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DENR+POL:

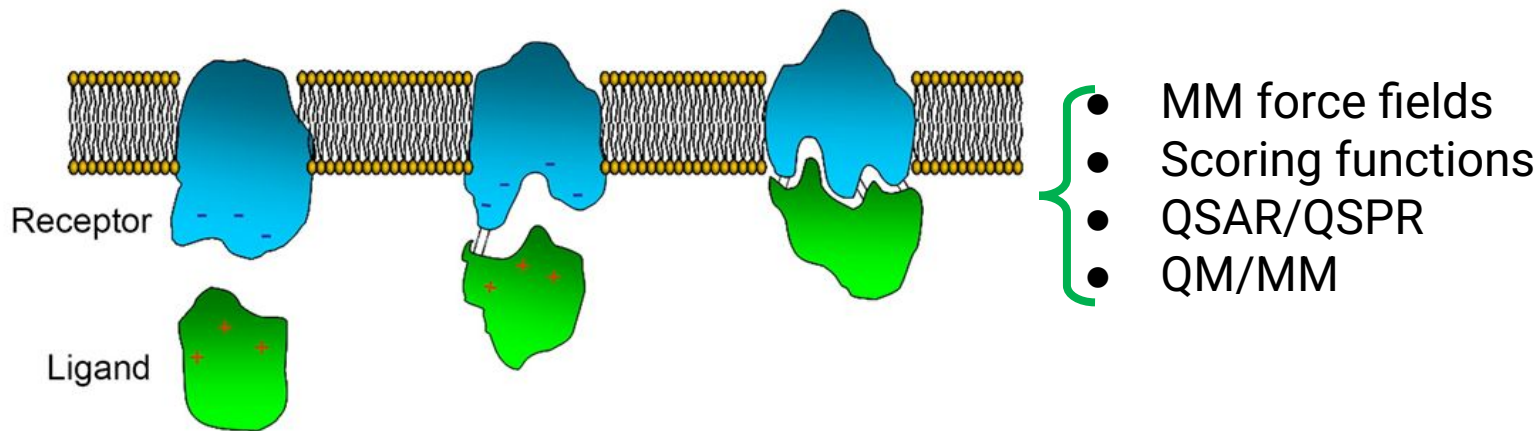
**Theoretically consistent polarizable empirical charges
for drug-like and biological molecules**

A.R. Shaimardanov, V.S. Frolov, D.A. Shulga, V.A. Palyulin

**XXXI Symposium on Bioinformatics and Computer-Aided Drug Discovery
(BCADD-2025)**

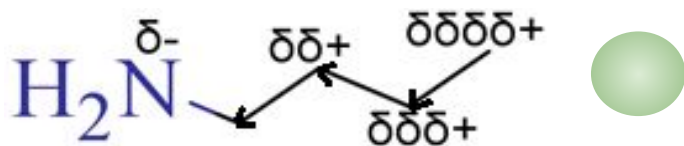
Empirical atomic charge models

- ❑ Atomic charges (q) – a popular approach to evaluate electrostatic interactions.
- ❑ No unambiguous definition => many charge calculation methods.
- ❑ For much medicinal chemistry goals - empirical atomic charge models.
- ❑ Advantages: high speed, moderate accuracy, interpretability.

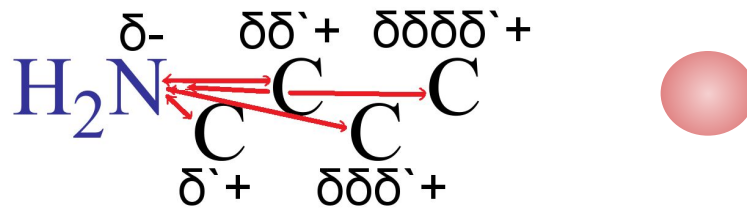


Electronic effects: an Essence of empirical charge models

- ❑ Electronic effects concept is a physical basis of empirical models.
- ❑ **An electronic effect** - hypothetic shift of electronic density due to mutual influence of some molecular fragments.
- ❑ An generalization of organic chemistry experience.
- ❑ Examples: inductive effect, polarization.
- ❑ The more correct the physics = the more accurate the method.
- ❑ Consistent description - the cornerstone of the effects world.



influence through σ -bonds



through electric field

Account for effects: Current situation

Charge model	Inductive effect	Polarization effect
EEM	-	+/-
AACT	?	+
SQE	?	+
SQE+Q0	?	+
ACKS2	?	+
PEOE	+	-
MMFF94 BCI	-	-
DENR	+	-
	+	+

Aim and objectives of the research

The aim:

To develop empirical charge calculation method, DENR+POL, that consistently describes inductive and polarization (Pol) effects in the drug-like molecules.

The objectives:

1. To study the polarization of intra- and intermolecular hydrogen bond (HB) in the drug-like molecules.
2. To determine the consistency of the effects description.
3. To identify the applicability of the model to drug-like molecules.
4. To reproduce QC molecular electrostatic potential (MEP) of the molecules.

Methods & Materials

Molecular set: 7K drug-likes -
PDBbind refined set v2020 +
subset of the MMFF94 molecule training set.

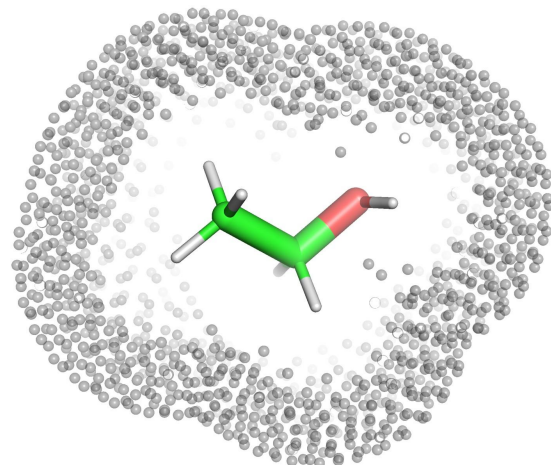
QC MEP as the reference.

Quality metric of a model:

$$\text{residual error}(m) = \text{MEP}_m^{\text{ref}} - \text{MEP}_m^{\text{emp}} = \text{MEP}_m^{\text{ref}} - \sum_i^{N_{\text{atoms}}} \frac{q_i}{R_{i,m}}, \mathbf{v}_m^{N_{\text{MEPpoints}}}$$

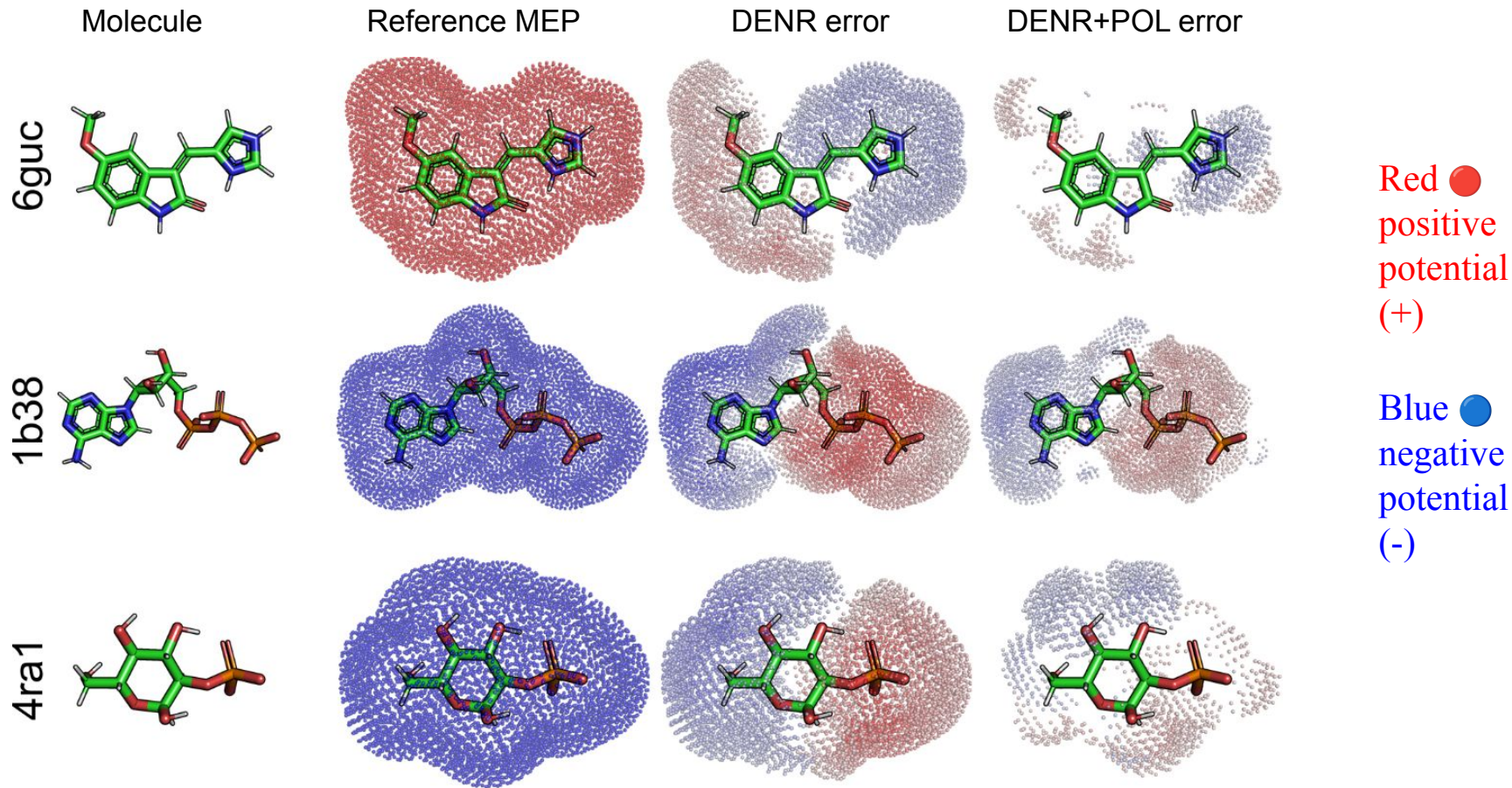
Selected molecules for analysis:

- 1) drug-like with intramolecular HB;
- 2) HB complex of two drug-like molecules.



