TUM seminar Theory and Computation of Materials and Molecules Electronic coarse-graining of semiconductors: methodology and applications

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Outline

- Introduction and illustrative examples
- Methodology
- Applications

Electronic Coarse-Graining (ECG) in examples

- When do we need ECG? When system is too complex for use of conventional electronic structure methods (e.g. polymers)
- What is the "price" for ECG? Only frontier orbitals are calculated (enough for majority of electronic phenomena)
- ECG example 1/4: biopolymers (localized states \Rightarrow trivial ECG)



ECG: more precise formulation

ECG idea/goal: Get minimal basis providing accurate description of a particular electronic property under molecular fluctuations (large-scale/low-energy electronic phenomena: UV-Vis spectra, transport)

Output: electronic basis + electronic Hamiltonian



All-electron density



CG density (2 MOs)

Requirements:

- Robustness of CG basis wrt molecular fluctuations
- Robustness and scalability of CG algorithm
- Quality control of CG basis and matrix elements
- Accurate extrapolation to infinite system (if needed)

ECG example 2/4: electronic transport in a molecular solid



- Coarse grain electrons to one site per molecule
- Simplify molecular motions to harmonic vibrations
- · Linearize coupling between electrons and molecular motions

$$\implies \sum_{ij} \boldsymbol{H}_{ij}^{1p} c_i^{\dagger} c_j + \sum_{\alpha} \hbar \omega_{\alpha} \left(b_{\alpha}^{\dagger} b_{\alpha} + \frac{1}{2} \right) + \sum_{ij\alpha} \hbar \omega_{\alpha} \boldsymbol{g}_{ij\alpha} \left(b_{\alpha}^{\dagger} + b_{\alpha} \right) c_i^{\dagger} c_j$$

Then solve this Hamiltonian (e.g. in small polaron hopping approximation) Annu Rev Phys Chem 66, 305 (2015) first review – Chem Rev 104, 4971 (2004)

ECG in examples 3/4: less trivial coarse-graining



*e.g. PBE-D3 is good for crystal structure but bad for e-properties JCTC 19, 8481 (2023)

ECG in examples 4/4: nontrivial coarse-graining (and applications)



energy storage - Chem Sci 13, 8161 (2022)

... and other structurally complex systems

Methodology of ECG

Requirements:

- Robustness of CG basis wrt molecular fluctuations
 Case study of A-D-A molecules (e.g. NFA for solar cells)
- Robustness and scalability of CG algorithm

 Implementation
- Quality control of CG basis and matrix elements
- Accurate extrapolation to infinite system (if needed)
 Efficient parametrization of tight-binding models

Robustness of CG basis wrt molecular fluctuations:

Case study of A-D-A molecules (non-fullerene acceptors in solar cells) JCP 159, 024107 (2023)



Determine qualitative and quantitative errors of inaccurate ECG:

- Is electron localized on acceptor or delocalized over molecule?
- How inaccurate ECG influences charge transport parameters?

• To be compared: 1-site vs 2-site, 2-LUMO vs 2-LMO models MO=Molecular Orbital, LUMO=Lowest Unoccupied MO, LMO=Localized MO

Main result





For molecular solids, the geometry is sampled by classical MD. For molecules, quantum effects are considered by sampling thermal vibrations. $_{9/32}$

More rigorously: 2-LMO model is the most appropriate

Statistical analysis of the basis deficiency in a coarse-grained description of electron MO and NO (i.e. LUMO and anion NO):



How inaccurate ECG influences calculated mobility?

In a solid A-D-A molecules form 2 intermolecular contacts per A

- \implies coordination number of electronic connectivity graphs is 3
- \implies either honeycomb (2D) or K4/Laves (3D) lattices



Analysis of effective masses and hopping amplitudes (\sim mobility) shows that the main effect is that 1-site model underestimates intermolecular couplings by factor of 2

Methodology of ECG

Requirements:

- Robustness of CG basis wrt molecular fluctuations
- Robustness and scalability of CG algorithm — Implementation
- Quality control of CG basis and matrix elements
- Accurate extrapolation to infinite system (if needed)

Robustness and scalability of ECG algorithm

Algorithm

- 1. Determination of coarse-grained (CG) basis
- 2. Parametrization of tight-binding Hamiltonian
- 3. Quality control (for new systems)

ECG might often look trivial but here "the devil is in the details", so there are many challenges in practical implementation:

- Large systems (large supercells)
- Entangled bands (overlapping in energy or hybridized)
- Cluster approximation to infinite systems
- Complex cases (e.g. excited states, no fixed SCF)
- Parametrization of complex models (e.g. electron-phonon)
- SCF and total energy in CG-basis

Determination of coarse-grained (CG) basis

In what follows we consider coarse-graining of molecular orbitals (MOs)

Algorithm

- 1. MOs of interest must be representable in CG basis \implies CG MOs are usually localized MOs (LMOs)
- 2. Tight binding parameters (ε_i and t_{ij}) are obtained by projection of Fock matrix of model fragments onto LMOs
- 3. Effective Hamiltonian of a large system is obtained by its fragmentation into the model fragments

