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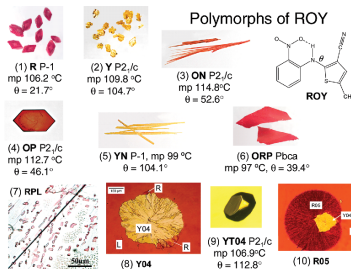


Improving the Description of Molecular Crystals via Multimer Embedding

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- Solids composed of molecules
- Non-covalent interactions
- Can exhibit polymorphism
- Small energy differences
- Properties differ between polymorphs

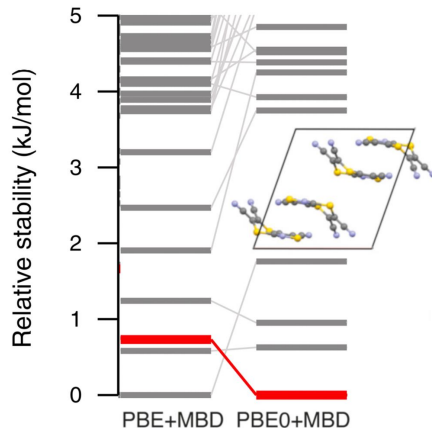


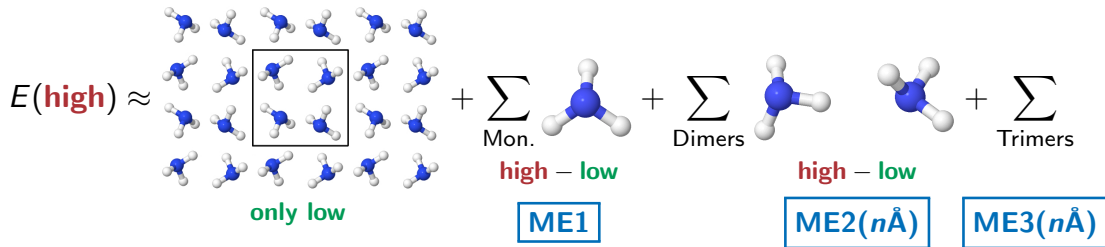
L. Yu, *Acc. Chem. Res.* 43, 1257 (2010).

Polymorphs can exhibit different

- Solubilities
- Kinetic stabilities
- Melting points
- THz spectra
- NMR chemical shifts
- Elastic constants
- Heat capacities
- Conductivities
- Colors

- Small lattice energy differences between polymorphs require accurate methods
- Vibrational free energies often needed for correct polymorph stabilities
- Hybrid DFT desirable
- Converged periodic calculations using hybrid functionals are often computationally too expensive





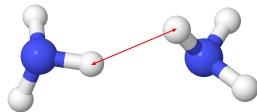
$$E_{\text{per}}^{\text{high}} \approx E_{\text{per}}^{\text{low}} + \sum_i n_i \Delta E_i + \sum_{i>j} \frac{n_{ij}}{2} \Delta E_{ij}^{\text{int}} + \sum_{i>j>k} \frac{n_{ijk}}{3} \Delta E_{ijk}^{\text{int}}$$

$$E_{\text{latt}}^{\text{high}} = \frac{E_{\text{per}}^{\text{high}}}{Z} - E_{\text{mon}}^{\text{high}}$$

$$\Delta E = E_{\text{high}} - E_{\text{low}} \quad E_{ij}^{\text{int}} = E_{ij} - E_i - E_j$$

$$E_{ijk}^{\text{int}} = E_{ijk} - E_{ij}^{\text{int}} - E_{ik}^{\text{int}} - E_{kj}^{\text{int}} - E_i - E_j - E_k$$

Multimers within cutoff distance:

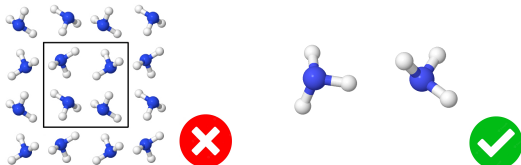


A. D. Boese, J. Sauer, *Cryst. Growth Des.* 17, 1636 (2017).

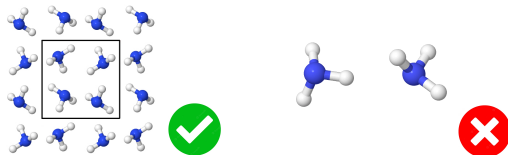
J. Hoja, A. List, A. D. Boese, submitted, arXiv:2209.02687 (2022).

