

*In silico, in vitro, in vivo* alternatives to the  
carcinogenicity bioassay

**Romualdo Benigni**

**Istituto Superiore di Sanita'  
Rome Italy**

**[rbenigni@iss.it](mailto:rbenigni@iss.it)**

# Mechanistic findings at the basis of the science and regulation of mutagens and carcinogens

- **Millers'** electrophilic reactivity theory of carcinogenesis  
(not including nongenotoxic carcinogens)
- **Chemical mutagenicity:**  
**Malling's** *in vitro* **S9** metabolic activation;  
**Salmonella, or Ames'** test for DNA-reactive chemicals
- **Ashby's** theoretical model of carcinogenicity (compilation of **Structural Alerts**)

# Mechanistic findings at the basis of the science and regulation of mutagens and carcinogens

- Because of the success **of Millers'** electrophilic reactivity theory of carcinogenesis, and of **Ames'** test:
  - major research efforts on the hypothesis **Mutation = Cancer**
- Later on, recognition of **nongenotoxic carcinogens**

# Looking for further mutagenicity Short-Term Tests (STT) to predict carcinogenicity

- Hypothesis:  
Different **genetic endpoints** (gene mutation, chromosomal damage),  
and different **cells** (bacterial, mammalian) to cover the spectrum of  
cancer-relevant factors
- Development of **> 100 STTs** based on:  
**mutagenicity** (e.g., gene mutation in mammalian cells, chromosomal  
aberrations, aneuploidy)  
  
other **genotoxic** events (e.g., DNA damage)

# Comparative carcinogenicity prediction exercise:

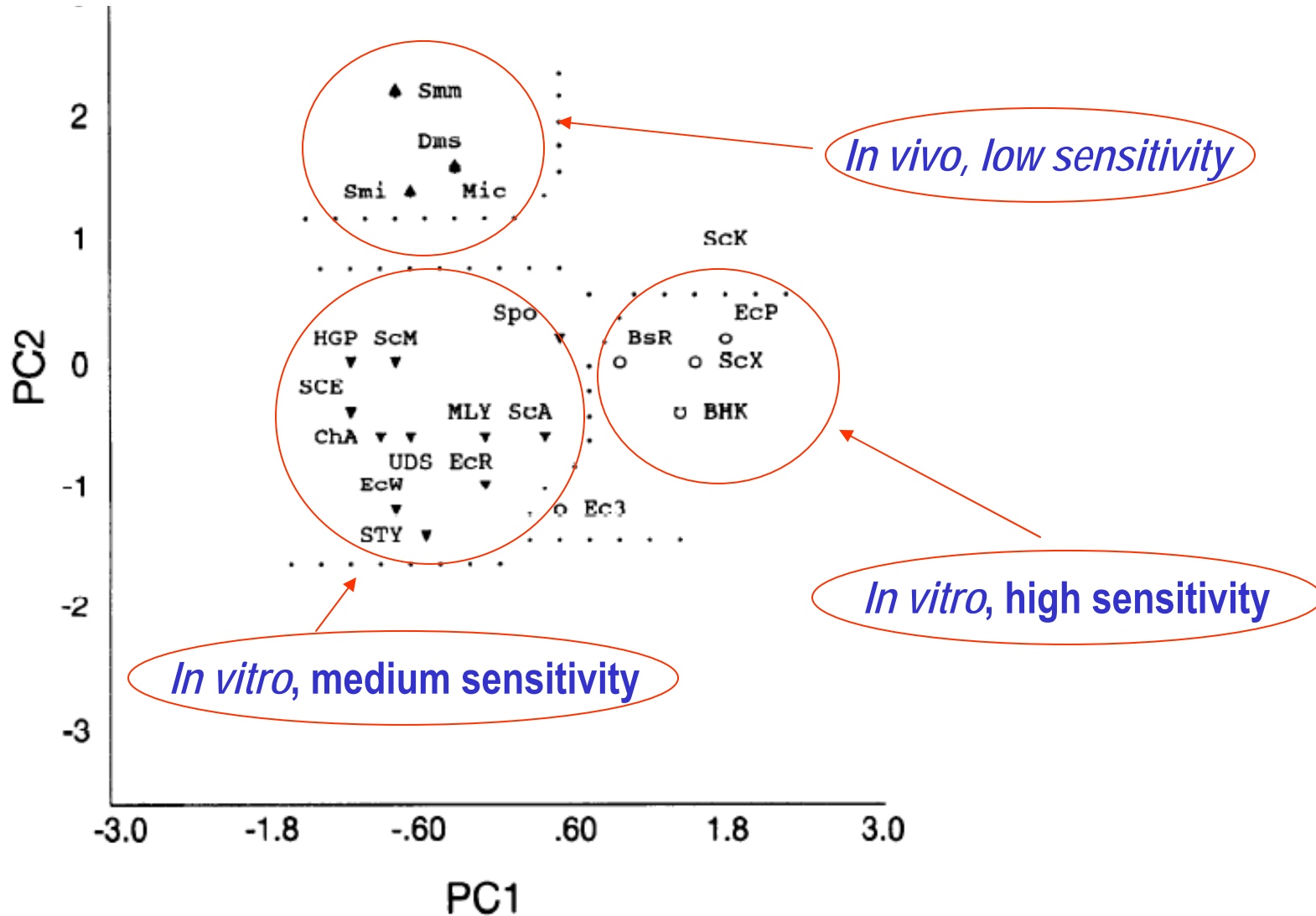
**22 mutagenicity STTs x 42 chemicals**

De Serres and Ashby, 1981, Evaluation of Short-Term Tests for Carcinogens,  
Progr. Mutat.Res. Vol 1

## Conclusion(s):

- Central role for *Salmonella*
- Continue with exploration of different **genetic endpoints** and different **cells**

