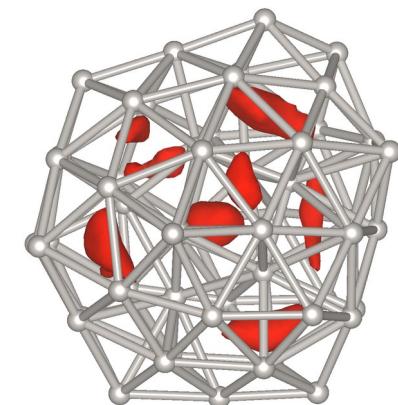
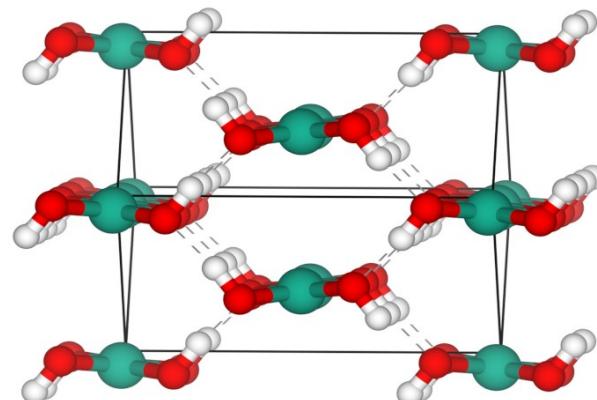


Global crystal structure optimization with ASE (and with a little help from DFTB)



Maxime Van den Bossche

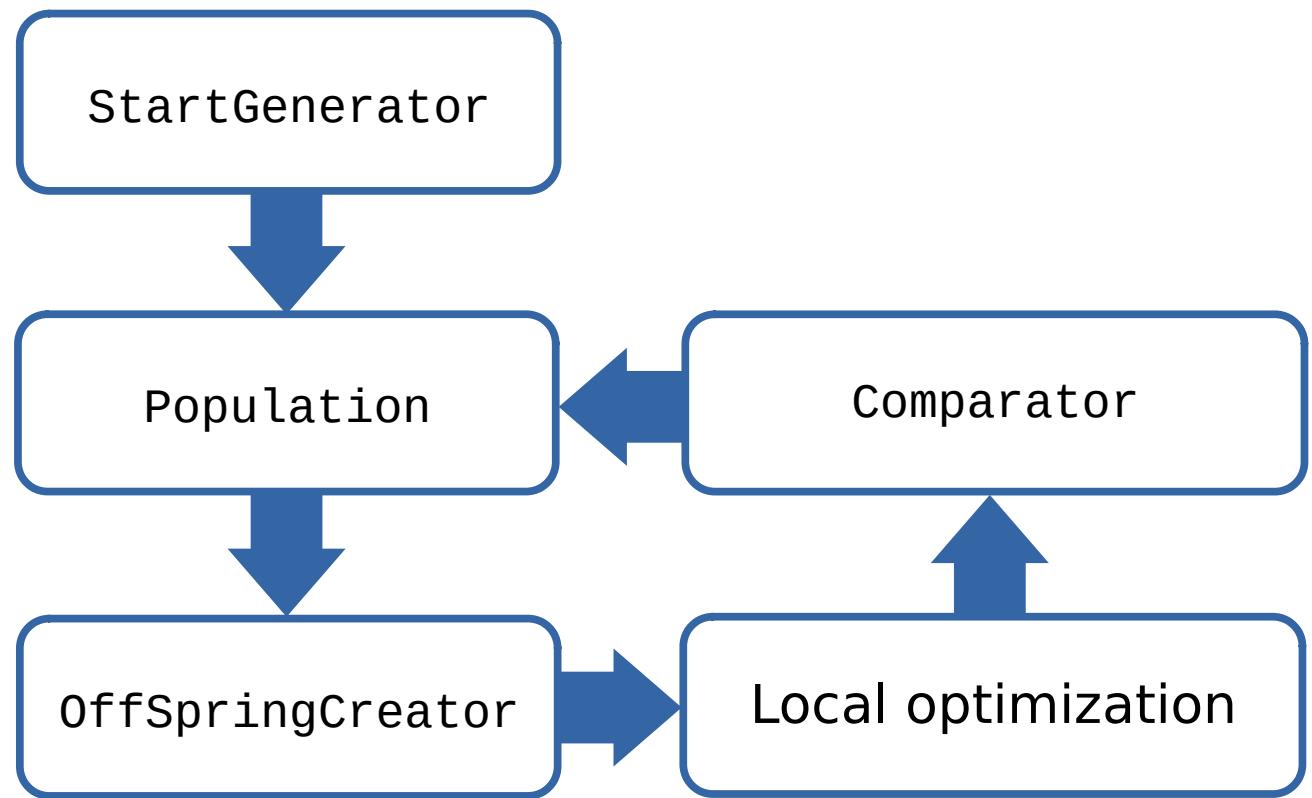
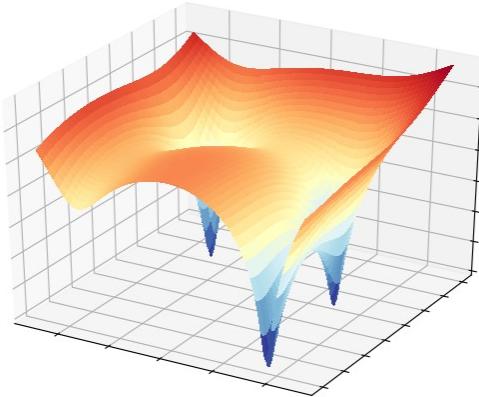
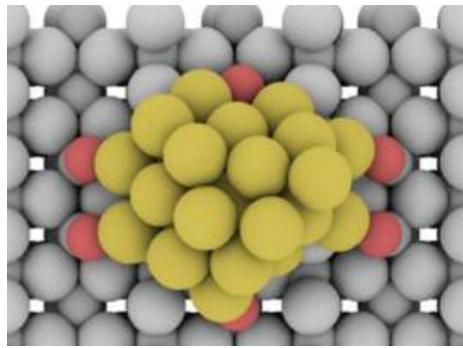