MEP on the grid

Results: residual MEP reproduction



Results: Polarization of intramolecular HB

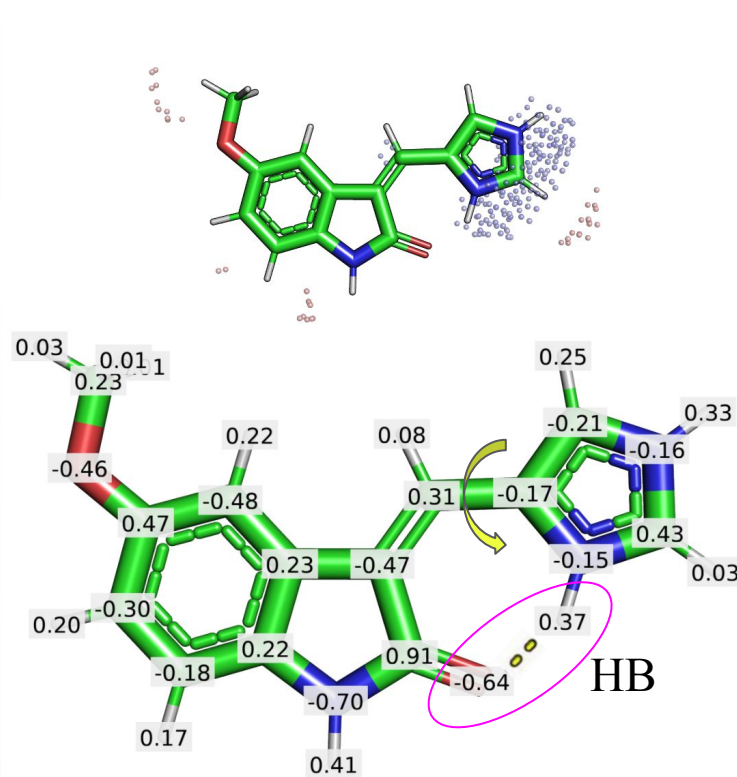
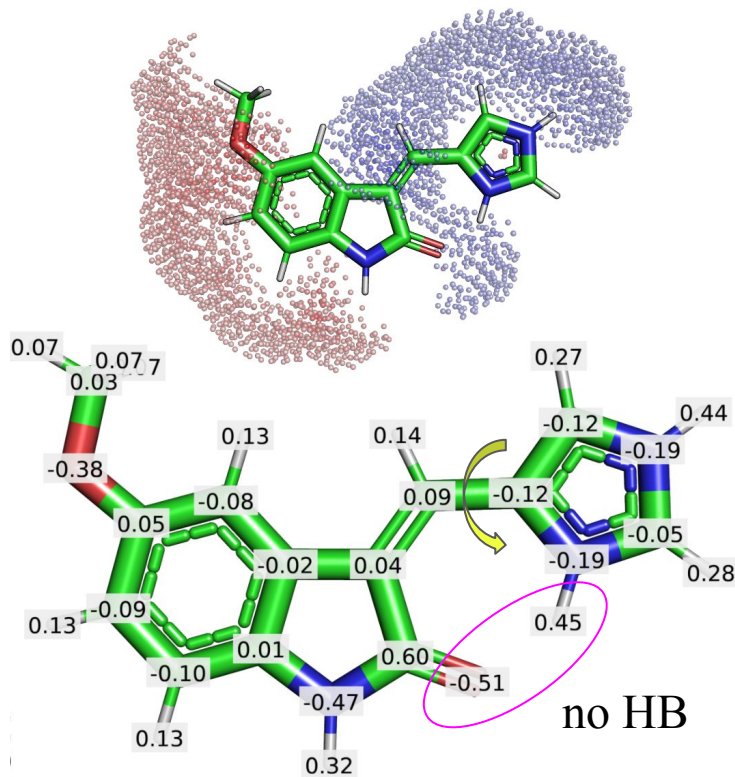
DENR

DENR+POL

0°

C=C-C-N

torsion



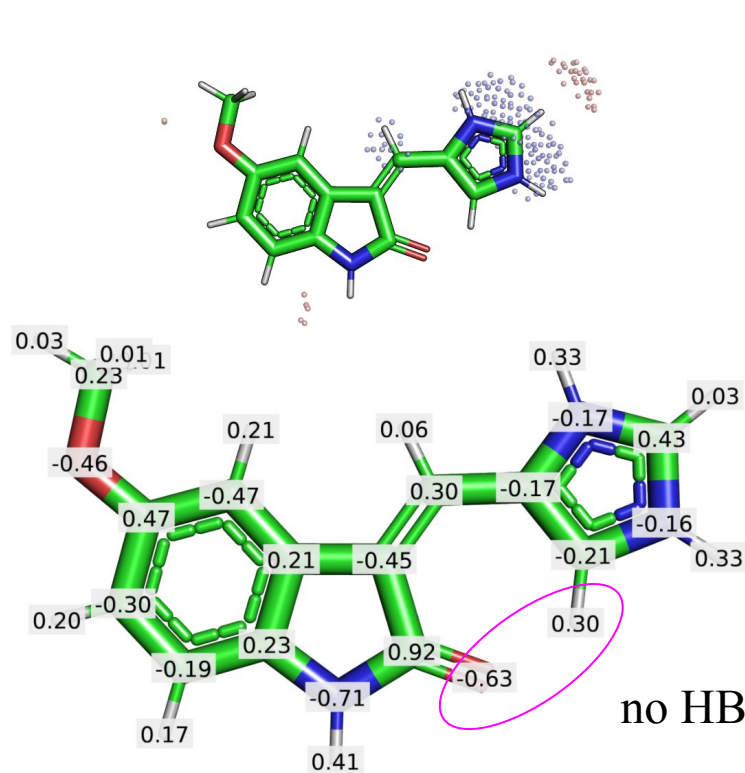
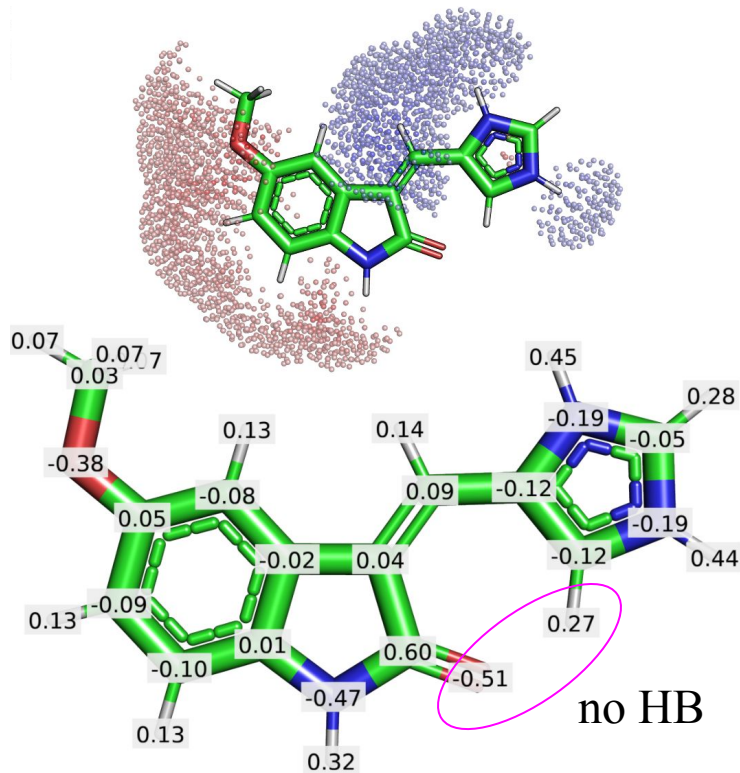
Results: Polarization of intramolecular HB

DENR

DENR+POL

C=C-C-N
torsion

180°

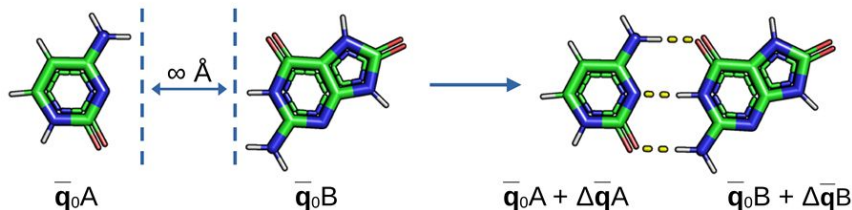


Results: Polarization of intermolecular HB

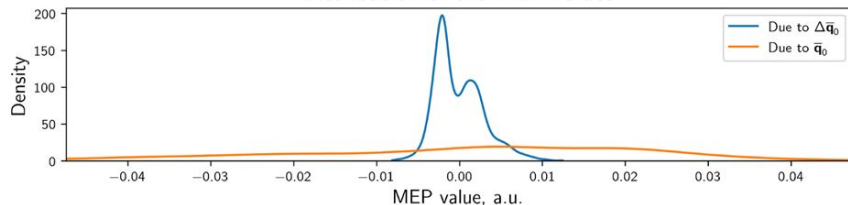
DENR+POL charges ($\bar{q}$)

Non-interacting monomers at $\infty \text{ \AA}$
intramolecular polarization only

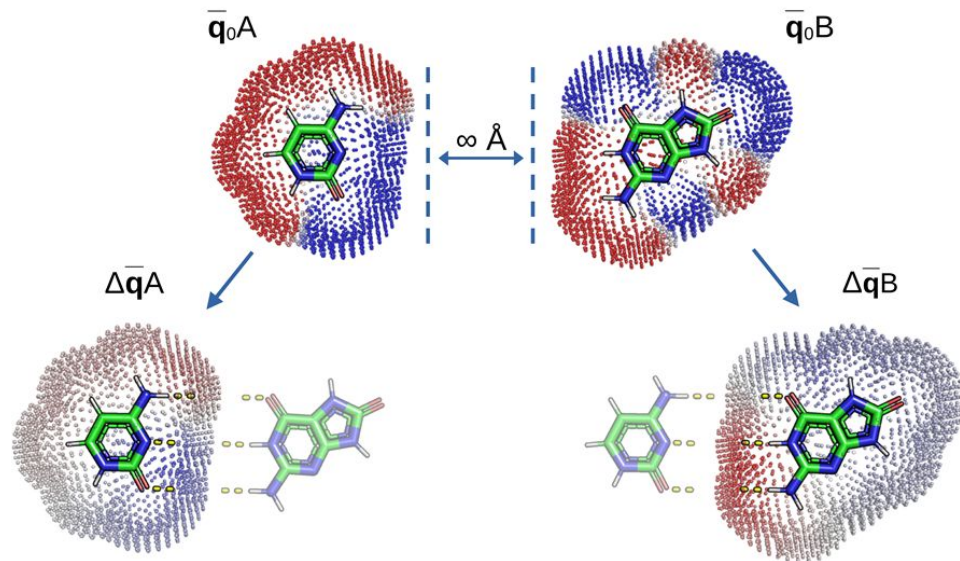
Hydrogen $\cdots$ -bonded complex
intra- + intermolecular polarization



Distribution of the MEP values



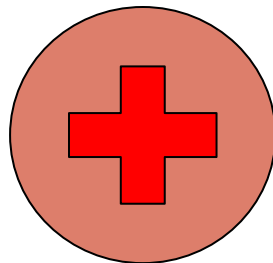
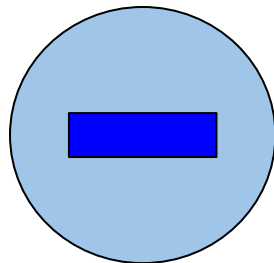
Molecular Electrostatic Potential by DENR+POL charges



Conclusions

1. We developed new empirical charge model, DENR+POL, and parameterized it using QC MEP.
2. The model consistently accounts for both inductive and polarization effects in medicinal chemistry-related molecules.
3. The DENR+POL well describes polarization in both intermolecular and intramolecular HB.
4. Theoretical consistency and low computational cost of the model => an appealing first choice method for drug-like molecules.

