



optibrium

# Confidently Targeting High Quality Hits from High-Throughput Screening

ACS Spring National Meeting. CINF, April 3<sup>rd</sup> 2017

Matthew Segall, Tamsin Mansley, Peter Hunt, Edmund Champness

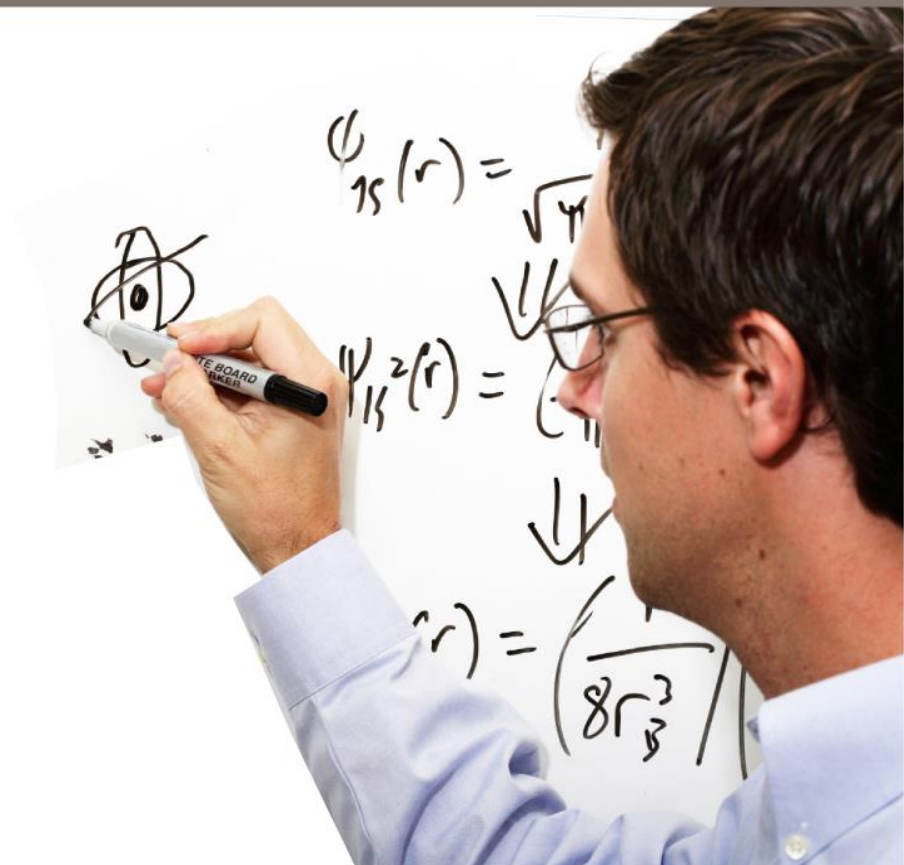
[matt.segall@optibrium.com](mailto:matt.segall@optibrium.com)

# Overview

---

- Goals of high-throughput screening (HTS) triage
- Mapping the chemical space of activity
- Understanding the activity landscape
- Targeting high quality hit series
- Conclusions

# Goals of HTS Triage

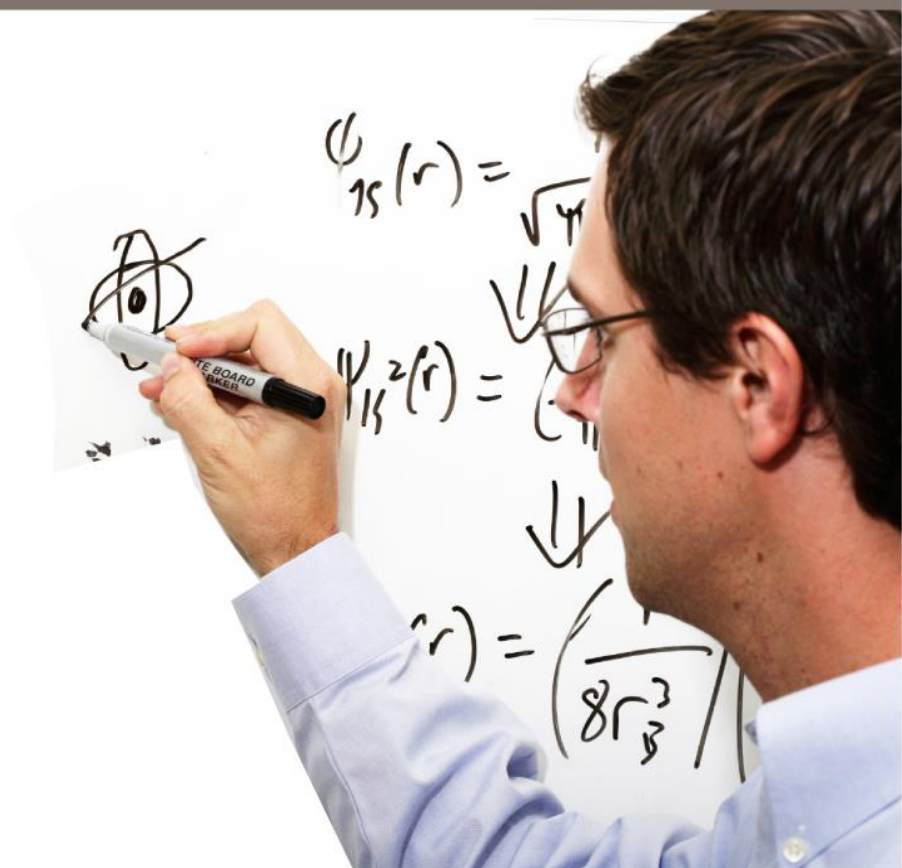


# Goals of HTS triage

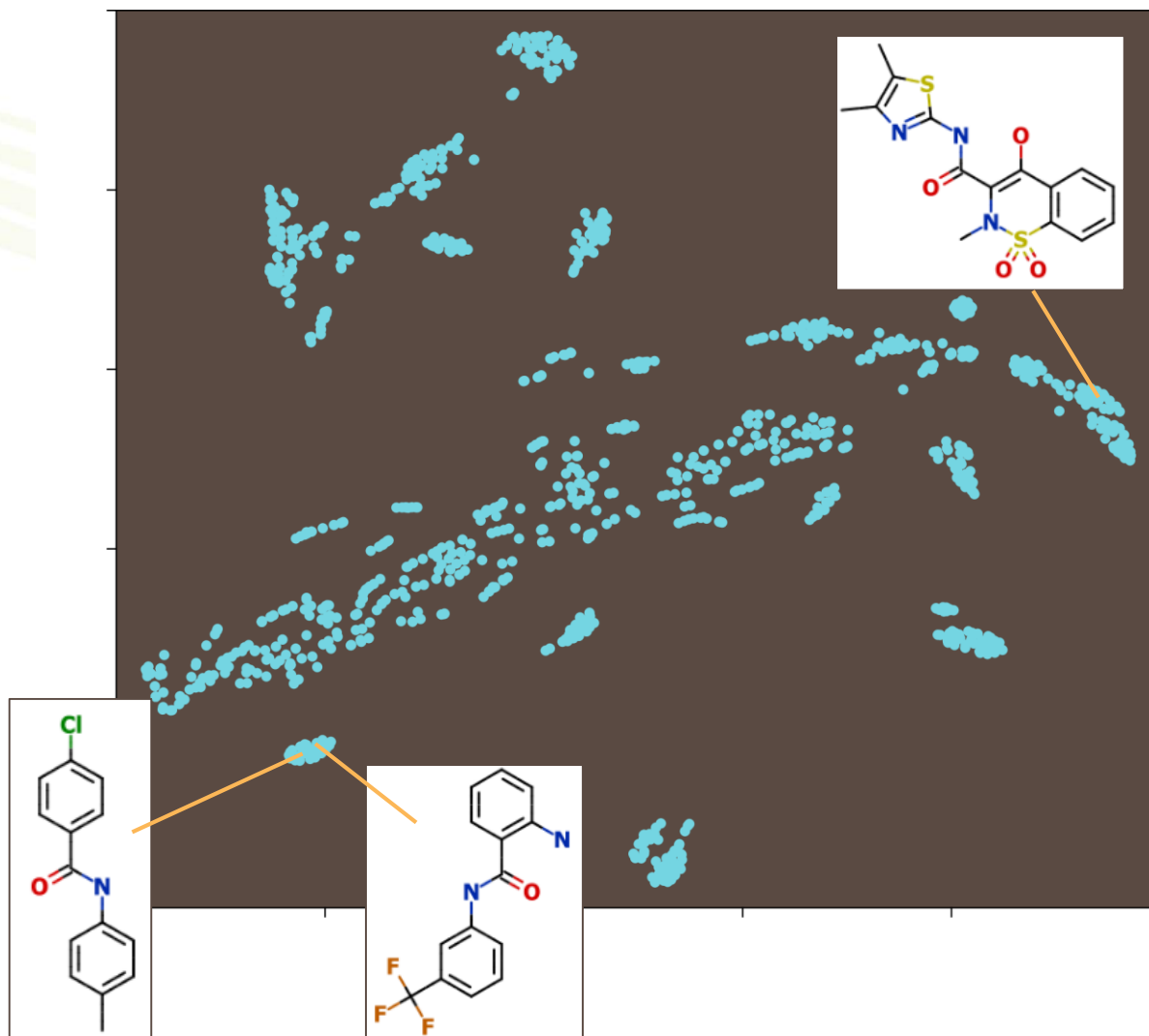
---

- One or more active series
  - Diversity is beneficial to provide backup series
- Good structure-activity relationships in series
  - Opportunities for optimisation
- High quality starting points for hit-to-lead
  - Appropriate physicochemical properties
  - Access to good absorption, distribution, metabolism and excretion (ADME) properties
  - Avoid frequent hitters (false positives) and high risk functionalities

