

Numerical Characterization of Structure-Activity Relationships from a Medicinal Chemists Point of View

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Assumptions

- ▶ Similar molecules will have similar activities
- ▶ Small changes in structure will lead to small changes in activity
- ▶ One implication is that SAR's are additive
- ▶ This is the basis for QSAR modeling

Structure Activity Landscapes

Melanocortin-4 receptor inhibitors

Defining & Using
Structure-Activity
Landscapes

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Background

Visualization

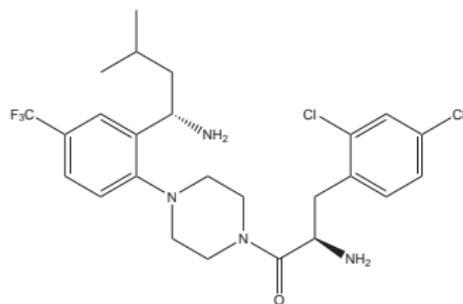
Utilization

Predictive Models

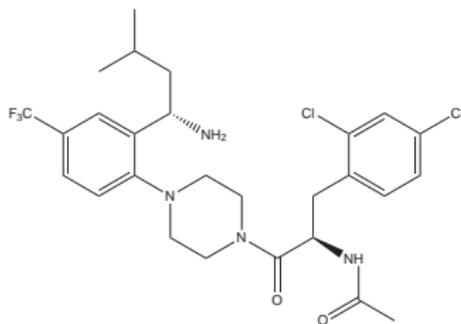
3D models

Chemical spaces

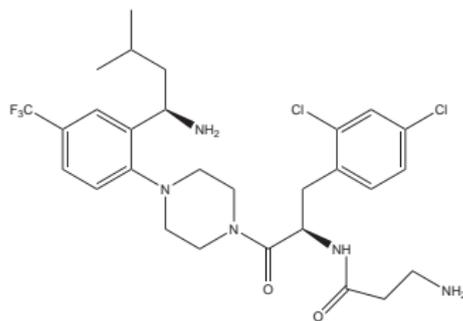
Summary



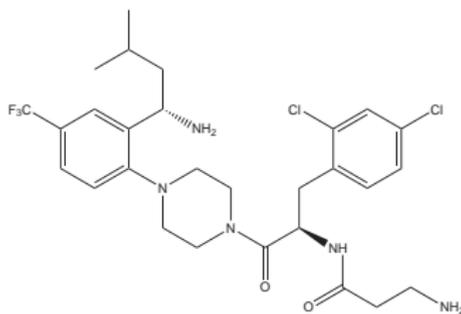
$K_i = 39.0 \text{ nM}$



$K_i = 1.8 \text{ nM}$



$K_i = 10.0 \text{ nM}$



$K_i = 1.0 \text{ nM}$

Structure Activity Landscapes



Rugged gorges or rolling hills?

- ▶ Small structural changes associated with large activity changes represent steep slopes in the landscape
 - ▶ **Activity Cliffs**
- ▶ But traditionally, QSAR *assumes* gentle slopes
- ▶ Machine learning is not very good for special cases

Converting activity cliffs to numbers

- ▶ A cliff can be numerically characterized
- ▶ Structure-Activity Landscape Index (SALI)

$$\text{SALI}_{i,j} = \frac{|A_i - A_j|}{1 - \text{sim}(i,j)}$$

- ▶ Cliffs are characterized by elements of the matrix with very large values

Visualizing the SALI Matrix

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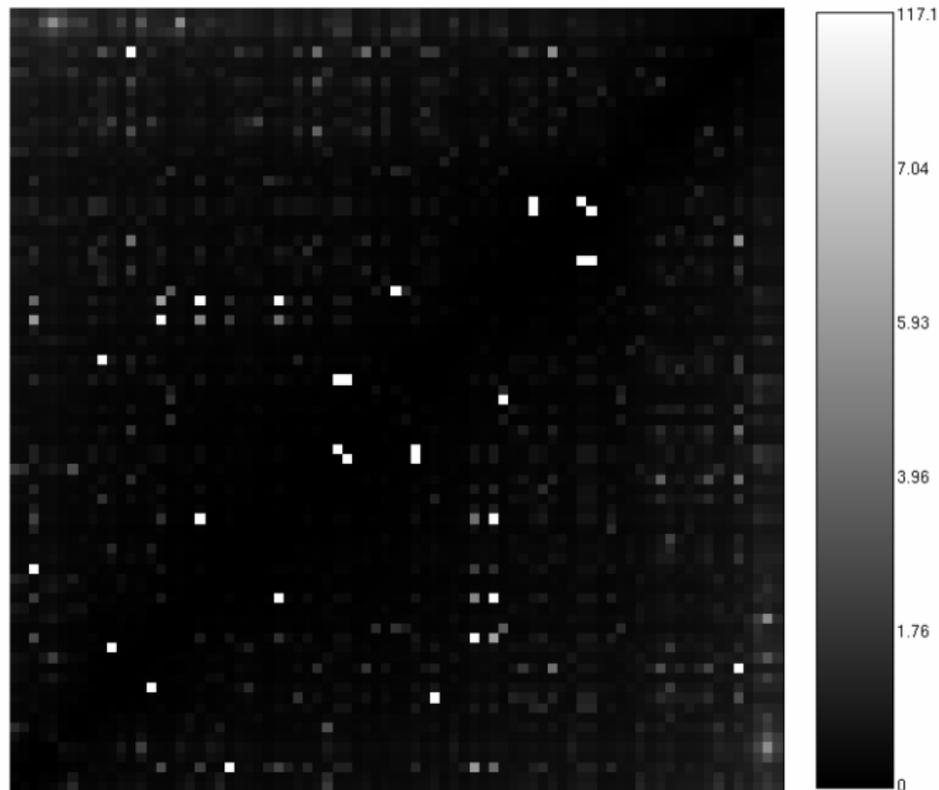
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Visualizing SALI Values