# Comparative carcinogenicity prediction: 22 mut. STTs x 42 chemicals

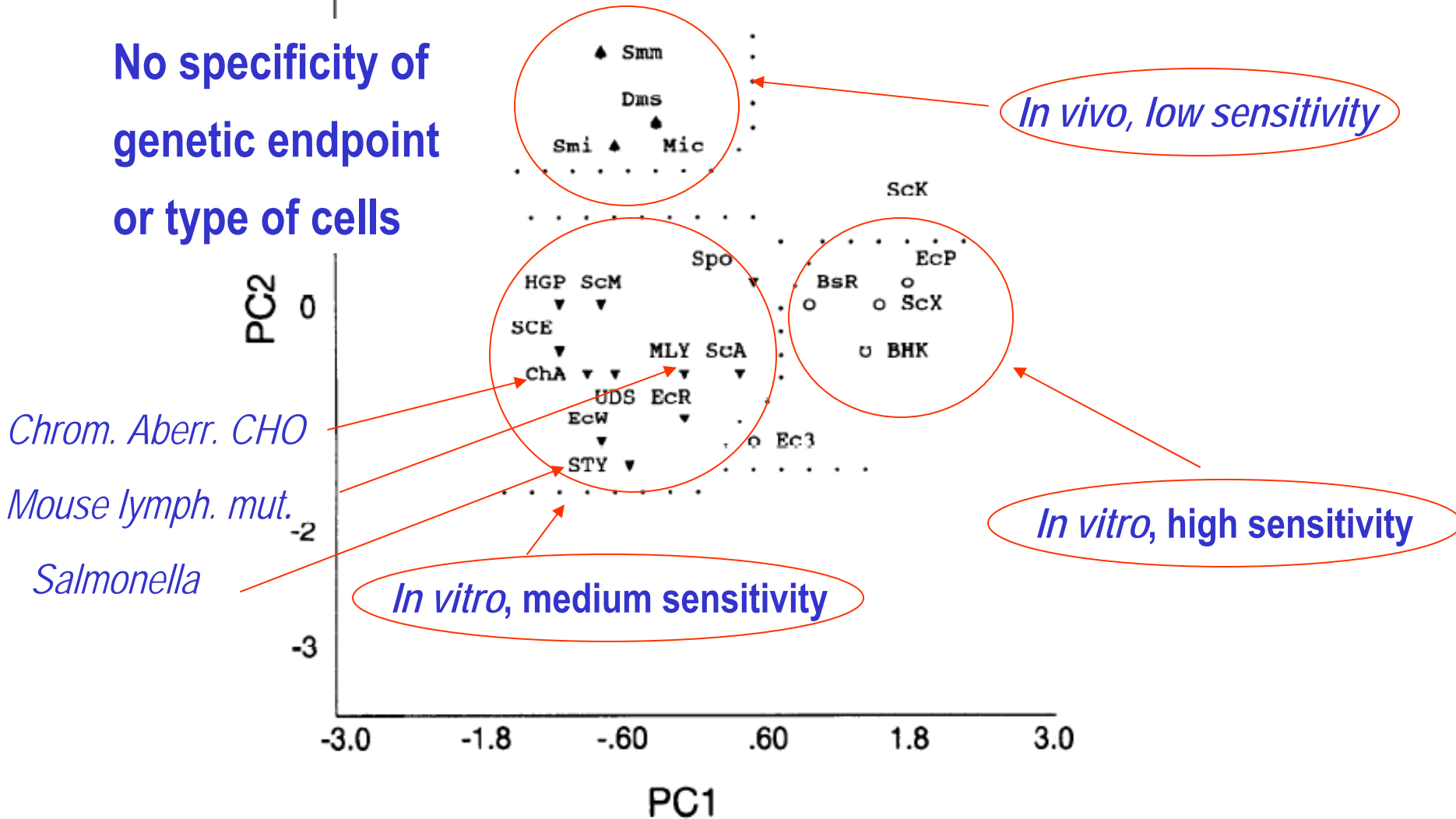


**Analysis:** Benigni and Giuliani, 1985, *Mutat.Res.*, 147: 139 - 151

**Data from:** De Serres and Ashby, 1981, *Evaluation of Short-Term Tests for Carcinogens*, *Progr. Mutat.Res.* Vol 1

# Comparative carcinogenicity prediction: 22 mut. STTs x 42 chemicals

No specificity of genetic endpoint or type of cells

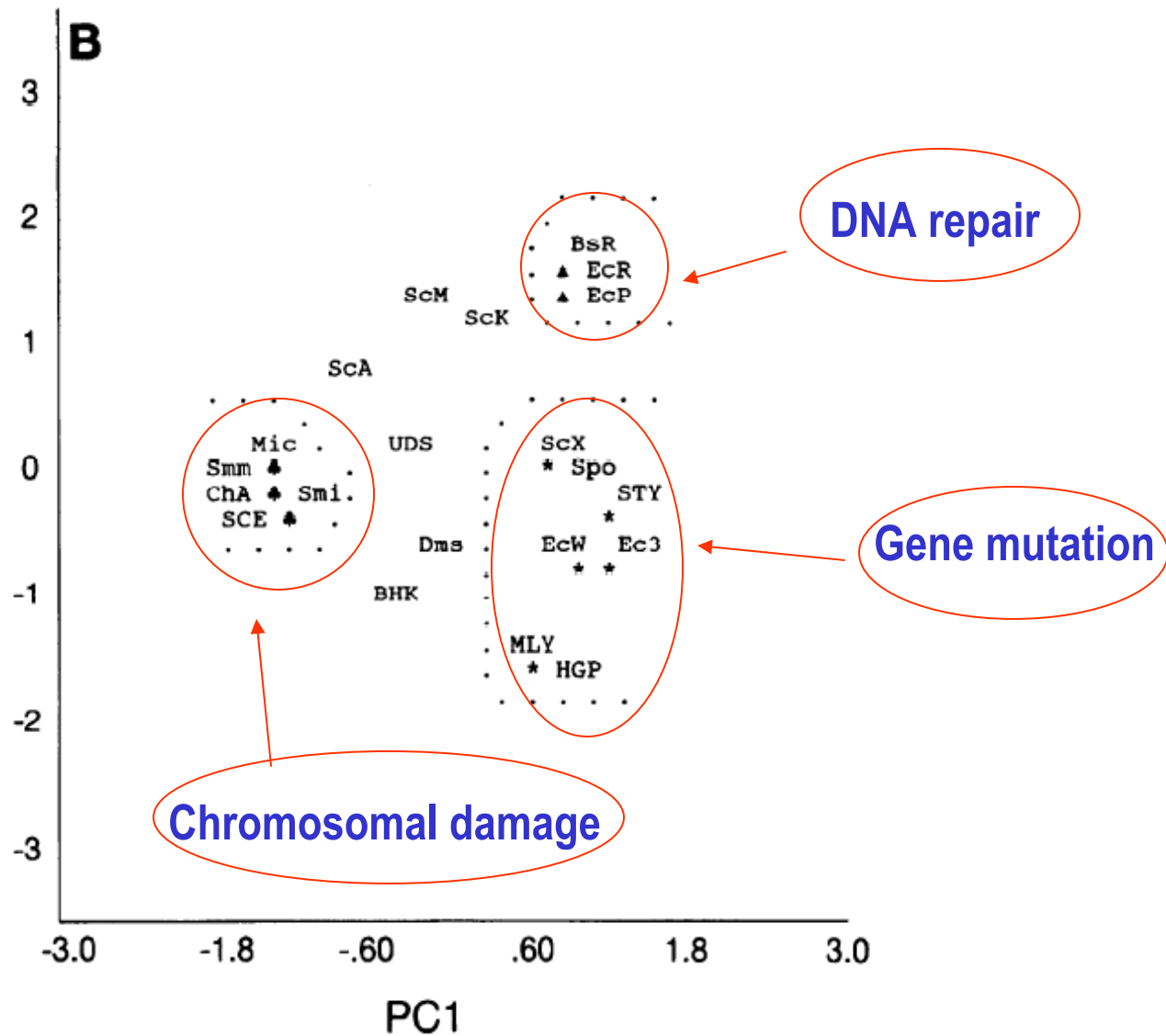


**Analysis:** Benigni and Giuliani, 1985, Mutat.Res., 147: 139 - 151

**Data from:** De Serres and Ashby, 1981, Evaluation of Short-Term Tests for Carcinogens, Progr. Mutat.Res. Vol 1

# A questionnaire for biologists:

how 22 mutagenicity STTs will respond to 42 chemicals ?