Thank you



for your attention!

Requirements of good empirical charge model

The important effects:

1. Ionized (formally charged) groups

2. Influence of nearest neighbors – directly bonded atoms

3. Inductive effect – remote atoms influence through σ -bonds

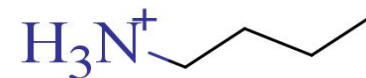
4. Polarization effect - through electric field

Molecular electrostatic potential (MEP) – unambiguously defined quantity.

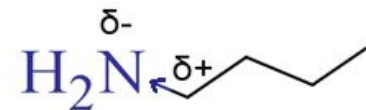
MEP reproduction by charges → **correct** description
electrostatic interactions.

Model X with wide applicability domain:

- ✓ Effects 1-4
- ✓ Correct account for polarization
- ✓ MEP reproduction



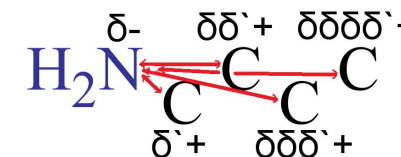
1



2



3

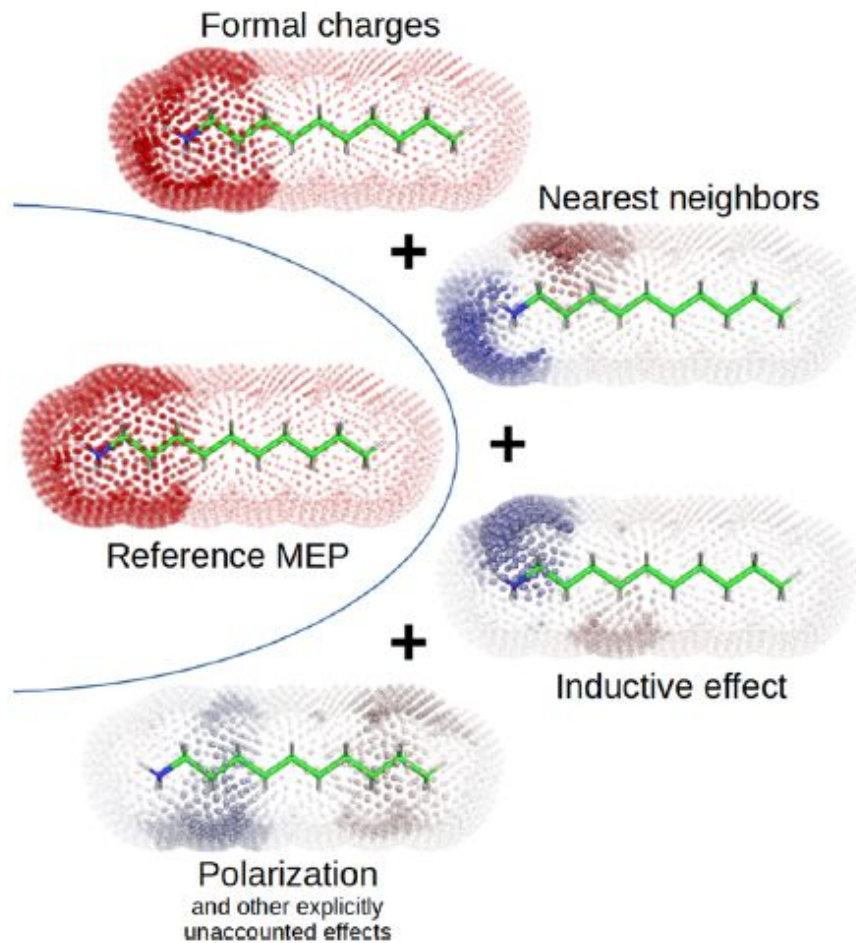


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Charge Model DENR (it exists):

- ✓ Effects 1-3 (no Pol)
- ✓ Background for Pol treatment
- ✓ MEP reproduction

Concept of consistent description



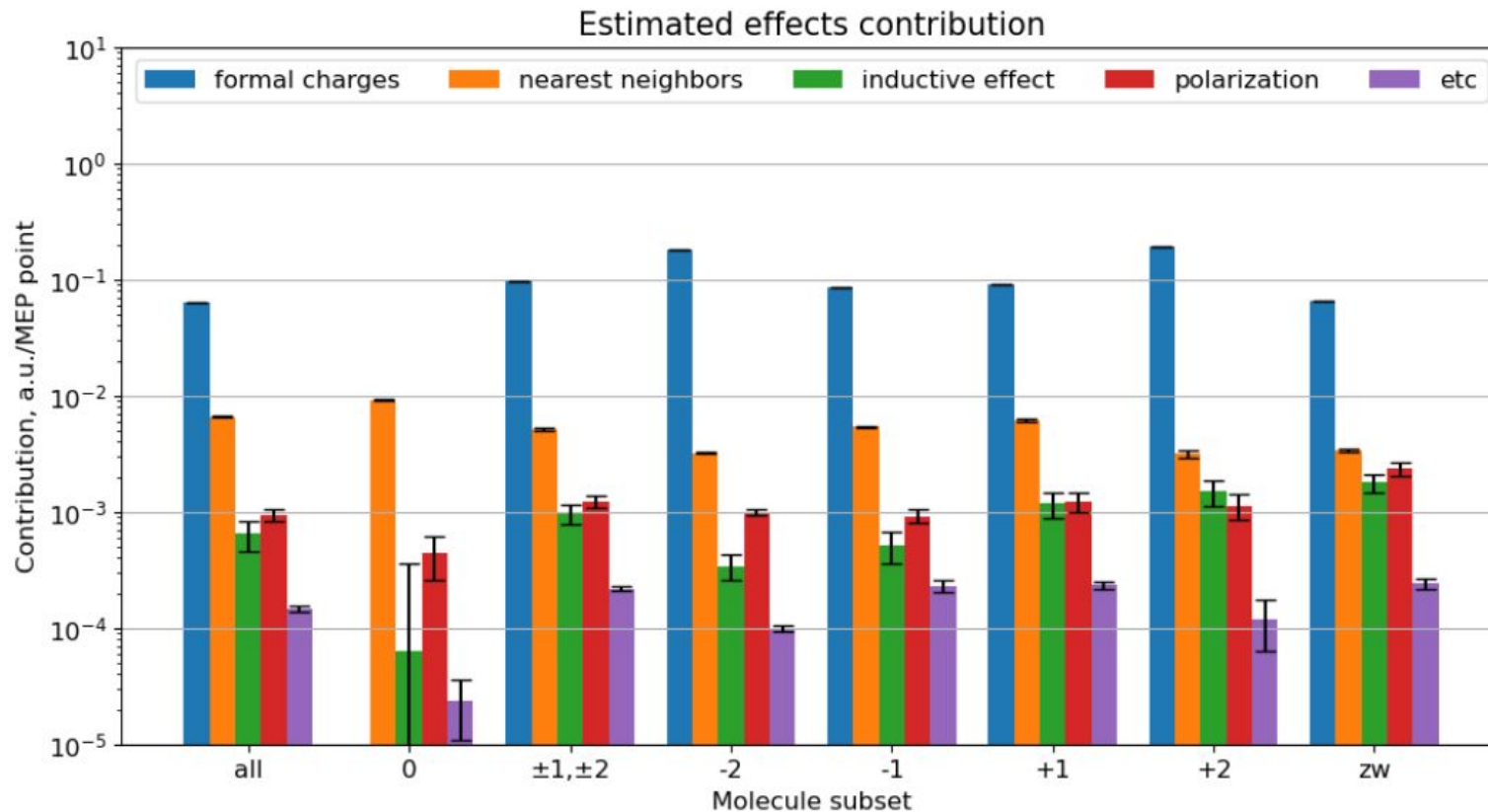
Consistent description =
hierarchy effects¹ principle:

*the description of more fundamental effect
should always precede
description of more subtle one.*

The example:
formal charges >
nearest neighbors >
inductive effect $\approx$ polarization

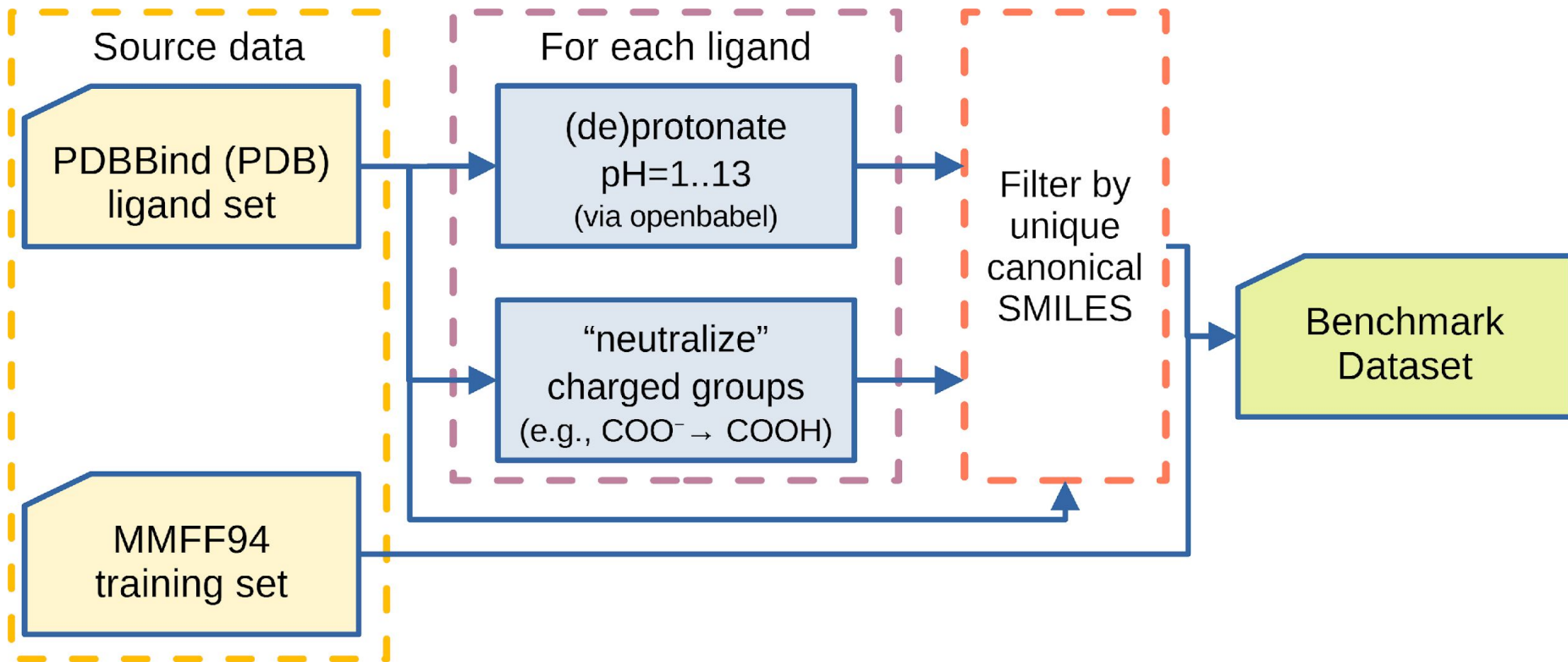
(1) Shaimardanov, A. R. et al. *J. Phys. Chem. A* **2022**, 126, 6278.

