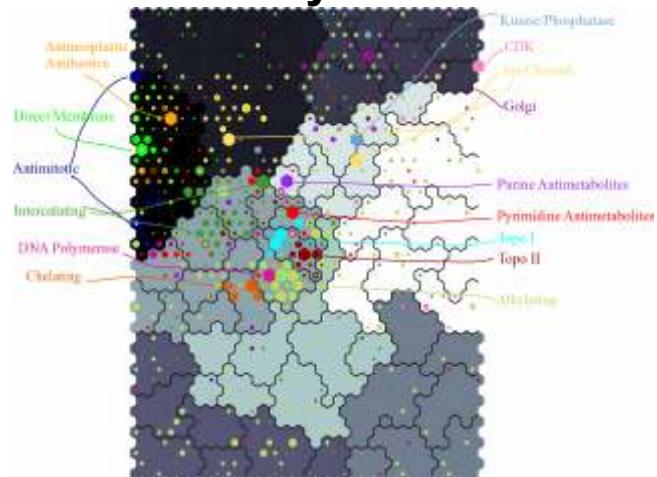
Measurements

- Tumor weight reduction (TW50)
- Survival time (ST150)
- Toxicity (survival _{control vs treatment})
- Therapeutic index (TW50,ST150/Tox)

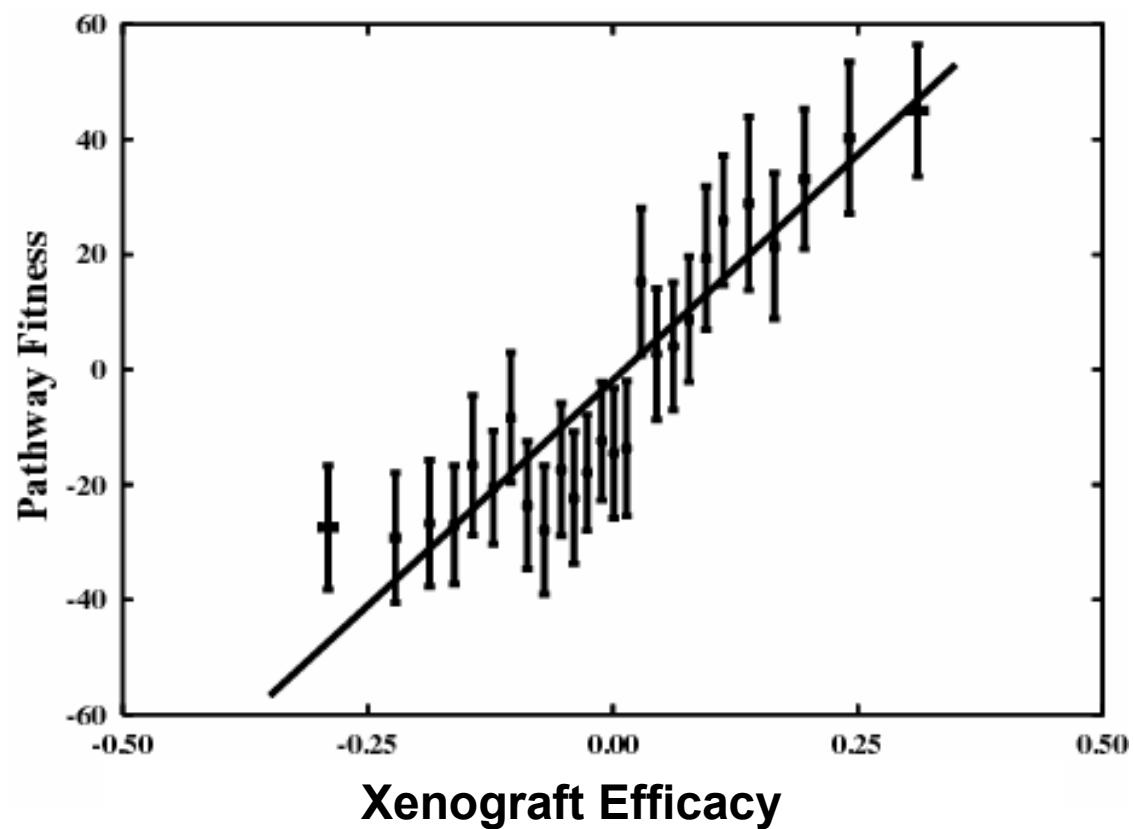
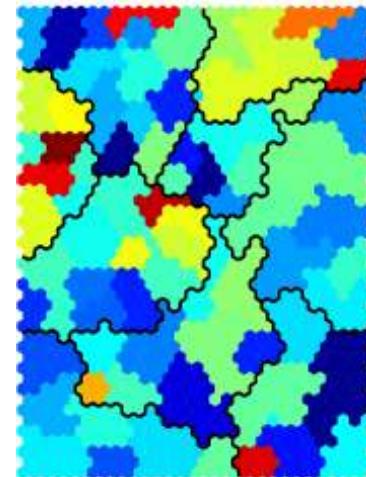