# Looking for further mutagenicity Short-Term Tests to predict carcinogenicity

- Looking for complements to *Salmonella (Ames)*:  
**National Toxicology Program evaluation of four *in vitro* STTs**  
(n. chemicals = 114)
- Different **genetic endpoints** (gene mutation, chromosomal damage), and different **cells** (bacterial, mammalian)

*Salmonella typhimurium* (Ames)

Chromosomal Aberrations in CHO cells

SCEs in CHO cells

Mouse Lymphoma mutation

*Tennant et al., 1987 Science 236: 933-941*

*Zeiger et al., 1990 Environ. Mut. Mutagen. 16(18): 1-14*

# Relevance of STTs to rodent carcinogenicity

Chi-square (p)

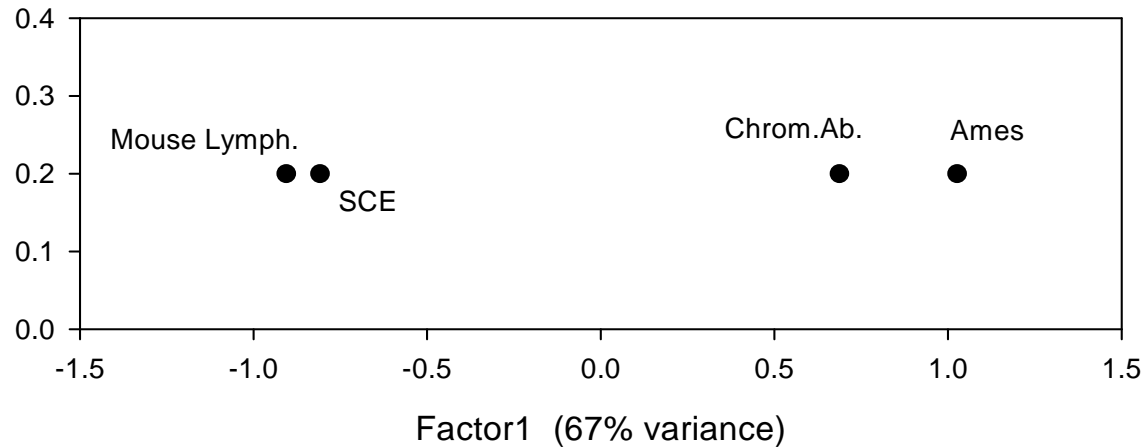
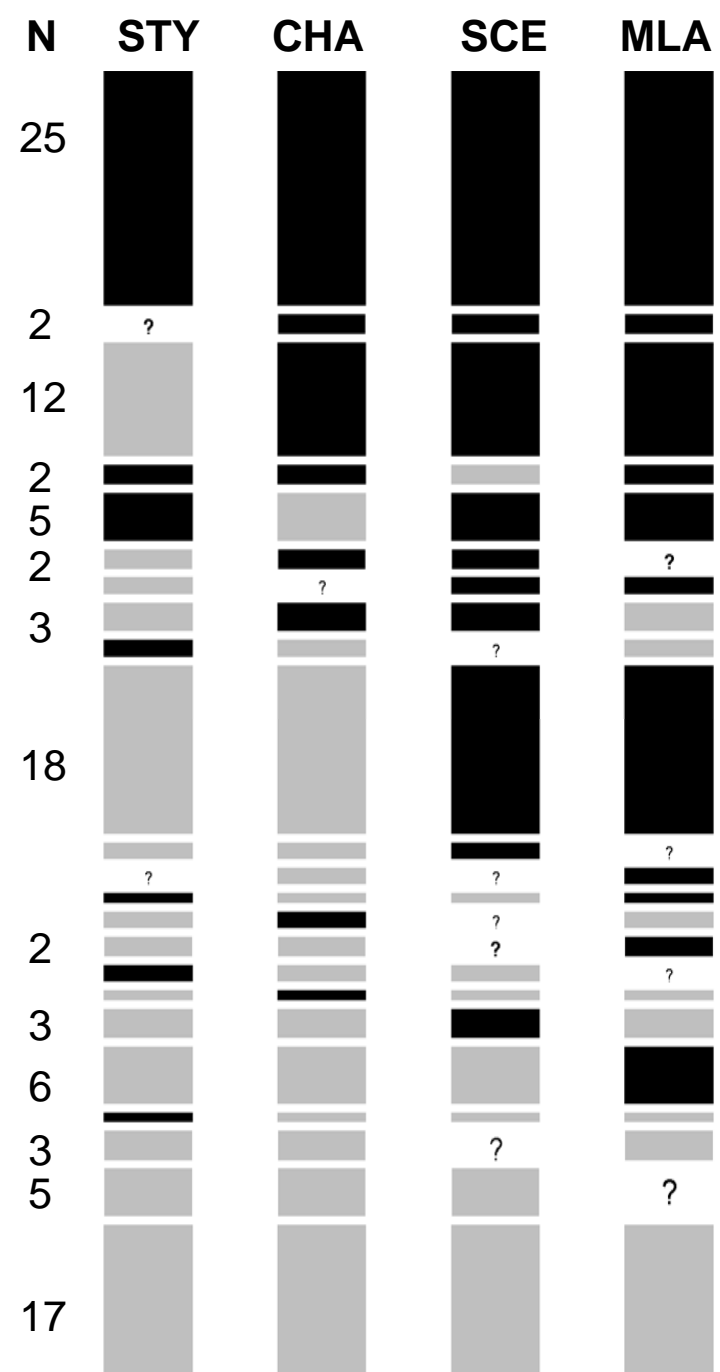
Ames	< 0.0001
Chrom. Ab (CHO)	0.011
Sce (CHO)	0.277
Mouse Lymphoma mut.	0.305