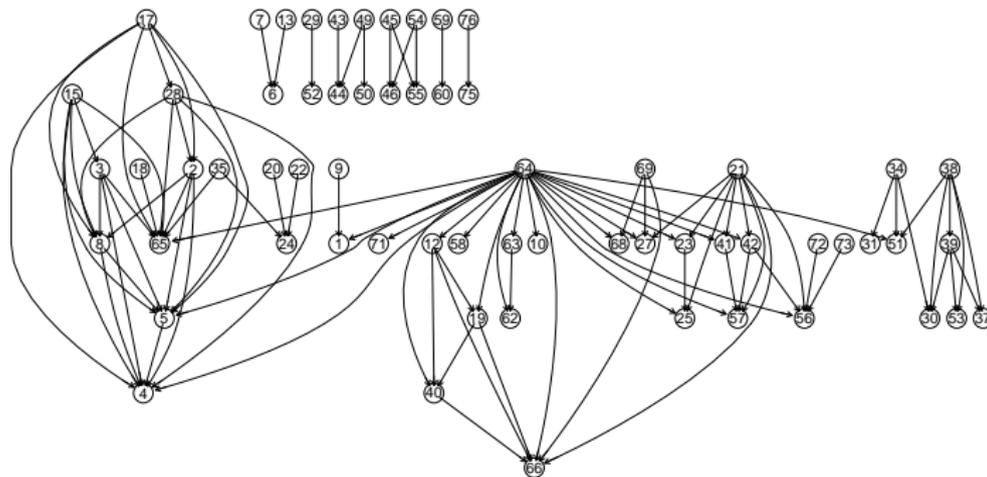
Alternatives?

- ▶ A heatmap is an easy to understand visualization
- ▶ Coupled with brushing, can be a handy tool
- ▶ A more flexible approach is to consider a network view of the matrix

The SALI graph

- ▶ Compounds are nodes
- ▶ Nodes i, j are connected if $SALI_{i,j} > X$
- ▶ Only display connected nodes

Visualizing the SALI Graph



- Nodes are ordered such that the tail node in an edge has lower activity than the head node

SALIViewer

- ▶ Java application for generating and visualizing SALI graphs
- ▶ Create SALI graphs from SMILES and activity data, using the CDK fingerprints
- ▶ Easily examine SALI graphs at different cutoffs
- ▶ Provides 2D depictions for nodes and edges
- ▶ Generate SALI curves

Better Visualization - SALIVIEWER

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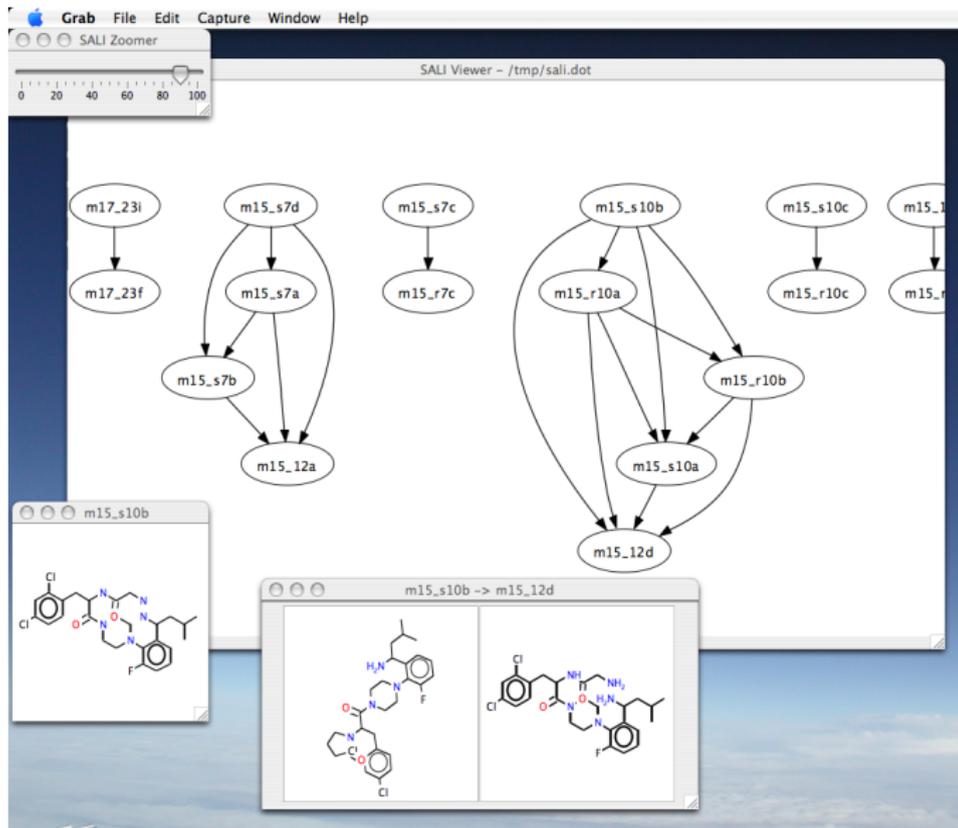
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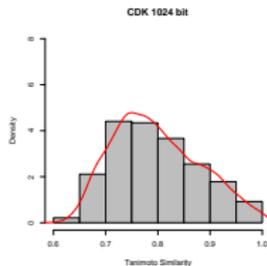
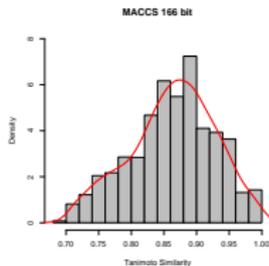
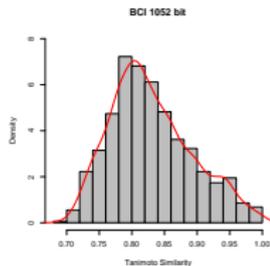
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Varying Fingerprint Methods

