

FORCE FIELDS

MD: interatomic interactions

$$\begin{aligned}\hat{H} &= \hat{T}_{\text{nuc}} + \hat{T}_{\text{ele}} + V(\mathbf{R}) + V(|\mathbf{R} - \mathbf{r}|) + V(\mathbf{r}) \\ &= \hat{T}_{\text{nuc}} + E_{\text{pot}}(\mathbf{R})\end{aligned}$$

Obtaining $E_{\text{pot}}(\mathbf{R})$ via *ab initio*: AIMD

$$\left[\hat{T}_{\text{ele}} + V(|\mathbf{R} - \mathbf{r}|) + V(\mathbf{r}) \right] \psi(\mathbf{r}; \mathbf{R}) = E_{\text{pot}}(\mathbf{R}) \psi(\mathbf{r}; \mathbf{R})$$

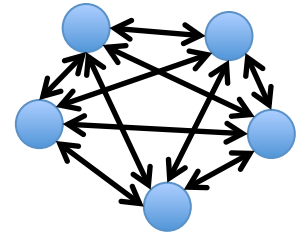
$$\mathbf{F} = M\mathbf{a} = - \left(\frac{\partial E_{\text{pot}}(\mathbf{R})}{\partial \mathbf{R}} \right)$$

Using model to simulate $E_{\text{pot}}(\mathbf{R})$: force field

MD: H_{ele}

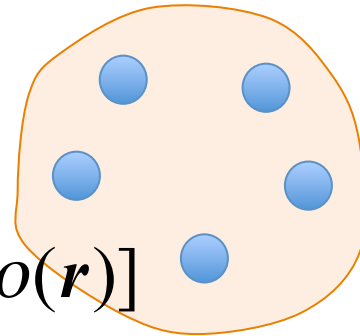
Hartree-Fock
and
post H-F

$$\begin{aligned}\hat{H}_{\text{ele}} &= \hat{T}_{\text{ele}} + V(|\mathbf{R} - \mathbf{r}|) + V(\mathbf{r}) \\ &= -\sum_{i=1}^{N_e} \frac{1}{2} \nabla_i^2 - \sum_{i=1}^{N_e} \sum_{I=1}^{N_n} \frac{Z_I}{|\mathbf{R}_I - \mathbf{r}_i|} + \sum_{i=1}^{N_e} \sum_{j=i+1}^{N_e} \frac{1}{|\mathbf{r}_i - \mathbf{r}_j|}\end{aligned}$$



Density Functional Theory

$$\begin{aligned}H_{\text{ele}}^{\text{DFT}}[\rho(\mathbf{r})] &= T_{\text{ele}}[\rho(\mathbf{r})] + V_{\text{ext}}[\rho(\mathbf{r})] + V_{\text{H}}[\rho(\mathbf{r})] + V_{\text{XC}}[\rho(\mathbf{r})] \\ &= T_{\text{ele}}[\rho(\mathbf{r})] - \sum_{I=1}^{N_n} \int \frac{Z_I \rho(\mathbf{r})}{|\mathbf{R}_I - \mathbf{r}|} d^3r + \int \int \frac{\rho(\mathbf{r}) \rho(\mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|} d^3r d^3r' + V_{\text{XC}}[\rho(\mathbf{r})]\end{aligned}$$



MD: force calculations

$$\begin{aligned} \mathbf{F}_I^{\text{HFT}} &= -\nabla_I \langle \Psi_0 | \hat{H}_e | \Psi_0 \rangle \\ &= -\langle \nabla_I \Psi_0 | \hat{H}_e | \Psi_0 \rangle - \langle \Psi_0 | \nabla_I \hat{H}_e | \Psi_0 \rangle - \langle \Psi_0 | \hat{H}_e | \nabla_I \Psi_0 \rangle \\ &= -\langle \Psi_0 | \nabla_I \hat{H}_e | \Psi_0 \rangle - E_0 \langle \nabla_I \Psi_0 | \Psi_0 \rangle - E_0 \langle \Psi_0 | \nabla_I \Psi_0 \rangle \end{aligned}$$

$$\nabla^\dagger = -\nabla$$

$$= -\langle \Psi_0 | \nabla_I \hat{H}_e | \Psi_0 \rangle$$

Hellman-Feynman theorem!
The wave functions here must be the solution of stationary Schrödinger equation.

Problem: must be complete basis, containing *infinite* number of basis functions. Ideal only.

MD: force calculations

In practice: using Slater determinant, and LCAO.

Use basis sets:

$$\Psi_0 = \frac{1}{\sqrt{N!}} \det \{ \phi_i \}$$

$$\phi_i = \sum_v c_{iv} f_v(\mathbf{r}; \mathbf{R})$$

$$\nabla_I \phi_i = \sum_v \left(\nabla_I c_{iv} \right) f_v(\mathbf{r}; \mathbf{R}) + \sum_v c_{iv} \left(\nabla_I f_v(\mathbf{r}; \mathbf{R}) \right)$$

So in practices, there will be additional terms appear.

MD: force calculations

$$\mathbf{F}_I^{\text{IBS}} = \sum_{i\nu\mu} \left(\left\langle \nabla_I f_\nu \left| H_e^{\text{NSC}} - \varepsilon_i \right| f_\mu \right\rangle + \left\langle f_\nu \left| H_e^{\text{NSC}} - \varepsilon_i \right| \nabla_I f_\mu \right\rangle \right)$$

Another name:

Pulay force

IBS: incomplete-basis-set correction

NSC: non-self-consistency correction

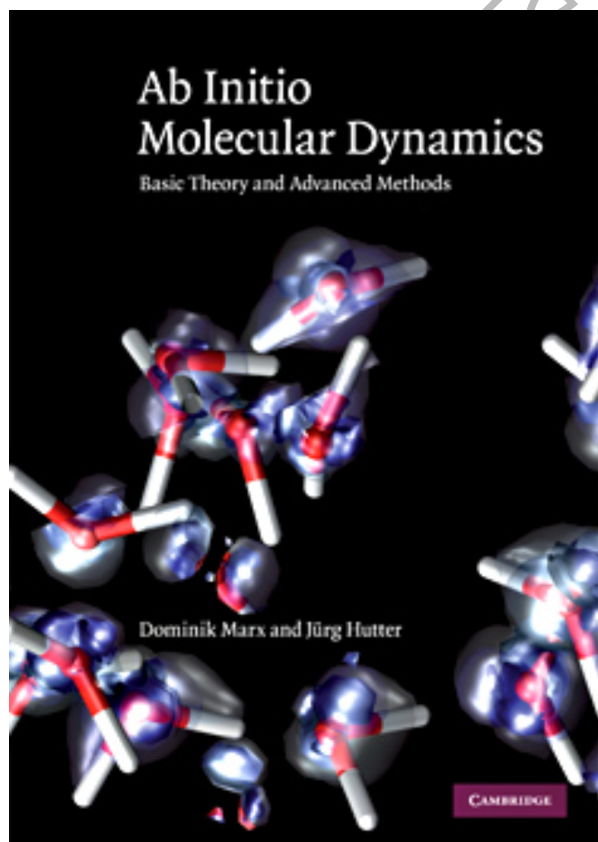
There is no Pulay force in plane wave calculations, but once one uses finite basis, it appears.

$$\mathbf{F}_I^{\text{NSC}} = - \int d^3r (\nabla_I n(\mathbf{r})) (V^{\text{SCF}} - V^{\text{NSC}})$$

$$\mathbf{F}_I = \mathbf{F}_I^{\text{HFT}} + \mathbf{F}_I^{\text{IBS}} + \mathbf{F}_I^{\text{NSC}}$$

MD: AIMD

本质上是电子结构计算的问题!