## **Batteries**

Ames + Chrom.Ab.	0.0004
Ames + Mouse Lymphoma	0.120

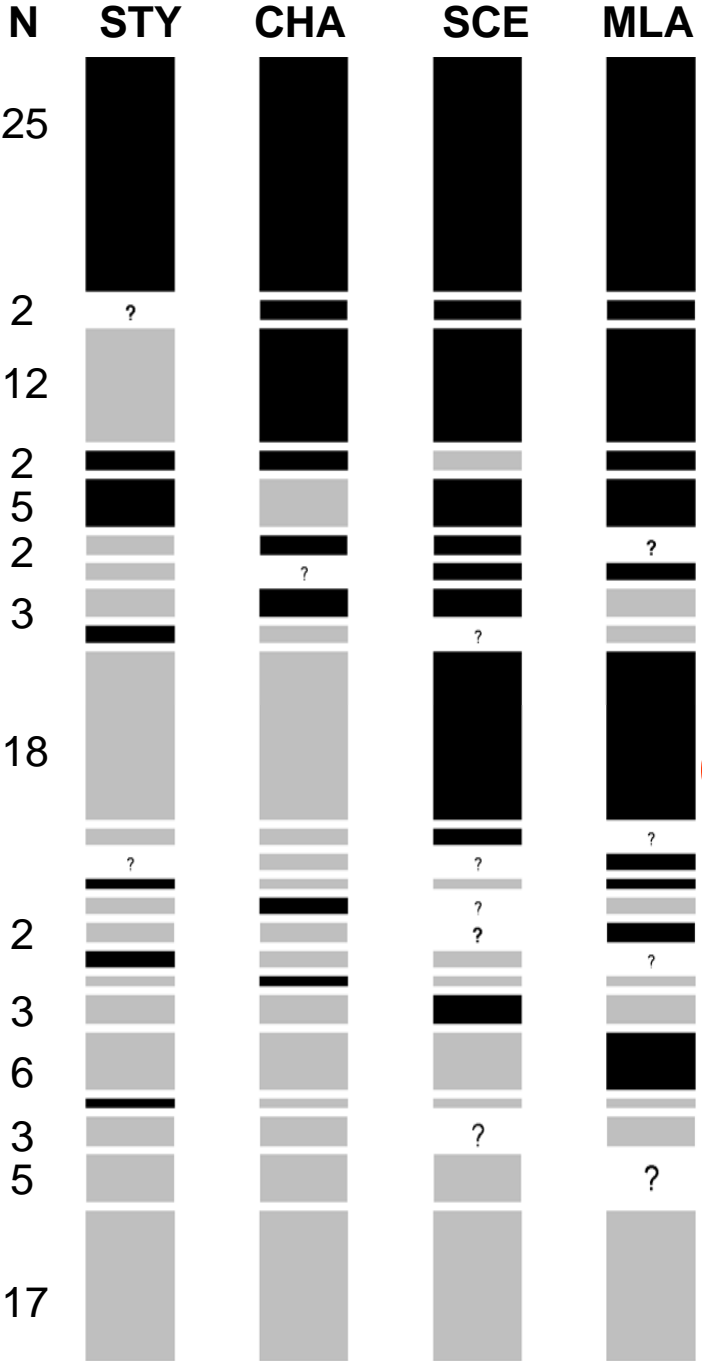
*(our elaborations)*

# Genetic Toxicity Profiles: 114 NTP Chemicals



positive  
 negative  
 ? equivocal

# Genetic Toxicity Profiles: 114 NTP Chemicals



Positive in Salmonella (and other tests):  
high correlation with carcinogenicity

Negative in Salmonella and positive in other tests:  
no correlation with carcinogenicity

- positive
- negative
- ? equivocal

# STTs to predict carcinogenicity

- **Mutagenicity = Carcinogenicity ?**

Only within a limited area of the chemical space, i.e., **DNA-reactive** chemicals

- DNA-reactive chemicals induce **cancer, together with a wide spectrum of mutations**
- Most predictive mutagenicity-based assay: **Ames test**
- **Non DNA-reactive chemicals**, mutagenic in other *in vitro* assays (e.g., clastogenicity) but Ames-negative: no correlation with carcinogenicity
- No reliable *in vivo* **STT** (e.g., micronucleus) is available

*Benigni R. et al., Exp.Opinion Drug Metab.Toxicol., 2010, 6: 1-11..*

*Zeiger E Regulat.Pharmacol.Toxicol. 1998;28:85-95.*

# Ames test *versus* rodent carcinogenicity

		Ames test	
		neg	pos
Carcinogenicity			
Carcinogens {	Non carcinogens	233	76
	Non DNA-reactive	136	34
	DNA-reactive	79	277

**Ames identifies DNA-reactive carcinogens**

*Results from 835 chemicals in ISSCAN v3a*

<http://www.iss.it/ampp/dati/cont.php?id=233&lang=1&tipo=7>

# Ames test *versus* rodent carcinogenicity

		Ames test	
		neg	pos
Carcinogenicity			
Non carcinogens		233	76
Carcinogens {	Non DNA-reactive	136	34
	DNA-reactive	79	277

**Ames mutagen: 80% probability of being a carcinogen**

*Results from 835 chemicals in ISSCAN v3a*

<http://www.iss.it/ampp/dati/cont.php?id=233&lang=1&tipo=7>

# Prediction of carcinogenicity by the STTs

## *Mechanistic classes*

*Ames*

*Chrom. Aberrations*

Chi-square (p values)

### Direct Alkylating Agents

All	0.0049	0.0001
Epoxides	0.0013	NS
Aliphatic halogens	NS	0.0055

### Indirect Alkylating

All	NS	NS
-----	----	----

too few data for most classes

### Intercalating

All	NS	---
-----	----	-----

### Aminoaryl DNA-adducts forming

Aromatic amines	0.0074	NS
Nitroarenes	0.0443	NS



Backing up the STTs with **Structure-Activity** concepts

## **Structure-activity relationship** concepts:

application to different issues, through different approaches

*Coarse-grain*

### **Structure Alerts**

(mechanistic classes, category formation, priorities)

*Fine-tuned*

Quantitative Structure-Activity Relationships (**QSAR**)  
of congeneric classes

*Hybrid (??)*

non-local, or global QSARs

# Toxtree: Rulebase for mutagens / carcinogens

Structure-based approach consisting of:

- New compilation of *Structure Alerts* (genotox and non-genotox)
- Three mechanistically-based *QSARs* for congeneric classes (aromatic amines, aldehydes)

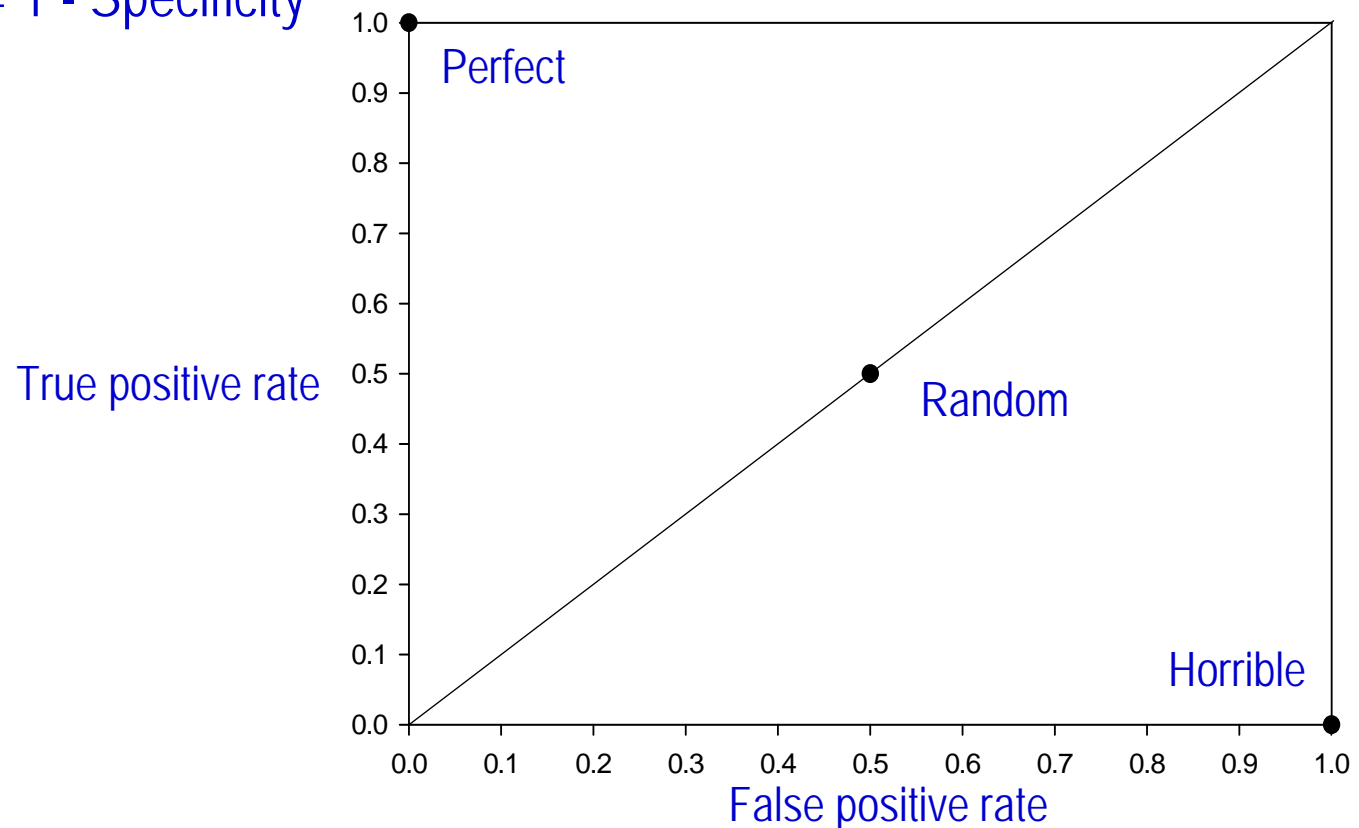
**Toxtree** (version 1.6)