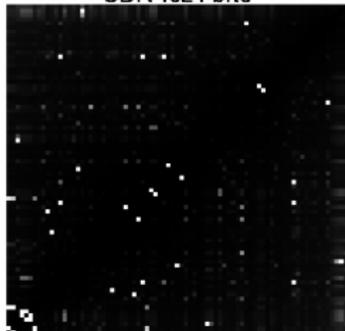
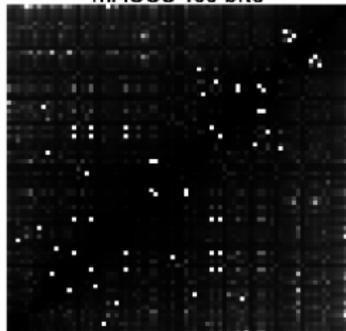
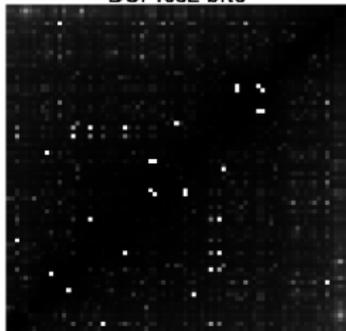


- ▶ Shorter fingerprints will lead to more “similar” pairs
- ▶ Requires a higher cutoff to focus on significant cliffs

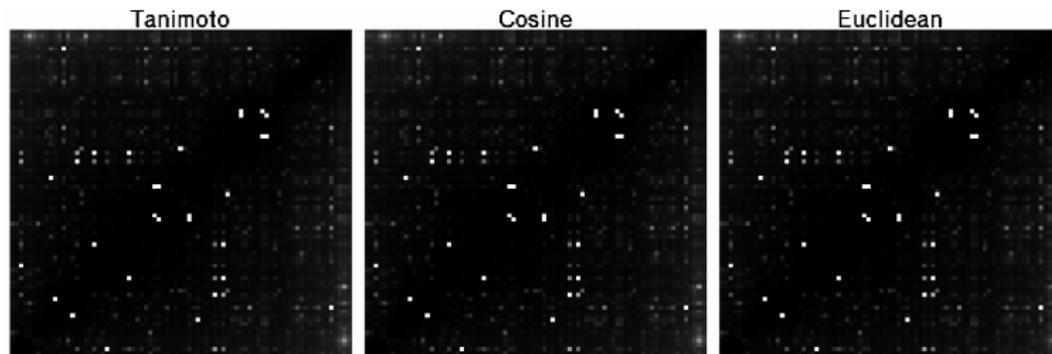
BCI 1052 bits

MACCS 166 bits

CDK 1024 bits



Varying the Similarity Metric



- ▶ The similarity metric does not affect the SALI values

SALI Graphs & Predictive Models

- ▶ The graph view allows us to view SAR's and identify trends easily
- ▶ The aim of a QSAR model is to encode SAR's
- ▶ Traditionally, we consider the quality of a model in terms of RMSE or R^2
- ▶ But in general, we're not as interested in RMSE's as we are in whether the model predicted something as more active than something else
 - ▶ What we want to have is the correct **ordering**
 - ▶ We assume the model is statistically significant

Measuring model quality

- ▶ A QSAR model should easily encode the “rolling hills”
- ▶ A good model captures the most significant cliffs
- ▶ Can be formalized as

How many of the edge orderings of a SALI graph does the model predict correctly?

