Quantum Machine Learning

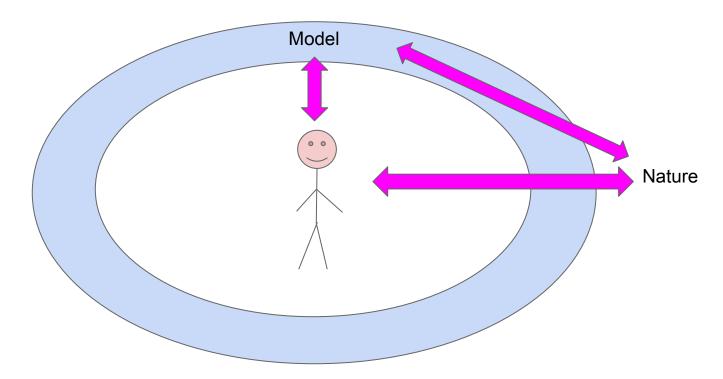
Learning curves, representations, training sets

Anatole von Lilienfeld

"Machine Learning, Quantum Mechanics, and Chemical Compound Space"
By Ramakrishnan and von Lilienfeld
published in: *Reviews in Computational Chemistry*edited by Abby L. Parrill and Kenny B. Lipkowitz
Volume **30**, Chapter 5, pages 225-256 (2017)



What's more important? The right skill set or the right mind set?



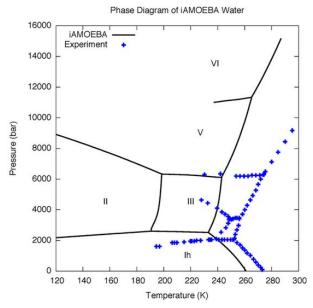
How do we do physics?

- 1. Guess a law
- 2. Build a model
- 3. Predict an outcome
- 4. If it does not compare to experiment it's wrong

Richard



Why do we do (chemical) physics?



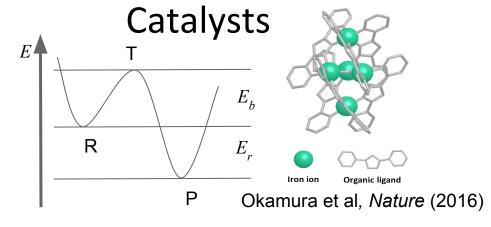
Pande et al, J. Phys. Chem B (2013)

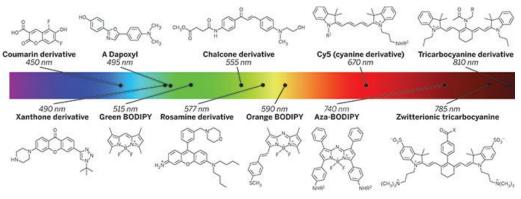
OLEDs



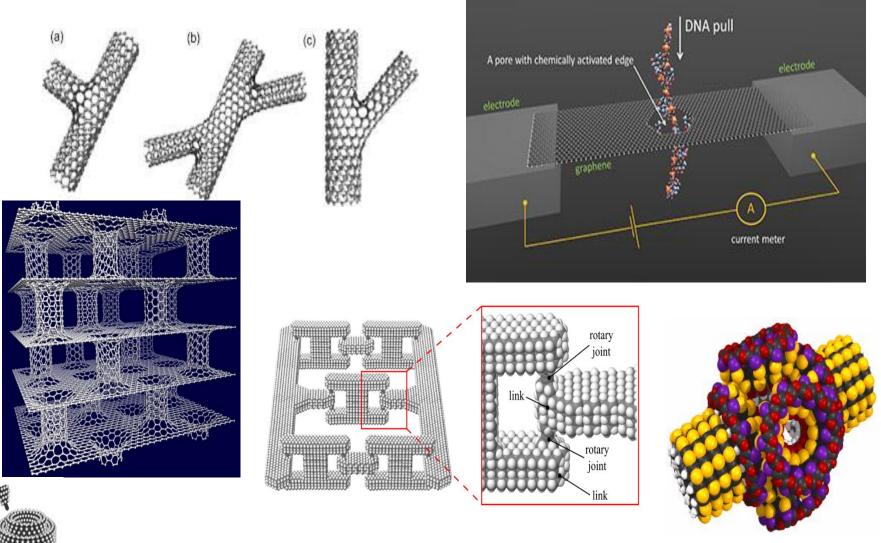
Understanding!!!

Structure



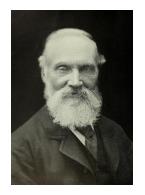


Complex Integrated Nanosystems



Theory to understand chemistry ... to help design experiments

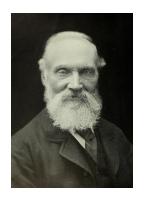
→ predictions that can be falsified



1. " ... when you cannot measure it, when you cannot express it in numbers, your knowledge is of a meagre and unsatisfactory kind." Lord Kelvin

Theory to understand chemistry ... to help design experiments

→ predictions that can be falsified



- 1. "... when you cannot measure it, when you cannot express it in numbers, your knowledge is of a meagre and unsatisfactory kind." Lord Kelvin
- 2. " ... we only understand molecules once we predict properties with quantitative accuracy" M. Quack, ETHZ (2000)

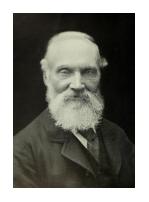


TABLE I. History of the accurate calculations of the ground state of helium atom.

Year	Ref.	Type	Energy (a.u.)
1929	Hylleraas (Ref. 2)	Hylleraas (three terms)	-2.902 43
1957	Kinoshita (Ref. 6)	Kinoshita type	-2.903 72 2 5
1966	Frankowski and Pekeris (Ref. 7)	Logarithm	-2.903 724 377 03 2 6
1994	Thakkar and Koga (Ref. 8)	Half-integer	-2.903 724 377 034 114 4
1998	Goldman (Ref. 9)	Polynomial	-2.903 724 377 034 119 594
1999	Drake (Ref. 10)	Double exponent	-2.903 724 377 034 119 596
2002	Sims and Hagstrom (Ref. 12)	Hylleraas-CI	-2.903 724 377 034 119 598 2 9 99
2002	Drake et al. (Ref. 11)	Triple exponent	-2.903 724 377 034 119 598 305
2002	Korobov (Ref. 13)	Slater geminal	-2.903 724 377 034 119
2006	Schwartz (Ref. 15)	Logarithm $(ln(s))$	-2.903 724 377 034 119
	5 (5)	3 2 W	194 404 44 0 049 5
2007	This work	ICI (new logarithm)	-2.903 724 377 034 119 598 311 159 245
			194 404 446 696 9 05 37

Theory to understand chemistry ... to help design experiments

→ predictions that can be falsified

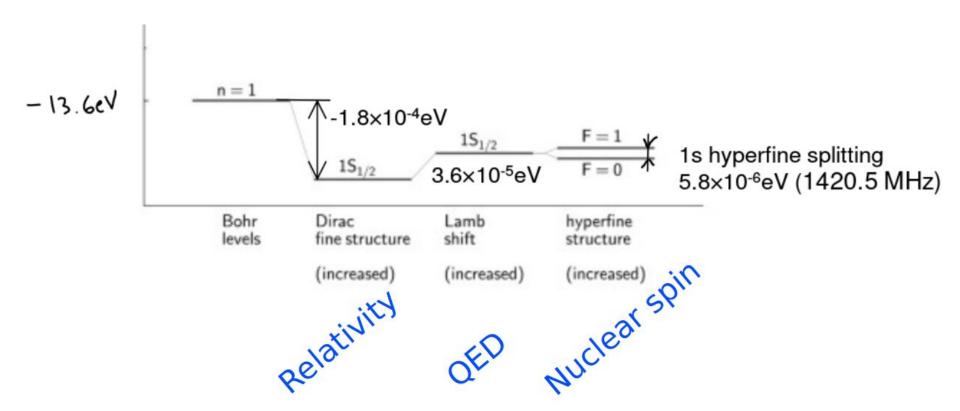


- 1. "... when you cannot measure it, when you cannot express it in numbers, your knowledge is of a meagre and unsatisfactory kind." Lord Kelvin
- 2. " ... we only understand molecules once we predict properties with quantitative accuracy" M. Quack, ETHZ (2000)
- 3. "... It is nice to know that the computer understands the problem.

 But I would like to understand it too." E. Wigner
- → compare to experiment (arbiter)



Energy of Hydrogen atom



1.0 1784

based on mechanical production equipment driven by water and steam power



2.0 1870

based on mass production enabled by the division of labor and the use of electrical energy



3.0 1969

based on the use of electronics and IT to further automate production



4.0 tomorrow

based on the use of cyber-physical systems



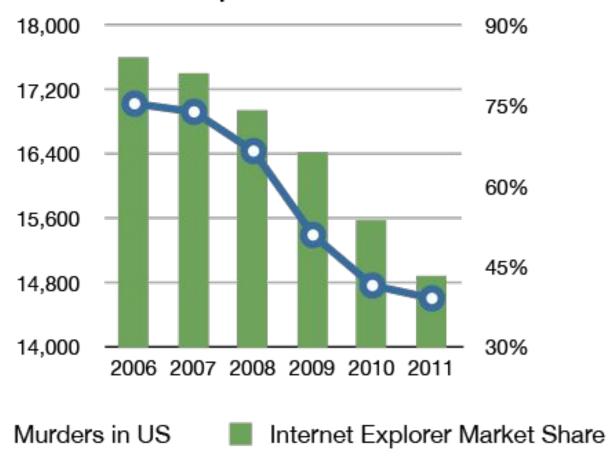




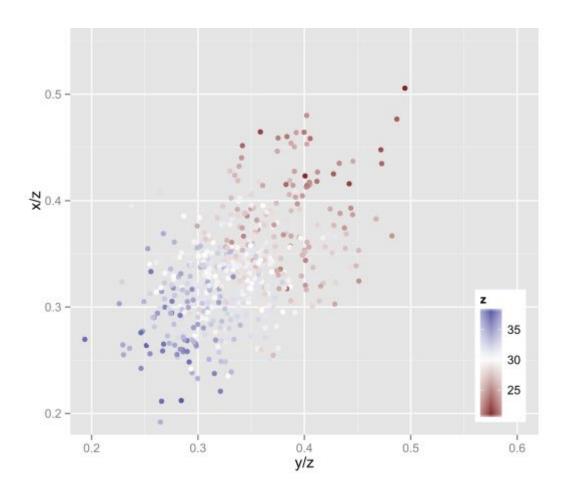


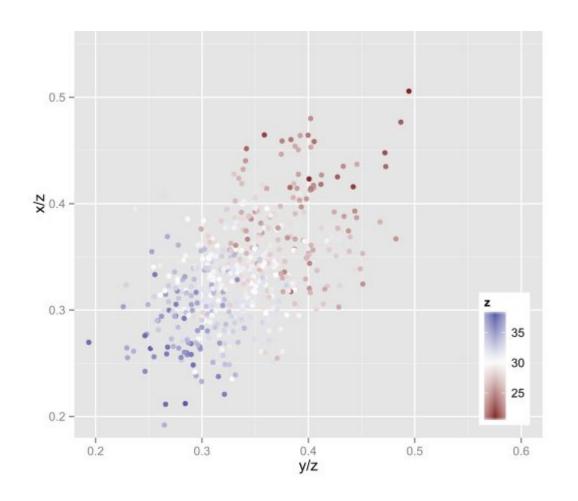


Internet Explorer vs Murder Rate



More at http://www.tylervigen.com/spurious-correlations





Example of spurious correlation for 500 draws of x,y,z with respective means of 10,10,30 and standard deviations 1, 1, and 9. From wikipedia

$$x, y \sim N(10, 1)$$
$$z \sim N(30, 9)$$

Correlation must *not* be used to infer a causal relationship, however if there is a causal relationship there must be a correlation ...

→ Correlation is a necessary but not sufficient condition.

Dangerous: Humans have cognitive bias ["Thinking, Fast and Slow" Tversky and Kahneman, "Fooled by Randomness", Nassim Taleb]

Spurious correlation can also be due to

- 1. chance (anything which varies simultaneously will correlate)
- 2. a common cause
- 3. identity relationships

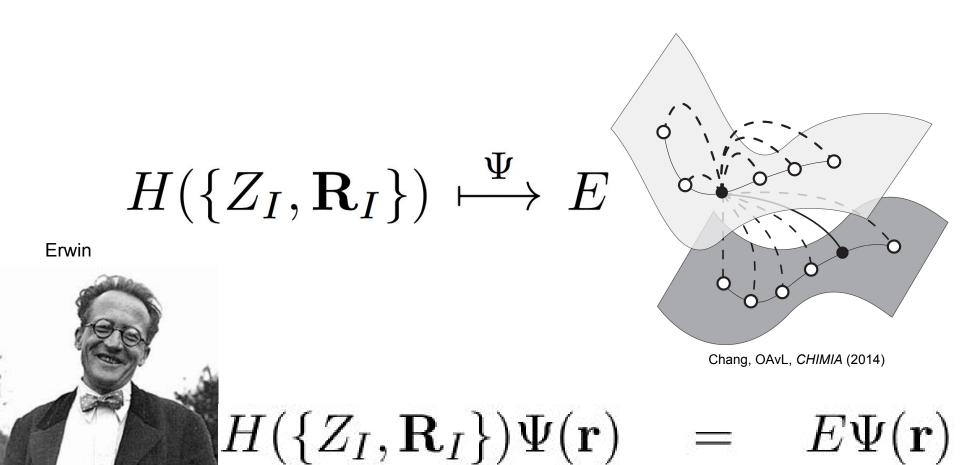
correlations (inductive) vs. law (deductive)

Erwin



$$H(\{Z_I, \mathbf{R}_I\})\Psi(\mathbf{r}) = E\Psi(\mathbf{r})$$

correlations (inductive) vs. law (deductive)



correlations (inductive) vs. law (deductive)

$$\{Z_I, \mathbf{R}_I\} \stackrel{\mathrm{ML}}{\longmapsto} E$$

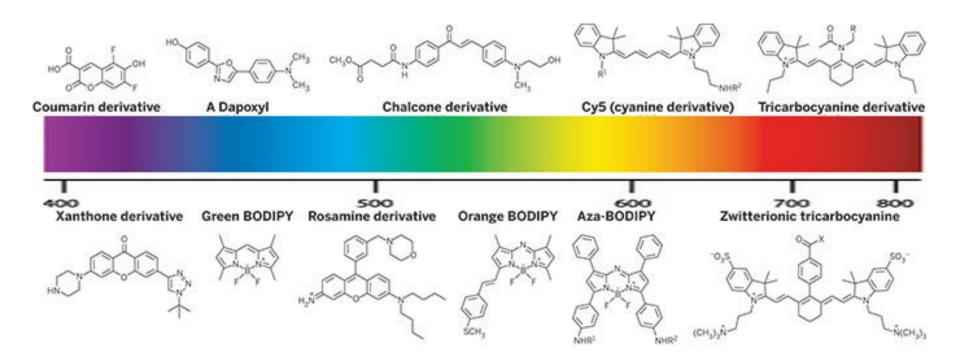
$$H(\{Z_I, \mathbf{R}_I\}) \stackrel{\Psi}{\longmapsto} E$$

