



UNC
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Applications of machine learning and artificial intelligence to designing chemicals and materials with the desired properties

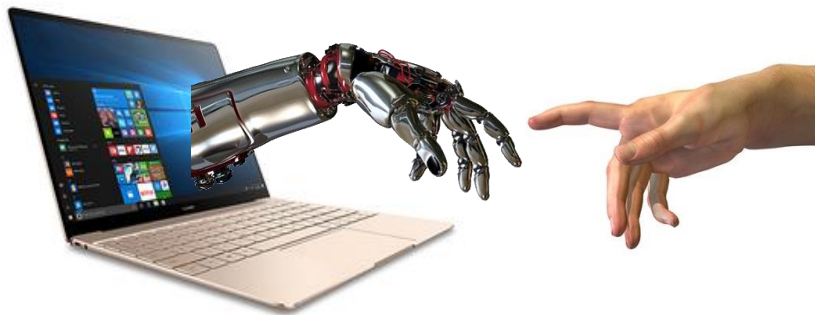
Alexander Tropsha

UNC Eshelman School of
Pharmacy

Outline

- Brief notes on machine learning/QSAR
- Materials Informatics and Materials Design
- Design, development and application of the Reinforcement Learning for Structural Evolution (ReLeaSE)*
- Summary and future work: QSAR without borders

Machine Learning Framework

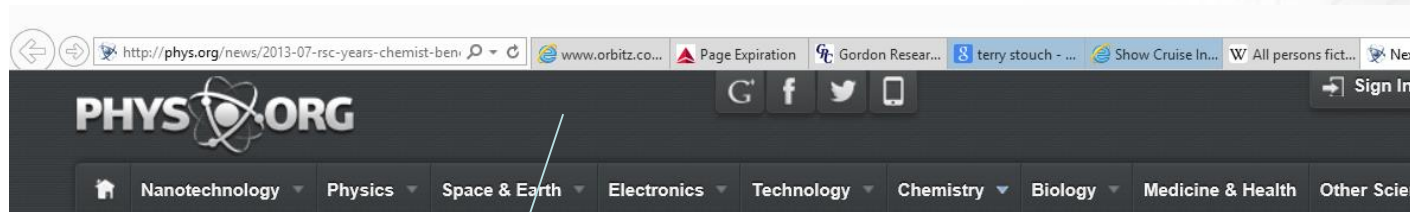


$$y = f(\mathbf{x})$$

output prediction function Molecular features

- **Training:** given a *training set* of labeled examples $\{(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)\}$, estimate the prediction function f by minimizing the prediction error on the training set
- **Testing:** apply f to a never before seen *test example* \mathbf{x} and output the predicted value $y = f(\mathbf{x})$

The growing appreciation of molecular modeling and informatics



Home > Chemistry > Materials Science > July 17, 2013

Next RSC president predicts that in 15 years no chemist will do bench experiments without computer-modelling them first

Jul 17, 2013

The newly-appointed President-Elect of the Royal Society of Chemistry today forecast the impact of advances in modelling and computational informatics on chemistry



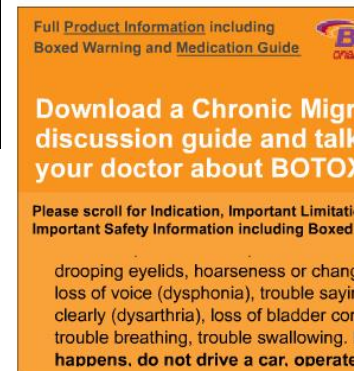
LIBERTY
UNIVERSITY
ONLINE

Christian counselors are needed to guide people through the toughest times of their lives.

Will you answer the call?

Professor Dominic Tildesley, who will become president in 2014, said: "The speed and development of computers is now so rapid, and the advances in modelling and informatics are so dramatic that in 15 years' time, no chemist will be doing any experiments at the bench without trying to model them first."

Professor Tildesley is a world-leading expert in large-scale computational modelling and



Full Product Information including Boxed Warning and Medication Guide

Download a Chronic Migraine discussion guide and talk your doctor about BOTOX

Please scroll for Indication, Important Limitations, Important Safety Information including Boxed Warning

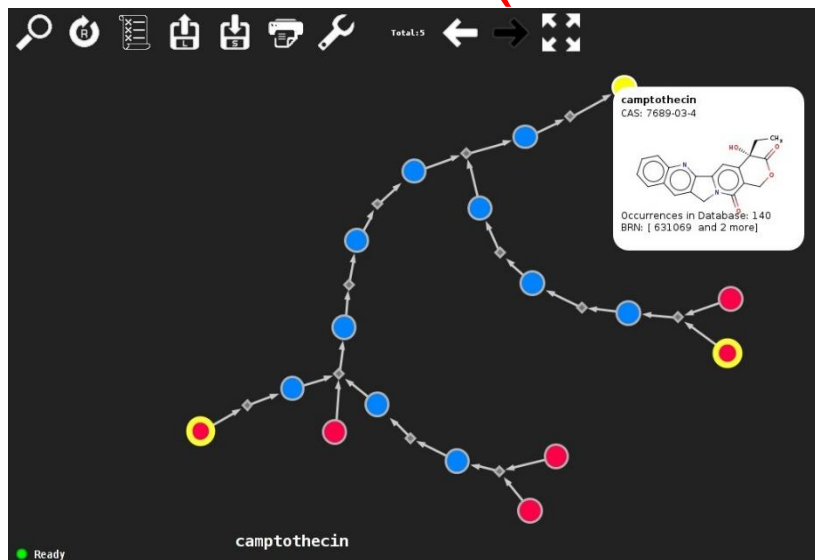
drooping eyelids, hoarseness or change in voice (dysphonia), trouble saying words clearly (dysarthria), loss of bladder control, trouble breathing, trouble swallowing. If these symptoms happen, do not drive a car, operate

4

Featured

Popular

Automated Retrosynthesis (Chematica)



Chem

Volume 4, Issue 3, 8 March 2018, Pages 390–398

CellPress

Backstory

Chematica: A Story of Computer Code That Started to Think like a Chemist

Bartosz A. Grzybowski, Sara Szymkuć, Ewa P. Gajewska, Karol Molga, Piotr Dittwald, Agnieszka Wolos, Tomasz Klucznik

[Show more](#)

<https://doi.org/10.1016/j.chempr.2018.02.024>

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Refers To

Tomasz Klucznik, Barbara Mikulak-Klucznik, Michael P. McCormack, Heather Lima, Sara Szymkuć, Manishabrata Bhowmick, Karol Molga, Yubai Zhou, Lindsey Rickershauser, Ewa P. Gajewska, Alexei Touthkine, Piotr Dittwald, Michał P. Starlek, Gregory J. Kirkovits, Rafał Roszak, Ariel Adamski, Bianka Sieredzińska, Milan Mrksich, Sarah L.J. Trice, Bartosz A. Grzybowski
Efficient Syntheses of Diverse, Medically Relevant Targets Planned by Computer and Executed in the Laboratory

Chem, Volume 4, Issue 3, 8 March 2018, Pages 522–532

[PDF \(2659 K\)](#) | [Supplementary content](#)

The growing appreciation of molecular modeling and informatics

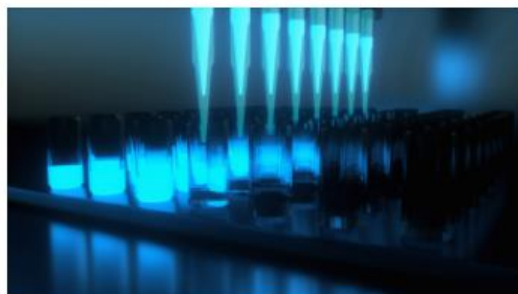
THE ROYAL SOCIETY OF CHEMISTRY [GB] | <https://www.chemistryworld.com/news/wanted-synthetic-chemists-humans-need-not-apply/3008401.article>

ToxPiWizard | ToxCast | file:///C:/Users/atrop... | Google | New Tab | Save to Mendeley | Savings & Investment | wtharvey.com/m8n2... | Серил Лист ожида... | GS_5

NEWS OPINION MATTER ENERGY EARTH LIFE CULTURE CAREERS PODCASTS WEBINARS LONG READS

NEWS

A brave new world of robot chemists and 'synthesiser farms' awaits



NEWS

Wanted: synthetic chemists (humans need not apply)

24 JANUARY 2018

Automation could free chemists from tedious lab work – if they're ready to think differently about research

Promise of dramatic acceleration of drug discovery



Pharma**VOICE**.com

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News	Blog	R&D	Commercial	Operations			
Magazine	PharmaVOICE 100	Resources	Events	Editorial	Advertise	Subscribe	

GSK Has Developed A New Analytics Platform That Can Reduce The Time It Takes To Analyze Clinical Data From Months To Clicks

Source:

Thomas Macaulay, CIO UK

March 12, 2018

The platform uses large-scale data analytics to drive better decisions about the drug discovery pipeline, by allowing the pharmaceuticals giant to test the potential for new drugs before it begins clinical trials.

Rise of the machines in legal industry



MML
UNC.EDU

legal  insider
www.legaltechnology.com



Deloitte Insight: Over 100,000 legal roles to be automated

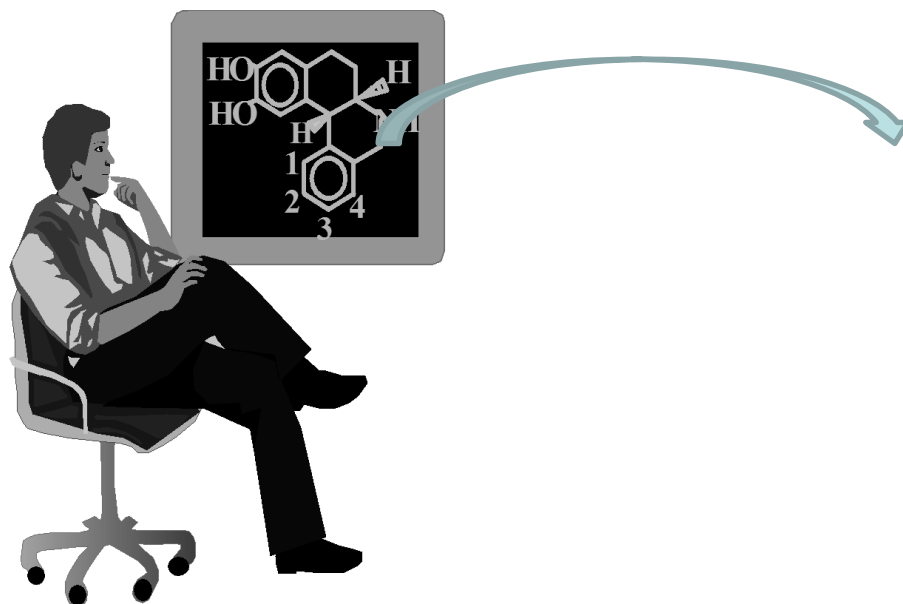
Added on the 16th Mar 2016 at 10:28 am



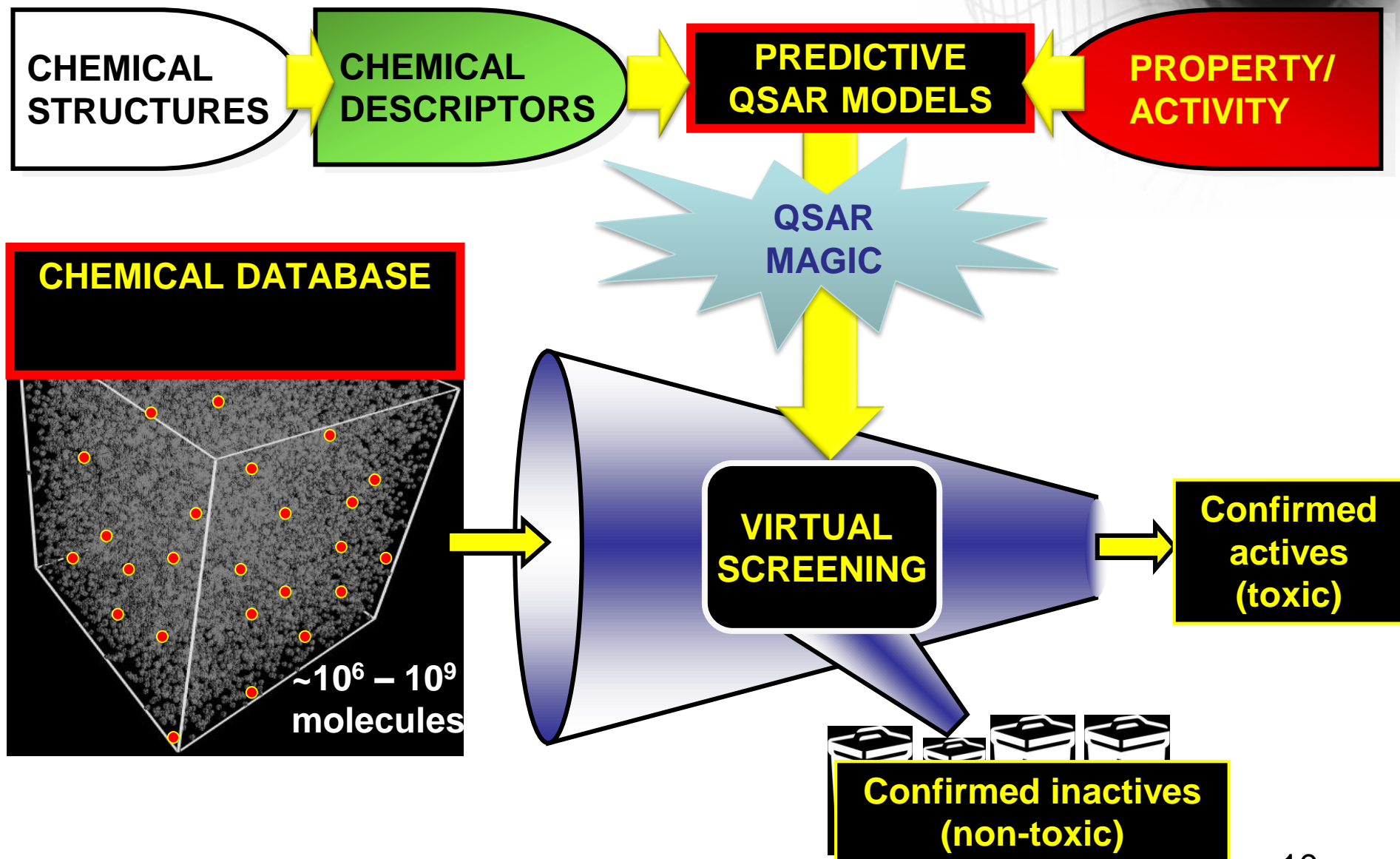
Over 100,000 jobs in the legal sector have a high chance of being automated in the next twenty years, according to extensive new analysis by Deloitte.

The Deloitte Insight report, which predicts “profound reforms” across the legal profession within the next 10 years, finds that 39% of jobs (114,000) in the legal sector stand to be automated in the longer term as the profession feels the impact of more “radical changes.”

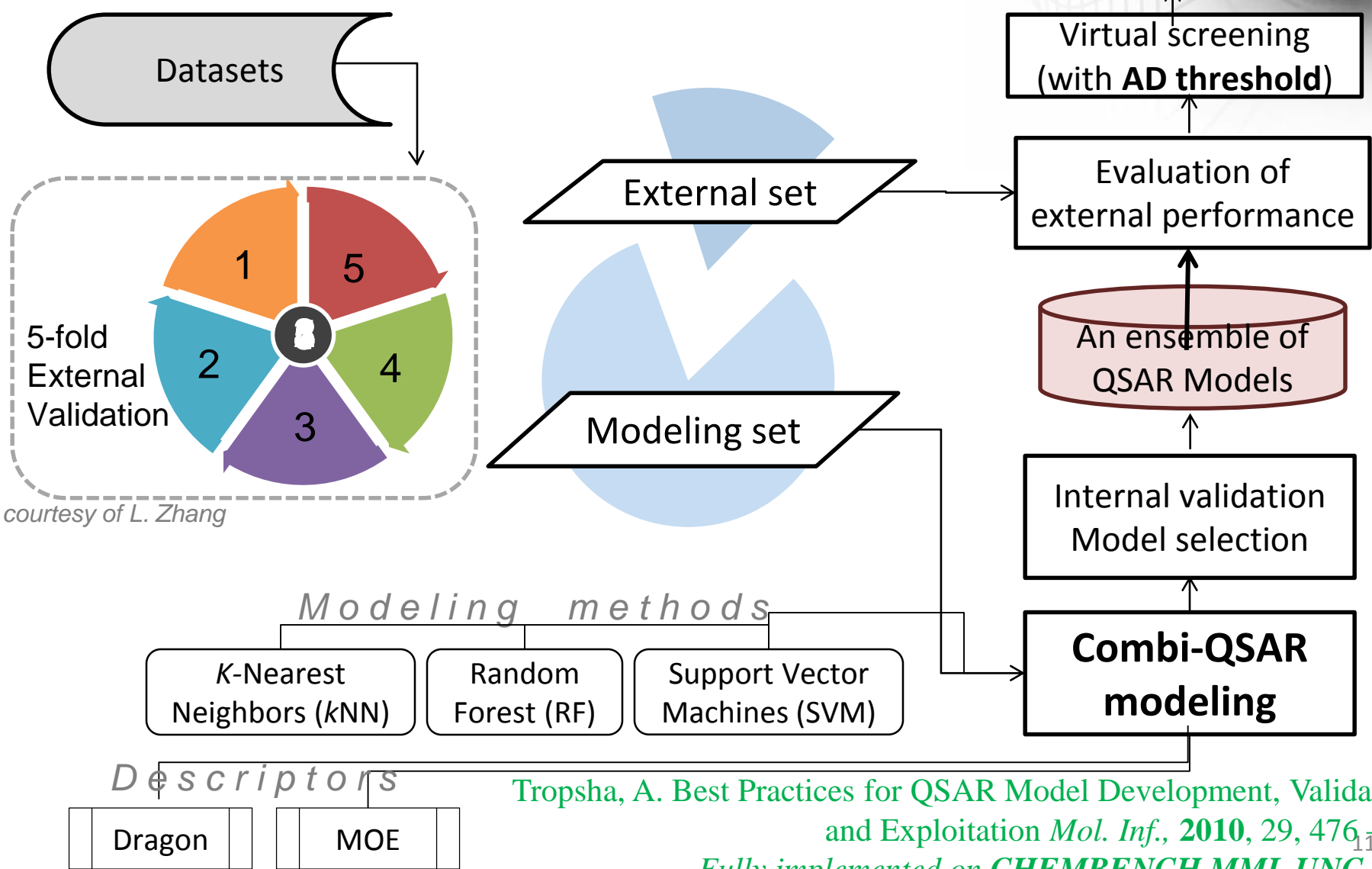
The ultimate dream of a computational chemist



The chief utility of computational models: Annotation of new compounds



QSAR Modeling Workflow: the importance of rigorous validation

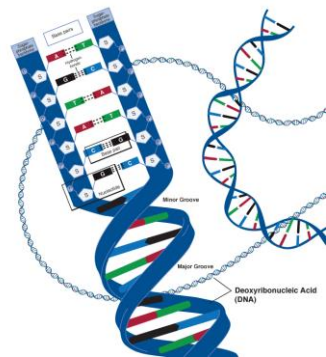


Tropsha, A. Best Practices for QSAR Model Development, Validation, and Exploitation *Mol. Inf.*, **2010**, 29, 476–488
Fully implemented on CHEMBENCH.MMLUNC.EDU