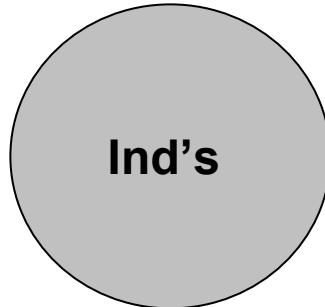


Activity Class



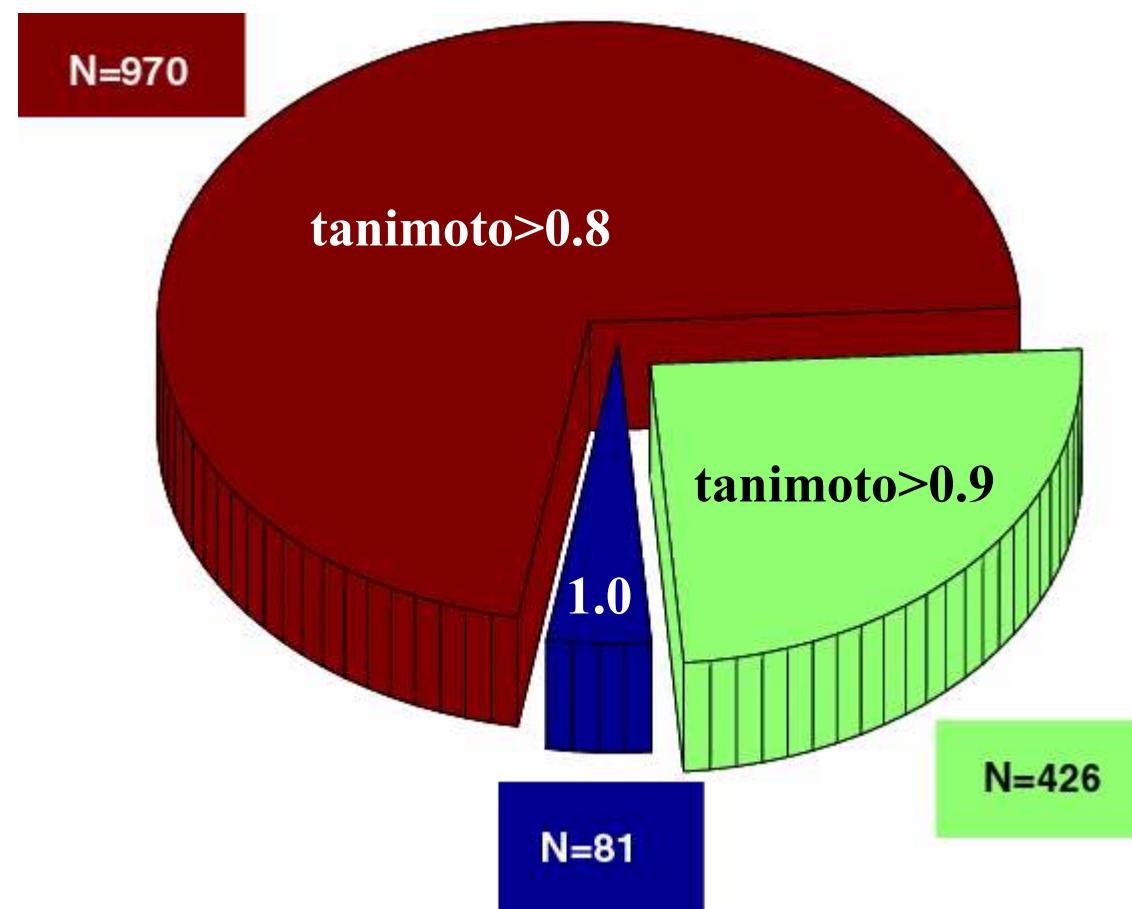
Fitness Scores





399 Anticancer Medicines in Development
(283 nonbiologicals) 123 (45%) have structural
analogs in NCI screening set www.phrma.org

success



Recommendations

Statistics:

- **beyond sorting**
- **clustering**
 SOM
- **decision trees**
 random forests
- **curse of dimensionality**
 false positives
 positive predictive value

Data Sharing

- **chemistry**
- **gene expression**
- **mutation**
- **SNP**
- **'cancer genes'**
- **negative results**

Reverse mining

- **retrospective testing**
 clinical trials
 preclinical data

microRNA

Toxicology

Clinical trials

Gene expression

SNP copy number

Cellular growth inhibition

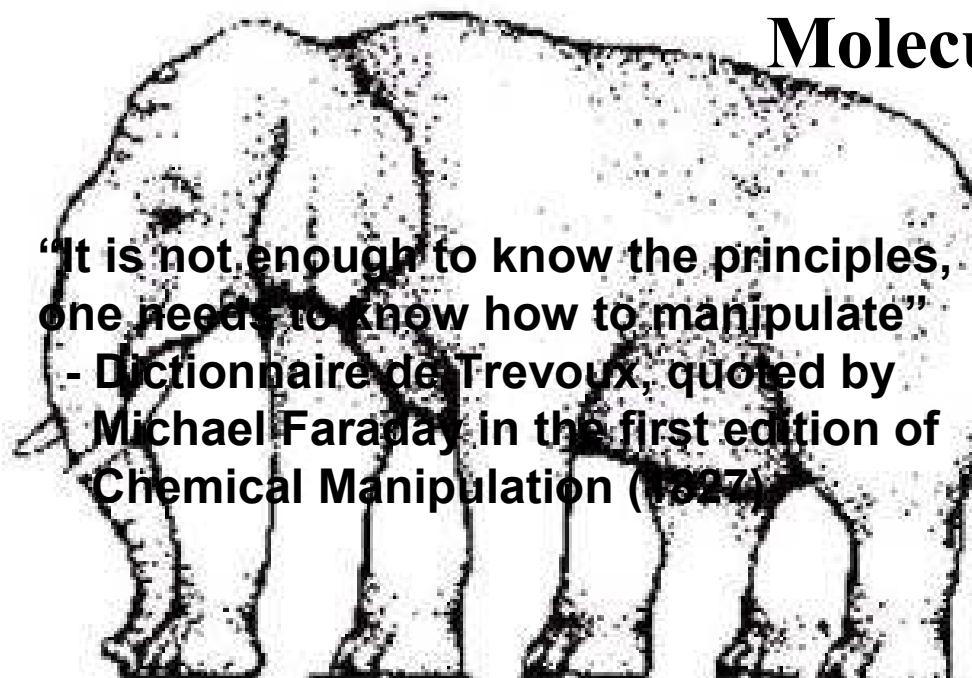
Molecular properties

Proteomics

Xenografts

Karyotype

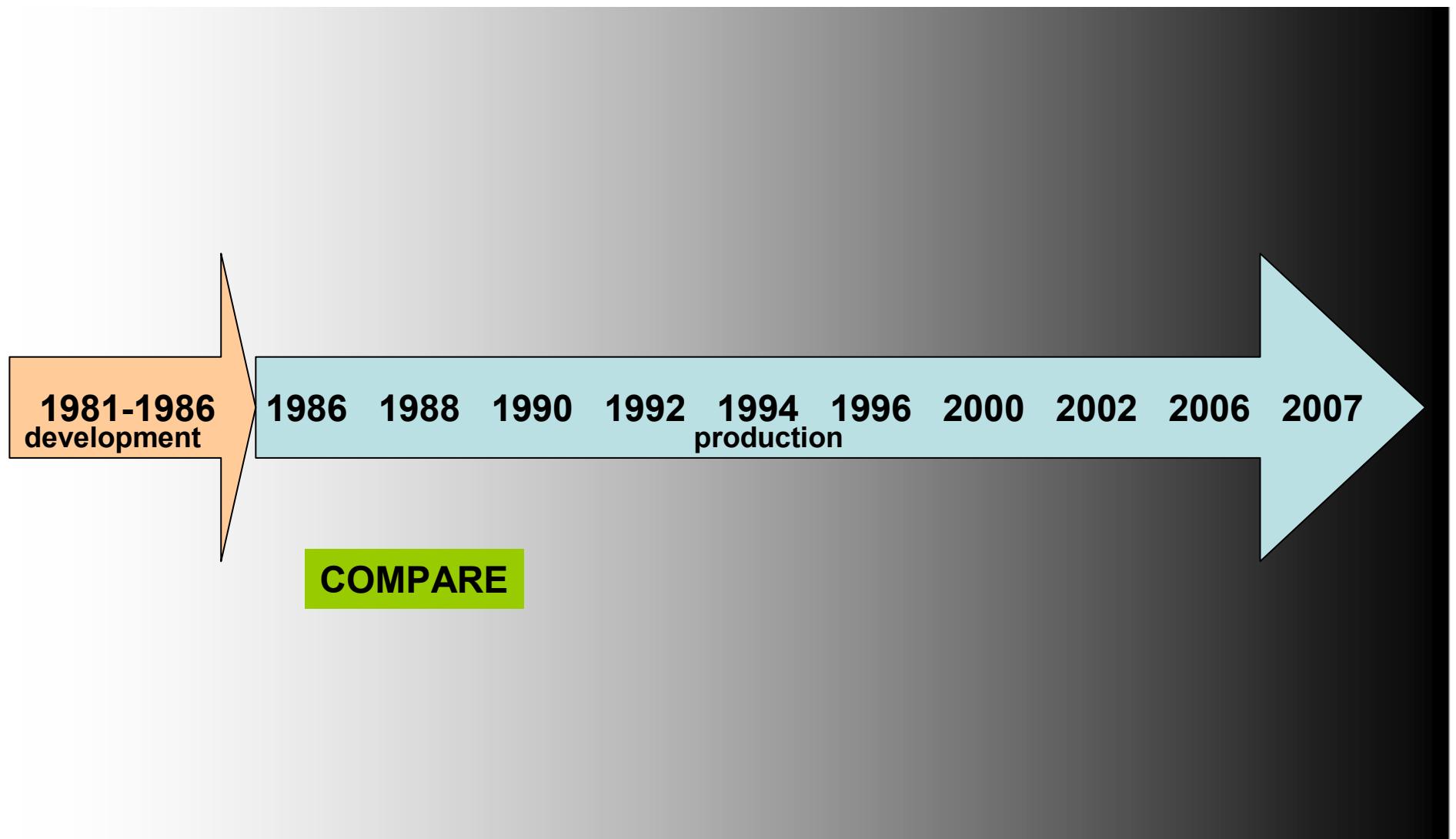
Methylation status



**'It is not enough to know the principles,
one needs to know how to manipulate'**
- Dictionnaire de Trevoux, quoted by
Michael Faraday in the first edition of
Chemical Manipulation (1827)

