

The SMEs' role in the area of systems biology approaches for medical applications

Alexander, Kel
Chief Scientific Officer
geneXplain GmbH

We can explain your genes and
put bricks together...

• Gene regulation • Biomarkers • Drug activity • Network modeling •

geneXplain

We aim to provide a comprehensive platform
of bioinformatics, cheminformatics
and systems biological tools
for personalized medicine and pharmacogenomics



The team participated in projects funded by EU Health – FP6, FP7

- **EuroDia (FP6: 01.03.2006-28.02.2010) - Diabetes**
 - ▶ Functional genomics of pancreatic beta cells and of tissues involved in control of the endocrine pancreas for prevention and treatment of type 2 diabetes. (BIOBASE)
- **VALAPODYN (FP6: 01.10.2006-30.03.2010) – Epilepsy**
 - ▶ Validated Predictive Dynamic Model of Complex Intracellular Pathways Related to Cell Death and Survival. (BIOBASE)
- **Net2Drug (FP6: 01.02.2007-31.07.2010) - Cancer**
 - ▶ From gene regulatory networks to drug prediction. (BIOBASE, ISB)
- **SysCo (FP6: 01.09.2007-27.02.2011) - Infection**
 - ▶ Systematic Functional analysis of Intracellular Parasitism as a model of genomes conflict. (BIOBASE, ISB)

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The team*) participated in projects funded by EU Health – FP6, FP7

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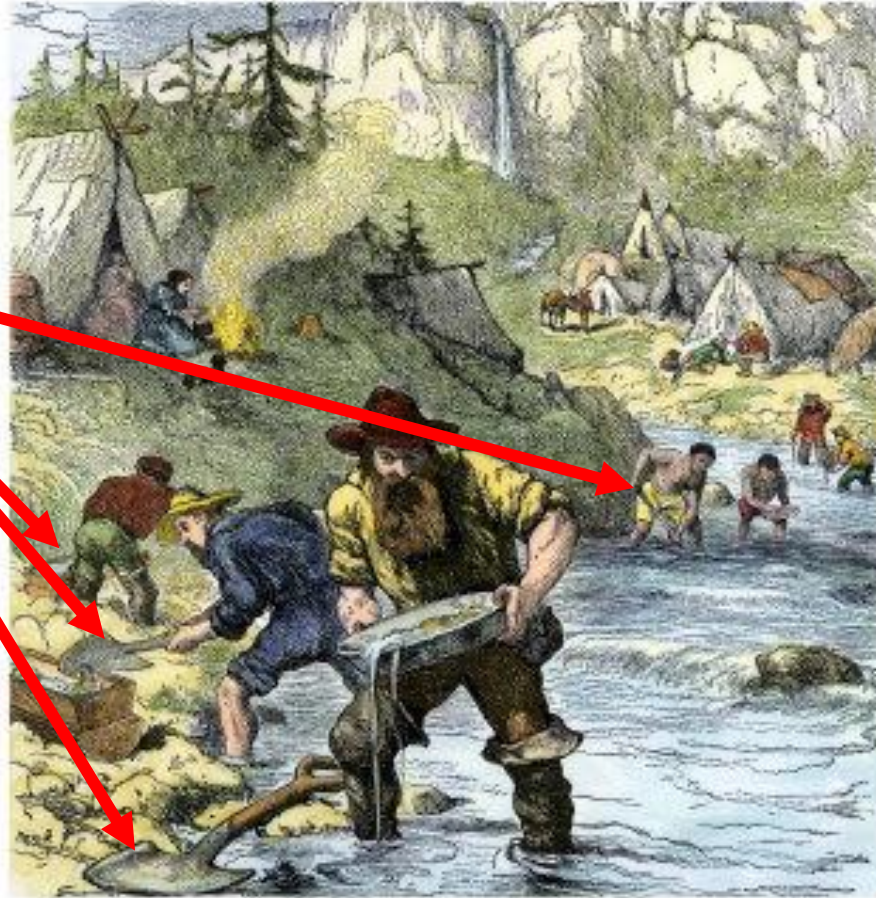
- **Gen2Phen (FP7: 01.01.2008-31.12.2012) - SNPs**
 - ▶ Genotype-To-Phenotype Databases: A Holistic Solution (BIOBASE)
- **LipidomicNet (FP7: 01.05.2008-30.04.2012) – Obesity**
 - ▶ Lipid droplets as dynamic organelles of fat deposition and release. Translational research towards human disease. (BIOBASE, ISB)
- **SysCol (FP7: 01.01.2011-31.12.2015) - Cancer**
 - ▶ Systems Biology of Colorectal Cancer (geneXplain)

*) The R&D team working in different time periods in SMEs: BIOBASE GmbH, ISB (Institute of Systems Biology, Russia) and geneXplain GmbH