A. List, J. Hoja, A. D. Boese, in preparation.

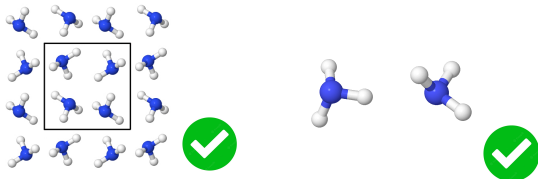
Gaussian basis set:



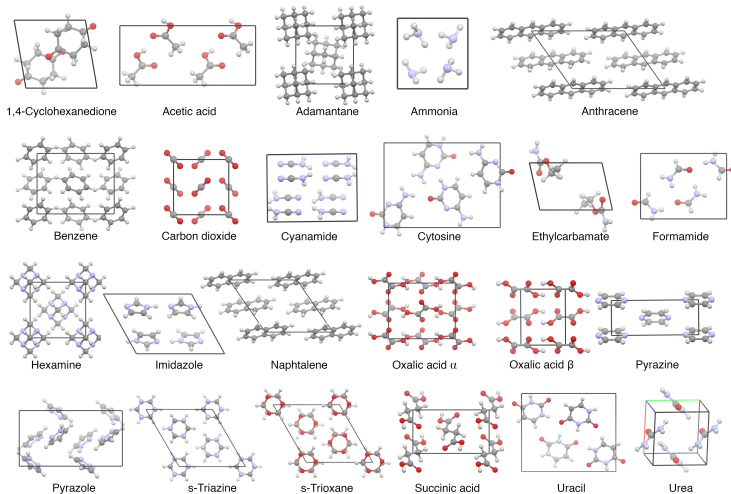
Plane waves:



Numeric atom-centered basis set:



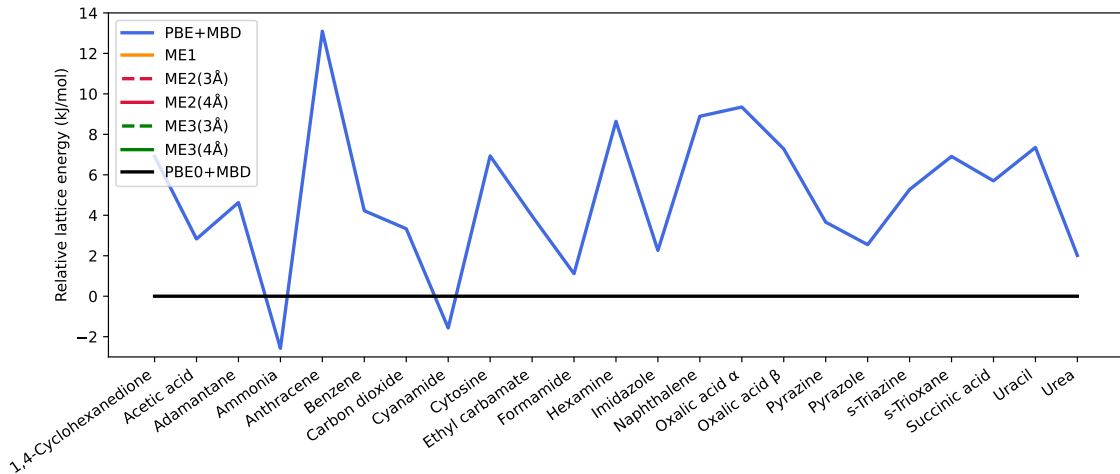
FHI-aims
The ab initio materials
simulation package