AB INITIO MOLECULAR DYNAMICS:
BASIC THEORY AND ADVANCED
METHODS

DOMINIK MARX
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University of Zürich



CAMBRIDGE
UNIVERSITY PRESS

MD: cMD

c: conventional, classical

Use a physical/mathematical model for $E_{\text{pot}}(\mathbf{R})$

$$E_{\text{pot}} \approx E^{\text{EF}}(\mathbf{R}) = \sum_{I=1}^{N_n} v_1(\mathbf{R}_I) + \sum_{I<J}^{N_n} v_2(\mathbf{R}_I, \mathbf{R}_J) \\ + \sum_{I<J<K}^{N_n} v_3(\mathbf{R}_I, \mathbf{R}_J, \mathbf{R}_K) + \dots$$

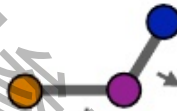
MD: cMD

c: conventional, classical

Use a physical/mathematical model for $E_{\text{pot}}(R)$

Typical energy function

$$E = \sum_{b \in \text{bonds}} k_b (r_b - r_b^0)^2 + \sum_{a \in \text{angles}} \kappa_a (\theta_a - \theta_a^0)^2 + \sum_{d \in \text{dihedrals}} \sum_n l_{dn} \cos(n\pi)$$


bond stretch
angle torsion
dihedrals

$$+ \sum_{i < j \in \text{atoms}} \frac{q_i q_j}{r_{ij}} + \sum_{i < j \in \text{atoms}} 4\epsilon_{ij} \left(\left(\frac{\sigma_{ij}}{r_{ij}} \right)^{12} - \left(\frac{\sigma_{ij}}{r_{ij}} \right)^6 \right)$$

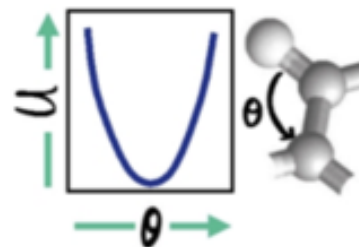
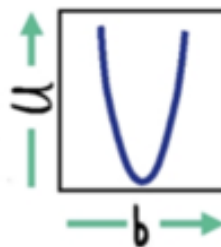

electrostatics
dispersion

Usually fixed charges

$$U = \sum_{\text{All Bonds}} \frac{1}{2} K_b (b - b_0)^2 + \sum_{\text{All Angles}} \frac{1}{2} K_\theta (\theta - \theta_0)^2$$

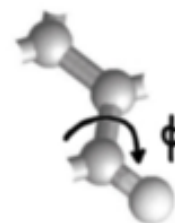
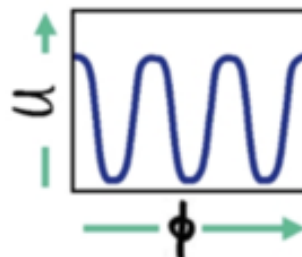
All Bonds

All Angles



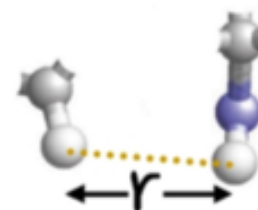
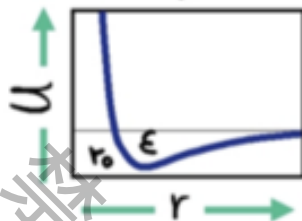
$$+ \sum K_\phi [1 - \cos(n\phi + \delta)]$$

All Torsion Angles



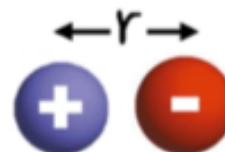
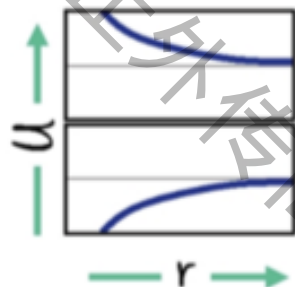
$$+ \sum \epsilon \left[\left(\frac{r_0}{r} \right)^{12} - 2 \left(\frac{r_0}{r} \right)^6 \right]$$

All nonbonded pairs



$$+ \sum 332 q_i q_j / r$$

All partial charges

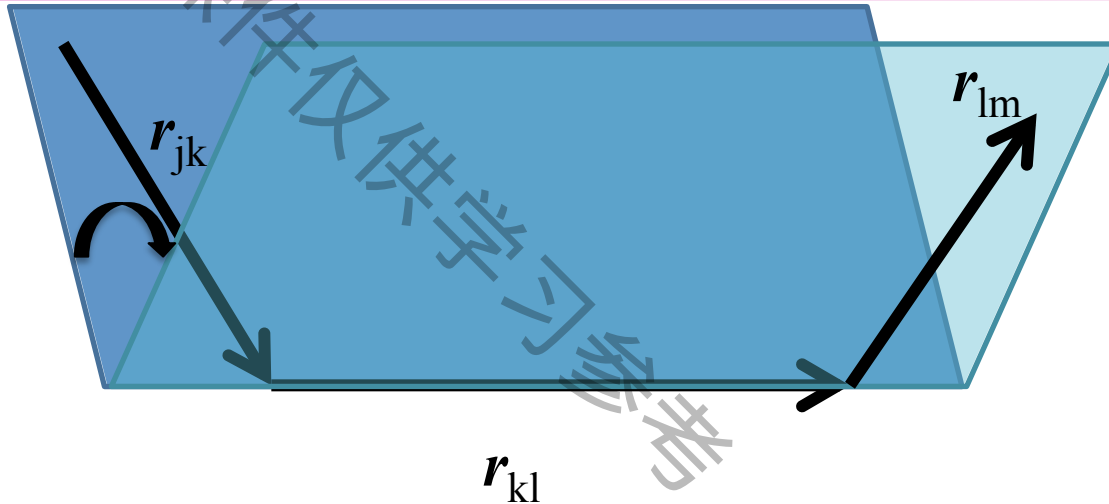


```

(base) leap:yongleli$ ls
cmd/      lib/      parm/     pixmaps/  prep/
(base) leap:yongleli$ grep "Mn" * -r
cmd/leaprc.water.spce: { "Mn2+" "Mn" "sp3" }
cmd/leaprc.DNA.OL15: { "Mn2+" "Mn" "sp3" }
cmd/leaprc.water.opc: { "Mn2+" "Mn" "sp3" }
cmd/leaprc.water.tip3p: { "Mn2+" "Mn" "sp3" }
cmd/leaprc.water.tip4pew: { "Mn2+" "Mn" "sp3" }
cmd/leaprc.water.fb3: { "Mn2+" "Mn" "sp3" }
cmd/leaprc.water.fb4: { "Mn2+" "Mn" "sp3" }
lib/atomic_ions.log:> i = createAtom MN Mn2+ 2.0
lib/atomic_ions.log:> set i element "Mn"
lib/atomic_ions.lib: "MN" "Mn2+" 0 1 131072 1 25 2.000000
lib/atomic_ions.lib: "MN" "Mn2+" 0 -1 0.0
lib/atomic_ions.cmd:i = createAtom MN Mn2+ 2.0
lib/atomic_ions.cmd:set i element "Mn"
parm/frcmod.ions234lm_hfe_spce:Mn2+ 54.94
parm/frcmod.ions234lm_hfe_spce:Mn2+ 1.338 0.00789684 HFE set for SPC\E water from Li et al., JCTC, 2013, 9, 2733
parm/frcmod.ions234lm_iod_spce:Mn2+ 54.94
parm/frcmod.ions234lm_iod_spce:Mn2+ 1.467 0.02960343 IOD set from Li et al., JCTC, 2013, 9, 2733
parm/frcmod.ions234lm_iod_tip4pew:Mn2+ 54.94
parm/frcmod.ions234lm_iod_tip4pew:Mn2+ 1.467 0.02960343 IOD set from Li et al., JCTC, 2013, 9, 2733
parm/frcmod.ions234lm_126_tip3p:Mn2+ 54.94
parm/frcmod.ions234lm_126_tip3p:Mn2+ 1.407 0.01686710 CM set for TIP3P water from Li et al., JCTC, 2013, 9, 2733
parm/frcmod.ions234lm_126_tip4pew:Mn2+ 54.94
parm/frcmod.ions234lm_126_tip4pew:Mn2+ 1.401 0.01586934 CM set for TIP4P\EW water from Li et al., JCTC, 2013, 9, 2733
parm/frcmod.ions234lm_1264_tip3p:Mn2+ 54.94
parm/frcmod.ions234lm_1264_tip3p:Mn2+ 1.485 0.03450196 12-6-4 set for TIP3P water from Li and Merz, JCTC, 2014, 10, 289
parm/frcmod.ions234lm_iod_tip3p:Mn2+ 54.94
parm/frcmod.ions234lm_iod_tip3p:Mn2+ 1.467 0.02960343 IOD set from Li et al., JCTC, 2013, 9, 2733
parm/frcmod.ions234lm_1264_spce:Mn2+ 54.94
parm/frcmod.ions234lm_1264_spce:Mn2+ 1.495 0.03745682 12-6-4 set for SPC\E water from Li and Merz, JCTC, 2014, 10, 289
parm/lj_1264_pol.dat:Mn2+ 0.534 Caculated from B3LYP/6-311++G(2d,2p) level of theory
parm/frcmod.ions234lm_126_spce:Mn2+ 54.94
parm/frcmod.ions234lm_126_spce:Mn2+ 1.406 0.01669760 CM set for SPC\E water from Li et al., JCTC, 2013, 9, 2733
parm/frcmod.ions234lm_1264_tip4pew:Mn2+ 54.94
parm/frcmod.ions234lm_1264_tip4pew:Mn2+ 1.485 0.03450196 12-6-4 set for TIP4P\EW water from Li and Merz, JCTC, 2014, 10, 289
parm/frcmod.ions234lm_hfe_tip4pew:Mn2+ 54.94
parm/frcmod.ions234lm_hfe_tip4pew:Mn2+ 1.276 0.00354287 HFE set for TIP4P\EW water from Li et al., JCTC, 2013, 9, 2733
parm/frcmod.ions234lm_hfe_tip3p:Mn2+ 54.94
parm/frcmod.ions234lm_hfe_tip3p:Mn2+ 1.339 0.00799176 HFE set for TIP3P water from Li et al., JCTC, 2013, 9, 2733
(base) leap:yongleli$

