The logo of the American College of Toxicology is a circular seal. The outer ring contains the text "AMERICAN COLLEGE OF TOXICOLOGY" in a serif font. Inside the ring is a sunburst design with a central circle and radiating lines. Below the sunburst is an open book. At the bottom of the seal is a banner with the Latin motto "educere ducere".

Welcome to the American College of Toxicology's Webinar Series

In Collaboration with the



British Toxicology Society

SOT

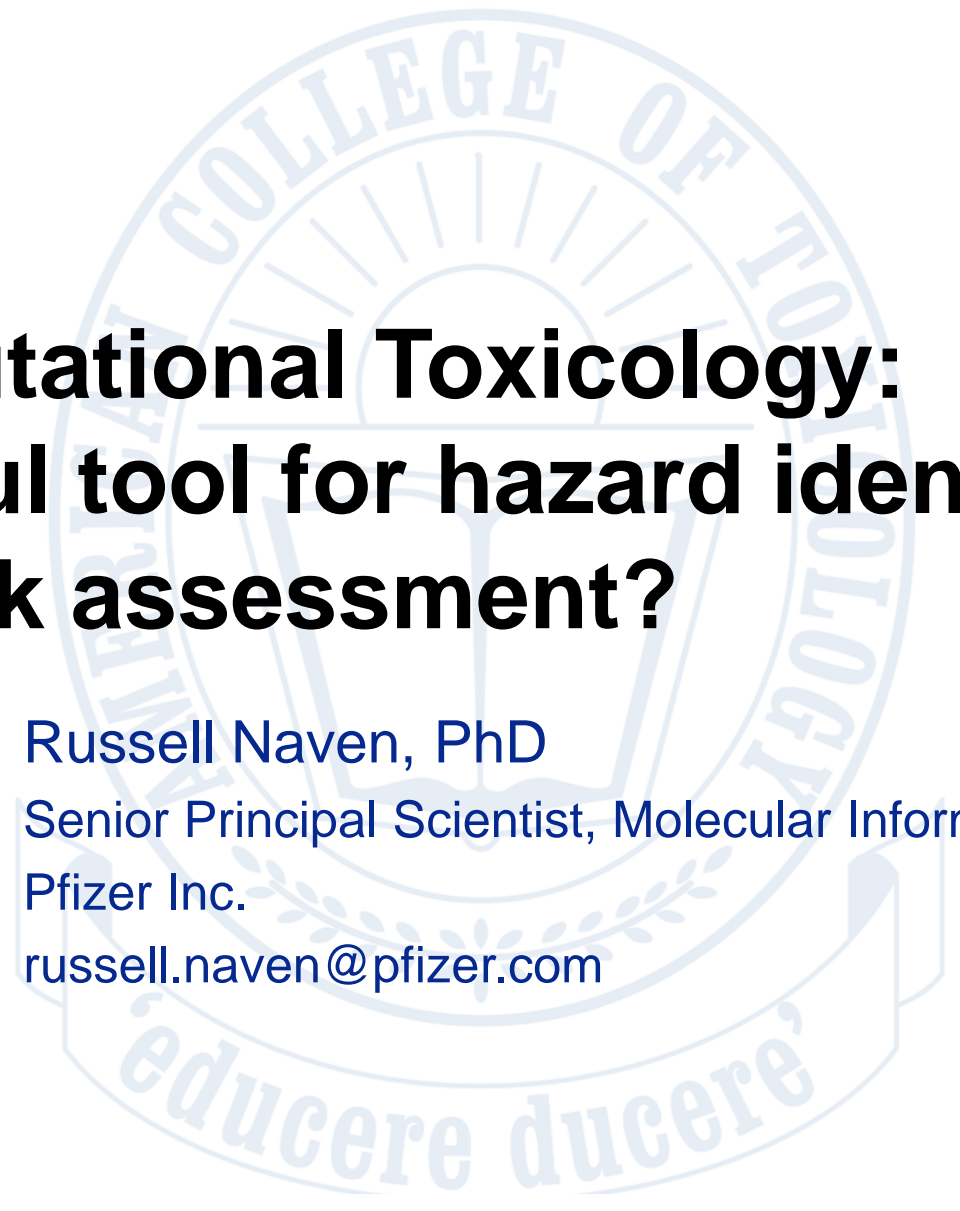
Biological Modeling
Specialty Section



Society of Toxicologic Pathology



Teratology Society



Computational Toxicology: A useful tool for hazard identification and risk assessment?

Russell Naven, PhD

Senior Principal Scientist, Molecular Informatics

Pfizer Inc.

russell.naven@pfizer.com

Presenter

Russell Naven, PhD

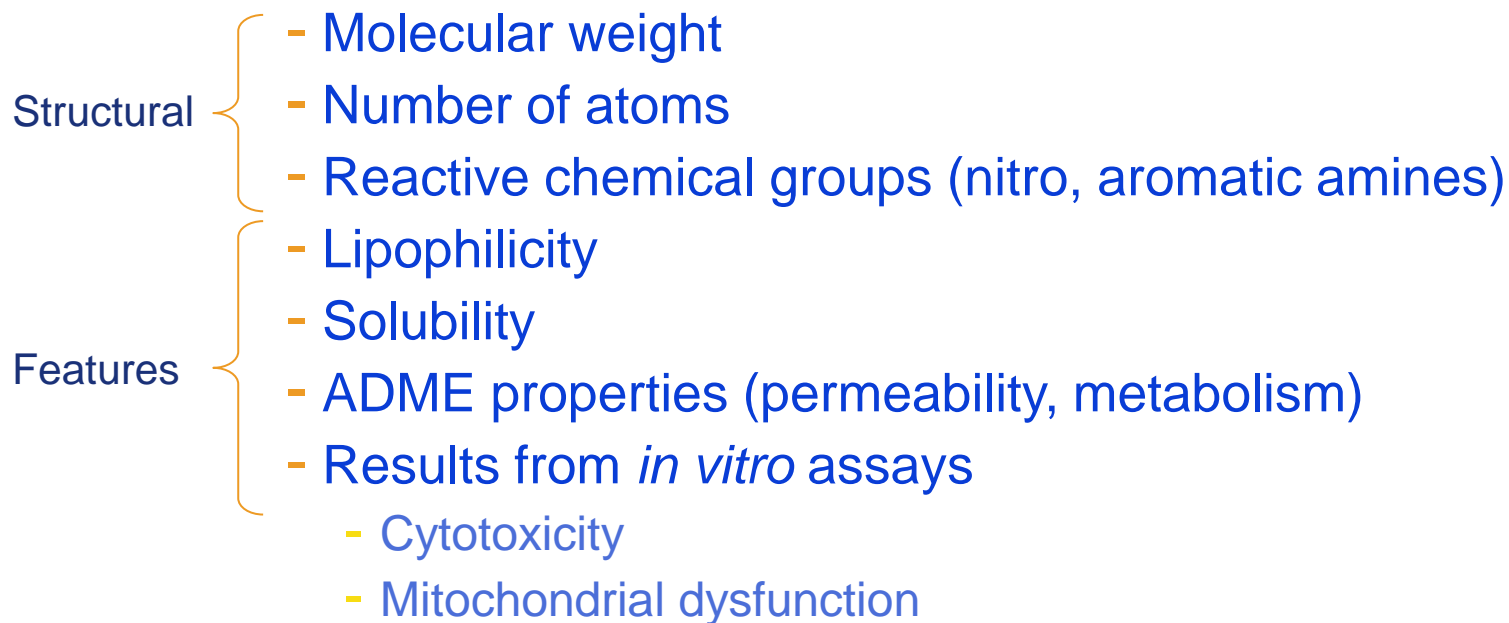
- Senior Principal Scientist, Molecular Informatics
Worldwide Medicinal Chemistry, Pfizer Inc., US
 - Identify safety risks early in the drug discovery process
 - Development of predictive *in silico* and *in vitro* models of *in vivo* toxicity
- Principal Scientist, Lhasa Limited, UK
 - Developed structure-activity relationships for various toxicological endpoints for inclusion in Derek for Windows (Derek Nexus)
- Senior Synthetic Chemist, AstraZeneca, UK
 - Design and synthesis in oncology and inflammation research

Presentation

- What is computational toxicology?
- Traditional applications e.g. prediction of mutagenicity
 - Challenges in modelling *in vitro* data
 - Assessing predictive performance
- The prediction of complex toxicological endpoints
- Development of robust safety screening paradigms
- Summary

What is Computational Toxicology?

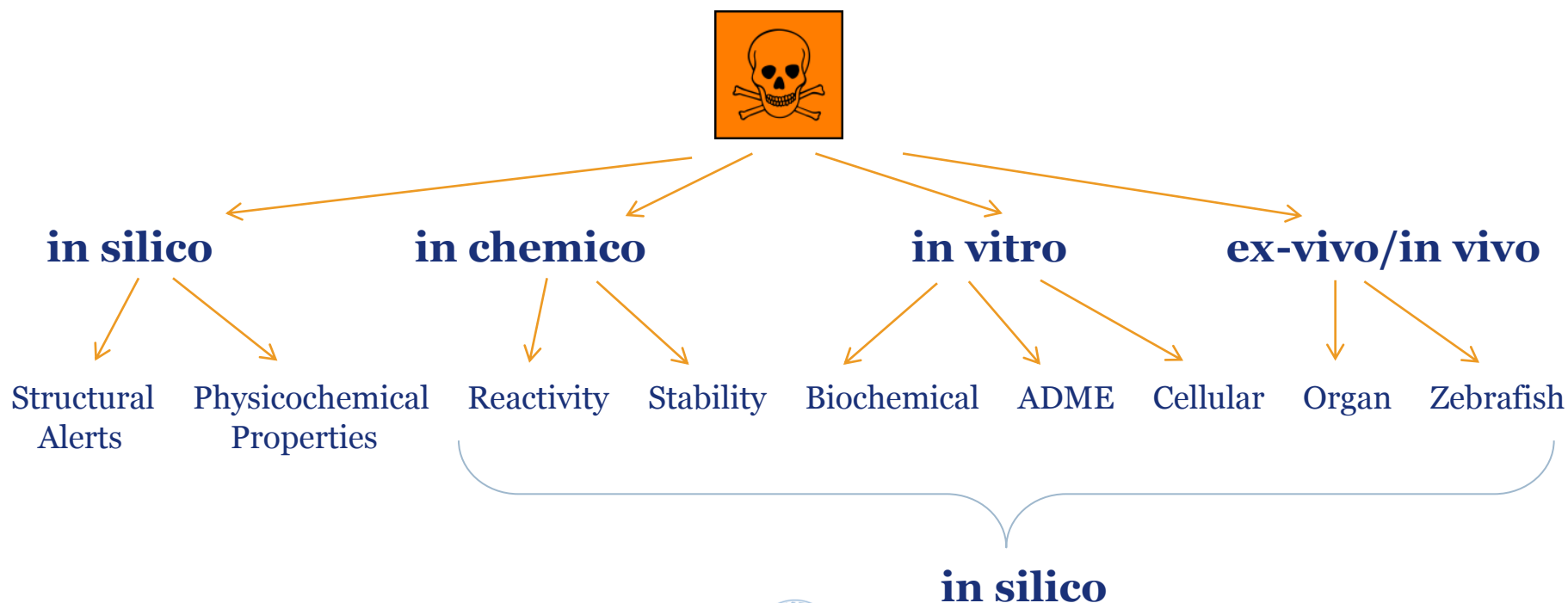
- Understanding the relationship between the properties of a compound and its toxicological activity



- Build predictive models for toxicity prediction

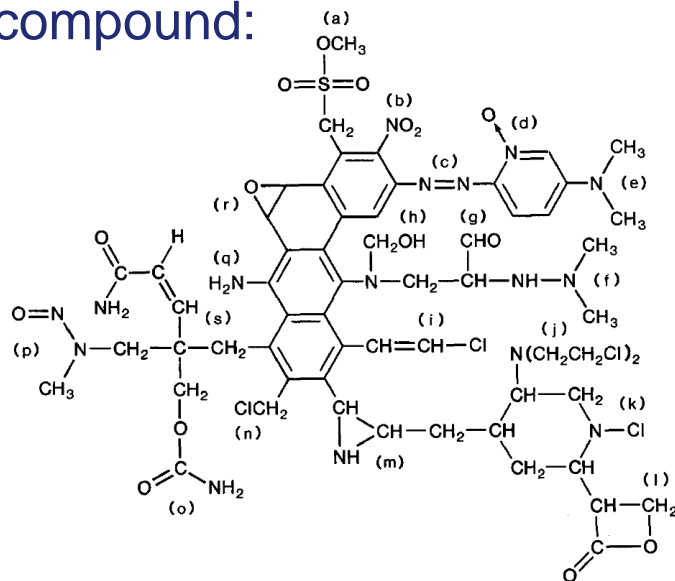
What is Computational Toxicology?