Erwin

Chang, OAvL, CHIMIA (2014)

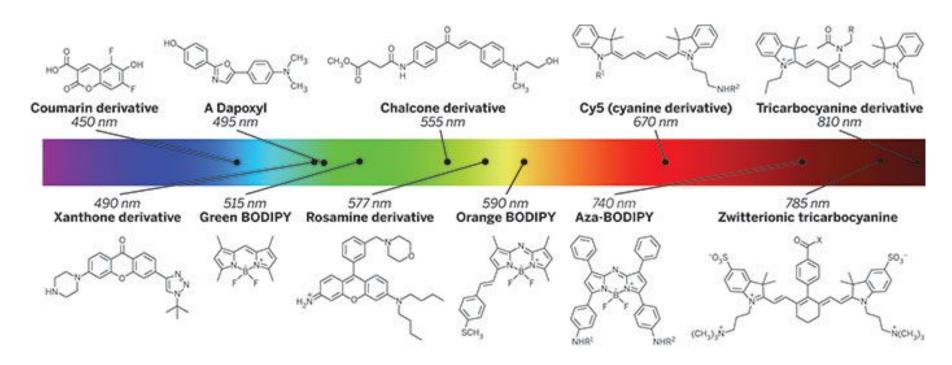
$$H(\{Z_I, \mathbf{R}_I\})\Psi(\mathbf{r}) = E\Psi(\mathbf{r})$$

Configuration + Composition → Chemical Space



Young-Tae Chang et al C&E News 93 (12) 39-40 (2015)

Configuration + Composition → Chemical Space

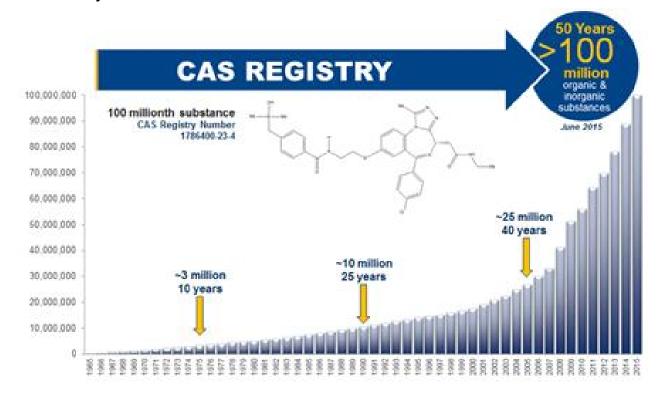


Young-Tae Chang et al C&E News 93 (12) 39-40 (2015)

How many are possible?

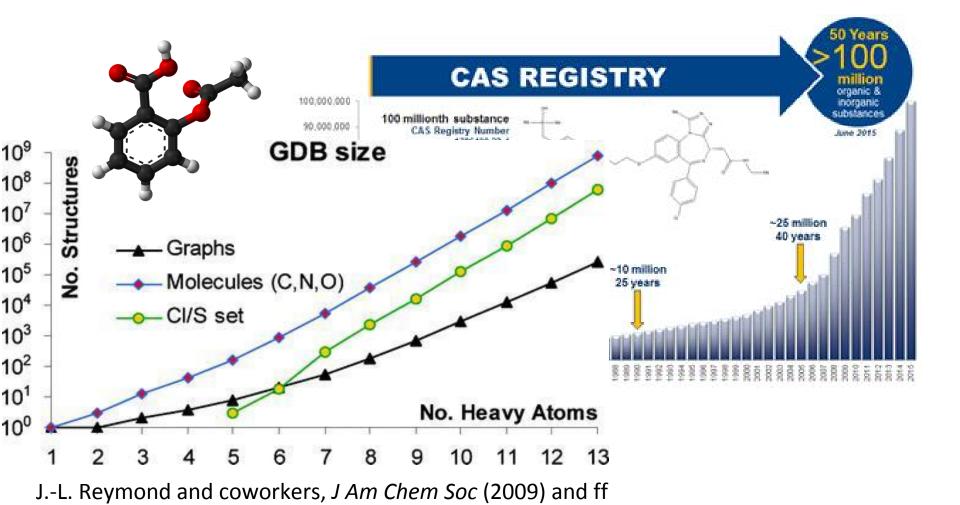
Differ in composition and constitution (no conformational isomers)

- ~120 M
- ~15 k are being added on a daily basis

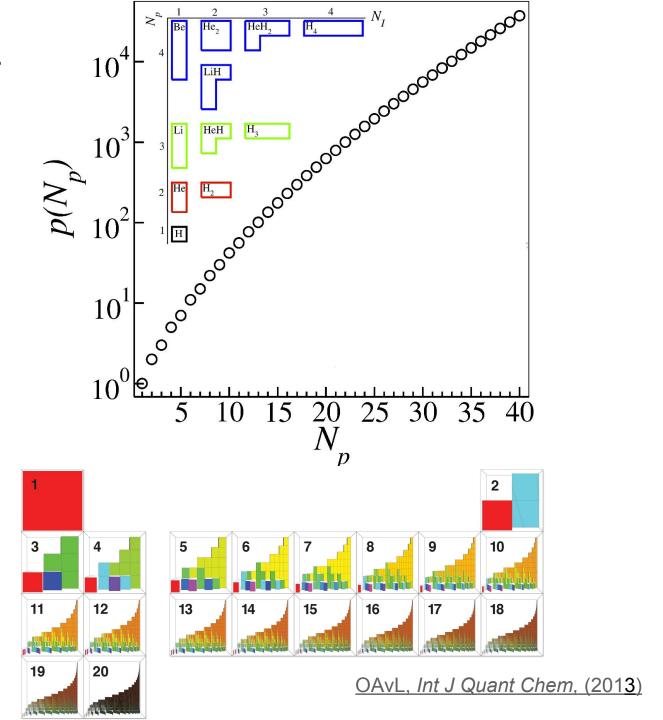


"The greatest shortcoming of the human race is our inability to understand the exponential function"

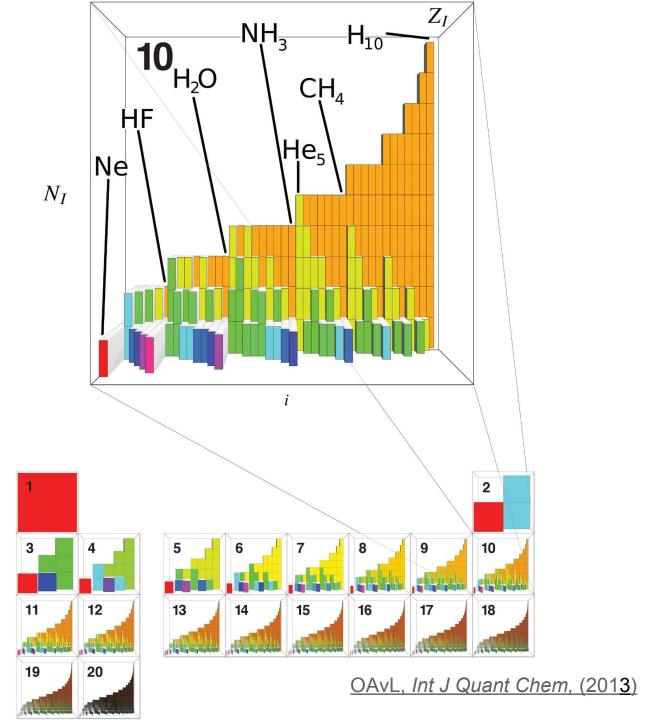
Al Bartlett, U of Colorado Boulder



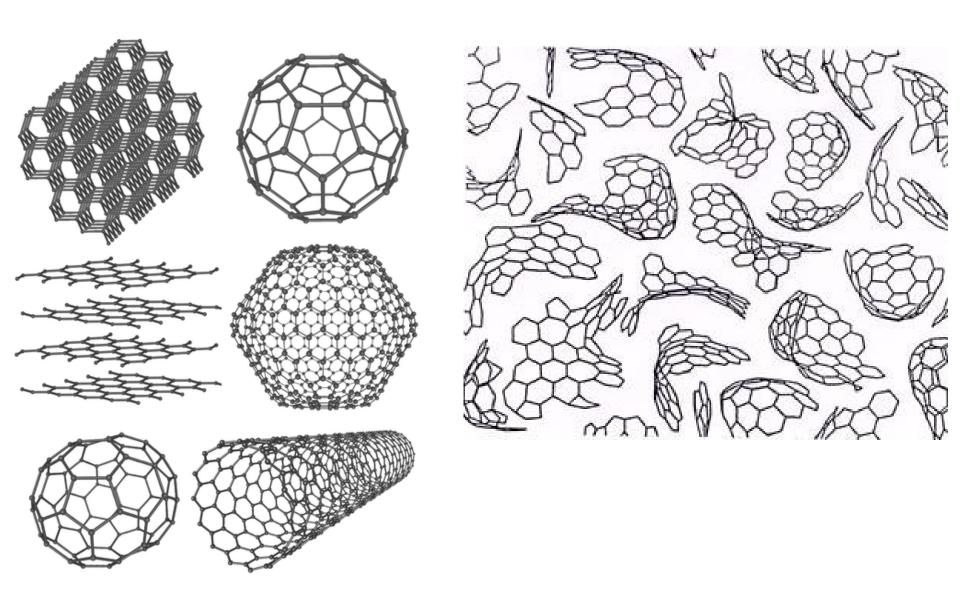
Composition

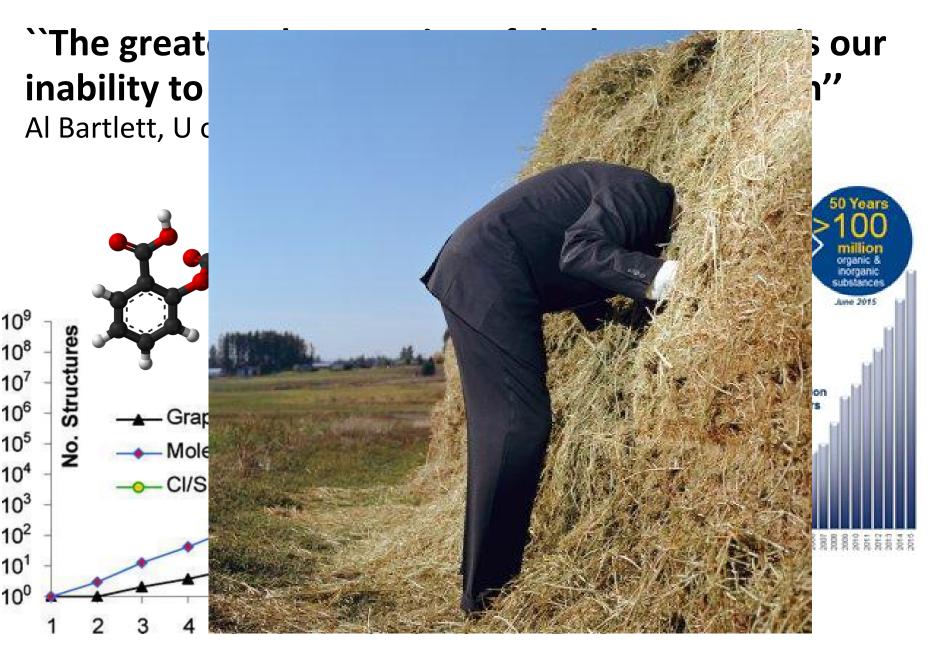


Composition 10 protons



Spatial configuration Carbon allotropes





J.-L. Reymond and coworkers, J Am Chem Soc (2009) and ff

Conclusions

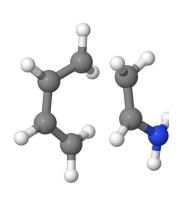
1. Instantaneous QM quality predictions

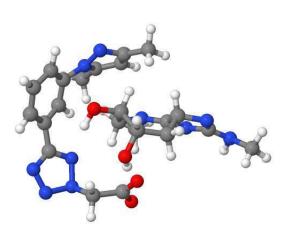
2. Learning curves reveal quality of ML model

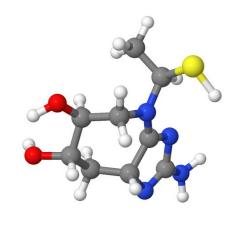
3. Representations

4. Data sets

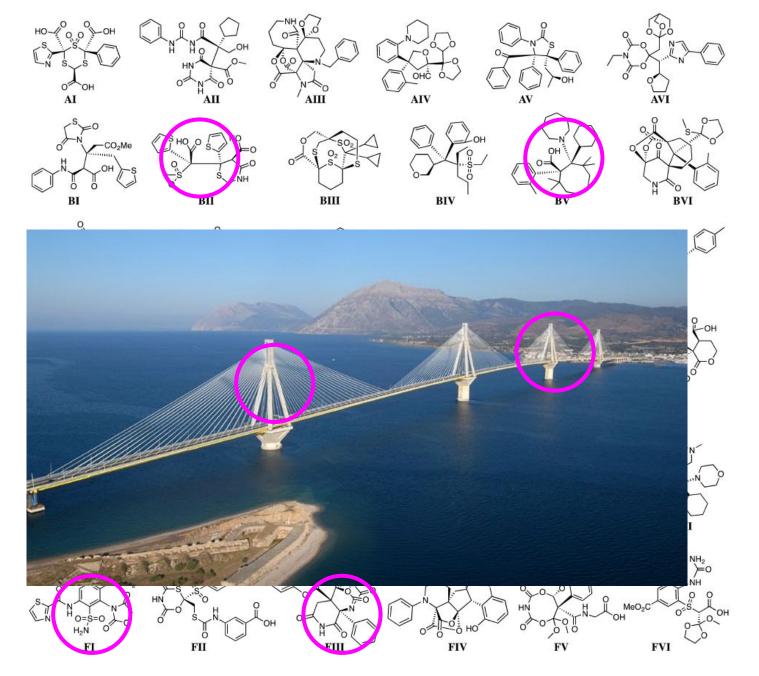
-70.160669 96288144_10











Virshup, Yang, Beratan et al J Am Chem Soc (2013)