Open-source, freely available: <http://ecb.jrc.it/qsar/qsar-tools/index.php?c=TOXTREE>

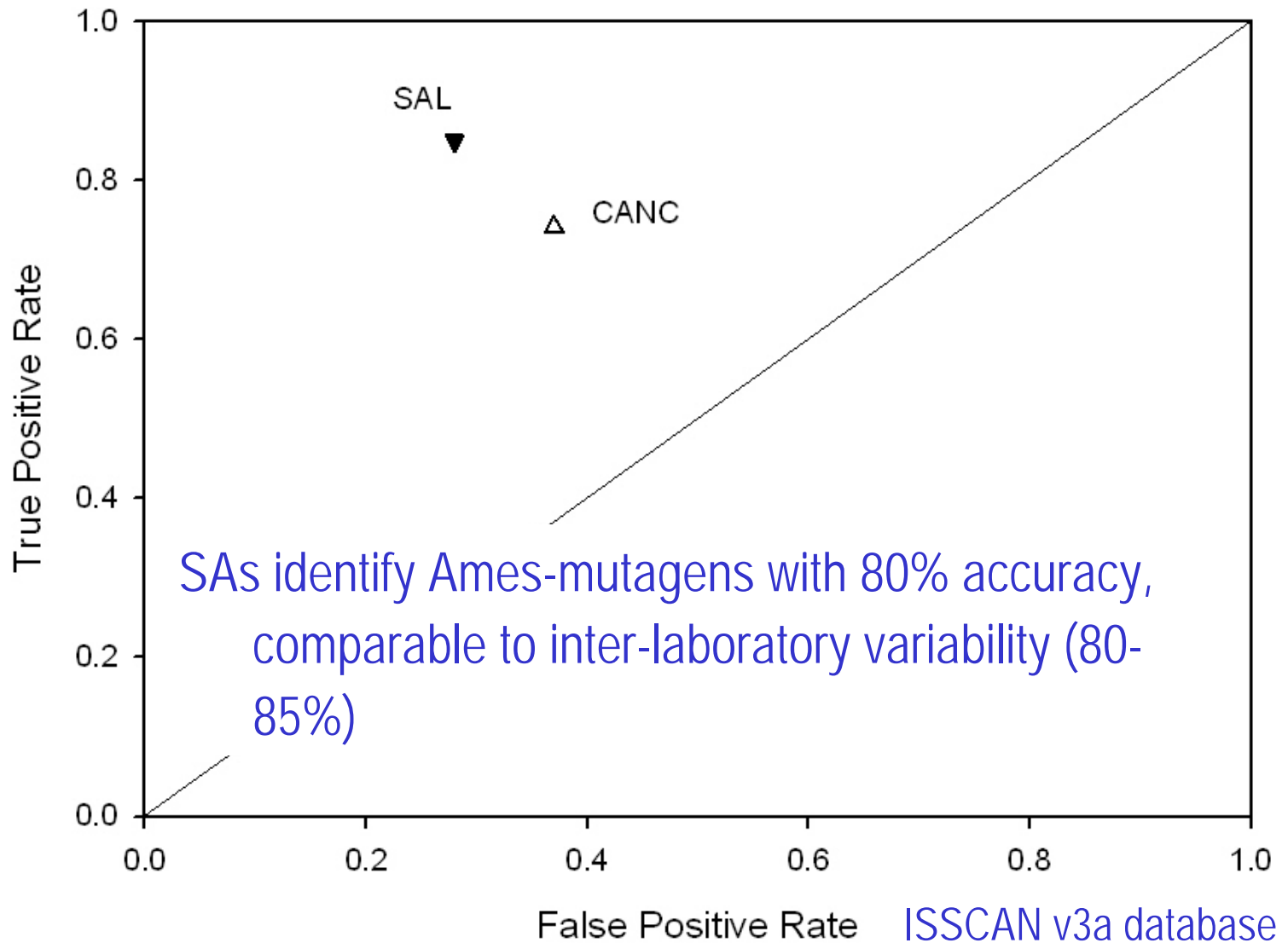
# ROC graph: A simple, graphical way of comparing predictions with results

True positive rate = (Positives predicted as positive) / (Real positives)  
= Sensitivity

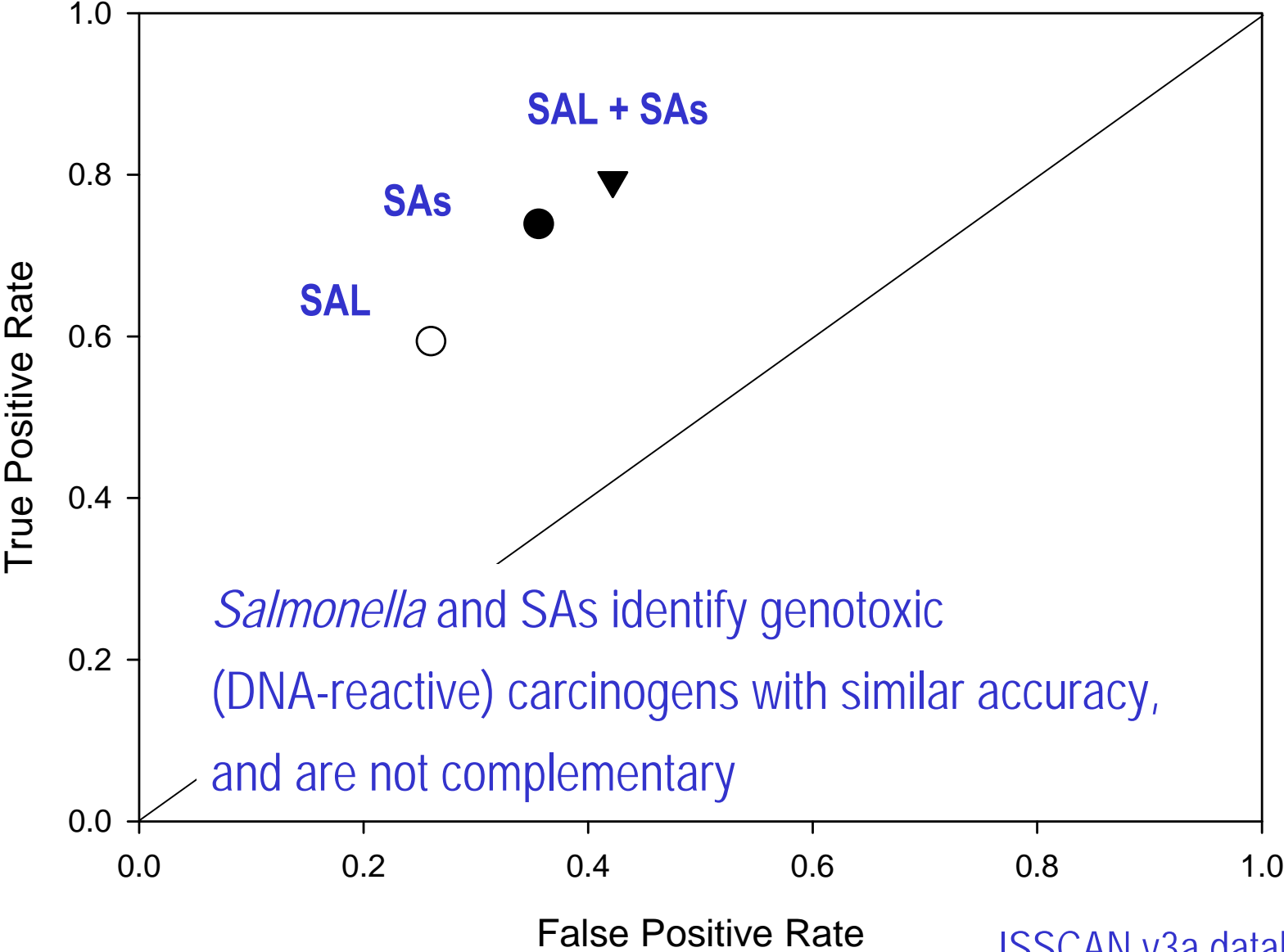
False Positive Rate = (Negatives predicted as positive) / (Real negatives)  
= 1 - Specificity