```

How to calculate dihedral angle



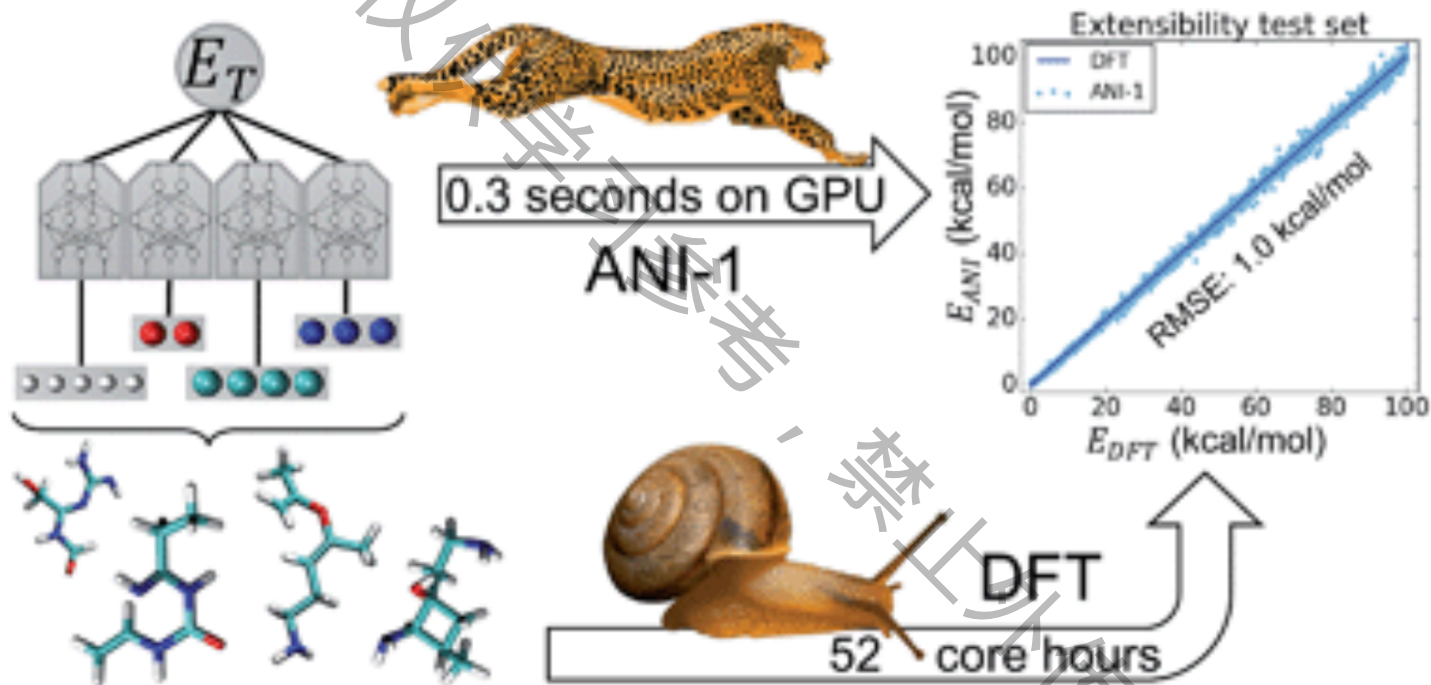
$$\phi_{jklm} = \arccos \{ u_{jklm} \}$$

$$u_{jklm} = \vec{e}_{jkl} \cdot \vec{e}_{klm}$$

$$\vec{e}_{jkl} = \frac{\vec{r}_{jk} \times \vec{r}_{kl}}{|\vec{r}_{jk} \times \vec{r}_{kl}|}, \quad \vec{e}_{klm} = \frac{\vec{r}_{kl} \times \vec{r}_{lm}}{|\vec{r}_{kl} \times \vec{r}_{lm}|}$$

simplemd/run

机器学习力场



J. S. Smith, O. Isayev and A. E. Roitberg
Chem. Sci., 2017, 8, 3192



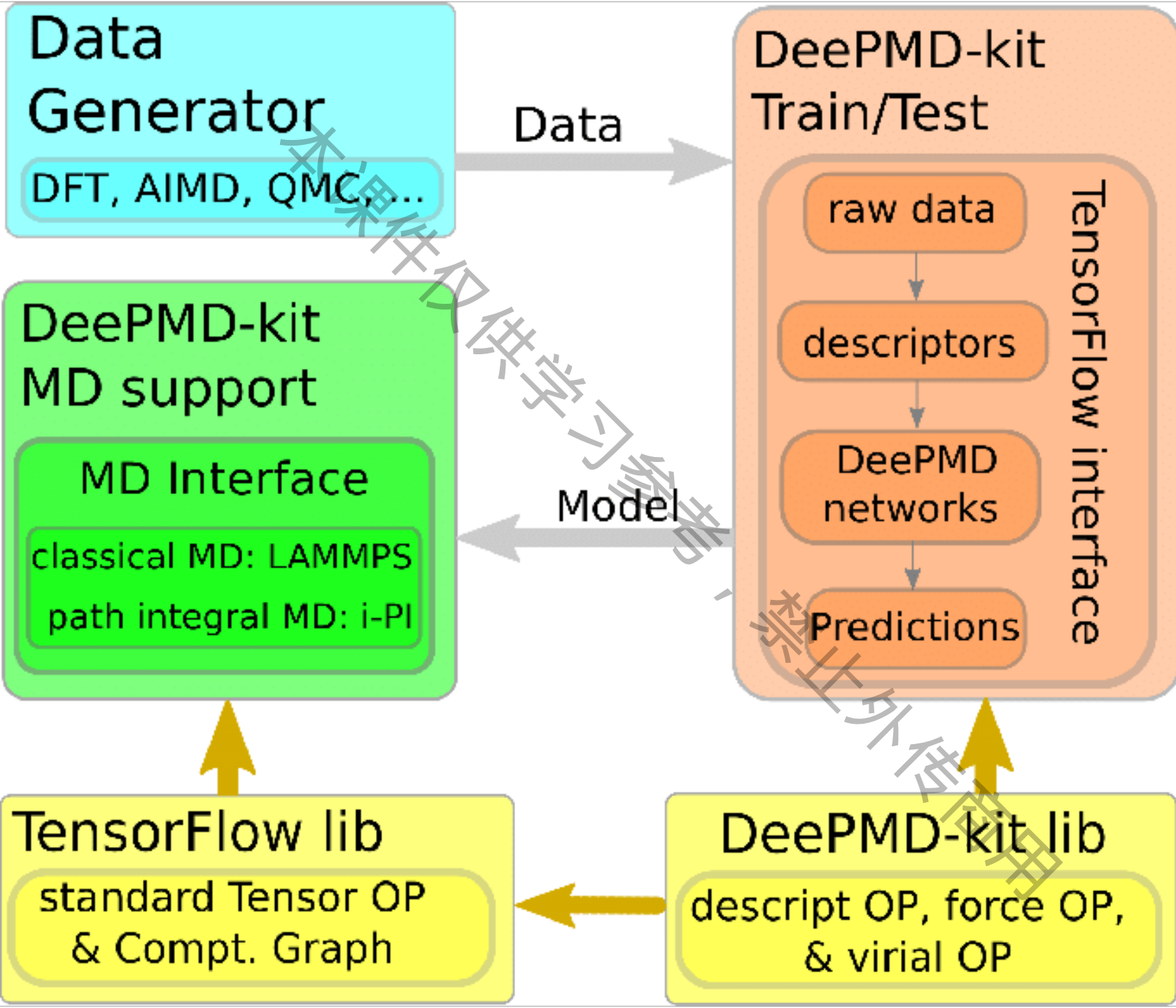
Computer Physics Communications

Volume 228, July 2018, Pages 178-184



DeePMD-kit: A deep learning package for many-body potential energy representation and molecular dynamics ☆

Han Wang ^{a, b} ✉, Linfeng Zhang ^c ✉, Jiequn Han ^c ✉, Weinan E ^{c, d, e} ✉



Neural Network Potential Software



ANI/Torchani

- Improved from BP descriptor
- Training using PyTorch, GPU accelerations
- Python interface. Better usability.