Kernel Ridge Regression

Kernel

$$E^{est}(\mathbf{M}) = \sum_{i}^{N} \alpha_i k(\mathbf{M}, \mathbf{M}_i)$$

e.g.
$$k(\mathbf{M}, \mathbf{M}') = \exp\left(-\frac{d(\mathbf{M}, \mathbf{M}')^2}{2\sigma^2}\right)$$

Regression

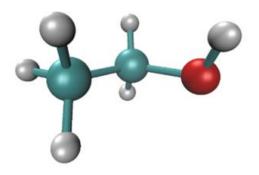
$$\min_{\alpha} \left(\sum_{i} \left(E^{est}(\mathbf{M}_{i}) - E_{i}^{ref} \right)^{2} + \lambda \sum_{ij} \alpha_{i} \alpha_{j} k(\mathbf{M}_{i}, \mathbf{M}_{j}) \right)$$

Solution

$$\alpha = (\mathbf{K} + \lambda \mathbf{I})^{-1} \mathbf{E}^{ref}$$

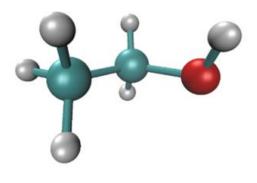
From molecule to representation

Molecule



From molecule to representation

Molecule

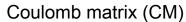


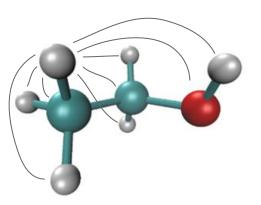


From molecule to Coulomb matrix (CM) to Bag of Bonds (BOB)

$$M_{IJ} = \begin{cases} 0.5Z_I^{2.4} & \forall I = J, \\ \frac{Z_I Z_J}{|\mathbf{R}_I - \mathbf{R}_J|} & \forall I \neq J. \end{cases}$$

Molecule



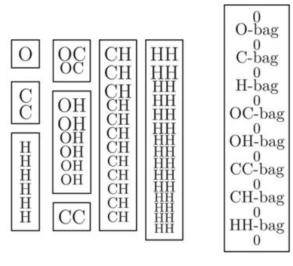


	0	C	C	H	H	H	H	H	H
0	0	OC	OC	OH	OH	OH	OH	OH	OH
C	OC	C CC	CC	CH	CH	CH	CH	CH	CH
C	OC	CC	C	CH	CH	CH	CH	CH	CH
H	OH	CH	CH	H	HH	HH	HH	HH	HH
H	OH	CH	CH	HH	H	HH	HH	HH	HH
\overline{H}	OH	CH	CH	HH	HH	H	HH	HH	HH
H	OH	CH	CH	HH	HH	HH	H	HH	HH
H	OH	CH	CH	HH	HH	HH	HH	H	HH
H		CH							

Rupp et al, Phys Rev Lett (2012)

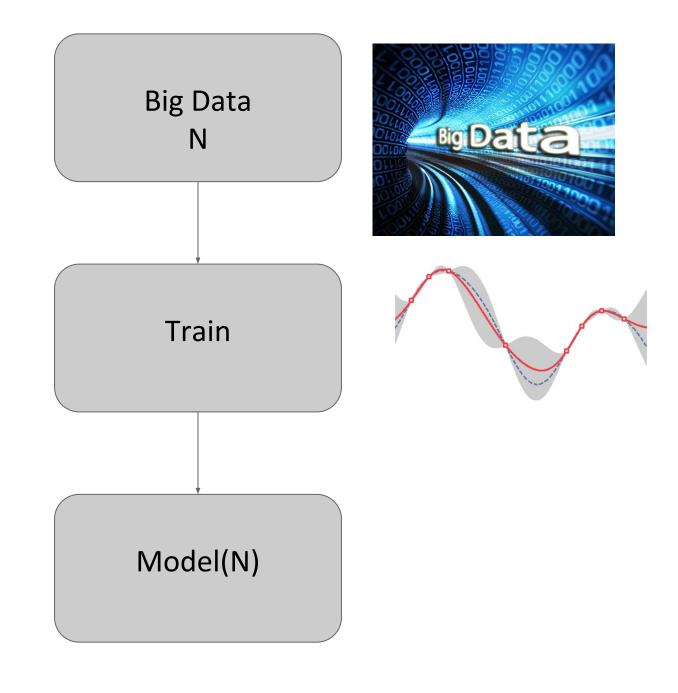
- Unique but overcomplete
- Invariances (Tra&Rot)
- Compact
- Physical meaning
- Fast
- Simple metrics are not smooth if sorted

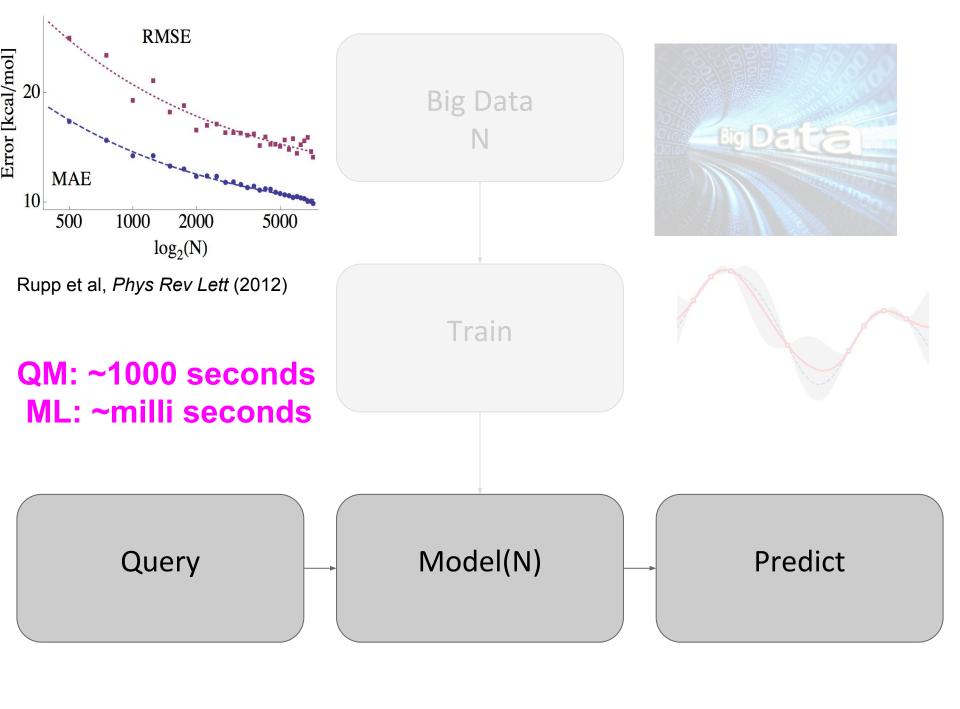
Bag of Bonds (BoB)

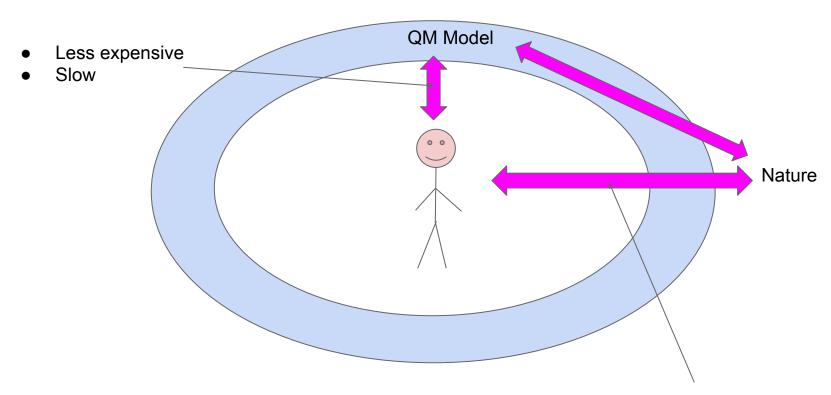


Hansen et al, J Phys Chem Lett (2015)

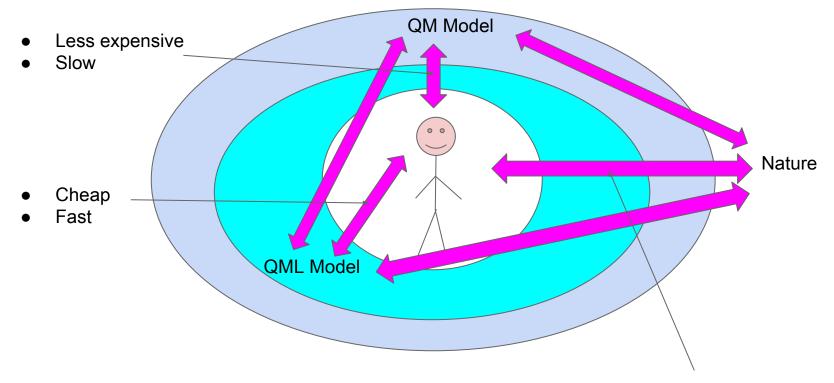
- Not unique (homometricity)
- Invariant (Tra&Rot)
- Compact
- Physical meaning
- Fast
- Simple metrics are smooth



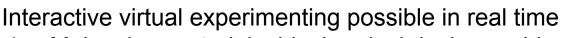




- Expensive unless fast and automatized
- Inaccessible (too small, too hot, too far, too slow)



- Expensive unless fast and automatized
- Inaccessible (too small, too hot, too far, too slow)



- 1. Molecular, materials, biochemical design problems
- 2. Discover new trends/relationships/rules/fill gaps
- 3. Enhance teaching, communication, and outreach

Conclusions

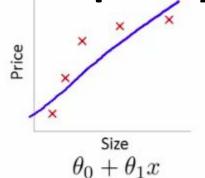
1. Instantaneous QM quality predictions

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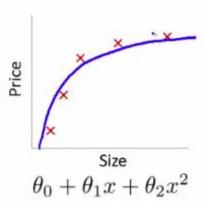
4. Data sets

Model quality

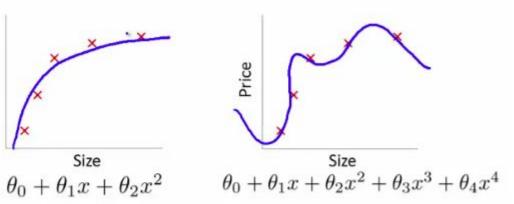


High bias (underfit)

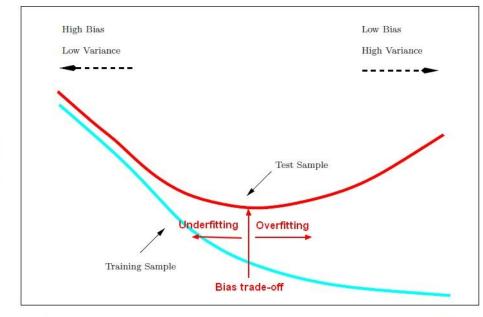
Prediction Error



"Just right"



High variance (overfit)

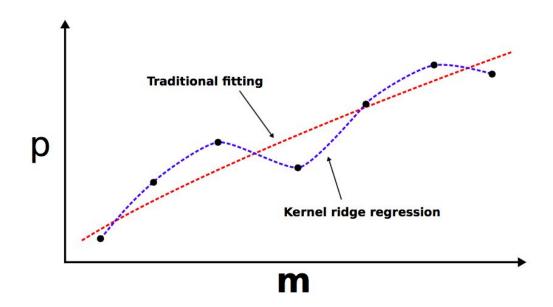


Low

High

Model quality

The bigger the data the better ...



†Model quality

Test

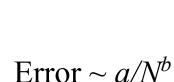
Traditional fitting

Kernel ridge regression

N

The bigger the data the better ...

"Learning curves in machine learning" Claudia Perlich, Encyclopedia of Machine Learning (Springer, 2011) pp. 577–580.

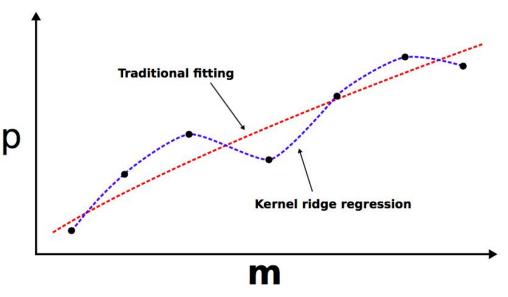


Train

Ε

K.-R. Mueller et al, Neural Comput (1996)

 $\rightarrow \log(\text{Error}) = \log(a) - b\log(N)$



†Model quality

Test

Traditional fitting

Kernel ridge regression

N

The bigger the data the better ...