- ▶ Define $S(X)$, representing the number of edges correctly predicted for a SALI network at a threshold X
- ▶ Repeat for varying X and obtain the *SALI curve*

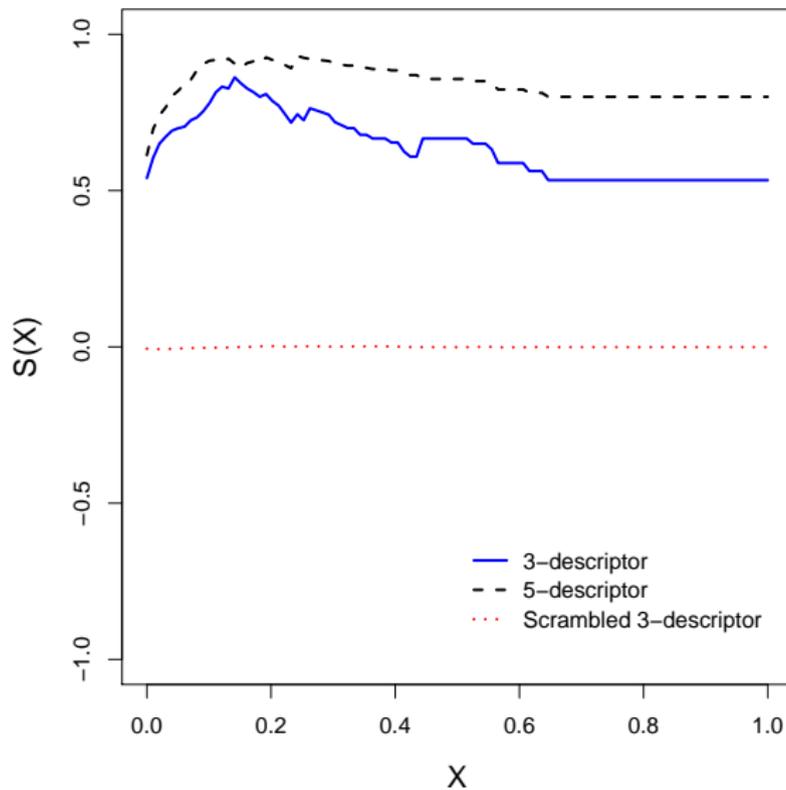
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SALI Curves - An Example



SALI Curves & Model Comparison

- ▶ Considered four datasets
- ▶ Developed linear regression models, using exhaustive search for feature selection
- ▶ Identify three models for each dataset
 - ▶ Minimum RMSE (“best”)
 - ▶ Median RMSE
 - ▶ Maximum RMSE (“worst”)
- ▶ Generate SALI curves for each model and summarize by dataset

SALI Curves & Model Comparison

Defining & Using
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Background

Visualization

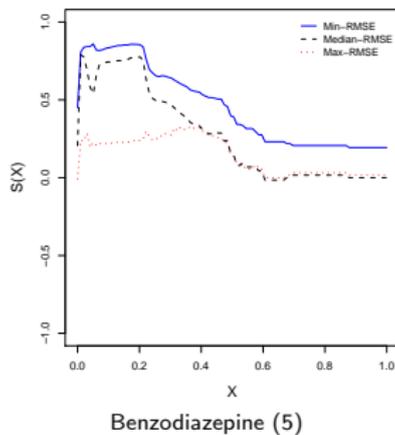
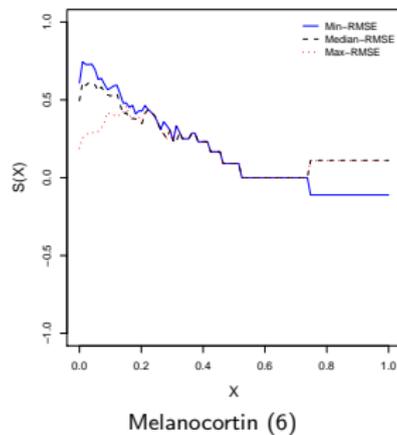
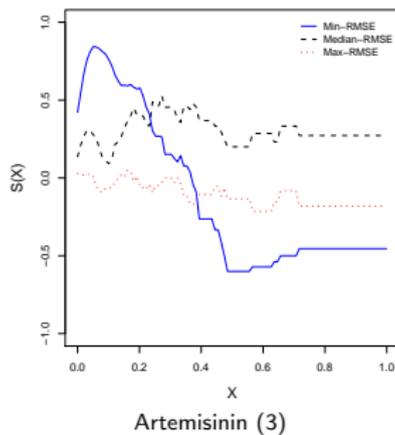
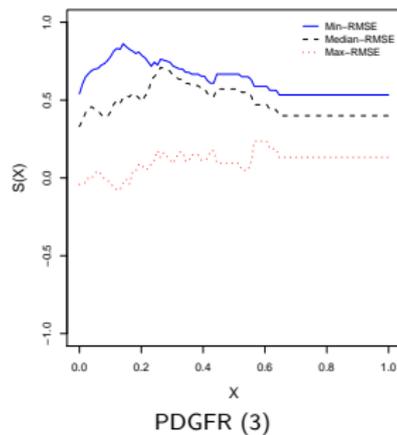
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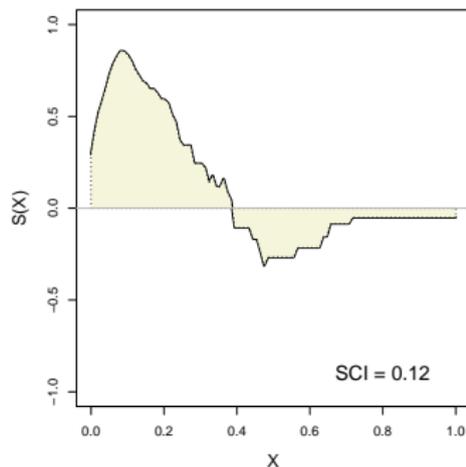
Chemical spaces