NCI-60 Timeline

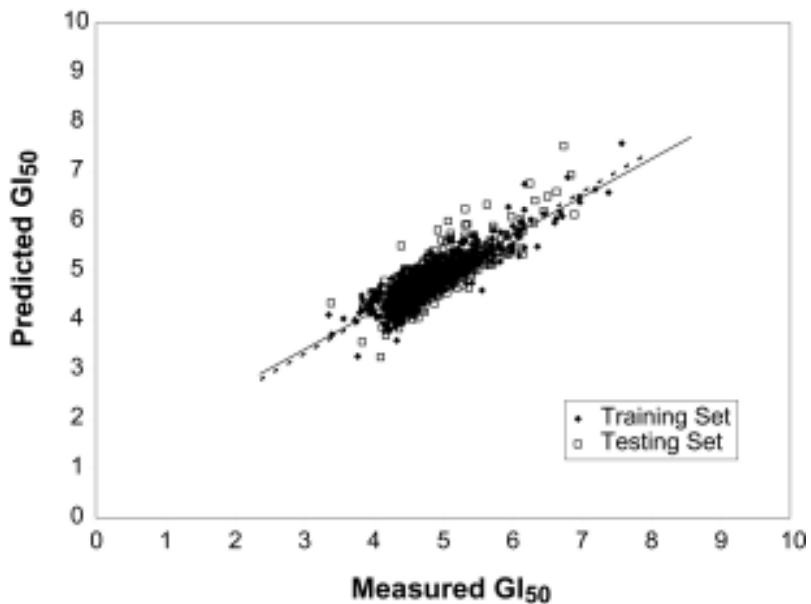
Shoemaker, Nat. Rev. Cancer, 2006



Chemistries: Modeling GI50

$$GI_{50} = F_1(\text{properties}) = c_1x_1 + c_2x_2 + \dots + c_Nx_N$$

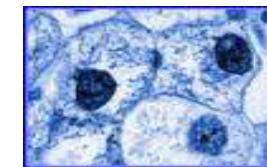
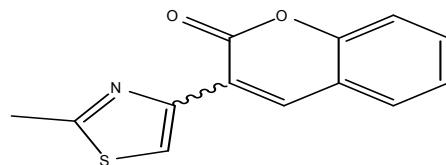
AlogP Mwt LSUFC



Training: $r^2 = 0.77$
Testing: $r^2 = 0.67$

Xenograft data

Rx



outcome = B × [(treatment) (chemistry)(cellular growth inhibition)]

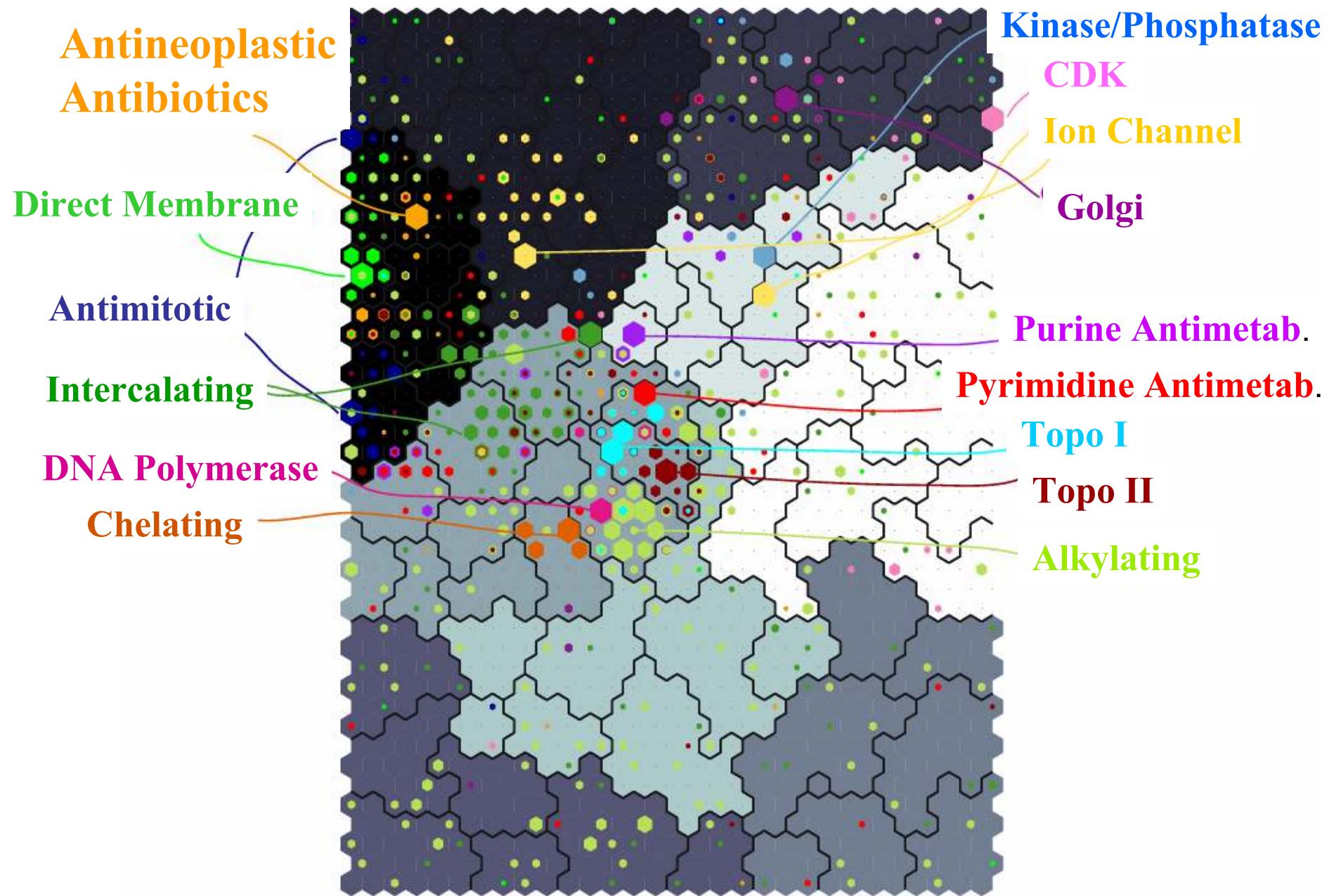
exptl design

properties

GI50

Treatment variations alone account for a log order of difference in efficacy outcome