RuNNer/N2P2

- Original / Improved BP descriptor
- Training using RuNNer by Behler, no GPU accelerations
- Lammmps interface

DeePMD-kit

- Other descriptor
- Training using TensorFlow, GPU accelerations
- Lammmps interface

Ref 1: Smith J. S., et. Al. Chemical Science, 2017, 8(4): 3192-3203.

Ref 2: Singraber, et al. Journal of chemical theory and computation, 2019.

Ref 3: Wang H., et al. Computer Physics Communications, 2018, 228: 178-184.

Accurate Deep Potential model for the Al–Cu–Mg alloy in the full concentration space^{*}

Wanrun Jiang(姜万润)^{1,2}, Yuzhi Zhang(张与之)^{3,4}, Linfeng Zhang(张林峰)^{5,†}, and Han Wang(王涵)^{6,‡}

MD: cMD

1. Hard sphere、Lennard-Jones、Morse、Hubbard、Heisenberg

Simple physical models, only related to pair interactions.
Used in theoretical physics, gas, etc.

2. FFMM、Compass

Gas, liquids and crystals composed by **small** molecules

3. CHARMM、AMBER、Gromos

Bio-molecules: proteins, RNA, DNA, membranes, molecular crystals
Used pair-additional assumptions

4. EAM, MEAM, inversion P(Chen-Mobius inversion), Tersoff P...

Inorganic compounds, graphene, alloy, carbon nanotubes, etc.

Common force fields

force field	originated by (year)	used for / coverage
AMBER	Kollman, UCSF (1994)	proteins, nucleic acids, carbohydrates
CHARMM	Karplus, Harvard (1983)	proteins, nucleic acids, lipids, organics
GROMOS	U. Groningen (1996)	proteins, nucleic acids, sugars, organics

OPLS	Jorgensen, Purdue (1988)	many liquids
ECEPP	Scheraga, Cornell (1975)	proteins
UFF	Rappe et al. (1992)	approximate for full periodic table
MM2/3/4	Allinger, U Georgia (1977)	small molecules and hydrocarbons

Assessment of GAFF2 and OPLS-AA General Force Fields in Combination with the Water Models TIP3P, SPCE, and OPC3 for the Solvation Free Energy of Druglike Organic Molecules

Dario Vasseti, Marco Pagliai, and Piero Procacci*

✓ **Cite this:** *J. Chem. Theory Comput.* 2019, 15, 3, 1983–1995

Publication Date: January 29, 2019 ✓

<https://doi.org/10.1021/acs.jctc.8b01039>

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Evaluation of nine condensed-phase force fields of the GROMOS, CHARMM, OPLS, AMBER, and OpenFF families against experimental cross-solvation free energies†



Check for updates

[Sadra Kashefolgheta](#), ^a [Shuzhe Wang](#), ^a [William E. Acree, Jr.](#) ^b and [Philippe H. Hünenberger](#) ^{*a}

$$V(r_{ij}) = \frac{q_i q_j}{r_{ij}} - \frac{C_i C_j}{r_{ij}^6} + D(B_i + B_j) \exp\left(\frac{A_i + A_j - r_{ij}}{B_i + B_j}\right)$$

Predicting Fracture Propensity in Amorphous Alumina from Its Static Structure Using Machine Learning

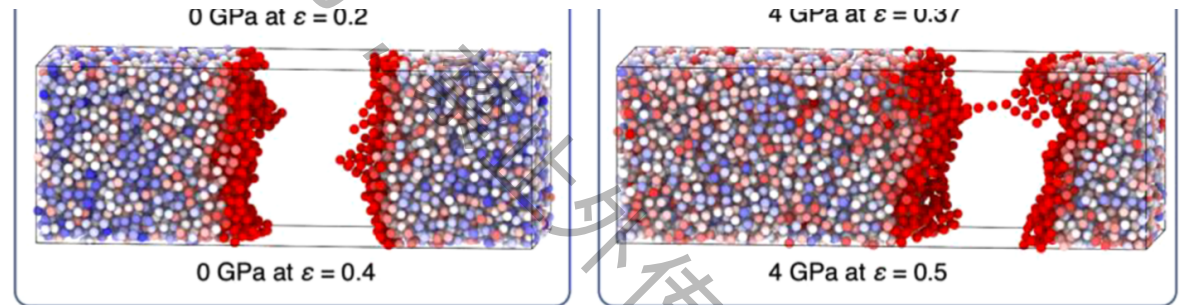
Tao Du, Han Liu, Longwen Tang, Søren S. Sørensen, Mathieu Bauchy,* and Morten M. Smedskjaer*



Cite This: *ACS Nano* 2021, 15, 17705–17716



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$$V(r_{ij}) = \frac{q_i q_j}{4\pi\epsilon_0 r_{ij}} - \frac{C_{OO}}{r_{ij}^6} + A_{OO} \exp\left(-\frac{r_{ij}}{\rho_{OO}}\right) + A_{MO} \exp\left(-\frac{r_{ij}}{\rho_{MO}}\right)$$

Sun, Jizhong, et al. "A simple transferable interatomic potential model for binary oxides applied to bulk α - Al_2O_3 and the (0001) α - Al_2O_3 surface." *Journal of crystal growth* **290** (2006): 235-240.

MD: interatomic interactions

$$\mathbf{F} = M\mathbf{a} = -\left(\frac{\partial E_{\text{pot}}(\mathbf{R})}{\partial \mathbf{R}}\right)$$

Gradients:

1. Analytic gradients

Analytical formula for $dE_{\text{pot}}(R)/dR$
AIMD, or Hellman-Feynman theorem,
or...

2. Numerical gradients: central difference

$$\left(\frac{\partial E_{\text{pot}}}{\partial R_i}\right) \approx \frac{E_{\text{pot}}(R_i + \delta R_i) - E_{\text{pot}}(R_i - \delta R_i)}{2\delta R_i}$$

Frontier for force field

1. Key: Accuracy + speed
2. Many-body interactions
(F. Paesani)
3. Quantum mechanics calculated data
thousands → tens or hundreds of thousands
several molecules → database of molecules
4. Machine Learning:
 - a. fitting the parameters
 - b. Directly learn the *ab initio* results

MD: polarizable force fields

Electrostatic functional forms

How detailed does the function need to be in order to describe reality?

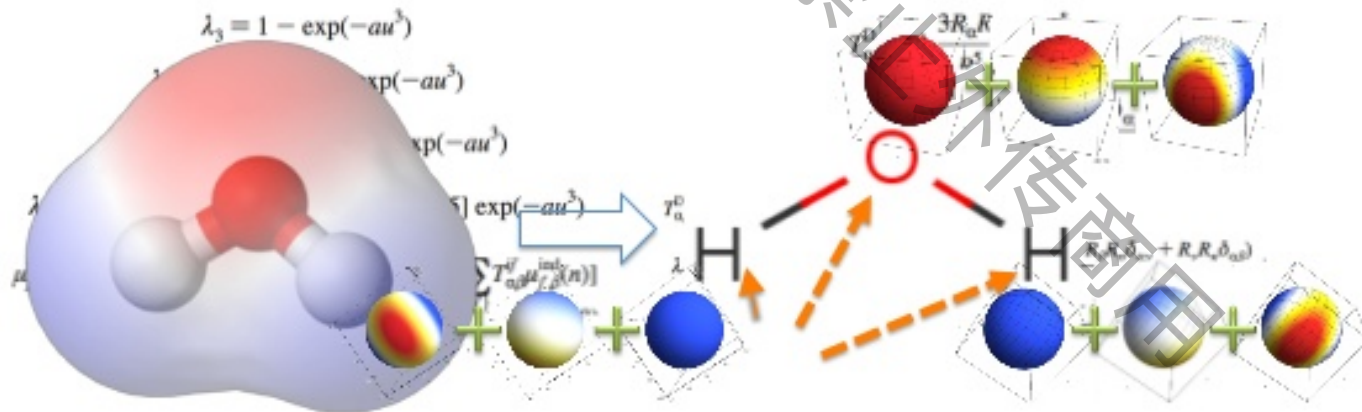
AMBER fixed-charge force field:

- Point charge on each atom

$$\sum_{i < j} \frac{q_i q_j}{r_{ij}}$$

AMOEBA polarizable force field:

- Point charge, dipole, and quadrupole on each atom
- Polarizable point dipole on each atom with short-range damping

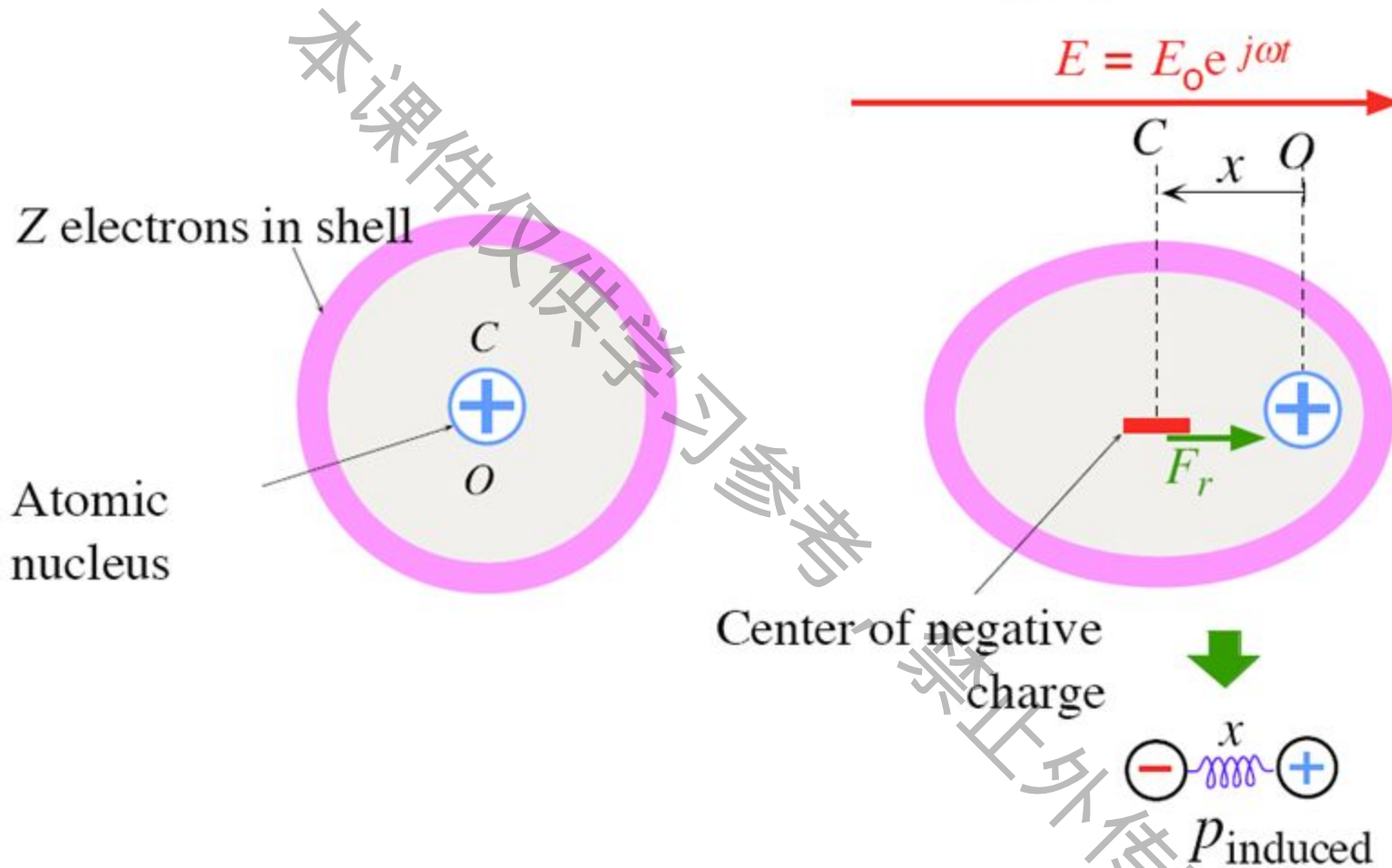


Evaluation of the AMOEBA force field for simulating metal halide perovskites in the solid state and in solution

J. Chem. Phys. **152**, 024117 (2020); <https://doi.org/10.1063/1.5131790>

 P. V. G. M. Rathnayake¹, Stefano Bernardi¹, and  Asaph Widmer-Cooper^{1,2,a)}

Electronic Polarization, ϵ_r and n



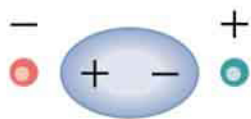
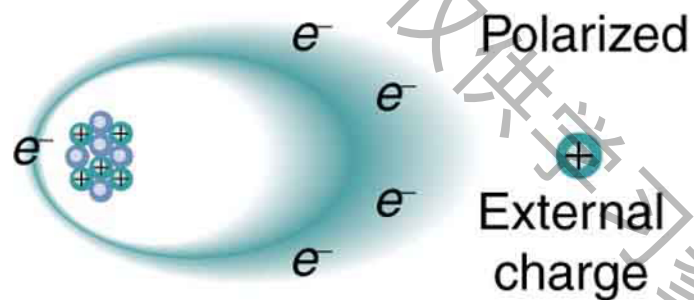
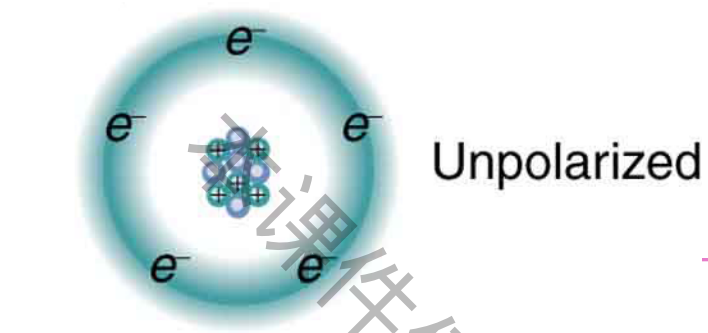
(a) A neutral atom in $E = 0$.

(b) Induced dipole moment in a field

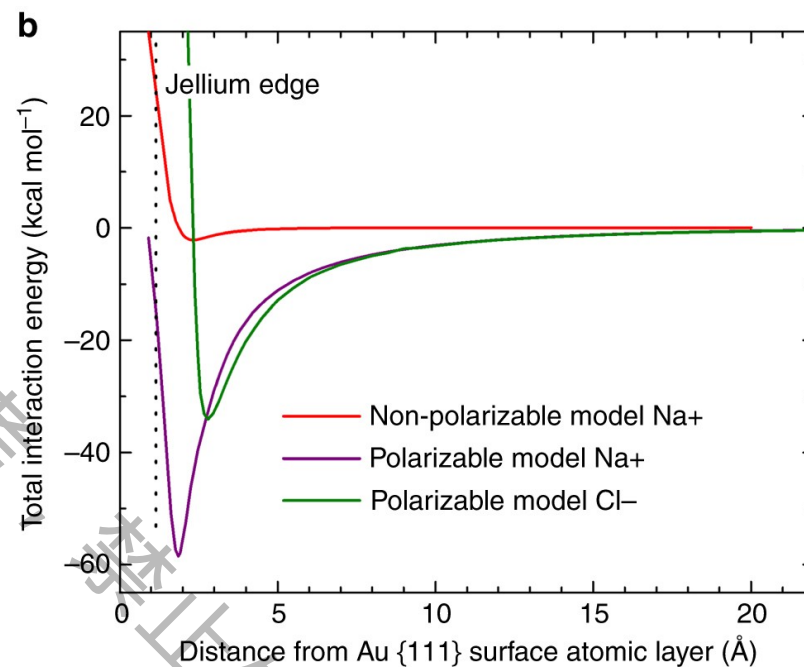
Electronic polarization of an atom. In the presence of a field in the $+x$ direction, the electrons are displaced in the $-x$ direction (from O), and the restoring force is in the $+x$ direction.