Summary



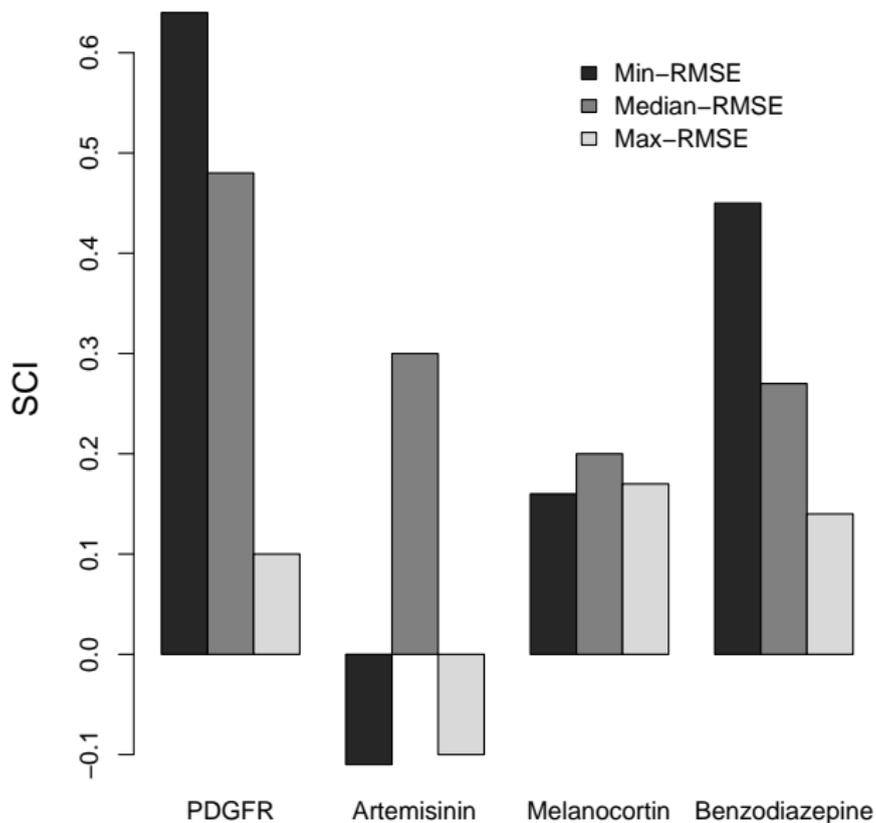
SALI Curves & Model Comparison

- ▶ The initial and final portions of the curve are of interest
- ▶ It's also useful to summarize the whole curve
- ▶ We evaluate the area between the curve and the X-axis (SCI)
 - ▶ $-1 \leq \text{SCI} \leq 1$



SALI Curve Integral

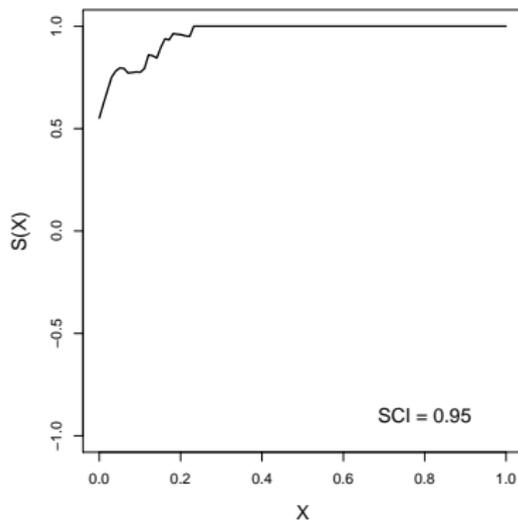
SALI Curves & Model Comparison



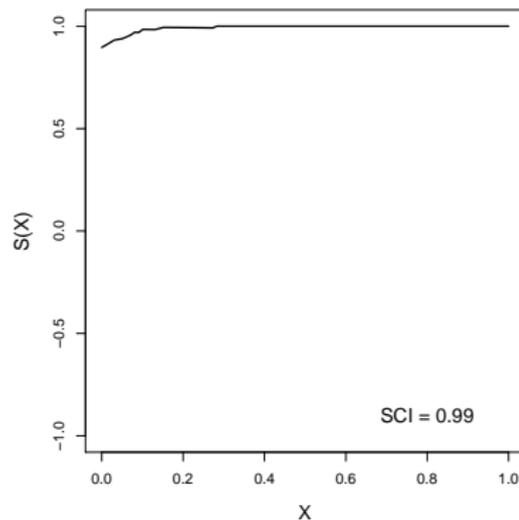
Examining Any Type of Model ...