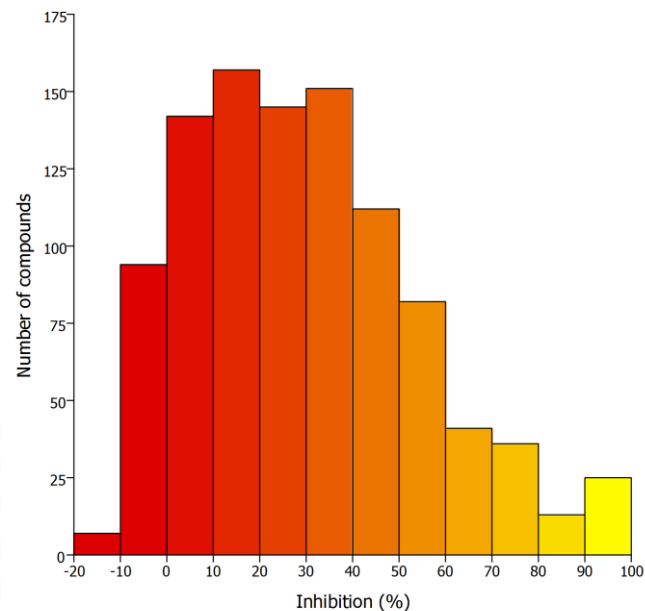
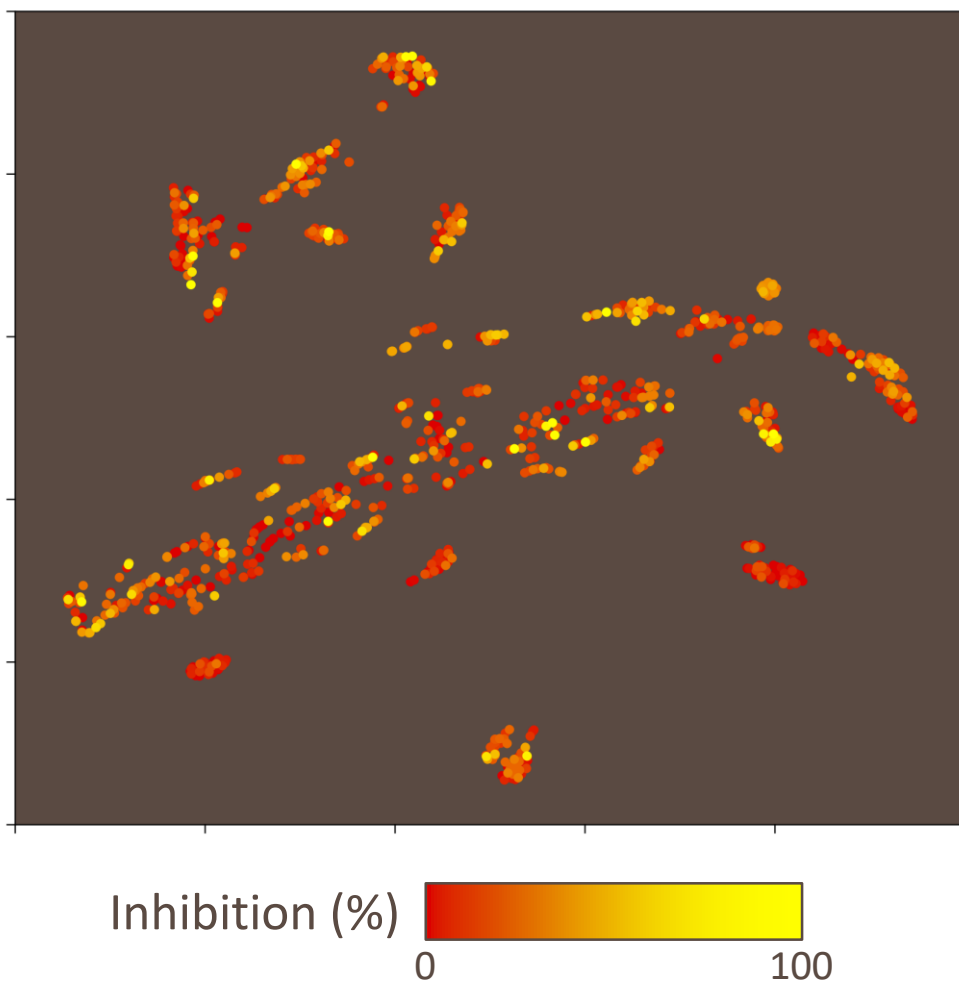
# Mapping the Chemical Space of Activity



# Example Screening Library

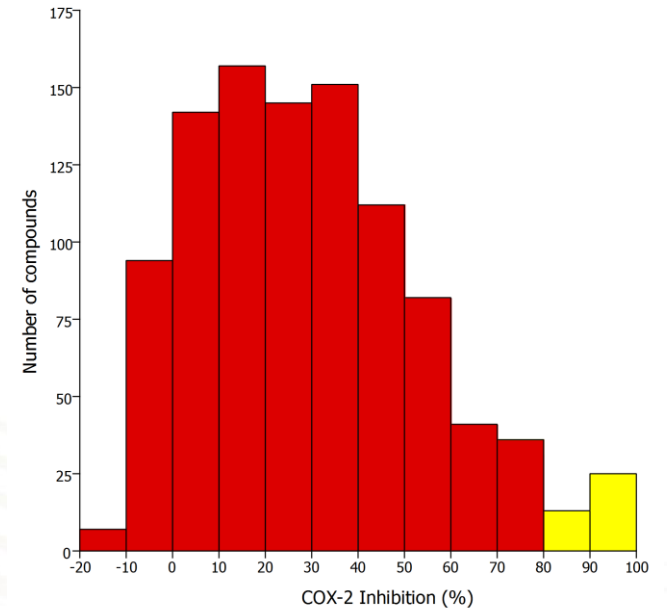
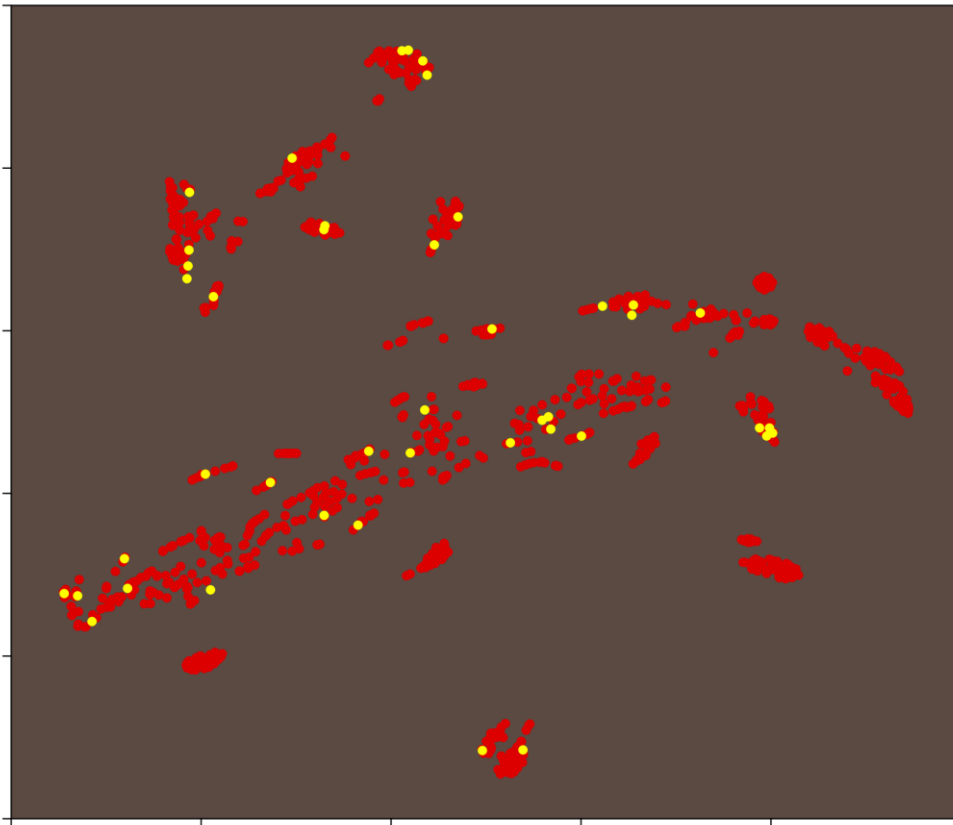