Hierarchy of the effects

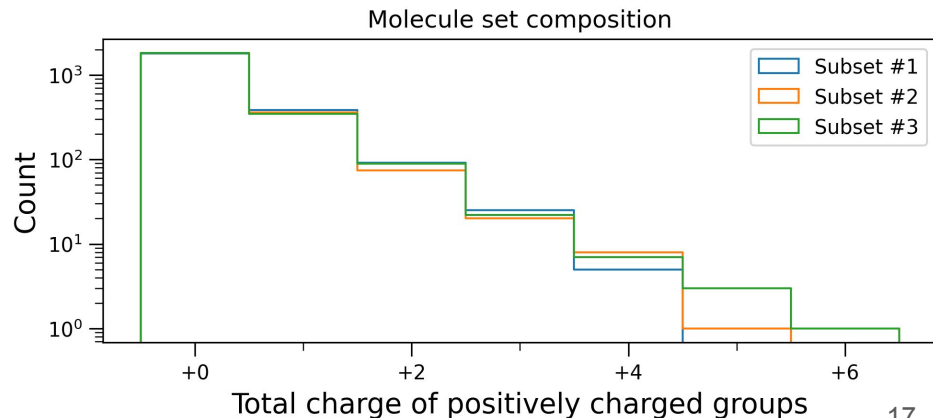
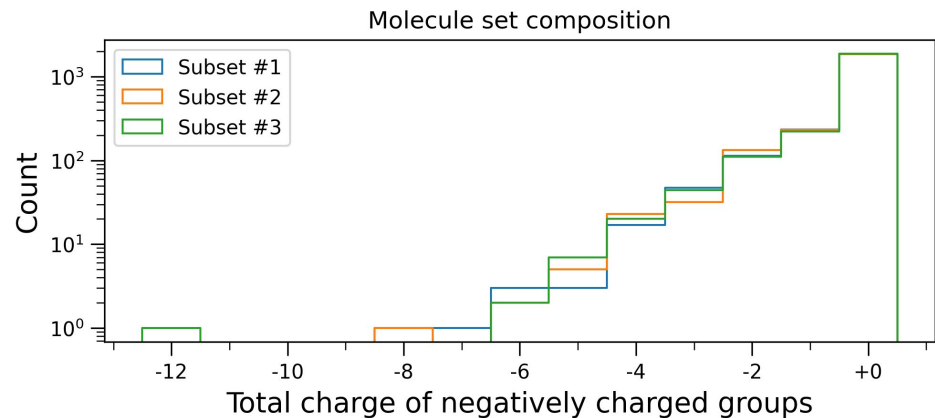
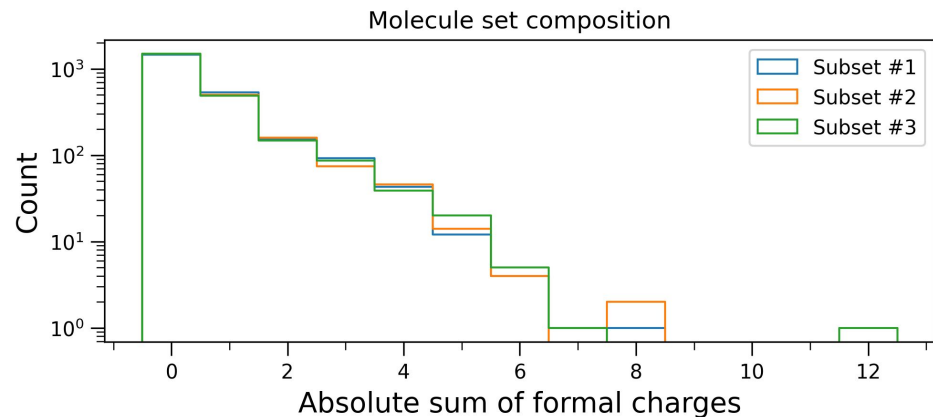
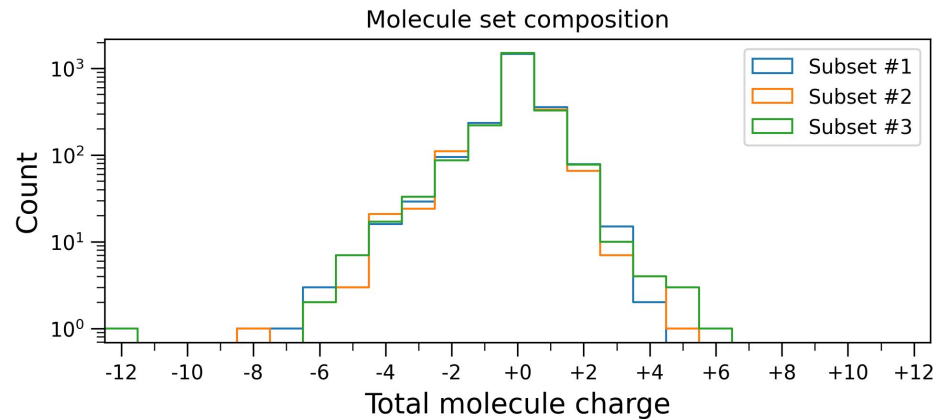


Methodology from (1) Shaimardanov, A. R. et al. *J. Phys. Chem. A* **2022**, 126, 6278.

Procedure of dataset preparation



Molecule set composition



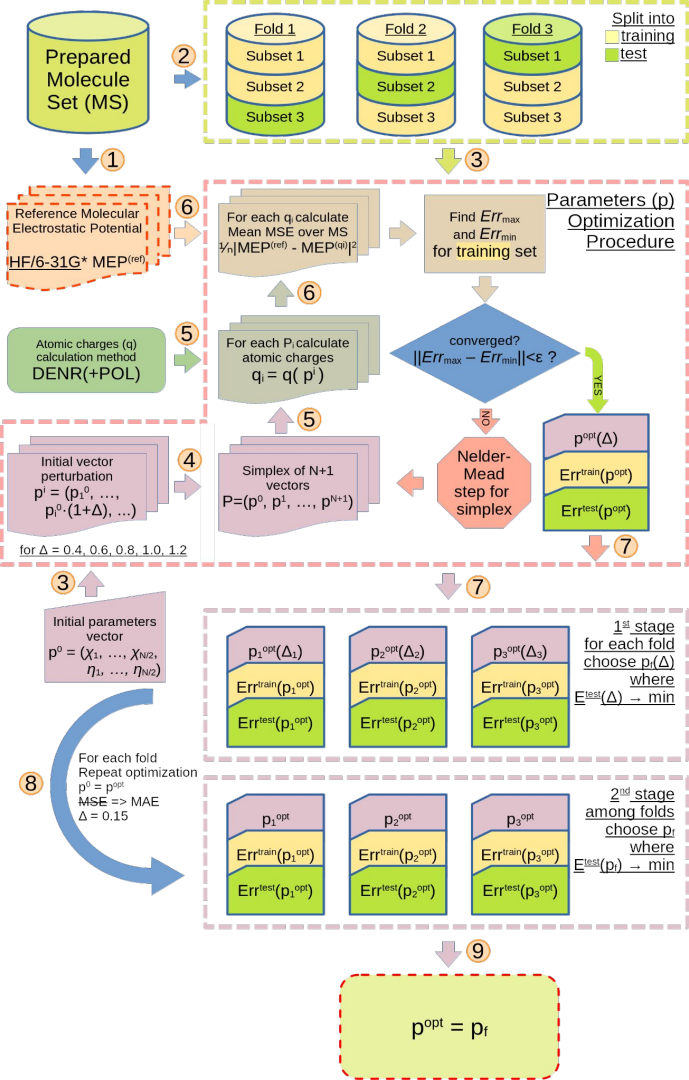
Calculation of the MEP reproduction error