Molecular Classes



	Act 1	Act 2	Act 3	Interm.	Act 4
GI50	✓	✓			
Chemistry		✓			
mRNA					
Pathways					
Xenograft					

	Act 1	Act 2	Act 3	Interm.	Act 4
GI50	✓	✓	✓	✓	✓
Chemistry		✓	✓		✓
mRNA			✓	✓	✓
Pathways				✓	✓
Xenograft					✓

	Act 1	Act 2	Act 3	Interm.	Act 4
GI50	✓	✓	✓		
Chemistry		✓	✓		
mRNA			✓		
Pathways					
Xenograft					

Chemistry Meets Biology

	Act 1	Act 2	Act 3	Interm.	Act 4
GI50	✓	✓	✓	✓	
Chemistry		✓	✓		
mRNA			✓	✓	
Pathways				✓	
Xenograft					

Pathway Fitness - Cohesiveness

- Relationship between the number of genes in a pathway that are shared with other pathways and the cohesiveness of the pathway

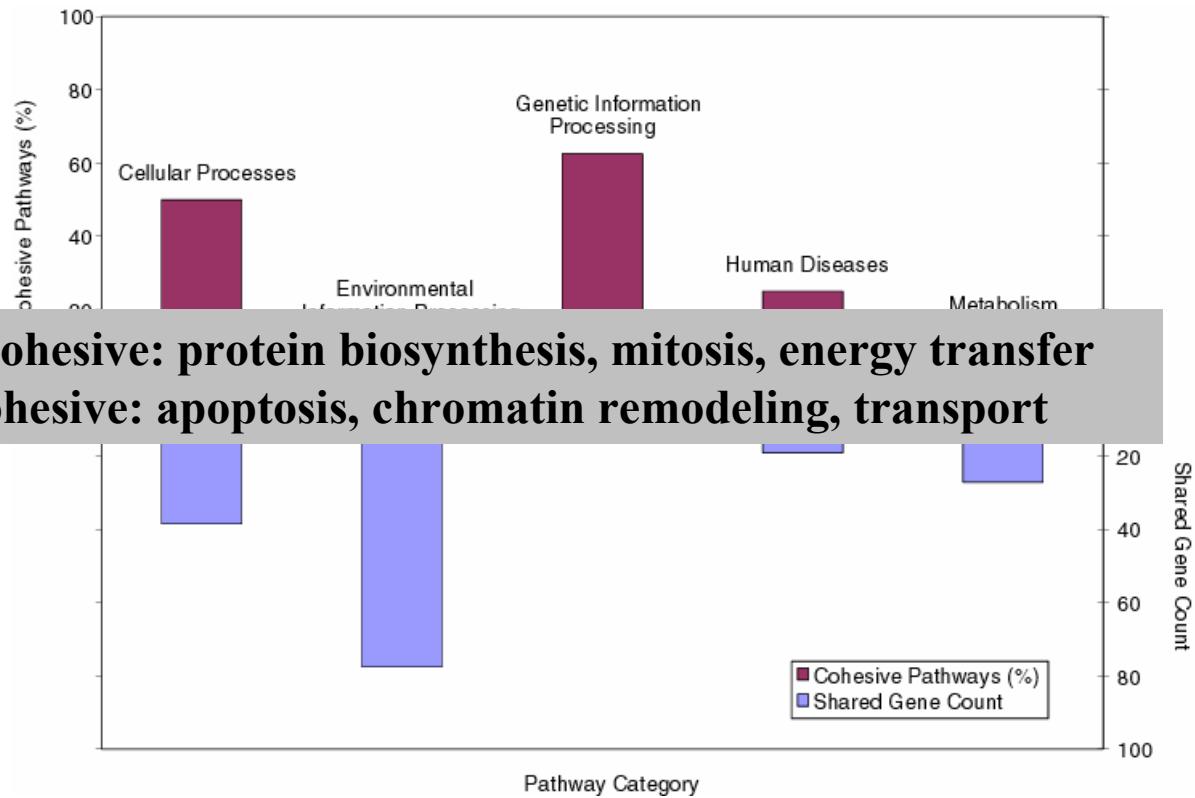
- Genetic Information Processing

- highest percentage of cohesive pathways
- least number of shared genes

More cohesive: protein biosynthesis, mitosis, energy transfer
Less cohesive: apoptosis, chromatin remodeling, transport

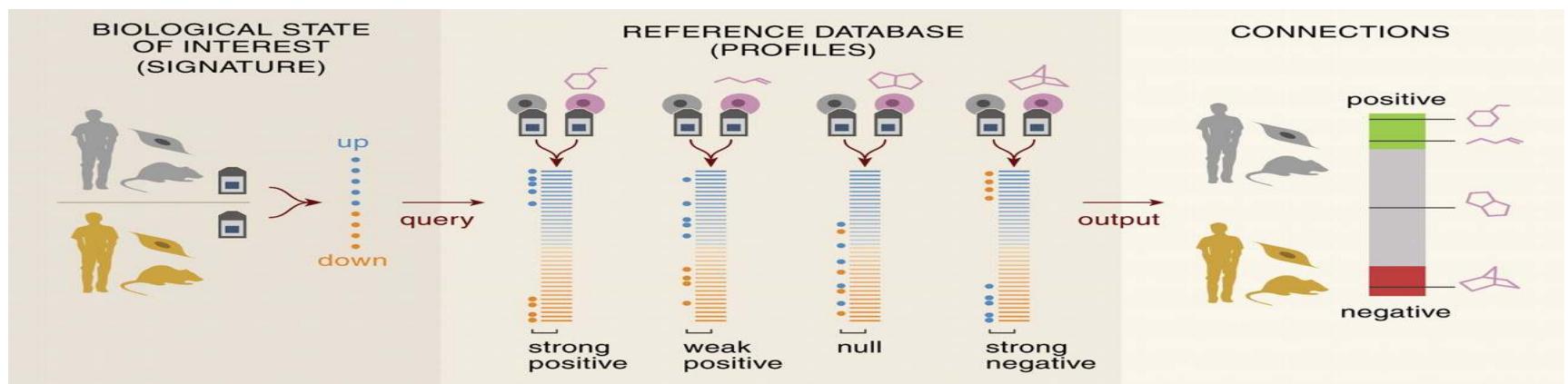
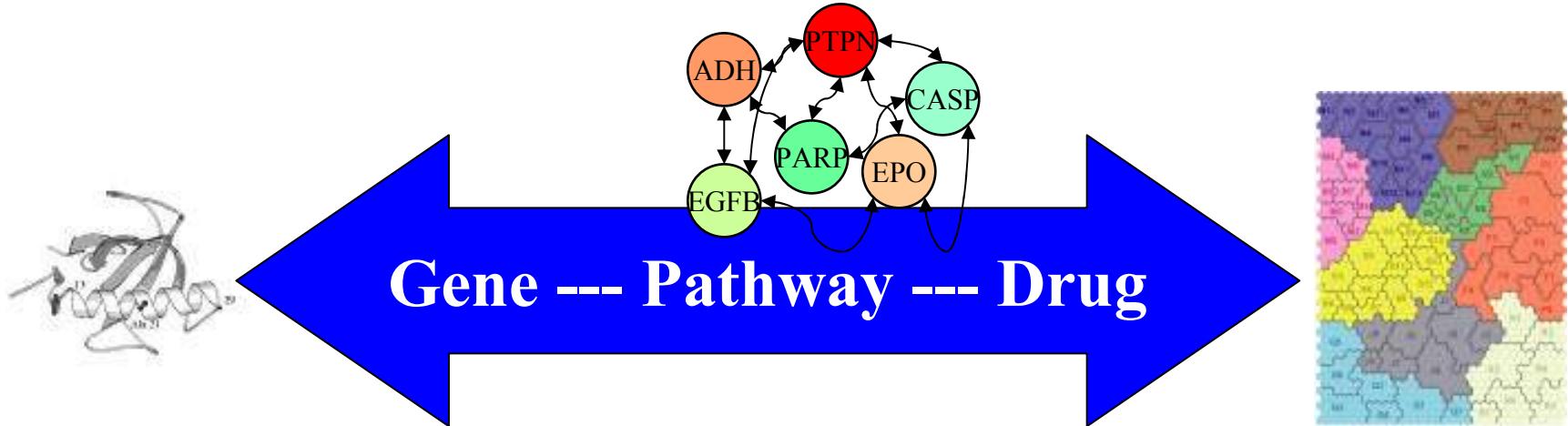
- Environmental Information Processing