Fig 9.4

External
charge



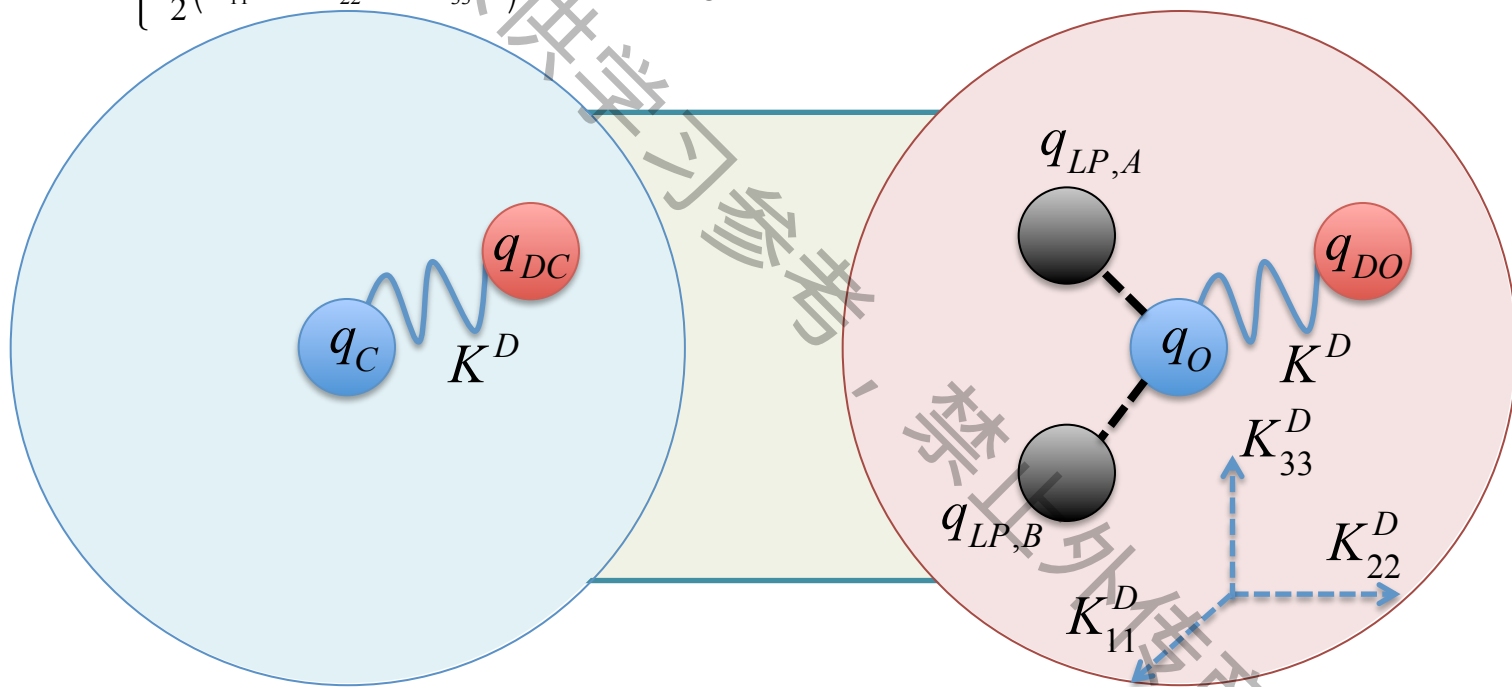
Large-scale view of polarized atom



I. L. Geada, et al. *Nat. Commun.* **2018**, 9, 716.

$$E_{\text{elec,Coulomb}} = \frac{1}{4\pi D} \left(\sum_{i \neq j} \frac{q_i q_j}{\| \mathbf{r}_i - \mathbf{r}_j \|} + \sum_{i \neq j} \frac{q_{D,i} q_j}{\| \mathbf{r}_{D,i} - \mathbf{r}_j \|} + \sum_{i \neq j} \frac{q_{D,i} q_{D,j}}{\| \mathbf{r}_{D,i} - \mathbf{r}_{D,j} \|} \right)$$

$$U_{\text{self}} = \begin{cases} K^D d^2, & \text{homogeneous} \\ \frac{1}{2} (K_{11}^D d^2 + K_{22}^D d^2 + K_{33}^D d^2), & \text{inhomogeneous} \end{cases}$$



Polarizability $\alpha = \frac{q_D^2}{K^D}$ dipole $d = \frac{q_D E}{K^D}$ induced dipole $\mu = \frac{q_D^2 E}{K^D}$

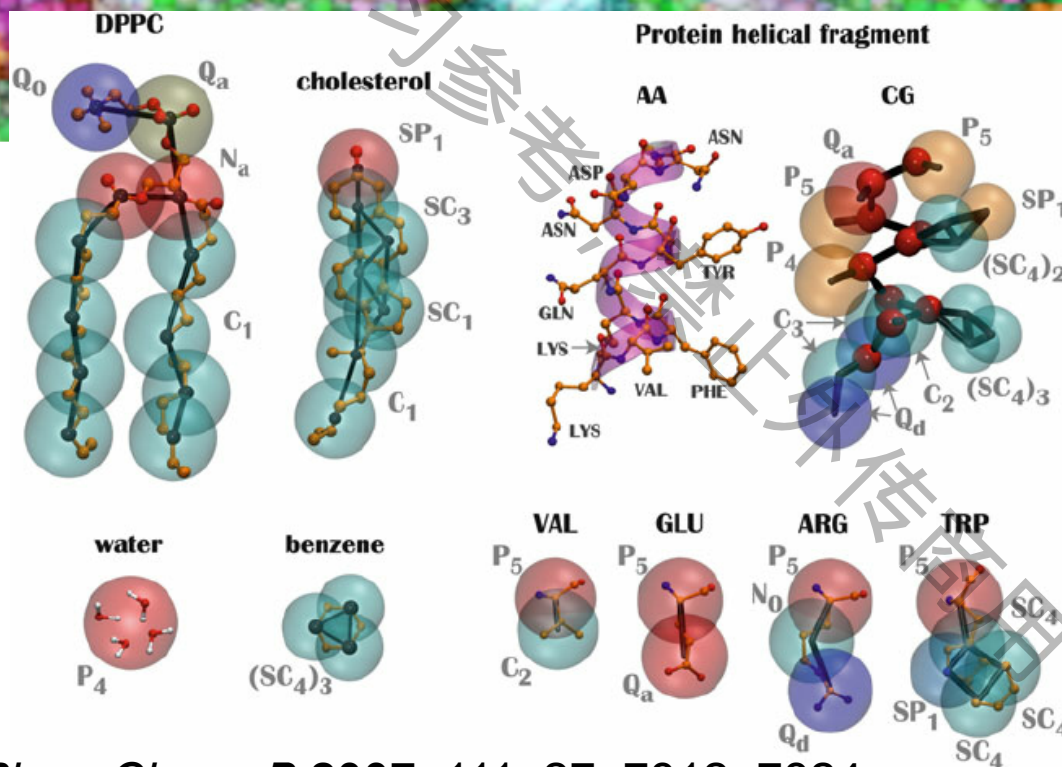
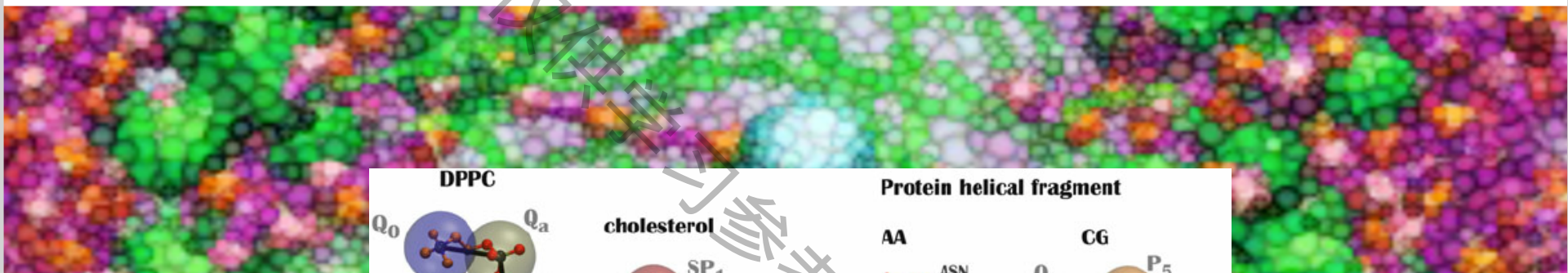
Table 1**The scope of available polarizable force fields**