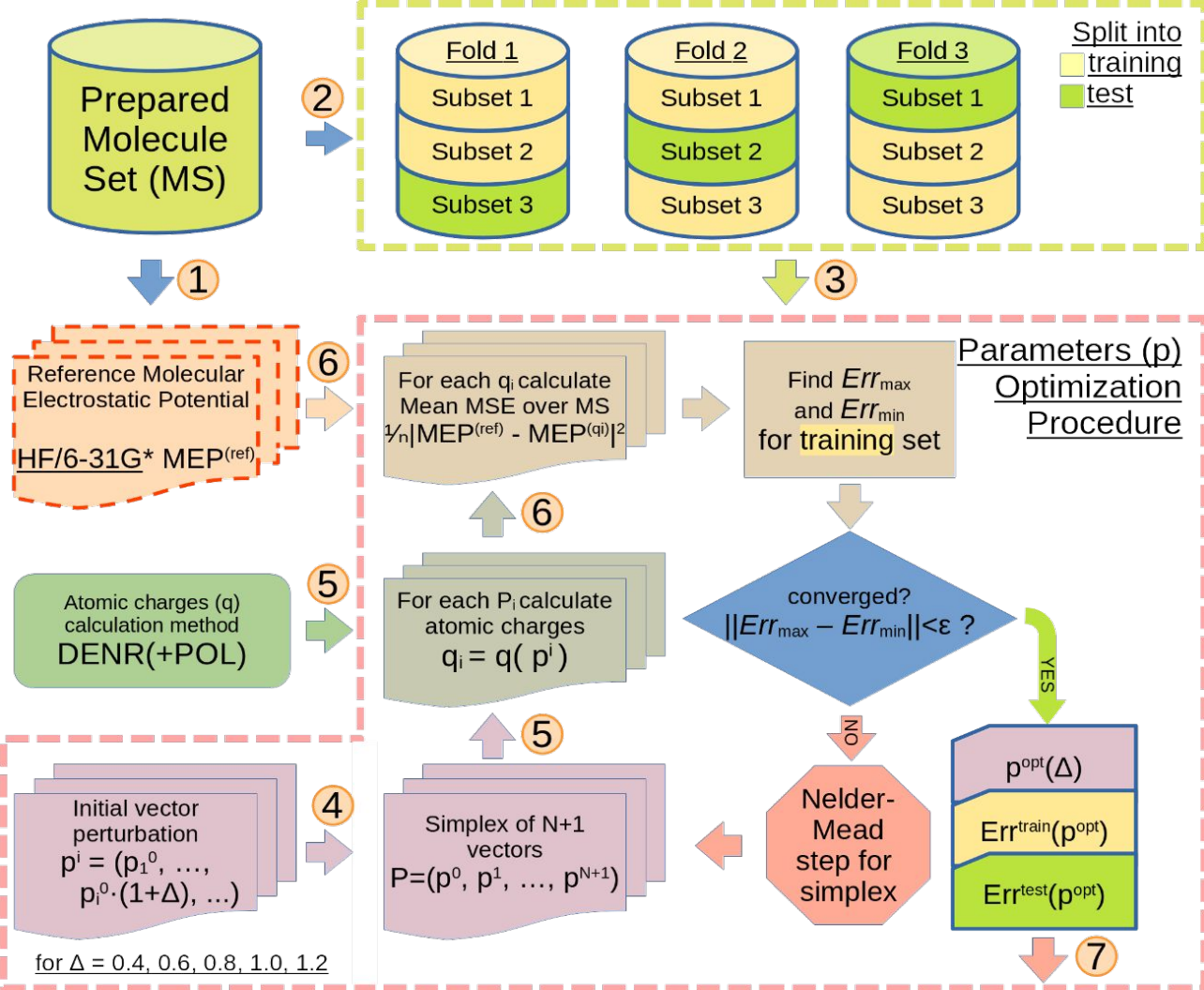
$$MMSE = \frac{1}{Nmols} \sum^{Nmols} \frac{1}{NMEPpoints} \sum_j^{NMEPpoints} \left[MEP_j - \sum_i^{Natoms} \frac{q_i}{R_{ij}} \right]^2 \quad (B1)$$

$$MMAE = \frac{1}{Nmols} \sum^{Nmols} \frac{1}{NMEPpoints} \sum_j^{NMEPpoints} \left| MEP_j - \sum_i^{Natoms} \frac{q_i}{R_{ij}} \right| \quad (B2)$$

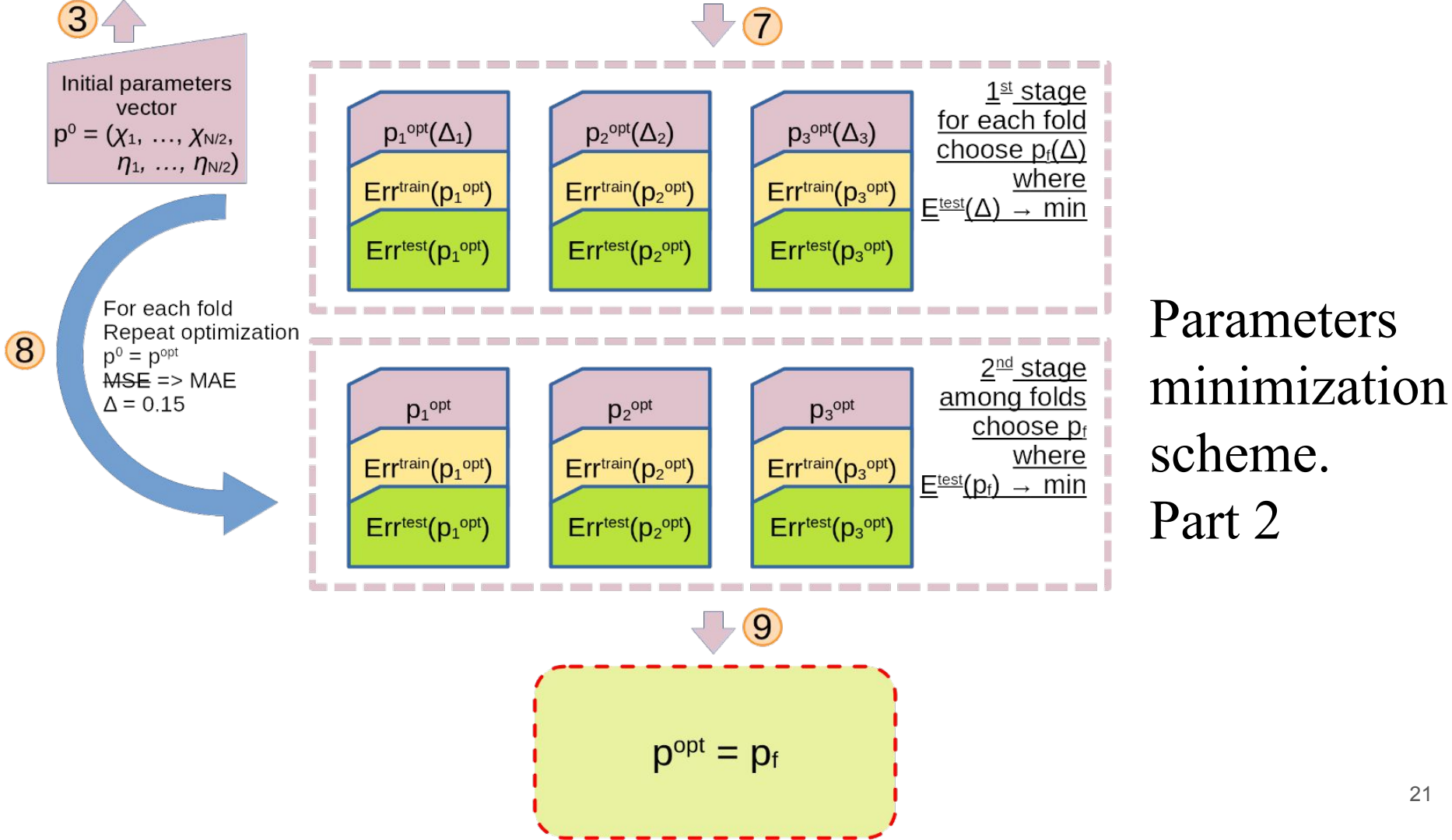
where MMSE stands for Mean (Mean Squared Error), MMAE stands for Mean (Mean Absolute Error), MEP_j is a value of a potential in j -th *MEP* point, q_i is charge value on i -th atom, R_{ij} is a distance between i -th atom and j -th *MEP* point.

Parameters minimization scheme



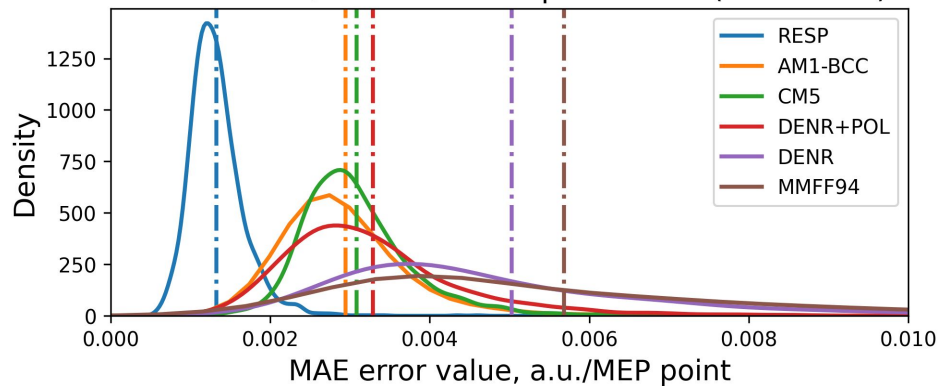


Parameters minimization scheme.
Part 1

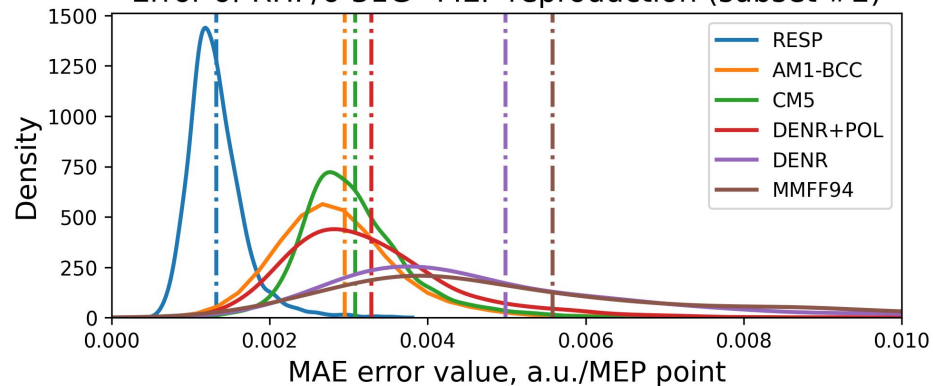


Cross-validation results

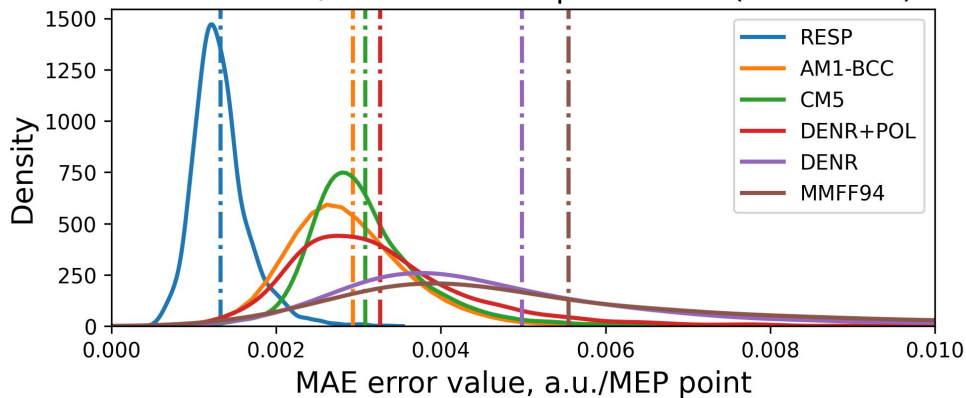
Error of RHF/6-31G* MEP reproduction (subset #1)