Why LMO-based approach is working:

- Slater determinant is invariant under rotations of 1e orbitals
- Locality of phenomena: locality of 1e-density matrix and Hamiltonian, often only local SCF is important [V Heine, Solid State Phys 35, 1 (1980)]

Localization of molecular orbitals for coarse-graining

- Localization procedure is not unique Acc Chem Res 47, 2758 (2014) (trade off between spatial and energy localization)
- Projection approaches are scalable and robust
- Localization to unit cell is performed by Wannier functions
- The largest problem is bands entanglement (e.g. π and σ)
- LMOs can be used to get SCF (MOZYME code in MOPAC, fragmentation methods Chem Rev 112, 632 (2012))

Algorithm

- 1. Reduce to unit cell (Wannier, or cluster in PBC or OBC)
- 2. Define projector (quality of the projector is critical)
- 3. Select initial MO subspace, project, and refine MOs
- 4. Convert LMOs to CG-basis (e.g. by block diagonalization)

Projecting MO to LMO

Open-source code MolMod/LocalizeMO

Project on/in/out by SVD of M^+SL or $M^+P'SPM$,

where S – overlap, M – MOs, L – pre-LMOs, P – projector

Possible projector: subset of AOs (geometrical fragmentation) Possible pre-LMOs: LMOs by other method ('ad hoc' approach) Example: benzene σ and σ^* LMOs (NBO/on/in/out/...)



Locality and fast convergence of tight-binding parameters... Illustrated by NBO analysis of the valence band of blue (A7) phosphorene



- There are two symmetry unique NBOs
- Error in estimation energy difference between them is 270 meV for the smallest cluster and 42 meV for the next one in series
- Error in estimation coupling between them is 18 meV for the smallest cluster and 3 meV for the next one in series

Already the smallest fragment can be used for transfer integrals

... small clusters are enough for parametrization Example of hydrogen-passivated graphene clusters – graphene nanoflakes



Methodology of ECG

Requirements:

- Robustness of CG basis wrt molecular fluctuations
- Robustness and scalability of CG algorithm
- Quality control of CG basis and matrix elements
- Accurate extrapolation to infinite system (if needed)
 Efficient parametrization of tight-binding models

Efficient parametrization of tight-binding (TB) models

Approaches to first-principles parametrization:

- Fit by one-electron energies (mainstream)
 - + Only energies are needed
 - Only energies are accurate for sure
 - Systematic accuracy improvement requires ML-fitting
 - Best for simple and minimal TB models
- Derive from Fock matrix (natural to ECG)
 - + Numerically exact in principle (no fitting)
 - + Transferable TB elements (e.g. cluster-to-solid)
 - More complicated in implementation
 - Wave-function is needed
 - Best for complex and accurate TB models

Parametrization of TB models from cluster calculations https://cmsos.github.io/tbm



tions (DFT) Output: Symmetry-unique TB elements

Input: Cluster calculations (DFT)

Technical challenges and solutions

https://cmsos.github.io/tbm

Use symmetry for TB model formulation, solution, and parametrization

- 10 bonds in diamond \implies 100 symmetry-unique dimers \implies 5000 dimers per primitive cell \implies 10⁷ comparisons
- FiniteGroups package identifies 1500 symmetry-unique dimers in diamond per minute
- Wave-function symmetry is currently supported only for 1 orbital per site models enough for valence *sp*-orbitals

Use library of electronic structure prototypes

- Structurally different materials might have the same TB model
 - Triangular lattice herringbone packing
 - Honeycomb lattice brickwork packing of A-D-A molecules
 - K4 lattice wiremesh packing of A-D-A molecules
- We can compare materials by their effective Hamiltonians rather than computed properties

Applications of ECG

Calculate electronic structure of infinite systems:

- Solids of large flexible molecules such as A-D-A molecules Chem Mater 33, 966 (2021); JCP 159, 024107 (2023)
- Polymers: electronic structure of a single polymer chain Chem Sci 8, 1146 (2017), Solar Energy 198, 605 (2020)
- Polymers: electronic structure of crystalline polymers work in progress

Analyze localized states in infinite systems:

Metal-organic polymers and frameworks

Chem Sci 13, 8161 (2022)

Polarons in non-polar semiconductors

JPCL 12, 4674 (2021)

Traps in organic semiconductors

work in progress

Summary

- Electronic Coarse-Graining is a powerful tool (sometimes the only one) for calculation and analysis of electronic structure of complex systems:
 - Anything that can be monomerized (polymers, COFs, MOFs)
 - Solids and clusters of large molecules
 - Defects in solids
 - Infinite systems from finite-size calculations

Project web-page:

- Electronic Coarse-Graining https://zhugayevych.me/research/ECG.html
- Computational Materials Science of Organic Semiconductors https://cmsos.github.io