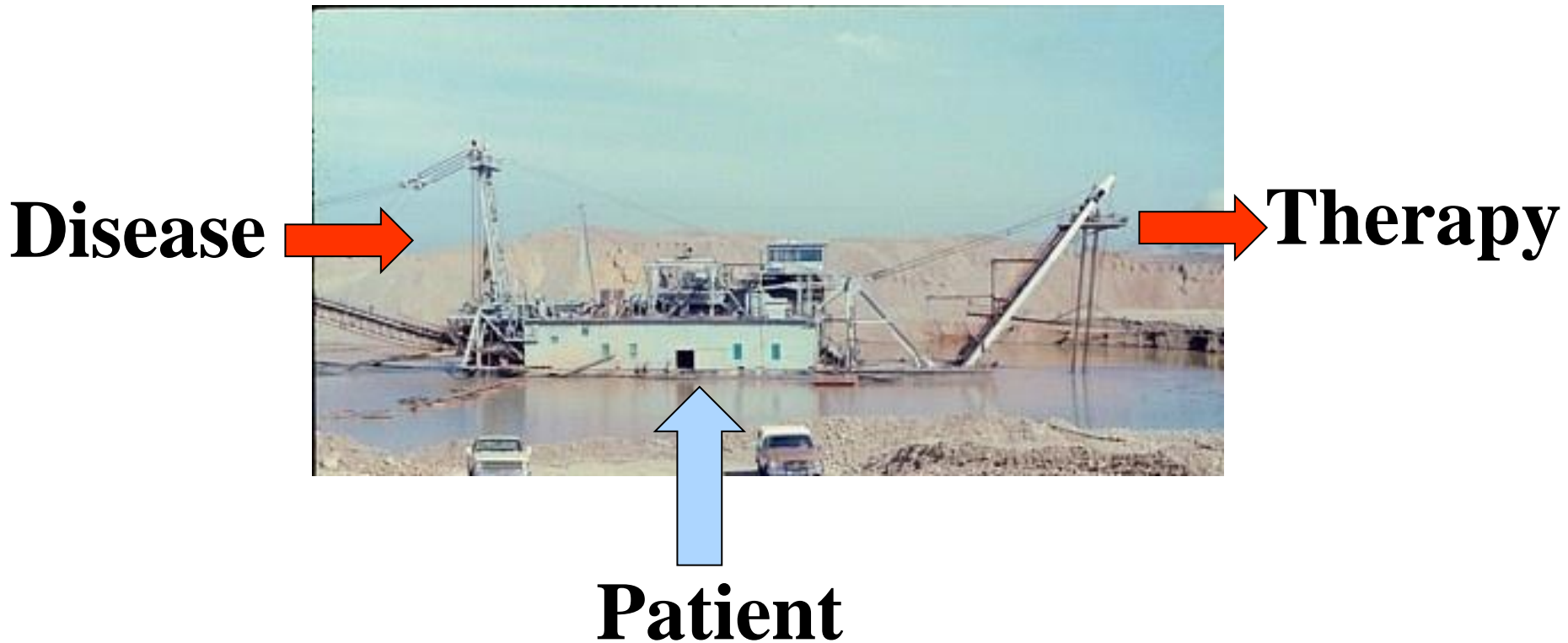


Drug discovery – the Gold Rush

Tools

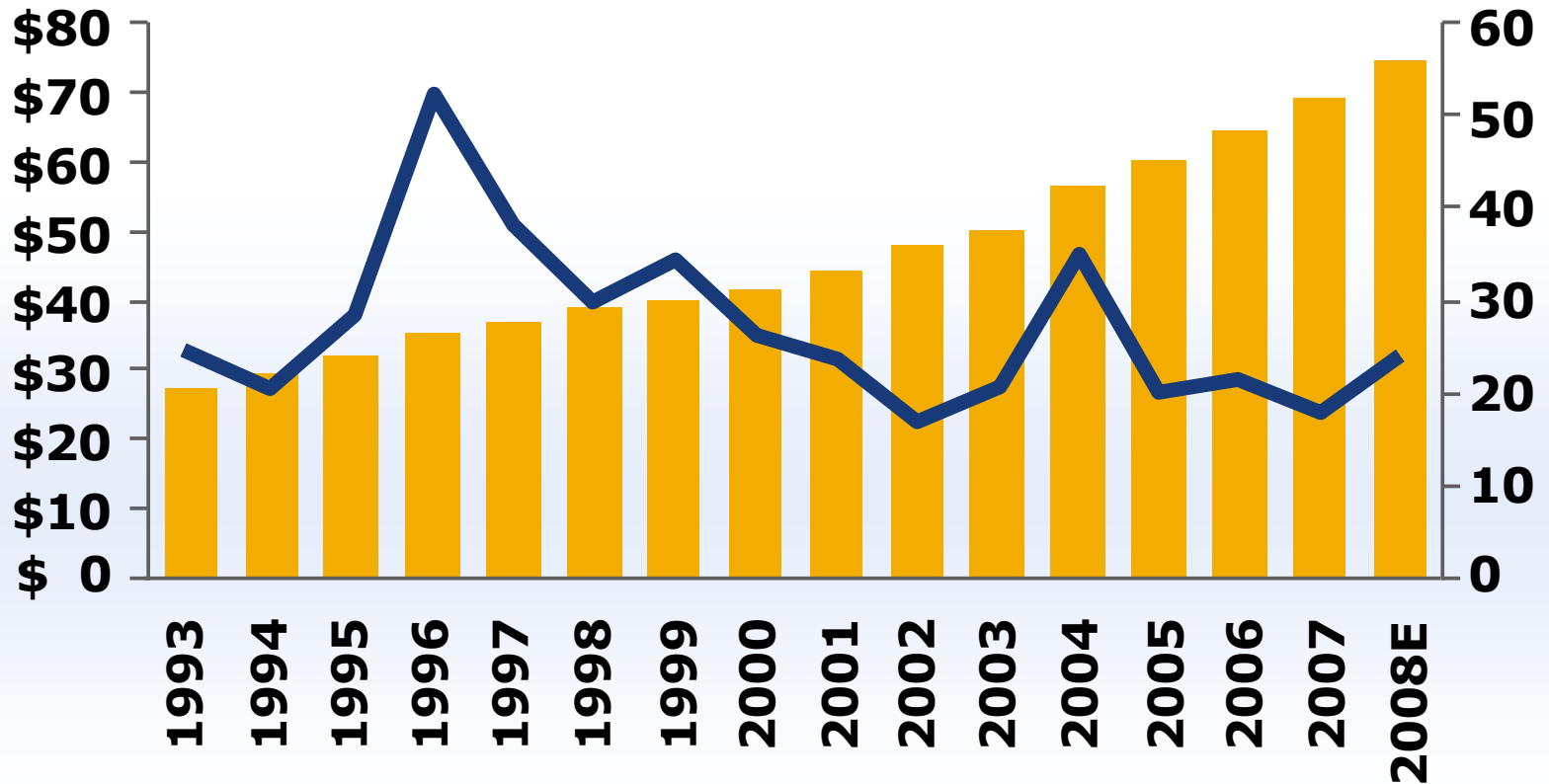


Drug discovery – should become a technology

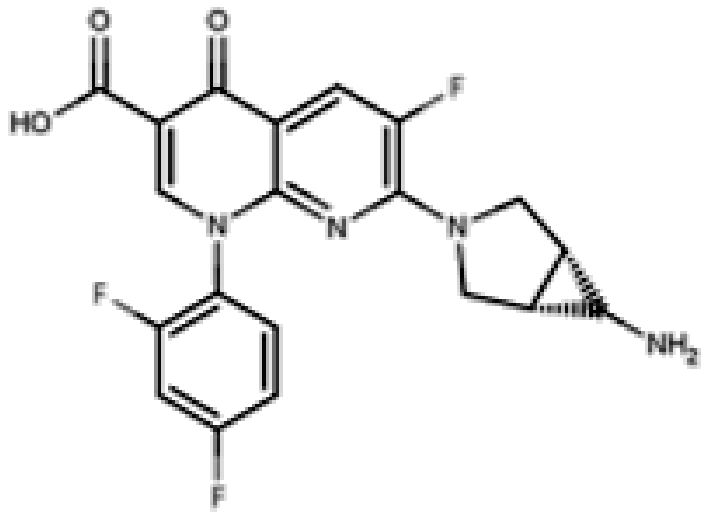


R&D Pipeline

■ Global R&D Spending — Drug approvals: NMEs/BLAs



Trovafloxacin - antibiotic



Withdrawn from market due to risk of idiosyncratic hepatotoxicity in 2001.

Systems medicine

Systems approaches will transform the way drugs are developed ... that will target multiple components of networks and pathways perturbed in diseases.

They will enable medicine to become predictive, personalized, preventive and participatory

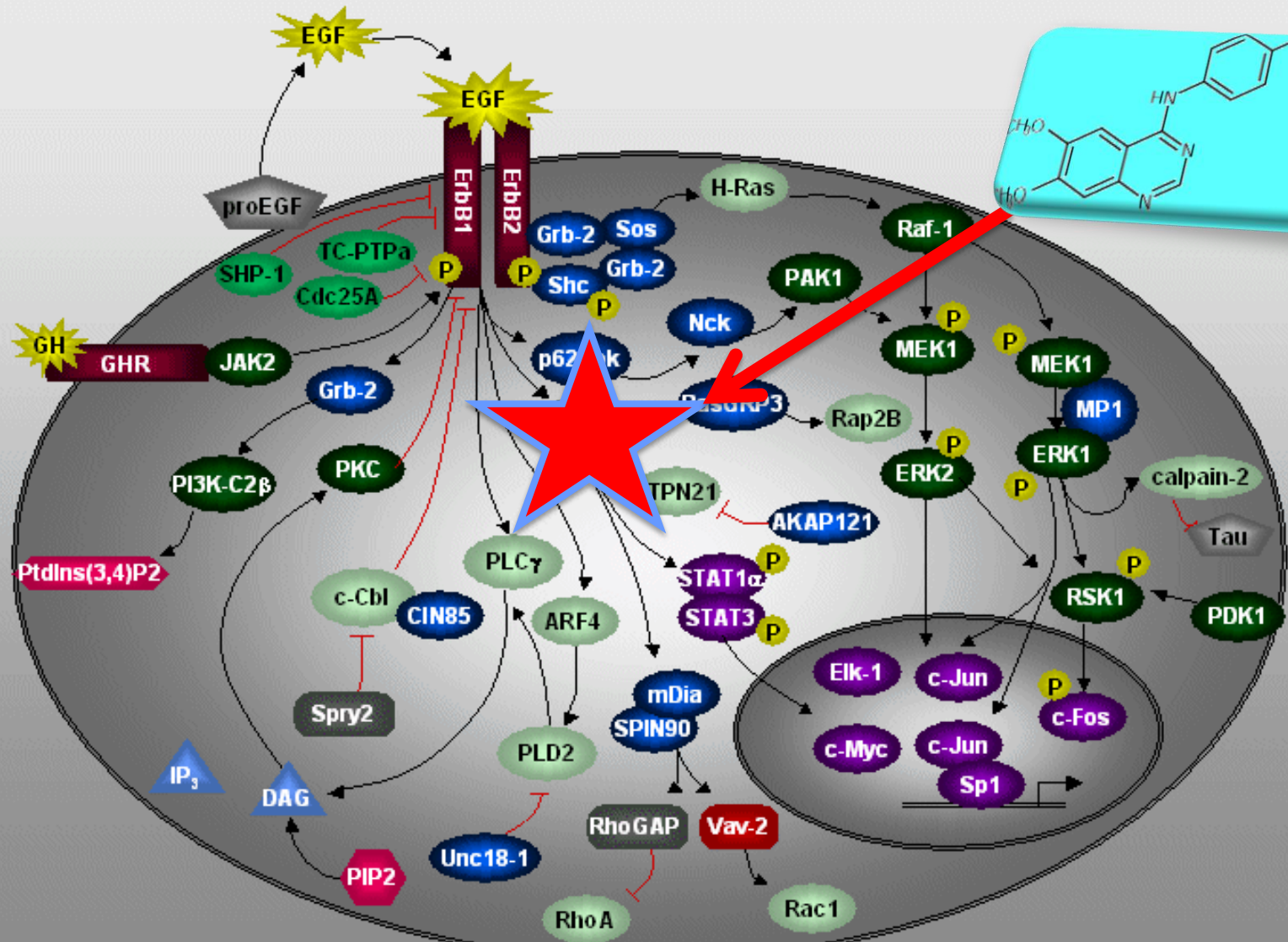
Systems medicine: the future of medical genomics and healthcare

Charles Auffray^{1*}, Zhu Chen² and Leroy Hood³

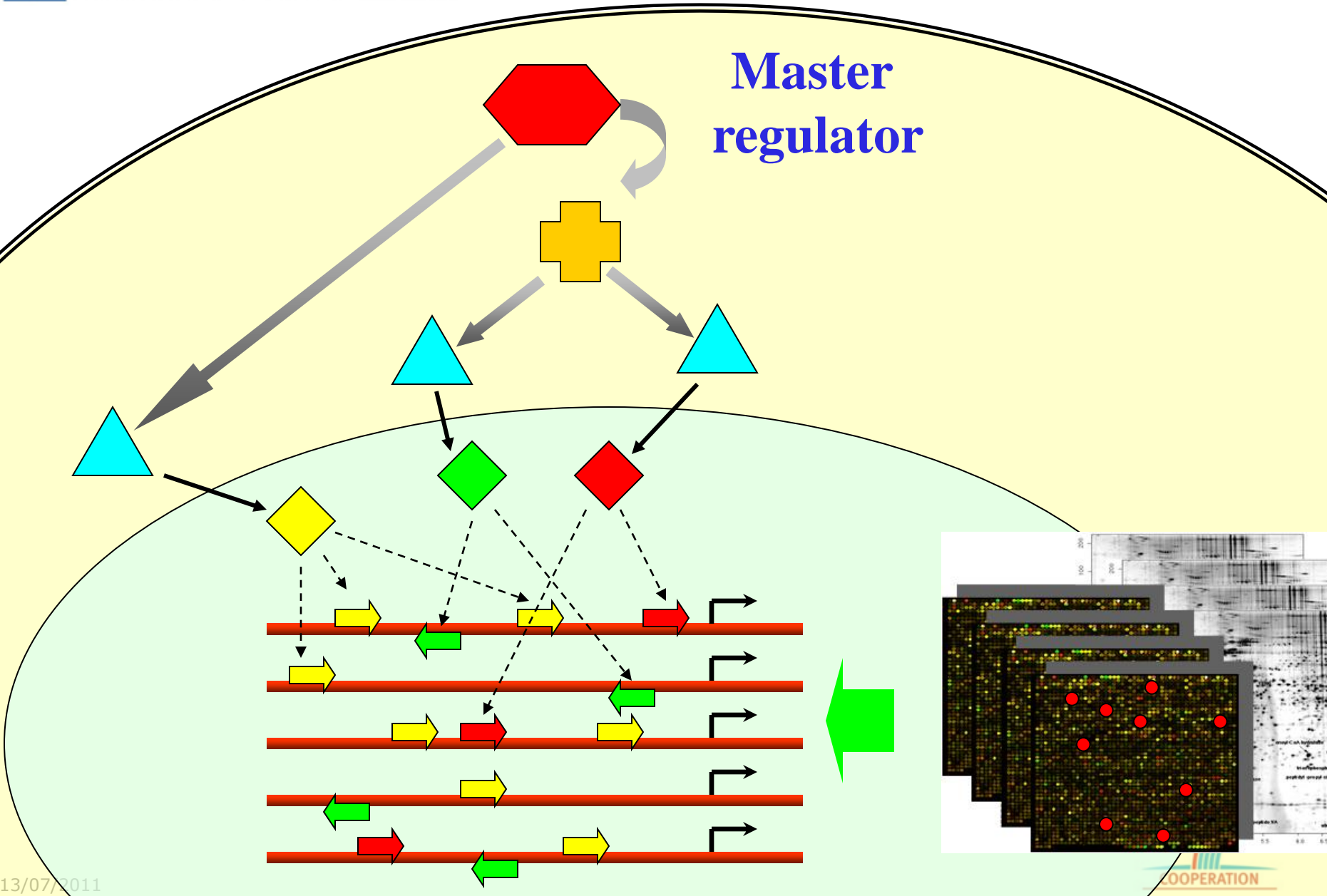
Genome Med 2009, 1:2



We should find a key pathway of a disease, select a good target and inhibit it.