# Distribution of Activity Inhibition (%)



	Mean	Max	Min	SD
Inhibition (%)	31	100	-17	24

# Identifying Hits

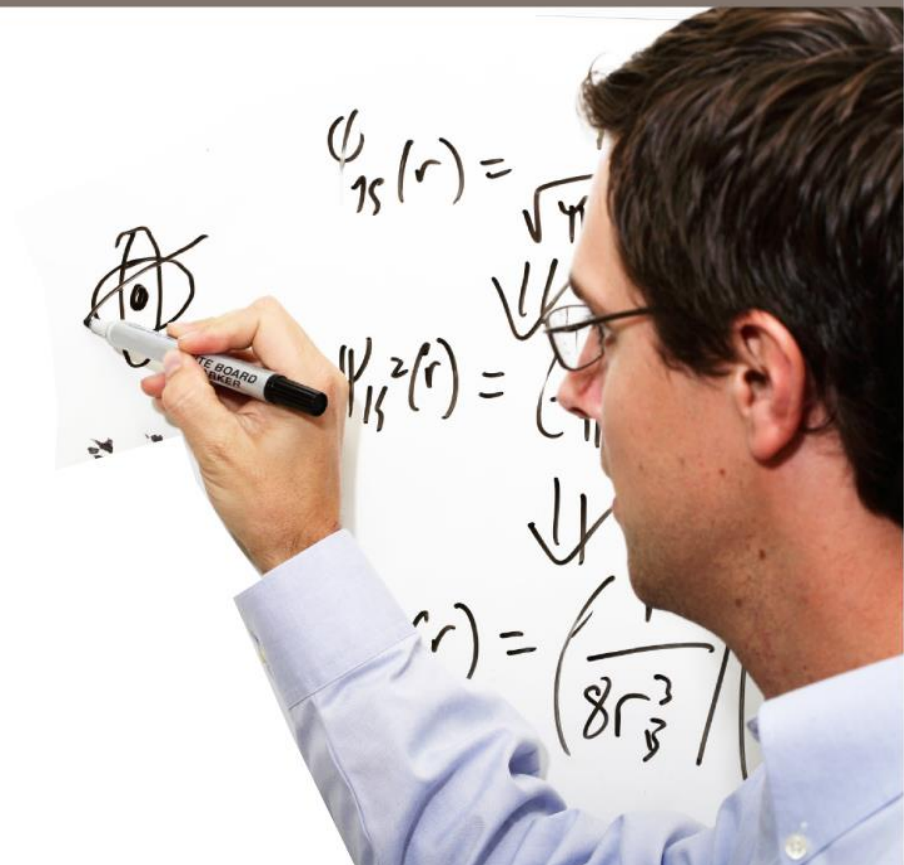


Hit (>80% Inhibition) ●

Miss ●

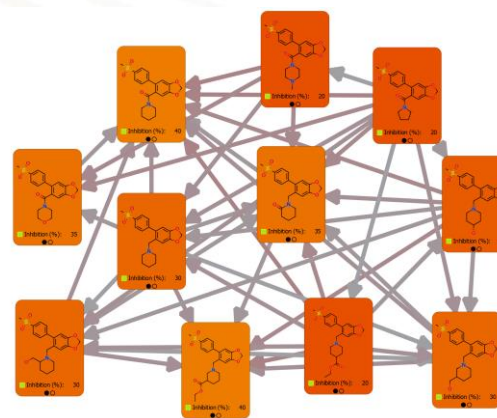
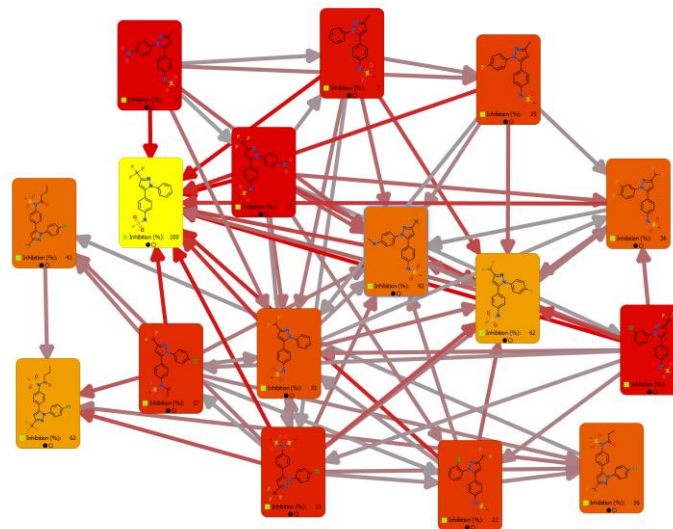


# Understanding the Activity Landscape

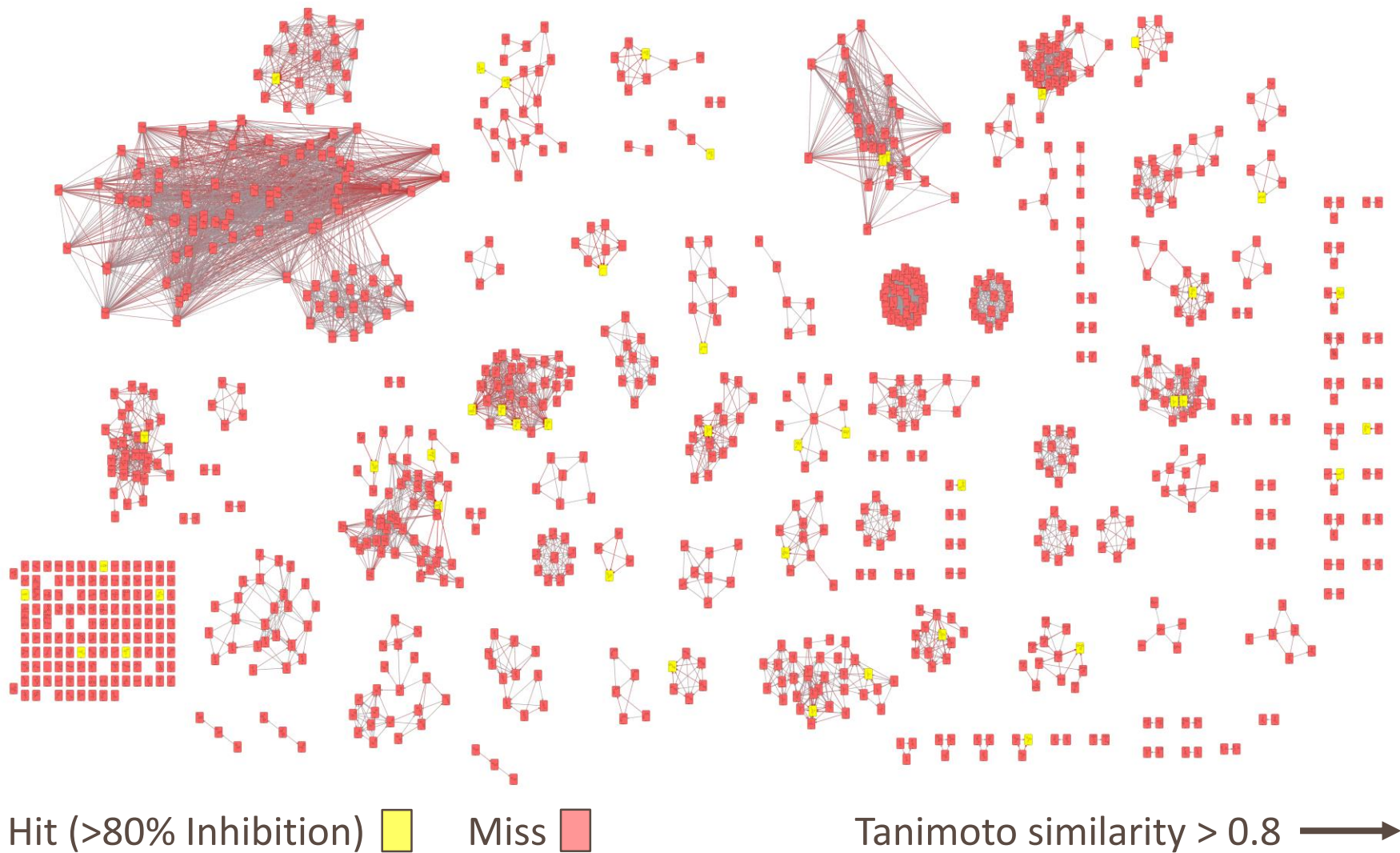


# Activity Landscapes

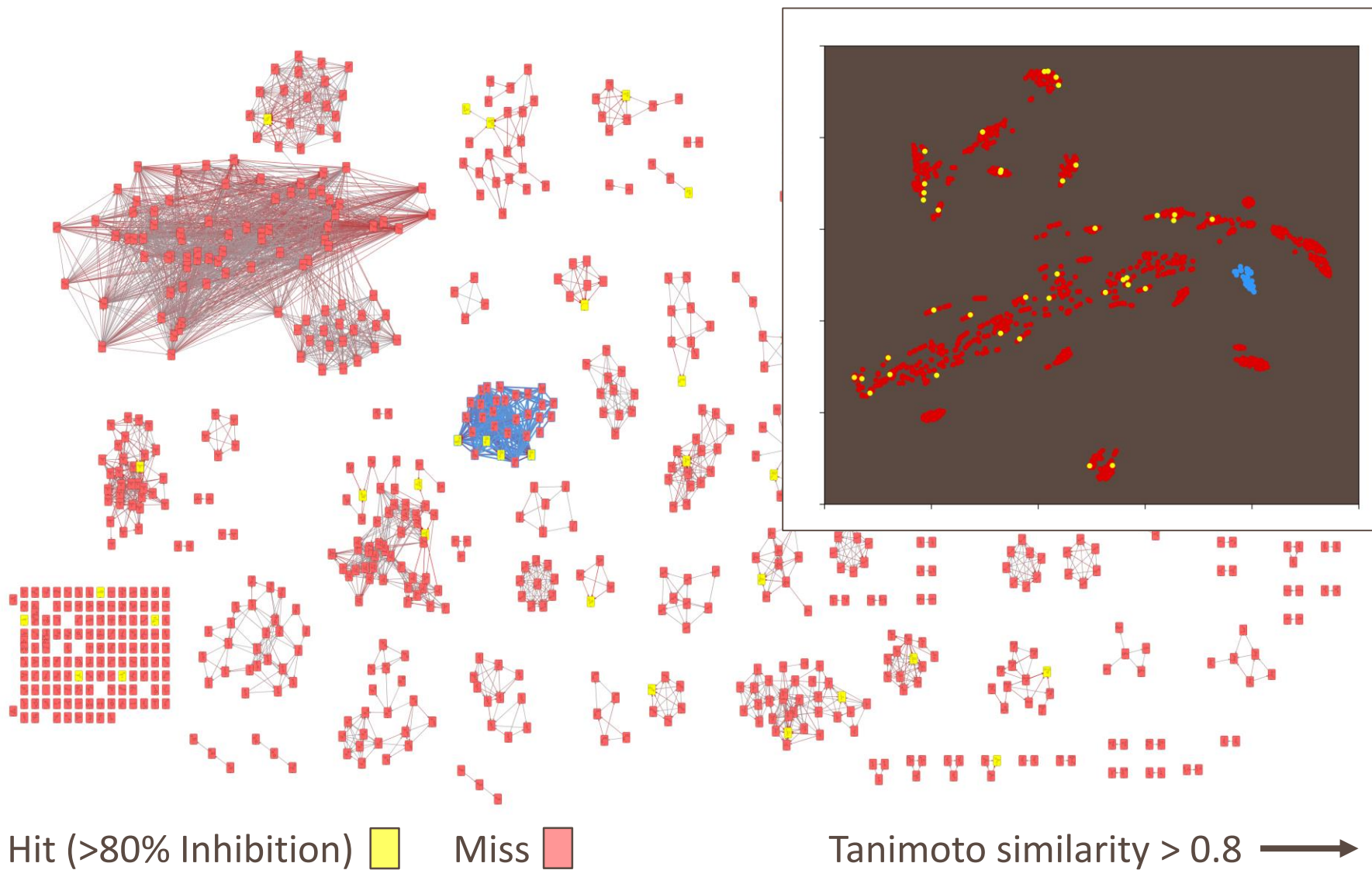
- All-by-all comparison
  - Identify groups of ‘similar’ compounds
- ‘Rough’ regions
  - large changes in activity result from small changes in structure
  - Interesting SAR
- ‘Flat spots’
  - Limited opportunity for optimisation of activity
  - Opportunity to optimise different property without having negative impact on activity