- lowest percentage of cohesive pathways
- largest number of shared genes

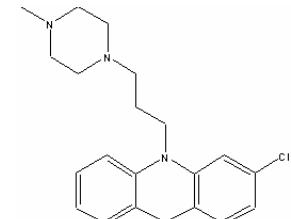
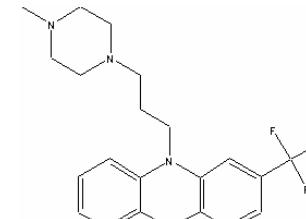
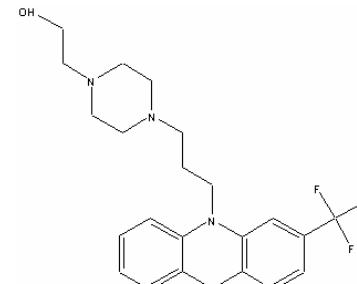
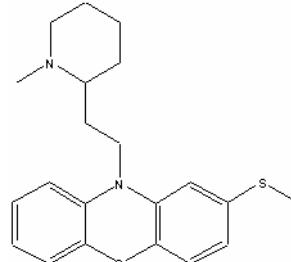
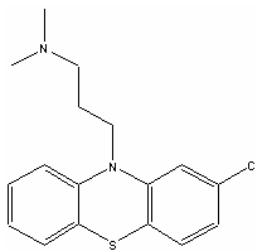


Huang et al. Genomics (2006)

Huang et al. Mol. Cancer Therapeutics (2006)



Connectivity Maps
Lamb et al., 2006



chlorpromazine

thioridazine

fluphenazine

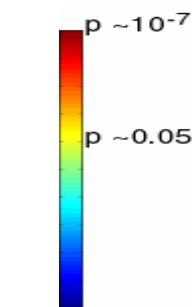
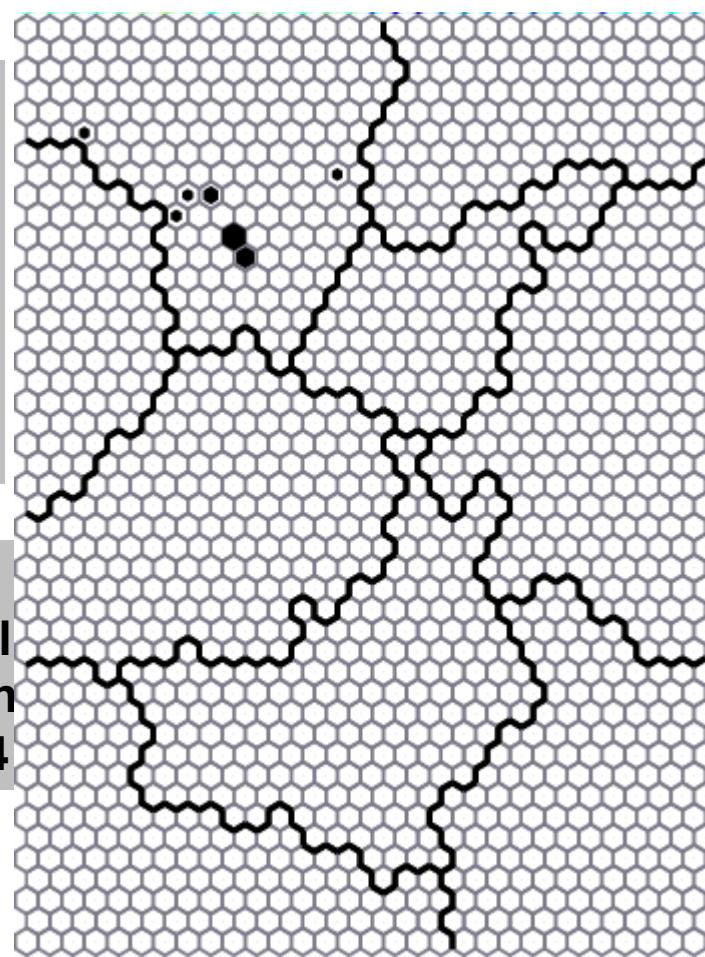
trifluoperazine

prochlorperazine

GO:3707 Steroid hormone receptor activity (PPARG, RXR, ESRR)

GO:199992 Diacylglycerol Binding (DAK, PKC)

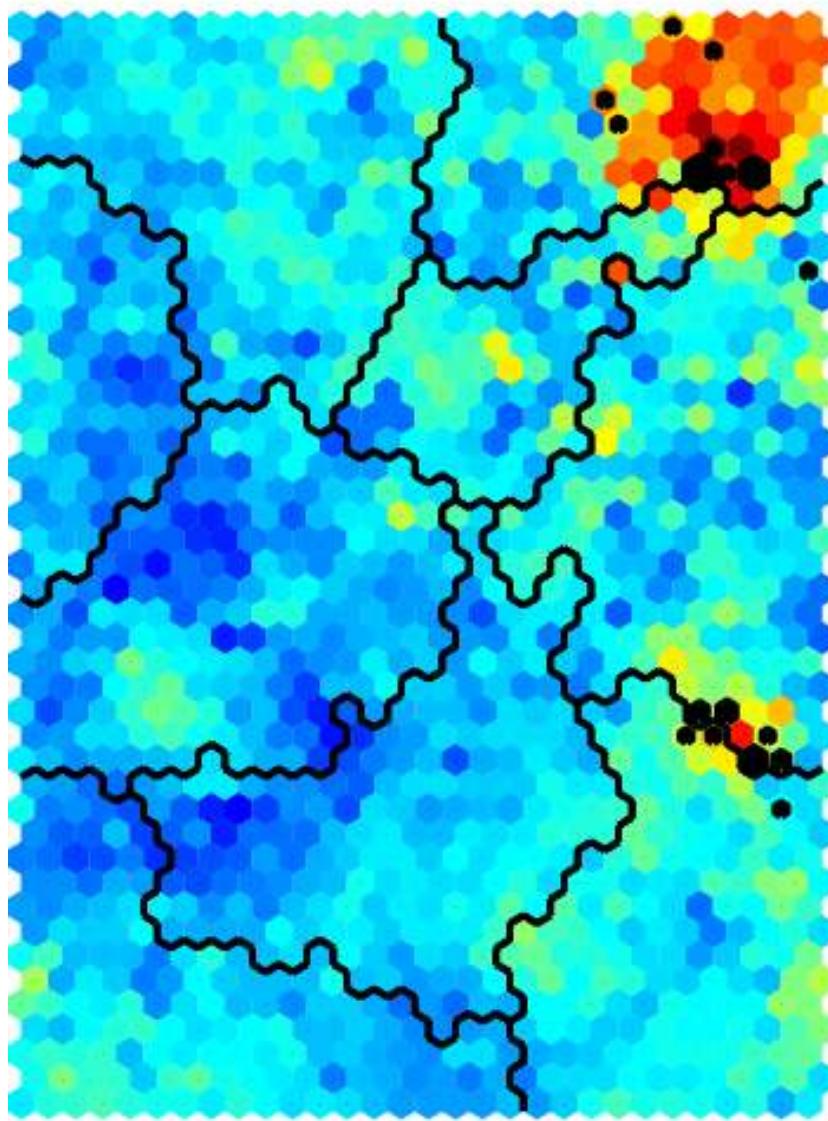
PPARgamma agonists ameliorate diacylglycerol-protein kinase Verrier et al. Circ. Res, 2004