Master regulator



GeneXplain platform – drug target discovery pipeline

Workflow Diagram:

```

    graph TD
      GS[Gene set] --> FT[Filter table]
      GS --> AS[Annotate table]
      AS --> SA[Set annotated]
      SA --> SSGS[Search sites on gene set]
      BS[Background set] --> SSGS
      SSGS --> S[Sites]
      S --> ST[summary]
      S --> CT2[Convert table(2)]
      ST --> MM[Matrices to molecules]
      CT2 --> GS2T[Gene set Transpath]
      MM --> M[Molecules]
      M --> RS[Regulator search]
      GS2T --> RS
      RS --> SH[Save hits]
      SH --> MU8[Molecules Upstream 8]
      SH --> MU8H2[Molecules Upstream 8 hits 2]
      MU8 --> SN[Save network]
      MU8H2 --> SN
      RankC[Rank column] --> SN
      SN --> MU8N2[Molecules Upstream 8 network 2]
      MU8N2 --> CT[Convert table]
  
```

History Table:

Type	Source	Time
analysis	Regulator search	2011.06.06 17:04:29
analysis	Convert table	2011.06.06 17:04:18
analysis	Regulator search	2011.06.06 17:03:28
workflow	Regulator analysis	2011.05.14 18:23:27

Description:
 In this analysis, gene expression was compared between lesion skin and uninvolved skin of the same 28 patients. The following were the steps of analysis shown here.

1. CEL file normalization. This step resulted in two files, Experiment normalized (MAS5) and Control normalized (MAS5).

GeneXplain platform – discovery pipeline

The screenshot displays the GeneXplain platform interface, which is divided into several functional areas:

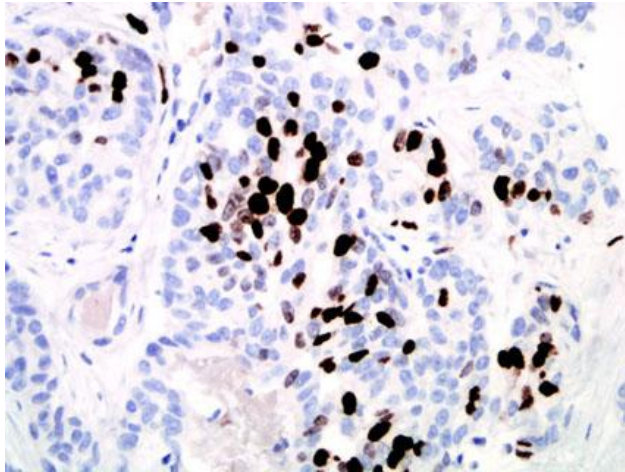
- Analyses Panel:** Shows a list of data files, including GSM409593.tbt.td through GSM409609.tbt, and folders for 'nutlin3' and 'nutlin3.zip'.
- Table View:** A table with columns 'ID', 'name', 'title', and 'SUM-p.value'. It lists various gene identifiers and their associated p-values, such as A_32_P190609 (ENSG00000111052, LIN7A, -93.26118912).
- Workflow Diagram:** A central flowchart showing the process: 'Gene set' leads to 'Sites on gene set', which then goes to 'Convert table(2)'. From there, it branches into 'summary' (leading to 'Matrices to molecules') and 'Gene set Transpath' (leading to 'Regulator search').
- Simulation Result:** A graph showing 'Quantity or concentration' over 'Time' (0 to 350). The graph is titled 'Complete 100%' and shows several curves for different components like E2F1, pRB, and pRBp.
- Network Diagram:** A complex network of nodes and edges, with nodes labeled E2F1, pRB, and pRBp. Some nodes are highlighted in yellow, and others in red.
- Regulator Search Results:** A hierarchical tree diagram starting with 'TMR6' at the top, branching down to various regulators like 'Sic1/Ink4a', 'Skr1/Ink4a', 'Skr2/Ink4a', 'p27/Ink4a', 'Skr3/Ink4a', 'NF-κappaB', 'Raf-1', 'ERK1', and 'ERK2'.

Description:
In this analysis, gene expression was compared between lesion skin and uninvolved skin of the same 28 patients. The following were the steps of analysis shown here.

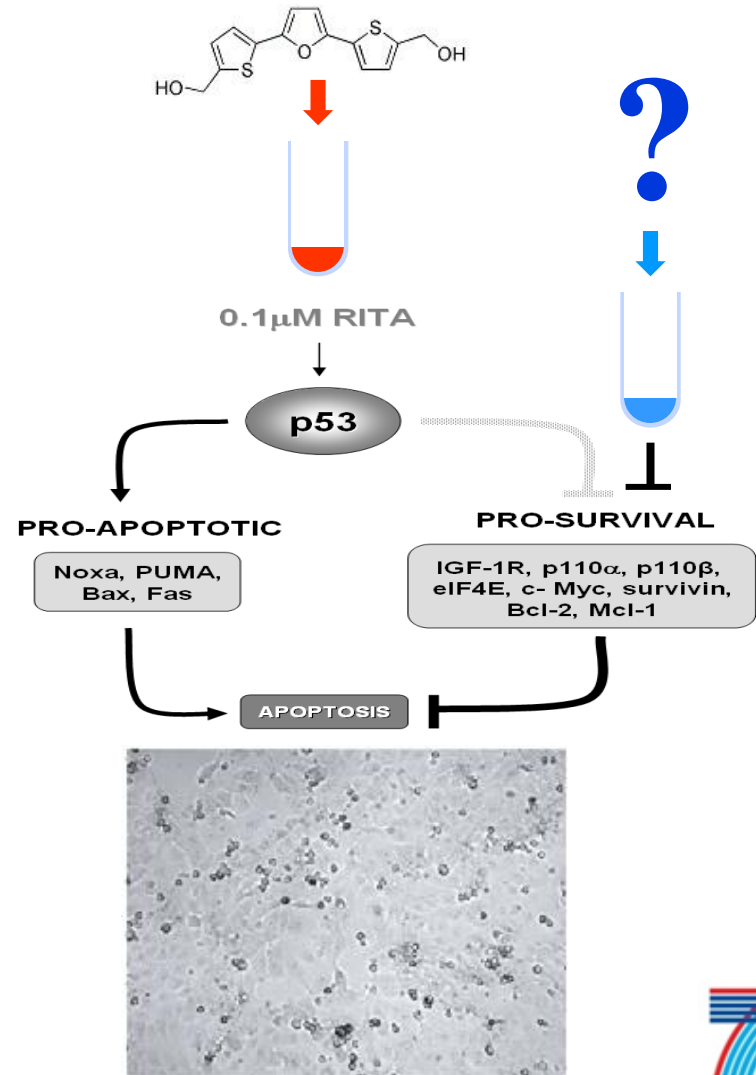
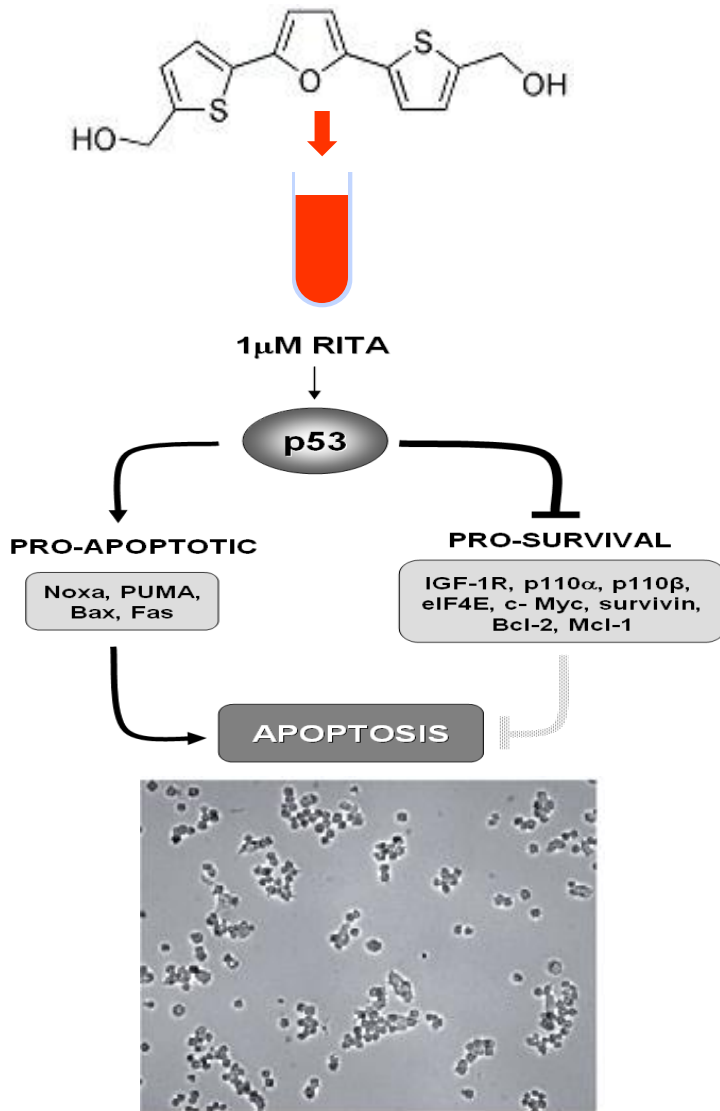
1. CEL file normalization. This step resulted in two files, Experiment normalized (MAS5) and Control normalized (MAS5).