Material Science and the Rise of Materials Informatics



- Explosive growth of materials data, both experimental databases and computational repositories.
 - Structural data: 160,000 entries in the Inorganic Crystal Structure Database (ICSD)
 - Experimental data: Numerous commercial and open experimental databases NIST, MatWeb, MatBase etc.
 - Computational data: Huge databases such as AFLOWLIB, Materials Project, and Harvard Clean Energy
 - Chemical space of possible materials is HUGE $\sim 10^{100}$ candidates [*Nat. Chem.* 7, 274-275 (2015)]
- Materials Genome Initiative or MGI (US Govt): Need for new high performance materials

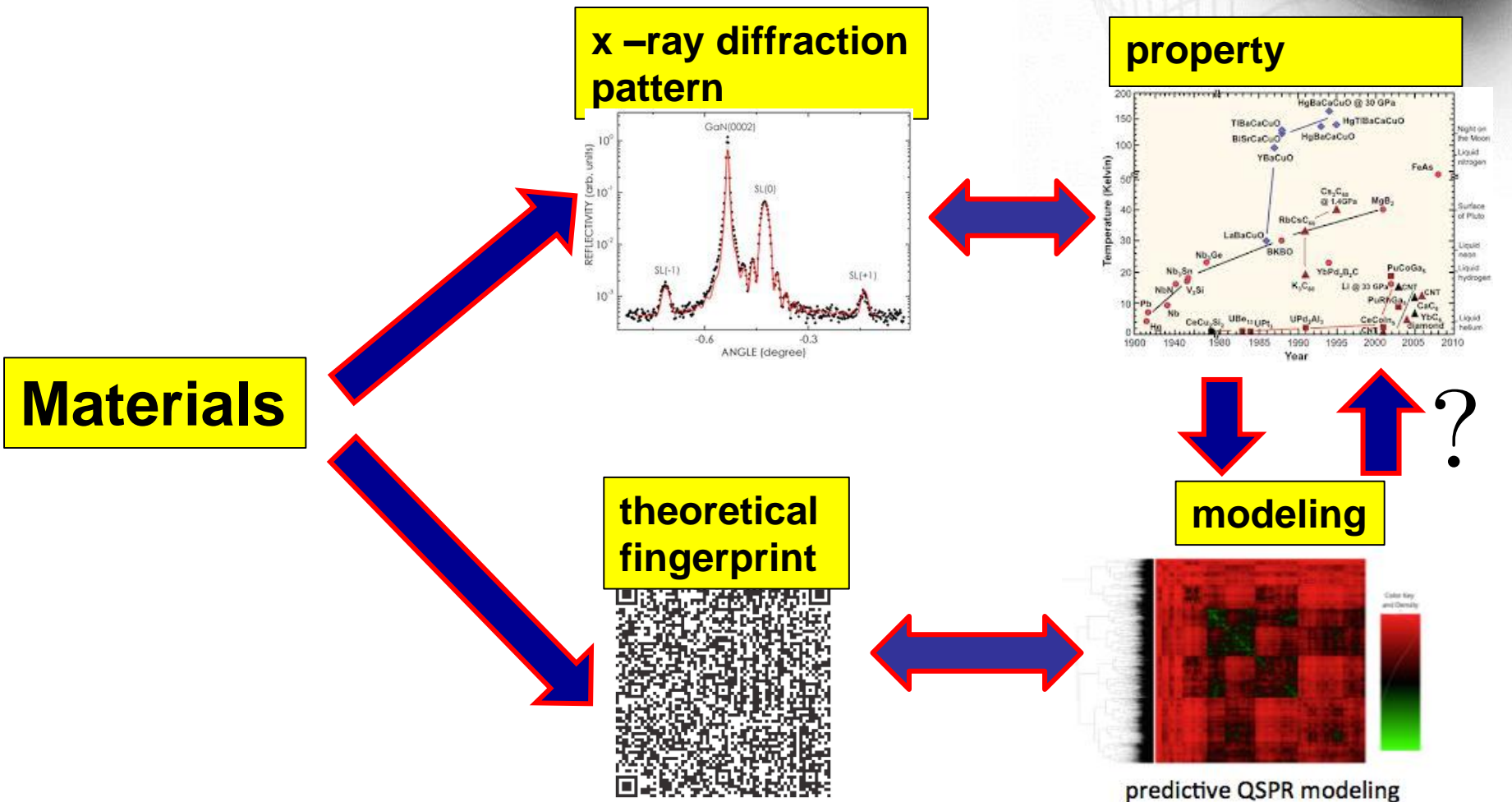


Periodic Table

For elements with no stable isotopes, the mass number of the isotope with the longest half-life is in parentheses.

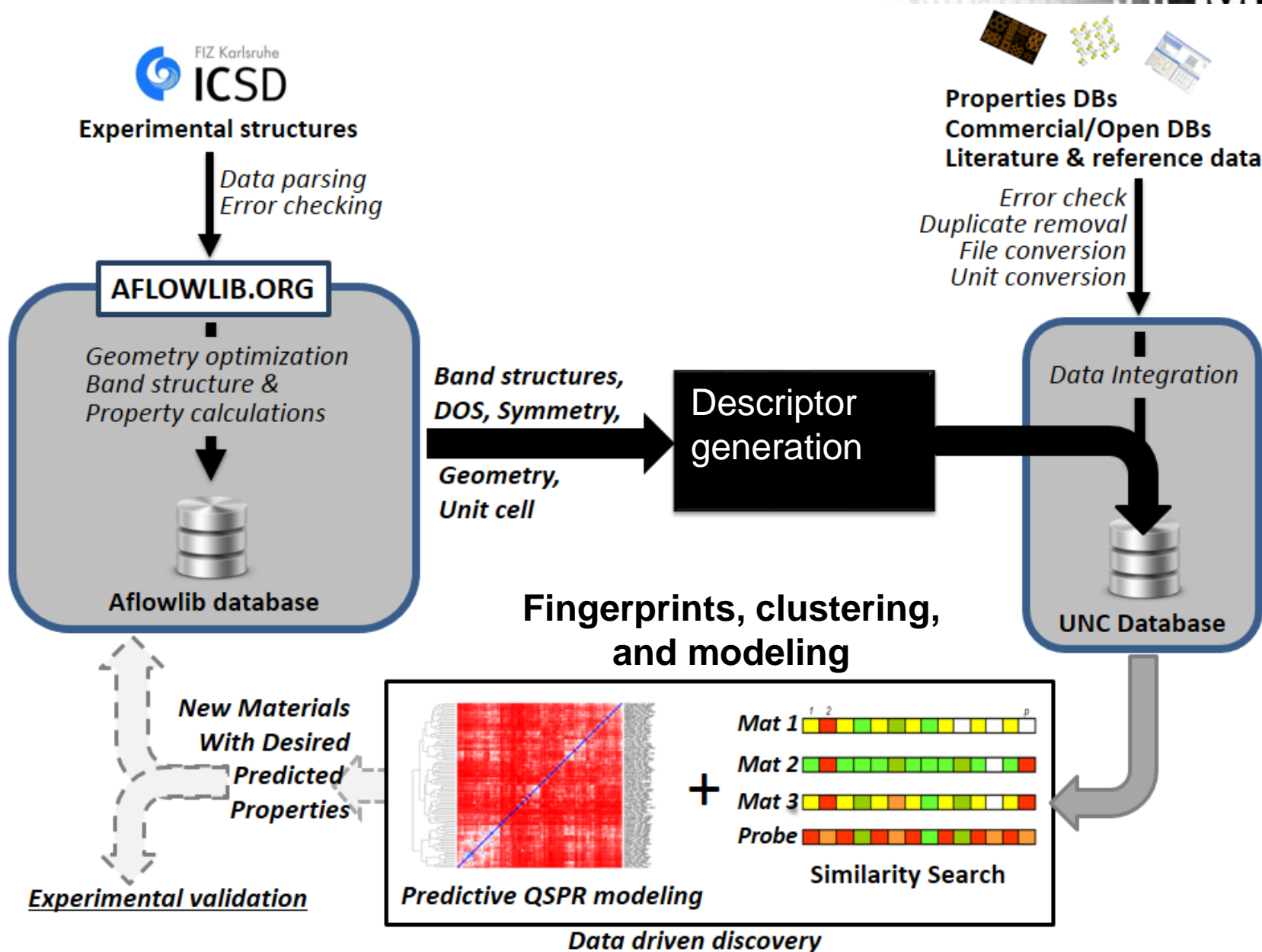
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	
1	H	Ds Darmstadtium (271) 2-8-18-32-32-17-1										He						
2	Li	Be	Ds										Ne					
3	Na	Mg	Ds										Ar					
4	K	Ca	Sc	Ti	V	Cr	Mn	Fe	Co	Ni	Cu	Zn	Ga	Ge	As	Se	Br	Kr
5	Rb	Sr	Y	Zr	Nb	Mo	Tc	Ru	Rh	Pd	Ag	Cd	In	Sn	Sb	Te	I	Xe
6	Cs	Ba	La-Lu	Hf	Ta	W	Re	Os	Ir	Pt	Au	Hg	Tl	Pb	Bi	Po	At	Rn
7	Fr	Ra	Ac-Lr	Rf	Db	Sg	Bh	Hs	Mt	Ds	Rg	Cn	Uut	Fl	Uup	Lv	Uus	Uuo
				La	Ce	Pr	Nd	Pm	Sm	Eu	Gd	Tb	Dy	Ho	Er	Tm	Yb	Lu
				Ac	Th	Pa	U	Np	Pu	Am	Cm	Bk	Cf	Es	Fm	Md	No	Lr

Closing the gap: materials structure-property relationships



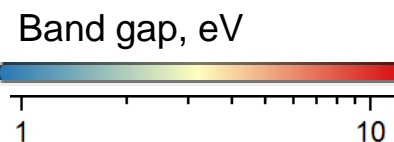
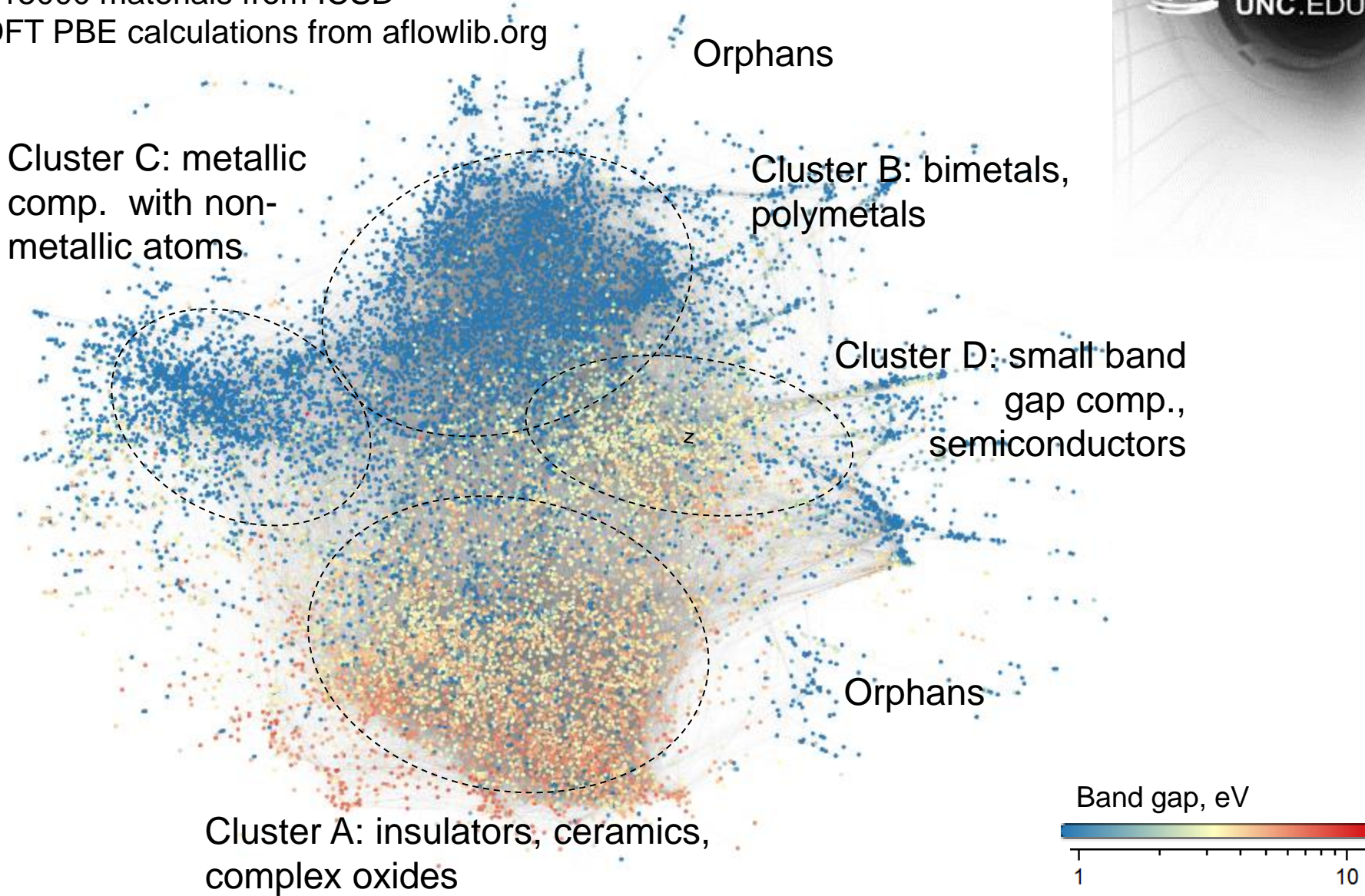
Quantitative Structure Activity Relationship approaches (QSAR)
Quantitative Structure-Property Relationships (QMSPR)

Material Informatics/MQSAR Workflow

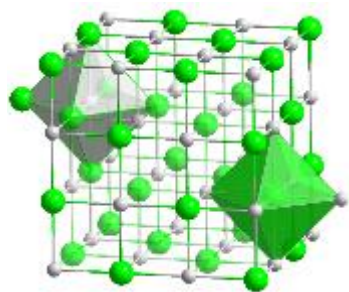


Material Map (B-Fingerprints)

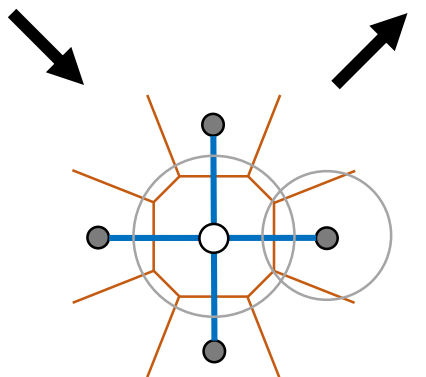
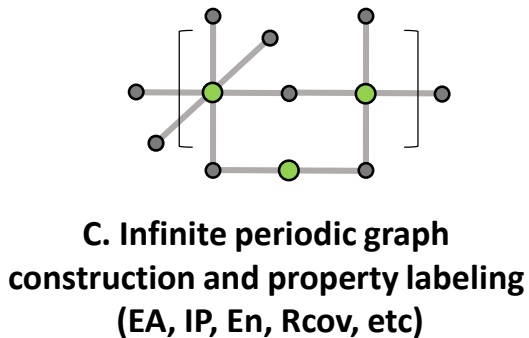
>15000 materials from ICSD
DFT PBE calculations from aflowlib.org



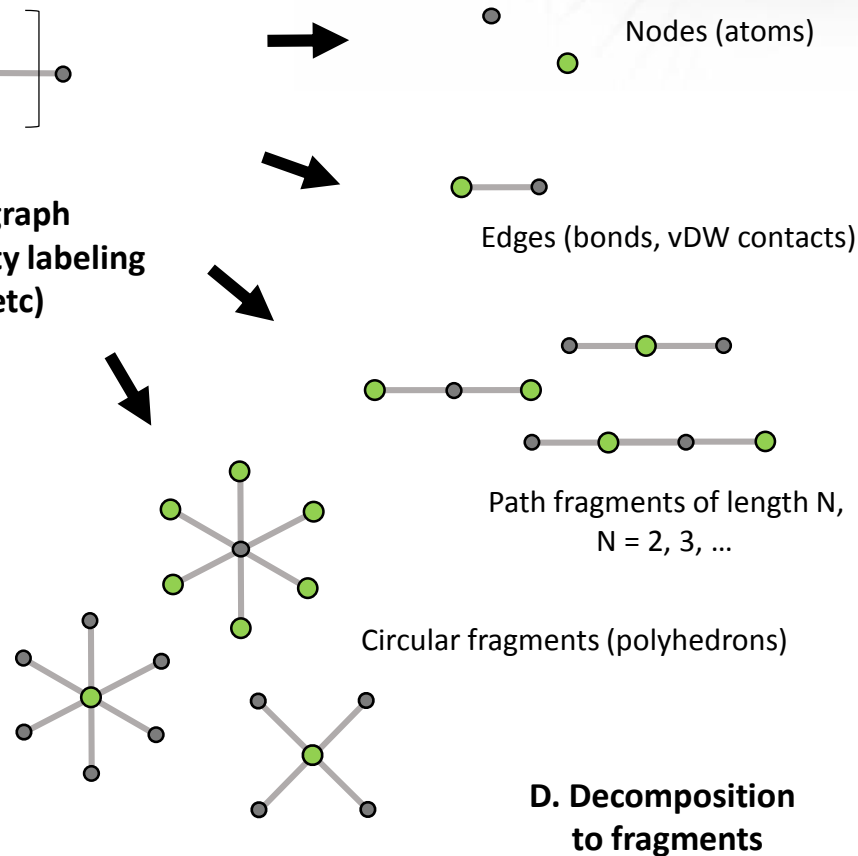
Systematic representation of materials using fragment descriptors



