Reliable approximation in electronic structure methods:

from real space partitioning to quantum Monte Carlo approaches

Amanda Dumi Seminar at Oak Ridge National Laboratory

When is DFT not enough?

- DFT citation/publications demonstrate it's applicability and often reliable performance
- In strongly correlated regimes, or when functional and system physics do not agree

Reduce cost through various system reductions

Sampling Configurations Quantum Monte Carlo: applications and developments

Hilbert Space

selected CI as approximate trial wave functions

Real Space

Determining molecular fragments with unsupervised machine learning (UML)

QMC application: periodic absorption¹

H on graphene

Method	Binding energy (meV)
DMC PW91	-691 ± 19 -810 to -830, -870
PBE	-790, -840, -980

- functional dependent binding energy
- a need for benchmark values

¹A. Dumi et al, 'The Binding of Atomic Hydrogen on Graphene from Density Functional Theory and Diffusion Monte Carlo Calculations', *The Journal of Chemical Physics*, 156.14 (2022), 144702DOI:10.1063/5.0085982.

Diffusion Monte Carlo

• Recast the Schrodinger equation in imaginary time

$$rac{\partial \left|\Psi
ight
angle }{\partial au}=-\hat{H}\left|\Psi
ight
angle$$

- A formal soltuion to this:

$$|\psi\left(\tau_{1}+\delta\tau\right)\rangle=e^{-\hat{H}\delta\tau}\left|\psi\left(\tau_{1}\right)\right\rangle$$

Anything non-orthogonal to the ground state is going to decay out exponentially

$$\lim_{\tau \to \infty} |\psi(\tau)\rangle = c_0 e^{-\epsilon_0 \tau} \, |\phi_0\rangle$$

$$-\frac{\partial \psi({\bf R},\tau)}{\partial \tau} =$$

$$\left[\sum_{i=1}^{N} -\frac{1}{2} \nabla_{i}^{2} \psi(\mathbf{R}, \tau)\right] \ \text{diffusion term}$$

+

$$(V(\mathbf{R})-E_T)\,\psi(\mathbf{R},\tau)$$
 branching term

Importance sampling

• use of trial wavefunction for efficient sampling.

Fixed-node approximation

- Antisymmetry of fermions causes sampling problems
- Solution: fix nodes of trial wavefunction
- necessitates accurate nodal surface of trial wavefunction

QMC application: periodic absorption

H on graphene

Method	Binding energy (meV)
PBE	-820
PBE	-871
PBE0	-851 (-800)
HSE	-794 (-743)
DMC	-691 ± 19
PW91	-810 to -830, -870
PBE	-790, -840, -980

- QMCPACK binding energies provide benchmark.
- Hybrid functionals are very close.
- Even with close binding energies, density seems to disagree in bonding region.

Reduce cost through various system reductions

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QMC application: non-valence correlation bound anion²

NVCB rely on accurate description of correlation to bind

$$E_{corr} = E_{true} - E_{HF}$$

model system: H_2O_4

Radial integration of Hartree-Fock orbital density, R=4 Angstroms

Selected-CI

$$\Psi_n = \sum c_I^{(n)} \left| D_I \right\rangle$$

$$e_{\alpha} = \frac{\langle \Psi^{(n)} | \hat{H} | \alpha \rangle^2}{E^{(n)} - \langle \alpha | \hat{H} | \alpha \rangle}.$$

 $\{|\alpha\rangle\}$

single & double excitations

$$\{|\alpha\rangle\}_*^{(n)}$$

 Ψ_n

 Ψ_{n+1}

${\rm H_2O_4}$ Results

	wave function	basis set	EBE (meV)	
D/HF		aug-cc	-pVDZ+7s7p	183 ± 10
SD/HF		aug-cc	-pVDZ	176 ± 12
SD/HF		cc-pV]	DZ	-528 ± 25
SD/B3LYP		aug-cc	-pVDZ+7s7p	212 ± 11
SD/HF(N)//S	SD/NO SDCI(A	aug-cc	-pVDZ+7s7p	205 ± 10
SD/HF(N)//2	MD/NO SDCI(A	A) aug-cc	-pVDZ+7s7p	202 ± 12

²S. Upadhyay et al, 'The Role of High-Order Electron Correlation Effects in a Model System for Non-Valence Correlation-Bound Anions', *The Journal of Chemical Physics*, 153.22 (2020), 224118DOI:10.1063/5.0030942.

	wave function	basis set	EBE (meV)	
MD/CIPSI N	O	aug-cc	-pVDZ+3s1p	190 ± 9

• Summary of selected-CI success/challenges and overall performance of other methods.

R = 7

- HF binds electron -> correlation not essential
- still orbital shape has noticeable changes
- DFT can recover those changes.

R = 4

- NCVB regime
- HF does not bind
- CIPSI shape changes slight, impacts energy.
- SDCI shape seems wrong

Follow-up Questions Ongoing projects at Sandia National Laboratories

- Can various selected CI approaches or later optimizations improve the compactness and quality of the trial wave function?
- Is there a balance in how well we are capturing dynamic vs static correlation in these systems? Especially in cases of energy differences.
- \hookrightarrow Aluminum systematic study: exploring various multideterminant generation and optimization schemes
 - Selected CI captures static correlation, but should we take a second look at how dynamic correlation is captured?