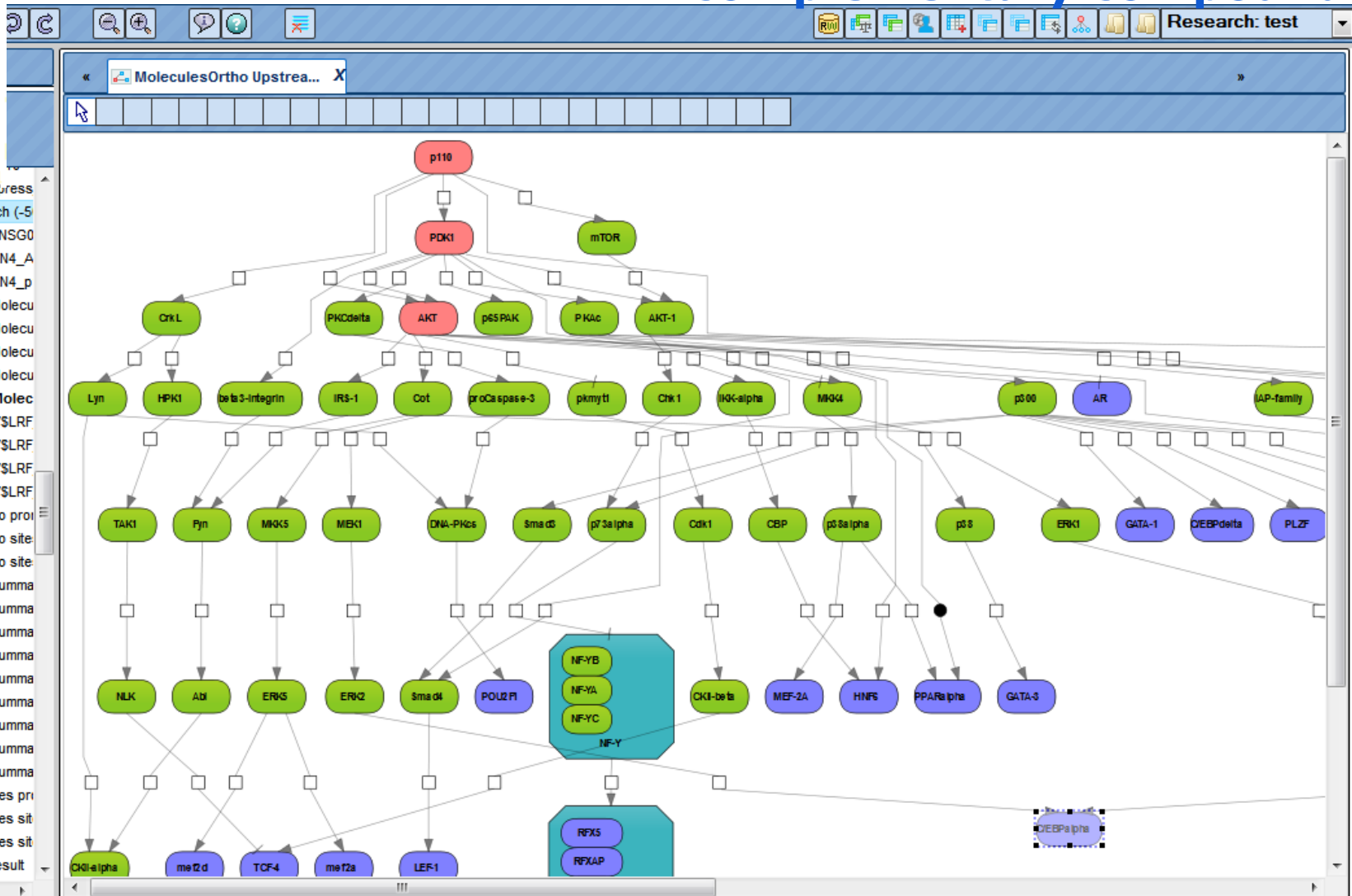
Cancer



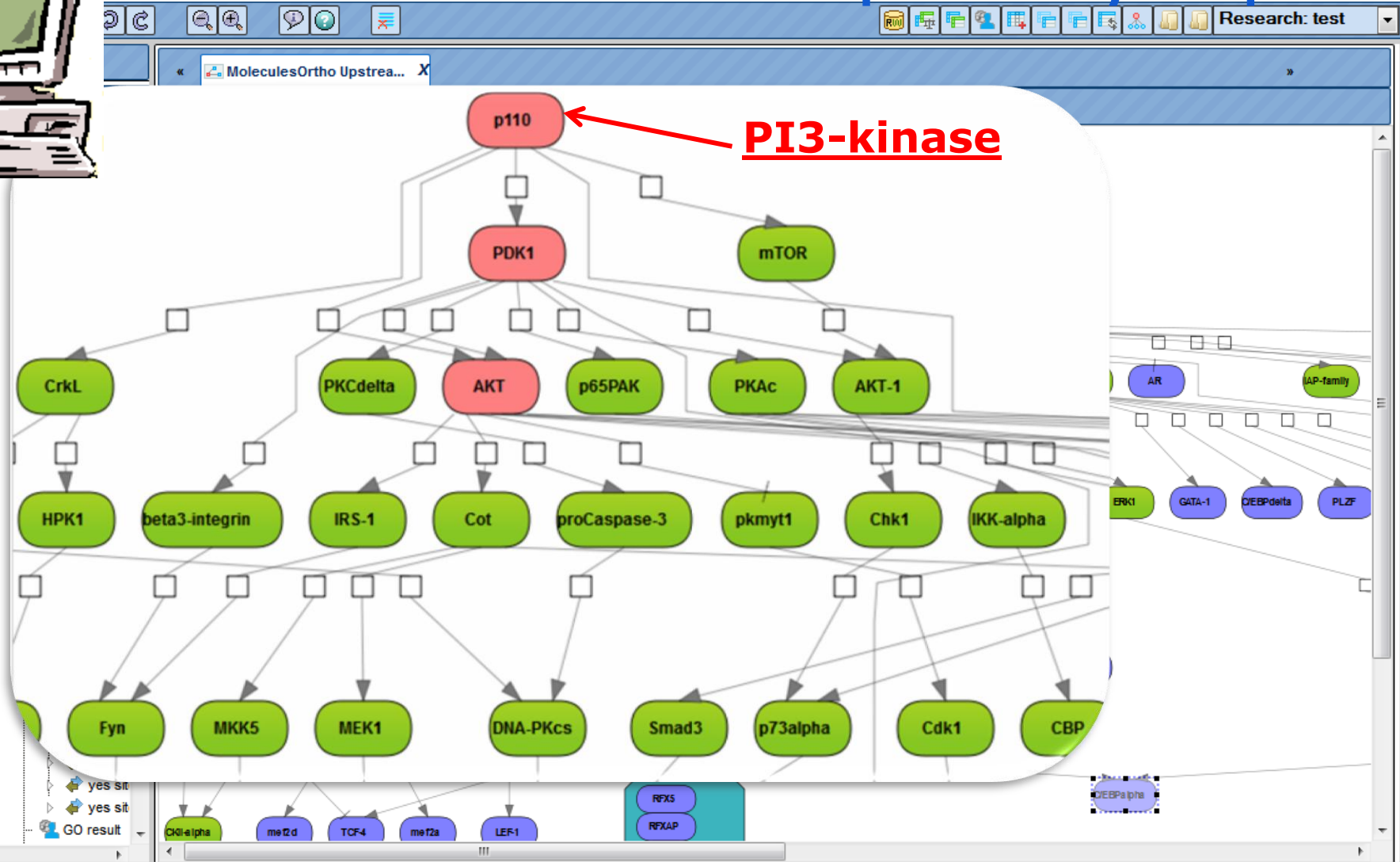
Apoptosis versus survival of cancer cells



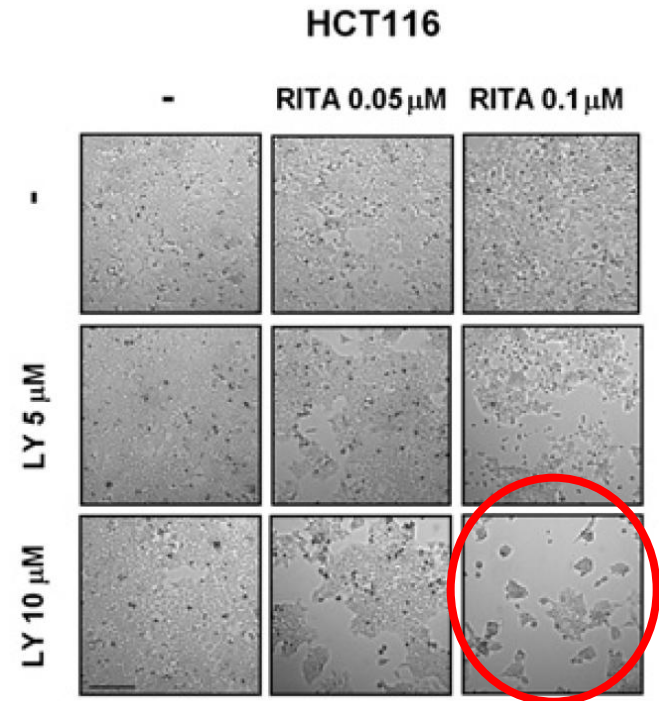
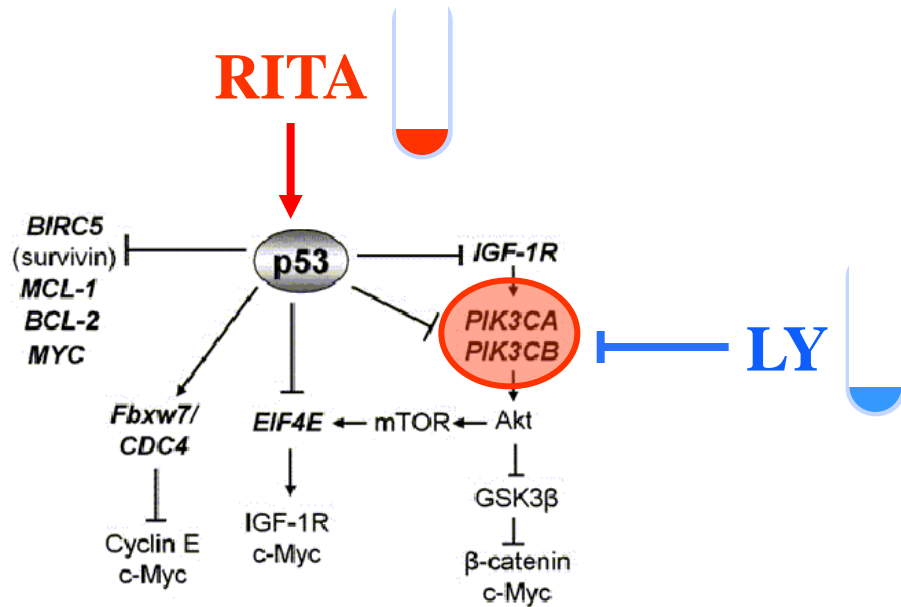
Survival mechanisms of cancer cells upon RITA treatment and potential target proteins for a complementary compound