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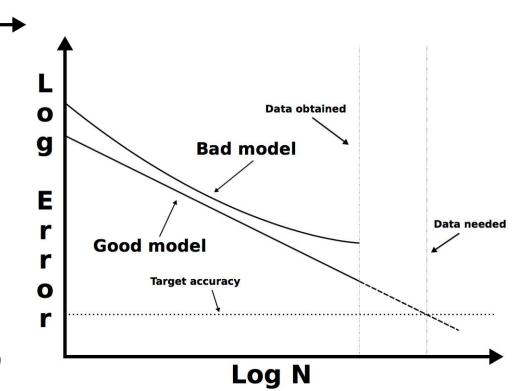
Ε

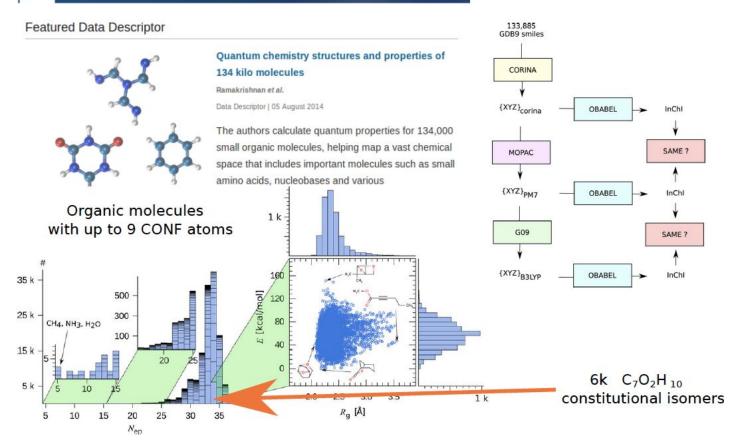
0

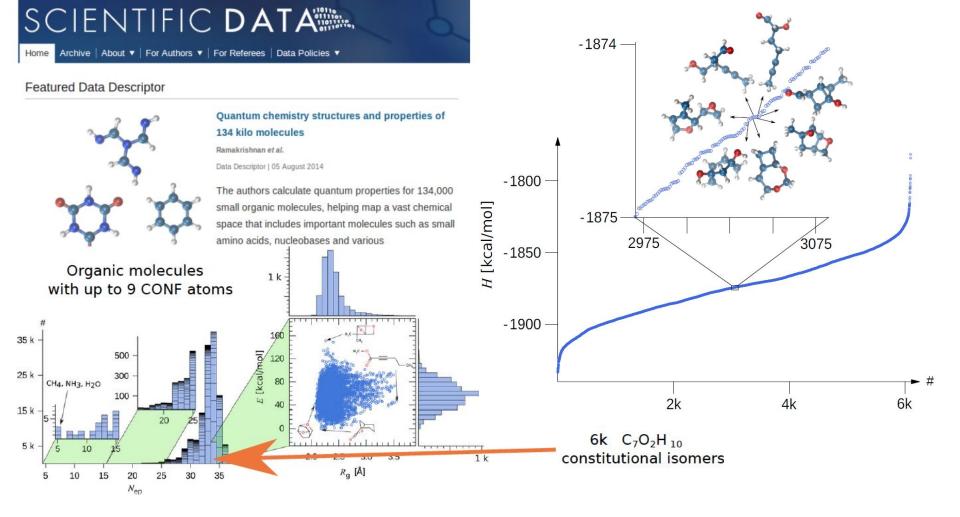
Train

K.-R. Mueller et al, Neural Comput (1996)

$$\rightarrow \log(\text{Error}) = \log(a) - b\log(N)$$

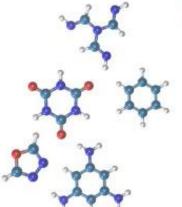






"Enumeration surpasses imagination"
J.-L. Reymond





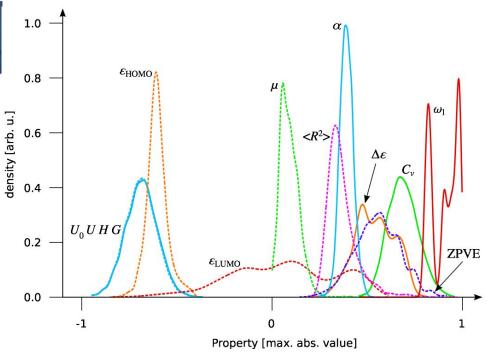
Quantum chemistry structures and properties of 134 kilo molecules

Ramakrishnan et al.

Data Descriptor | 05 August 2014

The authors calculate quantum properties for 134,000 small organic molecules, helping map a vast chemical space that includes important molecules such as small amino acids, nucleobases and various pharmaceutically-relevant organic building blocks.

These data can be used as a benchmark in the development of new methods in computational chemistry and molecular materials design



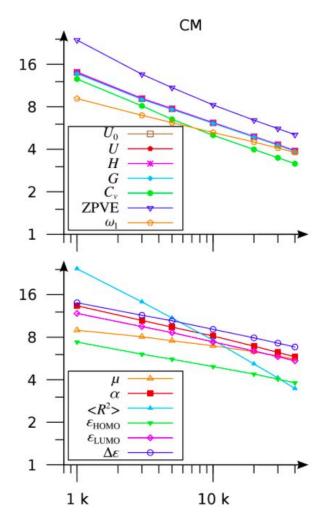
$$p_q = \sum_{t=1}^N c_t^p K_{qt}$$
 $\mathbf{c}^p = (\mathbf{K} + \lambda \mathbf{I})^{-1} \, \mathbf{p}^r$ We set $\lambda = 0 \dots$

$$\mathcal{L} = (\mathbf{p}^r - \mathbf{K}\mathbf{c}^p)^{\mathrm{T}} (\mathbf{p}^r - \mathbf{K}\mathbf{c}^p) + \lambda \mathbf{c}^{p\mathrm{T}}\mathbf{K}\mathbf{c}^p$$
$$[\mathbf{c}^{p_1}\mathbf{c}^{p_2}\dots\mathbf{c}^{p_n}] = \mathbf{K}^{-1} [\mathbf{p}_1^r\mathbf{p}_2^r\dots\mathbf{p}_n^r] \quad \Rightarrow \quad \mathbf{C} = \mathbf{K}^{-1}\mathbf{P}^r$$

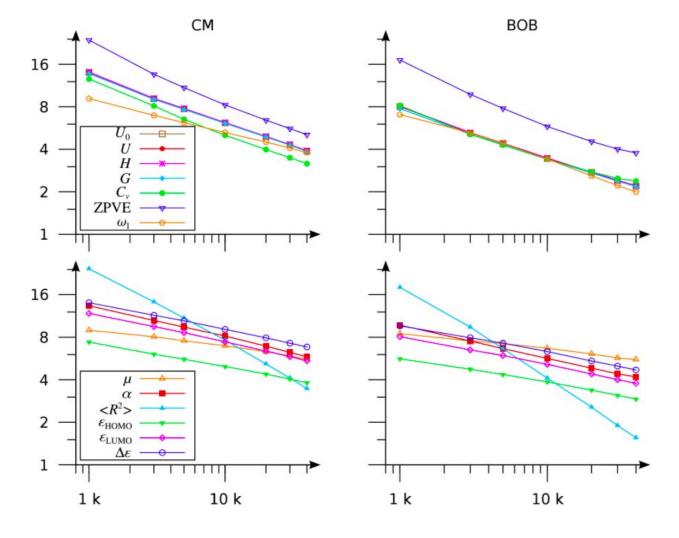
$$k_{ij} = e^{-D_{ij}/\sigma}$$

$$\frac{1}{2} \le k_{ij} \le 1$$

$$\sigma_{\text{opt}} = D_{ij}^{\text{max}}/\log(2)$$

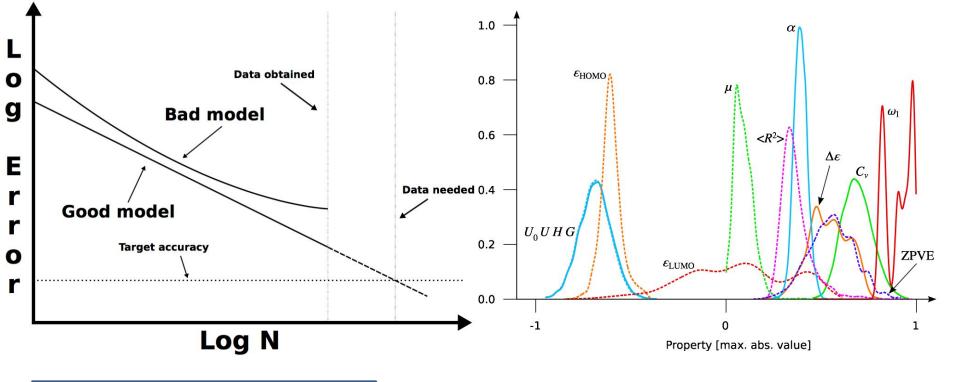


Tested on 134k-N organic molecules taken from: Ramakrishnan et al, Scientific Data (2014)



Tested on 134k-N organic molecules taken from: Ramakrishnan et al, Scientific Data (2014)

*BOB, Hansen et al, submitted (2015)



$$P^{\text{est}}(\mathbf{M}) = \sum_{i} \alpha_{i} k(\mathbf{M}, \mathbf{M}_{i})$$
$$\vec{\alpha} = \mathbf{K}^{-1} \vec{P}^{\text{ref}}$$
$$\sigma = \max \{ |\mathbf{d}_{i} - \mathbf{d}_{j}| \} / \log(2)$$

Error
$$\sim a/N^b$$

K.-R. Mueller et al, Neural Comput (1996)

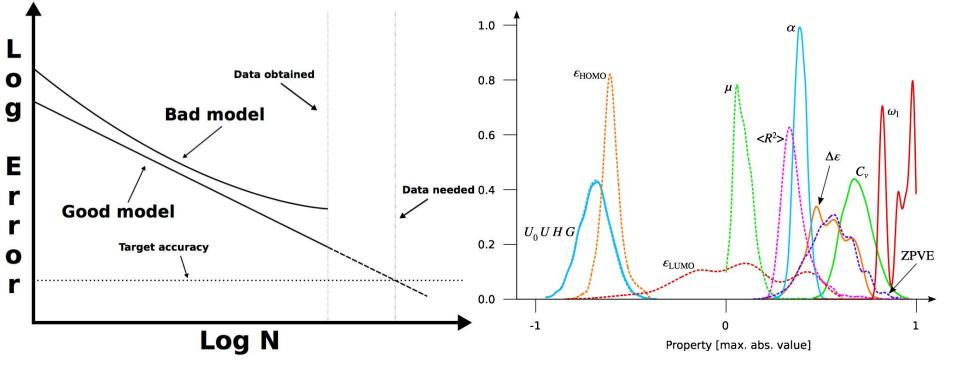
$$\rightarrow \log(\text{Error}) = \log(a) - b\log(N)$$

$$E_q = \langle \Psi_q | \hat{H} | \Psi_q \rangle$$

$$O_q = \langle \Psi_q | \hat{O} | \Psi_q \rangle$$

$$\alpha \sim \hat{O}$$

Ramakrishnan, OAvL, CHIMIA (2015)



$$P^{\text{est}}(\mathbf{M}) = \sum_{i} \alpha_{i} k(\mathbf{M}, \mathbf{M}_{i})$$
$$\vec{\alpha} = \mathbf{K}^{-1} \vec{P}^{\text{ref}}$$
$$\sigma = \max \{ |\mathbf{d}_{i} - \mathbf{d}_{j}| \} / \log(2)$$

Error
$$\sim a/(N')^b$$
, e.g. $N' = N/x$

K.-R. Mueller et al, Neural Comput (1996)

$$\rightarrow \log(\text{Error}) = \log(a) + 3x - b \log(N)$$

Possible reasons for large a and $b \rightarrow 0$

- 1. No cause and effect relationship (spurious)
- 2. Bad representation (no physics/uniqueness ...)
- 3. Bad data (noisy/not representative/ ...)
- 4. Bad regressor: Underfitting (too rigid)/Overfitting (``crazy'' assumptions)/Unconverged (Less coefficients than data points)
- 5. High dimensionality and curvature

More ways to be wrong than right

Conclusions

1. Instantaneous QM quality predictions

2. Learning curves reveal quality of ML model

3. Representations

4. Data sets

Kernel Ridge Regression

Kernel

$$E^{est}(\mathbf{M}) = \sum_{i}^{N} \alpha_i k(\mathbf{M}, \mathbf{M}_i)$$

e.g.
$$k(\mathbf{M}, \mathbf{M}') = \exp\left(-\frac{d(\mathbf{M}, \mathbf{M}')^2}{2\sigma^2}\right)$$

Regression

$$\min_{\alpha} \left(\sum_{i} \left(E^{est}(\mathbf{M}_{i}) - E_{i}^{ref} \right)^{2} + \lambda \sum_{ij} \alpha_{i} \alpha_{j} k(\mathbf{M}_{i}, \mathbf{M}_{j}) \right)$$

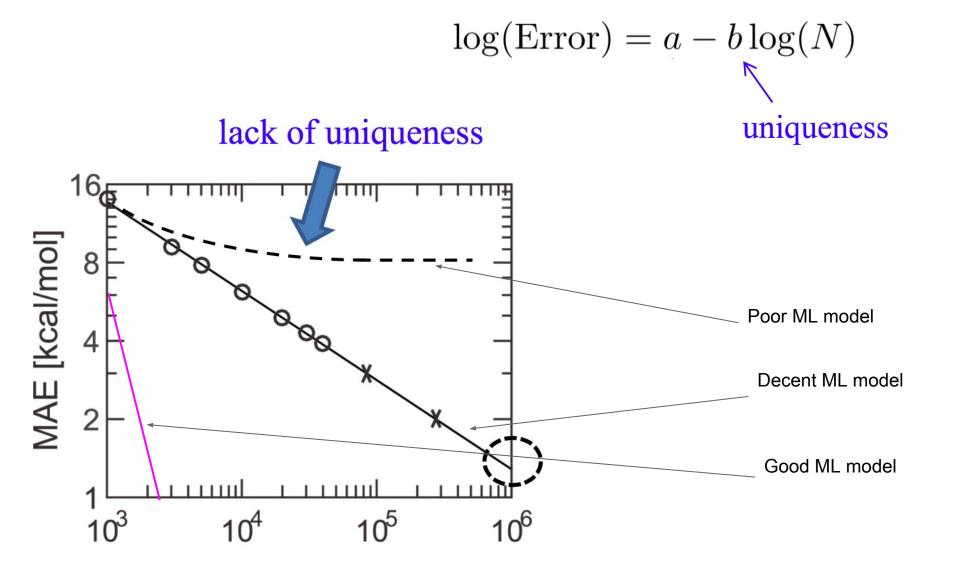
Solution

$$\alpha = (\mathbf{K} + \lambda \mathbf{I})^{-1} \mathbf{E}^{ref}$$

- i. Let D denote a descriptor, that is, not unique. Then, two systems $H_1 \neq H_2$ exist that differ in excess of the invariants, but they are mapped to the same descriptor value d, $H_1 \rightarrow d$ and $H_2 \rightarrow d$.
- ii. Because H_1 and H_2 differ by more than their property's invariances, they have different wave-functions, $\Psi_1 \neq \Psi_2$, yielding two different observables, $\mathcal{O}_1 = \langle \Psi_1 | \hat{O} | \Psi_1 \rangle$ and $\mathcal{O}_2 = \langle \Psi_2 | \hat{O} | \Psi_2 \rangle$. Here, we deliberately ingore the obvious exception and special situation of all observables which happen to be degenerate.
- iii. A trained statistical model predicts any observable \mathcal{O} solely based on descriptor input d leading to identical predictions $\mathcal{O}_1^{\text{pred}} = \mathcal{O}_2^{\text{pred}}$. In the limit of infinite training data, these predictions will be exact, implying $\mathcal{O}_1 = \mathcal{O}_2$, in contradiction to (ii).