# Toxtree SAs: agreement with Carcinogenicity and *Salmonella* (Ames)



# Carcinogenicity prediction: *Salmonella* (Ames) versus SAs



# Which use for the Structural concepts (Alerts) ?

## Success story: priority setting by human experts

Out of 400 chemicals tested by NCI/NTP:

- 2/3 selected as suspect carcinogens (n=267)  
68% carcinogenic (n=187)
- 1/3 selected on production/exposure considerations (n=133)  
20% carcinogenic (n=26); 6.8% positive in two species (n=9)

# QSARs of Aromatic amines

Carcinogenic **activity** in rodents

$$\text{Canc} = -1.16 \text{ HOMO} + 1.76 \text{ LUMO} - 2.86 \text{ L(R)} + 2.65 \text{ B5(R)} + 0.40 \text{ MR}_3 \\ + 0.58 \text{ MR}_5 + 0.54 \text{ MR}_6 - 1.55 \text{ I(An)} + 0.74 \text{ I(NO}_2) - 0.55 \text{ I(BiBr)}$$

n = 66 (- = 44; + = 73) Correct Classification = 87.9 %

Franke et al., 2001

Mutagenic **activity** in *Salmonella typhimurium* TA100 (+ S9)

$$\begin{array}{ccc} \text{Electronic} & & \text{Steric} \\ \downarrow & & \downarrow \\ \text{ActTA100} = 0.67 \text{ HOMO} - 0.75 \text{ LUMO} - 0.39 \text{ MR}_2 - 0.38 \text{ MR}_3 - 0.44 \text{ MR}_6 \\ & & - 0.62 \text{ Idist} \end{array}$$

n = 111 (- = 47; + = 64) Correct Classification = 87. %

Benigni et al., 2007



# European Chemicals Bureau - ISS Project

## Assessment of local QSARs for congeneric classes:

- scientifically (mechanistically) interpretable
- good internal statistics
- **real external predictivity** tested
- applicability domain checked:
  - functional group
  - parameters range
  - chemical similarity

## Conclusions on the local QSARs:

- scientifically interpretable, good internal statistics, but vary for their external predictivity
- **QSARs for potency: predictions 30 – 70 % correct**
- **QSARs for activity: predictions 70 – 100 % correct**
- Estimating intervals more reliable than estimating points
- **Internal validation measures do not correlate with external predictivity**

## Which use for local QSARs ?

- Local QSARs for activity: 70 – 100 % correct external predictions
- Intra-Assay (inter-laboratory) agreement for the Ames test: 80 – 85%