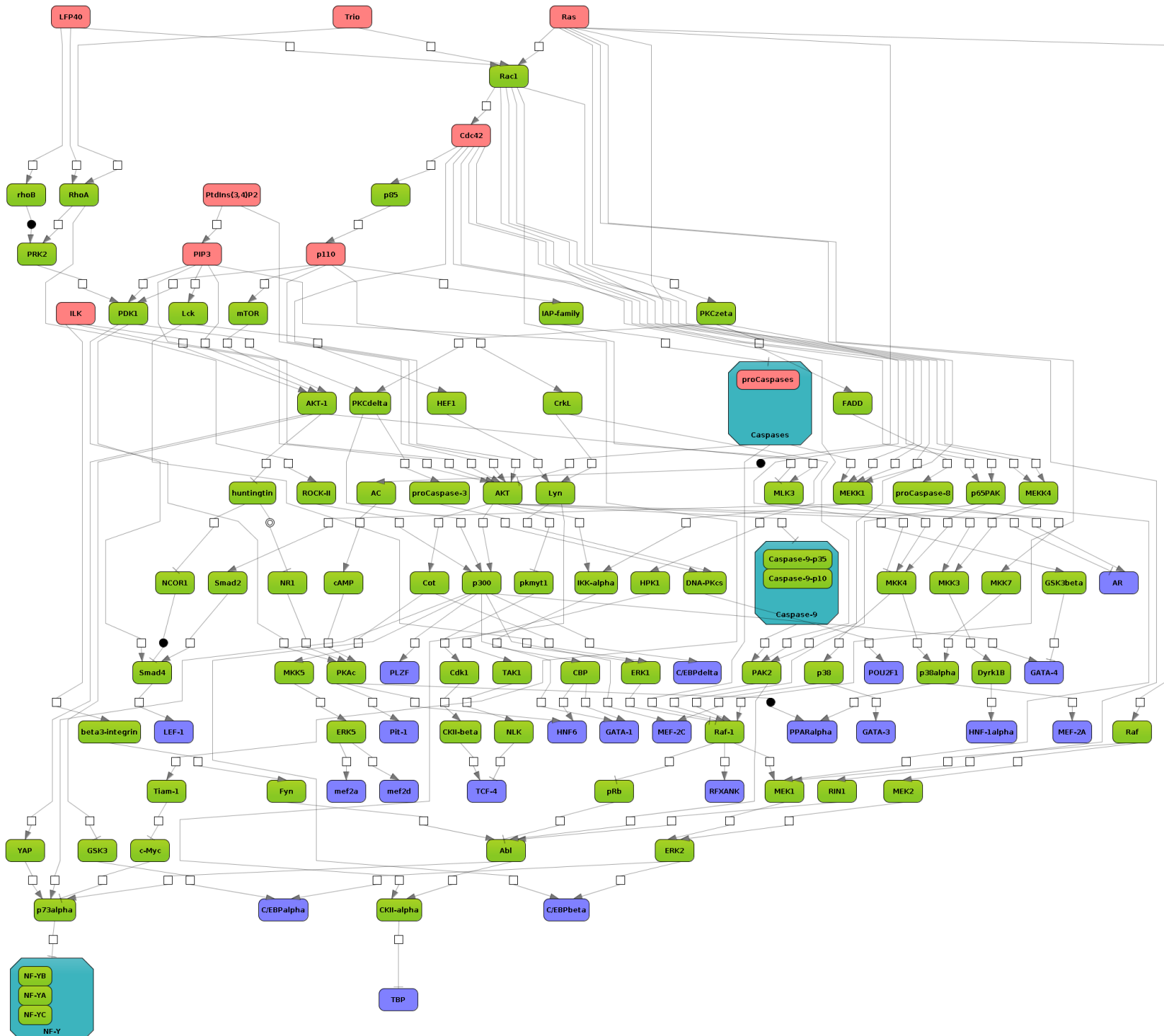


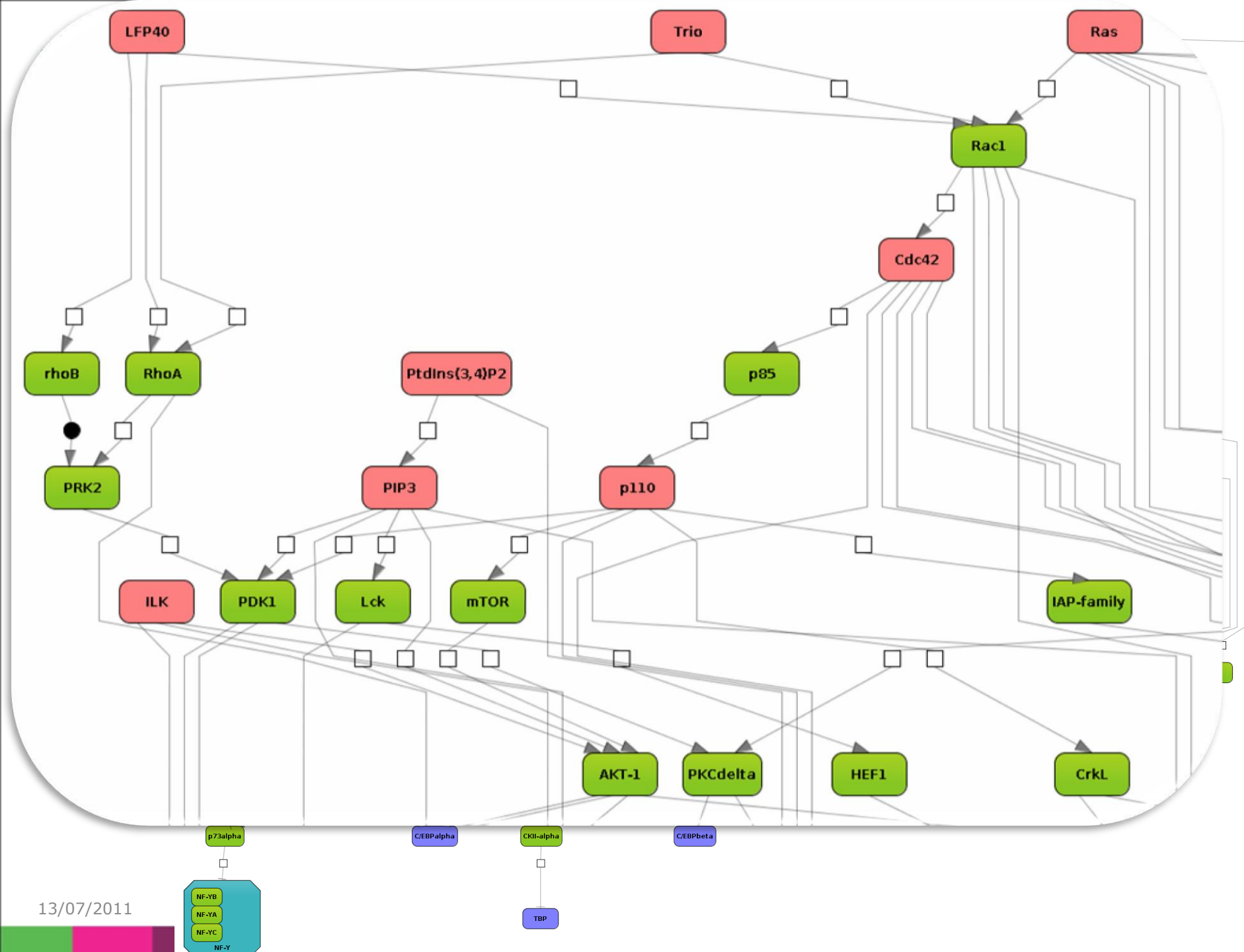
Survival mechanisms of cancer cells upon RITA treatment and potential target proteins for a complementary compound



Death of Cancer cells treated with 0.1 μM RITA and PI3-kinase inhibitor LY294002



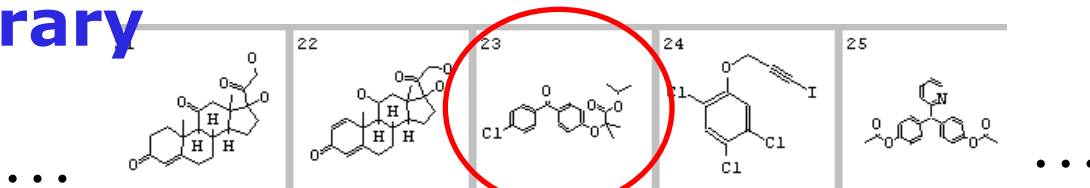




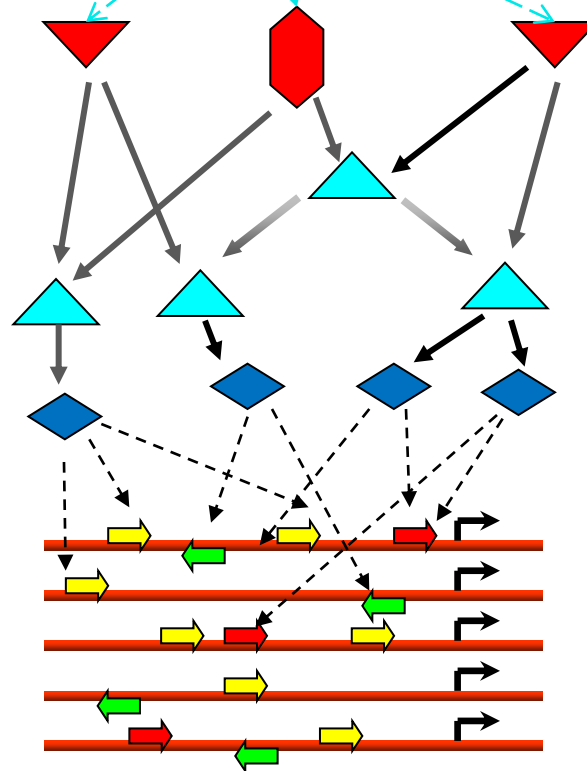
13/07/2011

Identified 64 novel compounds

ChemNavigator Library 24 million compounds



SAR/QSAR



PASS Prediction of Activity Spectra for Substances

Version 9.1 Professional

Copyright © 2009

V. Poroikov, D. Filimonov & Associates

<http://www.ibmc.msk.ru/PASS/>

48 Substructure
There are 6 kno
Drug likeness

Tested 16 compounds in a panel of several cancer cell lines.