- ▶ Previous examples make use of predicted values from QSAR models
- ▶ We can consider any “prediction” that is supposed to track observed activity
 - ▶ Ranks
 - ▶ Energies
- ▶ Allows us to apply this approach to *any* type of computational model that predicts something
 - ▶ Docking
 - ▶ CoMFA
 - ▶ Pharmacophore

Docking & CoMFA Models



Docking



CoMFA

- ▶ Not surprising that 3D models capture more cliffs
- ▶ The CoMFA model is nearly perfect!

Comparing Landscapes

- ▶ The SALI curve is a function of
 - ▶ dataset
 - ▶ descriptor space
- ▶ We can quantify a descriptor spaces ability to encode the structure-activity landscape using SALI graphs
 - ▶ What is the size of the graph as a function of SALI cutoff?
- ▶ The SALI approach allows us to investigate molecular representations that may not be directly accessible
- ▶ Work in progress

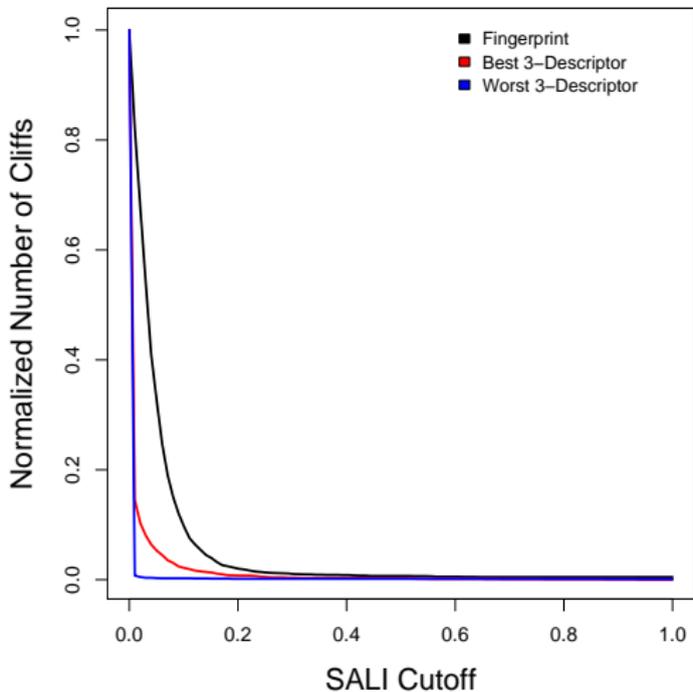
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Comparing Landscapes



A Type 2 SALI curve for the PDGFR dataset, comparing 3 different molecular representations

What's Next?

- ▶ SALI graphs and curves represent a *framework* for exploring structure-activity landscapes

Open questions

- ▶ Weighted SALI graphs (ADMET, synthetic feasibility)
- ▶ Is it correct to identify cliffs using fingerprints, and then predict cliffs using different descriptors?
- ▶ Can we use SALI curves to compare 3D and 2D descriptor spaces?
- ▶ Can we use SCI for feature selection?

Conclusions

- ▶ The SALI is an effective way to numerically encode activity cliffs
- ▶ The network view of these values allows us to explore SAR's in an intuitive way
- ▶ Using the SALI curve allows us to compare predictive models in a manner that is intuitive for a medicinal chemist

Acknowledgments

- ▶ John Van Drie

Defining & Using Structure-Activity Landscapes

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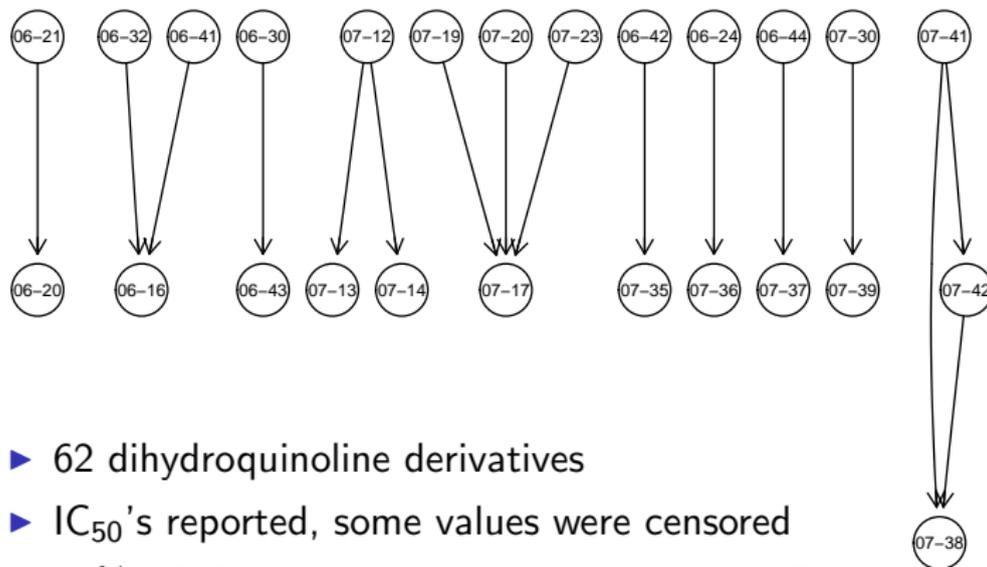
Chemical spaces