# Identifying Regions with Interesting SAR

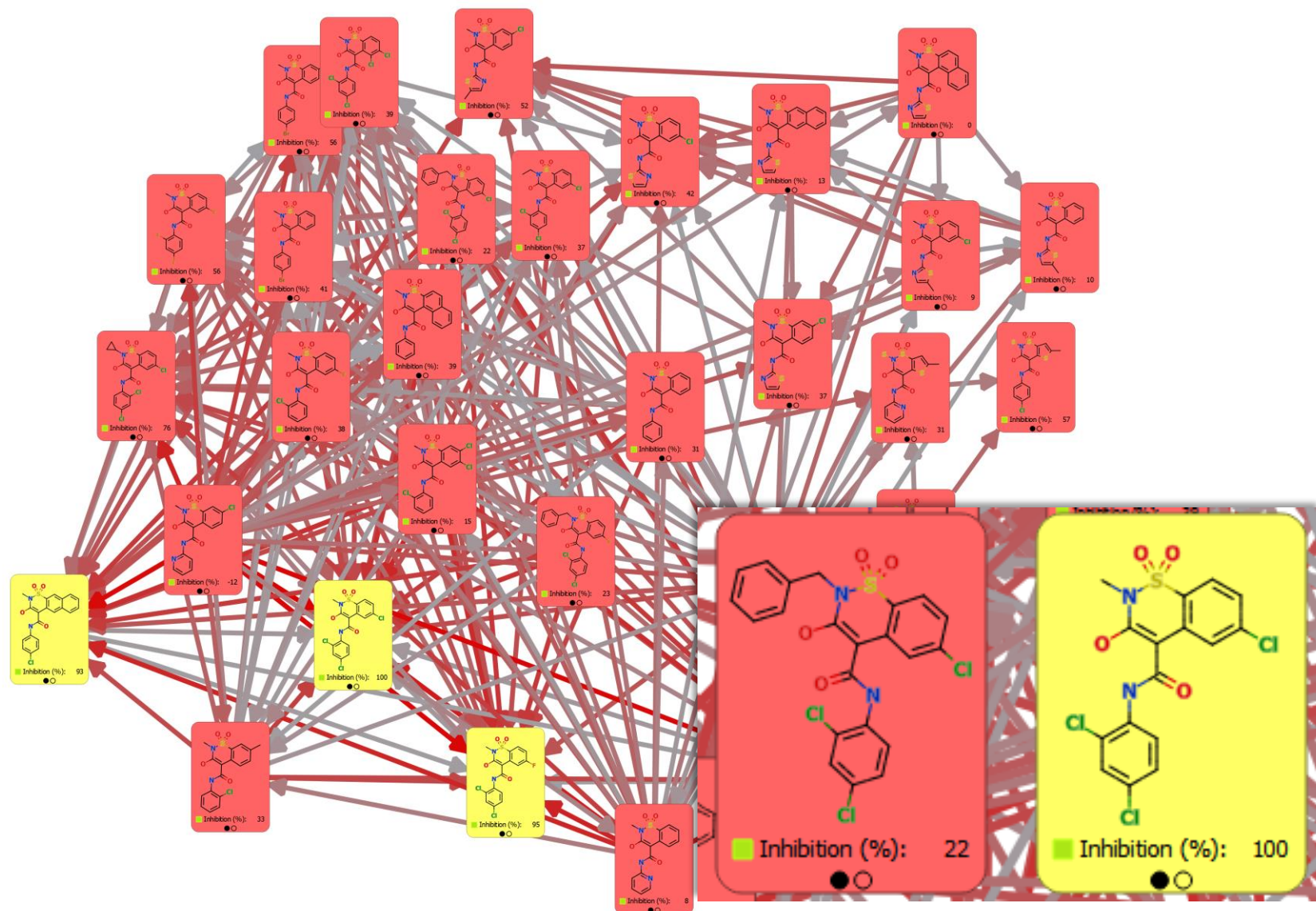


# Identifying Regions with Interesting SAR

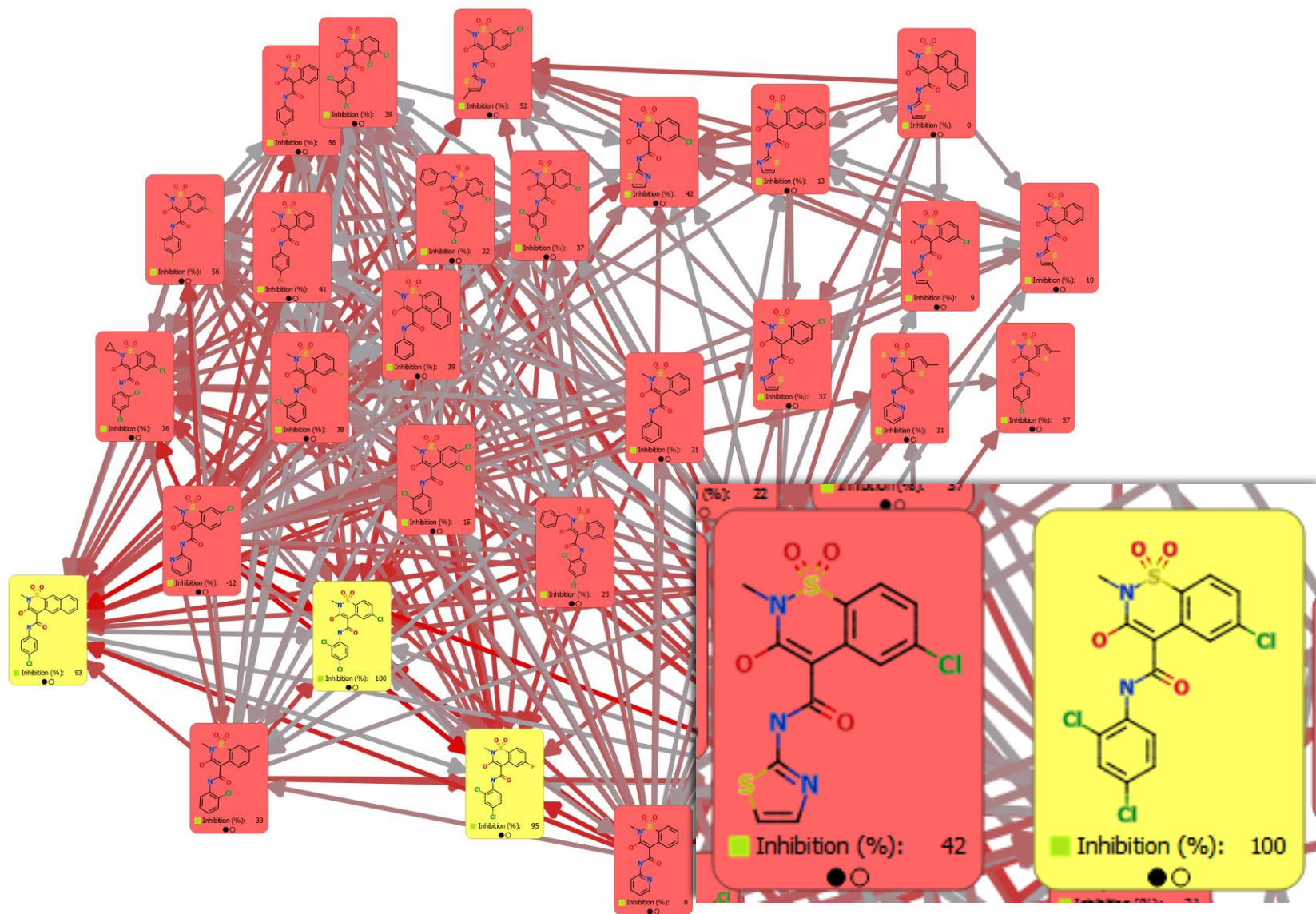




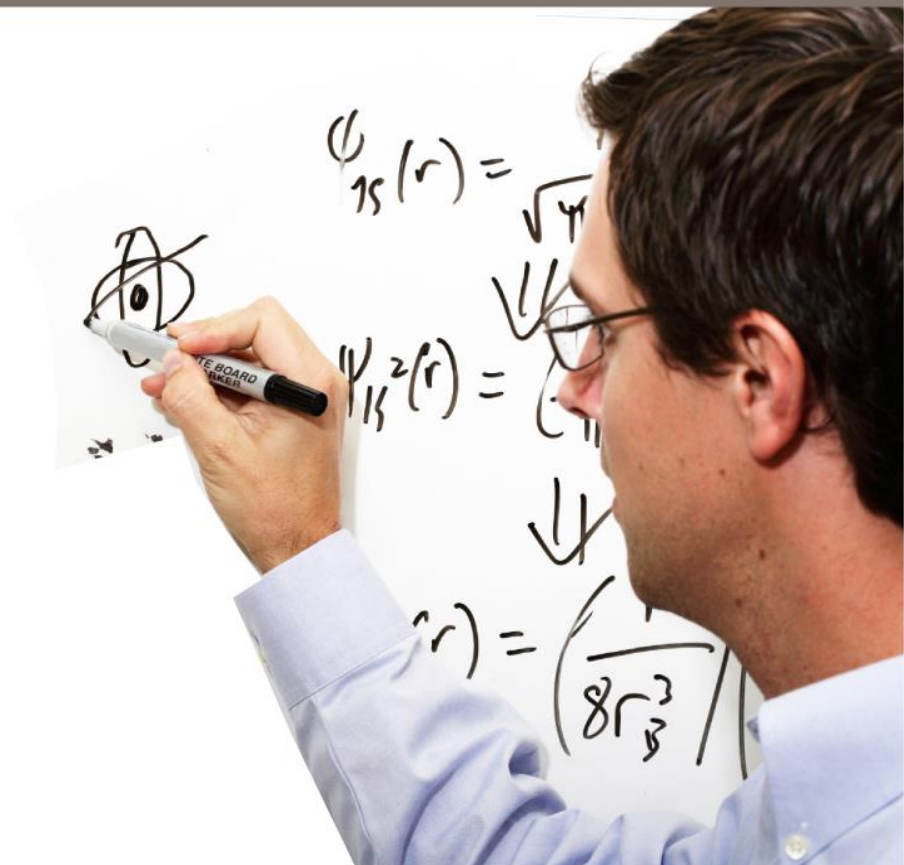
# Identifying Regions with Interesting SAR