*Piegorsch and Zeiger, 1990, in Statistical methods in Toxicology*

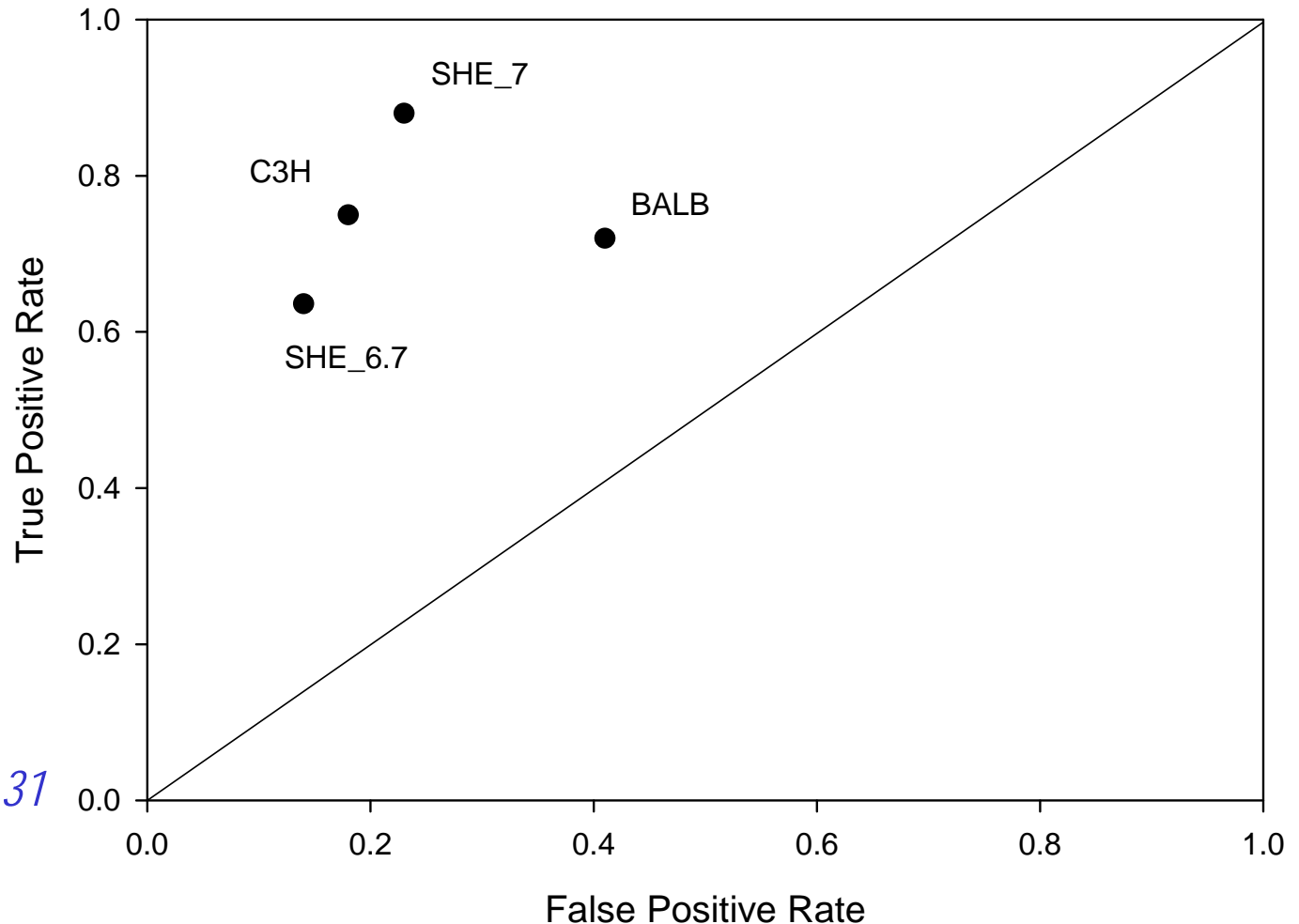
**Same range of uncertainty**

**Non-mutagenicity assays** for predicting carcinogenicity ?

# Cell transformation assays

in cultured cells, phenotypic alterations characteristic of tumorigenic cells

Cell Transformation: Carcinogenicity prediction



**Data:** *OECD*

*Series on Testing and  
Assessment, 2007, vol.31*

**Our elaboration**

# SHE pH $\geq$ 7 Cell transformation *versus* rodent carcinogenicity

Carcinogenicity	SHE	
	neg	pos
Negatives	33	11
Non DNA-reactive	6	28
DNA-reactive	5	58

A blue bracket on the right side of the table groups the 'Non DNA-reactive' and 'DNA-reactive' rows under the label 'Carcinogens'. A red oval highlights the 'Non DNA-reactive' and 'DNA-reactive' rows, and a red arrow points from this oval to the text below.

**Sensitive to DNA-reactive and non-DNA-reactive carcinogens**

*Chemicals n = 141, from OECD vol. 31*

## Great challenges:

overcoming the *in vitro* / *in vivo* gap

identifying “useful” steps in toxicity pathways

and more...

## *In vitro* alternatives to carcinogenicity

The equation *mutation = cancer* is not of general validity

Out of > 100 STTs, only *Salmonella* and **cell transformation** (to be confirmed on larger database) predict chemical carcinogenicity

Present testing strategies to be revised ?



## *in vitro* alternatives: state-of-the-art

- Toxicokinetic
- **Acute toxicity** *Reduction*
- **Skin irritation and corrosion** *Replacement*
- Skin sensitisation
- **Eye irritation** *Reduction*
- Acute systemic and local toxicity
- **Genotoxicity** *Reduction*
- **Carcinogenicity** *Reduction*
- Repeated dose toxicity
- Reproduction
- Developmental toxicity
- Ecotoxicity

“Existing approaches incorporating replacement, reduction and refinement of animal testing: applicability in food and feed risk assessment” *The EFSA Journal* (2009) 1052, 1-77

# Toxcast Phase 1

## *Pathway-based screening* approach

Perturbations of biochemical / biological pathways supposedly related to toxicity

## **Extensive use of new omics technologies**

Isolated, *in vitro* systems

**309 unique chemicals**

Mostly agrochemicals

- **76 in vivo bioassays**

- ToxRefDB

- Target organs (chronic), reproductive, developmental, **carcinogenicity**

## **524 in vitro assays**

- 9 *in vitro* assay providers

- 285 cell-based, 239 cell-free

- **can *in vivo* endpoints be predicted through *in vitro* ones ?**

# Averting chance correlations

## Cleaning Toxcast *in vitro* data

- **Statistical robustness:** 6 clusters of *in vitro* assays  
(Cluster and Principal Component Analysis of response profiles)
- The clusters are **biologically meaningful:**
  - Cluster 1: *Cell growth, cell adhesion and shape, etc... (immune-related)*
  - Cluster 2: *Signaling (cell cycle, apoptosis...)*
  - Cluster 3: *Nervous system factors*
  - Cluster 4: *Metabolic factors*
  - Cluster 5: *Cell growth, inflammation*
  - Cluster 6: *Transcription factors, gene activation*

# Prediction of Toxcast carcinogenicity through *in vitro*

- Discard erratic results
- Pick five representative assays from each cluster

Assays	Squared Canonical Correlation (sqcc)	
	<u>Mouse</u>	<u>Rat</u>
Cluster 1	0.02	0.03
Cluster 2	-	-
Cluster 3	-	-
Cluster 4	0.04	0.01
Cluster 5	0.01	-
Cluster 6	-	0.03

# Prediction of Toxcast carcinogenicity through *in vitro*

- All representative assays (n=30) are used

Assays	sqcc	
	<u>Mouse</u>	<u>Rat</u>
30 representatives	0.04	0.09

- Toxcast *in vitro* assays are **poorly correlated** with animal carcinogenicity

Benigni, submitted

[http://www.epa.gov/NCCT/toxcast/files/summit/ToxcastDataSummit\\_Poster\\_Benigni%20May2009.ppt](http://www.epa.gov/NCCT/toxcast/files/summit/ToxcastDataSummit_Poster_Benigni%20May2009.ppt)

## Great challenges:

overcoming the *in vitro* / *in vivo* gap

identifying “useful” steps in toxicity pathways

and more...

- **Never take the scientific constructs at face value**
- **Always contrast theories with data**

**The best laid schemes of mice and men go often askew**

*(R. Burns, 1785)*

# Acknowledgements

- Istituto Superiore di Sanita'

**Alessandro Giuliani**

**Cecilia Bossa**

**Olga Tcheremenskaia**





# Structural Alerts: predictive ability

	N	Actives	%
<b><u>Alkylating, direct acting</u></b>			
<b>SA_2: alkyl (C &lt; 5) or benzyl ester of sulphonic or phosphonic acid</b>	<b>13</b>	<b>12</b>	<b>92</b>
SA_5: S or N mustard	10	10	100
SA_7: epoxides and aziridines	28	20	71
SA_8: aliphatic halogens	75	51	68
SA_10: a,b-unsaturated carbonyls	55	37	67
SA_11: simple aldehyde	9	8	89
<b>SA_12: quinones</b>	<b>15</b>	<b>13</b>	<b>87</b>
<b><u>Alkylating, indirect acting</u></b>			
<b>SA_13: hydrazine</b>	<b>68</b>	<b>57</b>	<b>84</b>
SA_14: aliphatic azo and azoxy	8	8	100
SA_16: alkyl carbamate and thiocarbamate	8	7	88
<b>SA_21: alkyl and aryl N-nitroso groups</b>	<b>120</b>	<b>105</b>	<b>88</b>