Pathway Fitness (coherence)

amelioration via inhibition of diacylglycerol kinase.

Rapamycin Family

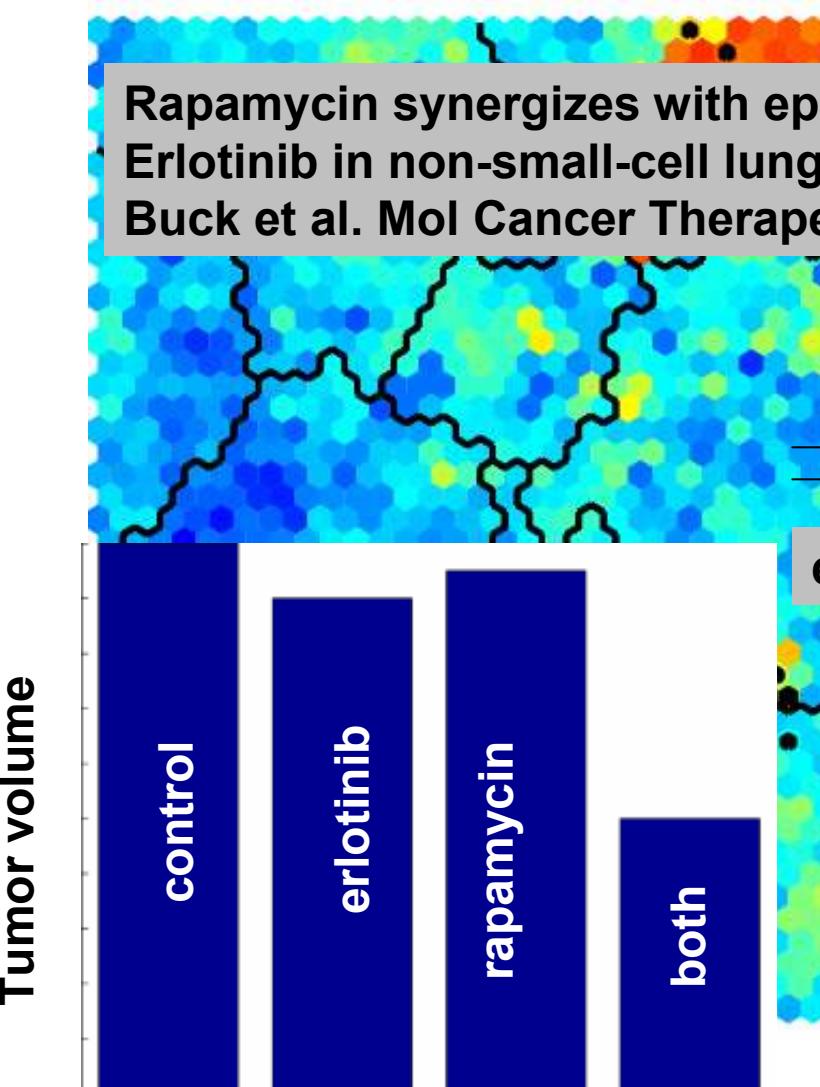


h_vegf
h_ires
h_ran
GO:74 cell cycle
GO:1525 angiogenesis
GO:6099 tca cycle

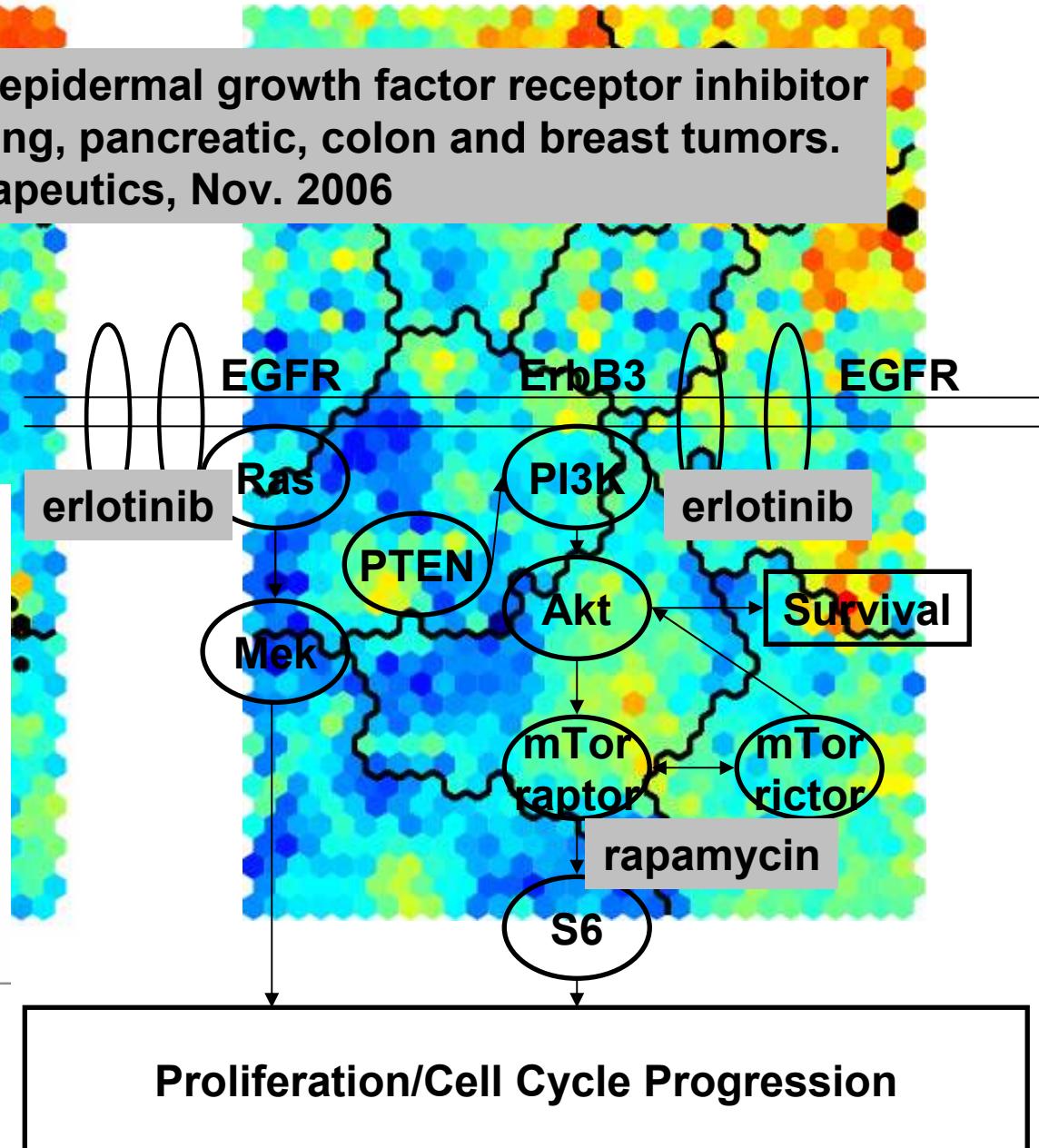
h_ctc
GO:3724 RNA helicase
GO:6631 fatty acid metabolism