Error of RHF/6-31G* MEP reproduction (subset #2)



Error of RHF/6-31G* MEP reproduction (subset #3)



DENR formalism

Molecular energy as a function of atomic charges:

$$E_{mol}(q) = E_0 + \sum_i^{N_{atoms}} \left[\chi_i^0 q_i + \eta_i^0 q_i^2 \right]$$

Electronegativity: $\chi_i = \frac{\partial E_{mol}(q)}{\partial q_i}$

Hardness: $\eta_i = \frac{\partial^2 E_{mol}(q)}{\partial q_i^2} = \frac{\partial \chi_i}{\partial q_i}$

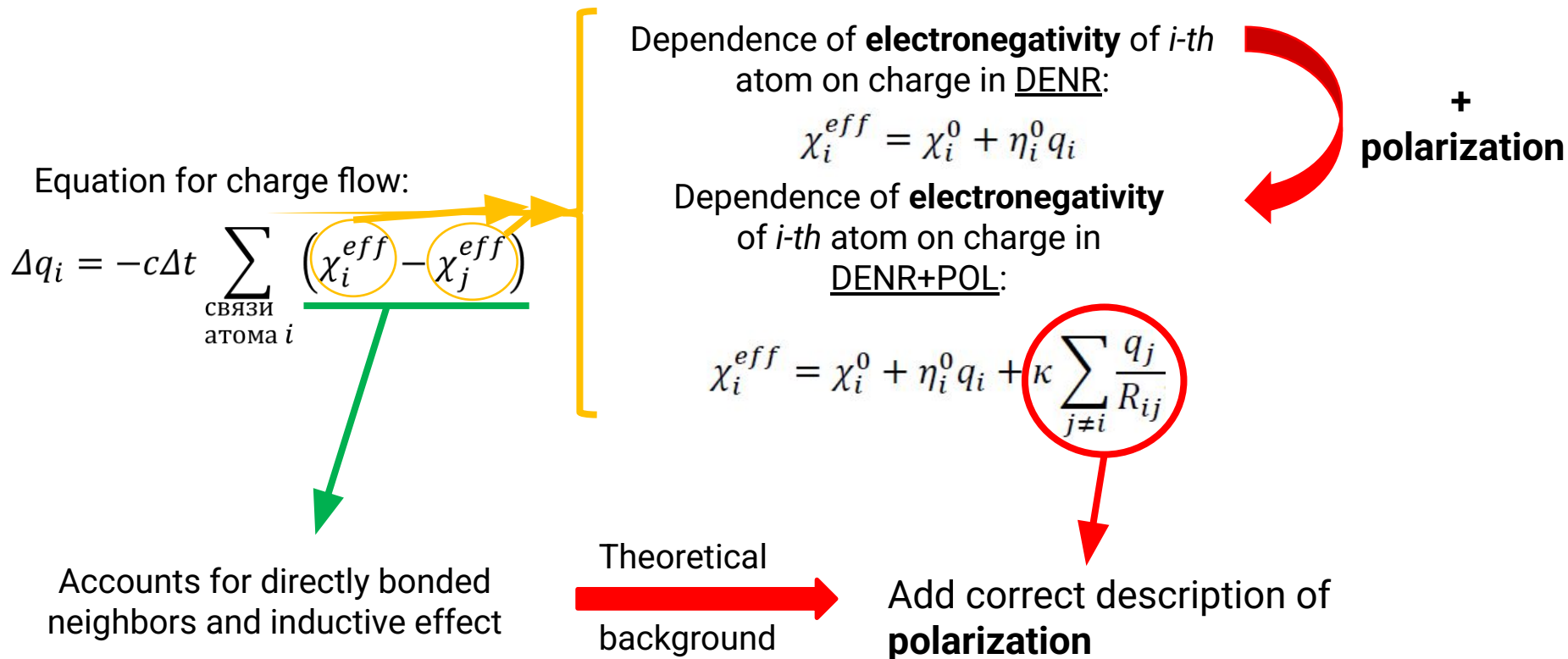
Charge flow: $\frac{dq_i}{dt} = -c \cdot \sum_j^{bonds} \left(\chi_j^{eff}(q_j^{(n)}) - \chi_i^{eff}(q_i^{(n)}) \right)$

$$\chi_i^{eff}(q_i) = \chi_i^0 + \eta_i^0 \cdot q_i$$

for $k = 1..n$
for $i = 1..N$

Charge on k-th iteration: $q_i^{(k)} = q_i^{(k-1)} + \frac{dq_i}{dt} \Delta t$

Derivation of formulas for description of polarization



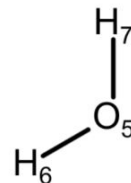
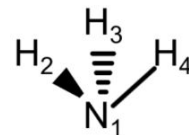
Laplace matrix formalism

$$\frac{dq_i}{dt} = -c \cdot \sum_{j \in \{i,j\}}^{\text{bonds of } i} \left(\chi_i^{eff}(\mathbf{q}) - \chi_j^{eff}(\mathbf{q}) \right)$$

$$\frac{d\mathbf{q}}{dt} = -c \cdot L\boldsymbol{\chi}^{eff}(\mathbf{q})$$

$$L_{ij} = \begin{cases} \text{deg}(i), & \text{if } i = j \\ -1, & \text{if } i \text{ is bonded to } j \\ 0, & \text{otherwise} \end{cases}$$

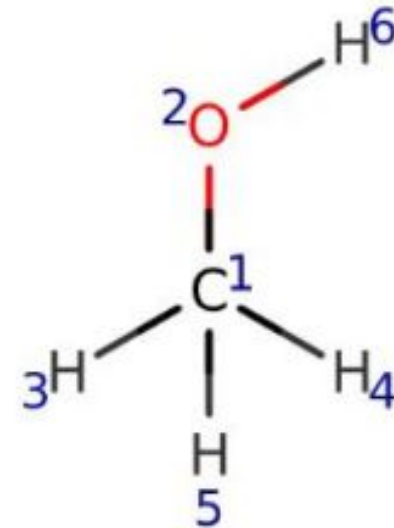
Laplace matrix



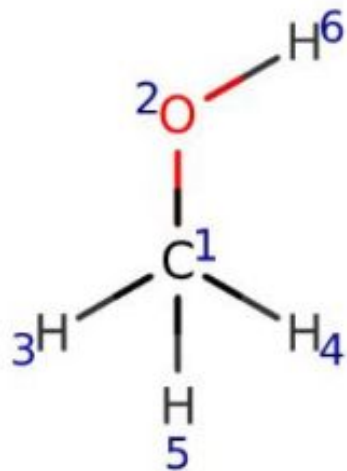
	N_1	H_2	H_3	H_4	O_5	H_6	H_7
N_1	3	-1	-1	-1	0	0	0
H_2	-1	1	0	0	0	0	0
H_3	-1	0	1	0	0	0	0
H_4	-1	0	0	1	0	0	0
O_5	0	0	0	0	2	-1	-1
H_6	0	0	0	0	-1	1	0
H_7	0	0	0	0	-1	0	1

Hardness matrix for DENR

$$\begin{array}{c}
 \text{C} \\
 \text{O} \\
 \text{H}^3 \\
 \text{H}^4 \\
 \text{H}^5 \\
 \text{H}^6
 \end{array}
 \begin{pmatrix}
 \text{C} & \text{O} & \text{H}^3 & \text{H}^4 & \text{H}^5 & \text{H}^6 \\
 \eta_{\text{C}}^0 & 0 & 0 & 0 & 0 & 0 \\
 0 & \eta_{\text{O}}^0 & 0 & 0 & 0 & 0 \\
 0 & 0 & \eta_{\text{H}^3}^0 & 0 & 0 & 0 \\
 0 & 0 & 0 & \eta_{\text{H}^3}^0 & 0 & 0 \\
 0 & 0 & 0 & 0 & \eta_{\text{H}^3}^0 & 0 \\
 0 & 0 & 0 & 0 & 0 & \eta_{\text{H}^6}^0
 \end{pmatrix}$$