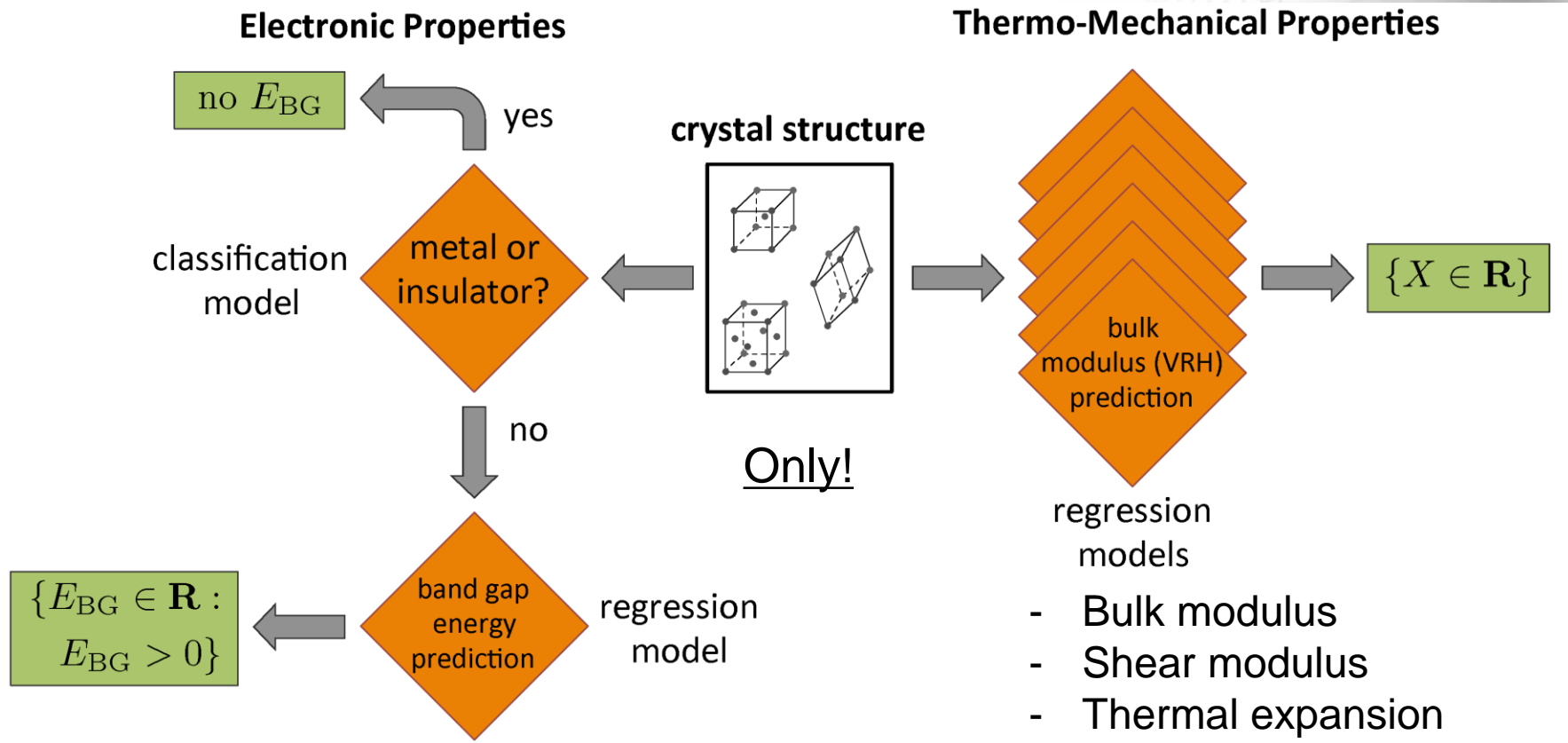
A. Crystal Structure



B. Voronoi tessellation and
neighbors search



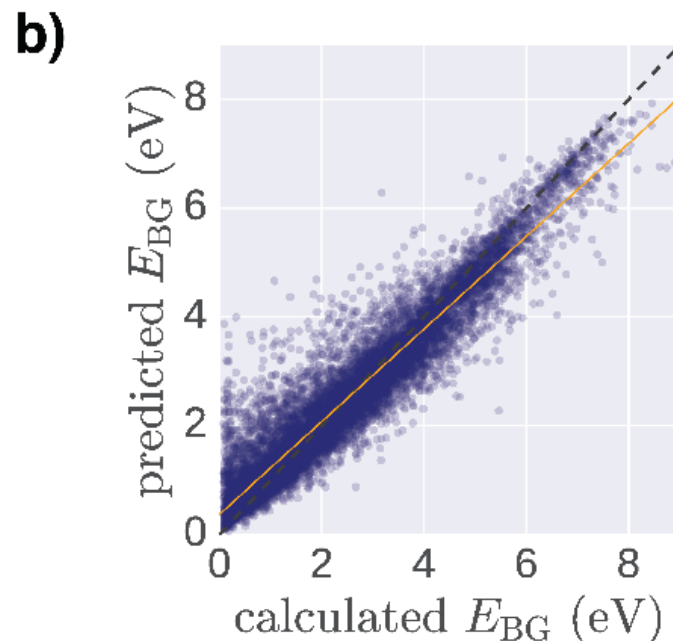
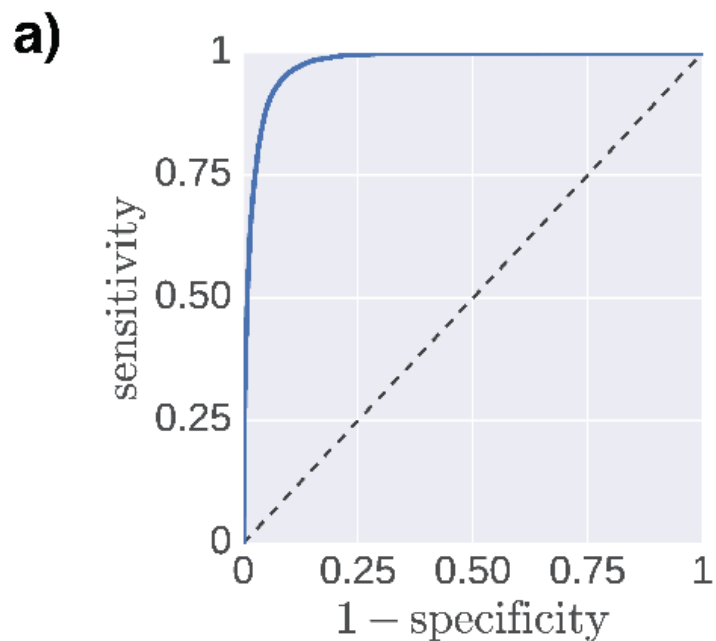
ML Workflow for Materials Property Prediction



All models are trained based on DFT-computed properties (VASP s/w from U. Vienna)

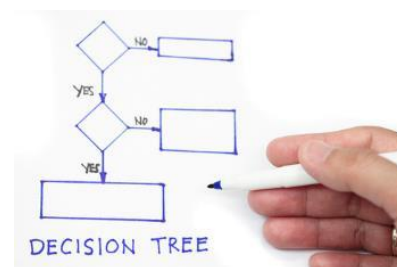
Prediction of Electronic Properties

electronic properties



Classification accuracy **95%**
ROC Curve (AUC) 0.98

**Learning approach for all models:
Gradient Boosting Decision Trees (GBT)**



Prediction of Thermomechanical Properties



(E_{BG} - band gap energy; B_{VRH} - bulk modulus; G_{VRH} - shear modulus; θ_D - Debye temperature; C_P - heat capacity at constant pressure; C_V - heat capacity at constant volume; α_V - thermal expansion coefficient)

property	RMSE	MAE	r^2
E_{BG}	0.51 (eV)	0.35 (eV)	0.90
B_{VRH}	14.25 (GPa)	8.68 (GPa)	0.97
G_{VRH}	18.43 (GPa)	10.62 (GPa)	0.88
θ_D	56.97 (K)	35.86 (K)	0.95
C_P	2.31 (k_B /cell)	0.84 (k_B /cell)	0.99
C_V	2.01 (k_B /cell)	0.70 (k_B /cell)	0.99
α_V	1.47×10^{-5} (K) $^{-1}$	5.69×10^{-6} (K) $^{-1}$	0.91

TABLE I. Statistical summary of the *five-fold cross-validated predictions* for the seven regression models (Figure 3).

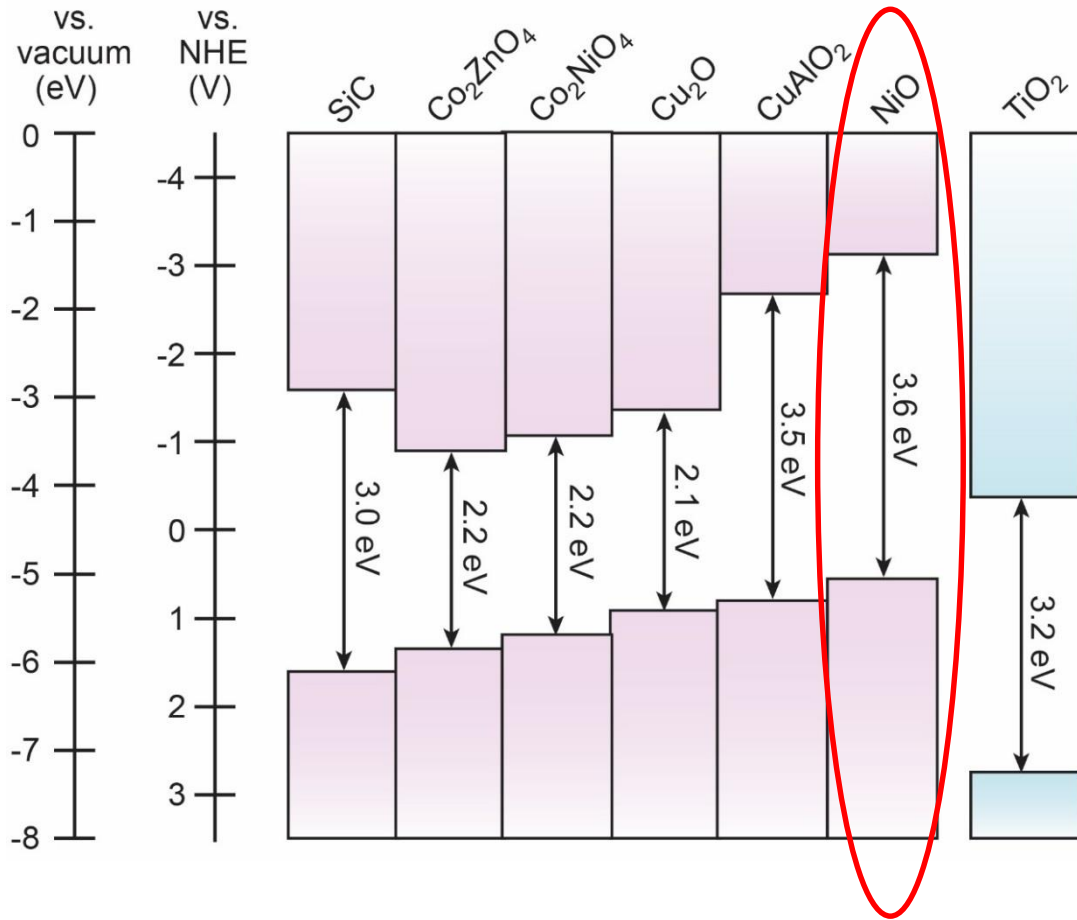
Summary of Materials Informatics: Methods



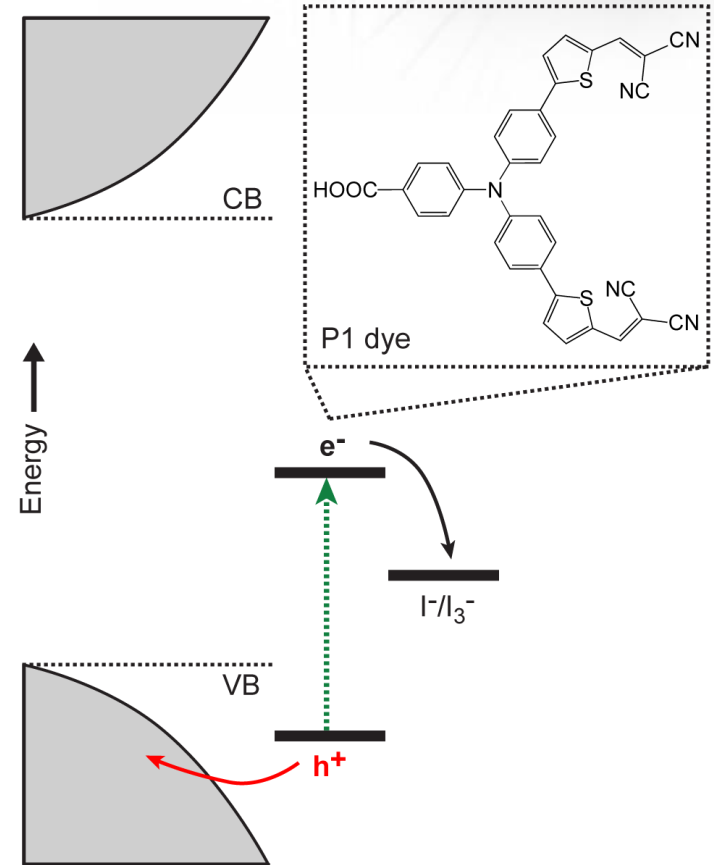
- Fast, accurate general purpose machine learning methods for material's property prediction. **Milliseconds on laptop vs. days on HPC cluster**
- Universal applicability to different materials: currently covered **85 elements** (H – Pu, without noble gases, Tc, Fr, Ra). All types of crystal lattices and symmetries.
 - Most competing approaches are specific to one prototype/family of materials or single property
- Works for other properties: elastic, thermoelectric, etc.
- Possible to gain *some* chemically/physically interpretable insight into “black box” model.
- Possible to derive materials design rules
- User friendly web app and RESTful API (<http://aflow.org/aflow-ml/>)

Photocathode materials

Evaluated as DSSCs

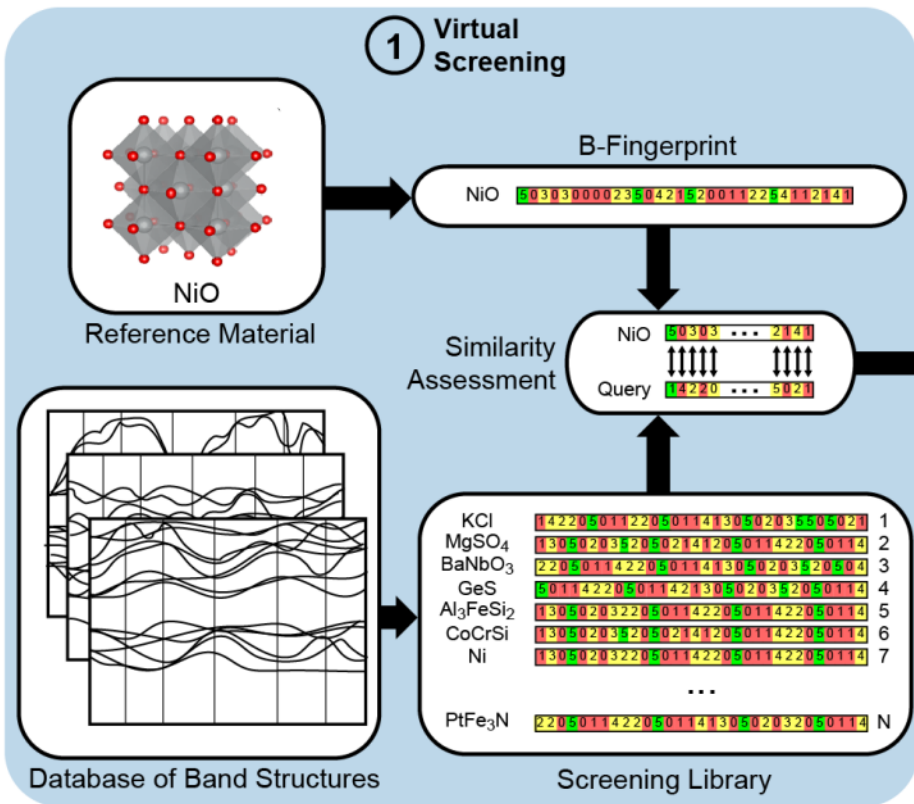


Dye-sensitized solar cells (DSSCs)



Design of alternate photocathodes

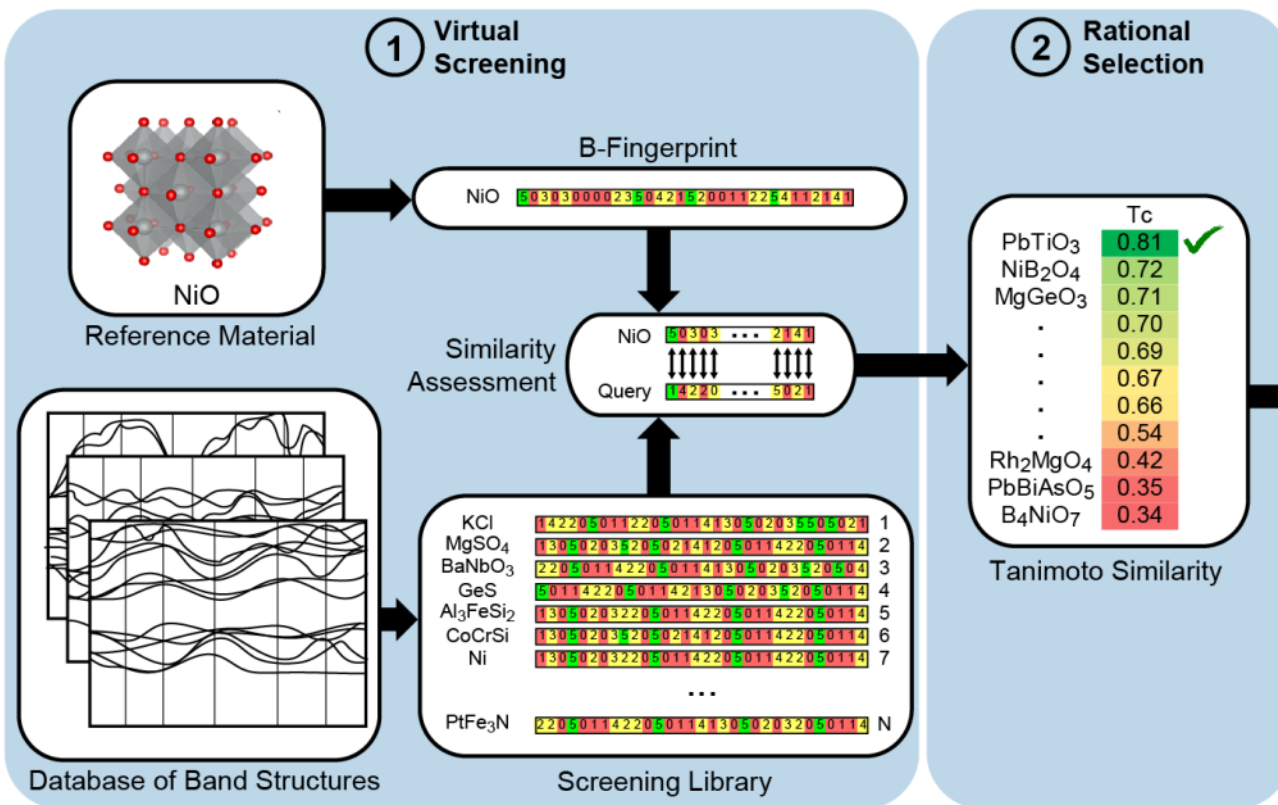
A materials informatics approach



(AFLOWLIB)

Design of alternate photocathodes

A materials informatics approach



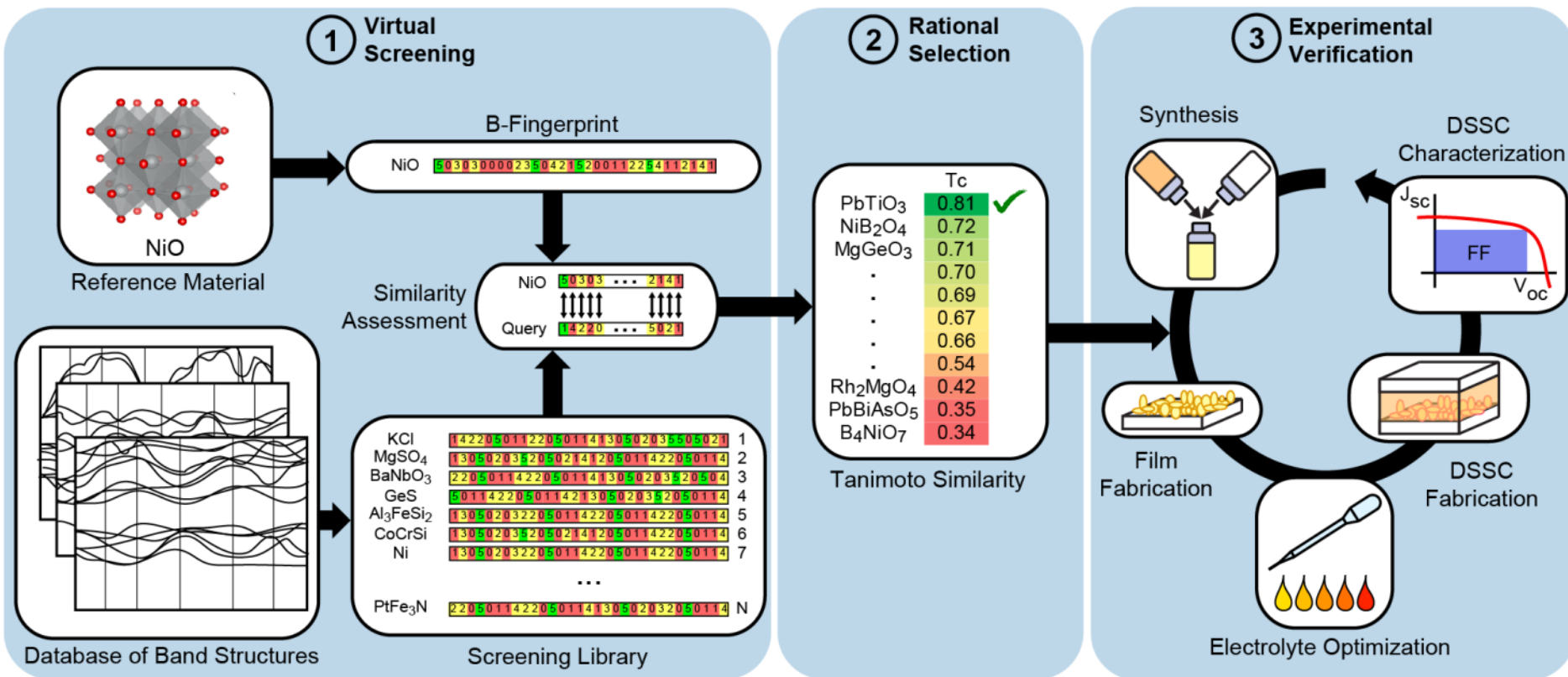
(AFLOWLIB)