Force fields	Polarizable models	Scope of biomolecules
AMBER ff02	Induced dipole	Proteins [99, 100] Nucleic acids [99] Atomic ions [96, 98]
AMOEBA	Induced dipole	Proteins [112, 127] Nucleic acids ^a Small molecules [128, 129] Atomic ions [130, 131]
CHARMM-FQ	Fluctuating charge	Proteins [92, 93] Lipids [132] Carbohydrates [133] Atomic ions [134]
CHARMM Drude	Classical Drude Oscillator	Proteins [135] Nucleic acids [136–139] Lipids [124, 140] Carbohydrates [141–144] Small molecules [119, 145–157] Atomic ions [47]

^aParameters have not been reported in the literature, but they are available through the website <http://dasher.wustl.edu>

FF: Coarse Grained Martini

Coarse Grain Forcefield for Biomolecules



On-the-fly updated polarized charge MD

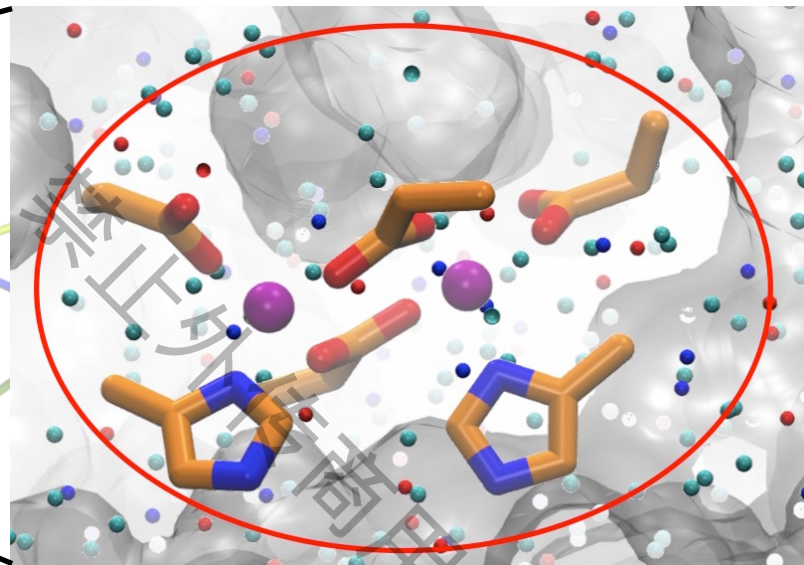
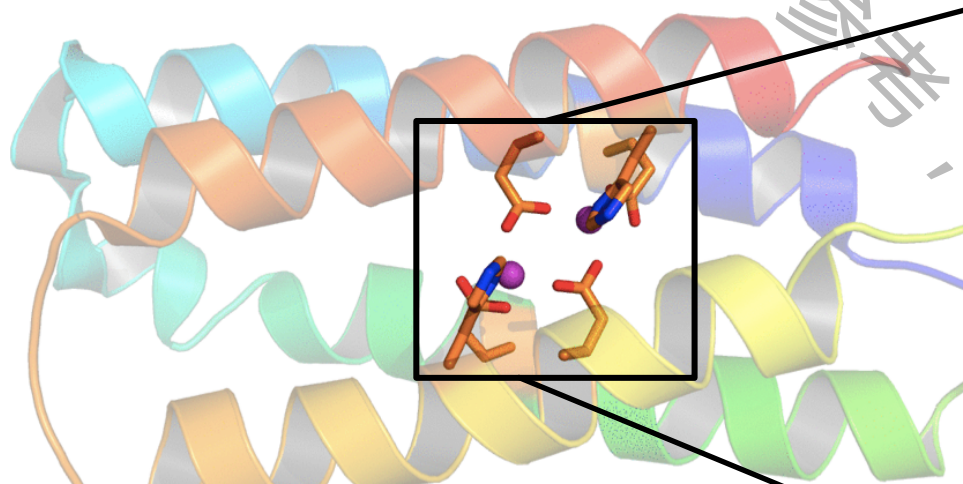
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Structure and Dynamics of a Dizinc Metalloprotein: Effect of Charge Transfer and Polarization

Yong L. Li,[†] Ye Mei,^{*†§} Da W. Zhang,^{||} Dai Q. Xie,[⊥] and John Z. H. Zhang^{*†§¶}