- Computational Toxicology is essential to improving the risk assessment process and identifying safety hazards across many industries



Modelling Mutagenic Activity

- Mutagenicity is hereditary DNA damage and can be observed in the Ames test
- In 1991 – Published a structural alert based electrophilic model compound:
- 2014 – ICH M7 guidelines for use of *in silico* models
- Predictions may be accepted *in lieu* of experimental data



INTERNATIONAL CONFERENCE ON HARMONISATION OF TECHNICAL
REQUIREMENTS FOR REGISTRATION OF PHARMACEUTICALS FOR HUMAN USE

ICH HARMONISED TRIPARTITE GUIDELINE

ASSESSMENT AND CONTROL OF DNA REACTIVE (MUTAGENIC)
IMPURITIES IN PHARMACEUTICALS TO LIMIT POTENTIAL
CARCINOGENIC RISK

M7

Current Step 4 version
dated 23 June 2014

Tennant and Ashby, Mut. Res. 1991, 257, 209-227



Modelling Mutagenic Activity

- Relatively simple mechanisms of activity enable the development of robust models

1. Guanine is an electron-rich base that can react with many electron-poor compounds

2. Bases may oxidize in the presence of reactive oxygen species

3. Intercalation can disrupt DNA synthesis, repair and enhance mechanisms 1&2

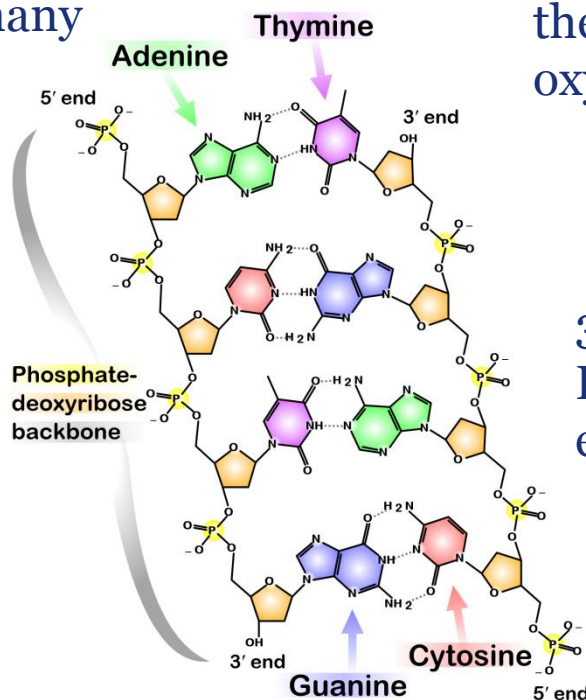
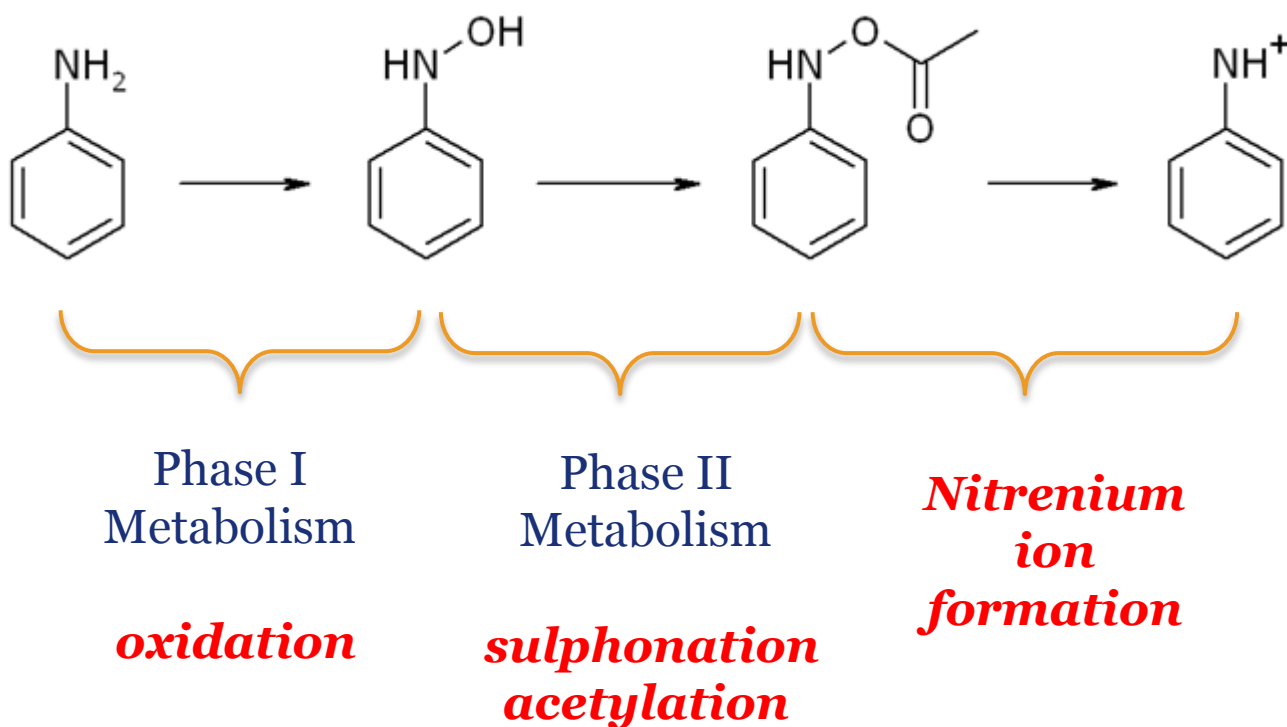


Fig: Madprime (wiki)

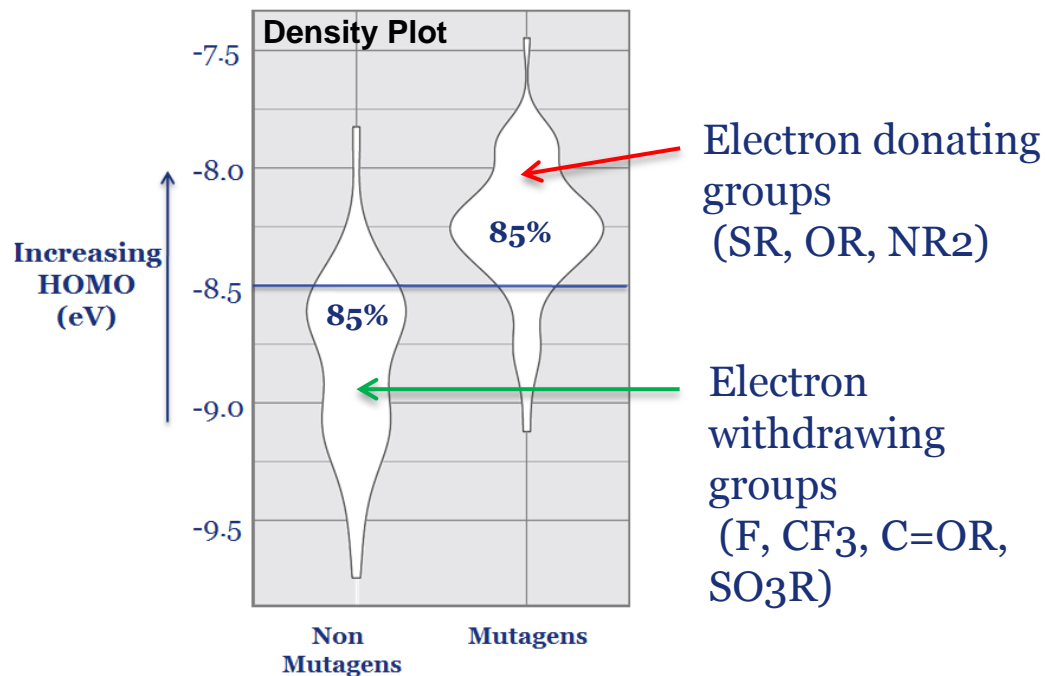
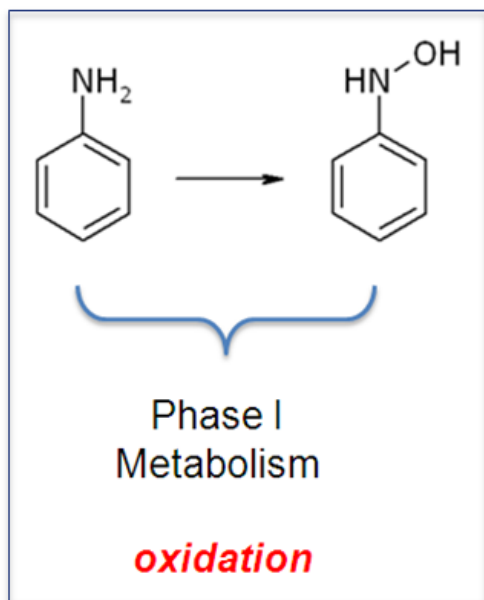
Aromatic Amine Mutagenicity

- General mechanism involves formation of reactive intermediates:



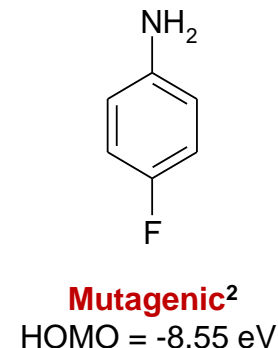
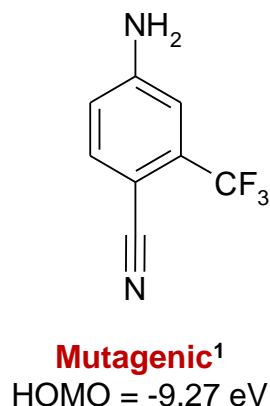
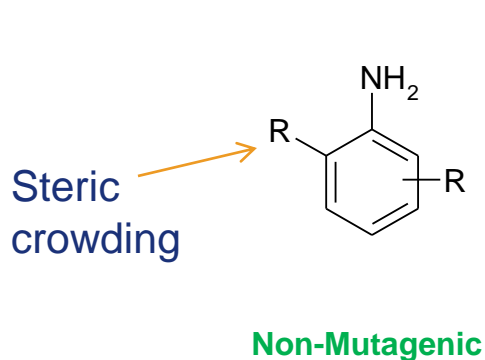
What Features Drive Aromatic Amine Mutagenicity

- Not all aromatic amines are mutagenic
- Mutagenicity correlates strongly with **H**ighest **O**ccupied **M**olecular **O**rbital (HOMO)
- HOMO reflects ability of amine to be oxidized by Cytochrome P450s



Mispredicted Compounds

- Despite >85% predictive performance, HOMO energy does not describe mutagenic activity of all chemical space



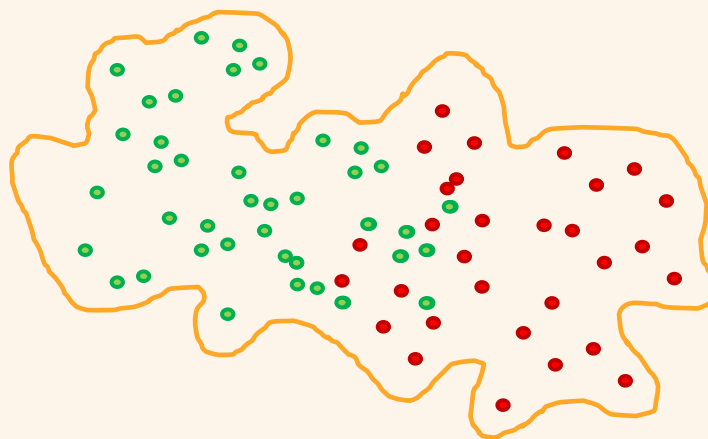
- These compounds require additional features within the model to describe their activity
- Should be investigated if we are to fully understand the predictive performance of the model