PbTiO₃ was identified as very similar to NiO
AND

It has a dielectric constant > 100

Design of alternate photocathodes

A materials informatics approach



(AFLOWLIB)

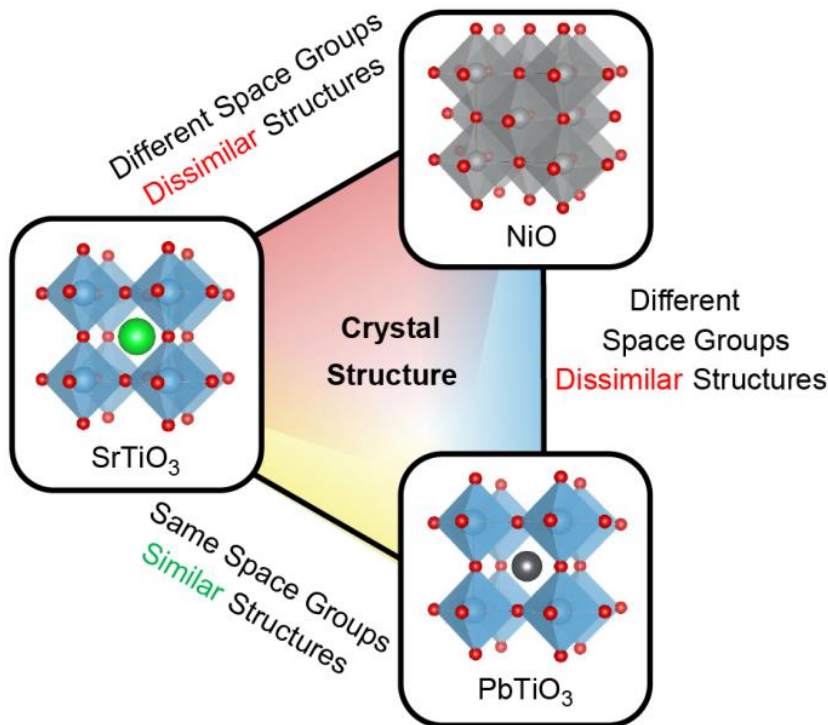
PbTiO₃ was identified as very similar to NiO
AND

It has a dielectric constant > 100

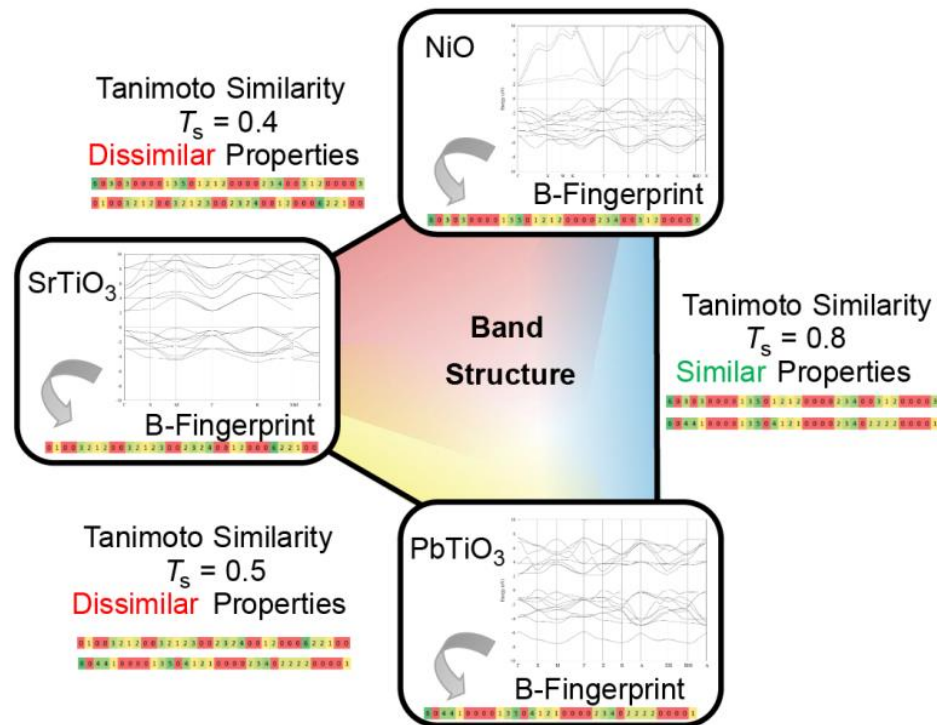
Materials informatics

Identifying top hit: PbTiO_3

Structure



Properties



PbTiO_3 was identified as very similar to NiO in terms of electronic properties despite different crystal structures

Summary of Materials Informatics: Supporting Experimental Discovery

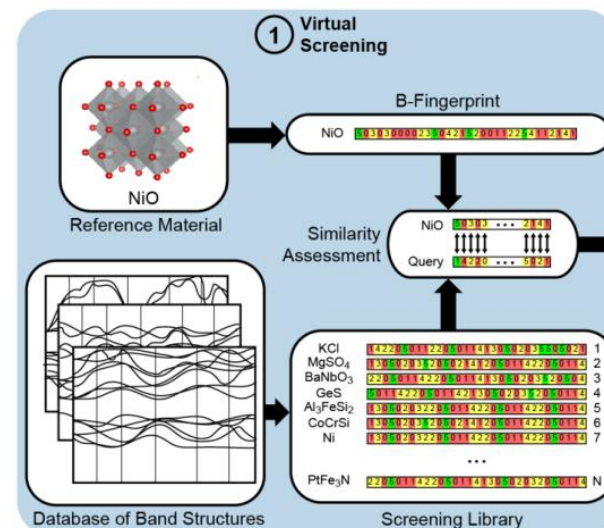
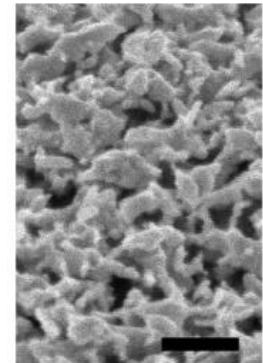
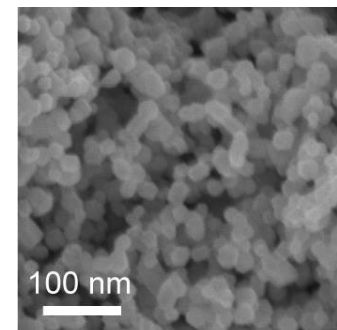
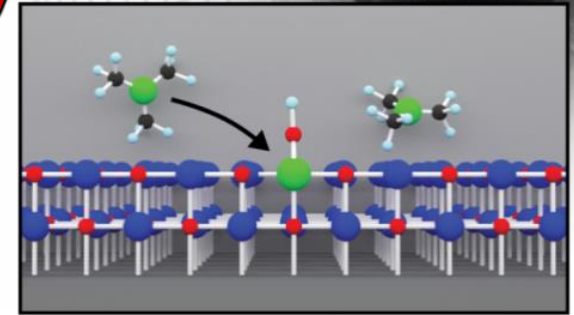
PbTiO₃ is identified as a new photocathode material.

Successful experimental validation

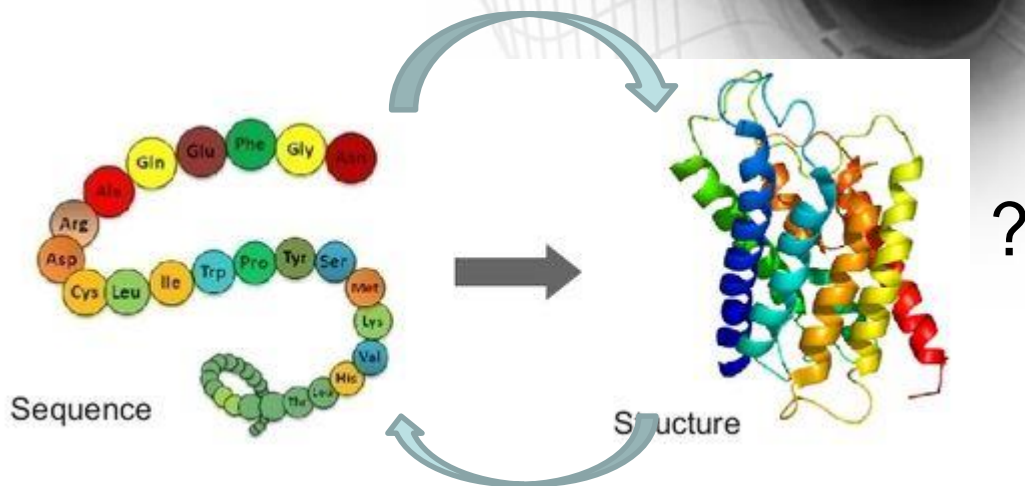
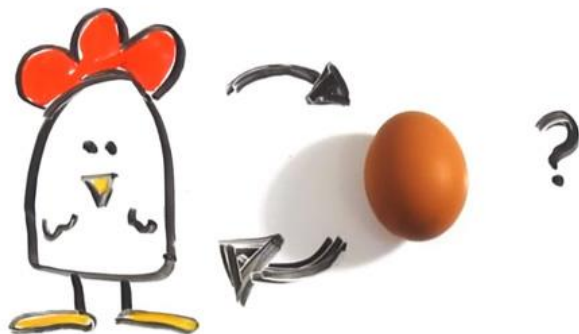
Record fill factors of >50

First fully aqueous DSSC device

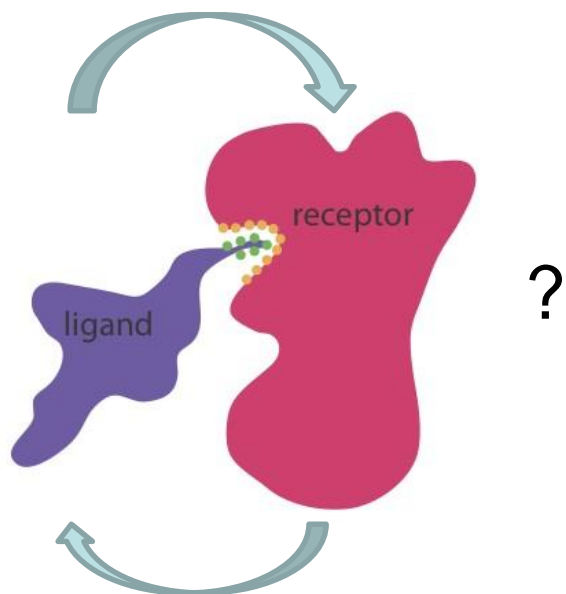
Currently, device performance is low;
possible improvement by designing a
new dye



The eternal philosophical question: Which came first?



R or R² ?



Strasbourg Summer
School in
Chemoinformatics -
2018



The eternal question: Which came first?



In the beginning was the Word...

And the Word was... **embedded**

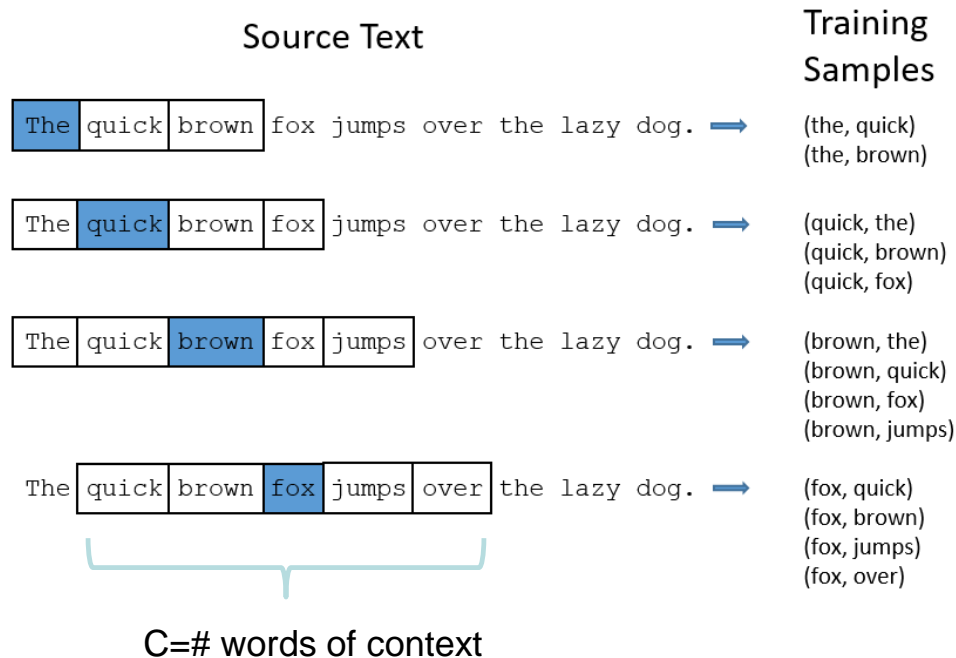
(freely adopted from the Gospel of John)

“You should know a word by the company it keeps”
J.R.Firth 1957



British linguist; formulated the notion of the “context-dependent nature of meaning”

Learning semantic context with Word2Vec



Can be used to learn:

CBOW:

- $Pr(word_k | words_context)$

Skip-Gram:

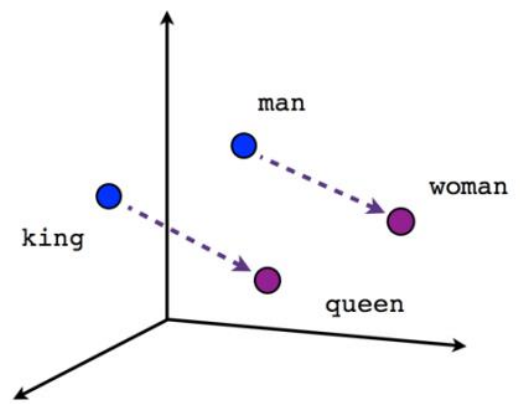
- $Pr(words_context | word_k)$

- Mikolov, Tomas; et al. "Efficient Estimation of Word Representations in Vector Space". [arXiv:1301.3781](https://arxiv.org/abs/1301.3781)

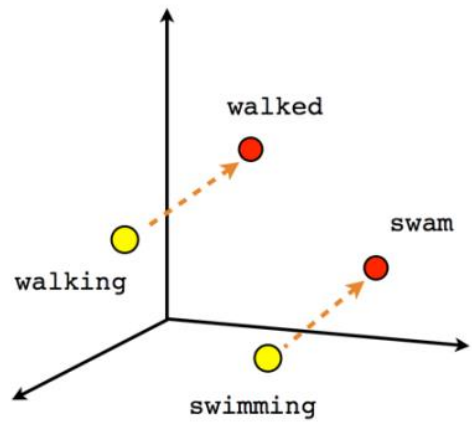
Word2Vec Images courtesy of Chris McCormick:

<http://mccormickml.com/2016/04/19/word2vec-tutorial-the-skip-gram-model/>

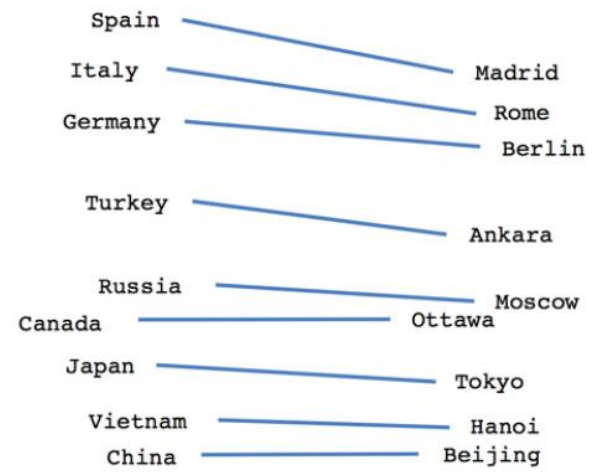
Word embedding and similarity in the semantic space



Male-Female



Verb tense



Country-Capital

SMILES are words that uniquely describe sentence-molecules!

Aspirin, also known as O=C(C)Oc1ccccc1C(=O)O, is a medication used to treat pain, fever, and inflammation.

CC(=C)C1(C=CC(=O)O1)O is a mycotoxin that is produced by *Aspergillus flavus* and *Penicillium roqueforti* mold.