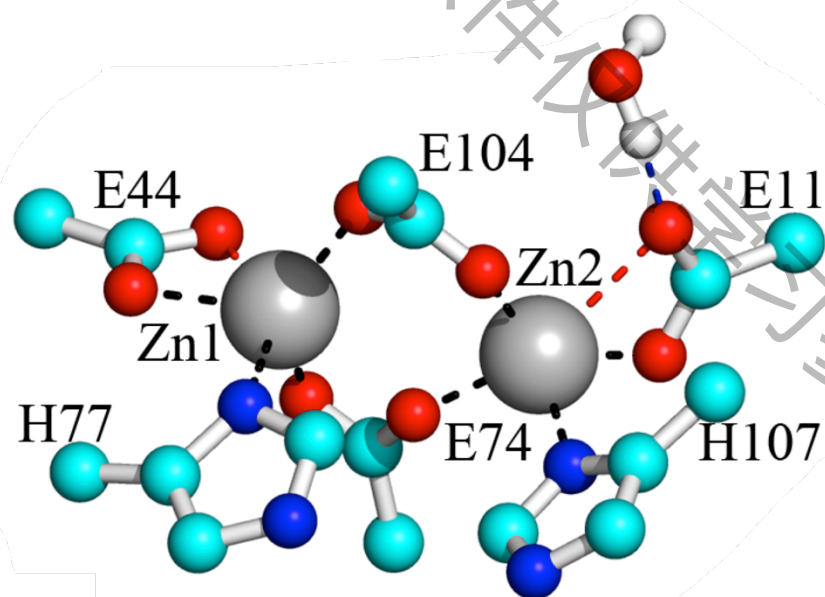


分子动力学模拟过程中实时更新电荷
成功获得**不对称**的配位中心结构

研究体系：双锌蛋白DFsc，作用：金属离子缓冲、胆固醇代谢、DNA合成

J. R. Calhoun, F. Nastro, O. Maglio, *et al. Structure*. 2008, 16, 210-215

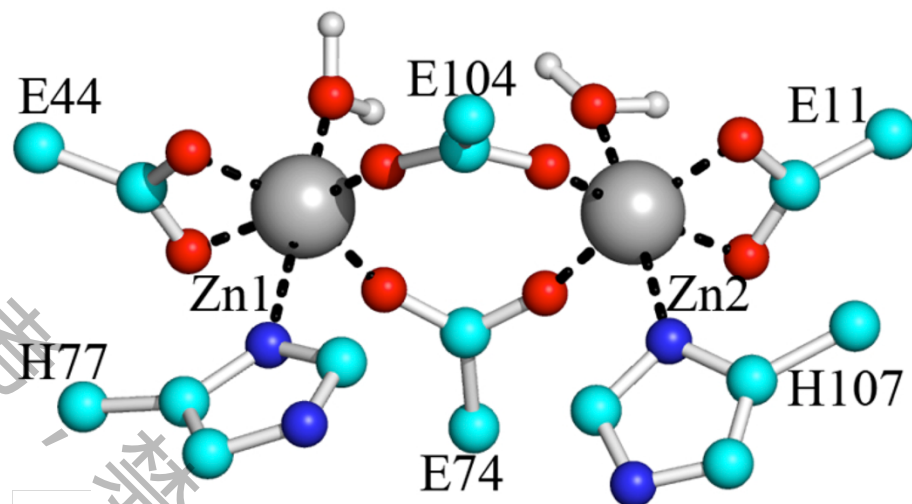
APPC模拟结果反映出体系的不对称性



APPC structure

锌离子电荷: $1.19e$ $1.36e$

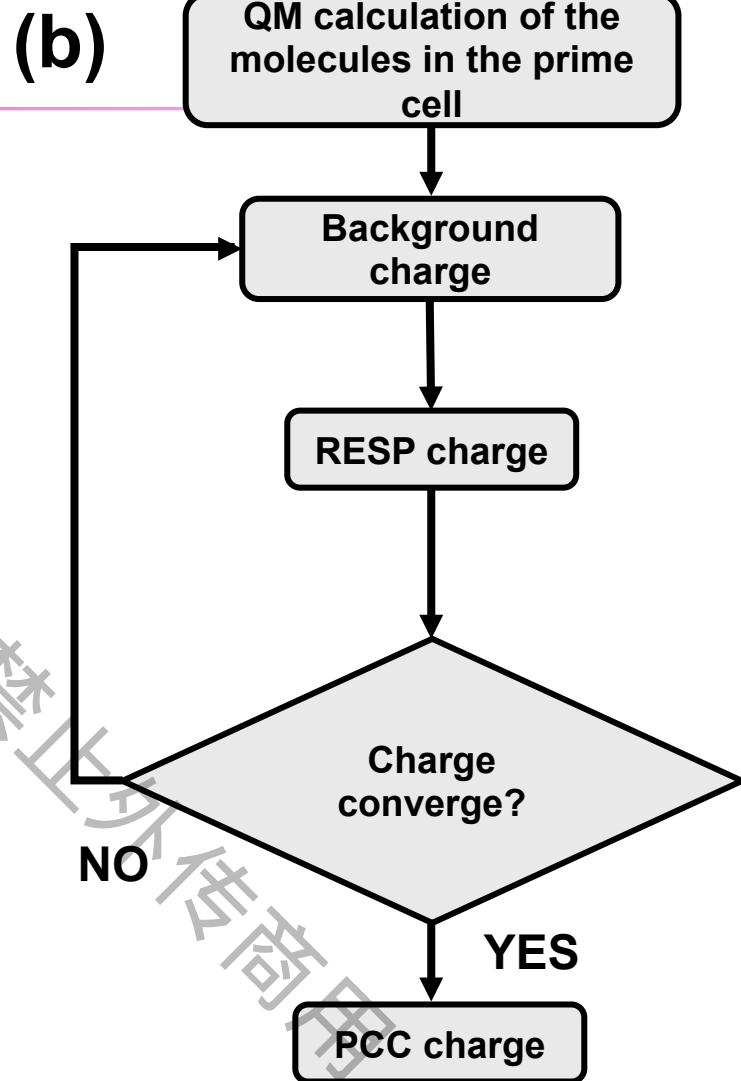
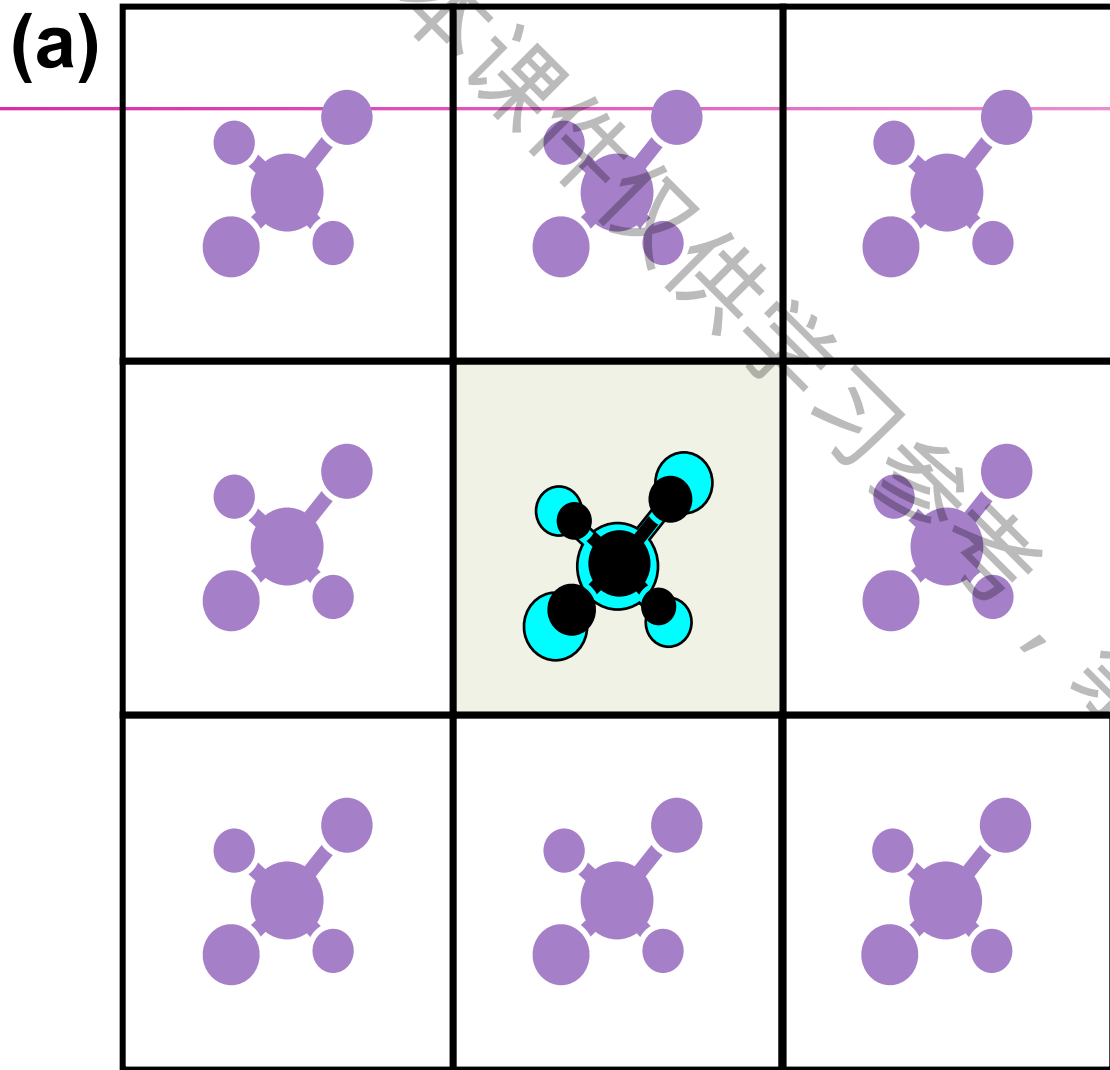
两锌离子不对称, 配位数为5/4,
不与水配位



AMBER99SB structure

$2.0e$ $2.0e$

两锌离子对称, 配位数为6/6,
与水配位



X. He et al. *Acc. Chem. Res.* **2014**, 47 (9), 2748-2757.
Y. Li et al. *J. Phys. Chem. B* **2011**, 115 (33), 10154-10162.

Polarized Crystal Charge (PCC)

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Publication Date: June 2, 2021 ✓

<https://doi.org/10.1021/acs.jpcc.1c03291>

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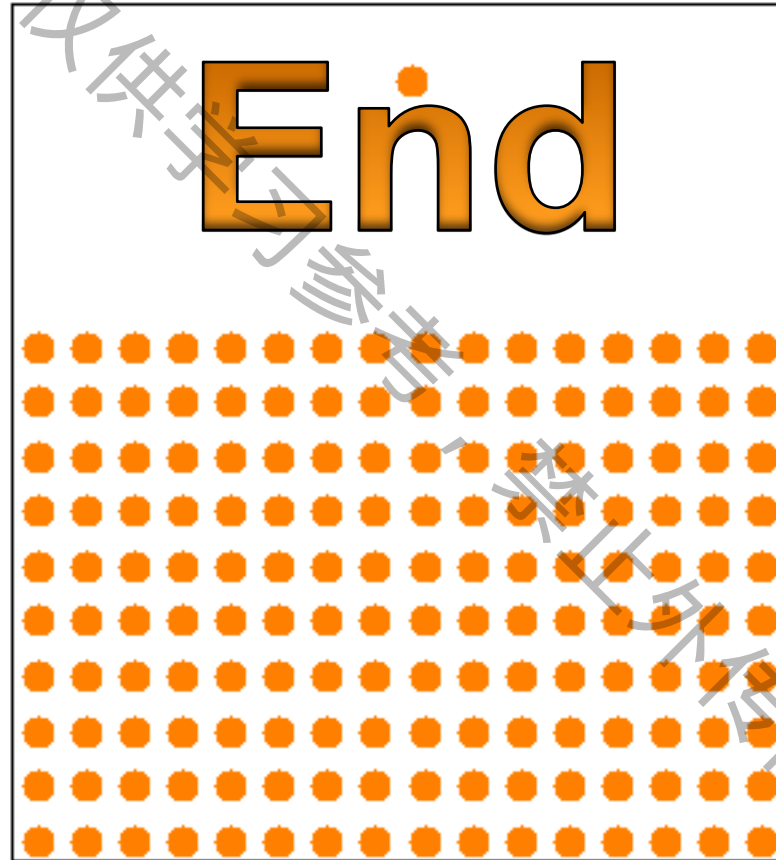
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