Overview

1. Functionality for crystal structure prediction (`ase.ga`)
2. Calculators: DFTB+ and QuantumEspresso
3. Miscellaneous
4. Wrap-up

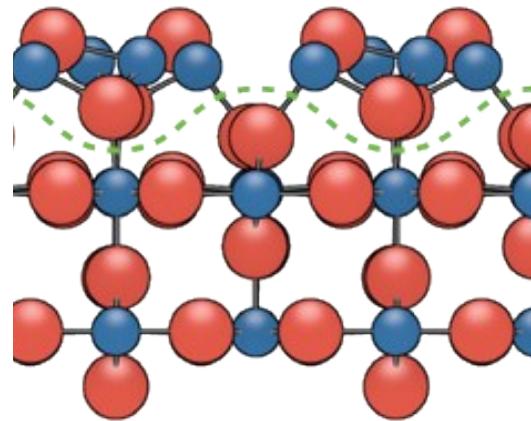
ASE-GA crystal structure prediction

Genetic algorithm workflow with ase.ga:



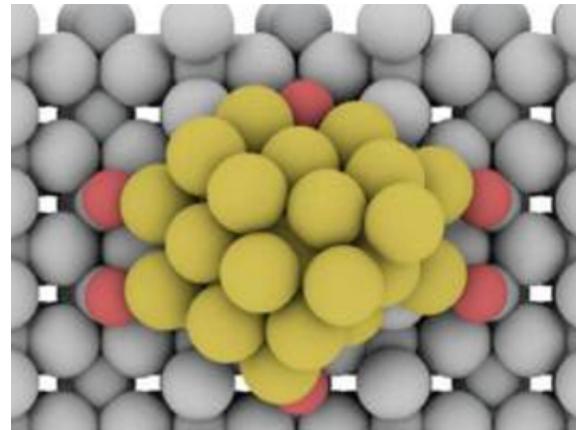
ASE-GA crystal structure prediction

$\text{SnO}_2(110)\text{-(4x1)}$



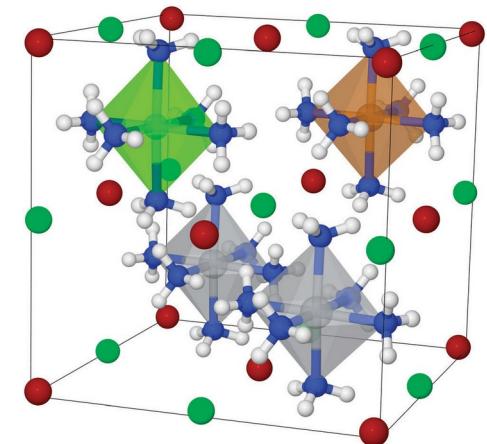
Merte et al.,
PRL 119 (2017)