C O M P O U N D S	<chem>O=C(C)Oc1ccccc1C(=O)O</chem>	Active	1	A C T I V I T Y
	<chem>CCOc1cc(C)ccc1OCC=CF</chem>	Inactive	0	
	<chem>COc1ccccc1OCCO</chem>	Inactive	0	
	<chem>CC(N)Sc1ccc(Cl)nc1</chem>	Inactive	0	
	<chem>COC(=O)NCc1ccccc1Cl</chem>	Active	1	

ReLeaSE* design principles: learning and exploiting structural linguistics of SMILES notation



- SMILES notations reflect rules of Chemistry
- SMILES notation embeds linguistic rules
- Neural nets could learn both of the above types of rules
- This knowledge can be transformed into the generation of new SMILES corresponding to novel chemically feasible molecules (generative model)
- One can build QSAR models based solely on SMILES notation (predictive model)
- QSAR models can be used as a reward function for reinforcement learning to bias the design of novel libraries

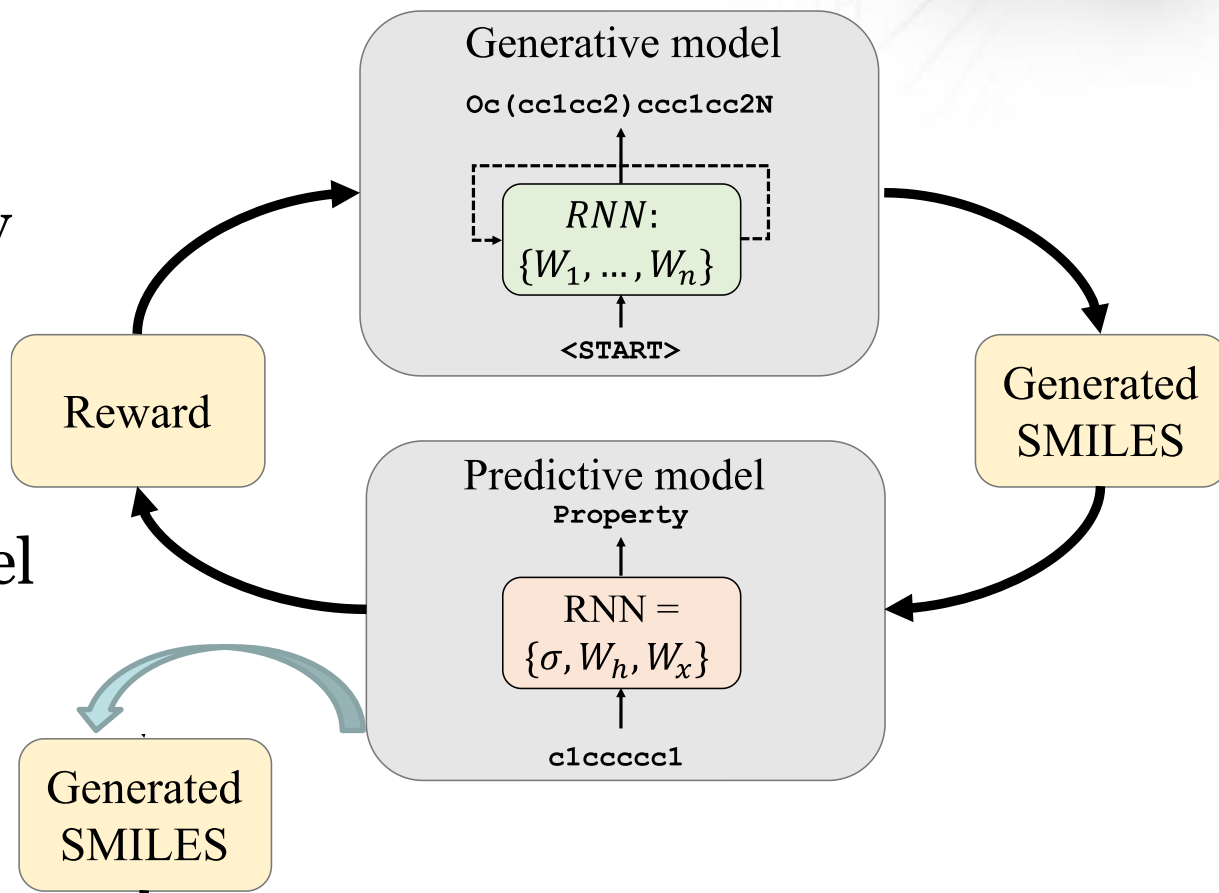
Design of the ReLeaSE* method

(Reinforcement Learning for Structural Evolution)



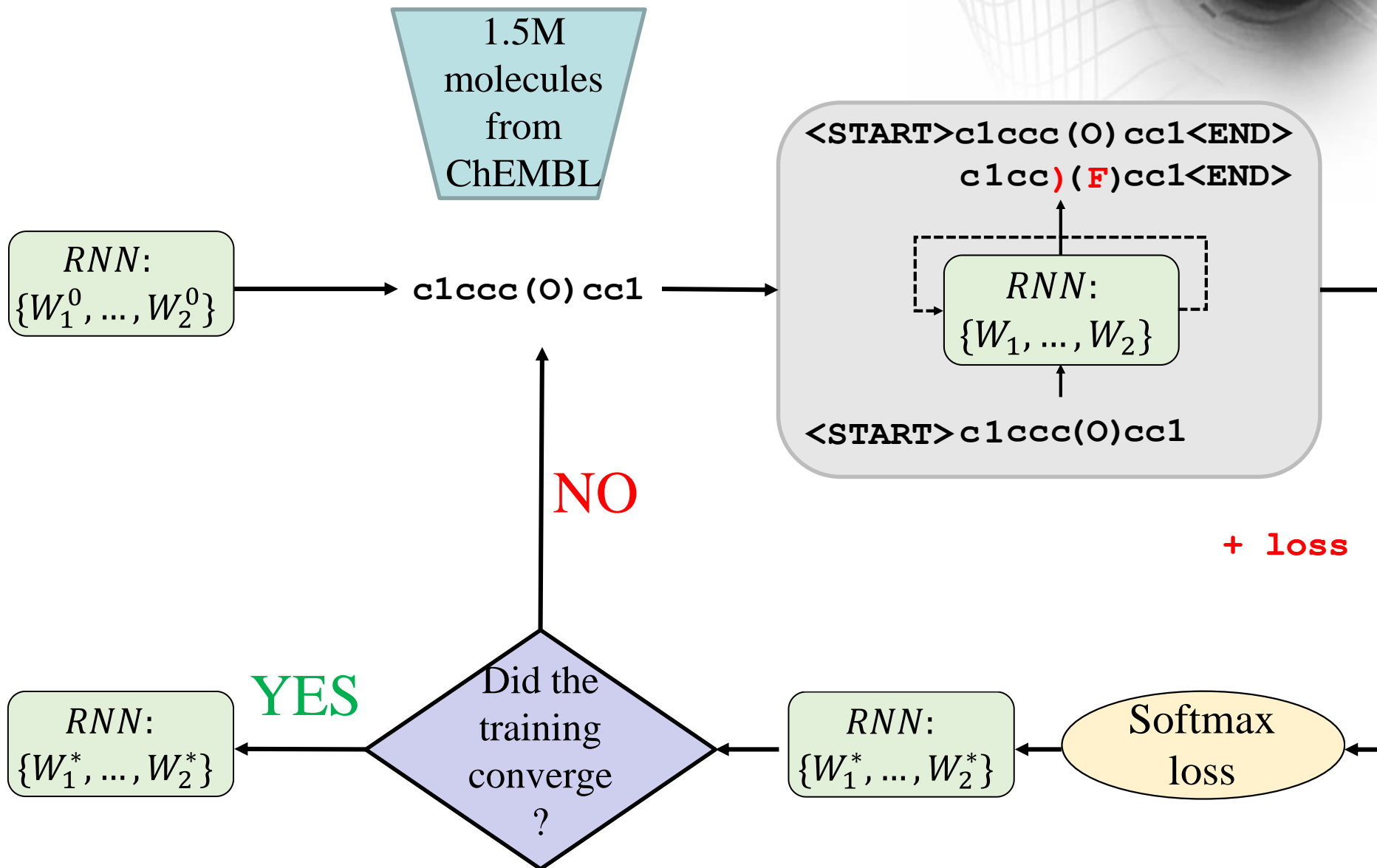
Elements of the thought cycle (molecules->models->molecules):

- Generate chemically feasible SMILES
- Develop SMILES-based QSAR model
- Employ QSAR model to bias library generation
- Produce new SMILES



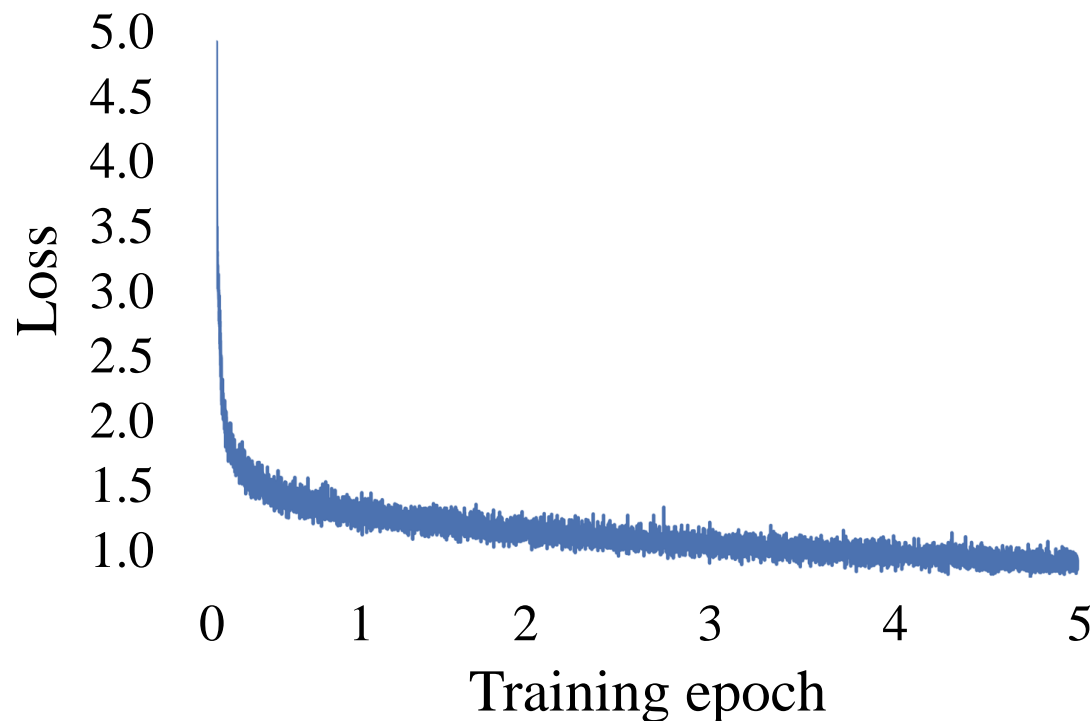
*Popova, Mariya, Olexandr Isayev, and Alexander Tropsha. "Deep reinforcement learning for de-novo drug design." *arXiv preprint arXiv:1711.10907* (2017); *Science Advances* (in press).

Generative model: training mode



Generative model: training mode

- Training continues until convergence
- Every SMILES from ChEMBL is used as training example ~ 3-5 times

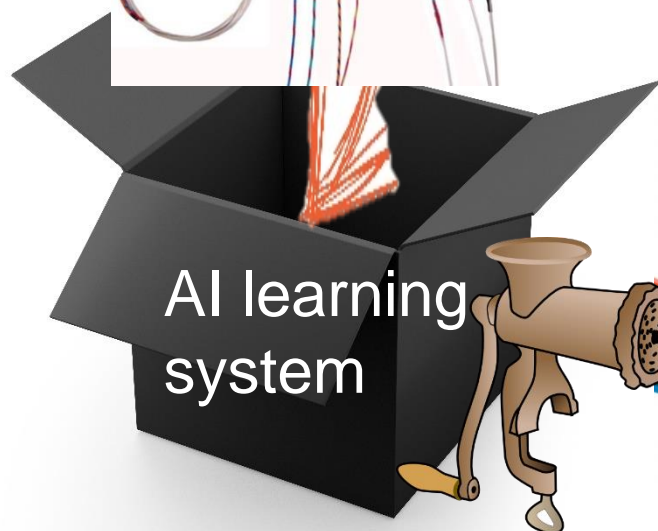


Are we making legitimate Smiles?

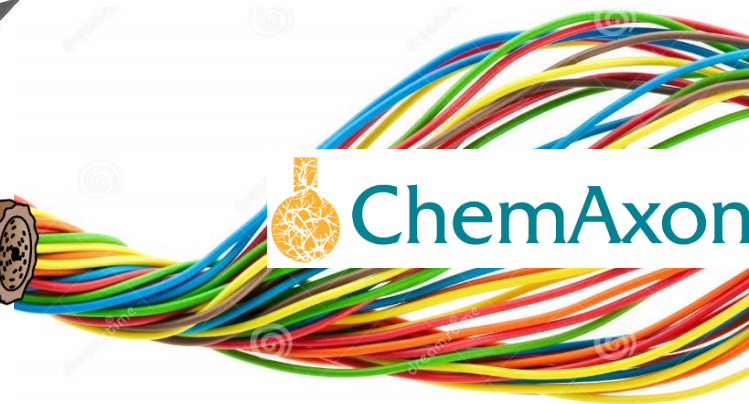
PubChem / ChEMBL



Smiles strings



AI learning
system



ChemAxon

95% Valid
Chemically-feasible
molecules

Smile-ification of QSAR!

C
O
M
P
O
U
N
D
S

O=C(C)Oc1ccccc1C(=O)O
CCOc1cc(C)ccc1OCC=CF
COc1ccccc1OCCO
CC(N)Sc1ccc(Cl)nc1
COC(=O)NCc1ccccc1Cl

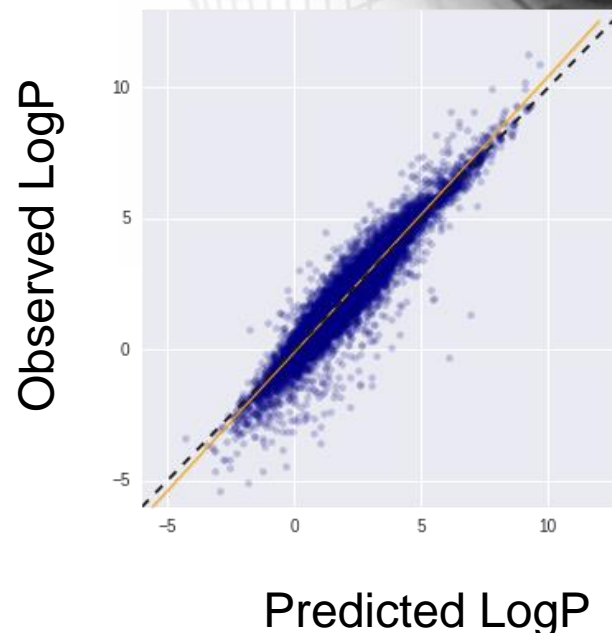
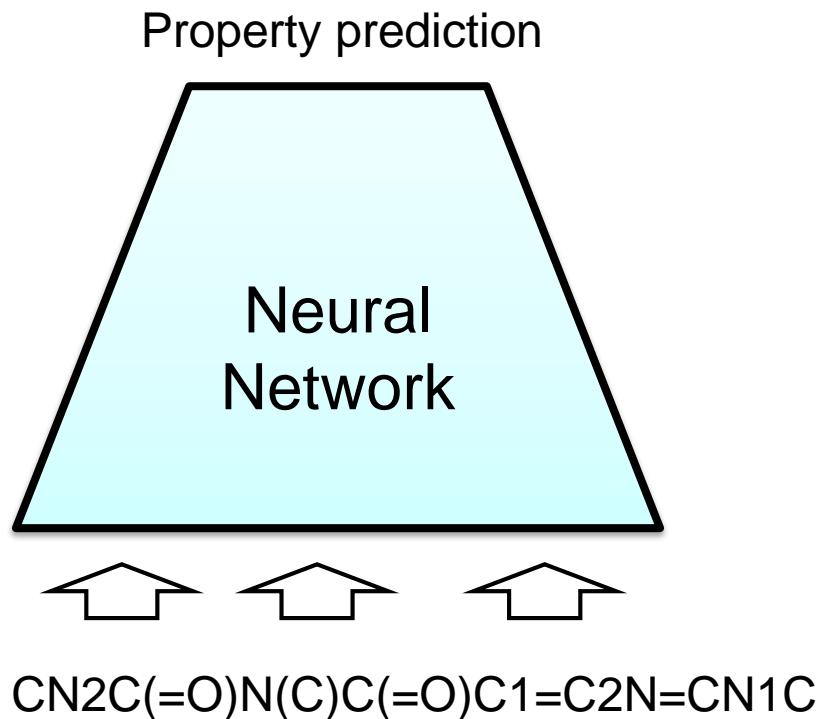


0.531
1.299
0.946
-0.218
0.017

A
C
T
I
V
I
T
Y

Quantitative Smiles – Activity Relationships

QSAR modeling using Smiles strings only*

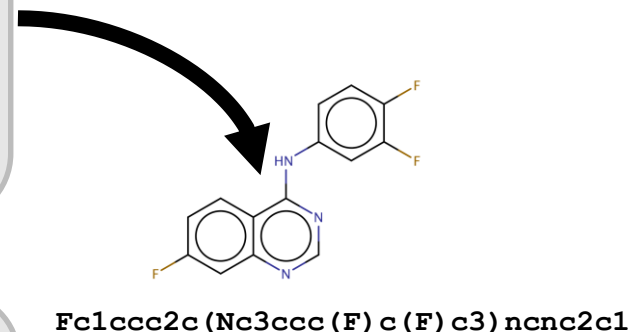
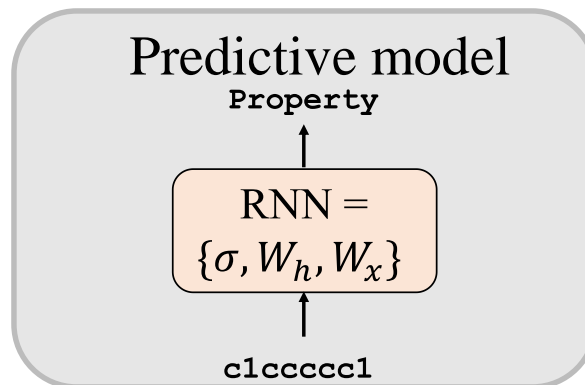
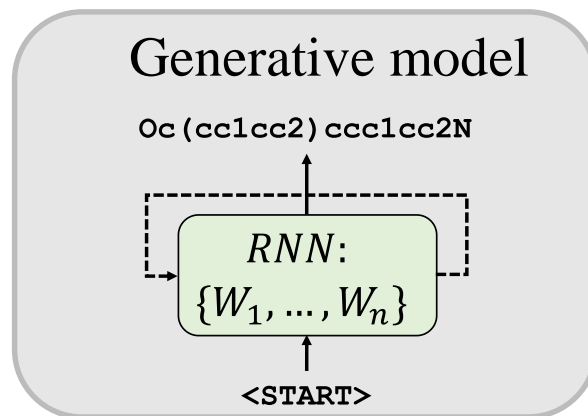


5CV RF model with DRAGON7 Descriptors 5CV NN model with SMILES directly

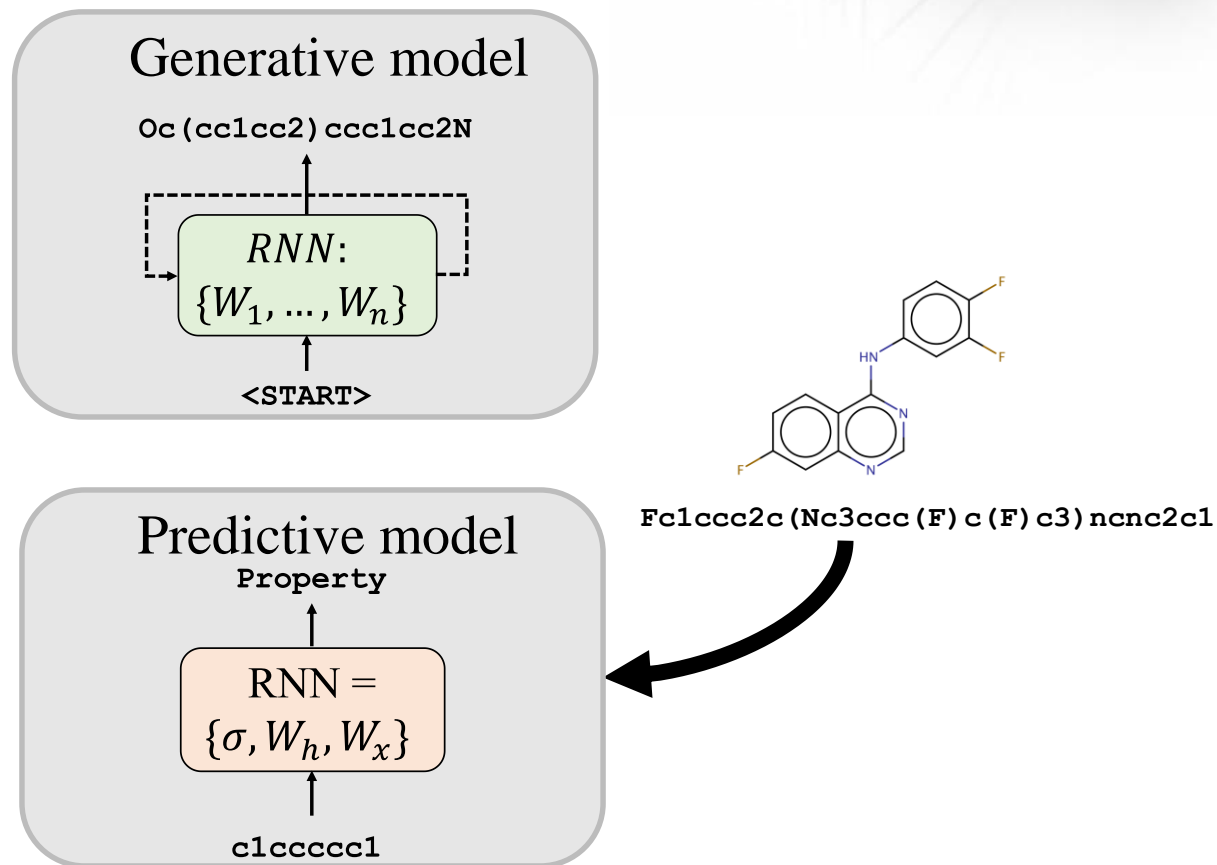
RMSE:	0.57	0.53
MAE:	0.37	0.35
R^2_{ext} :	0.90	0.91

*LogP data for ~16K molecules from PHYSPROP (srcinc.com), Toxcast Dashboard (<https://comptox.epa.gov/dashboard>), and others.