lack of uniqueness → absurd results → noise in training

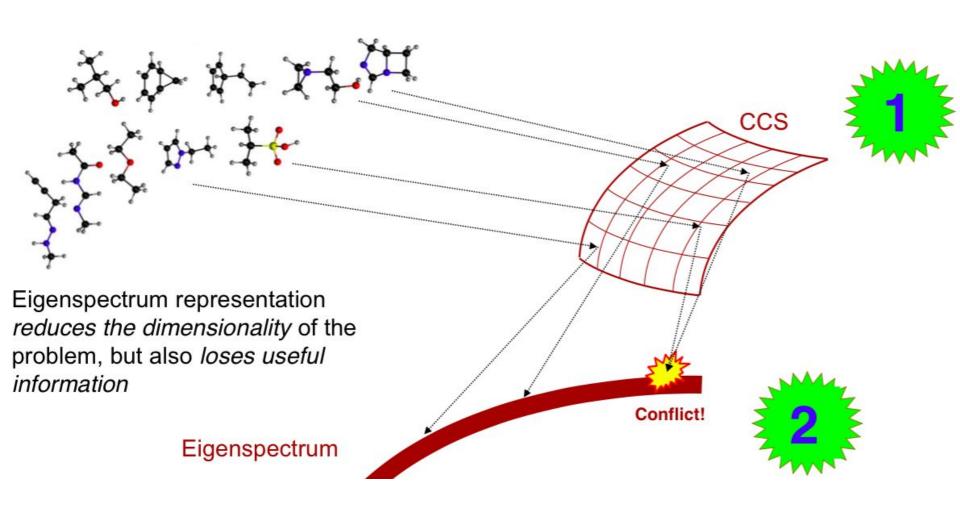
OAvL et al, IJQC (2013)

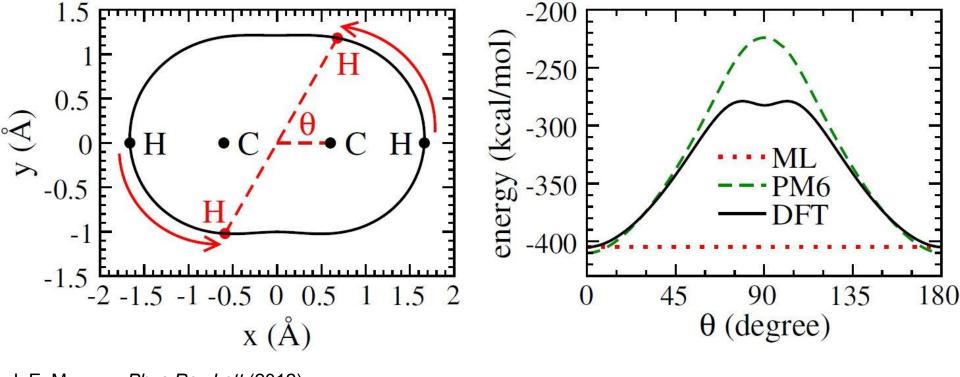


lack of uniqueness \rightarrow absurd results \rightarrow noise in training

OAvL et al, *IJQC* (2013)

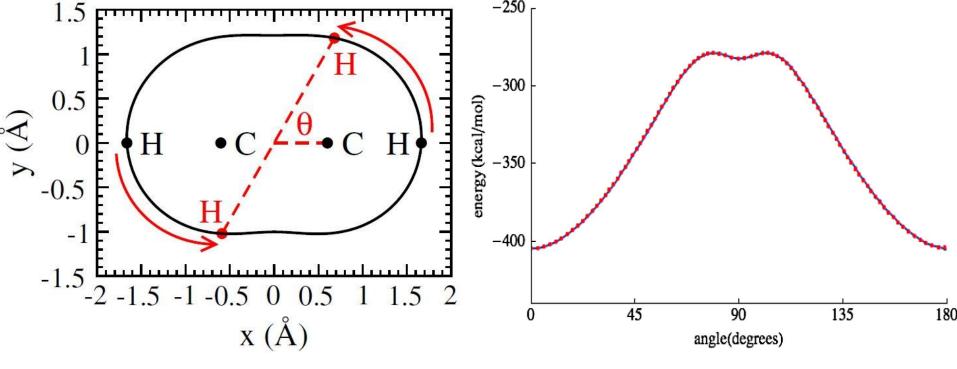
Huang, OAvL, J Chem Phys Comm (2016) arxiv.org/abs/1608.06194





J. E. Moussa, *Phys Rev Lett* (2012)

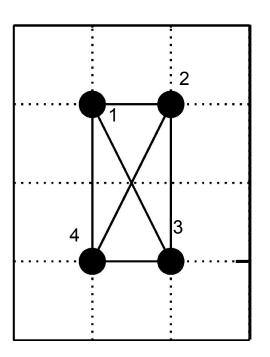
$$M_{IJ} = \begin{cases} 0.5Z_I^{2.4} & \forall \quad I = J, & \text{Coulomb-matrix} \\ \frac{Z_IZ_J}{|\mathbf{R}_I - \mathbf{R}_J|} & \forall \quad I \neq J. \end{cases}$$
 Coulomb-matrix • unique • translation • rotation • rotation • symmetry • diagonalize • fill up w zeros

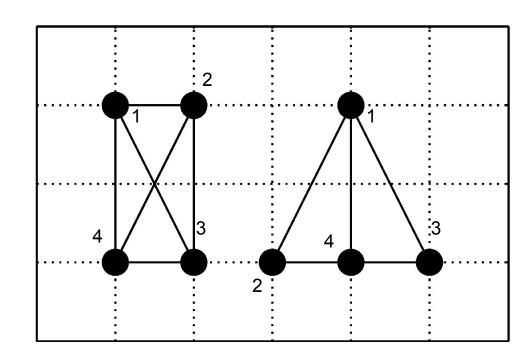


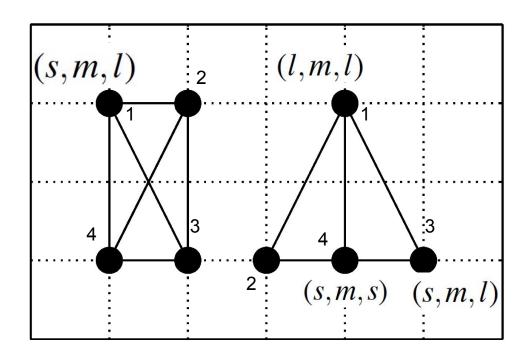
J. E. Moussa, Phys Rev Lett (2012)

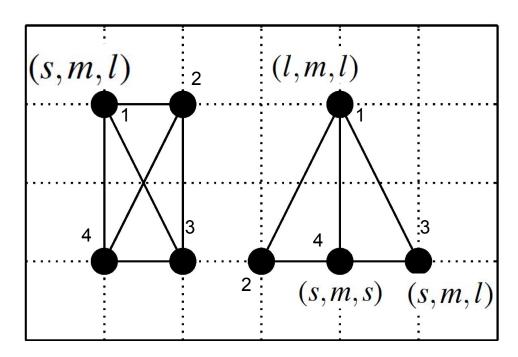
$$M_{IJ} = egin{cases} 0.5Z_I^{2.4} & orall & I = J, & \text{Coulomb-matrix} \\ \frac{Z_IZ_J}{|\mathbf{R}_I - \mathbf{R}_J|} & orall & I \neq J. & \text{unique} \\ N = 4 & \text{otanslation} \\ -> 3*N-6 = 6 \text{ degrees of freedom} & \text{otanslation} \\ & \text{$$

- rotation
- symmetry
- diagonalize sort
- fill up w zeros







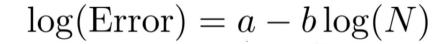


$$M_{IJ} = \begin{cases} 0.5Z_I^{2.4} & \forall I = J, \\ \frac{Z_I Z_J}{|\mathbf{R}_I - \mathbf{R}_J|} & \forall I \neq J. \end{cases}$$

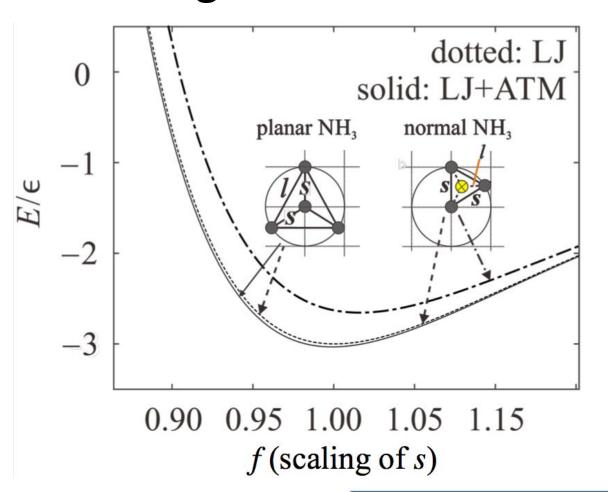
		s	-	m
	S		m	I
	I	m		S
	m	I	s	

	I	I	m
		m	S
I	m		S
m	s	S	

Learning curves

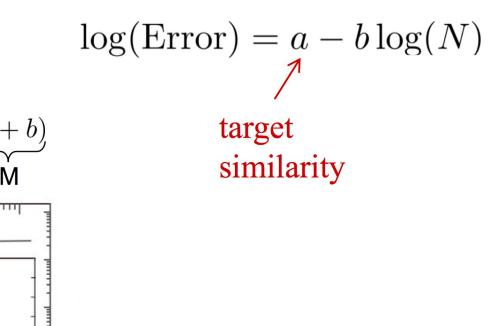


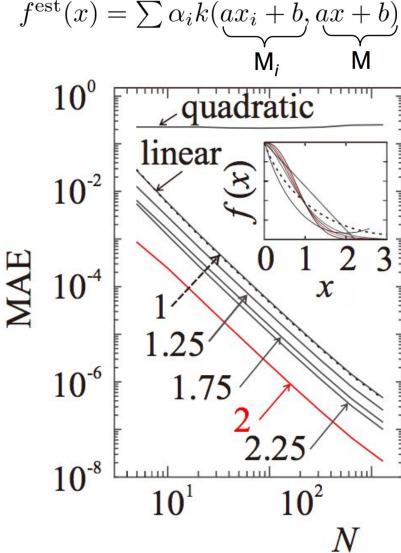




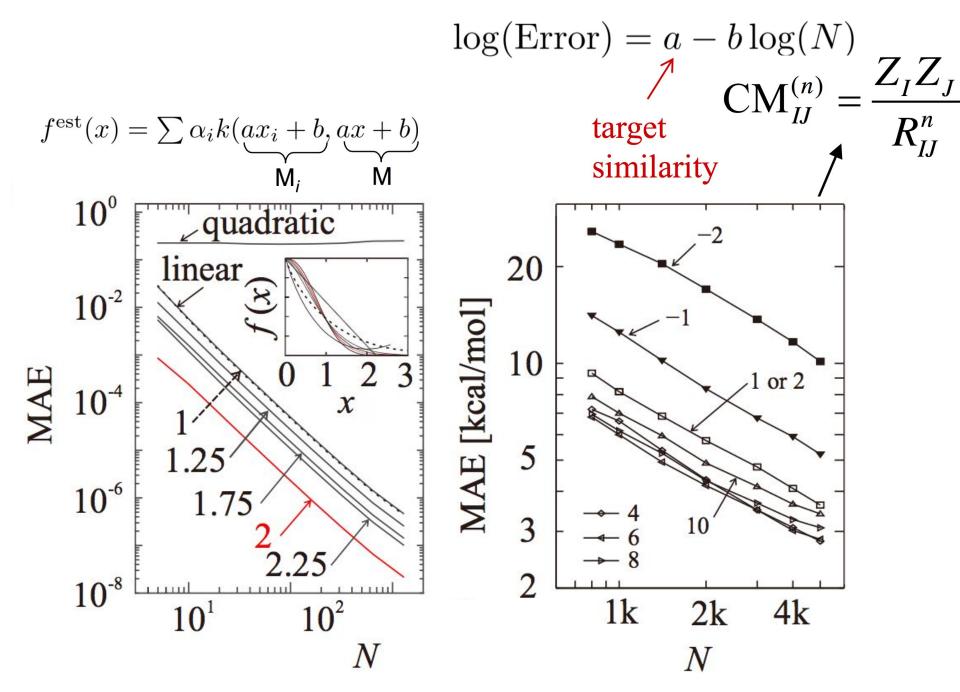
LJ: Lennard-Jones 2-body vdW potential

ATM: Axilrod-Teller-Muto 3-body vdW potential





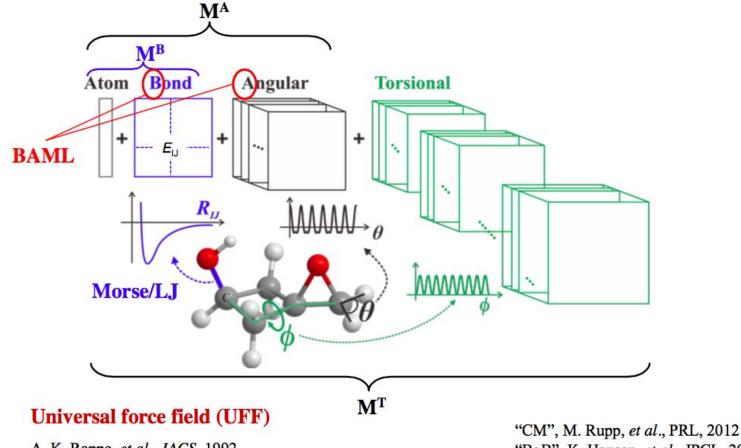
Huang, OAvL, J Chem Phys Comm (2016) arxiv.org/abs/1608.06194