Au/TiO_2



Vilhelmsen, Hammer,
ACS Catal. (2014)

$\text{M}_a\text{X}_b(\text{NH}_3)_c$



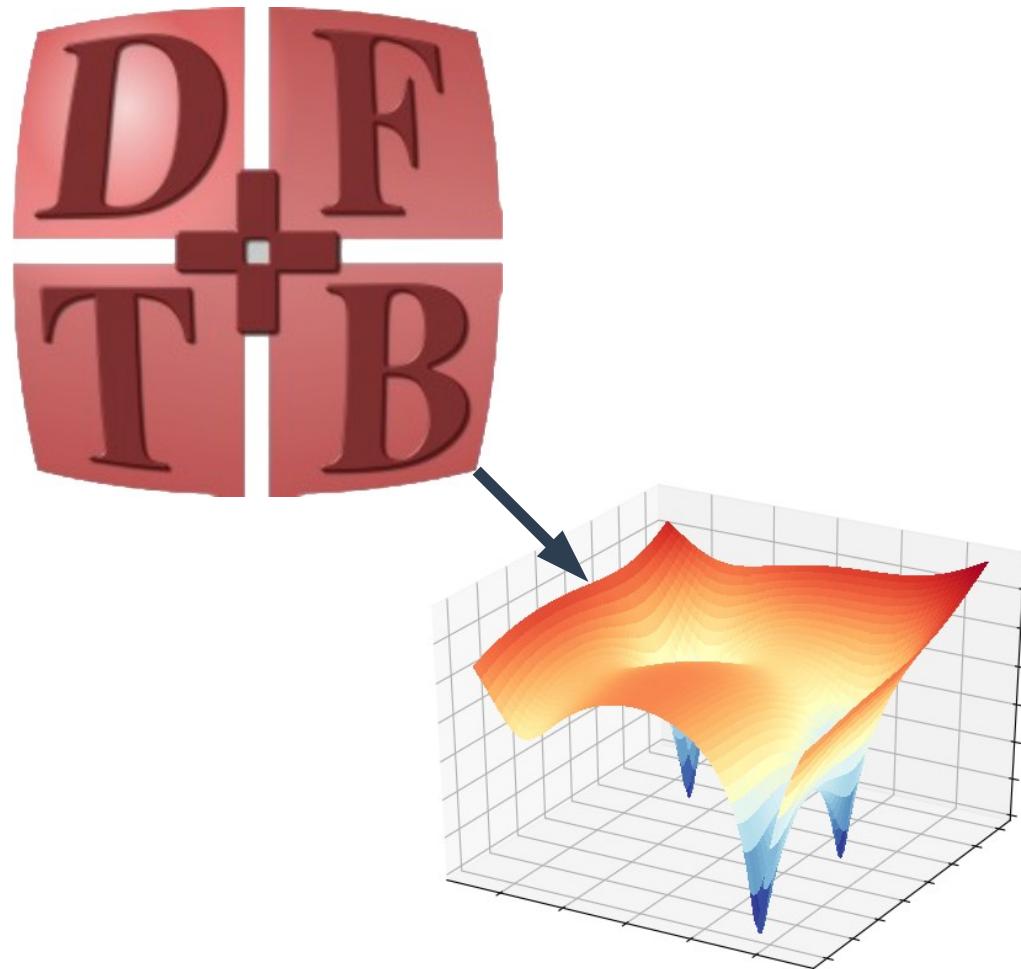
Jensen et al.,
PCCP (2014)

Previous implementation = without ‘global’ cell vector optimization

ASE-GA crystal structure prediction

Motivation:

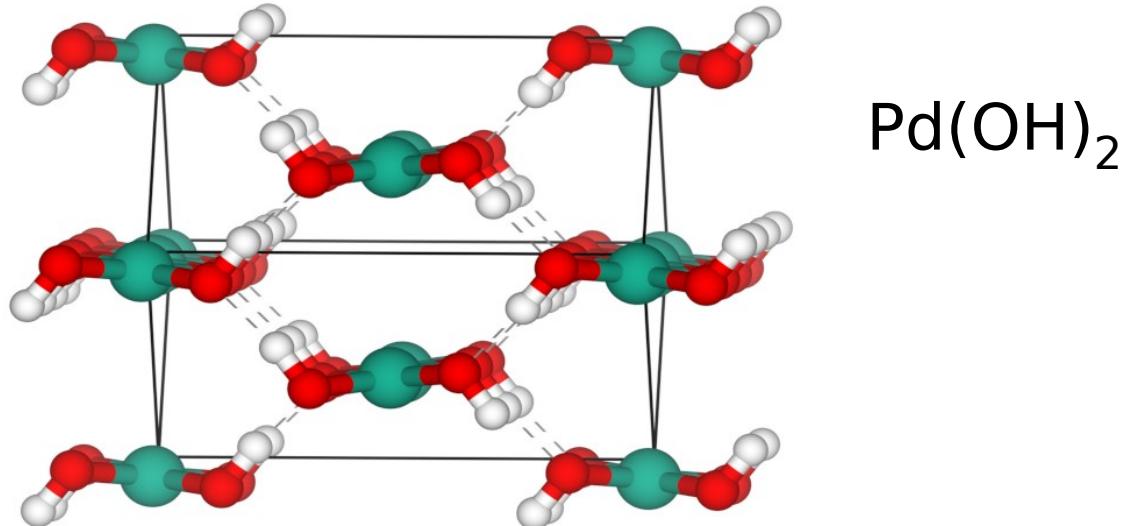
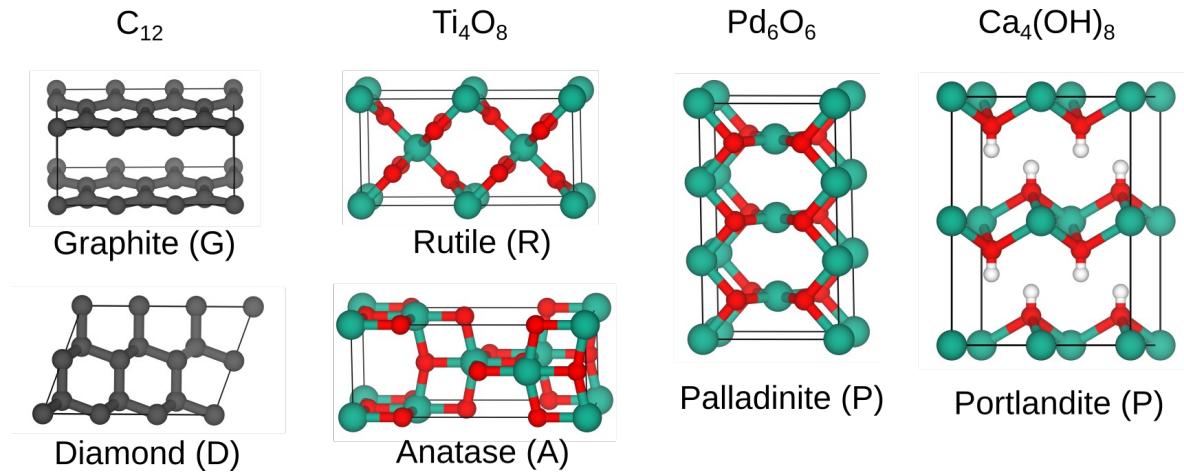
Development of
Tight-binding
Approximation
eNhanced
Global
Optimization