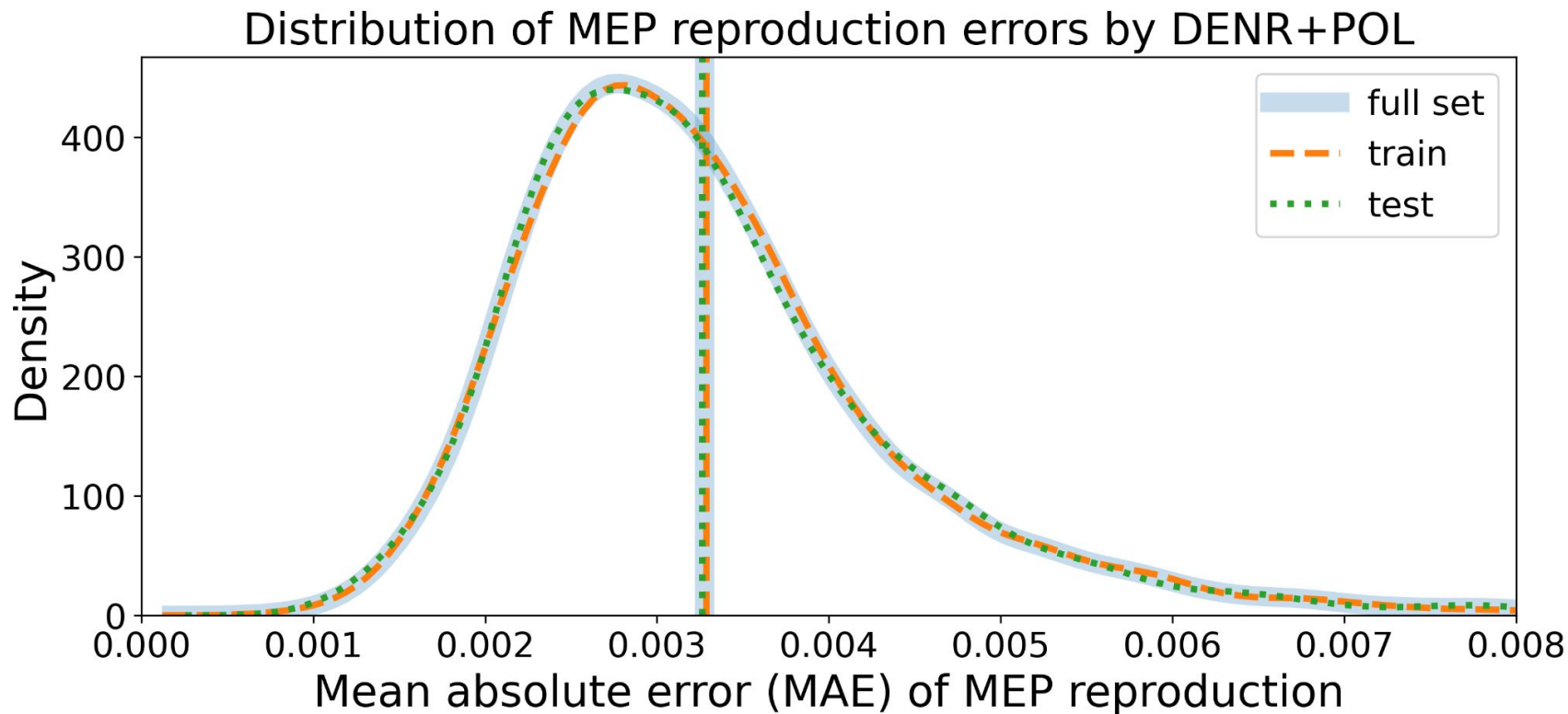


Hardness matrix for DENR+POL



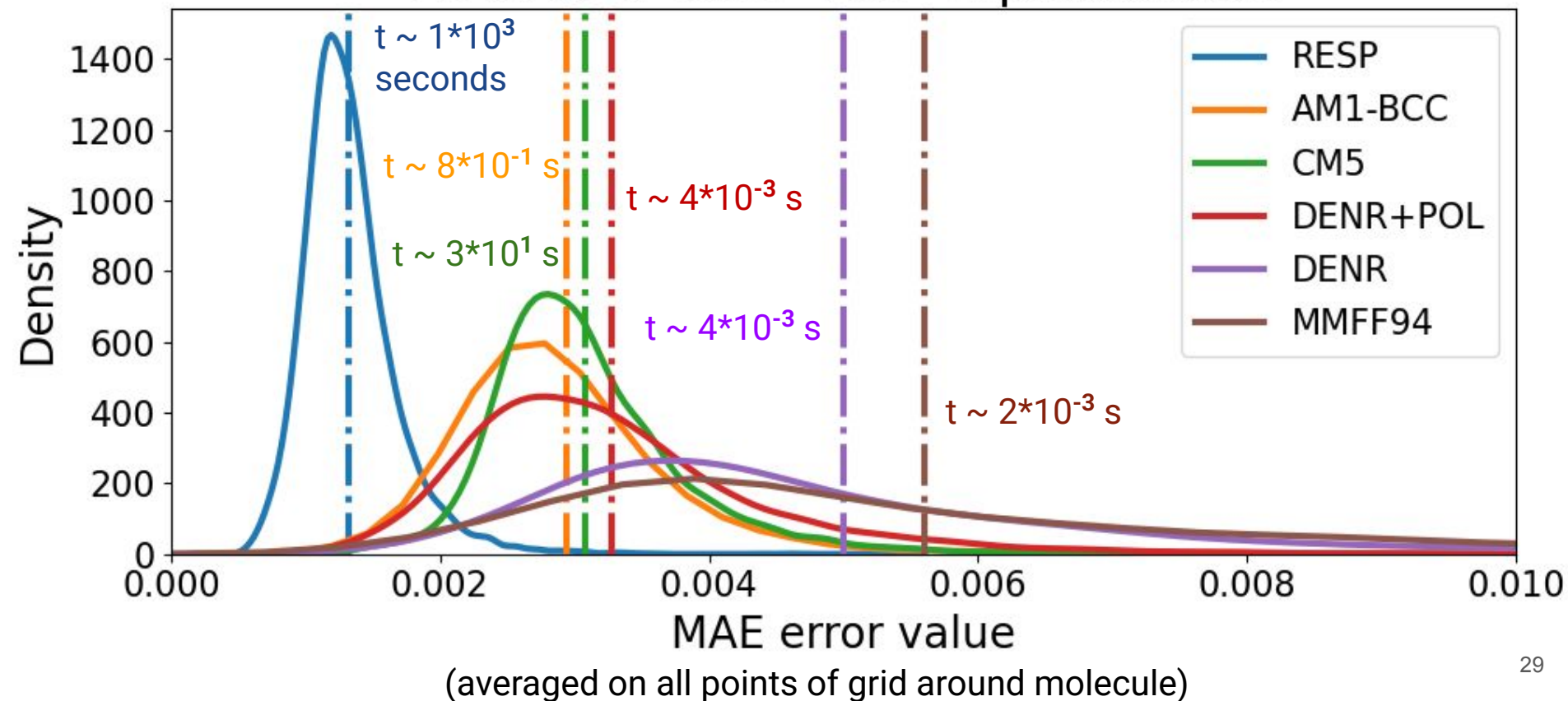
$$\begin{array}{c}
 \mathbf{C} \\
 \mathbf{O} \\
 \mathbf{H}^3 \\
 \mathbf{H}^4 \\
 \mathbf{H}^5 \\
 \mathbf{H}^6
 \end{array}
 \begin{pmatrix}
 \eta_{\mathbf{C}}^0 & r_{\mathbf{CO}}^{-1} & r_{\mathbf{CH}^3}^{-1} & r_{\mathbf{CH}^4}^{-1} & r_{\mathbf{CH}^5}^{-1} & r_{\mathbf{CH}^6}^{-1} \\
 r_{\mathbf{CO}}^{-1} & \eta_{\mathbf{O}}^0 & r_{\mathbf{OH}^3}^{-1} & r_{\mathbf{OH}^4}^{-1} & r_{\mathbf{OH}^5}^{-1} & r_{\mathbf{OH}^6}^{-1} \\
 r_{\mathbf{CH}^3}^{-1} & r_{\mathbf{OH}^3}^{-1} & \eta_{\mathbf{H}^3}^0 & r_{\mathbf{H}^3\mathbf{H}^4}^{-1} & r_{\mathbf{H}^3\mathbf{H}^5}^{-1} & r_{\mathbf{H}^3\mathbf{H}^6}^{-1} \\
 r_{\mathbf{CH}^4}^{-1} & r_{\mathbf{OH}^4}^{-1} & r_{\mathbf{H}^3\mathbf{H}^4}^{-1} & \eta_{\mathbf{H}^3}^0 & r_{\mathbf{H}^4\mathbf{H}^5}^{-1} & r_{\mathbf{H}^4\mathbf{H}^6}^{-1} \\
 r_{\mathbf{CH}^5}^{-1} & r_{\mathbf{OH}^5}^{-1} & r_{\mathbf{H}^3\mathbf{H}^5}^{-1} & r_{\mathbf{H}^4\mathbf{H}^5}^{-1} & \eta_{\mathbf{H}^3}^0 & r_{\mathbf{H}^5\mathbf{H}^6}^{-1} \\
 r_{\mathbf{CH}^6}^{-1} & r_{\mathbf{OH}^6}^{-1} & r_{\mathbf{H}^3\mathbf{H}^6}^{-1} & r_{\mathbf{H}^4\mathbf{H}^6}^{-1} & r_{\mathbf{H}^5\mathbf{H}^6}^{-1} & \eta_{\mathbf{H}^6}^0
 \end{pmatrix}$$