Huang, OAvL, J Chem Phys Comm (2016) arxiv.org/abs/1608.06194

Approach: best M is unique AND good model

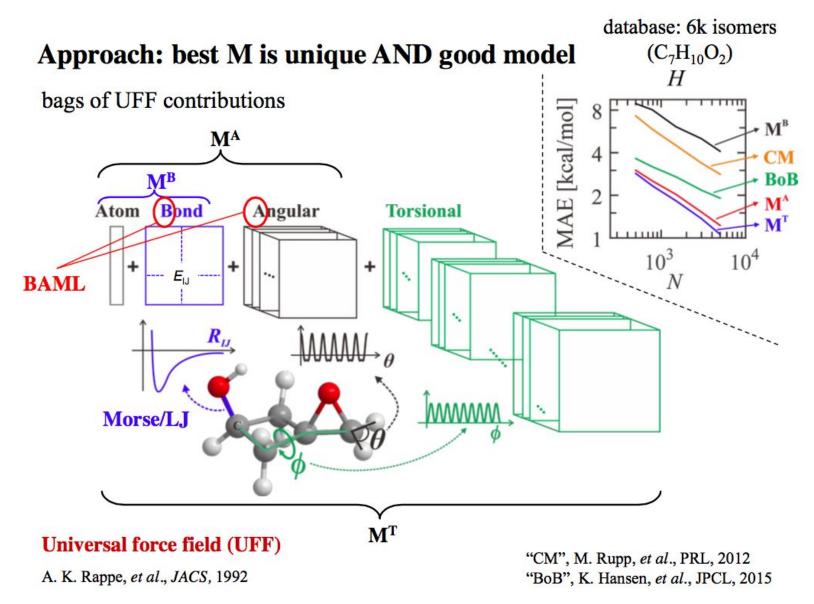
bags of UFF contributions



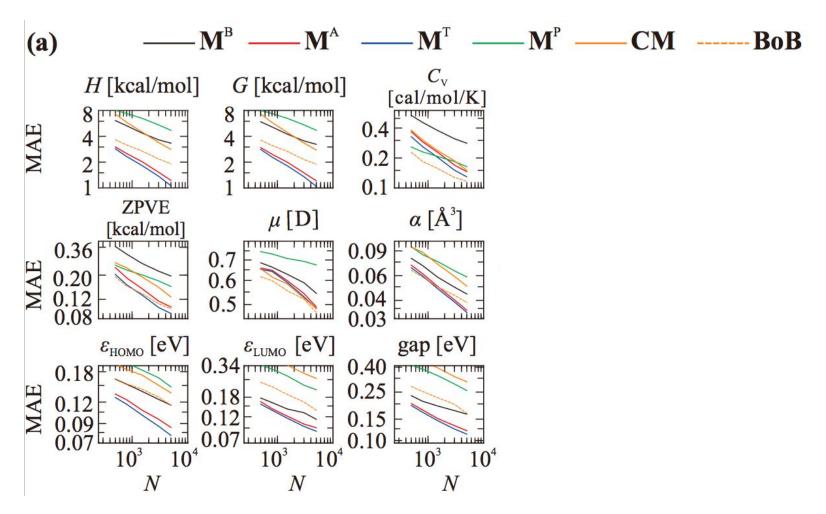
A. K. Rappe, et al., JACS, 1992

"BoB", K. Hansen, et al., JPCL, 2015

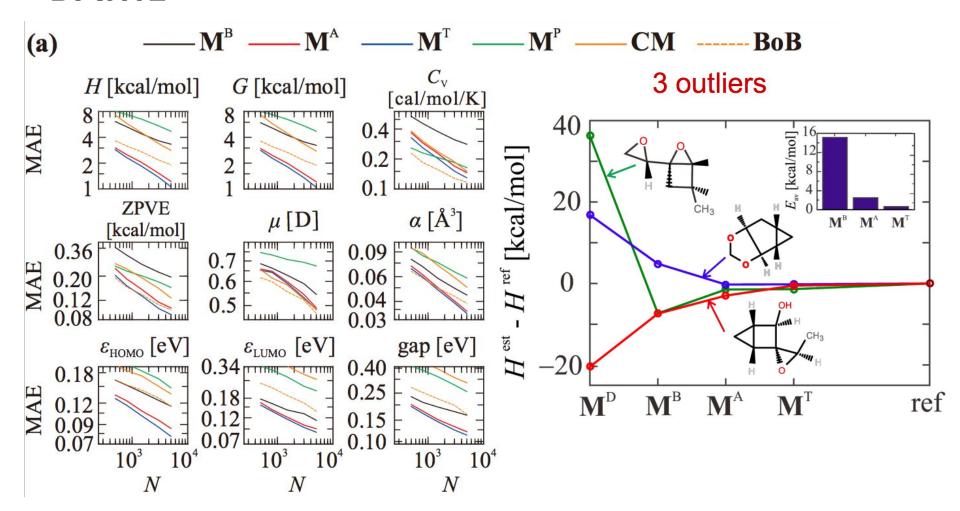
Huang, OAvL, J Chem Phys Comm (2016) arxiv.org/abs/1608.06194



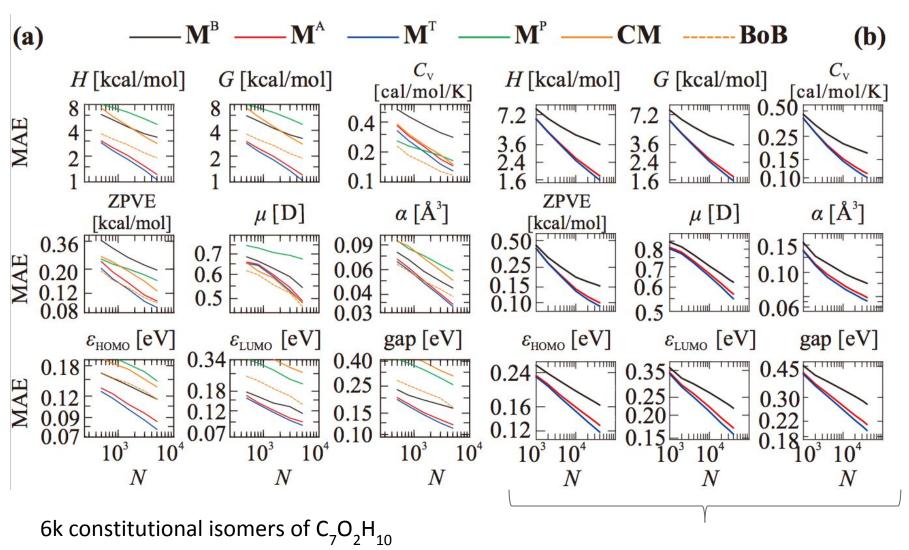
Huang, OAvL, J Chem Phys Comm (2016) arxiv.org/abs/1608.06194



6k constitutional isomers of $C_7O_2H_{10}$



6k constitutional isomers of $C_7O_2H_{10}$



QM9 (134k molecules)

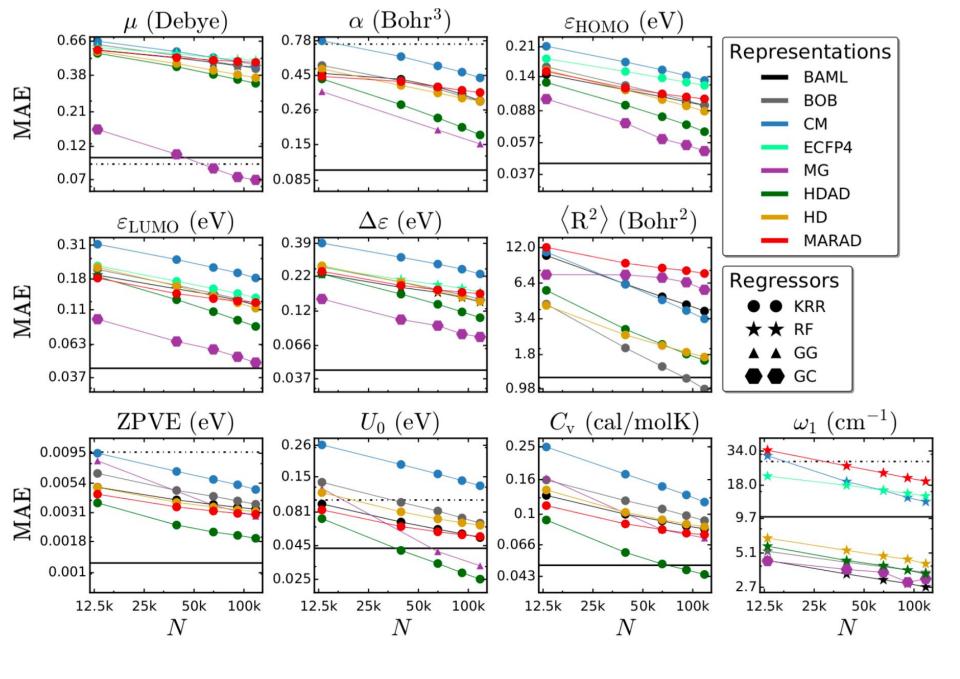
QM7b database (size: 7211)

MAE (5k out-of-sample)

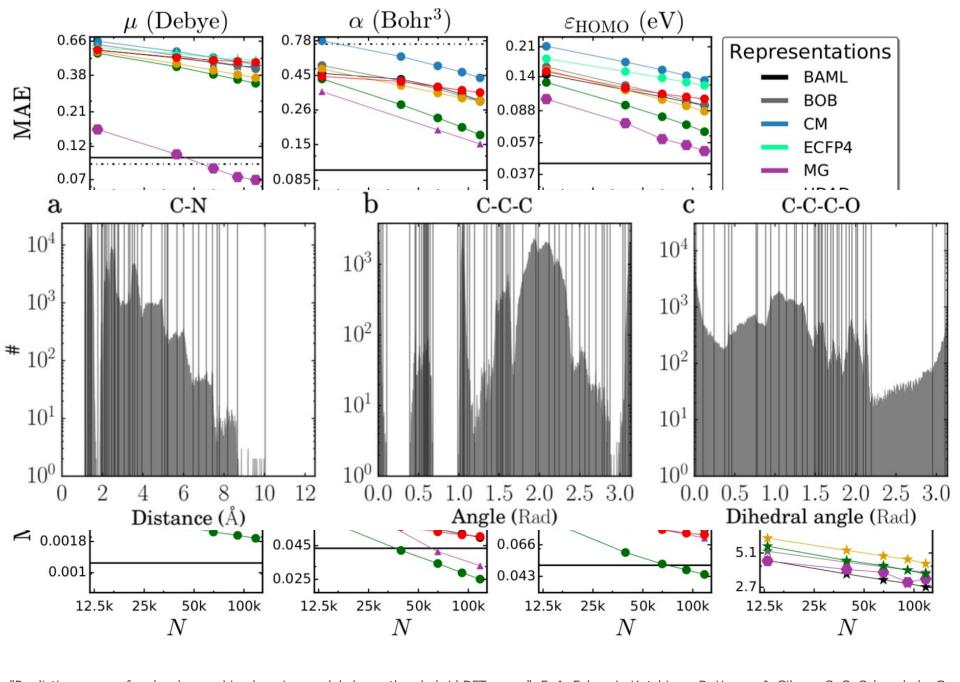
	BAML	BoB	SOAP ^a	CM ^b	$accuracy^b$
E (PBE0)/eV	0.05	0.08	0.04	0.16	0.15, 0.23, 0.09-0.22
α (PBE0)/ Å ³	0.07	0.09	0.05	0.11	0.05-0.27, 0.04-0.14
HOMO (GW)/eV	0.10	0.15	0.12	0.16	-
LUMO (GW)/eV	0.11	0.16	0.12	0.16	1.00
IP (ZINDO)/eV	0.15	0.20	0.19	0.17	0.20, 0.15
EA (ZINDO)/eV	0.07	0.17	0.13	0.11	0.16, 0.11
E_{1st} * (ZINDO)/eV	0.13	0.21	0.18	0.13	0.18, 0.21

^a S. De, et al., PCCP, 2016

^b G. Montavon, et al., NJP, 2013



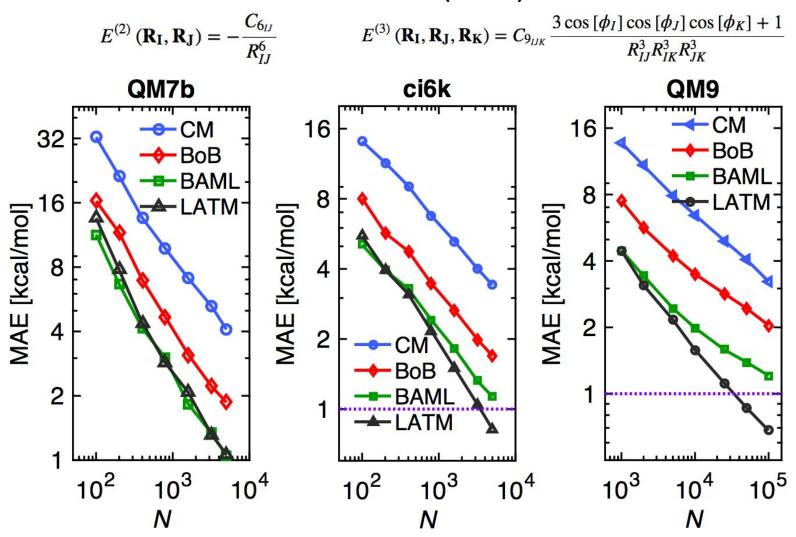
"Prediction errors of molecular machine learning models lower than hybrid DFT errors", F. A. Faber, L. Hutchison, B. Huang, J. Gilmer, S. S. Schoenholz, G. E. Dahl, O. Vinyals, S. Kearnes, P. F. Riley, OAvL arxiv.org/abs/1702.05532



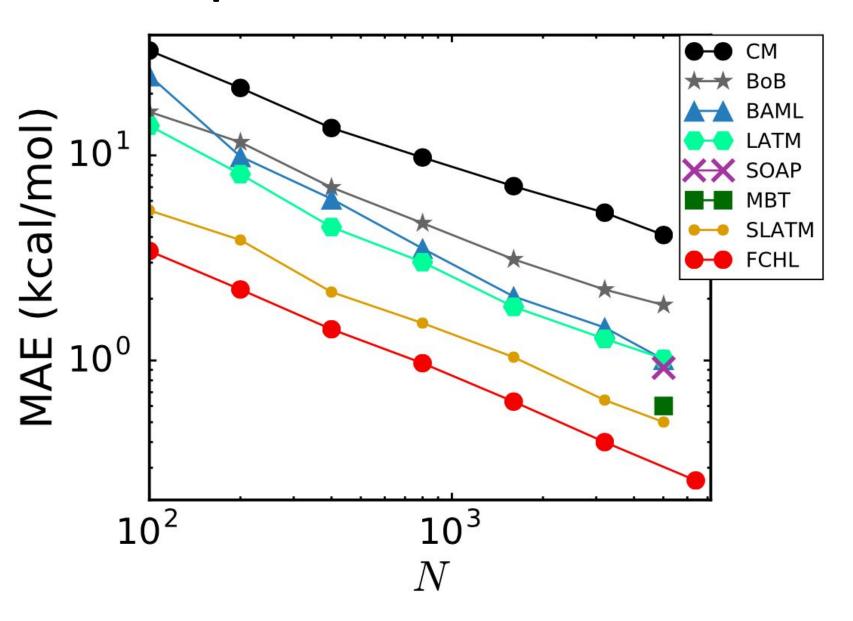
"Prediction errors of molecular machine learning models lower than hybrid DFT errors", F. A. Faber, L. Hutchison, B. Huang, J. Gilmer, S. S. Schoenholz, G. E. Dahl, O. Vinyals, S. Kearnes, P. F. Riley, OAvL arxiv.org/abs/1702.05532

LATM

Atoms + London + Axilrod-Teller-Muto (LATM)



Current performance on QM9



Faber, Christensen, Huang, OAvL, to be submitted (2017)

Representation leading to low a and b

1. Unique

2. Similar to target

3. Efficient

4. ...