# Identifying Regions with Interesting SAR



# Targeting High Quality Hit Series





# What Does “High Quality” Mean?

---

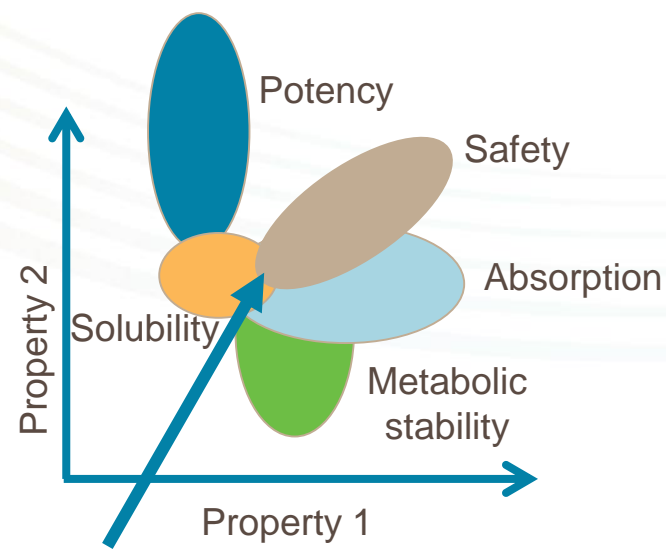
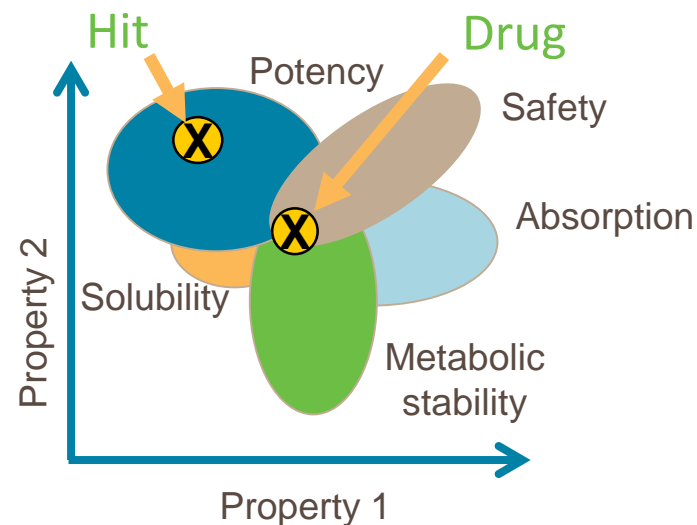
- Low molecular weight?
  - More room for optimisation
- Low logP?
  - Better opportunity for optimisation
  - Reduce risk of off-target effects
  - Better chance of good solubility/permeability
- Avoid pan-assay interference compounds (PAINS)\*?
  - Maybe promiscuous binders
  - Undesirable functionalities
- Appropriate ADME properties
  - Depends on project’s therapeutic objectives



# Guiding Decisions in Compound Optimisation

## Multi-parameter optimisation

- Identify chemistries with an optimal **balance** of properties
- Quickly identify situations when such a balance is not possible
  - Fail fast, fail cheap
  - Only when **confident**
  - Avoid **missed opportunities**



No good drug

# Filtering

Inhibition

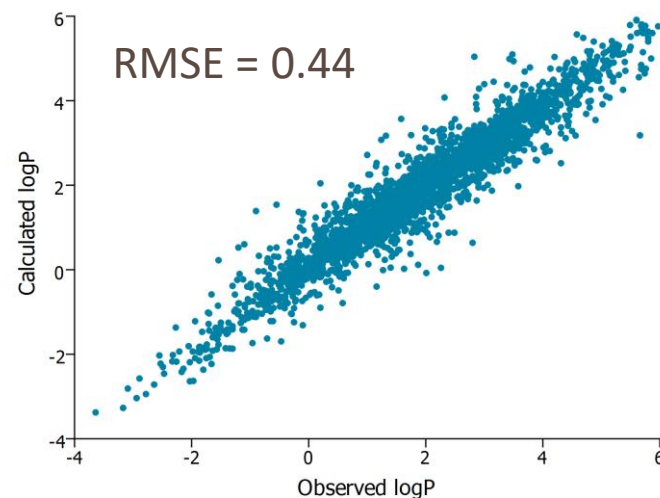
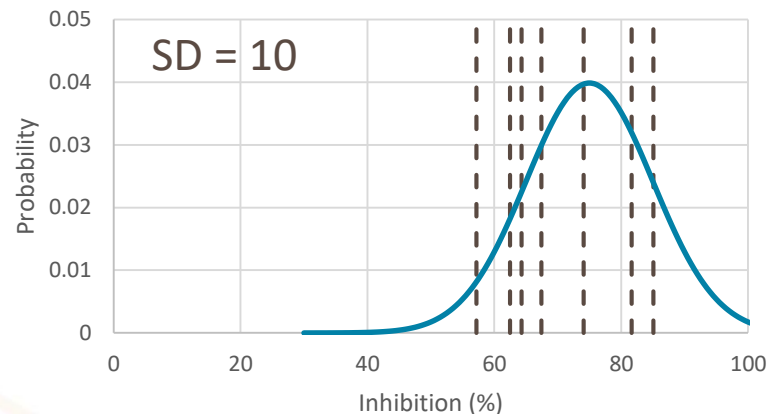
log<sub>10</sub>

PAINS



# Sources of Uncertainty







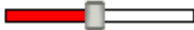
- Experimental variability
- Statistical uncertainty, e.g. logP
- Relevance, e.g. PAINS\*
  - Many compounds with PAINS alerts are not frequent hitters
  - Several successful drugs contain PAINS alerts



\*Capuzzi *et al.* JCIM DOI: 10.1021/acs.jcim.6b00465

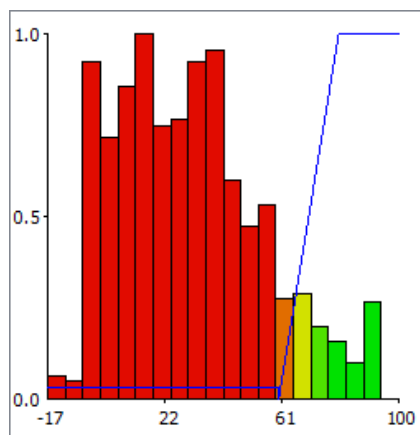
# Probabilistic Scoring

## Scoring Profile

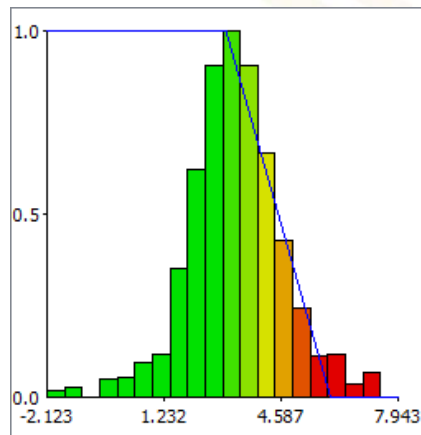
Property	Desired Value	Importance
Inhibition (%)	80 -> inf 	
logP	-inf -> 3 	
MW	-inf -> 300 	
PAINS count	0	

## Desirability Functions

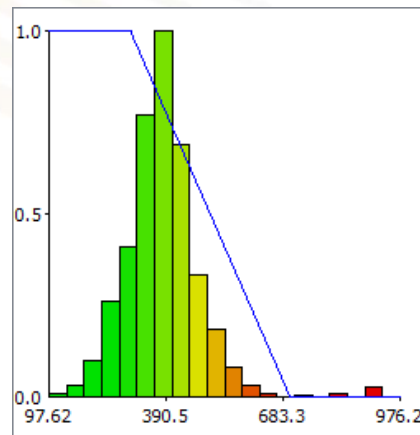
Inhibition (%)



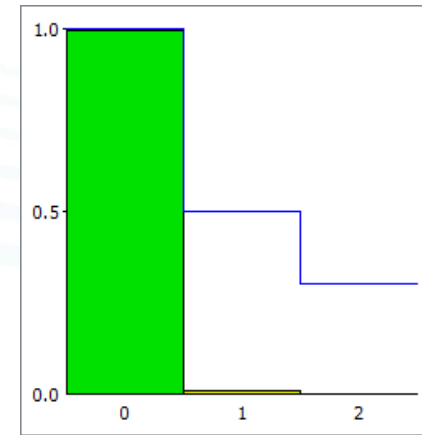
logP



MW



PAINS count

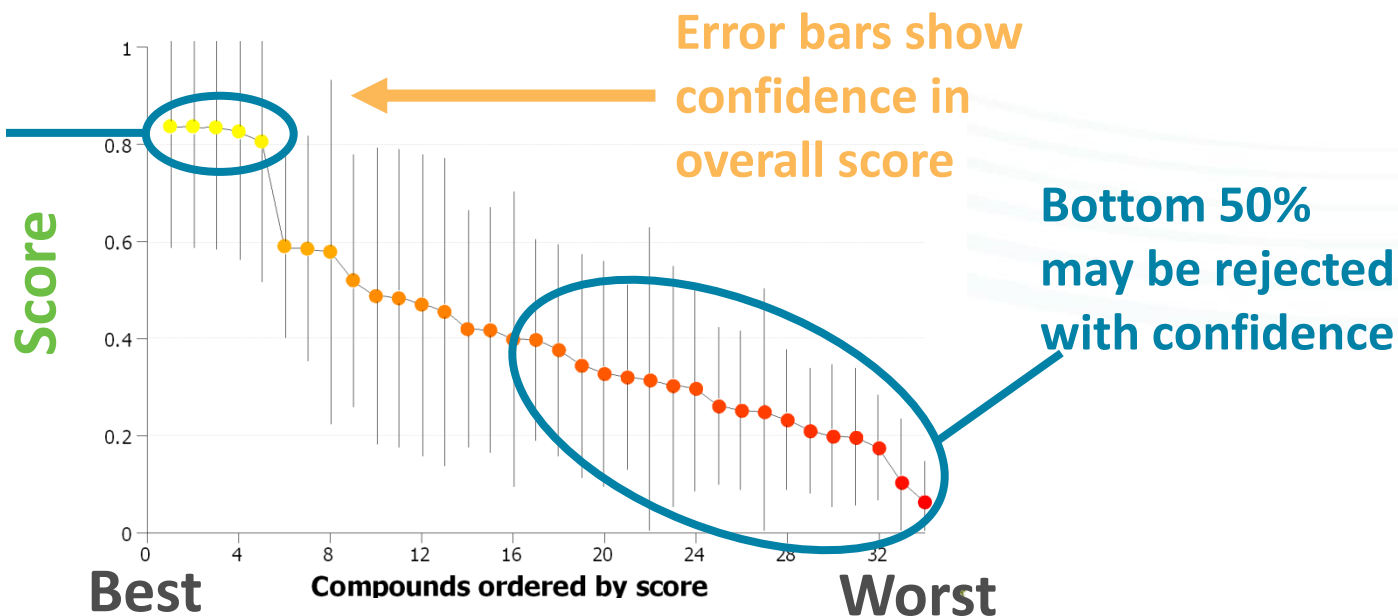


# Probabilistic Scoring

- **Property data**
  - Experimental or predicted
- **Criteria for success**
  - Relative importance
- **Uncertainties in data**
  - Experimental or statistical