ASE-GA crystal structure prediction

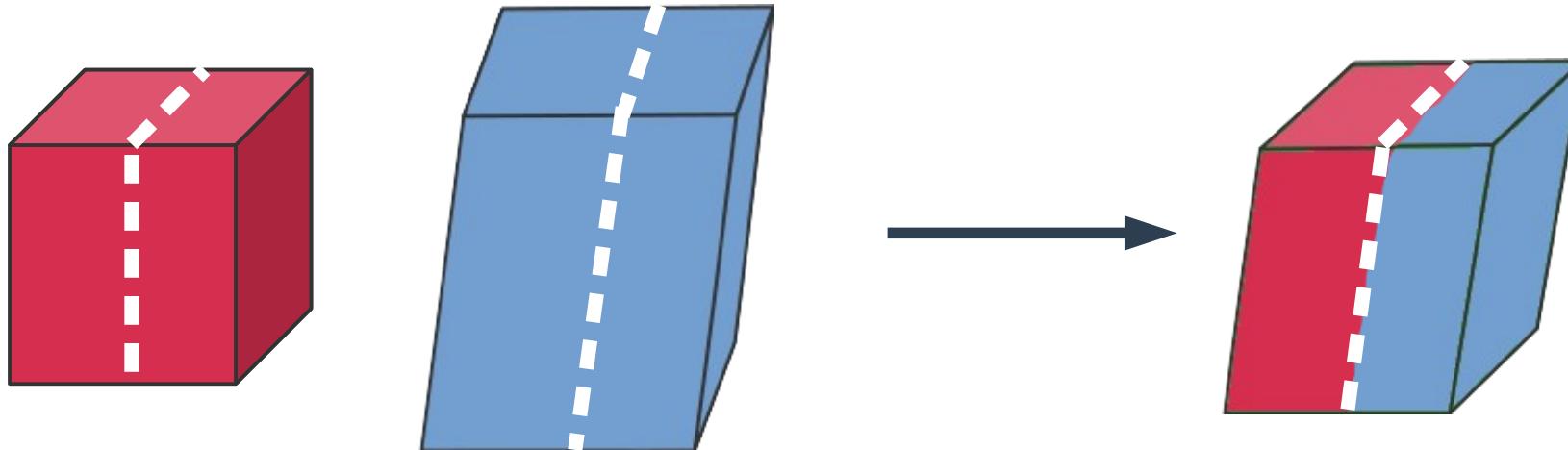
Motivation:

Development of
Tight-binding
Approximation
e**N**hanced
Global
Optimization

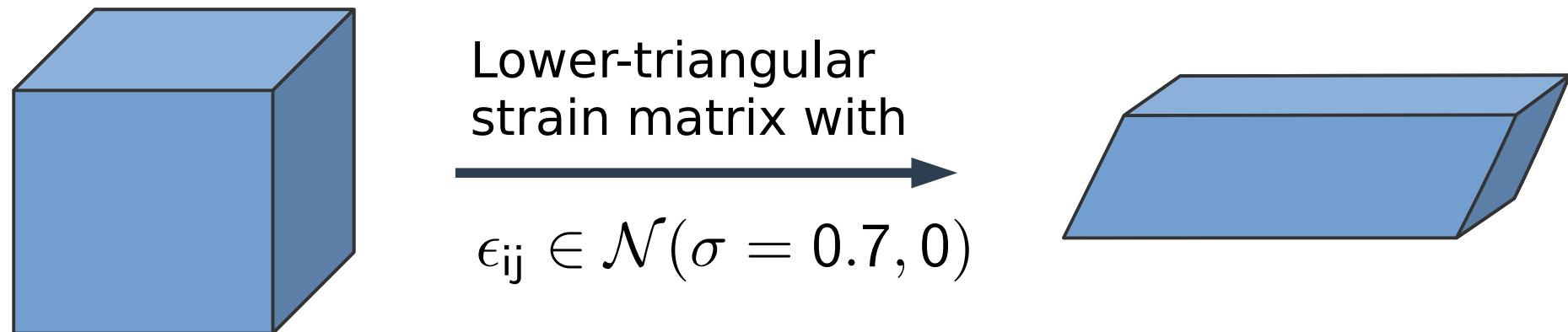


ASE-GA crystal structure prediction

* “*cut-and-splice*” crossover:

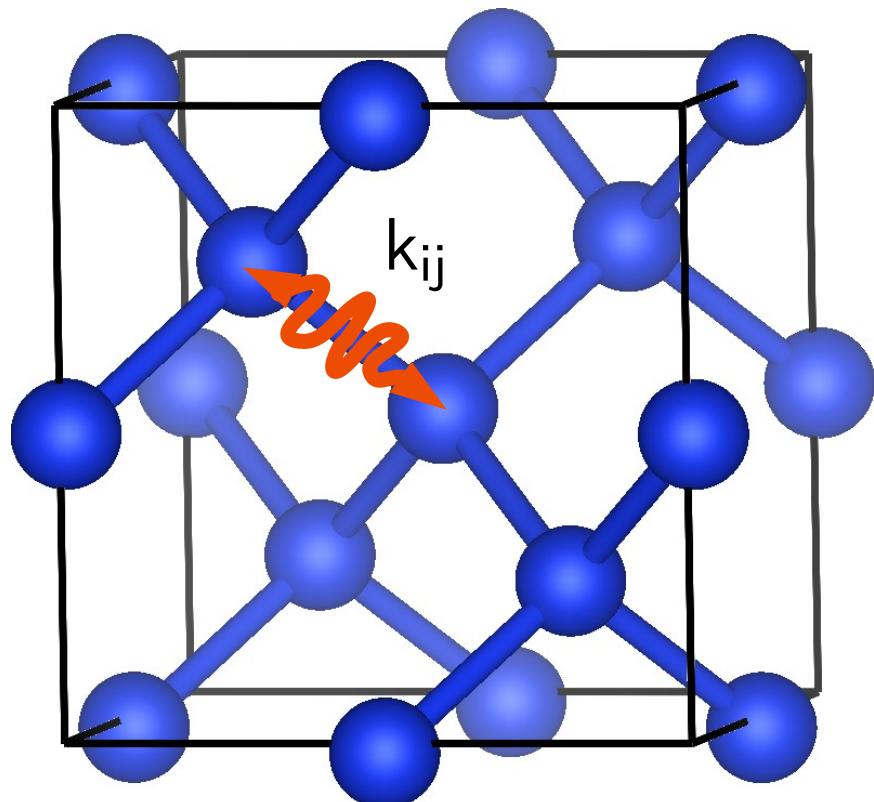


* “*strain*” mutation:



ASE-GA crystal structure prediction

* “soft” mutation:



Displacement along softest non-zero vibrational mode

Model hessian inspired by bond valence method

ASE-GA crystal structure prediction

GA-operator-related code in ase.ga:

cutandslicepairing.py
element_crossovers.py
element_mutations.py
particle_comparator.py
particle_crossovers.py
particle_mutations.py
slab_operators.py
standard_comparators.py
standardmutations.py
startgenerator.py
utilities.py

New:

bulk_crossovers.py
bulk_mutations.py
bulk_startgenerator.py
bulk_utilities.py
ofp_comparator.py

Future work (I):

Generalization
⇒ less redundancy
⇒ also 1D and 2D-vc-opt

ASE-GA crystal structure prediction

GA-operator-related code in `ase.ga`:

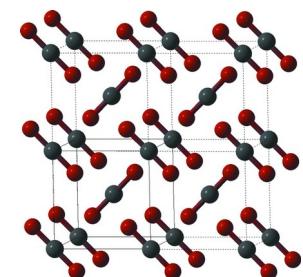
- `cutandslicepairing.py`
- `element_crossovers.py`
- `element_mutations.py`
- `particle_comparator.py`
- `particle_crossovers.py`
- `particle_mutations.py`
- `slab_operators.py`
- `standard_comparators.py`
- `standardmutations.py`
- `startgenerator.py`
- `utilities.py`

New:

- `bulk_crossovers.py`
- `bulk_mutations.py`
- `bulk_startgenerator.py`
- `bulk_utilities.py`
- `ofp_comparator.py`

Future work (II):