 \hookrightarrow Jastrow form study

Jastrow Factor forms

$$\Psi(\{r_i\}\{r_I\}) = \exp(\mathcal{J})D(\{r_i\})$$

standard: parameterize separate 1,2, and 2-body terms. optimize parameters with VMC

A possible alternative form:

- Using framework of SNAP atomic potential
- represent the particle densities as bispectrum components.³
- A perspective of particle neighborhoods which require fewer parameters for VMC optimization

Projection of the ρ particle density on the surface of the 4-D sphere:

$$E_{SNAP}^{i}\left(\mathbf{B}^{i}\right)=\beta_{0}^{\alpha_{i}}+\sum_{k=1}^{K}\beta_{k}^{\alpha_{i}}B_{k}^{i}=\beta_{0}^{\alpha_{i}}+\beta^{\alpha_{i}}\cdot\mathbf{B}^{i}$$

$$\begin{split} B_{j_1,j_2,j} &= \sum_{m_1',m_1 = -j_1}^{j_1} \sum_{m_2',m_2 = -j_2}^{j_2} \sum_{m',m = -j}^{j} \left(c_{m'm}^j\right)^* C_{j_1m_1j_2m_2}^{jm} \\ &\times C_{j_1m_1'j_2m_2'}^{jm'} c_{m_1'm_1}^{j_1} c_{m_2'm_2}^{j_2} \end{split}$$

TO DO:

initial python sandbox for testing unit tests ensure single and multiple species work accessing LAMMPS API object interface with QMCPack

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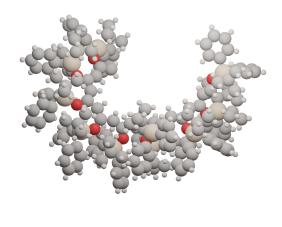
Real Space

Determining molecular fragments with unsupervised machine learning (UML)

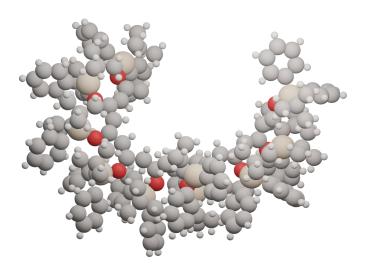
³A. P. Bartók, R. Kondor, and G. Csányi, 'On Representing Chemical Environments', *Phys. Rev. B*, 87 (2013), 184115DOI:10.1103/PhysRevB.87.184115.

⁴,A. P. Thompson et al, 'Spectral Neighbor Analysis Method for Automated Generation of Quantum-Accurate Interatomic Potentials', *Journal of Computational Physics*, 285 (2015), 316–30DOI:https://doi.org/10.1016/j.jcp.2014.12.018.

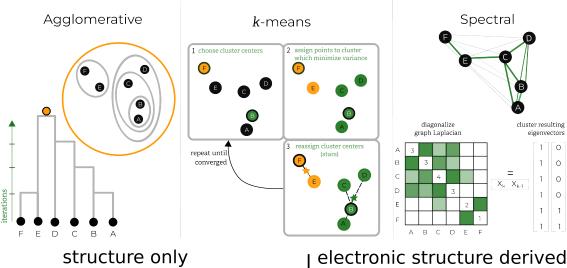
Fragmenting with unsupervised machine learning $\mbox{\sc Problem}$



Fragmenting with unsupervised machine learning Problem



Approach



coarse graining information

Cartesian

$$D_{AB} = \frac{1}{R_{AB}}$$

covalent radii (cr)

$$D_{AB} = \begin{cases} 1 \text{ if } R_{AB} \leq 1.1(A_{cr} + B_{cr}) \\ 0 \text{ otherwise} \end{cases}$$

Mayer bond matrix (Mbm)

$$B_{\mathrm{AB}}^{\mathrm{Mayer}} = \sum_{s}^{\mathrm{on}} \sum_{t}^{\mathrm{A}} \quad (\mathbf{PS})_{st} (\mathbf{PS})_{ts}$$

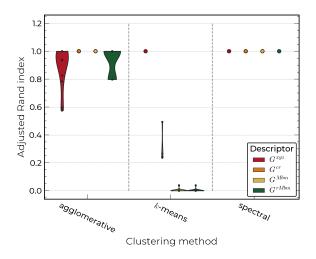
rounded Mbm (rMbm)

$$D_{AB} \ = \begin{cases} \lceil B_{AB} \rceil \text{ if } \{B_{AB}\} \ >= 0.5 \\ \lfloor B_{AB} \rfloor \text{ otherwise} \end{cases}$$

Performance

Systems:

- water clusters
- methylthiophenes
- small polymer systems



Performance:

Summary:

- UML methods are able to resolve fragments
- spectral clustering performs well and is insensitive to descriptor

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Conclusions

In regimes of strong electron correlation, mean field methods may be problematic and require need alternatives

Presented on three approaches:

- QMC: application to $\mathrm{H}_2\mathrm{O}_4$ and H on graphene \hookrightarrow systems where DFT varies.
- SCI methods in H_2O_4 as trial wave function
- UML MF: offer an approach to determining molecular fragmentation automatically, with very little information

References

Albert P. Bartók, Risi Kondor, & Gábor Csányi, 'On Representing Chemical Environments', *Phys. Rev. B*, 87 (2013), 184115DOI:10.1103/PhysRevB.87.184115

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