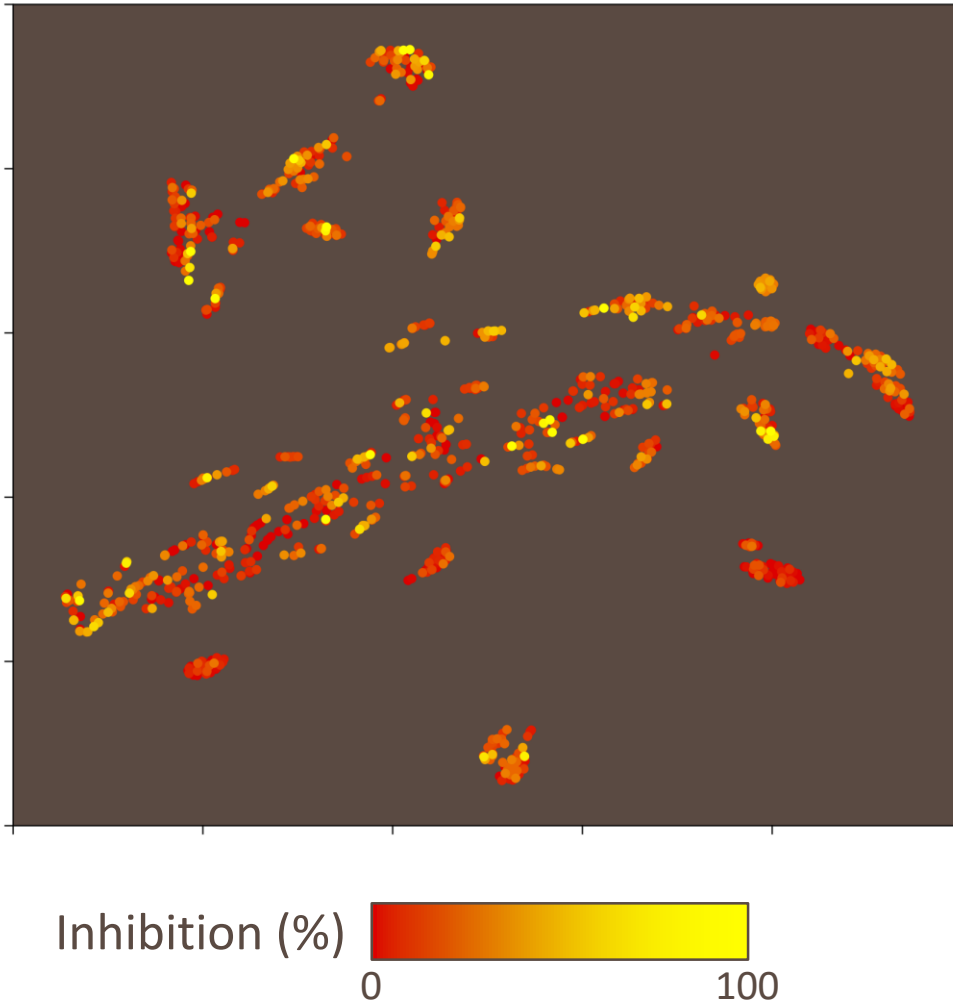
- **Score (Likelihood of Success)**
- **Confidence in score**

Data do not separate these as error bars overlap

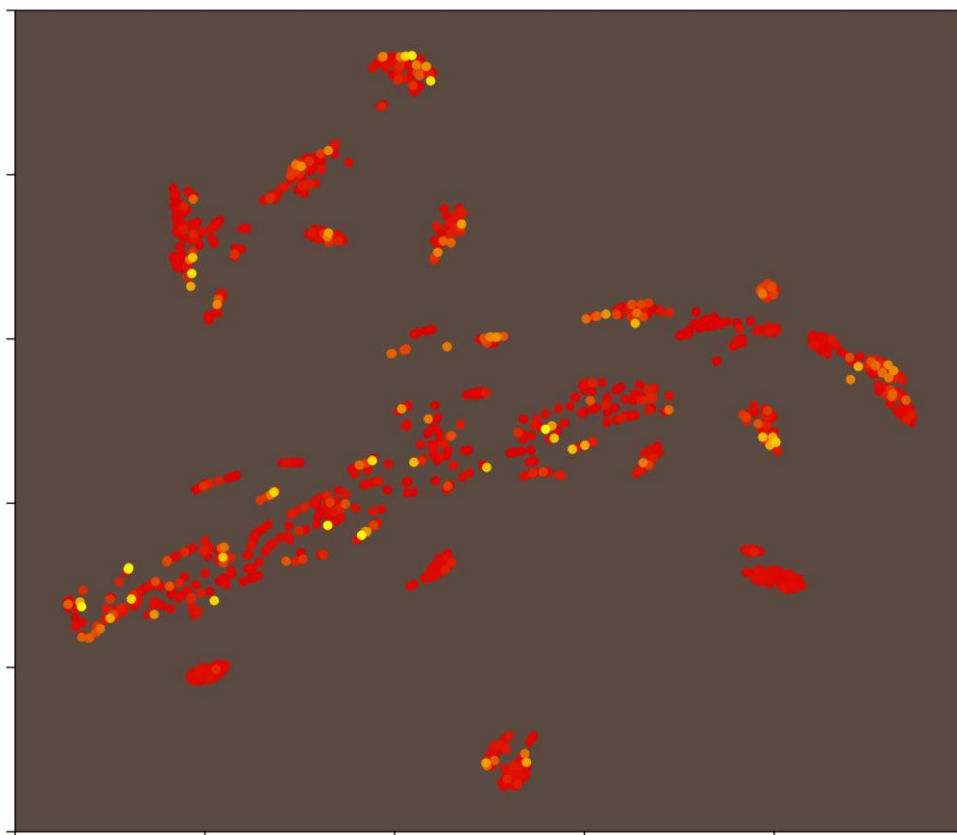










# Mapping Activity vs Score

---



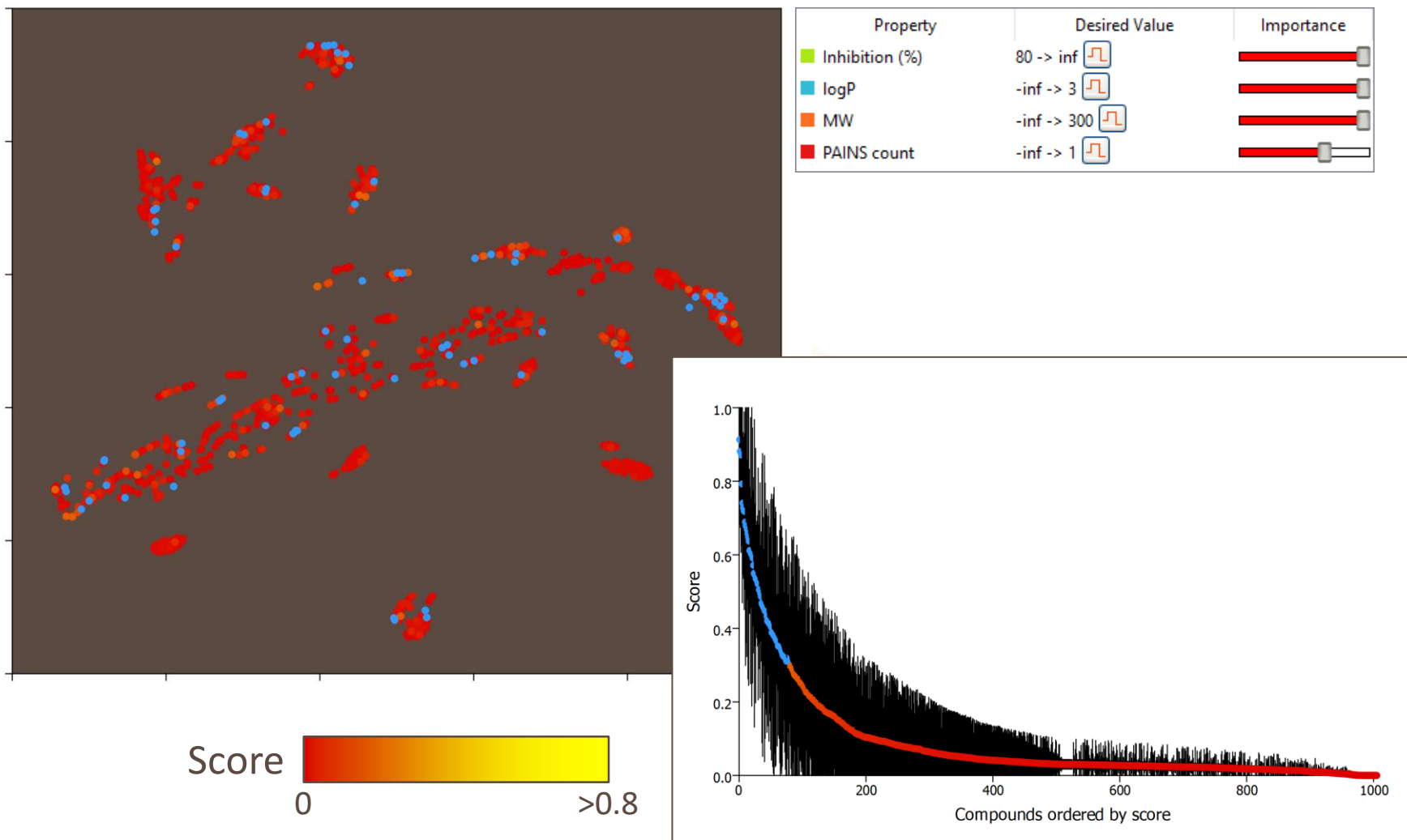
# Mapping Activity vs Score



Property	Desired Value	Importance
Inhibition (%)	80 -> inf 	
logP	-inf -> 3 	
MW	-inf -> 300 	
PAINS count	-inf -> 1 	