¹Sutter *et al.* Reg Toxicol Pharm, 2013, 67, 39-52

²Bentzien *et al.* J Chem Inf & Mod, 2010, 50, 274-297

The Applicability Domain of a Model

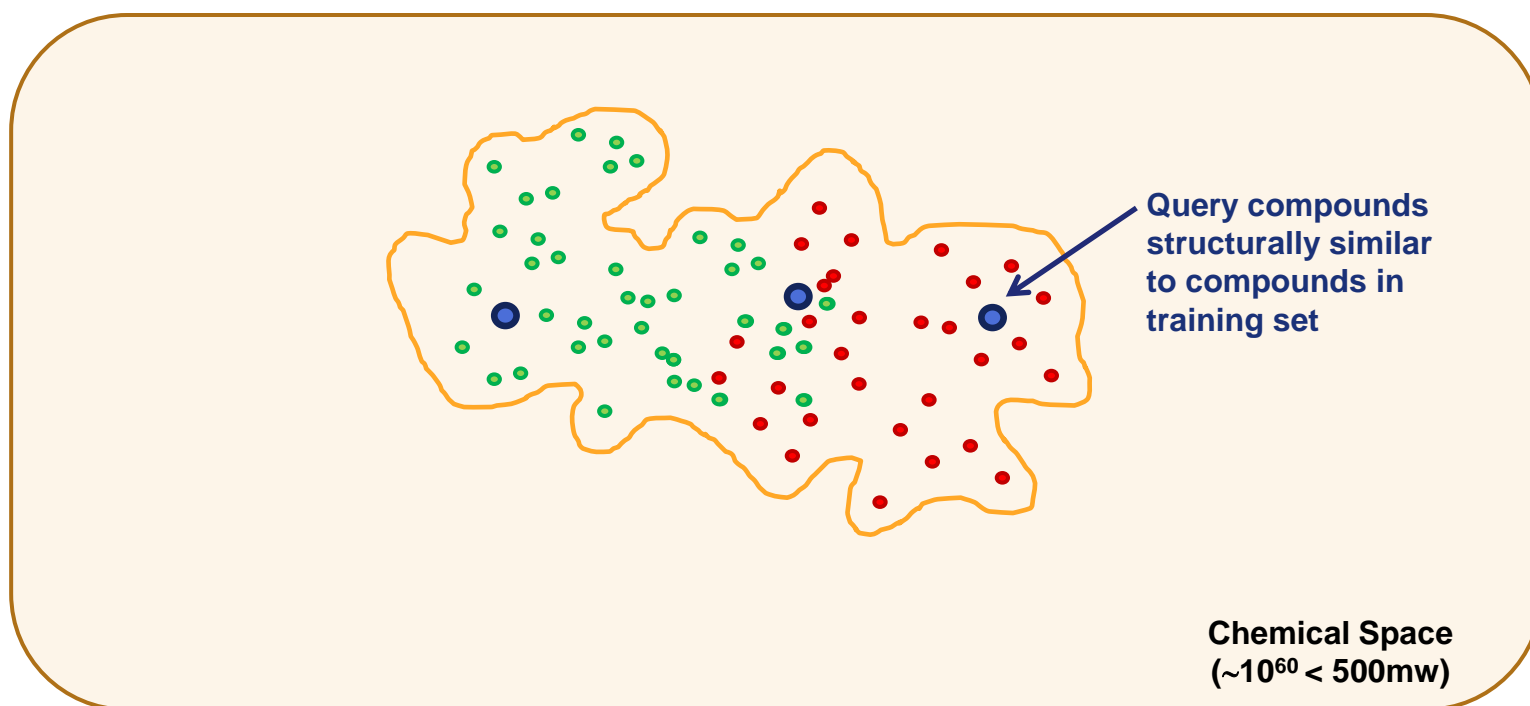
- Describes the chemical space upon which the model has been created



Chemical Space
($\sim 10^{60} < 500mw$)