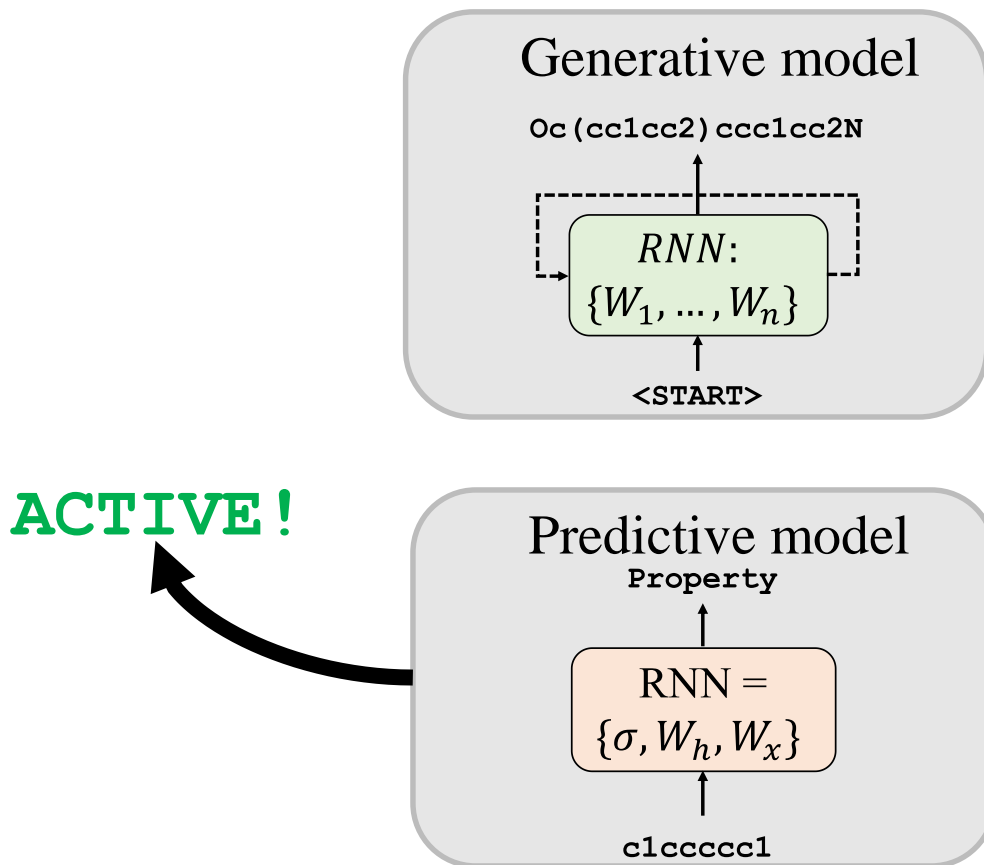
Reinforcement learning for chemical design



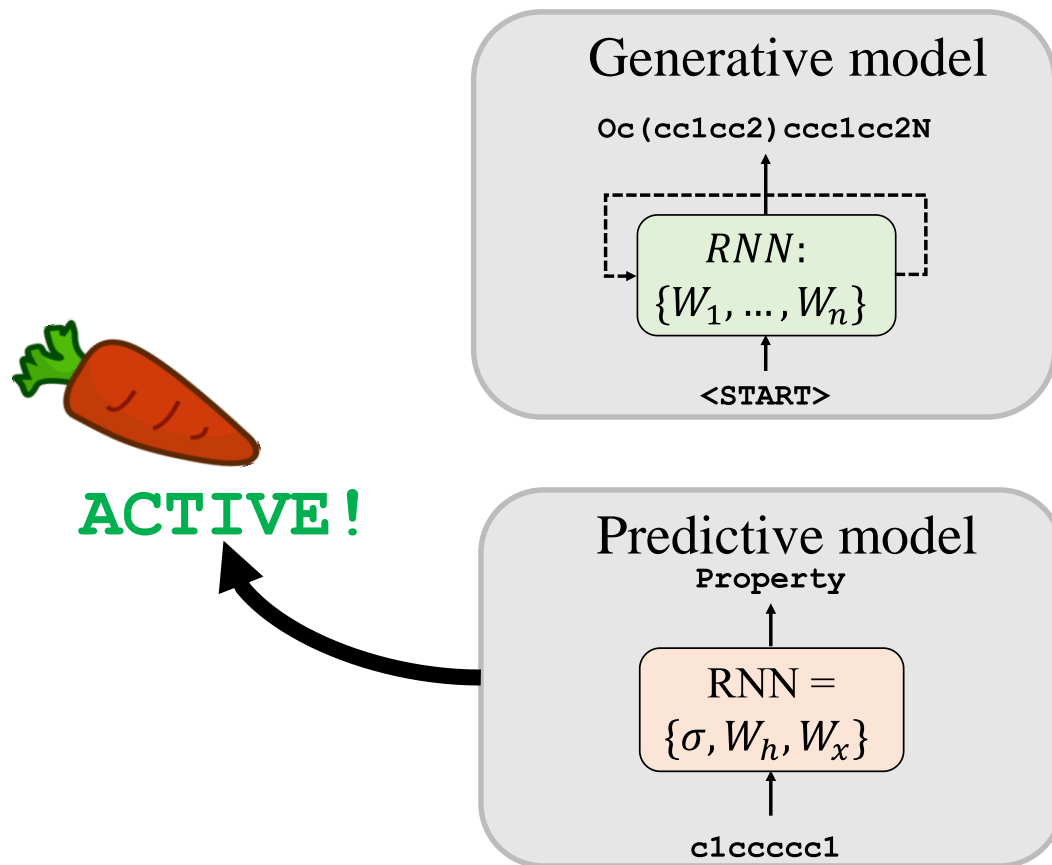
Reinforcement learning for chemical design



Reinforcement learning for chemical design



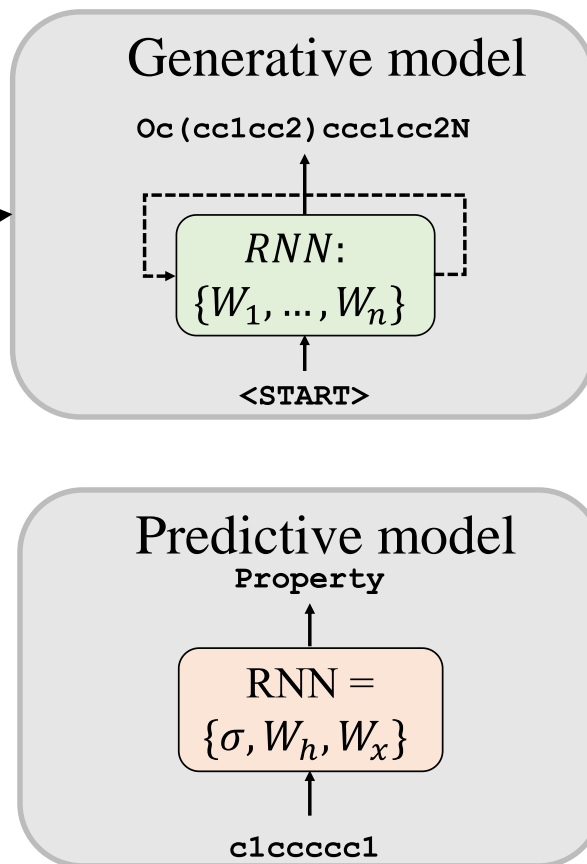
Reinforcement learning for chemical design



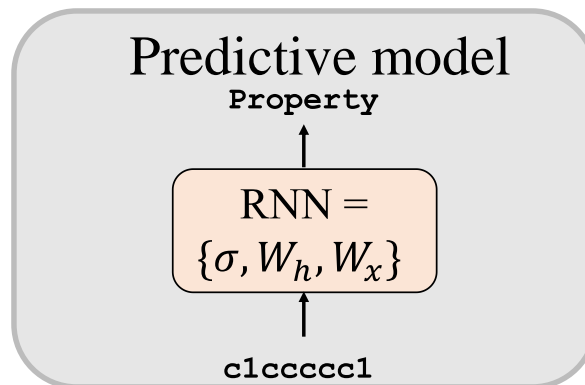
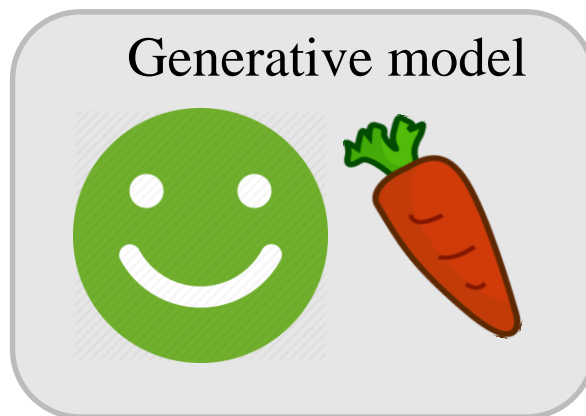
Reinforcement learning for chemical design



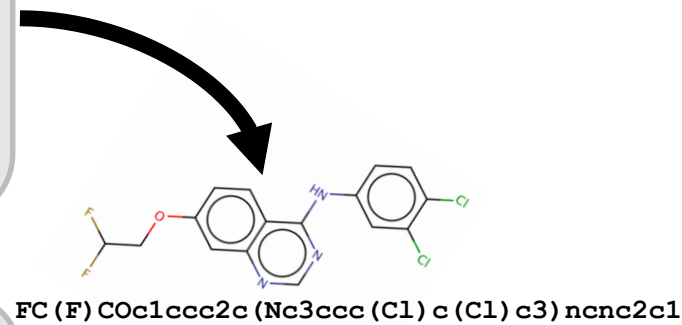
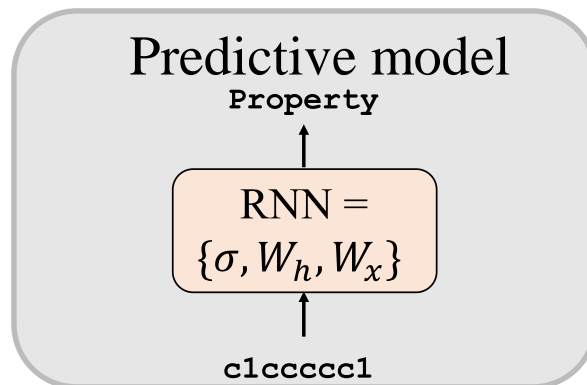
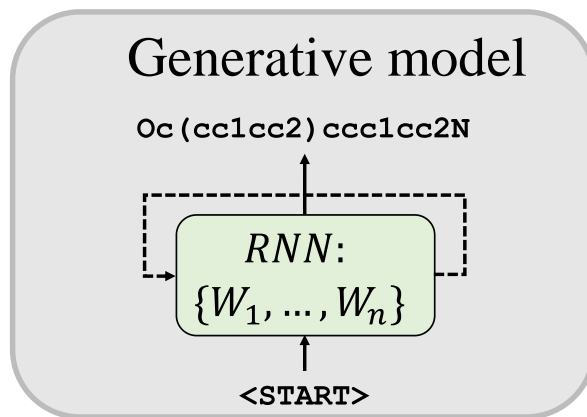
ACTIVE!



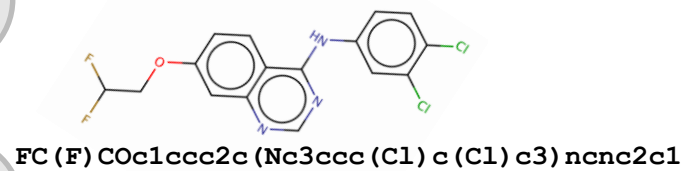
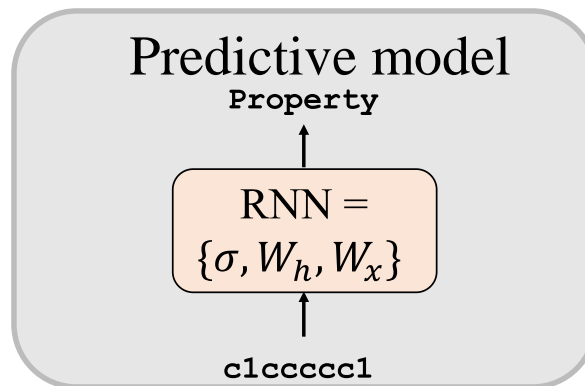
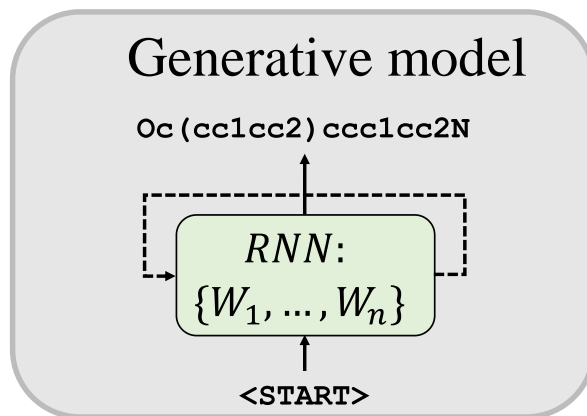
Reinforcement learning for chemical design



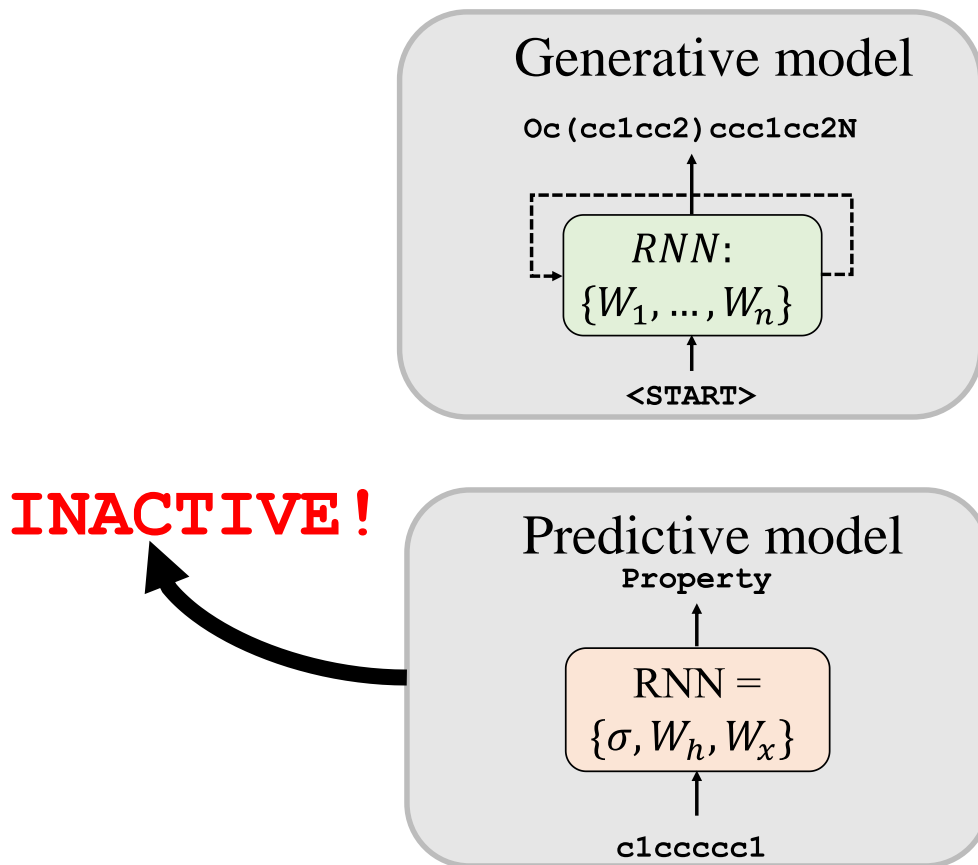
Reinforcement learning for chemical design



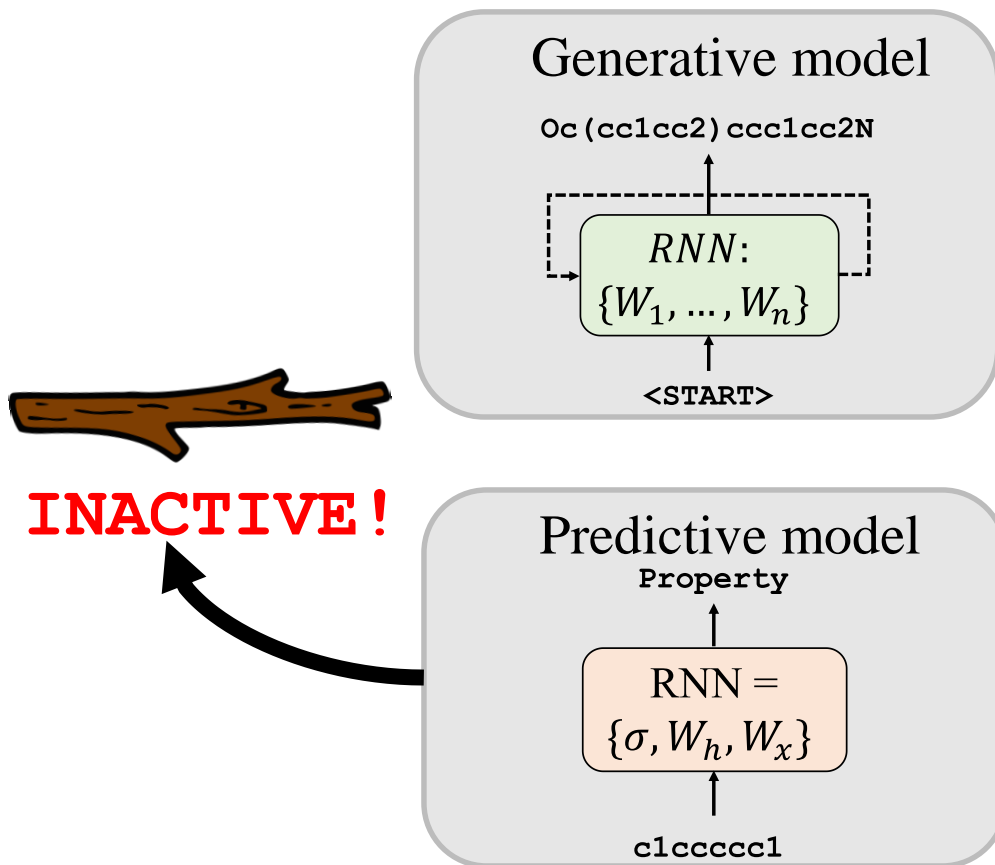
Reinforcement learning for chemical design