- Structurally similar to temsirolimus (N=24)
179 gene expressions are correlated with these 24 NSCs

Rapamycin

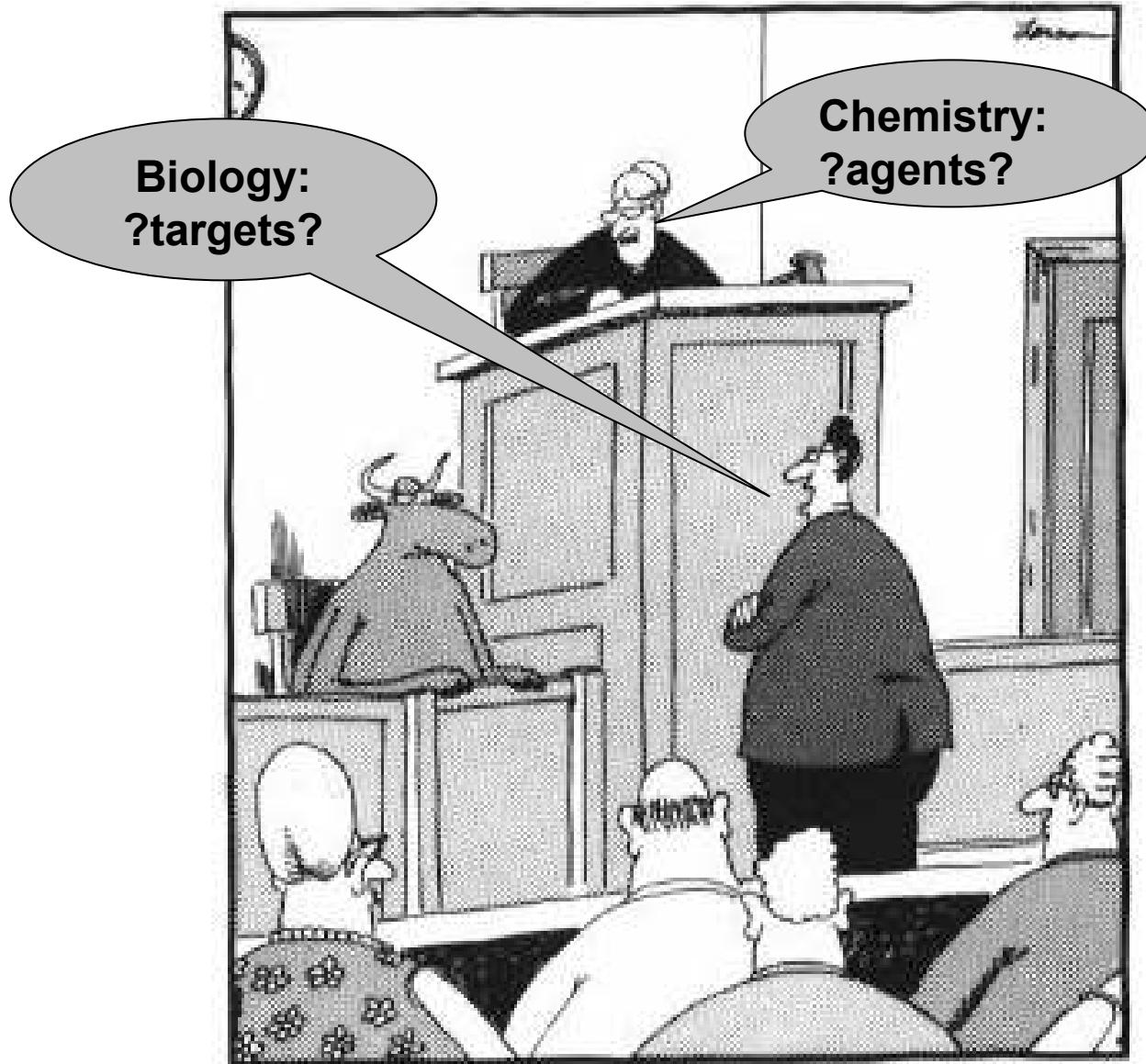


Erlotinib

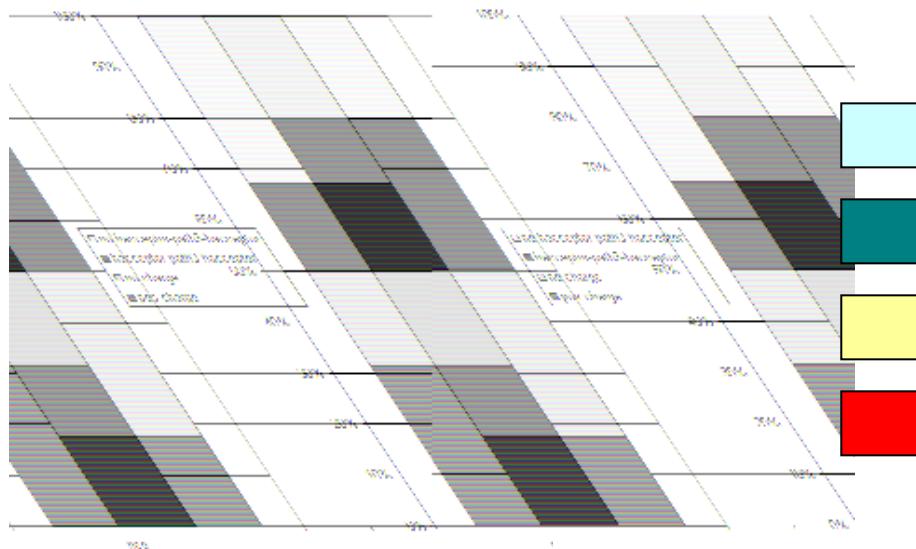
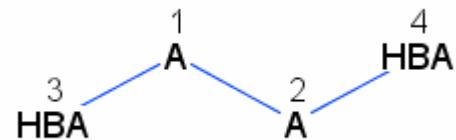


Chemistry Meets Biology

	Act 1	Act 2	Act 3	Interm.	Act 4
GI50	✓	✓	✓	✓	✓
Chemistry		✓	✓		✓
mRNA			✓	✓	
Pathways				✓	✓
Xenograft					✓