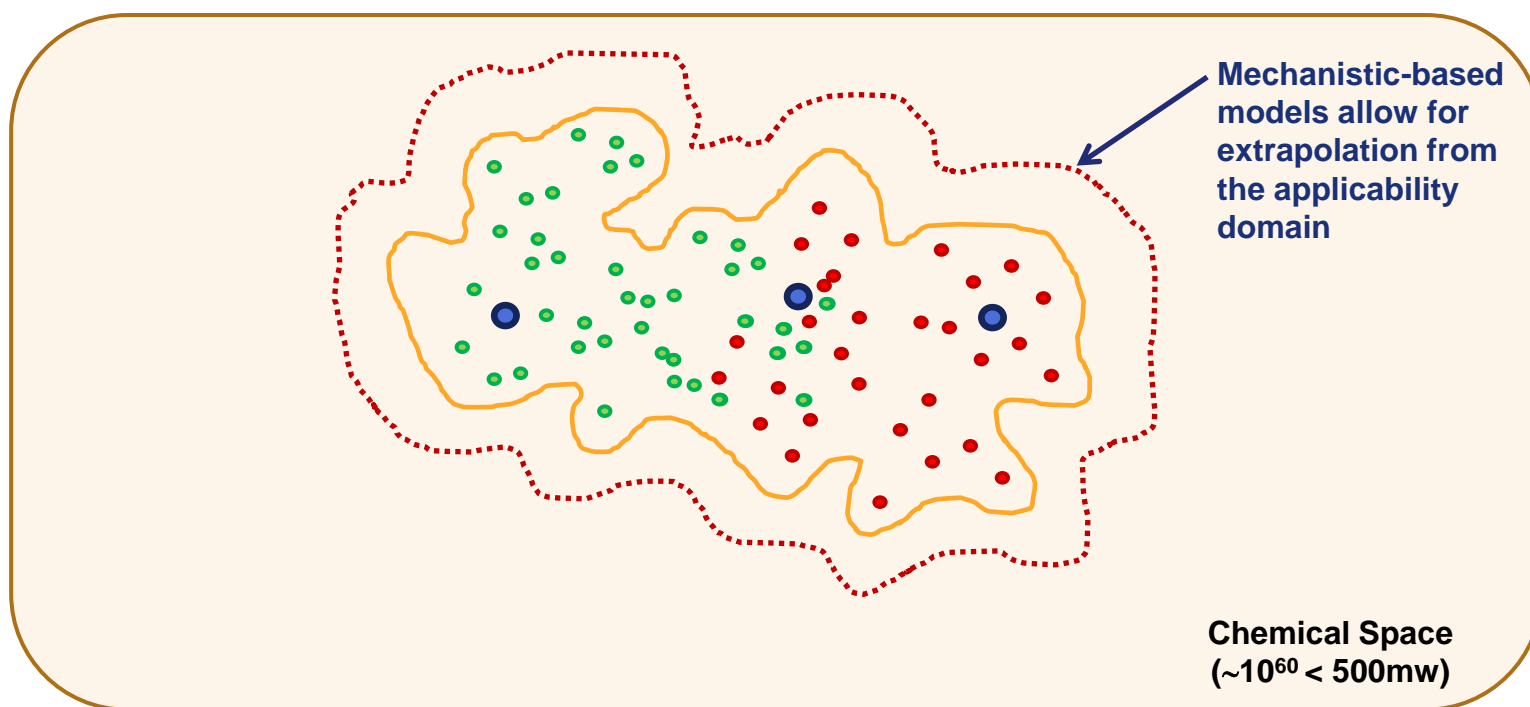
The Applicability Domain of a Model

- Describes the chemical space upon which the model has been created



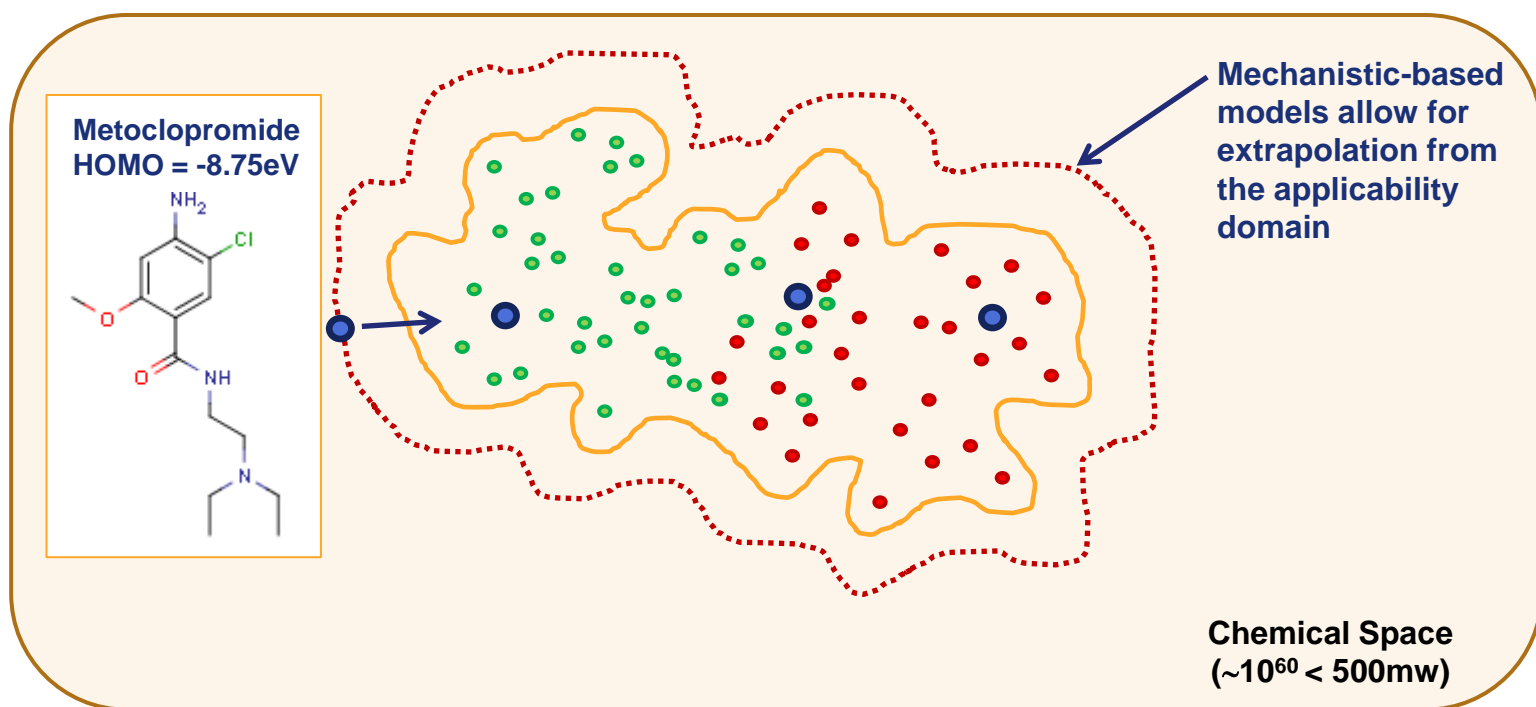
The Applicability Domain of a Model

- Describes the chemical space upon which the model has been created



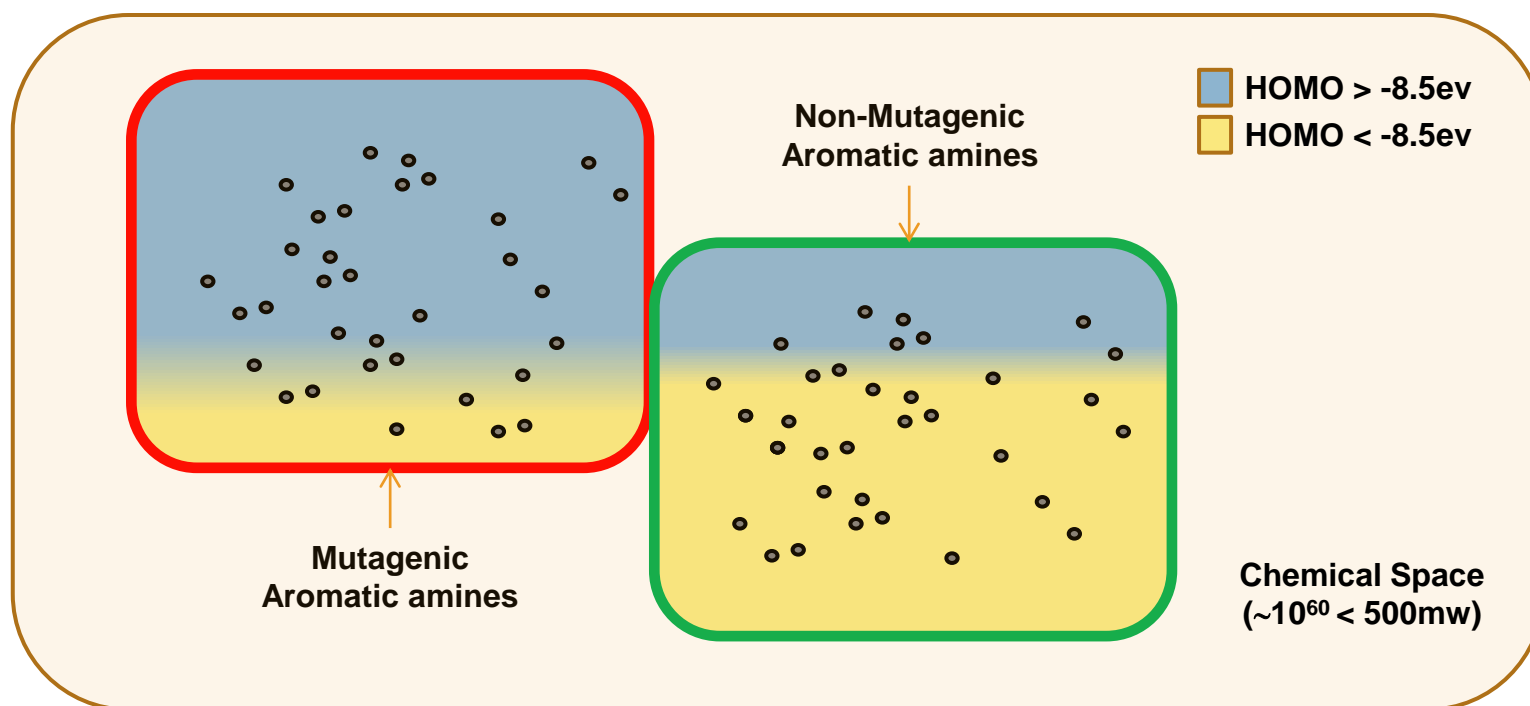
The Applicability Domain of a Model

- Describes the chemical space upon which the model has been created



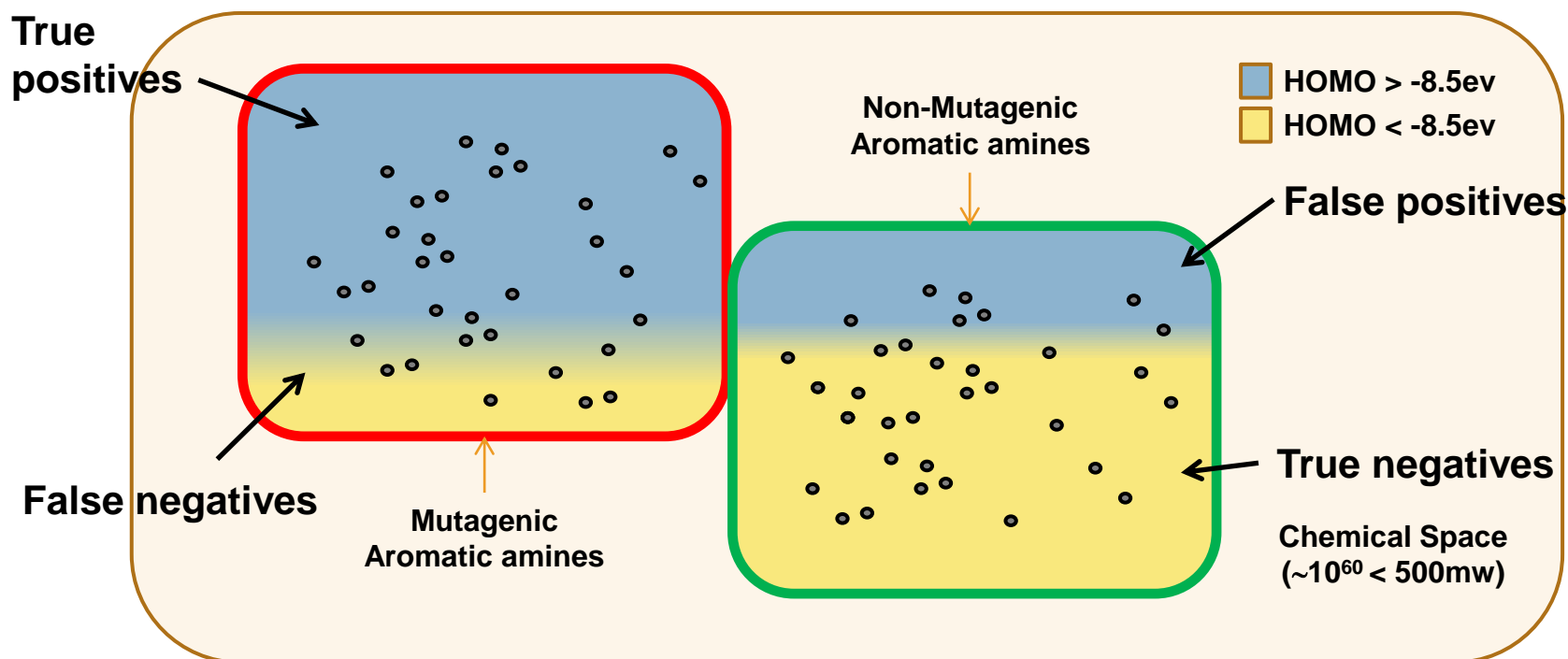
Model Validation

- Hypothetical pictorial view of aromatic amine dataset:



Model Validation

- Hypothetical pictorial view of aromatic amine dataset:



Truth Tables

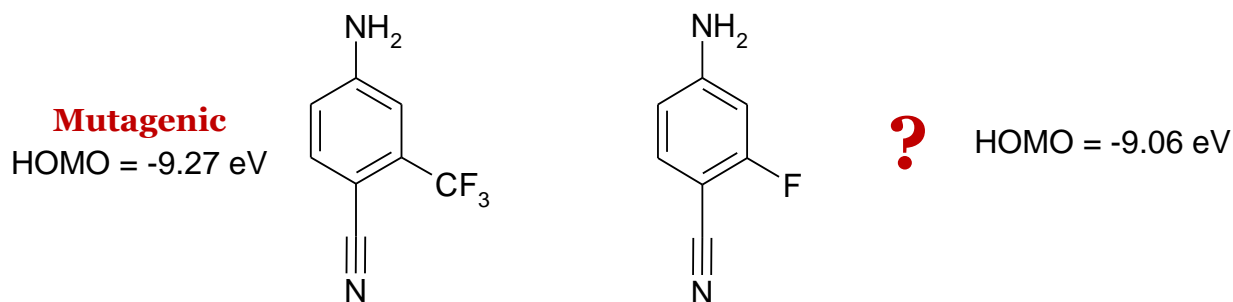
	Ames POS	Ames NEG
Model POS	True POS	False POS
Model NEG	False NEG	True NEG

- Statistics help in assessing how good the model is at predicting the training set (applicability domain)
- Real value is highlighting mispredicted compounds!



Understanding Mispredicted Compounds

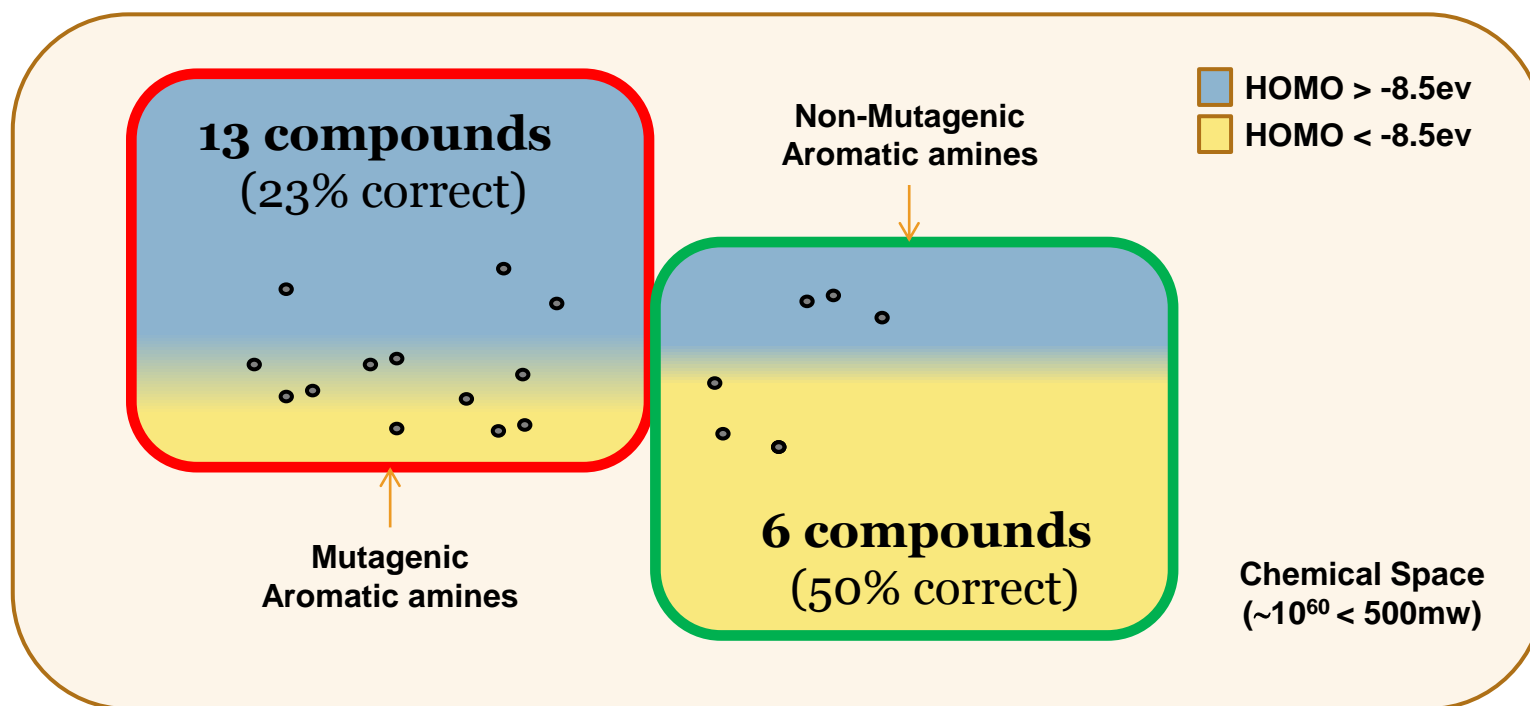
- Outliers, false positives and false-negatives highlight *potential* knowledge gaps within models



- Focused testing in the future must prioritize these areas
- Understanding the applicability of a model is essential to understanding relevance of external validation statistics

Model Validation

- Performance is dependent upon chemical space of test set
- e.g. test set of 19 compounds



The Real Value of Truth Tables

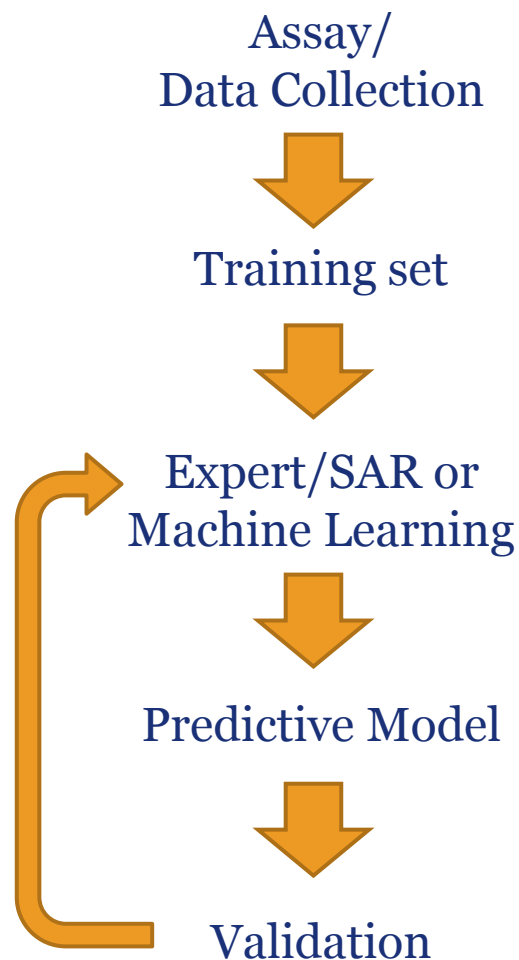
	Ames POS	Ames NEG
HOMO POS	Correlative	1) Steric crowding 2) Cyp-deactivation 3) Diverted metabolism
HOMO NEG	1) Another mechanism 2) Impurity?	useful in SAR determination

How Useful is My *in silico* Model?

- Performance is in the eye of the test set holder
 - Performance depends upon chemical space of the test set
- Transparency is key
 - How was your prediction derived?
 - Do you have access to the training set?
 - Is the model based on a mechanism that enables confident extrapolation outside of the applicability domain?
 - Aware of the limitations of the model?

in silico Summary

- This process largely works for toxicological endpoints based on structure-related mechanisms of action, e.g.
 - Mutagenicity
 - Skin sensitization
 - Genotoxic carcinogenicity
- Improvements can be made through investigating toxicological knowledge gaps within our datasets

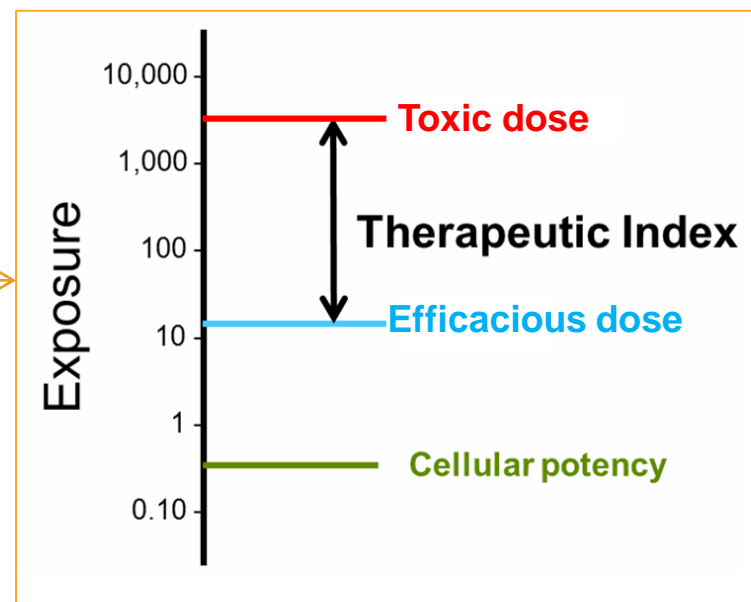


Can we apply computational toxicology to predict *in vivo* toxicity?