Found active: Compound N15

Hypoxia inducible factor 1 alpha inhibitor	Phosphatidylinositol 3-kinase beta inhibitor
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Targets

Showed growth suppression in 3 different breast cancer cell lines. The effect appears to be p53-independent (kills p53-null colon cancer cells) and it does not affect the growth of non-transformed mammary epithelial cells

Found active: Compound N6

Cyclin-dependent kinase 2 inhibitor	Myc inhibitor
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Targets

Out of panel of 7 different cancer lines it killed only melanoma cells without any effects in other cell lines and on control non-transformed mammary epithelial cells.

Tested 16 compounds in a panel of several cancer cell lines.

Found active:

Hypoxia inducible factor 1 alpha

Showed growth suppression in several cancer cell lines (The effect appears to be more pronounced in breast cancer cells) and it did not affect normal mammary epithelial cells.

Found active:

Cyclin-dependent kinase 2 inhibitor

Out of panel of 7 different compounds tested, only this one showed growth suppression in transformed mammary epithelial cells.

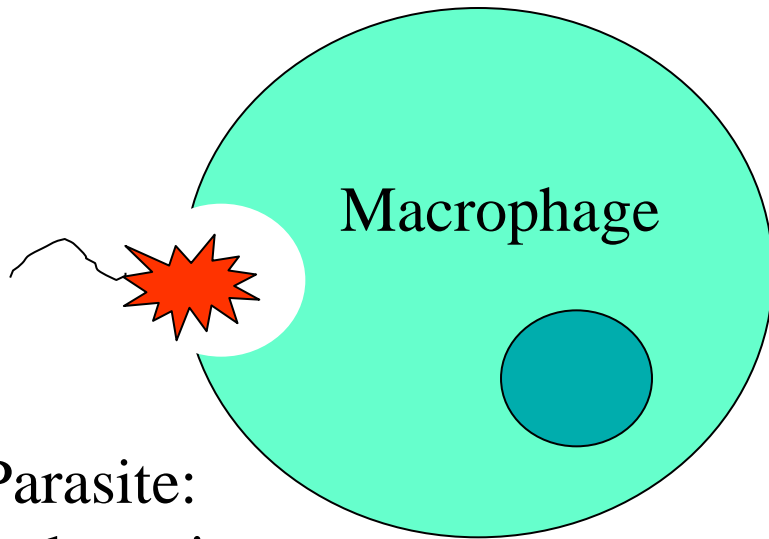


S

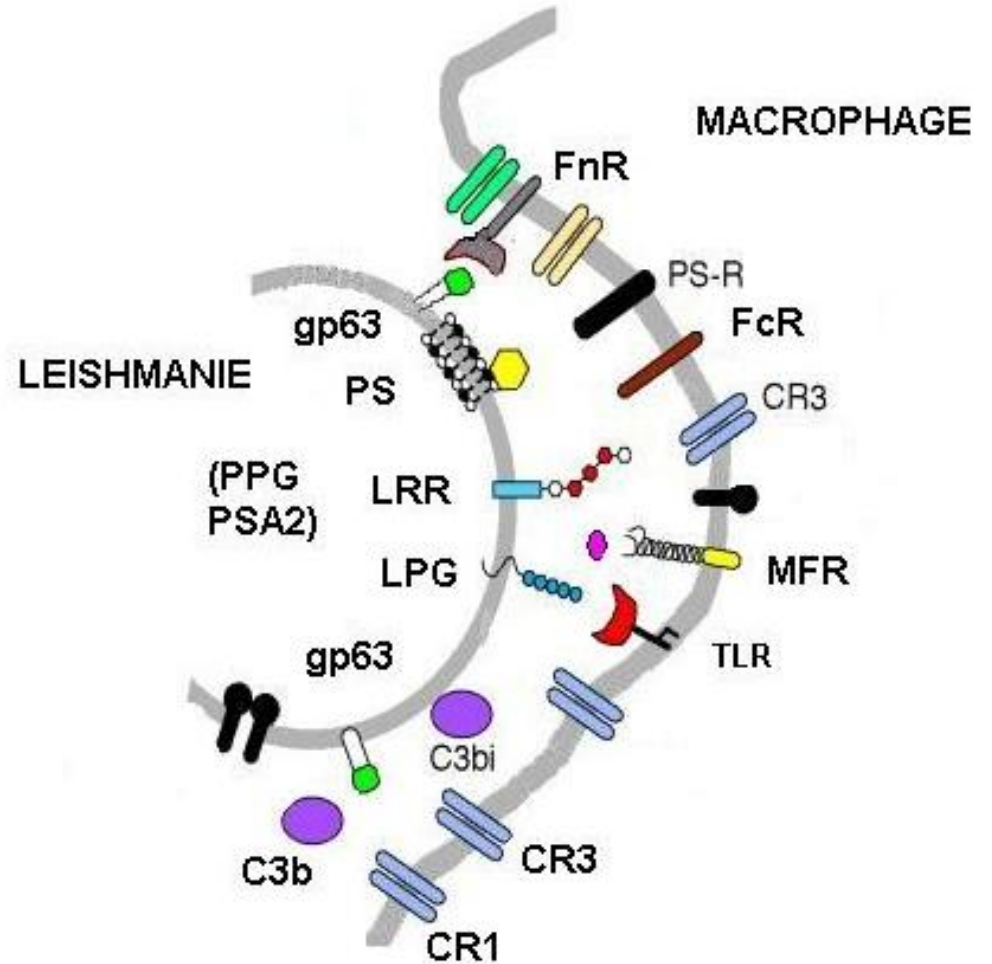
S



Infection project: SysCo



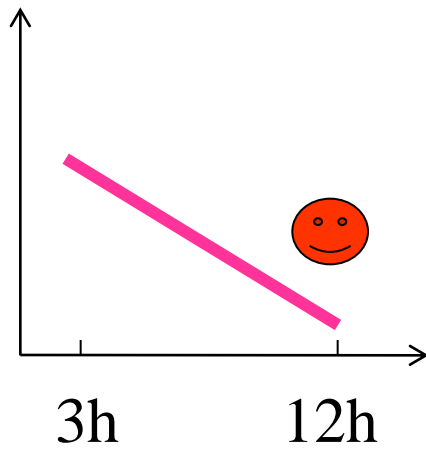
Parasite:
Leshmania



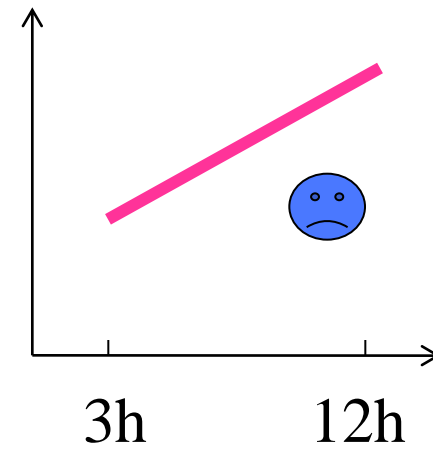
Parasite



Killed parasite



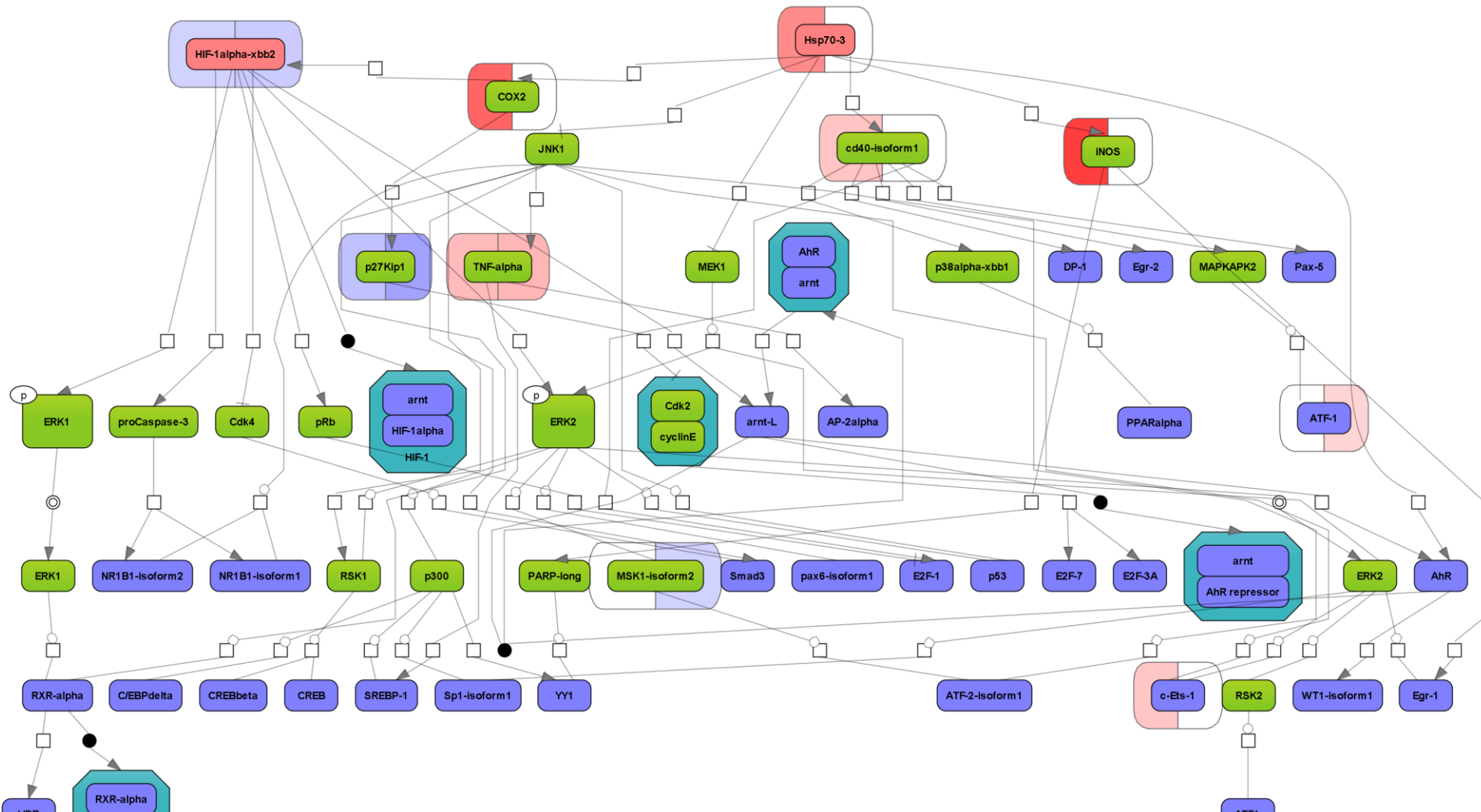
**Death of
macrophages
by
apoptosis**



Parasite

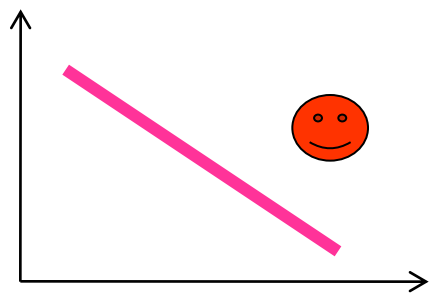
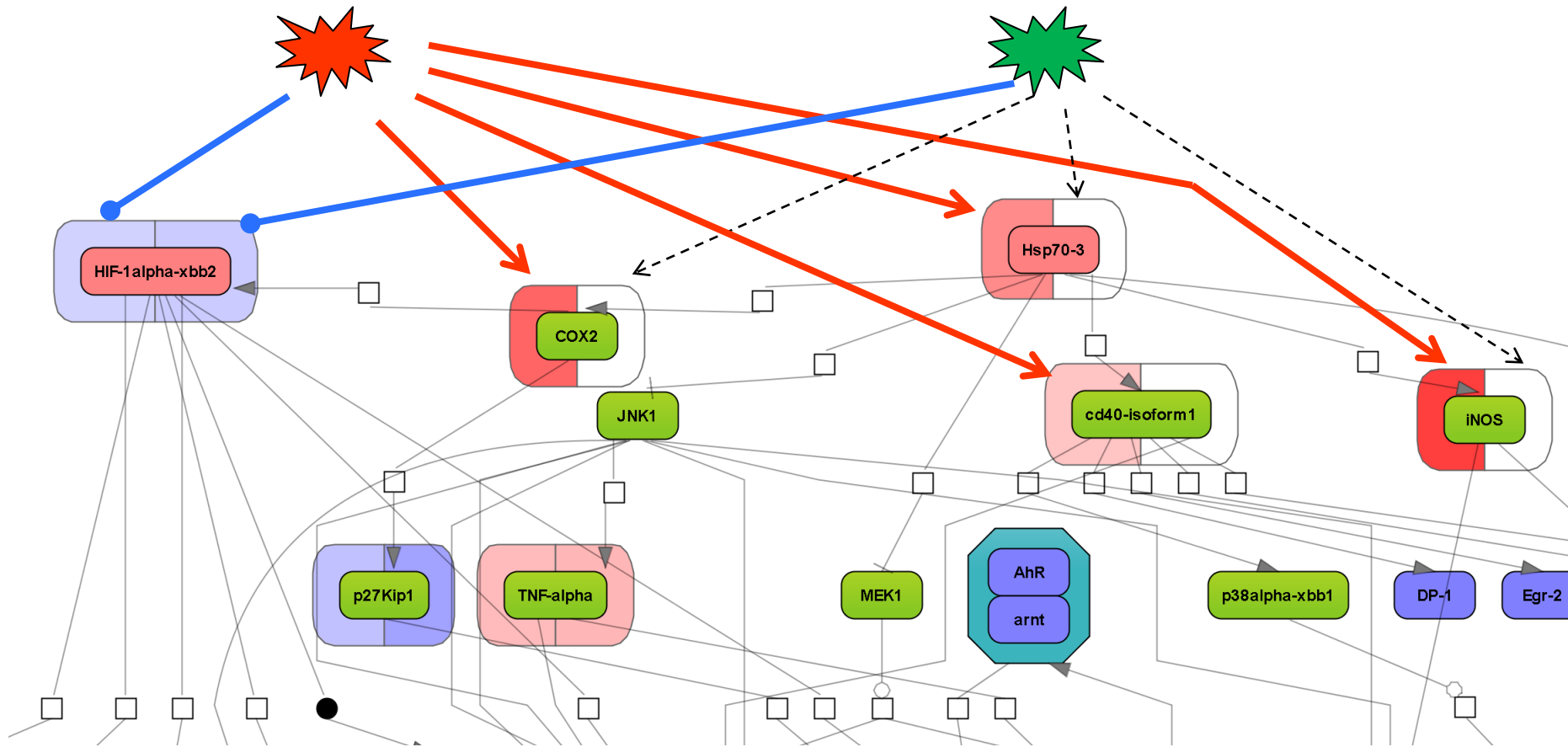


Killed parasite

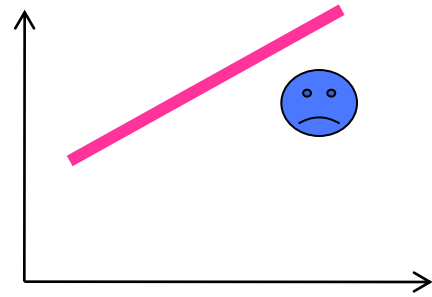


Parasite

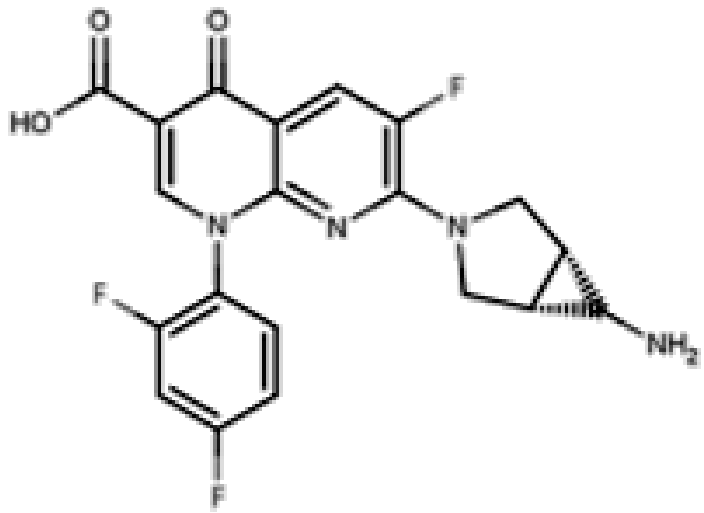
Killed parasite



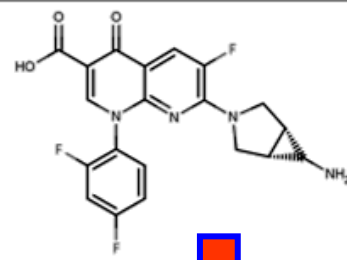
Death of macrophages



Trovafloxacin - antibiotic



Withdrawn from market due to risk of idiosyncratic hepatotoxicity in 2001.

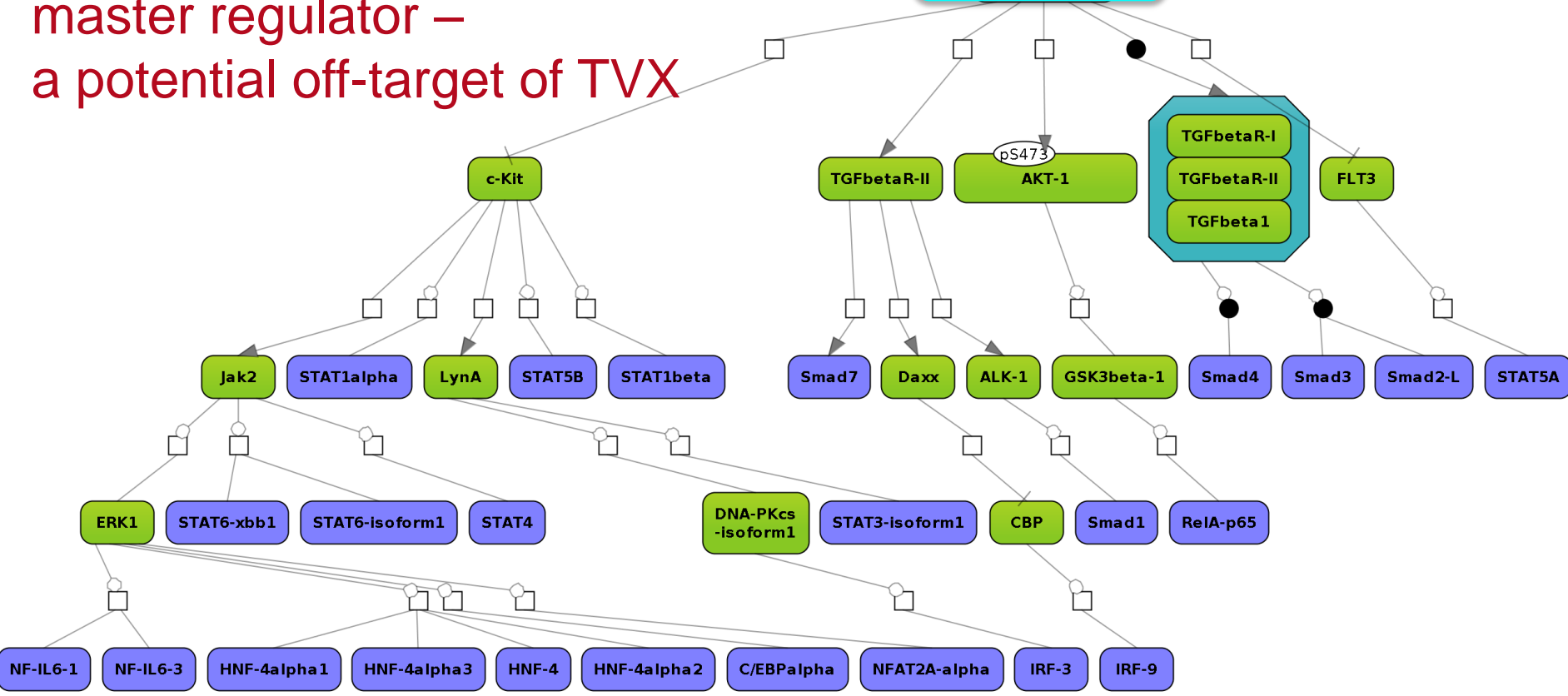


Trovafloxacin (TVX)