Summary

Making Use of the SALI Graph

- ▶ A little difficult with a non-interactive graph
- ▶ We can investigate a series of transformations that increase (or decrease) activity
- ▶ Identify two types of SAR's
 - ▶ Broad
 - ▶ Detailed
 - ▶ Depends on what cutoff we choose
- ▶ These correspond somewhat to the continuous and discontinuous SAR's described by Peltason et al.

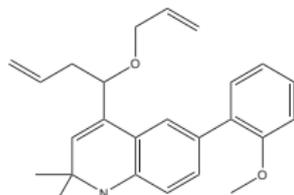
Glucocorticoid Inhibitors



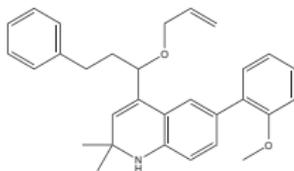
- ▶ 62 dihydroquinoline derivatives
- ▶ IC_{50} 's reported, some values were censored
- ▶ 50% SALI graph generated using 1052 bit BCI fingerprints

Glucocorticoid Inhibitors

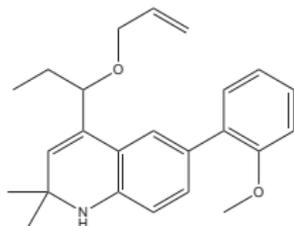
- ▶ Moving from ally or phenylethyl to ethyl causes a 6-fold increase in activity
- ▶ Reducing bulk at this position seems to improve activity
 - ▶ Pretty broad conclusion
- ▶ But ethyl is not much smaller than allyl
- ▶ We need more detail