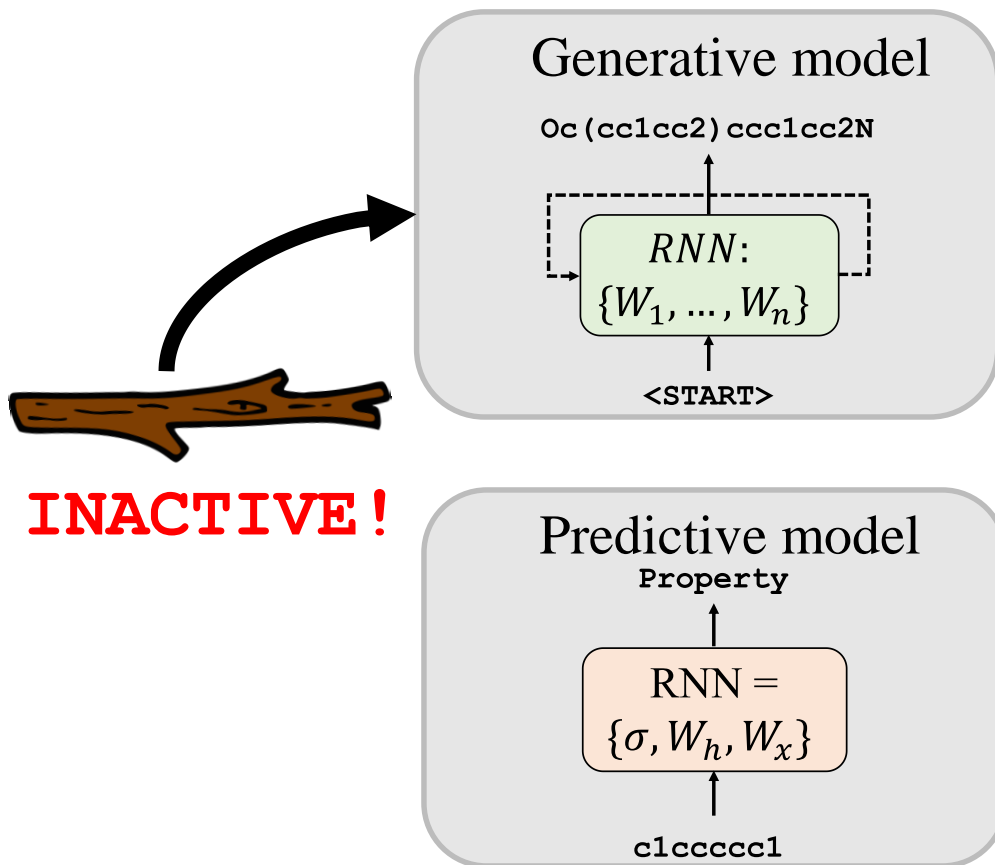
Reinforcement learning for chemical design



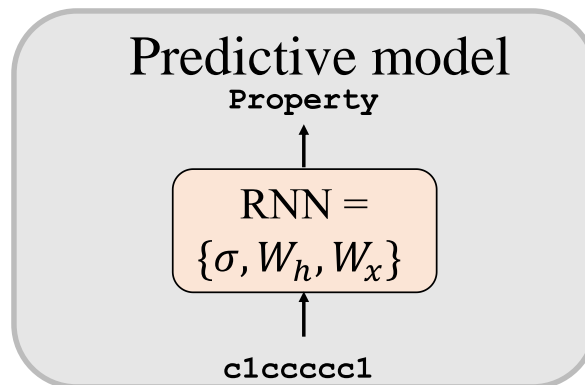
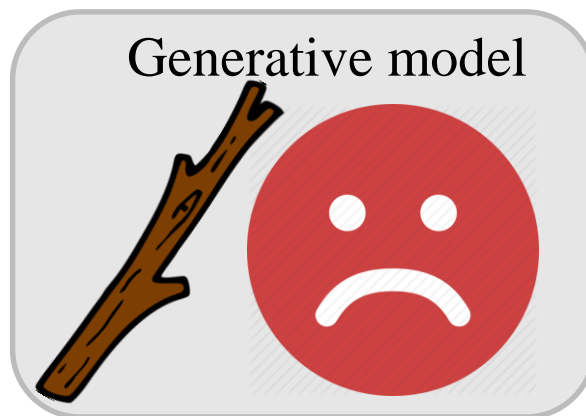
Reinforcement learning for chemical design



Reinforcement learning for chemical design



Reinforcement learning for chemical design

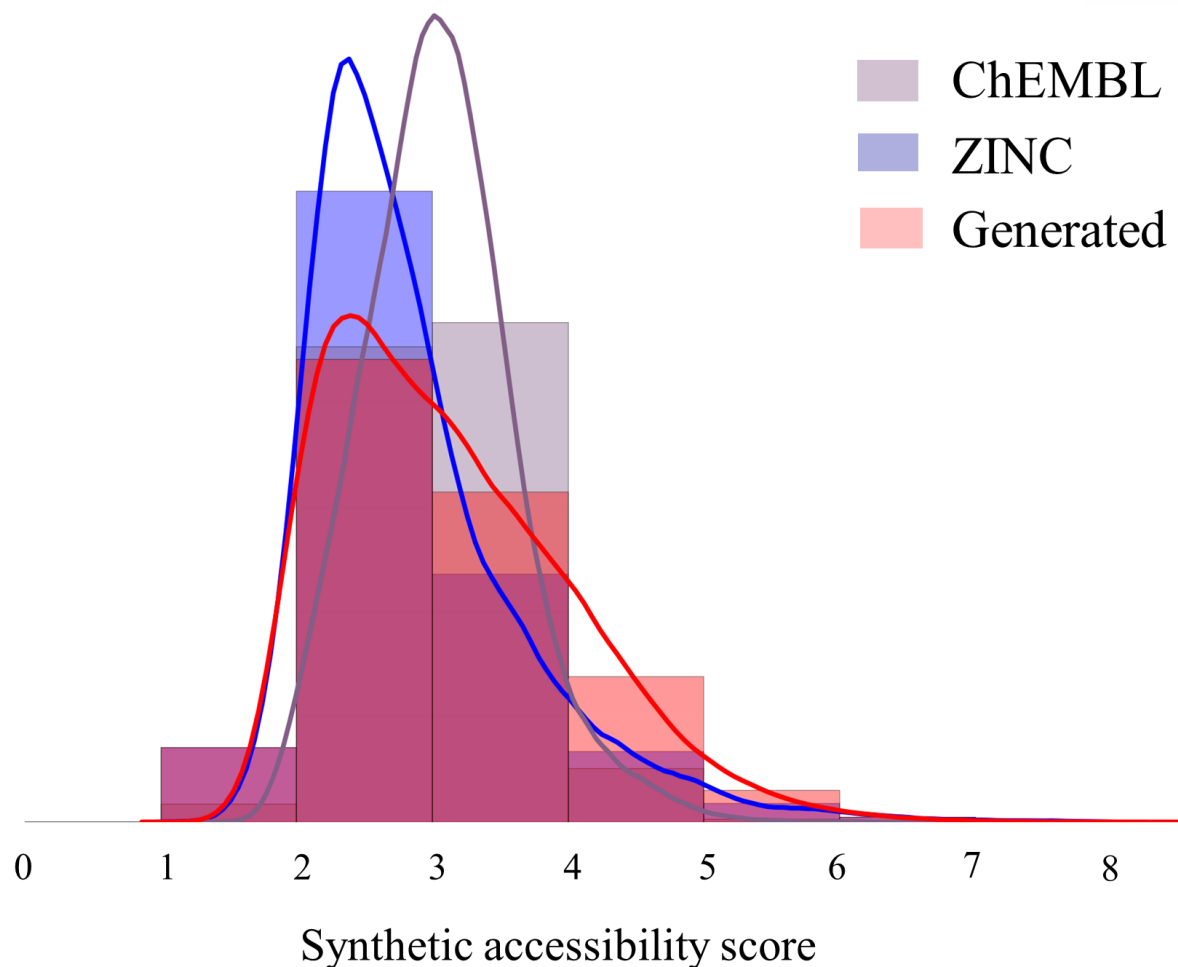


Technical details



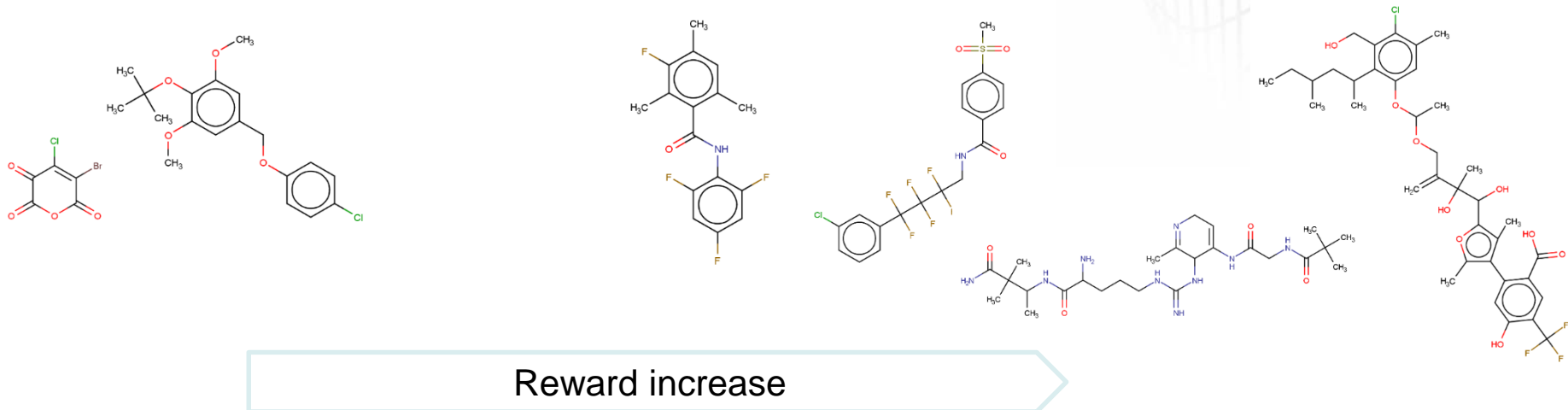
- Models were trained on Nvidia Titan X and Titan V GPUs
- Training the generative model on ChEMBL took ~ 25 days
- Training of predictive models took ~ 2 hours
- Biasing the generative model with reinforcement learning for one property ~ 1 day
- Generative model produces 1000 compounds per minute

Results: Synthetic accessibility score* of the designed libraries

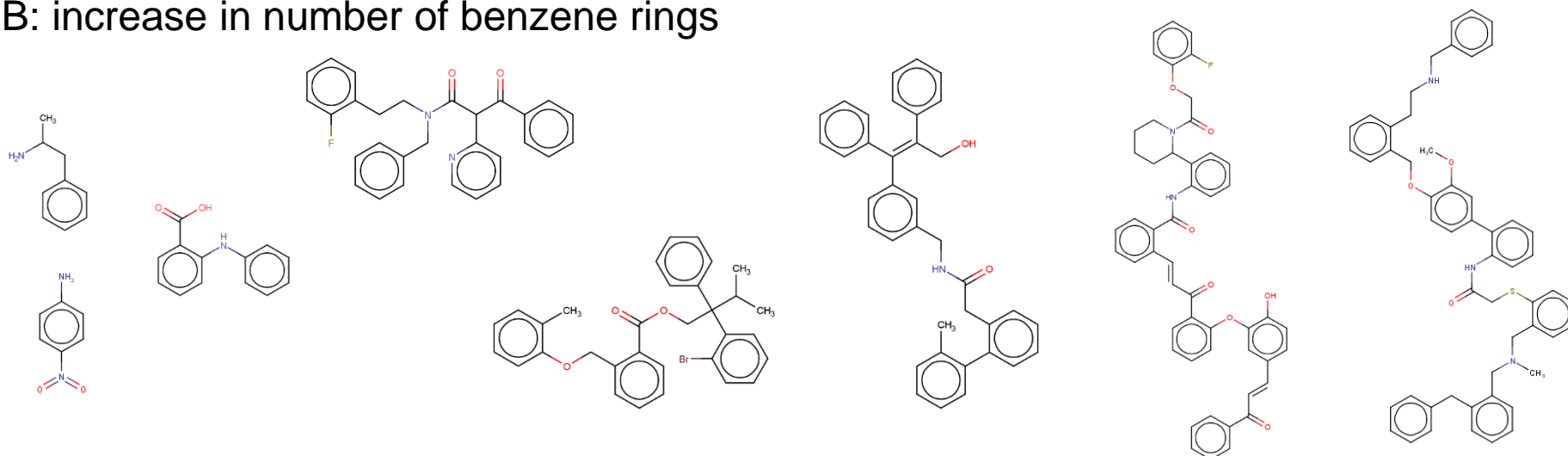


PoC: Structural Bias

A: increase in number of substituents

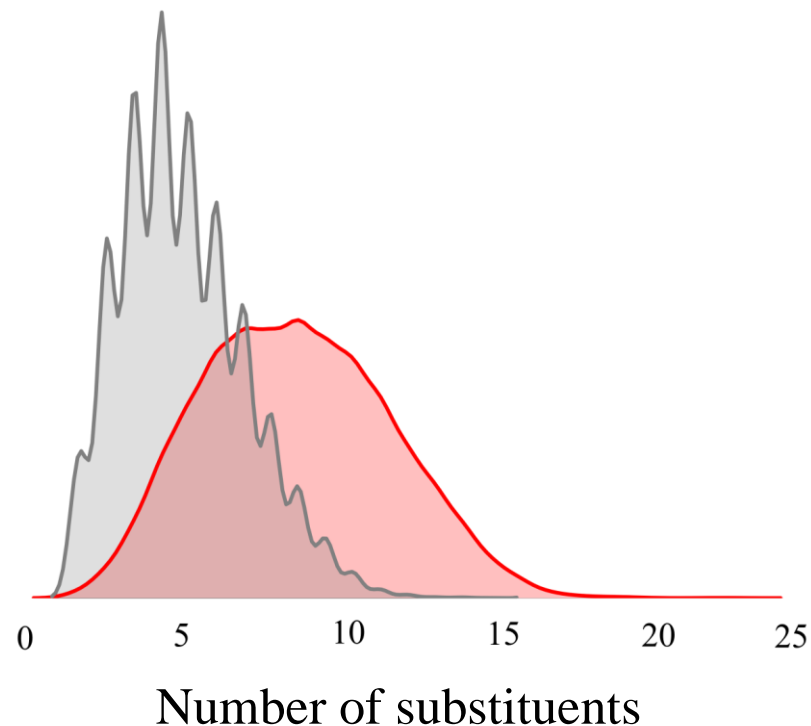
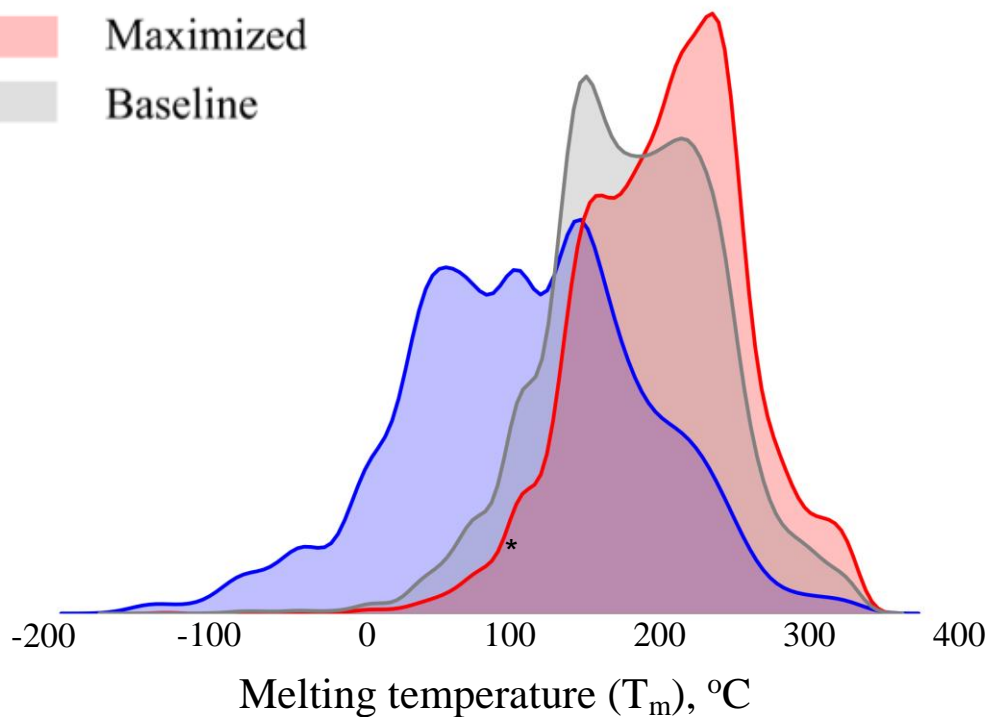


B: increase in number of benzene rings

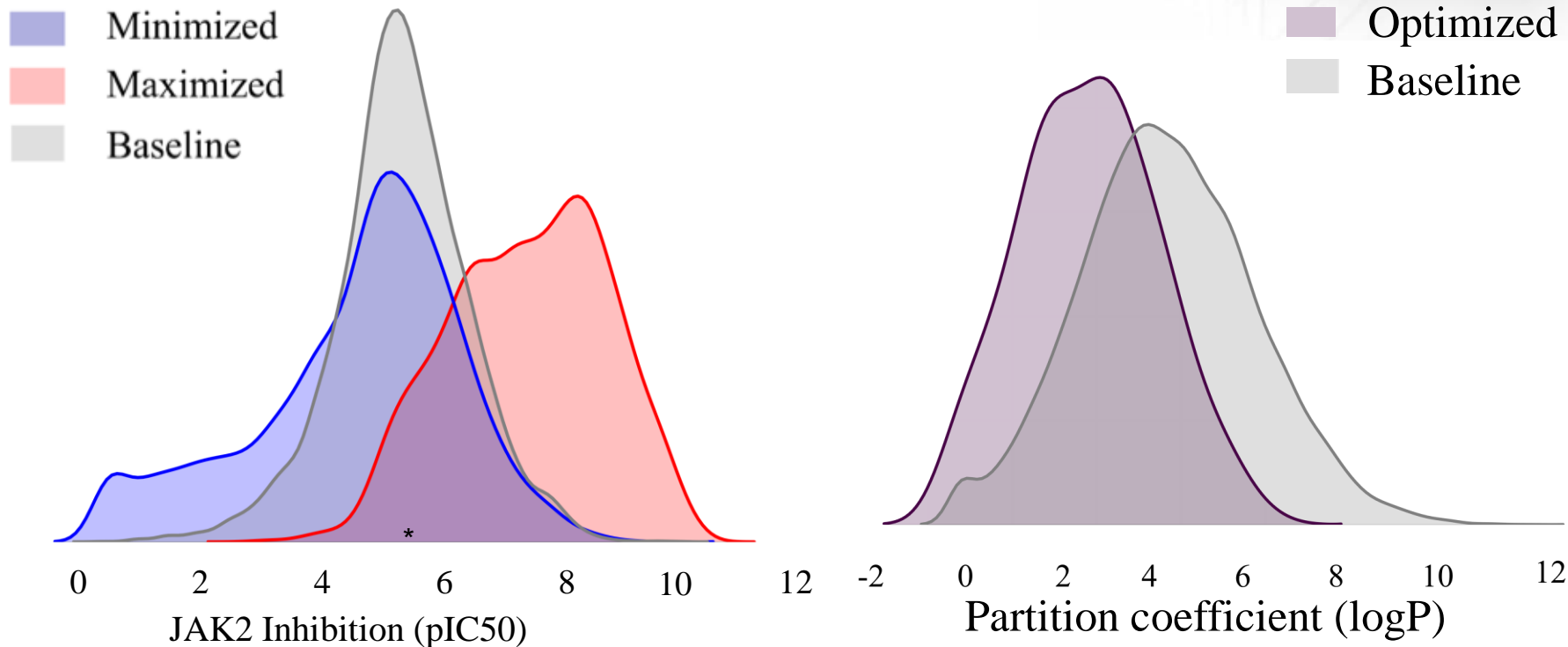


Results: Biasing target properties in the designed libraries

■ Minimized
■ Maximized
■ Baseline

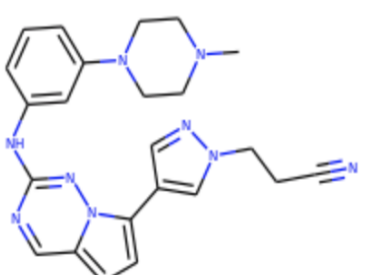


Results: Biasing target properties in the designed libraries



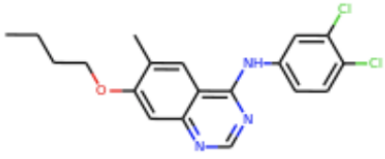
Target predictions for generated compounds using SEA*



Query	Target Key	Target Name	Description	P-Value	MaxTC
	NPM_HUMAN+5	NPM1	Nucleophosmin	3.118e-74	0.49
	CCNH_HUMAN+5	CCNH	Cyclin-H	2.571e-32	0.38
	PAK1_HUMAN+5	PAK1	Serine/threonine-protein kinase PAK 1	5.277e-24	0.39
	ALK_HUMAN+5	ALK	ALK tyrosine kinase receptor	3.714e-23	0.54
	JAK2_HUMAN+5	JAK2	Tyrosine-protein kinase JAK2	1.136e-21	0.61
	INSR_HUMAN+5	INSR	Insulin receptor	2.36e-17	0.54
	CCNB1_HUMAN+5	CCNB1	G2/mitotic-specific cyclin-B1	2.22e-16	0.38

*Keiser MJ, Roth BL, Armbruster BN, Ernsberger P, Irwin JJ, Shoichet BK. Relating protein pharmacology by ligand chemistry. *Nat Biotech* **25** (2), 197-206 (2007).