Can We Predict *in vivo* Toxicity?

- Yes - all compounds are toxic
- Calculated human LD50 values:¹
 - Water – 6 liters
 - Caffeine – 118 coffees
 - Alcohol – 13 shots
- Focus on toxicity observed at therapeutically relevant levels



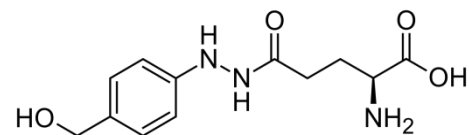
¹www.compoundchem.com/2014/07/27/lethaldoses

What Drives *in vivo* Toxicity?

- Exposure



- Toxic Potential

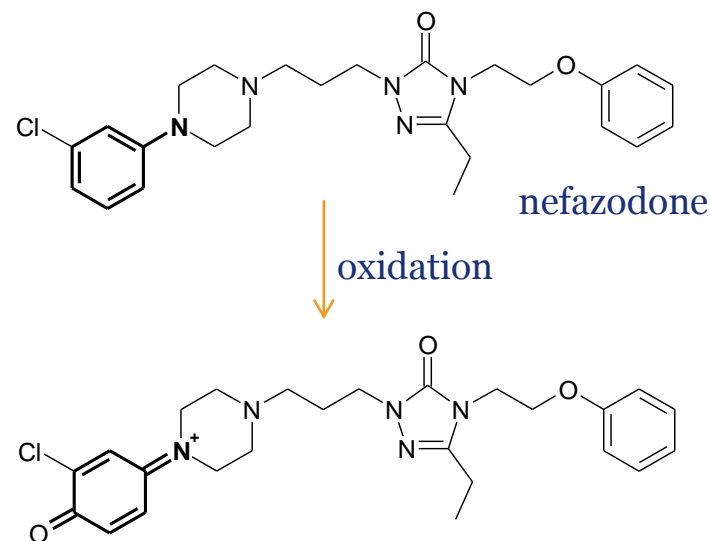


agaritine



Example - Nefazodone

- Potent 5-HT_{2A} receptor antagonist and antidepressant
- Withdrawn in 2003 owing to very rare, but severe, liver toxicity
- Has multiple safety liabilities
 - Contains structural alert (aniline)¹
 - Metabolic liabilities²
 - Inhibitor bile-salt export pump³
 - Cytotoxic⁴
 - Mitochondrial dysfunction⁴
 - High dose: >200mg/day

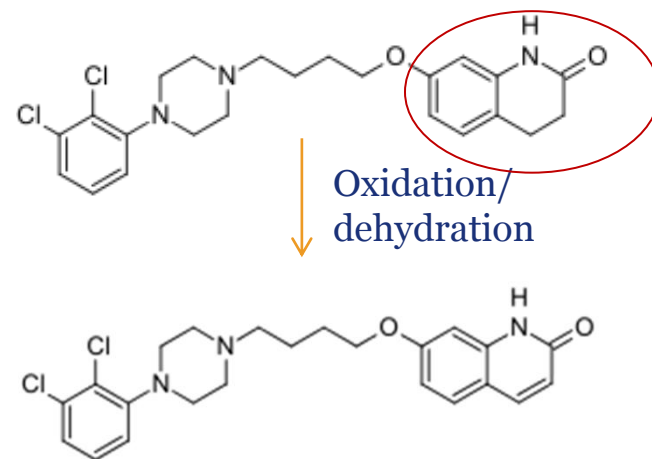


Refs

1. Stepan *et al.*, Chem. Res. Toxic., 2011, 24, 1345-1410.
2. Kalgutkar *et al.*, Drug Metab. Disp., 2005, 33, 243-253
3. Kostrubsky *et al.*, Toxicol. Sci., 2006, 90, 451-459
4. Dykens *et al.*, Toxicol. Sci., 2008, 103, 335-345.

What About Aripiprazole?

- Structurally similar, yet successfully marketed drug
 - **No** reports of acute hepatotoxicity
- Has multiple liabilities
 - Contains structural alert (aniline) ¹
 - Metabolic liabilities²
 - Cytotoxic and lysosomotropic³
 - *Low dose: 10-20 mg/day*



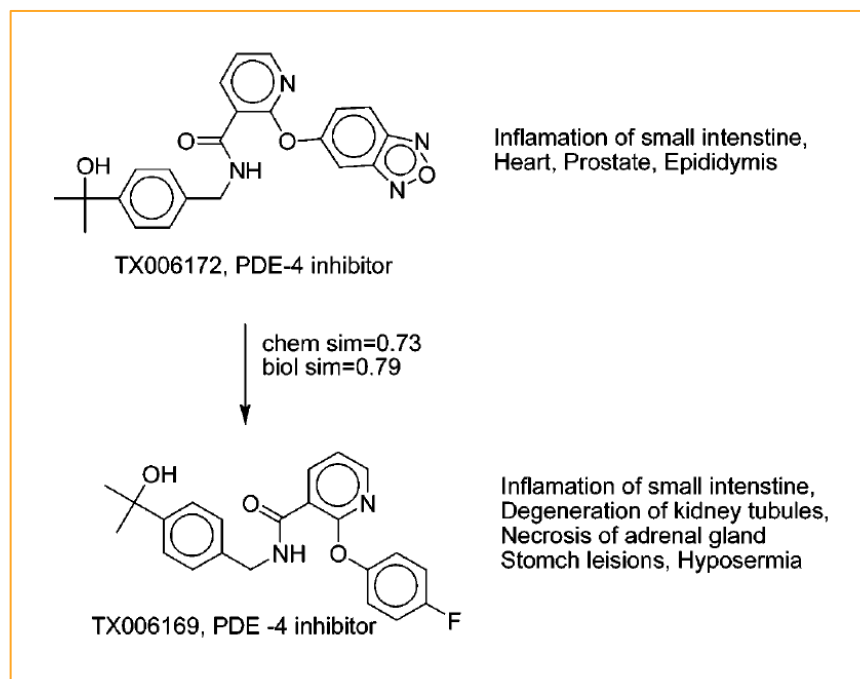
- Why is aripiprazole not hepatotoxic?
 - Related to different pharmacological profile?
 - Different metabolic profile?
 - Low dose?

Refs

1. Stepan *et al.*, Chem. Res. Toxic., 2011, 24, 1345-1410.
2. Bauman *et al.*, Drug Metab. Disp., 2008, 36, 1016-1029.
3. Nadanaciva *et al.*, Toxicol. in Vitro, 2011, 25, 715-723.

Do Similar Compounds Have Similar Toxicological Profiles?

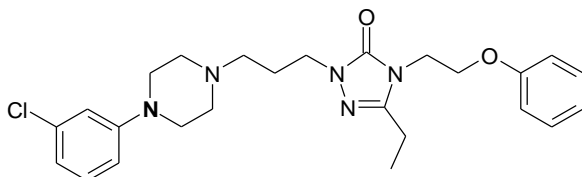
- Similar compounds with a similar *in vitro* toxicity profile may not express similar *in vivo* findings...



Shah and Greene, Chem Res &Tox, 2014, 27, 86-98

Challenges of Modelling *in vivo* Toxicology Data

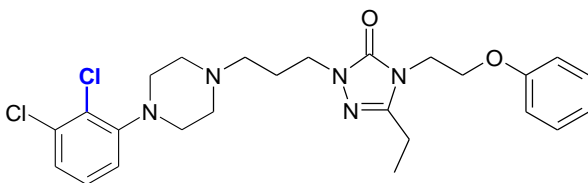
- Nefazodone



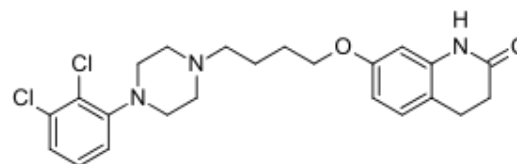
'hepatotoxic'

Dose = 200-400 mg/day

- Hepatotoxic at 20mg/day?
- Can we predict:



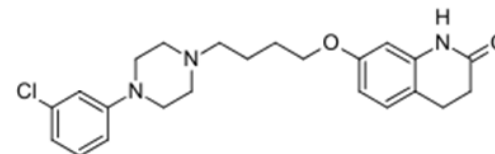
- Aripiprazole



'non-hepatotoxic'

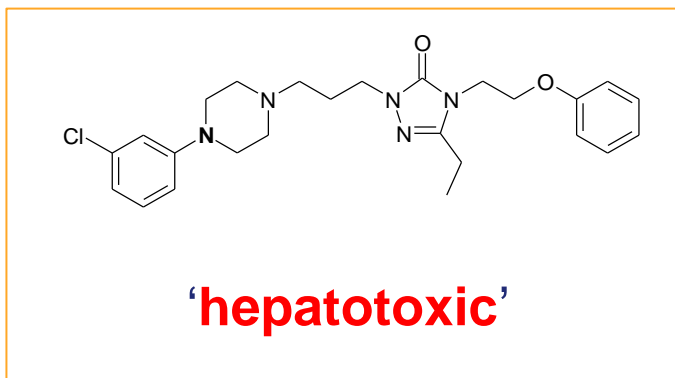
Dose = 10-30 mg/day

- Hepatotoxic at 200 mg/day?
- Can we predict:



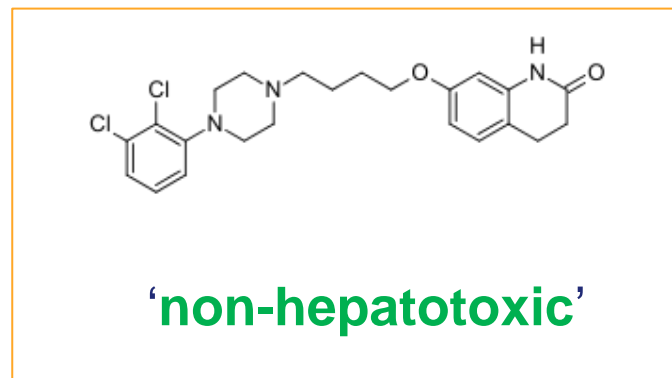
Challenges of Validating *in silico/in vitro* Models

- Nefazodone



- Aniline structural alert for hepatotoxicity?
 - True positive ✓
- Cytotoxicity assay
 - True positive ✓

- Aripiprazole



- Aniline structural alert for hepatotoxicity?
 - False positive ✗
- Cytotoxicity assay
 - False positive ✗

Validating Models of *in vivo* Data

- Performance is in the eye of the test set holder
- For *in vivo* endpoints, performance is dependent upon:
 - Appropriate annotation of toxicological data
 - Understanding of exposure and pharmacokinetic profile

	<i>in vivo</i> toxic	<i>in vivo</i> clean
Model/assay positive	True POS	False POS
Model/assay negative	False NEG	True NEG



Validating Models of *in vivo* Data

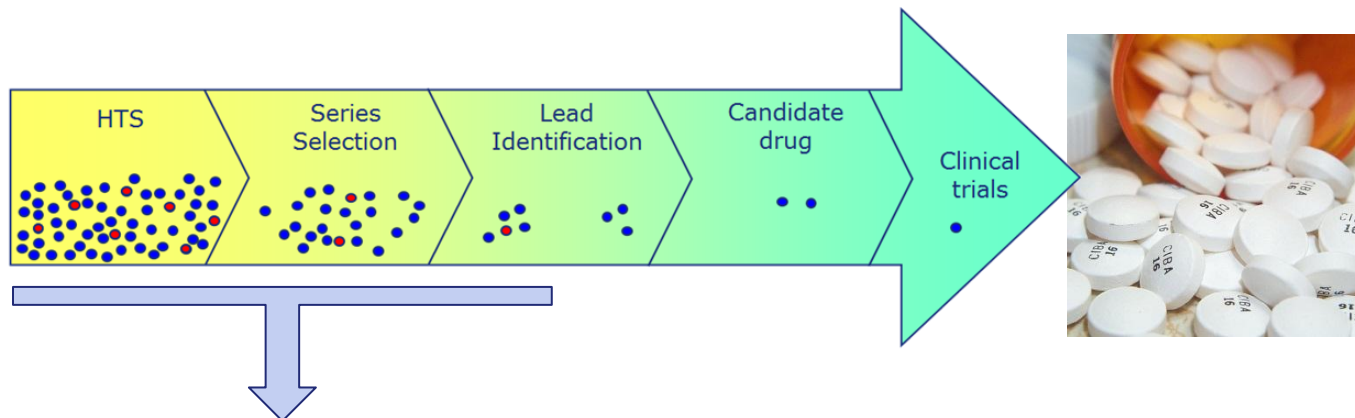
- Performance is in the eye of the test set holder
- For *in vivo* endpoints, performance is dependent upon:
 - Appropriate annotation of toxicological data
 - Understanding of exposure and pharmacokinetic profile