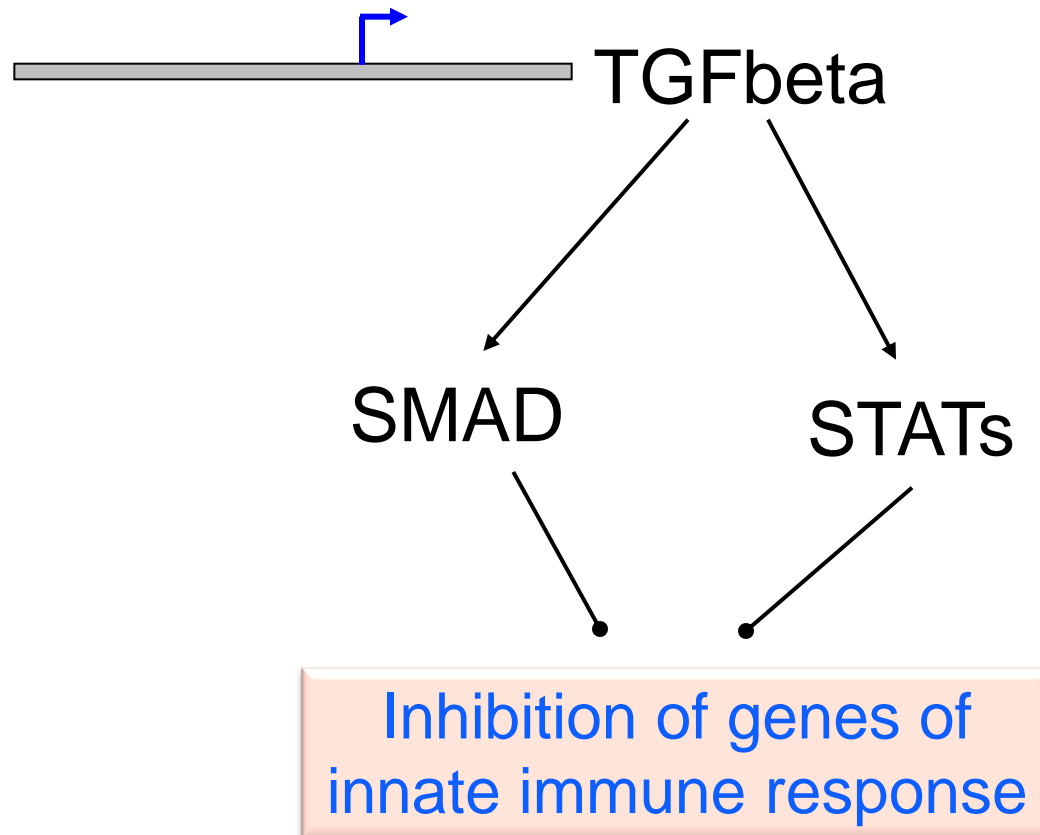


TGF-beta1

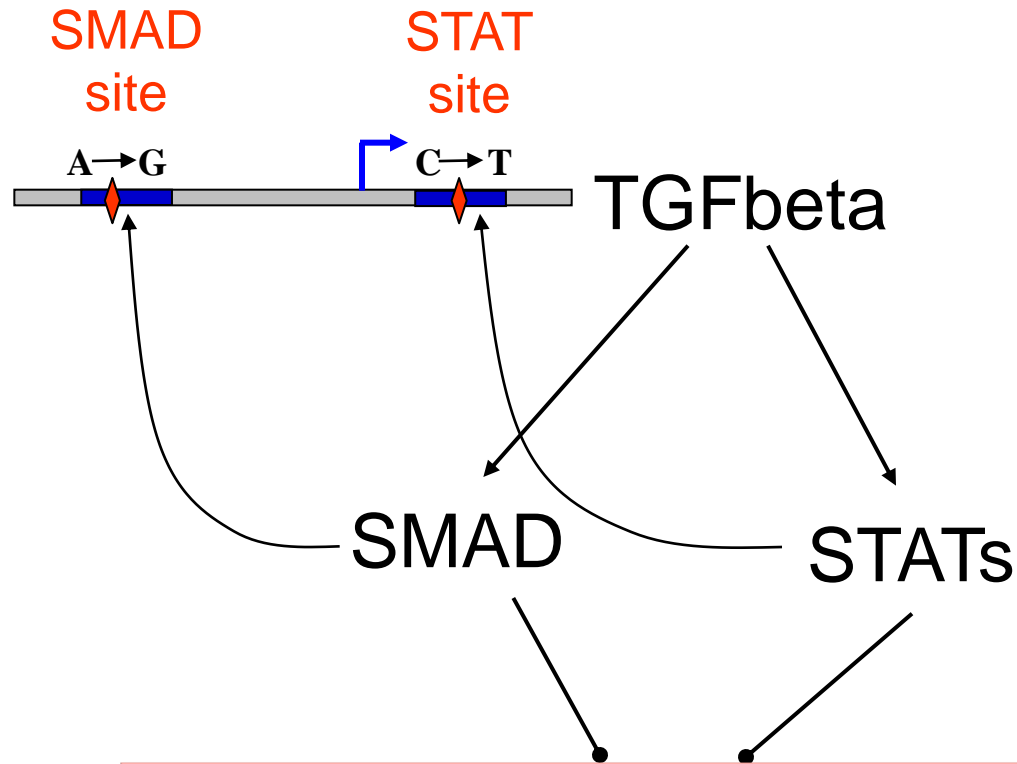
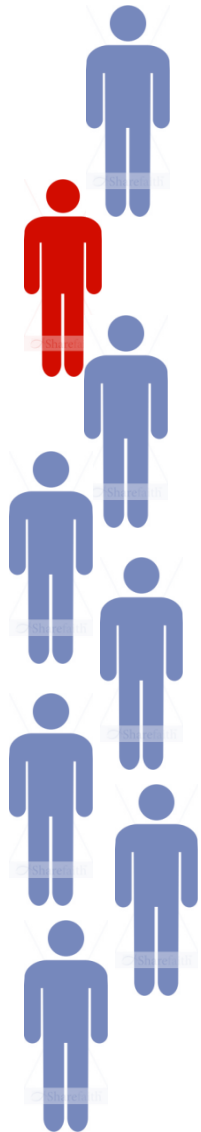
- We found TGF-beta1 as a master regulator – a potential off-target of TVX



TGF-beta dependent positive feedback



TGF-beta dependent positive feedback



Inhibition of genes of innate immune response

The successes achieved so far by the SMEs as partner in EU Health collaborative projects, which would not be possible without the EU collaborative grants

- **Creation of innovative tools became possible**
 - ▶ Effective drug target search algorithms.
 - ▶ Upstream algorithm for microarray data analysis.
 - ▶ Support for creation of geneXplain platform.
- **World-wide recognition of the company**
 - ▶ In academia.
 - ▶ In pharma and biotech.
- **Generation of a portfolio of know-hows.**
 - ▶ Disease specific drug targets/biomarkers.
 - ▶ Effective computational algorithms.

Potential for exploitation of the results achieved so far by the SME in EU Health collaborative projects.

- **Bioinformatics and systems biology software (Algorithms are created with EU help)**
 - ▶ Global market value of about 500 Mio €.
 - ▶ GeneXplain platform share is about 50 Mio €.
 - ▶ Potential customers: 50 large pharmaceutical and biotech companies + 500 middle-sized and smaller biotech companies + 3000 academic work groups .
- **Services (Expertise is created with EU help)**
 - ▶ Data analysis services and consulting.
 - ▶ Creation of tailor-made solutions (software, models)
- **Exploitation of identified drug targets and lead compounds.**
 - ▶ Cancer leads compounds should be further validated.

The importance, benefits and impact of Health collaborative research funding for SMEs in the area of systems biology for medical applications.

- **EU funding is ultimately important for systems biology SMEs**
 - ▶ Supporting innovations/ideas that do not show yet return of investments history.
 - ▶ Help young companies to start up
- **Benefits**
 - ▶ Collaboration synergism
 - ▶ Inventor owns the IPs.
- **Impact**
 - ▶ Creation of new company was possible only upon EU grant
 - ▶ R&D in SMEs in Europe are directed towards innovations
 - ▶ Gives great competitive advantage in future.

What kind of research opportunities could attract Industry and in particular biotechnology/healthcare SMEs in the field of systems medicine?

- **Creation of innovative tools – sustainable business**
 - ▶ Tools to support “omics” machines.
 - ▶ Tools to study disease mechanisms.
 - ▶ Tools for medical practice (virtual human/patient).
- **Participation in main stream research - marketing**
 - ▶ Learning from the best teams in the world.
 - ▶ Publications in the consortia – best marketing mechanism.
- **Opportunity to generate and co-own IPs on targets, biomarkers, drugs – opportunity business**
 - ▶ Various disease areas.
 - ▶ Increased chances due to synergy effect.

Explain genes to find new drugs



www.geneXplain.com