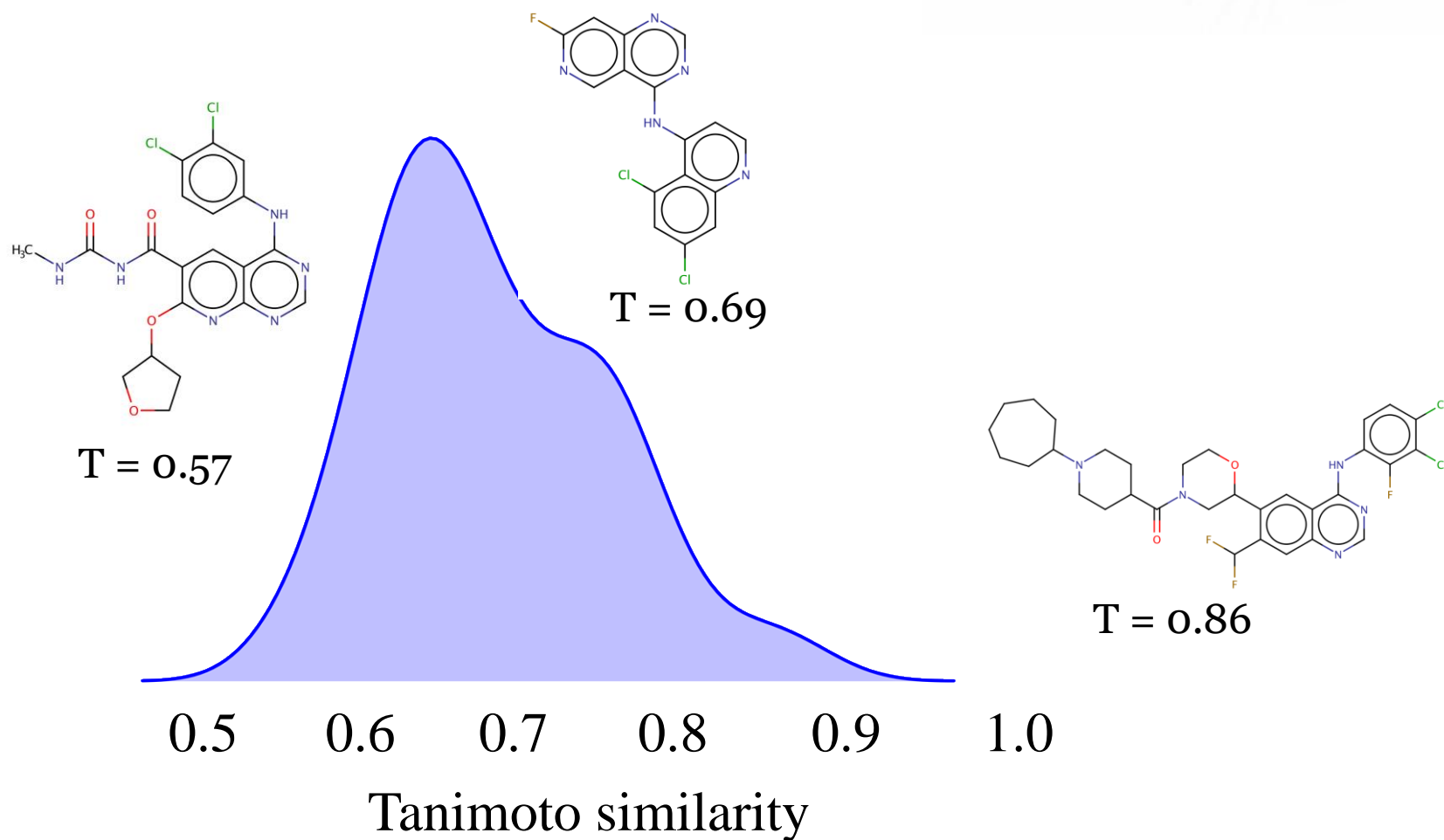
Target predictions for generated compounds using SEA*

Query	Target Key	Target Name	Description	P-Value	MaxTC
	EGFR_HUMAN+5	EGFR	Epidermal growth factor receptor	8.688e-244	0.61
	ERBB2_HUMAN+5	ERBB2	Receptor tyrosine-protein kinase erbB-2	8.544e-169	0.55
	ERBB2_RAT+5	ErbB2	Receptor tyrosine-protein kinase erbB-2	5.893e-87	0.42
	VGFR2_HUMAN+5	KDR	Vascular endothelial growth factor receptor 2	6.294e-65	0.58
	ERBB4_HUMAN+5	ERBB4	Receptor tyrosine-protein kinase erbB-4	1.354e-64	0.49

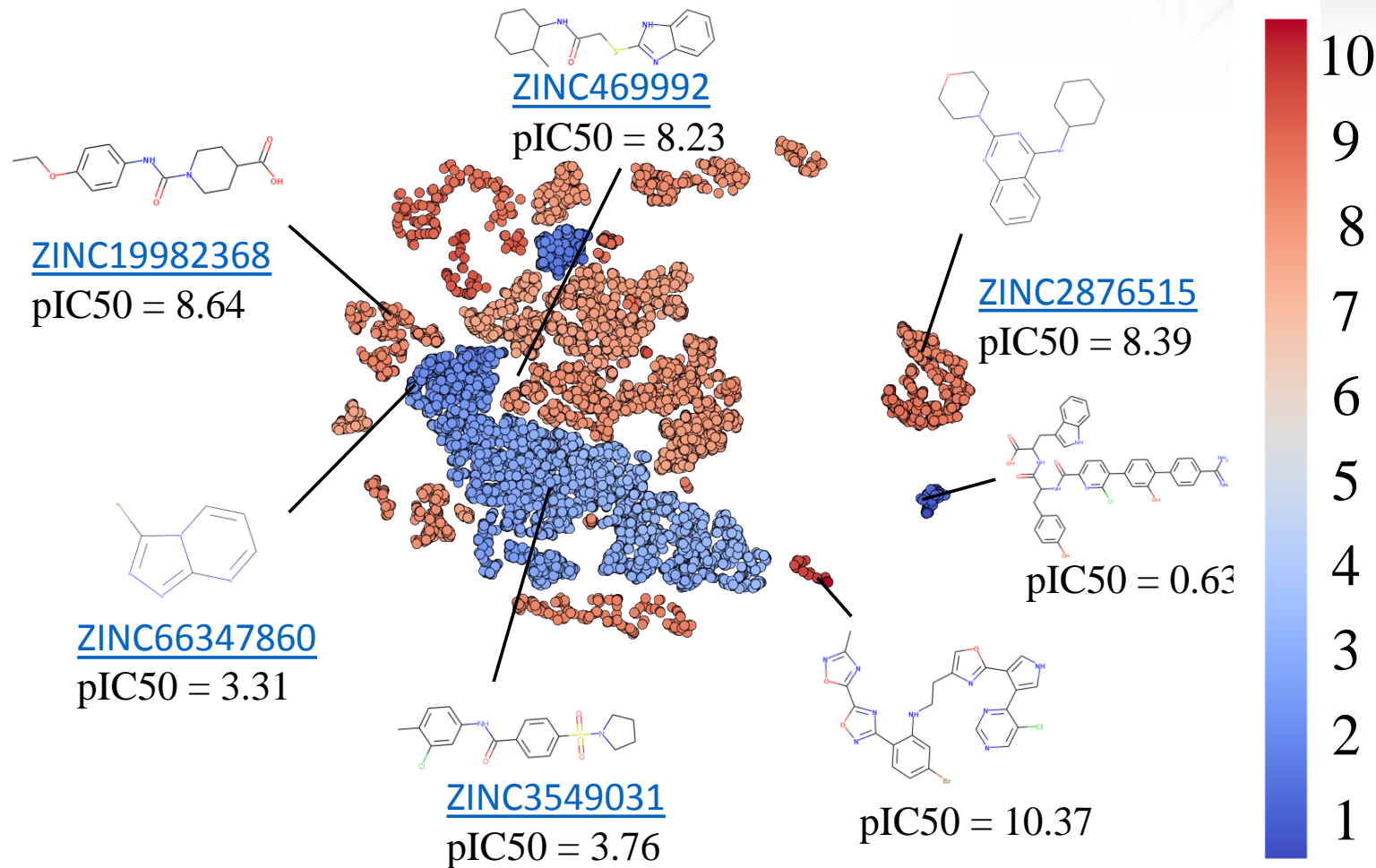
*Keiser MJ, Roth BL, Armbruster BN, Ernsberger P, Irwin JJ, Shoichet BK. Relating protein pharmacology by ligand chemistry. *Nat Biotech* **25** (2), 197-206 (2007).

Results: analysis of similarity

Distribution of Tanimoto similarity to the nearest neighbor in training dataset for compounds predicted to be active for EGFR by consensus of QSAR models:



Model visualization for putative JAK2 inhibitors (projection using t-SNE)



Summary

- AI methods coupled with SMILES representation (only!) afford biased library generation
- The system naturally embeds reinforcement learning to produce novel structures with the desired property
- The system can be tuned to bias libraries towards specific property ranges
- Next phase is experimental validation of hits

Summary of recent AI-based studies on chemical library design



Molecular representations	Generative models	Method of biasing generated compounds
<ul style="list-style-type: none">• Fingerprints• SMILES• Graphs	<ul style="list-style-type: none">• Autoencoders• Generative adversarial models• Recurrent neural networks• Convolutional neural networks	<ul style="list-style-type: none">• None• Latent space optimization• Fine-tuning on small subset of molecules with the desired property• Reinforcement Learning

An example of experimental validation of AI-based models*

- First training on large dataset
- Then fine-tuning on small subset of active compounds
- “These observations corroborate the ability of the generative AI model to **produce novel chemical entities within the training data domain**”.

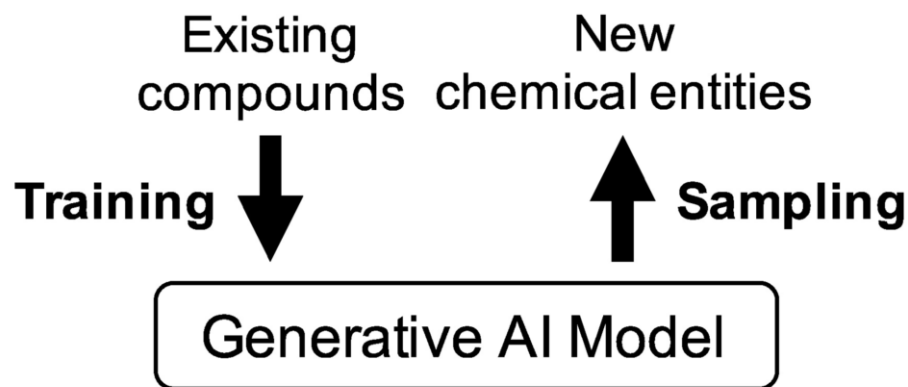


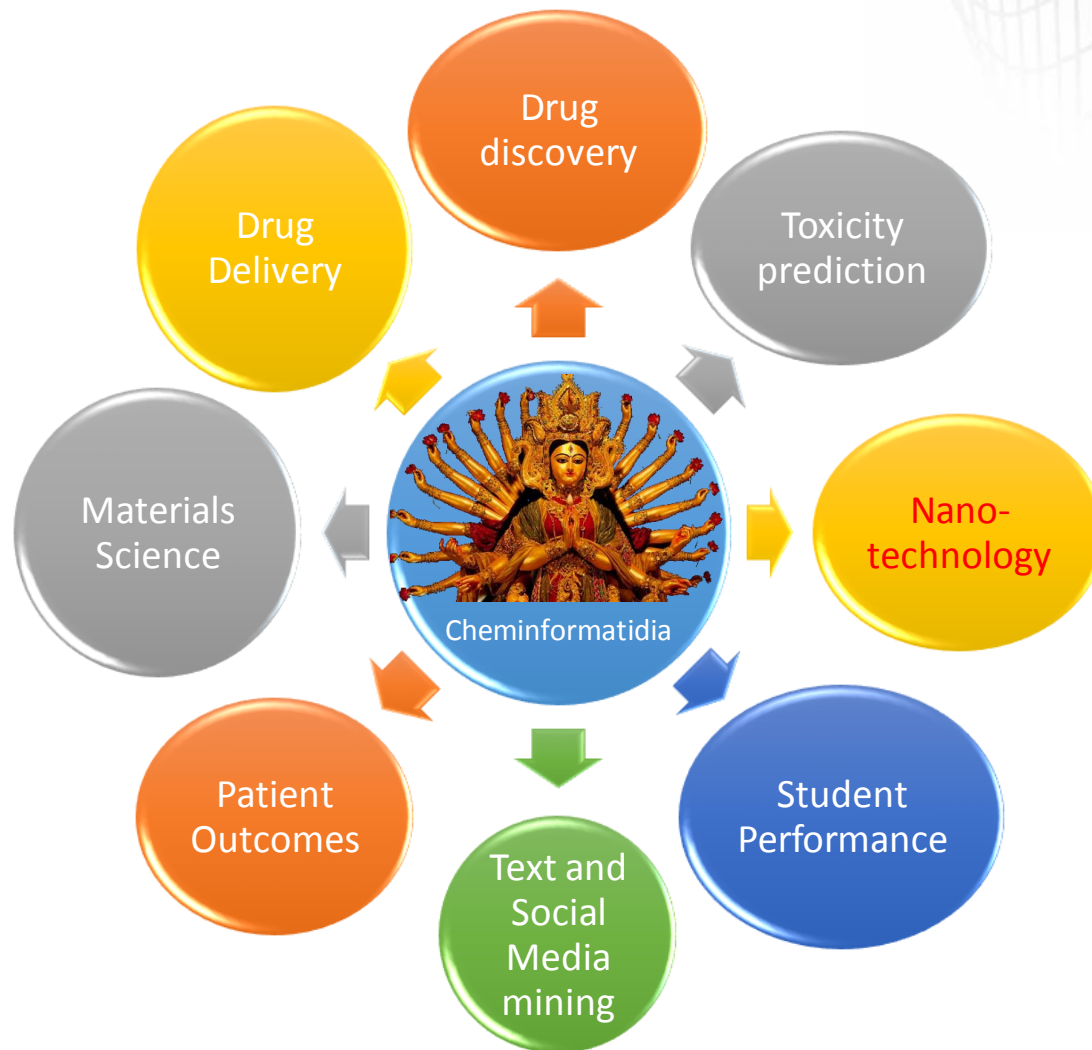
Table 1. *In vitro* activity of designs 1–5 on RXRs and PPARs (EC₅₀ values \pm SEM [μ M]; $n=2$ (when inactive) or 4 (when active) independent experiments in duplicates; *inactive*, no statistically significant reporter transactivation at a compound concentration of 30 μ M).

Compound no.	RXR α	RXR β	RXR γ	PPAR α	PPAR γ	PPAR δ
1	0.13 \pm 0.01	1.1 \pm 0.3	0.06 \pm 0.02	<i>inactive</i>	2.3 \pm 0.2	<i>inactive</i>
2	13.0 \pm 0.1	9 \pm 2	8.0 \pm 0.7	<i>inactive</i>	2.8 \pm 0.3	<i>inactive</i>
3	<i>inactive</i>	<i>inactive</i>	<i>inactive</i>	4.0 \pm 1.0	10.1 \pm 0.3	<i>inactive</i>
4	<i>inactive</i>	<i>inactive</i>	<i>inactive</i>	<i>inactive</i>	9 \pm 3	14 \pm 2
5	<i>inactive</i>	<i>inactive</i>	<i>inactive</i>	<i>inactive</i>	<i>inactive</i>	<i>inactive</i>
reference agonists ^{a)}	0.033 \pm 0.002	0.024 \pm 0.004	0.025 \pm 0.002	0.006 \pm 0.002	0.6 \pm 0.1	0.5 \pm 0.1

^{a)} Reference agonists, literature data: bexarotene^[17] for RXRs, GW7647^[18] for PPAR α , pioglitazone^[19] for PPAR γ , L165,041^[19] for PPAR δ

* D. Merk, L. Friedrich, F. Grisoni, G. Schneider, *Mol. Inf.* **2018**, 37, 1700153.

Many virtues of Cheminformatics



Acknowledgements



Principal Investigator

Alexander Tropsha

Research Professors

Alexander Golbraikh

Olexander Isayev

Eugene Muratov

Postdoctoral Fellows

Vinicius Alves

Stephen Capuzzi

Joyce Borba

Graduate students

Sherif Faraq

Kyle Bowers

Maria Popova

Andrew Thieme

- **Duke University**
 - Stefano Curtarolo
 - Corey Oses
- **UNC Chemistry**
 - Jim Cahoon
 - Taylor Moot
 - Aaron Taggart

MAJOR FUNDING

NIH

- 1U01CA207160
- R01-GM114015
- 5U54CA198999
- 1OT3TR002020

ONR

- N00014-16-1-2311