	<i>in vivo</i> toxic	<i>in vivo</i> clean
Model/assay positive	Correlative (not causative)	Mitigation through ADME?
Model/assay negative	Different mechanism?	True NEG



The Role of Early Screening Paradigms

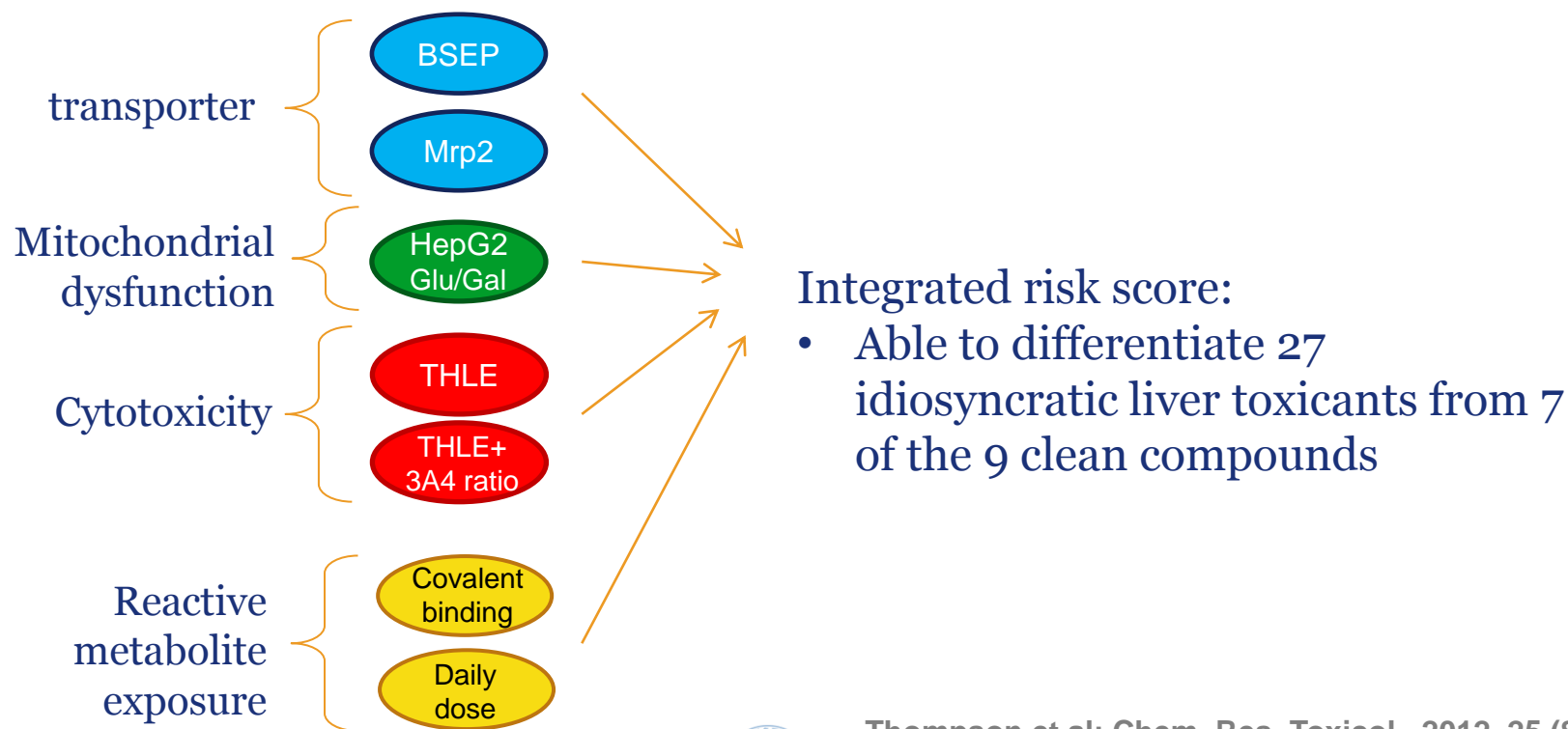
- *in vivo* Toxicology is complex to predict



- Identify potential risk early using *in silico* and *in vitro* models
- Recognize that *in vitro-in vivo* translation may not be possible without in-depth, costly, exposure-related studies

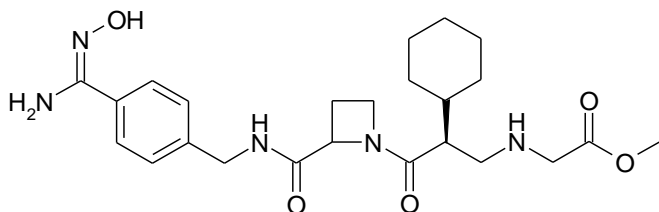
Early Screening Cascades: Example

- Platform of assays developed to identify risk of idiosyncratic drug reactions for 36 compounds with liver toxicity profiles



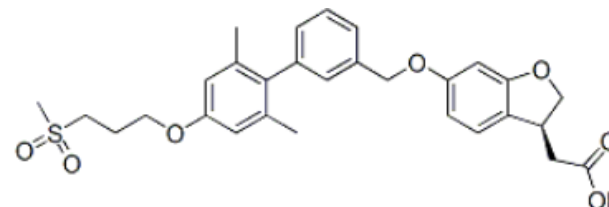
Recently Discontinued Drugs

- Withdrawn owing to liver safety signals in Phase III



Ximelagatran

- Thrombin Inhibitor
- Withdrawn 2006

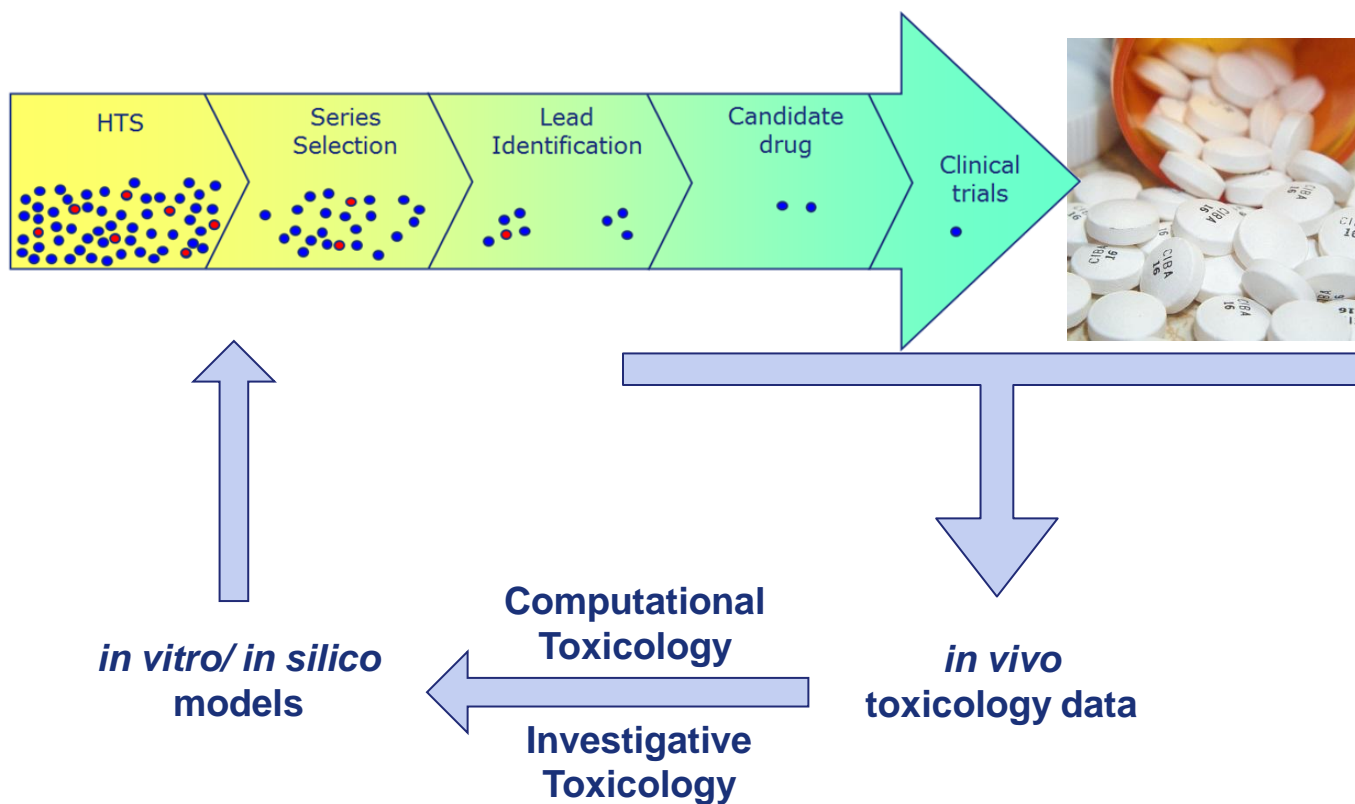


Fasiglifam

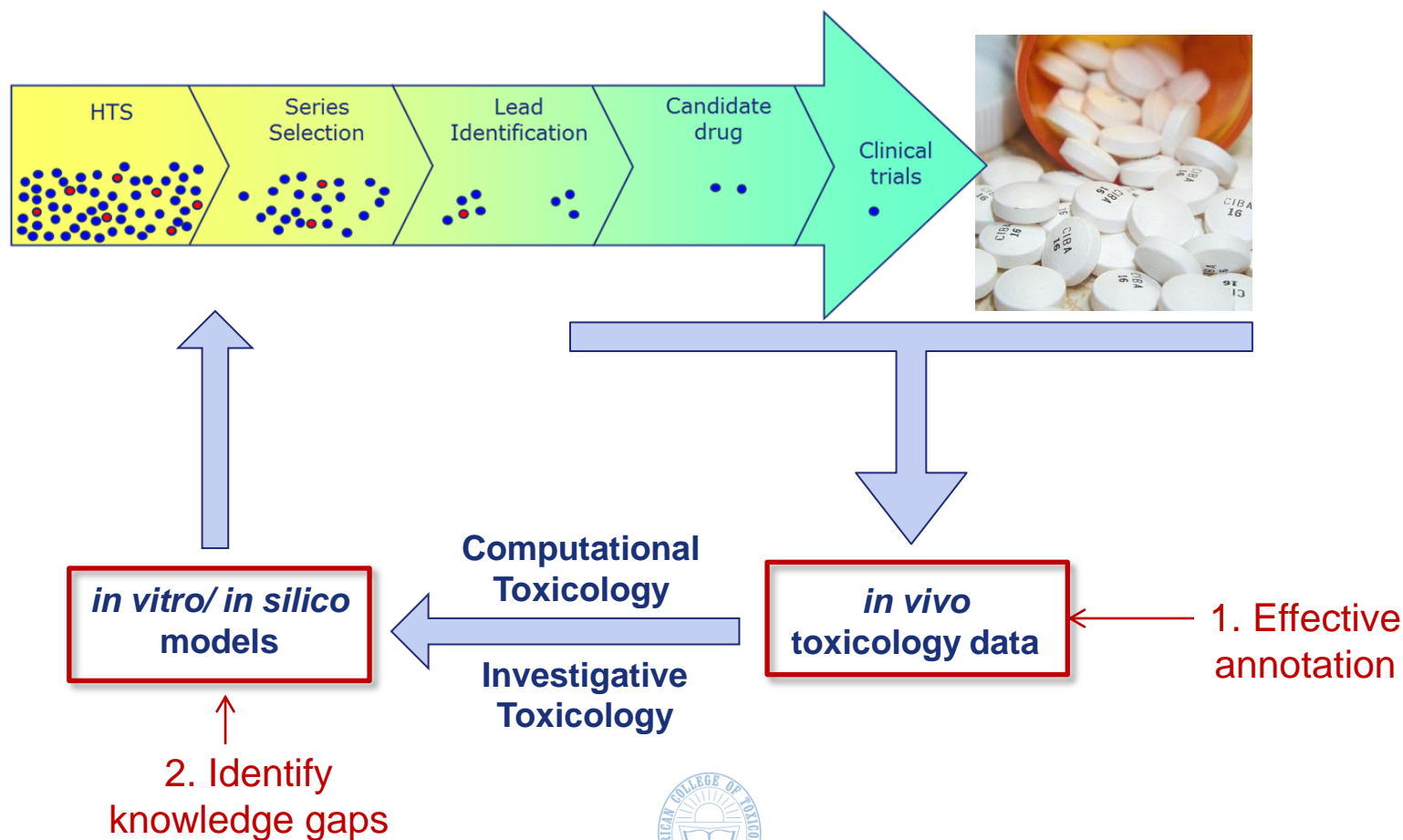
- GPR40 agonist
- Withdrawn 2013

- Why were these safety liabilities not caught early?