Tutorial for global optimization of molecular crystals

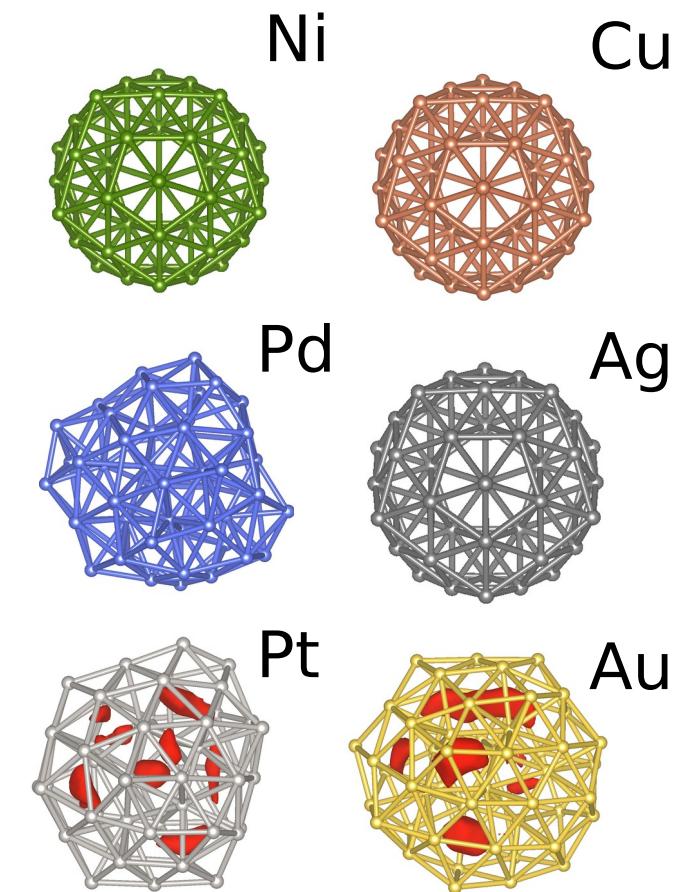
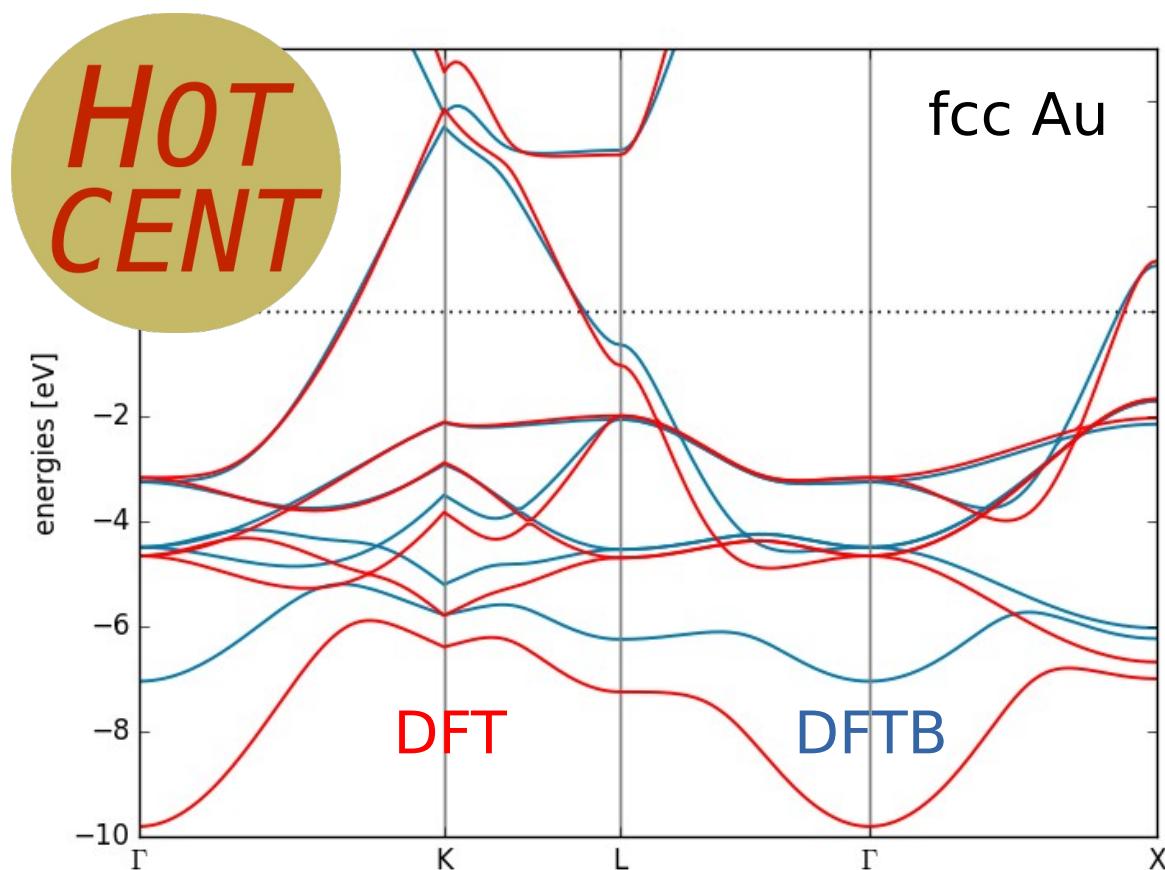


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Calculators: DFTB+ and QuantumEspresso

Motivation: calculating and plotting the band structure
e.g. for tight-binding parameter optimization



Calculators: DFTB+ and QuantumEspresso

Changes in both calculators:

1. “Advanced” k-point input
2. Parsing the eigenvalues, Fermi energy
& irreducible BZ k-points