07-20, 2000 nM

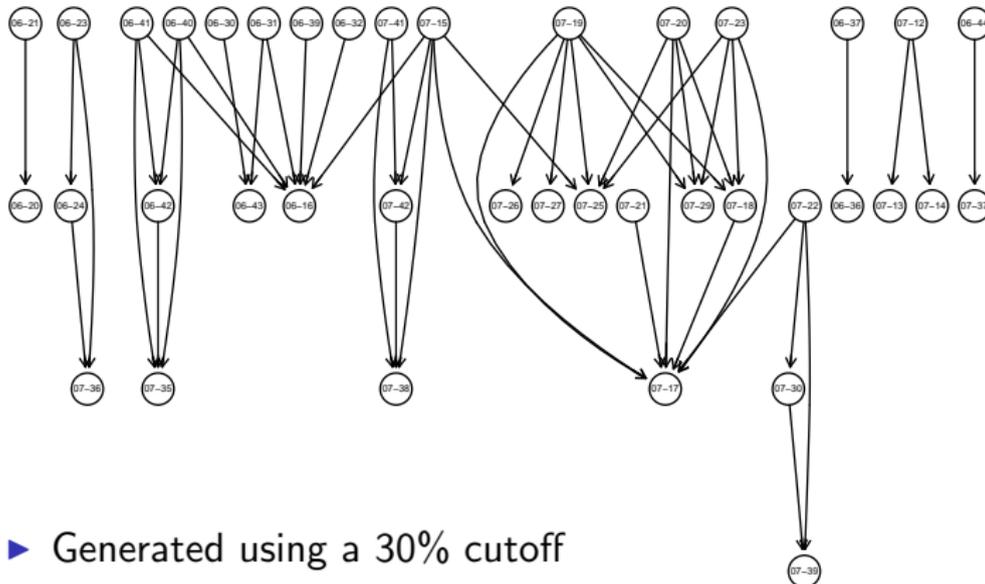


07-23, 2000 nM



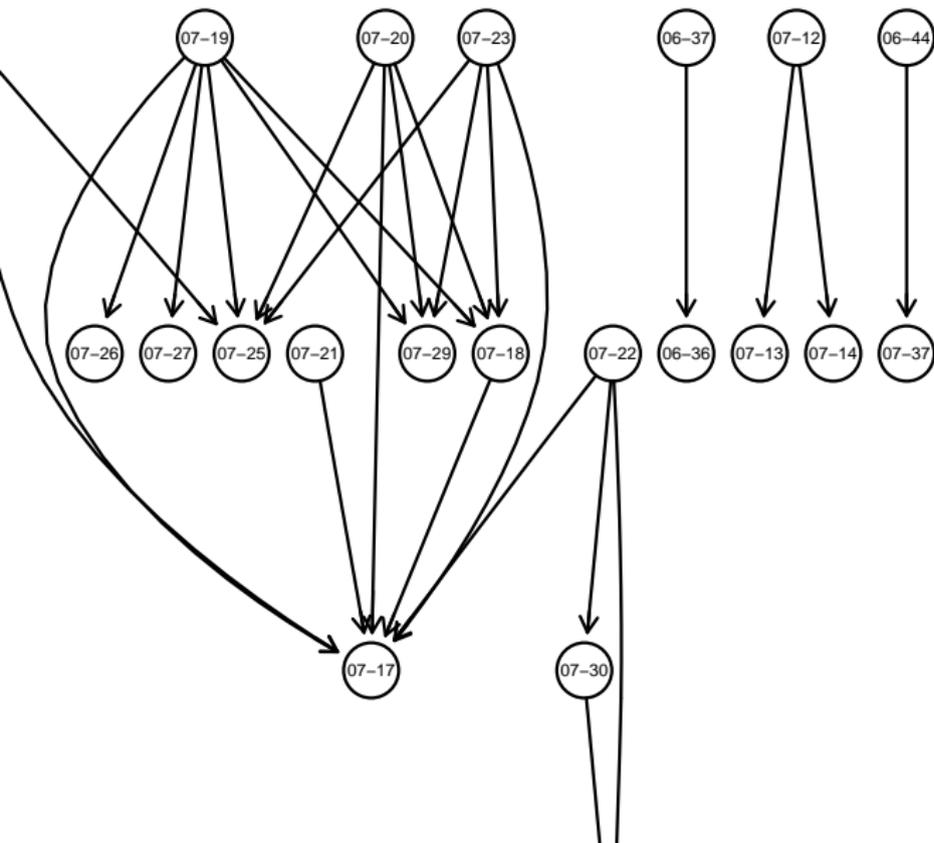
07-17, 355 nM

Glucocorticoid Inhibitors

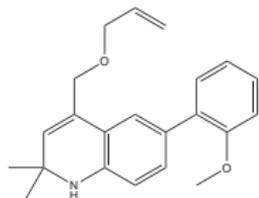


► Generated using a 30% cutoff

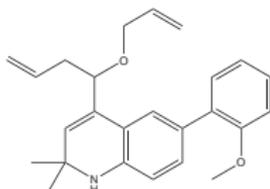
Glucocorticoid Inhibitors



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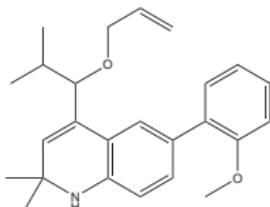


07-15, 2000 nM

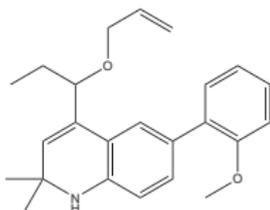


07-20, 2000 nM

- ▶ Suggests that electron density is also important
- ▶ Lower π density possibly correlates to increased activity
- ▶ Confirmed by **07-23** \rightarrow **07-18**
- ▶ **07-15** \rightarrow **07-17** is interesting since the change *increases* the bulk



07-18, 710 nM

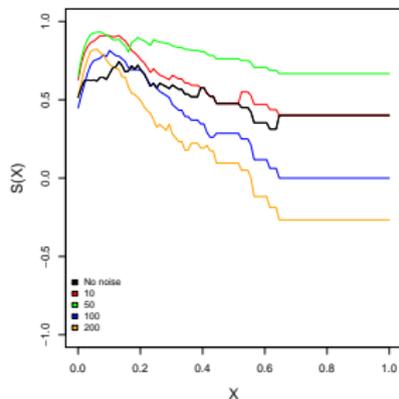
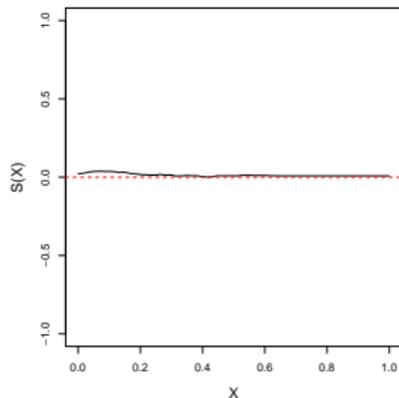


07-17, 355 nM

Glucocorticoid Inhibitors

- ▶ These observations match those made by Takahashi et al.
- ▶ More detailed graphs exhibit longer paths that focus on the bulk of side chains at the C4- α position
- ▶ A number of paths consider changes to the epoxide substitution
 - ▶ Usually of length 1
 - ▶ Highlights the fact that bulk at the C4 α has greater impact on activity than epoxide substitutions
- ▶ The SALI graph stresses the non-linearity of the SAR

SALI Curves - Control Experiments



Scrambling

- ▶ Scramble the Y-variable and rebuild the model
- ▶ Evaluate the SALI curve
- ▶ Repeat 50 times and take the mean of the counts for a given cutoff

Noise

- ▶ Add uniform noise to each descriptor, rebuild the model
- ▶ We expect little variation in the plateau