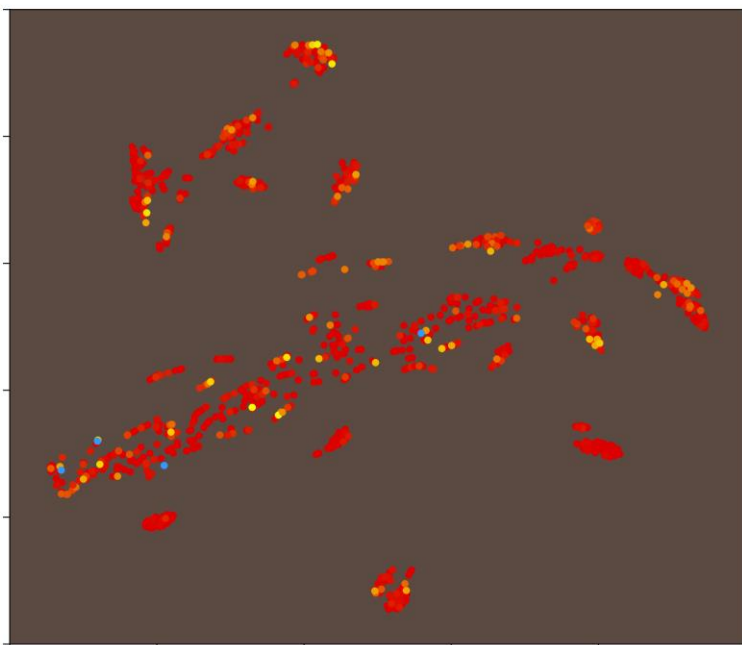
# Compound Selection





# Comparison with Filters

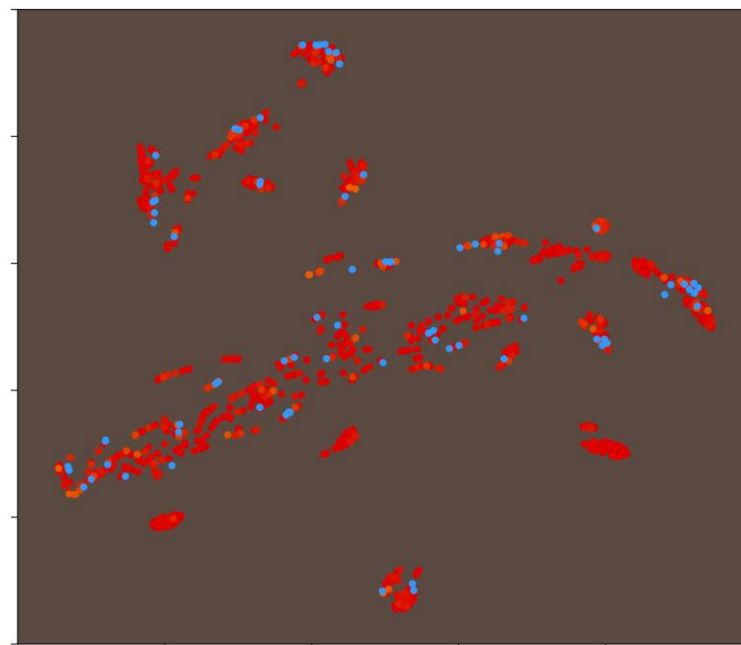
## Filtered Compounds











Inhibition (%) > 80  
MW < 300  
logP < 3  
No PAINS hits

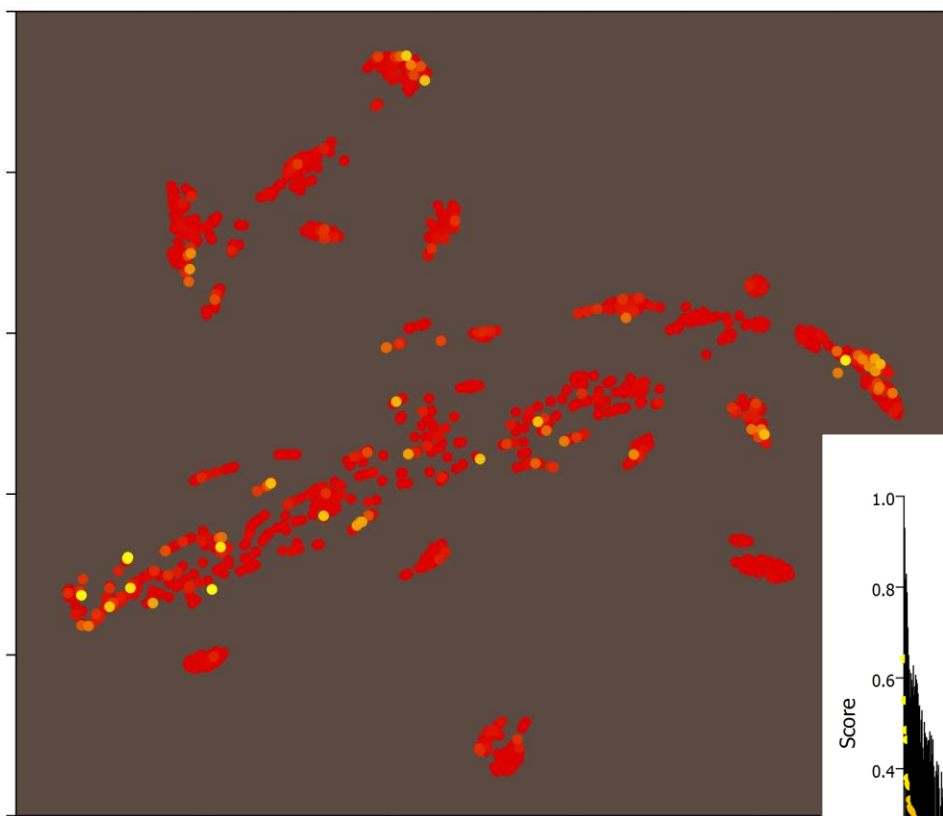
} Only 4 compounds!

## Scoring Selection

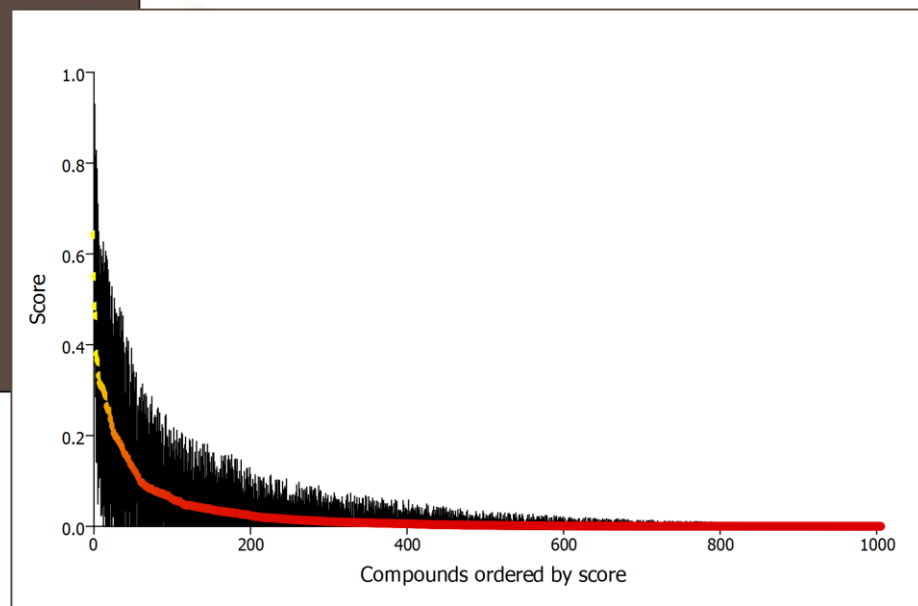


Property	Desired Value	Importance
Inhibition (%)	80 -> inf 	
logP	-inf -> 3 	
MW	-inf -> 300 	
PAINS count	-inf -> 1 	