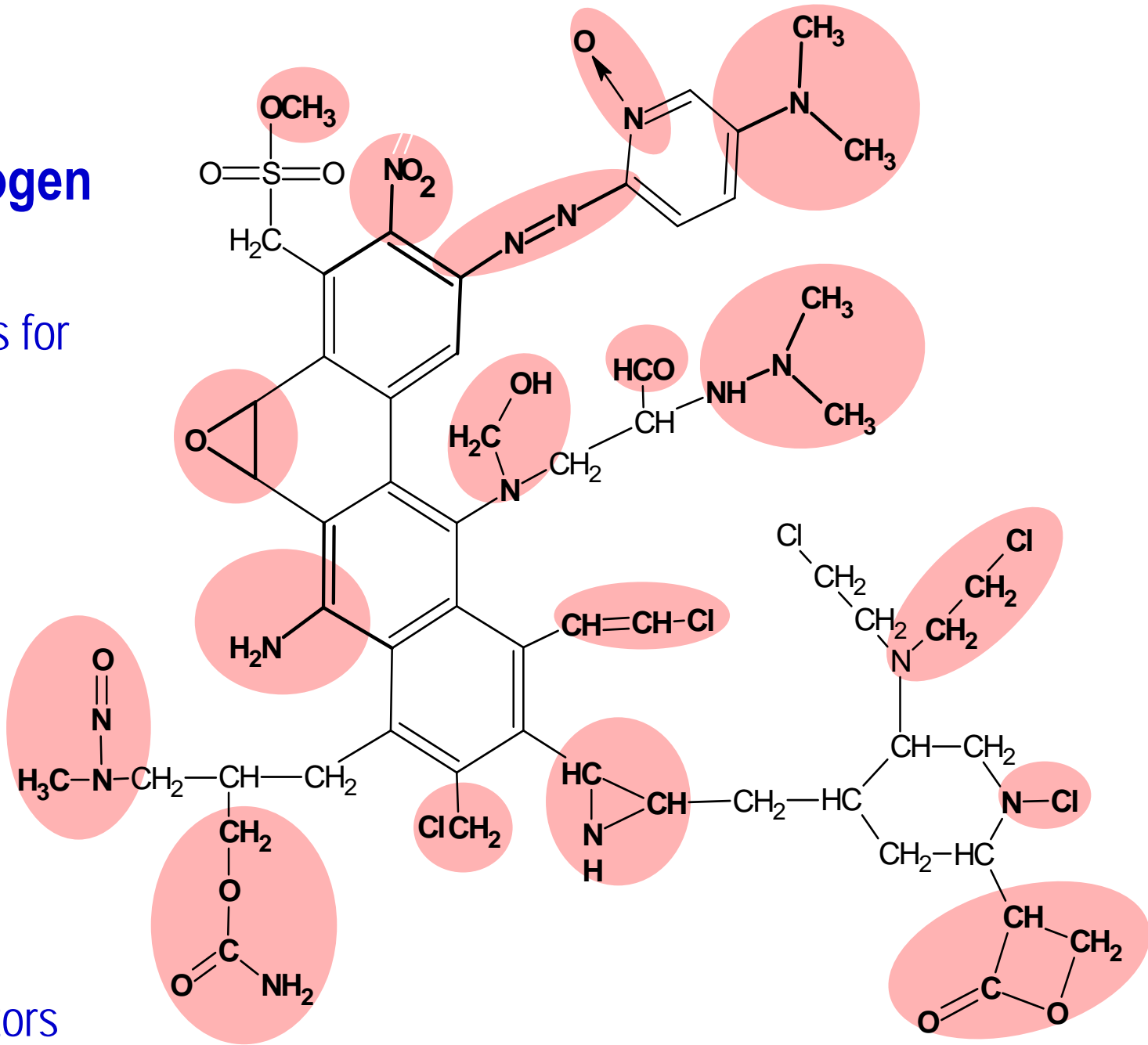
# Structural Alerts: predictive ability

	N	Actives	%
<b><u>Intercalating and DNA-adducts forming, indirect acting</u></b>			
SA_18: polycyclic aromatic hydrocarbons	14	11	79
<b>SA_19: heterocyclic polycyclic aromatic hydrocarbons</b>	<b>14</b>	<b>12</b>	<b>86</b>
<b><u>Aminoaryl DNA-adducts forming, indirect acting</u></b>			
SA_27: nitro-aromatic	88	63	72
<b>SA_28: primary aromatic amine, hydroxyl amine and its derived esters</b>	<b>107</b>	<b>87</b>	<b>81</b>
SA_28bis: aromatic mono- and dialkylamine	15	10	67
SA_28ter: aromatic N-acyl amine	17	13	76
SA_29: aromatic diazo	26	20	77
<b><u>Nongenotoxic</u></b>			
SA_20: (poly) halogenated cycloalkanes (nongenotoxic)	24	13	54
SA_17: thiocarbonyl (nongenotoxic)	18	14	78
SA_31a: halogenated benzene (nongenotoxic)	16	5	31
SA_31b: halogenated PAH (nongenotoxic)	10	8	80

# Ashby's Poly-carcinogen

- Structural alerts for carcinogenicity;
- DNA-reactive functionalities

Some alerts  
accompanied by  
detoxifying  
(modulating) factors



## ECB Activities

[Biocides](#)
[Classification & Labelling](#)
[Computational Toxicology](#)
[Existing Chemicals](#)
[Export-Import](#)
[New Chemicals](#)
[REACH](#)
[Testing Methods](#)
[EDEXIM](#)
[ESIS](#)
[IUCLID 5](#)
[Contacts](#)
[Documentation](#)
[Legislation](#)
[Links](#)
[Newsletter](#)
[Search](#)
[Site Map](#)
[What's New](#)

## Toxtree

Toxtree is a flexible and user-friendly open-source application that places chemicals into categories and predicts various kinds of toxic effect by applying decision tree approaches, such as the [Cramer classification scheme](#), the Verhaar scheme for aquatic modes of action, and a rulebase for skin irritation and corrosion based on rules developed by the German Federal Institute for Risk Assessment (BfR) and collaborators.

Toxtree was developed by IdeaConsult Ltd (Sofia, Bulgaria) under the terms of an ECB contract. The software is made freely available by ECB as a service to scientific researchers and anyone with an interest in the application of computer-based estimation methods in the assessment of chemical toxicity.

### Toxtree (Version 1.20) - Download area

Following the original release of Toxtree (v1.00) in March 2007, a new version with additional functionalities was released in March 2007.

## Toxtree (Estimation of Toxic Hazard - A Decision Tree Approach)

File Edit Chemical Compounds Toxic Hazard Method Help

<< >> Enter SMILES:

### Available structure attributes

Names	Created from SMILES
SMILES	CCCCC

Toxic Hazard

Structural Alert for g

Alert for ne

or carcinog

. typhimur

be a 5. typ

SAR

arcinogen

be a carcin

r assessm

explanation

### Select a tree

Available decision trees

Load from file

Cramer rules

Verhaar scheme

Benigni / Bossa rulebase (for mutagenicity and carcinog...)

Skin irritation / skin corrosion

[Benigni / Bossa rulebase (for mutagenicity and carcinogenicity)]

Demo substructure tree

Predicts the possibility of carcinogenicity and mutagenicity by discriminant analysis and structural rules. See The Reference guide.

# Non-local, “global” QSARs demonstrated problems in prediction exercises of external chemicals, e.g.,

## NTP-1 carcinogenicity prediction exercise

*Benigni,R. (1997): The first US National Toxicology Program exercise on the prediction of rodent carcinogenicity: definitive results. Mutat.Res., 387:35-45.*

## NTP-2 carcinogenicity prediction exercise

*Benigni,R. and Zito,R. (2004): The second National Toxicology Program comparative exercise on the prediction of rodent carcinogenicity: definitive results. Mutat.Res.Revs., 566:49-63.*

## Predictive Toxicology Challenge 2000-2001

*Helma,C. and Kramer,S. (2003): A survey of the Predictive Toxicology Challenge 2000-2001. Bioinformatics, 19:1179-1182.*

## Miscellaneous independent assessments

*Benigni,R. and Bossa,C. (2008): Predictivity of QSAR. J.Chem.Inf.Model., 48:971-980.*

# Future

## QSAR modeling:

- Identify further alerts for nongenotoxic carcinogens
- Build further local QSARs for congeneric classes

## More work on *in vitro* methods

Integrate *in vitro* and QSAR –to overcome uncertainties on both sides-