Calculators: DFTB+ and QuantumEspresso

```
calc = Espresso/DftbPlus( . . . )
atoms.set_calculator(calc)
atoms.get_potential_energy()
efermi = calc.get_fermi_level()

<change settings for non-self-consistent run>
calc.set(kpts={'path':'GKLGX', npoints=100})
calc.calculate(atoms)

bs = calc.band_structure()
bs.reference = fermi_energy
bs.plot()
```

Calculators: DFTB+ and QuantumEspresso

Future work:

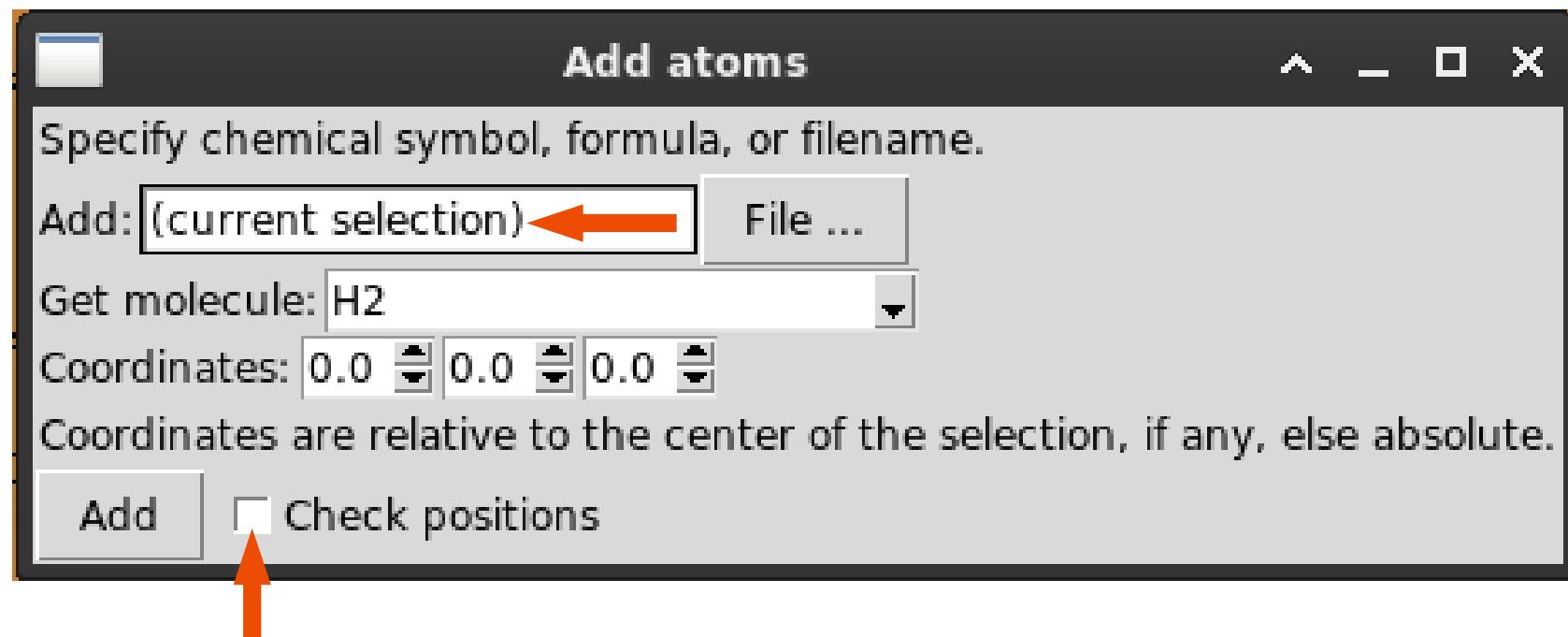
- More extensive and easy-to-run test suite
- Deal with changes in newest DFTB+ release (19.2)
(Fermi energy, eigenvalues, ... no longer in results.tag)

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Miscellaneous

Copy-pasting back in the GUI



Miscellaneous



Tutorial

**DFTB with ASE:
From Zero to Hero**

Friday at 9:00

FT4011

Overview

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Conclusions

1. Functionality for crystal structure prediction (`ase.ga`)
2. Calculators: DFTB+ and QuantumEspresso



gitlab.com/mvdb

Acknowledgements



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Bjørk
Hammer
group



Ask Hjorth Larsen