More ways to be wrong than right

Conclusions

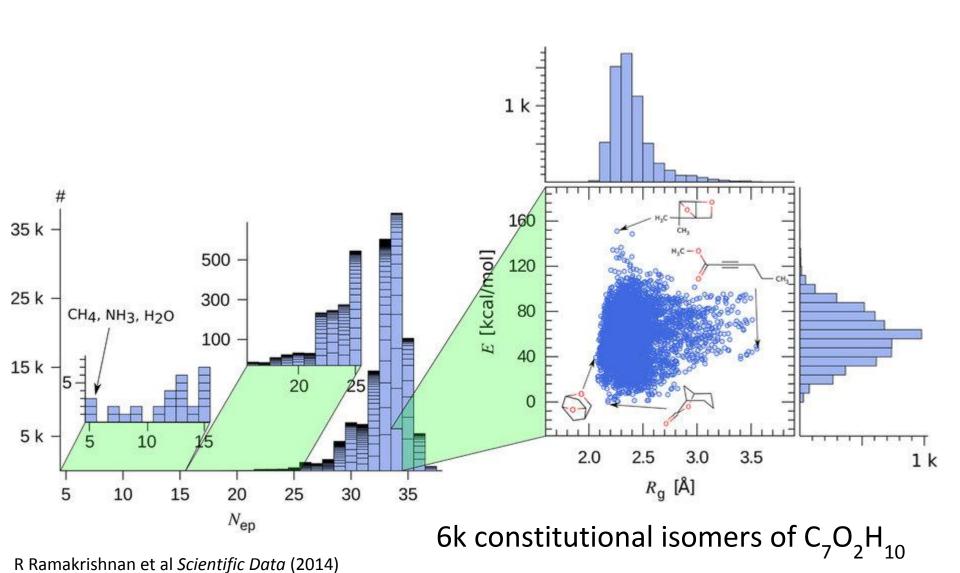
1. Instantaneous QM quality predictions

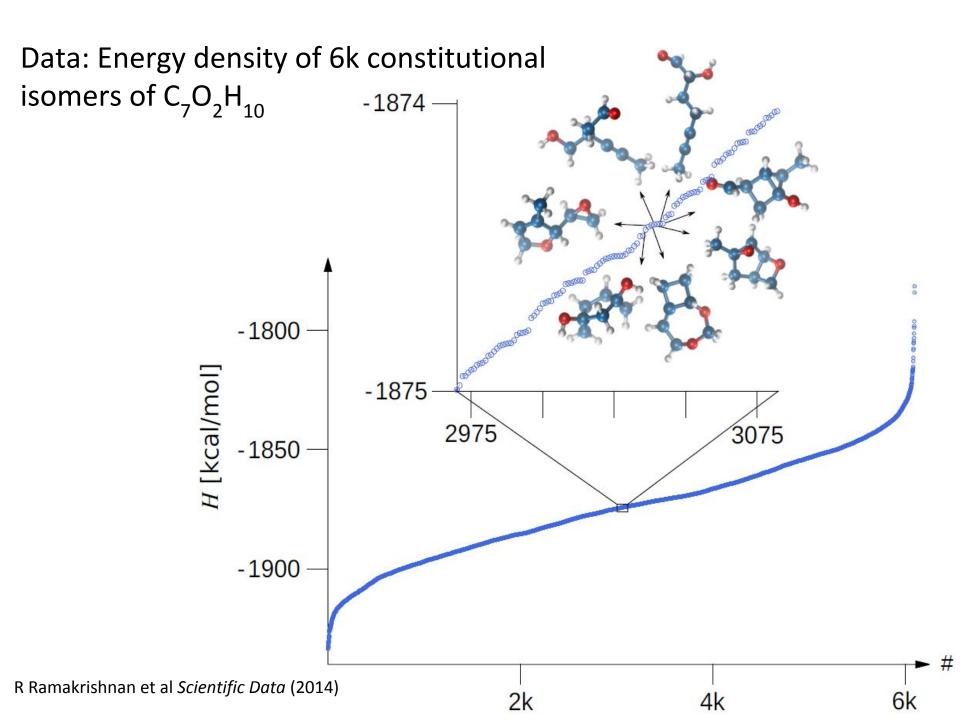
2. Learning curves reveal quality of ML model

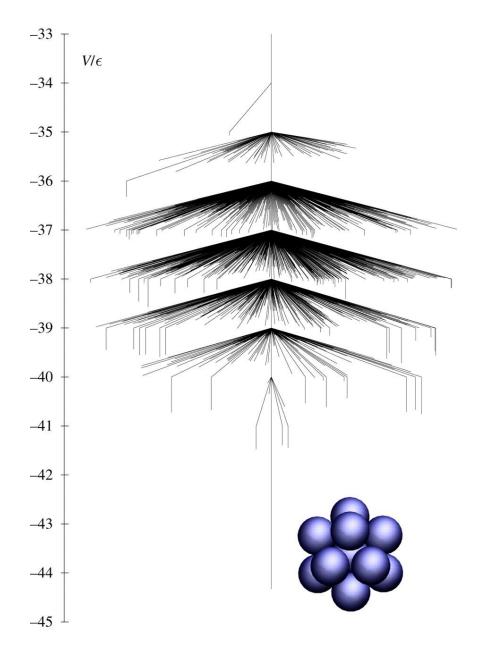
3. Representations

4. Data sets

Data: Smallest 134k organic molecules in GDB







D J Wales Philosophical Transactions A (2012)

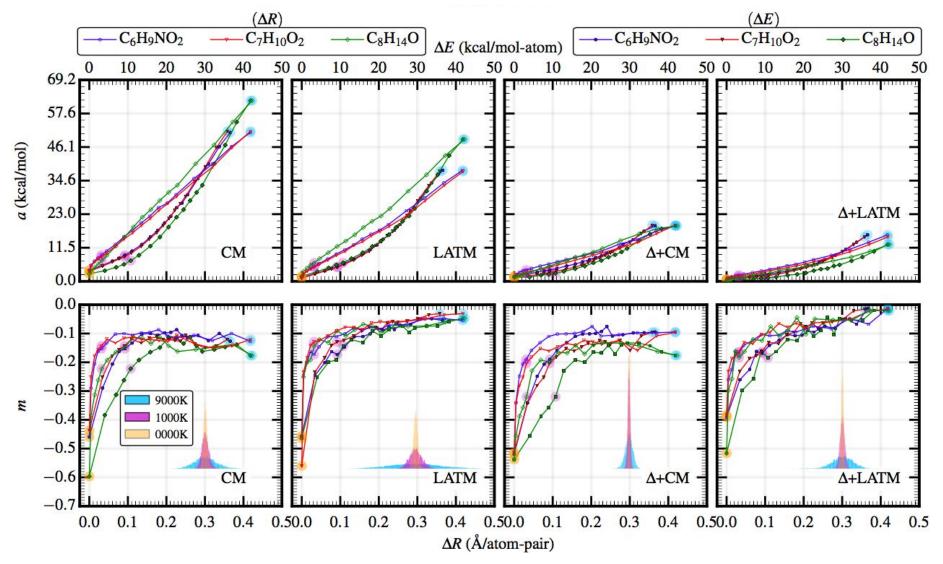
$\log(\epsilon) = \log(a) + m\log(N)$ **Temperature** 0.25 A 69.2 0.20 B E (kcal/mol) (3) 0.15 -92.3 $a(\varepsilon)$ -138.40.10 (log decal/mol) (a (kcal/mol) (a (kcal/mol) (b (kcal/mol) 23.1 0.05 0.00 0.0 3 5 d (Å) -0.2230.69 -0.423.07 2.307 MAE (kcal/mol) -0.6m 0.231 -0.80.023 0.002 Ar₆₀ LJ cluster -1.0Potential A Potential B -1.20.000 10 20 30 40 50 60 7 8 9 10 Number of perturbed atoms

R. Sarmiento-Perez, F. Faber, B. Huang, OAvL, to be submitted (2017)

Training set size

Temperature

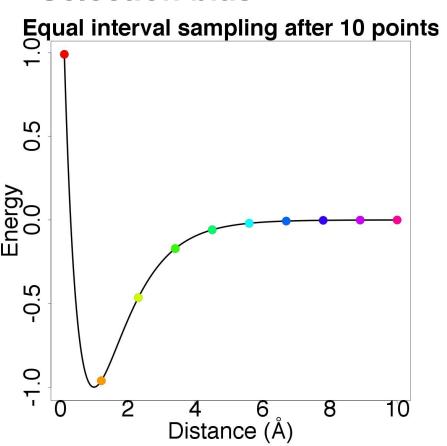
$$\log(\epsilon) = \log(a) + m\log(N)$$

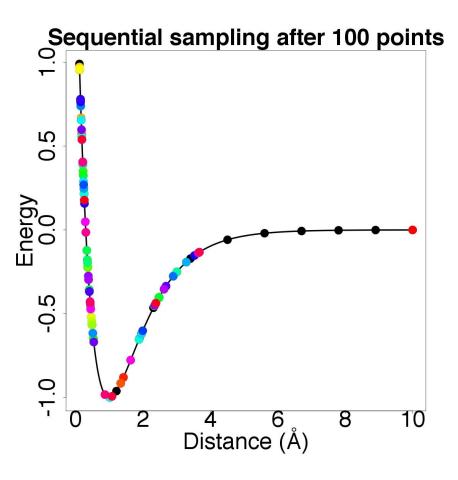


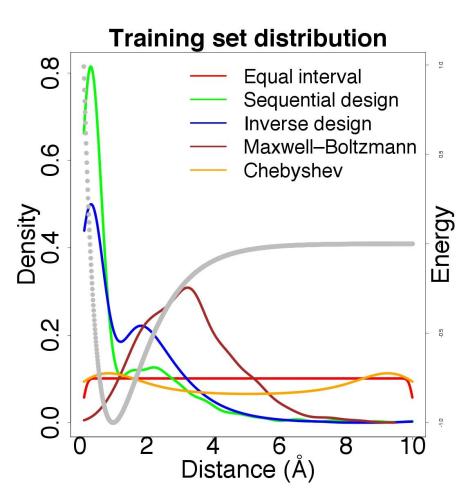
R. Sarmiento-Perez, F. Faber, B. Huang, OAvL, to be submitted (2017)