Improving *in vivo* Toxicity Prediction



Improving *in vivo* Toxicity Prediction



Features That Are Predictive of *in vivo* Toxicity

- Study 1: 207 preclinical candidates investigated
 - Compounds were annotated against the observation of any *in vivo* toxicity findings at 10 μ M (total plasma exposure)
 - Odds of toxicity established for various physicochemical properties

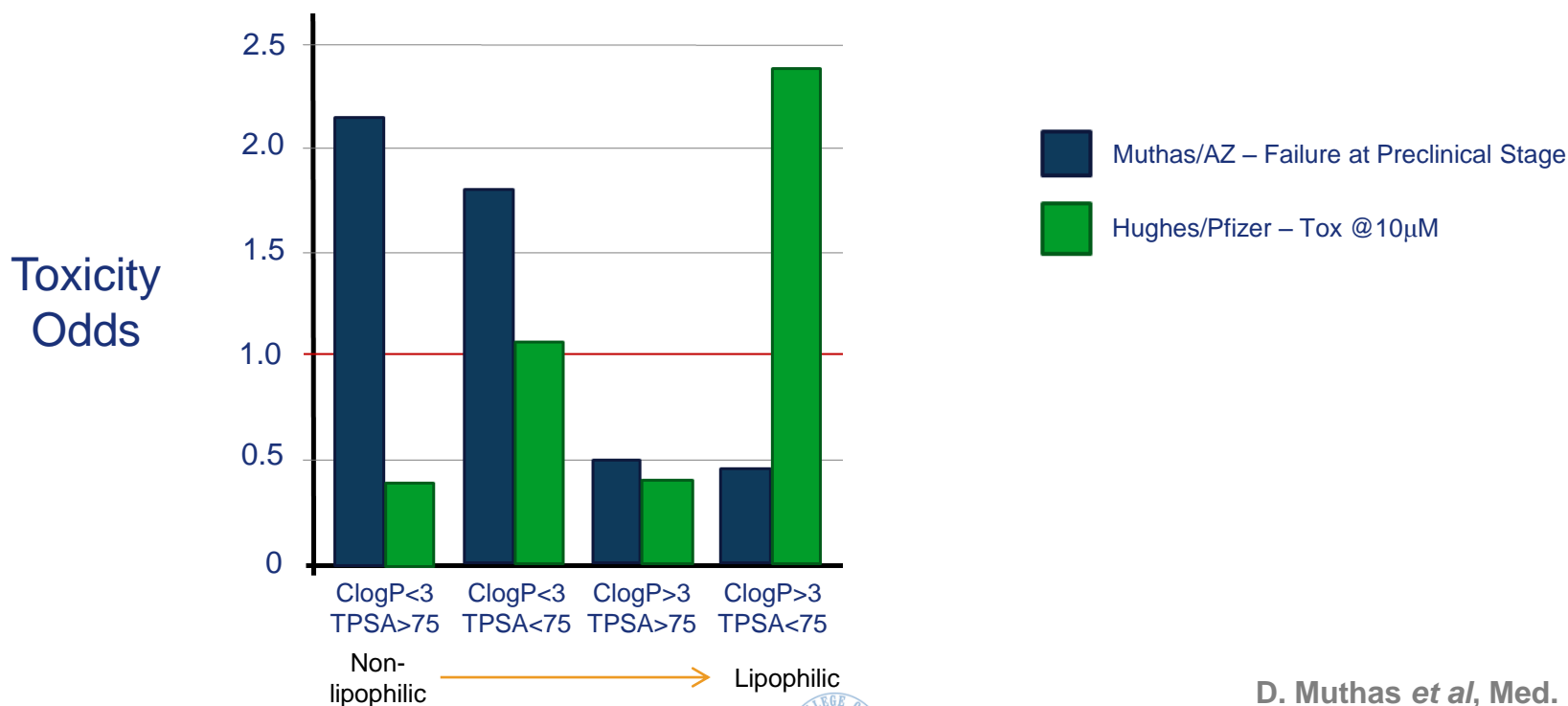
Tox@10 μ M	TPSA>75	TPSA<75
ClogP<3	0.39 (57)	1.08 (27)
ClogP>3	0.41 (38)	2.4 (85)

TPSA and ClogP are calculated measures of lipophilicity

⇒ Study conclusions: likelihood of toxicity increases with lipophilicity

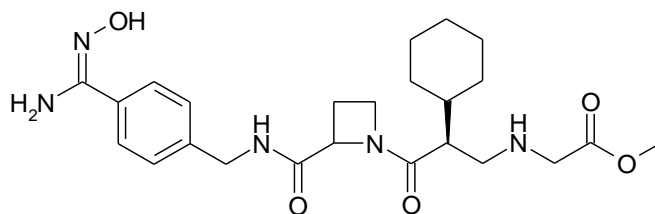
Features That Are Predictive of *in vivo* Toxicity

- Study 2: Odds of failure in preclinical/Phase 1 studies
 - Authors found that compounds were more likely to fail due to toxicity in non-lipophilic space:



D. Muthas *et al*, Med. Chem. Commun., 2013, 4, 1058

Which Thresholds Can You Believe In?



Ximelagatran

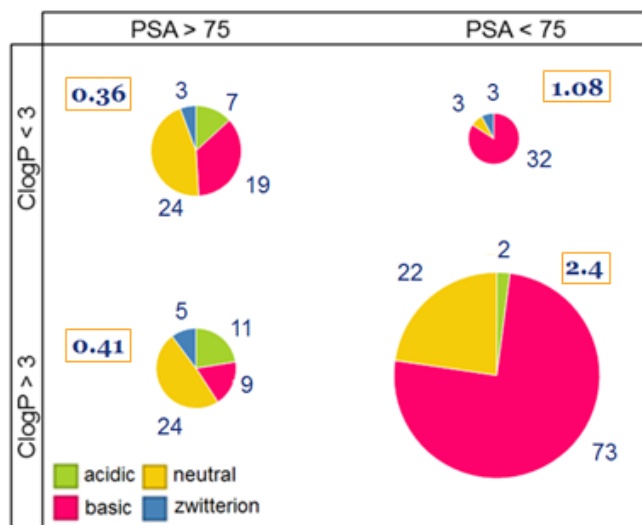
- ClogP = **1.8**
 - PSA = **144**
- ✓ - Hughes
✗ - Muthas

- All models are useful... but *only* for a portion compounds in the training set
- It is essential to understand in which chemical space the model works and where it doesn't



Understanding The Applicability Domain

- Hughes training set is dominated by lipophilic basic drugs



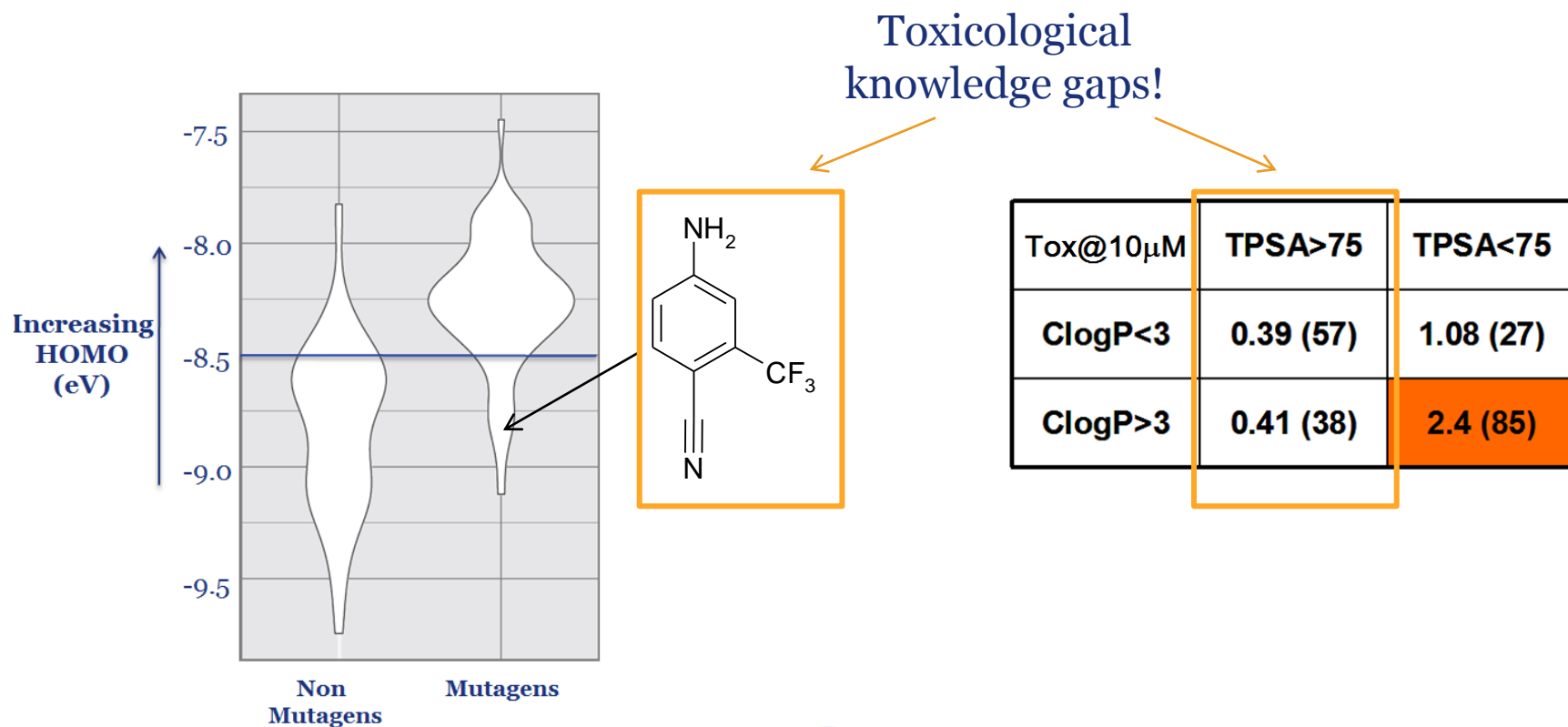
- Lipophilic basic drugs cause general toxicity, e.g. through lysosomal dysfunction and disruption of ion channels
- What factors drive toxicity of neutral +acidic compounds?



Hughes *et al*, Bioorg&Med Chem Lett, 2008,18, 4872–4875

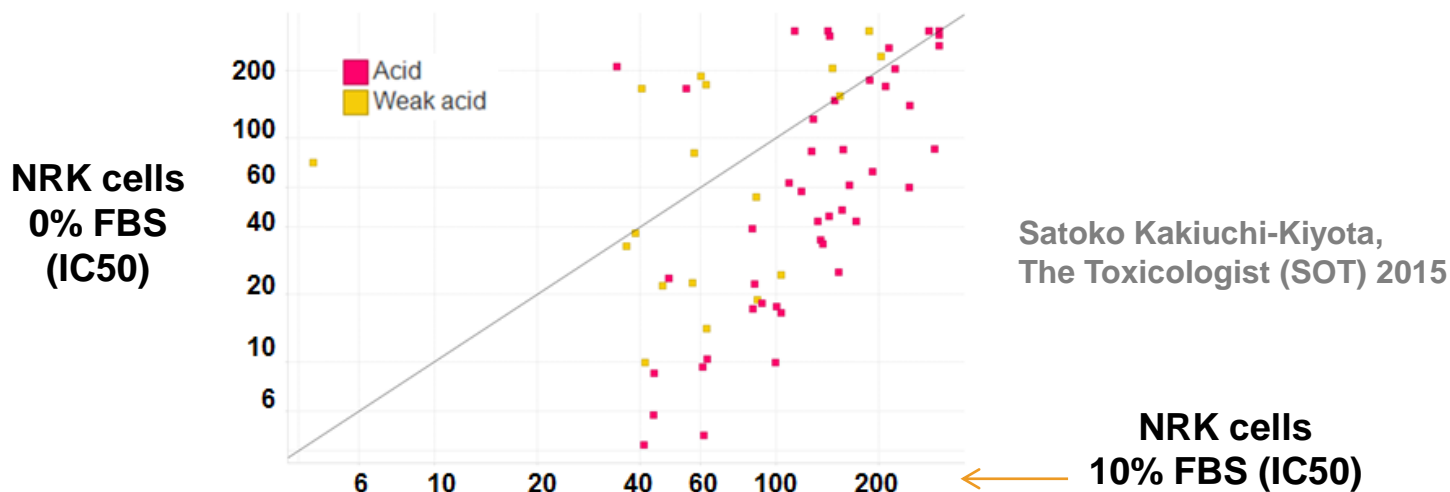
Identifying Toxicological Knowledge Gaps