Distribution of the MEP errors reproduction by DENR+POL



Comparison with popular models: MEP reproduction

Error of 6-31G* MEP reproduction

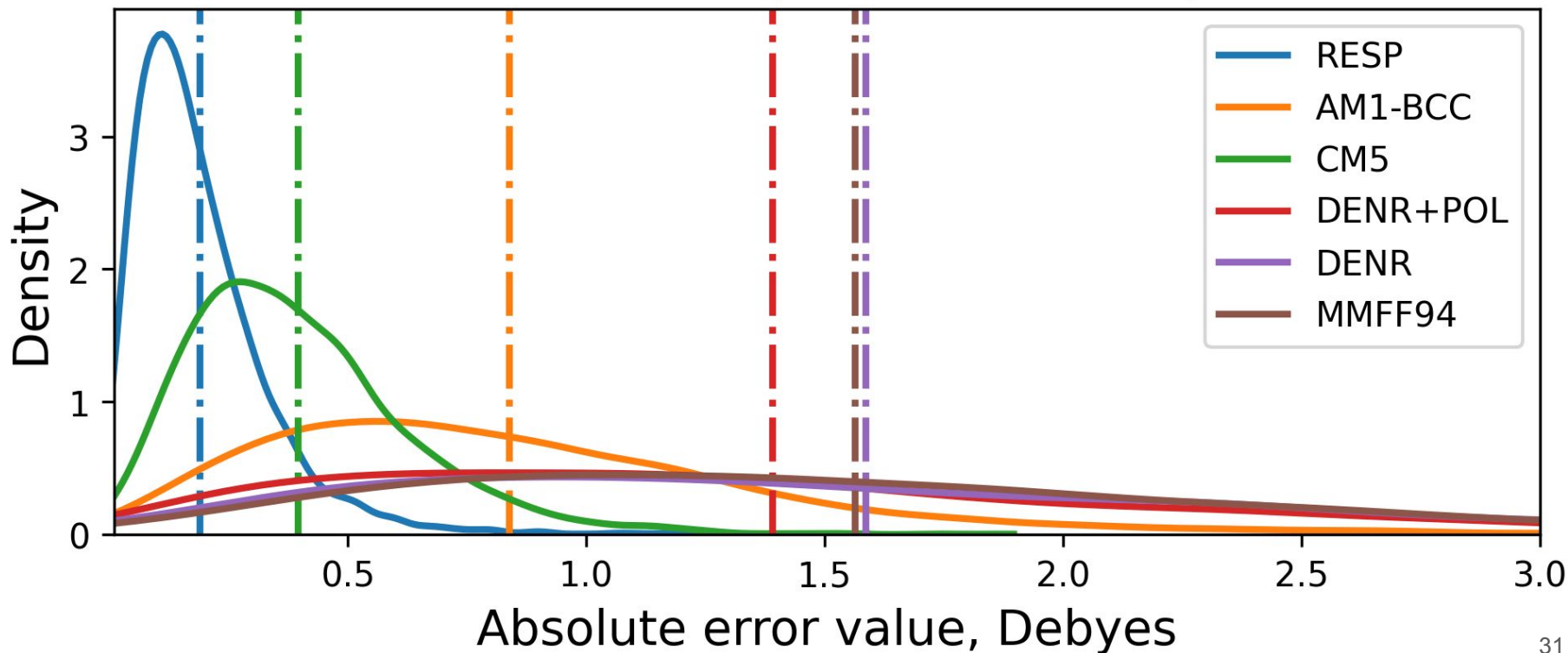


Comparison with popular models: Runtime

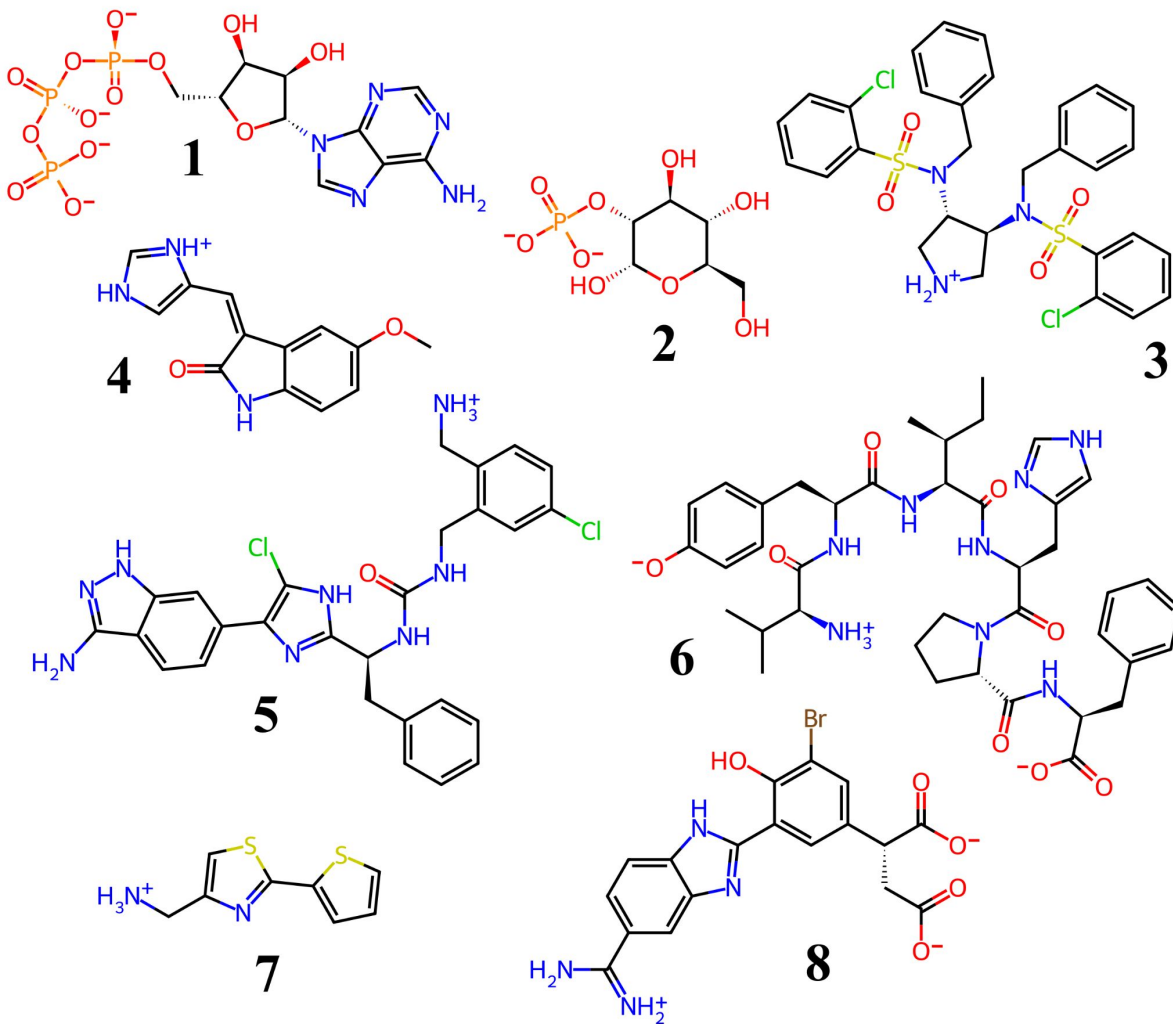
	Runtime, sec		
Method	mean	σ	mean runtime relatively to the DENR+POL
RESP	1.2×10^3	1.0×10^3	2.3×10^5
CM5	30	22	5.9×10^3
AM1-BCC	0.8	1.1	1.6×10^2
DENR+POL	5.1×10^{-3}	1.8×10^{-3}	1
DENR	4.5×10^{-3}	4.2×10^{-4}	0.88
MMFF94	1.8×10^{-3}	4.3×10^{-5}	0.35

Comparison with popular models: dipole reproduction

Error of RHF/6-31G* dipole moment reproduction

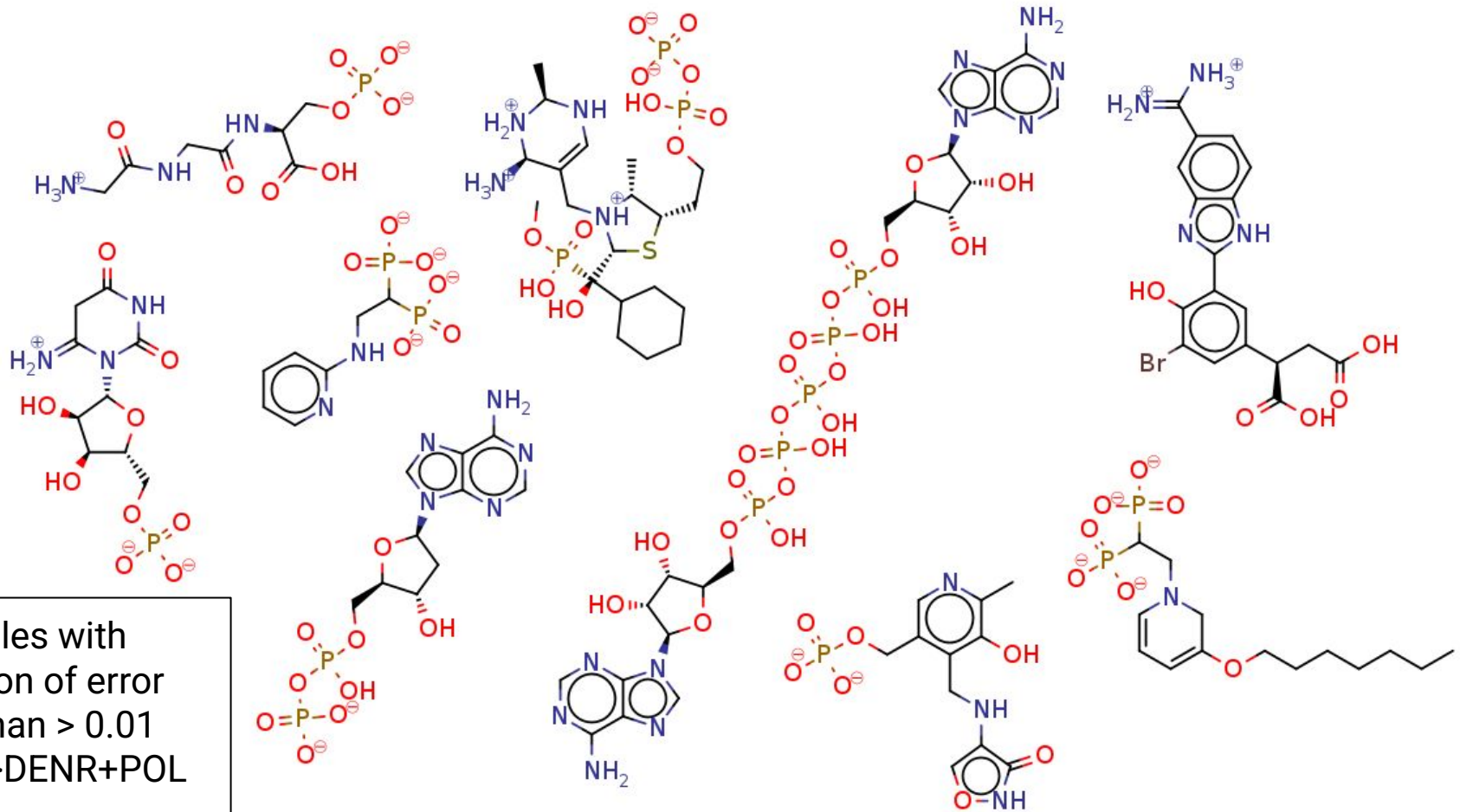


DENR+POL “wins” DENR in MEP reproduction



ID	MEP reproduction error, a.u. $\times 10^{-3}$	
	DENR +POL	DENR
1	6.7	18.9
2	3.7	11.9
3	3.6	8.6
4	3.2	8.3
5	3.2	8.0
6	6.3	15.1
7	4.9	9.9
8	8.4	18.9

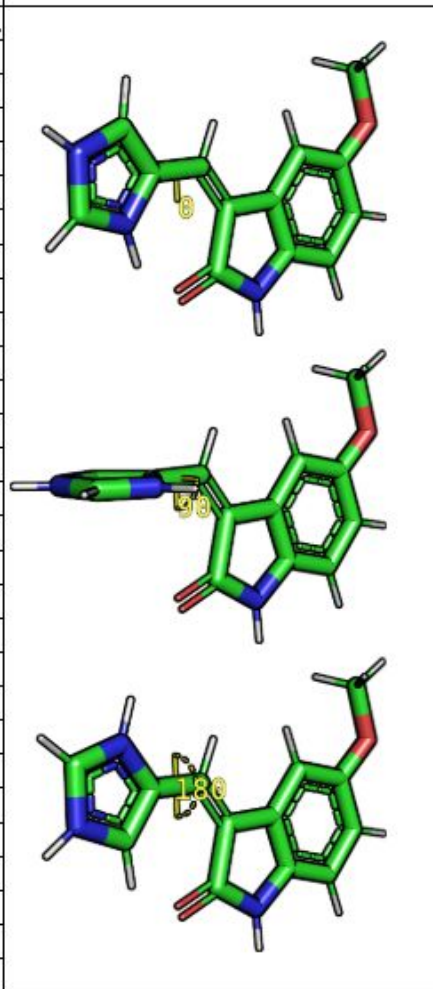
Biomolecules: perspectives



Molecules with
reduction of error
more than > 0.01
DENR->DENR+POL

Table G1. Dependence of the MEP reproduction error on the molecule conformation.

deg	MAE, a.u. $\times 10^{-3}$	
	DENR	DENR+POL
0	9.0	3.1
15	8.8	3.1
30	8.5	2.9
45	8.3	2.8
60	8.3	2.9
75	8.3	2.9
90	8.3	2.9
105	8.3	2.9
120	8.2	2.9
135	8.1	2.9
150	8.1	2.9
165	8.3	2.8
180	8.4	2.8
195	8.3	2.8
210	8.1	2.9
225	8.1	2.9
240	8.2	2.9
255	8.3	2.9
270	8.3	2.9
285	8.3	2.9
300	8.3	2.8
315	8.3	2.8
330	8.5	2.9
345	8.8	3.1
360	9.0	3.1
mean	8.4	2.9
avedev	0.2	0.1
avedev/mean, %	2.31%	2.17%



Dependence of the MEP reproduction error on the molecular conformation.