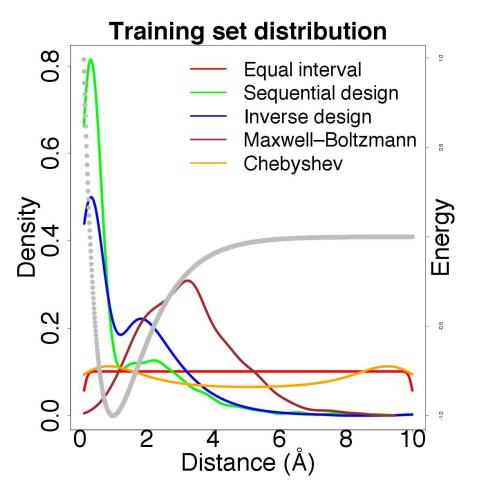
Selection bias

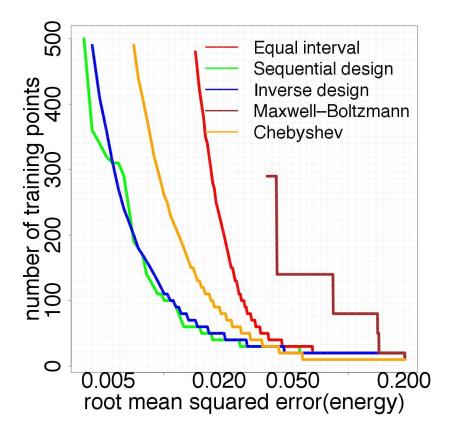


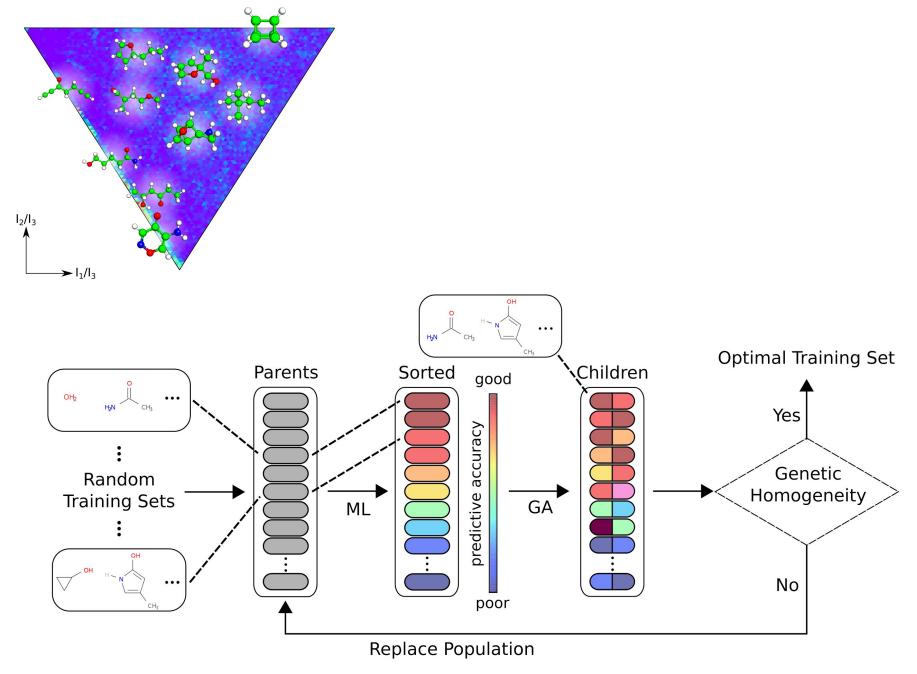












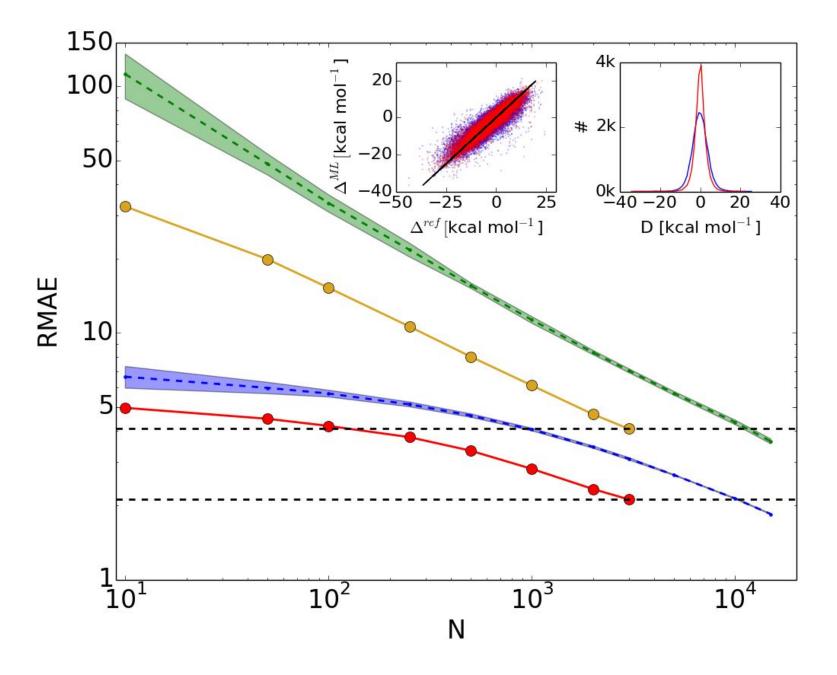
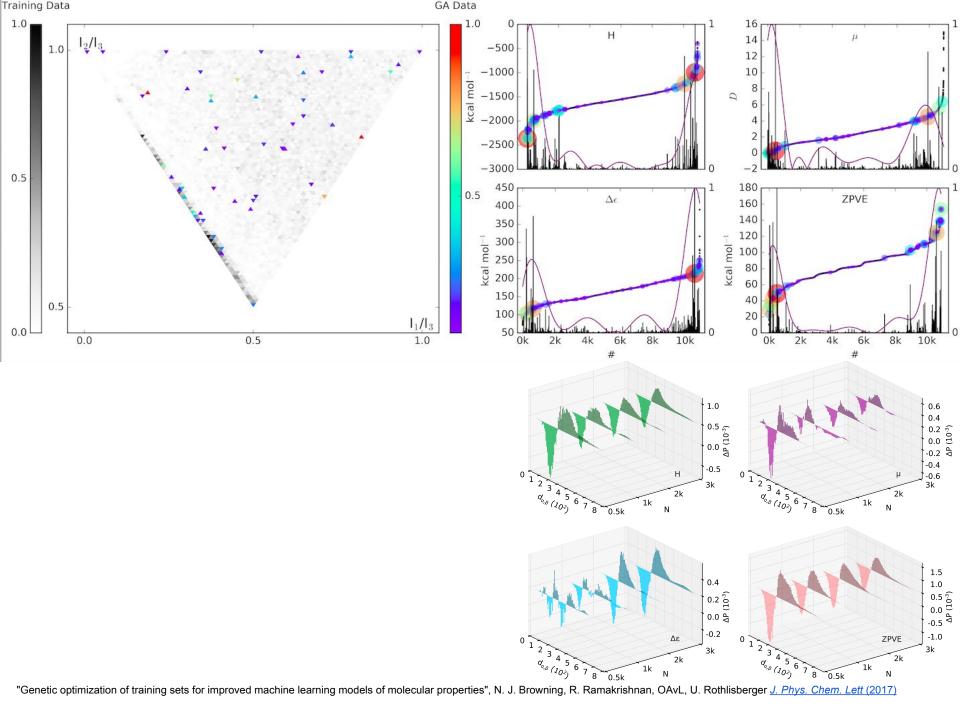
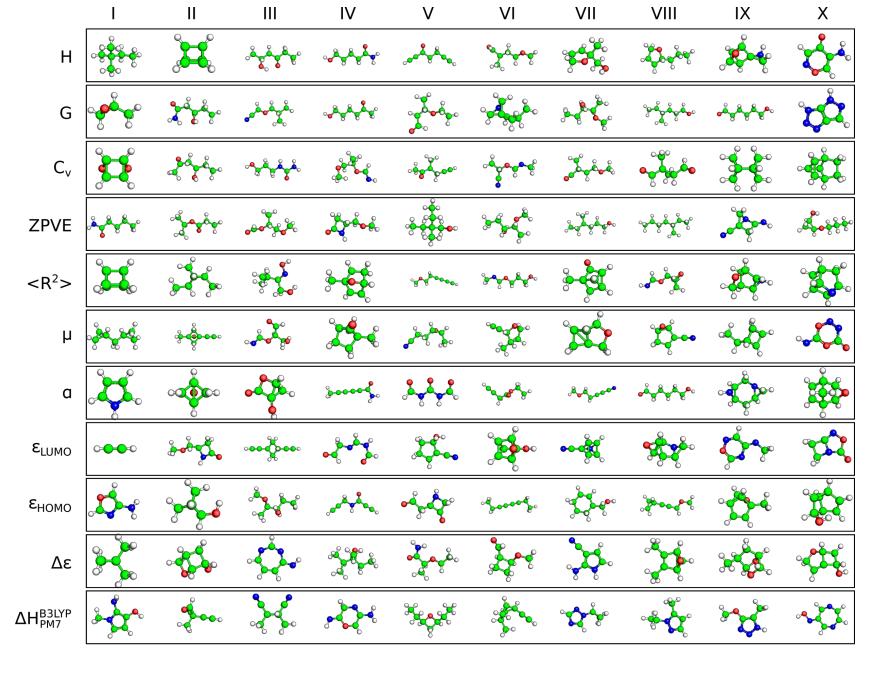
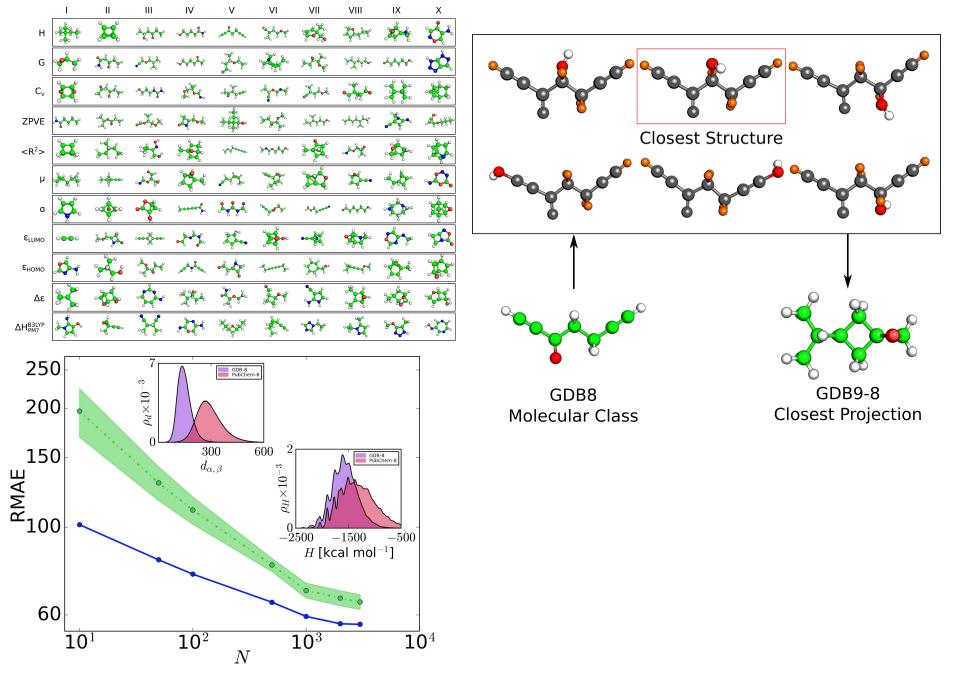


Table I. Randomized and GA-optimized out-of-sample relative mean absolute errors (RMAEs) for all properties. All target chemical accuracies are 1 kcal/mol, except for ZPVE, dipole moment and isotropic polarizability, which target accuracies of 10cm^{-1} , 0.1D and $0.1a_0^3$ respectively. GA-optimized RMAEs are denoted by P^{GA} while randomly generated training set MAEs are denoted as P^{ML} . Final row corresponds to out-of-sample RMAEs for enthalpy of atomization H using Δ_{PM7}^{B3LYP} -learning.

$P_{ m rand} \; (P_{ m GA})$	N						
	10	50	100	500	1k	2k	3k
H	113.0 (31.6)	48.0 (18.3)	33.3 (14.3)	14.8 ((7.5) 10.2	(5.8) 6.8	(4.5) 5.1 (3.9)
\overline{G}	101.8 (28.8)	44.0 (17.7)	31.4 (14.1)	14.3 ((7.5) 9.9	(5.6) 6.7	(4.3) 5.0 (3.9)
C_v	27.3 (14.5)	18.2 (9.4)	14.6 (7.8)	7.4 ((4.0) 5.2	(2.9) 3.4	(2.3) 2.5 (2.0)
ZPVE	10.1 (2.4)	4.3 (1.1)	2.8 (0.8)	0.9 ((0.4) 0.6	(0.3) 0.4	(0.2) 0.3 (0.1)
<r<sup>2></r<sup>	168.5 (92.2)	117.0 (44.2)	85.6 (33.2)	35.7 (1	9.1) 25.7 ((15.5) 18.3	(12.7) 14.5 (11.6)
$\overline{\mu}$	11.3 (8.5)	10.3 (7.7)	9.9 (7.4)	8.4 ((6.3) 7.5	(5.7) 6.2	(5.1) 5.2 (4.7)
α	40.8 (16.3)	23.1 (12.0)	18.5 (10.8)	11.8 ((7.8) 9.6	(6.5) 7.2	(5.4) 5.8 (4.9)
$\epsilon_{ m HOMO}$	13.0 (9.0)	11.2 (8.1)	10.4 (7.3)	7.7 ((5.2) 6.3	(4.5) 4.9	(3.8) 4.0 (3.5)
$\epsilon_{ m LUMO}$	22.3 (15.8)	18.8 (12.7)	17.0 (11.1)	11.9 ((8.0) 9.7	(6.7) 7.4	(5.6) 5.9 (5.0)
gap	24.0 (17.8)	20.8 (15.0)	19.5 (13.5)	14.3 ((9.8) 11.8	(8.1) 9.0	(6.8) 7.3 (6.2)
ΔH	6.6 (5.0)	6.0 (4.4)	5.7 (4.1)	4.6 ((3.2) 4.1	(2.6) 3.4	(2.1) 3.1 (1.9)



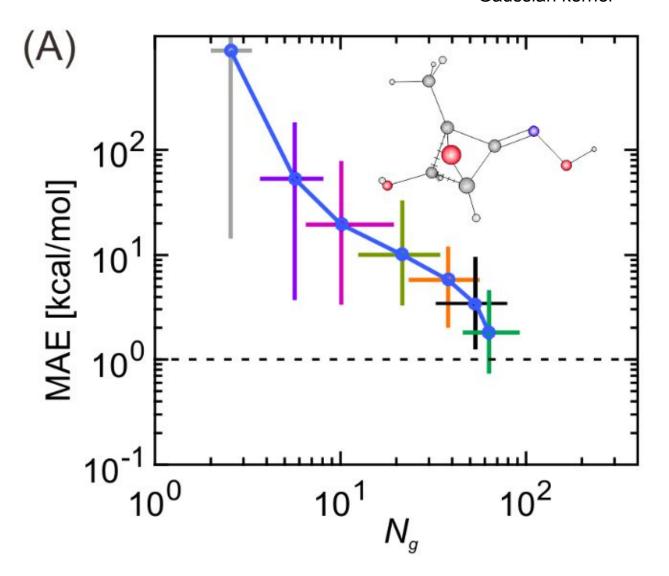




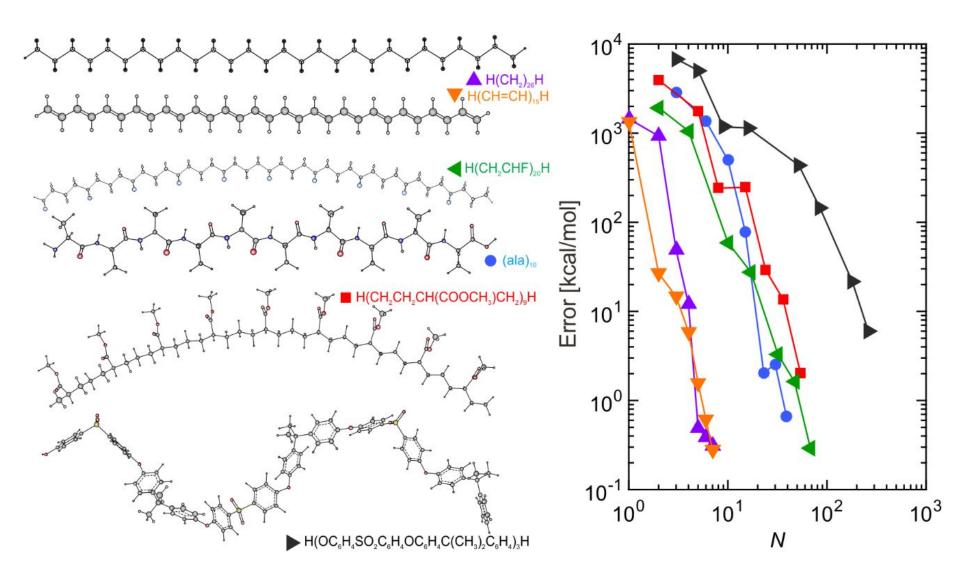
"Genetic optimization of training sets for improved machine learning models of molecular properties", N. J. Browning, R. Ramakrishnan, OAvL, U. Rothlisberger J. Phys. Chem. Lett (2017)

Effective

Errors on 1k QM9 predictions LATM representation Gaussian kernel



Effective



Data affects a and b

- 1. High dimensional function
- 2. Redundancy bias
- 3. Effective dimensionality
- 4. ...

More ways to be wrong than right

Conclusions

1. Instantaneous QM quality predictions

2. Learning curves reveal quality of ML model

3. Representations

4. Data sets

Conclusions II

Scientific method - proper way to gain knowledge

Inductive (Data)

- 1. Assume a law
- 2. Metric
- 3. Examples
- 4. Infer
- 5. New combination

Fast (ms)
Arbitrary reference
Automatic improvement

Transferable?
Minimally condensed

Deductive (Laws)

- 1. Assume a law
- 2. Approximate
- 3. Solve
- 4. Predict
- 5. New regimes

Slow (depending on approx.)
Approximation dependent
Human improvement

Transferable?

Maximally condensed