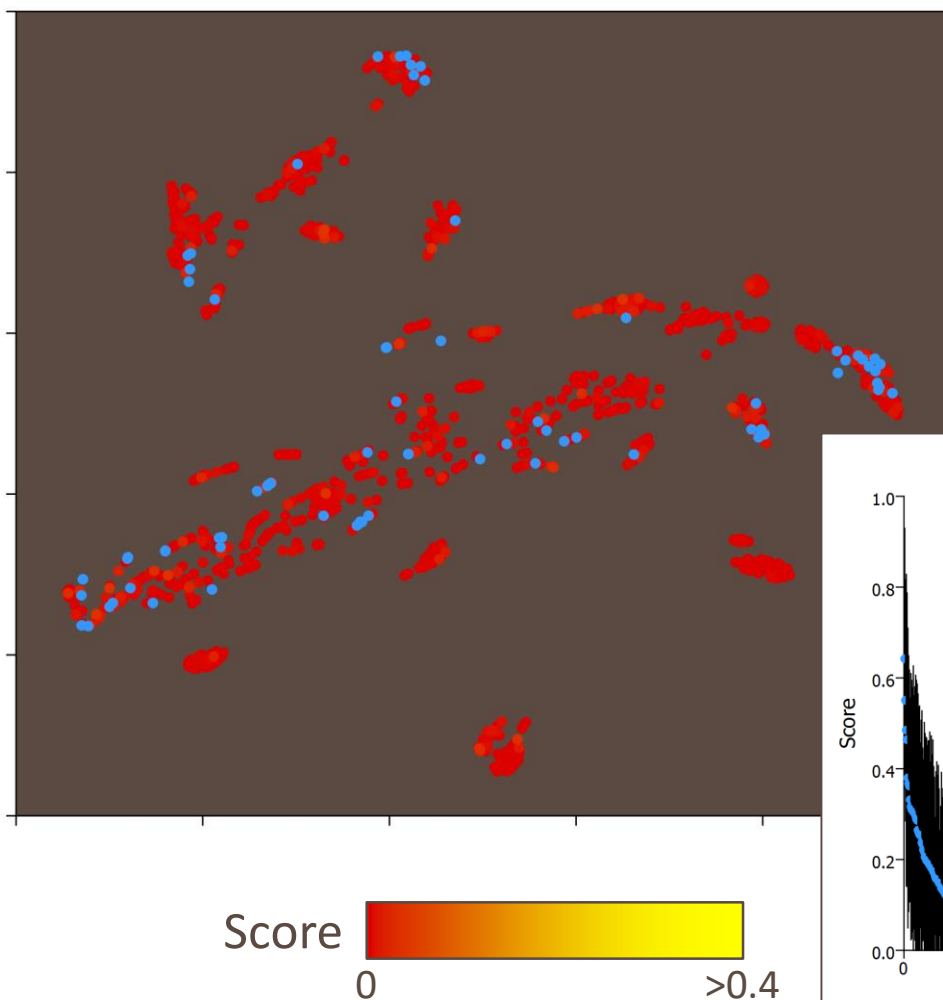
# ADME and Potency Profile



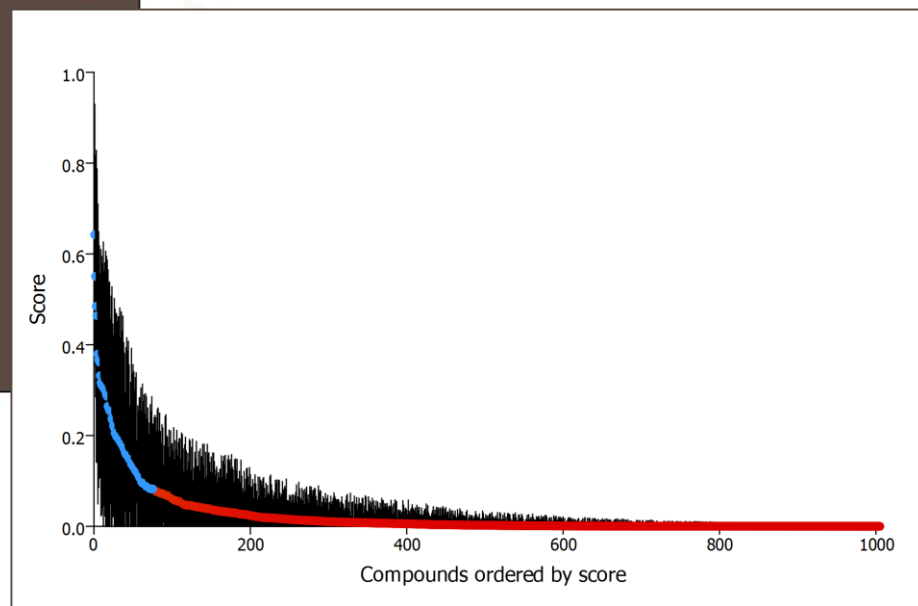
Property	Desired Value	Importance
Inhibition (%)	80 -> inf <input type="checkbox"/>	<input type="checkbox"/>
logS	> 1	<input type="checkbox"/>
HIA category	+	<input type="checkbox"/>
logP	0 -> 3.5 <input type="checkbox"/>	<input type="checkbox"/>
hERG pIC50	≤ 5	<input type="checkbox"/>
2D6 affinity category	low medium <input type="checkbox"/>	<input type="checkbox"/>
2C9 pKi	≤ 6	<input type="checkbox"/>
P-gp category	no	<input type="checkbox"/>
PPB90 category	low	<input type="checkbox"/>
BBB category	-	<input type="checkbox"/>
BBB log([brain]:[blood])	≤ -0.5	<input type="checkbox"/>



# ADME and Potency Profile Selection

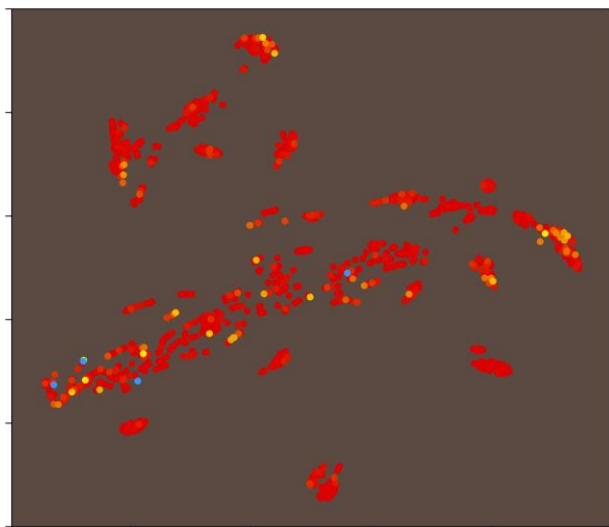


Property	Desired Value	Importance
Inhibition (%)	80 -> inf <input type="checkbox"/>	<input type="checkbox"/>
logS	> 1	<input type="checkbox"/>
HIA category	+	<input type="checkbox"/>
logP	0 -> 3.5 <input type="checkbox"/>	<input type="checkbox"/>
hERG pIC50	≤ 5	<input type="checkbox"/>
2D6 affinity category	low medium <input type="checkbox"/>	<input type="checkbox"/>
2C9 pKi	≤ 6	<input type="checkbox"/>
P-gp category	no	<input type="checkbox"/>
PPB90 category	low	<input type="checkbox"/>
BBB category	-	<input type="checkbox"/>
BBB log([brain];[blood])	≤ -0.5	<input type="checkbox"/>



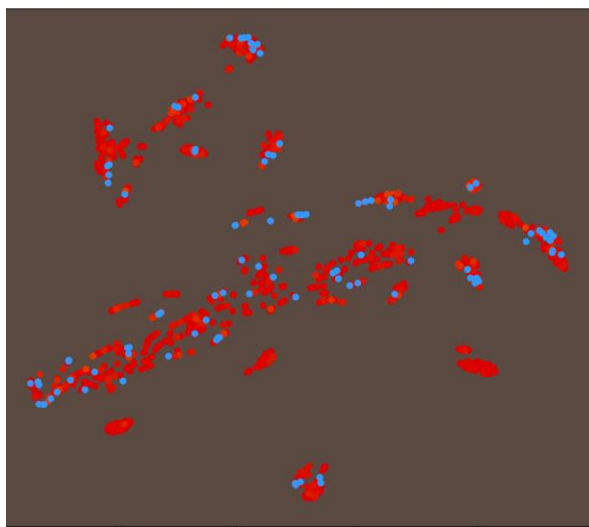
# Comparing Selection Strategies

## Filtered Compounds



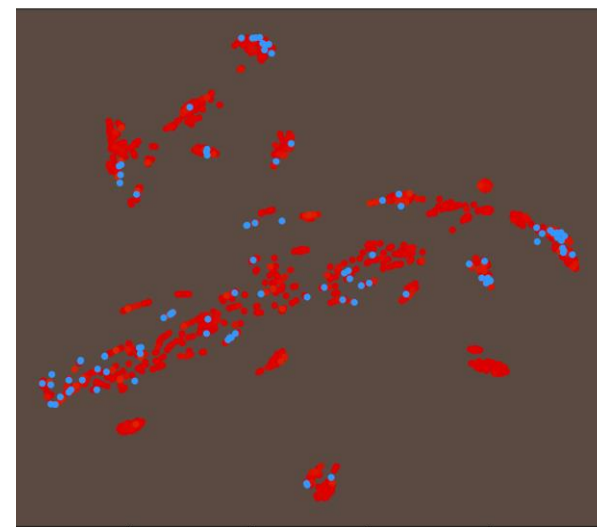
Inhibition (%) > 80  
 MW < 300  
 logP < 3  
 No PAINS hits

## Simple Profile



Property	Desired Value	Importance
Inhibition (%)	80 -> inf	High
logP	-inf -> 3	High
MW	-inf -> 300	High
PAINS count	-inf -> 1	Medium

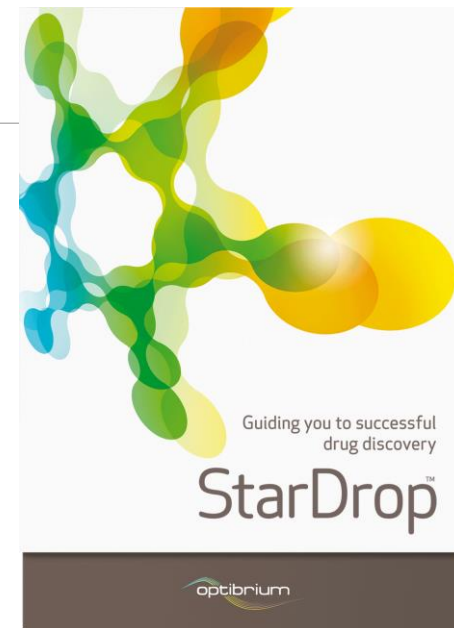
## ADME + Potency Profile



Property	Desired Value	Importance
Inhibition (%)	80 -> inf	High
logS	> 1	High
HIA category	+	High
logP	0 -> 3.5	High
hERG pIC50	≤ 5	High
2D6 affinity category	low medium	High
2C9 pKi	≤ 6	High
P-gp category	no	High
PPB90 category	low	High
BBB category	-	High
BBB log([brain];[blood])	≤ -0.5	High

# Conclusions

- When prioritising compounds/series from HTS, we should consider:
  - Activity
  - SAR of active series
  - Quality of hits
  - Novelty
- In assessing quality:
  - Be careful of ‘hard’ filters
  - Optimise the *balance* of properties appropriate for your project
  - Consider the uncertainties in your data – avoid missed opportunities
- Publications can be downloaded from [www.optibrium.com/community](http://www.optibrium.com/community)
- For more information, please visit Booth #1420, go to [www.optibrium.com/stardrop](http://www.optibrium.com/stardrop) or contact [info@optibrium.com](mailto:info@optibrium.com)



# Acknowledgements

---

Colleagues at Optibrium, including:

- Chris Leeding
- James Chisholm
- Nick Foster
- Alex Elliott
- Fayzan Ahmed
- Rasmus Leth
- Coran Hoskin
- Mario Öeren
- Aishling Cooke