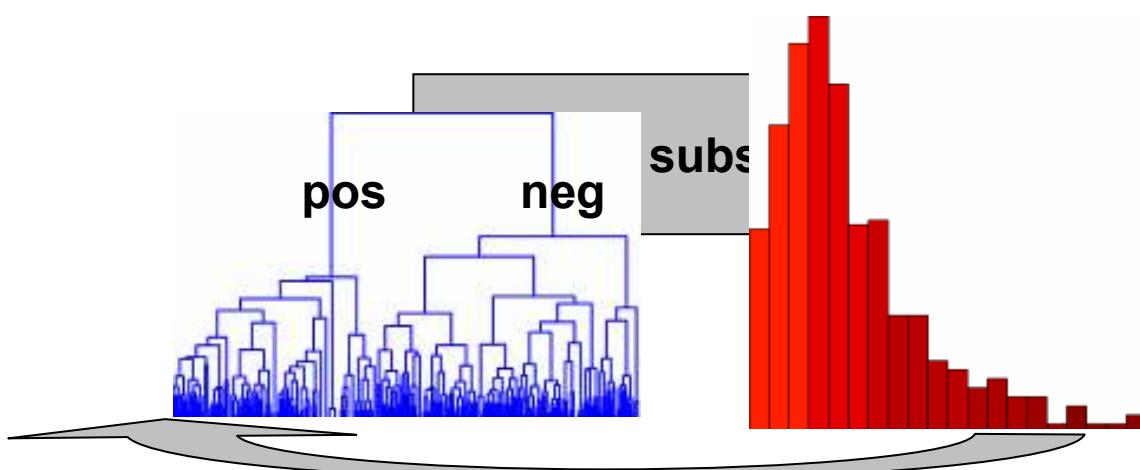


Look. We know that it works ---- that is no longer the question. What we now want to know is how... How now brown cow?"

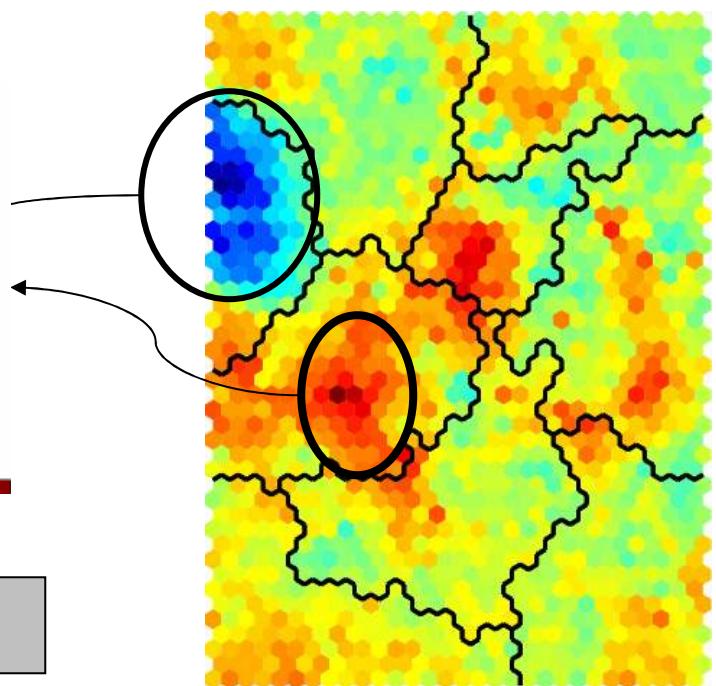


H-acceptor path3

+ charge

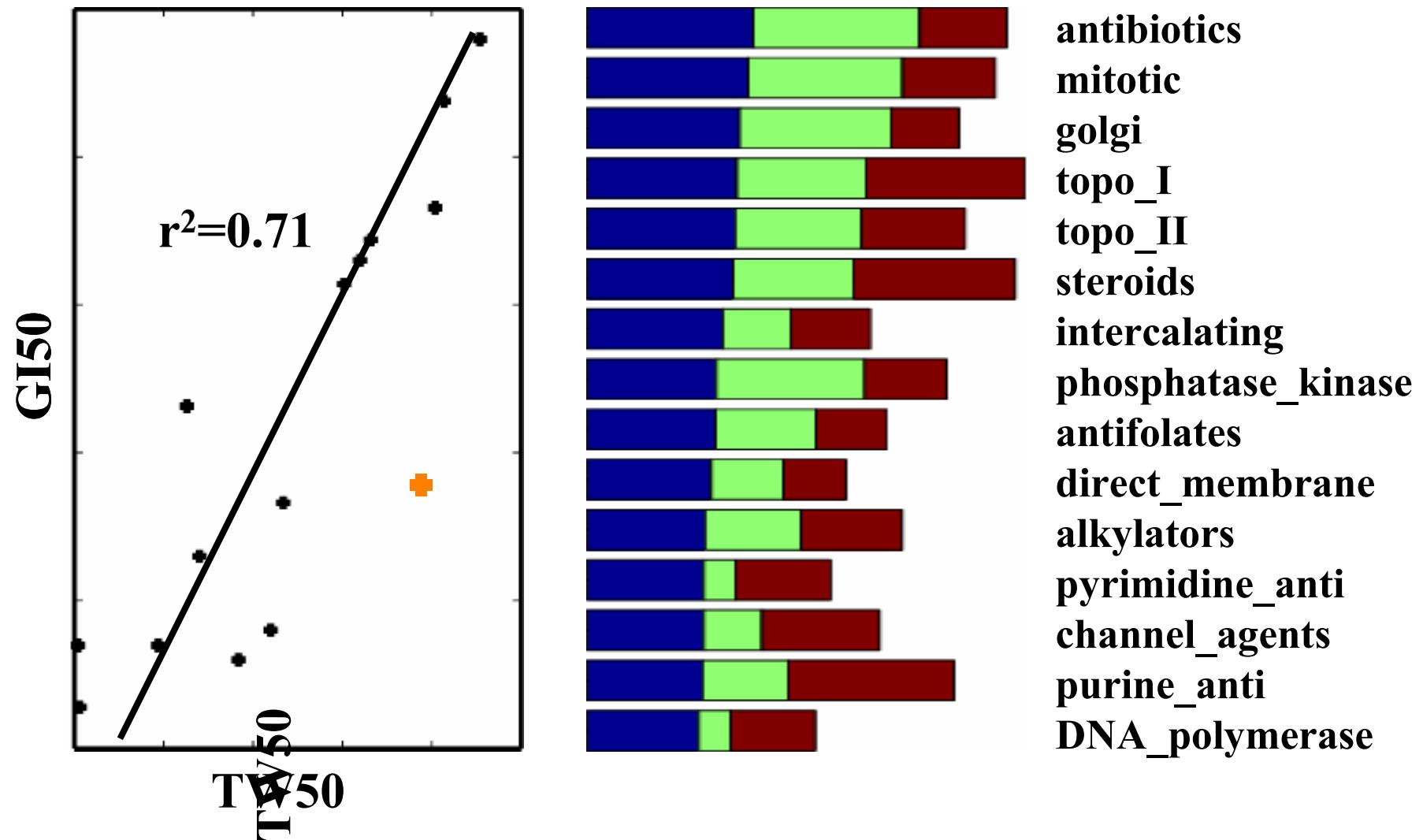


Thiosemicarbazone NSC73306



Xenograft Data

II



Phosphatase_kinase agents produce near maximal tumor weight reduction for modest values in GI50 and Therapeutic Index