- Applicability enables the identification of knowledge gaps in your assay or model



Toxicity and Acidic Compounds

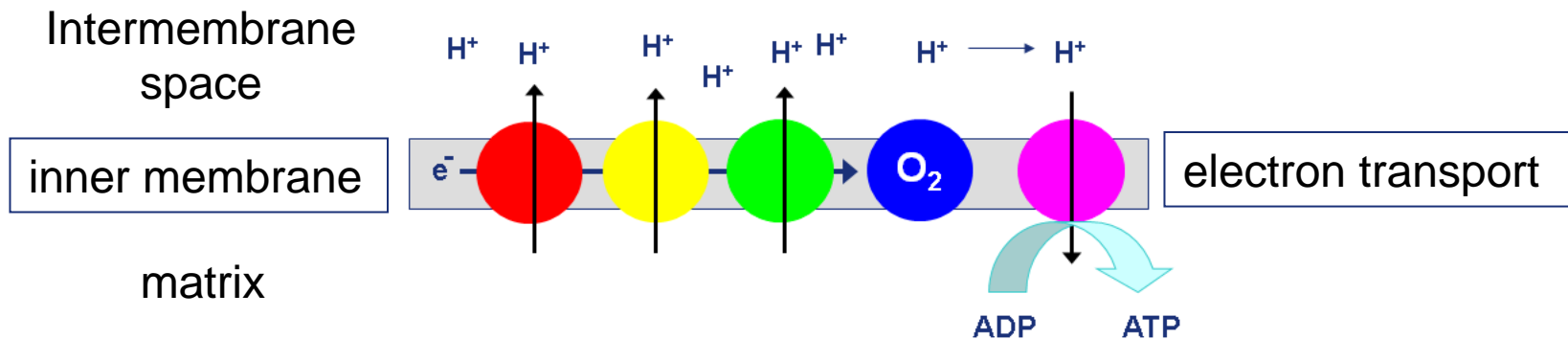
- Acidic compounds tend to have low cytotoxicity in cytotoxicity assays
- Acids tend to be highly protein bound
- Is toxicity mitigated by high protein binding to assay serum?



- The impact of this result is not clear without an assessment of the toxicological and ADME profile of compounds in the dataset

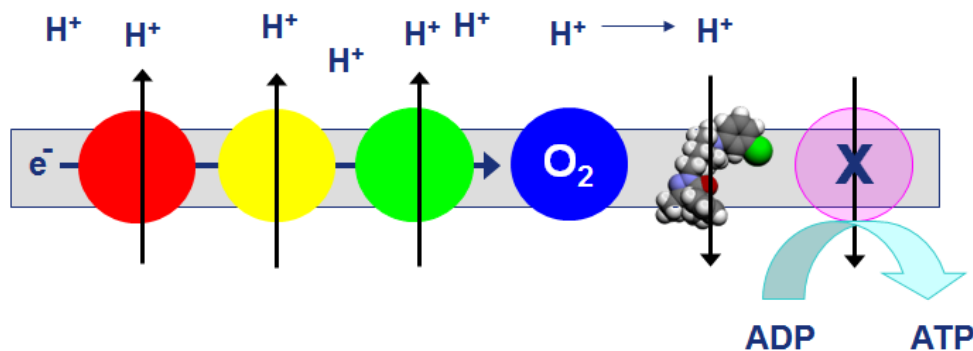
Disruption of Oxidative Phosphorylation

- Oxidative Phosphorylation occurs in mitochondria
 - Provides cellular energy (ATP)
 - Disruption linked to idiosyncratic organ toxicity

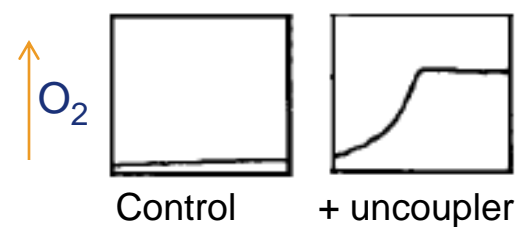


Mechanism of Uncoupling

- Modelling suggests lipophilicity and acidity is essential for protonophoric uncoupling
 - The uncoupler needs to reside in the membrane (be lipophilic)
 - Be able to shuttle protons across the membrane (have an acidic group)



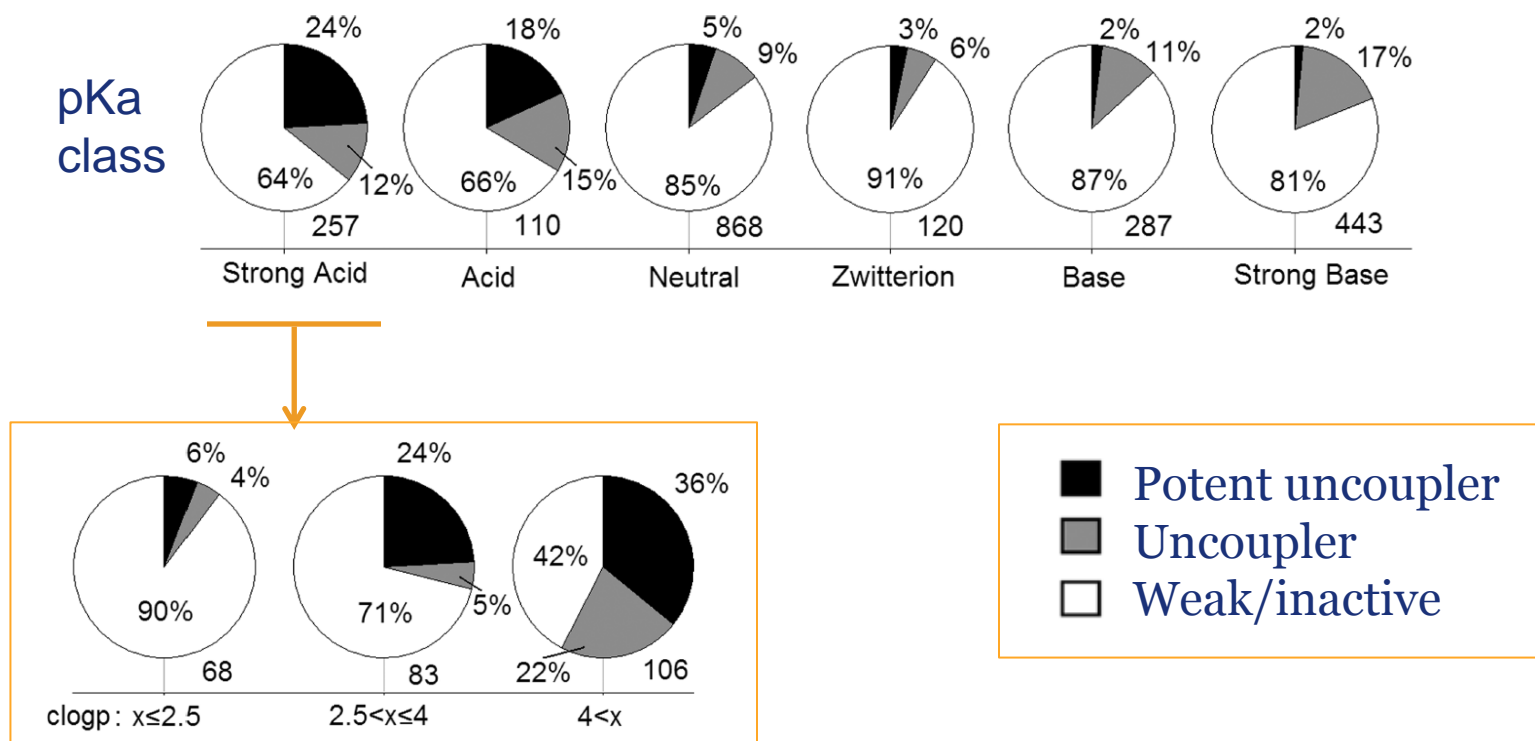
Detect uncoupling by measuring oxygen consumption



Hynes *et al*, Toxicol. Sci., 2006,92, 186-200

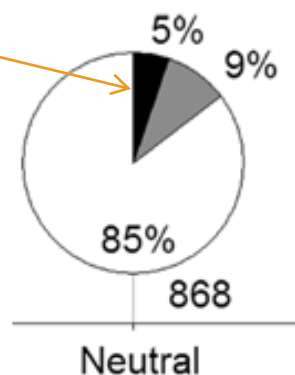
SAR Studies on 2000+ Compounds

- Uncoupling is highly dependent upon lipophilicity and acidity



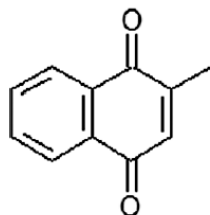
Outliers and Falsely-Predicted Compounds

- “False Negatives”

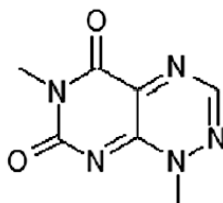


- Redox Cyclers

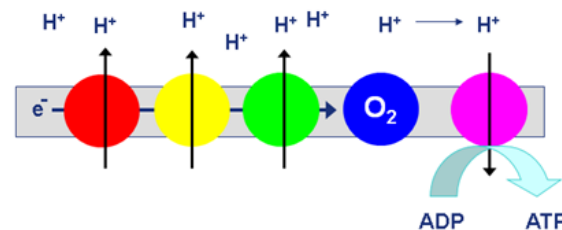
- Neutral, non-lipophilic compounds



Menadione

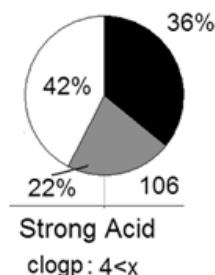


Toxoflavin

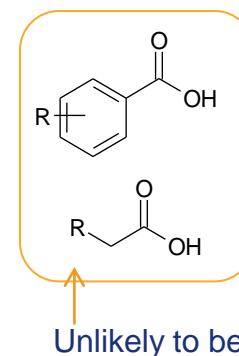
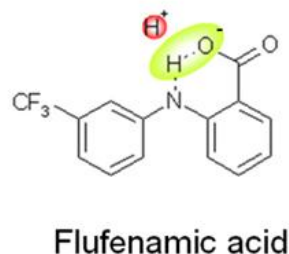
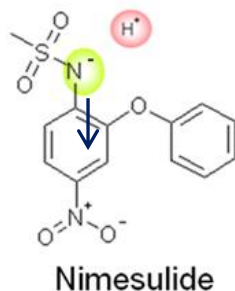


Outliers and Falsely-Predicted Compounds

- 42% of lipophilic, acidic compounds were not uncouplers



- Potent uncoupling requires stabilization of the negative charge:

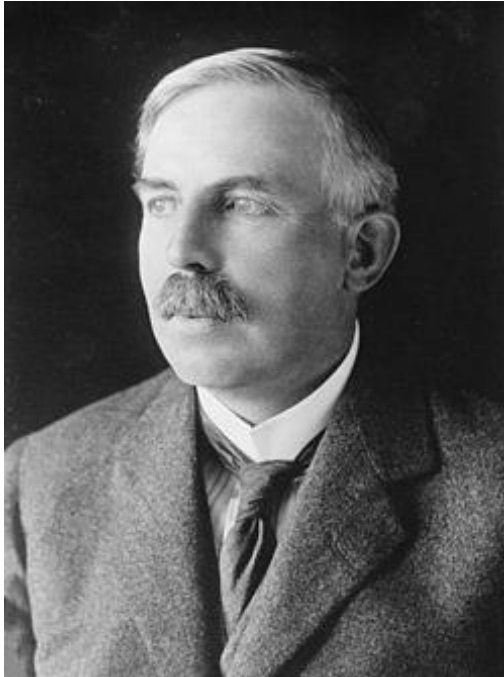


Key Points Summary

- Computational Toxicology is essential to developing effective risk assessment strategies and models
- *in vivo* Toxicology is complex and we need to understand what our assays and models are telling us
- Recognize the utility and limitations of current predictive tools
 - *in silico*
 - *in vitro*
- Move beyond broad annotations of *in vivo* toxicology data and include exposure assessment, if possible



Identify and address toxicological knowledge gaps



Ernest Rutherford

- If your experiment needs statistics, you ought to have done a better experiment

Acknowledgements

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 - Serum Free Assay
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Thank you for your participation in the American College of Toxicology Webinar!

*We hope to see you at the 36th Annual Meeting of the
American College of Toxicology
Red Rock Resort, Summerlin, Nevada,
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