- High: PBE0+MBD
- Low: PBE+MBD
- Light basis set

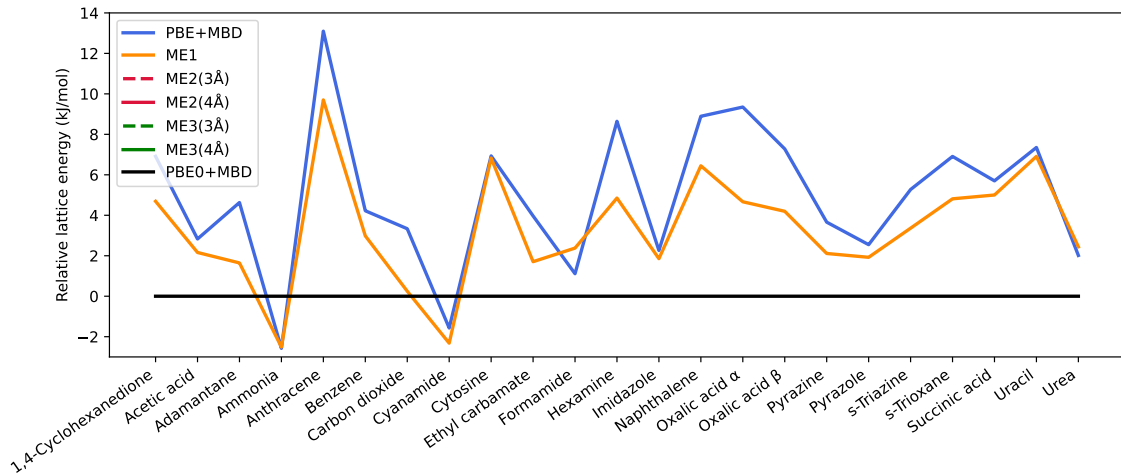
A.M. Reilly and A. Tkatchenko, *J. Chem. Phys.* 139, 024705 (2013).

Calculated on top of PBE0+MBD-optimized structures (light settings)



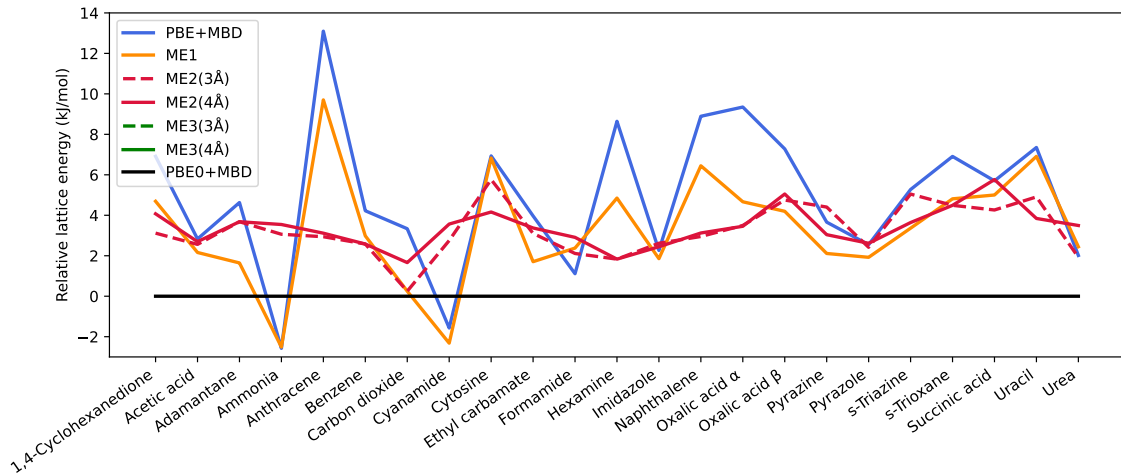
J. Hoja, A. List, A. D. Boese, submitted, arXiv:2209.02687 (2022).

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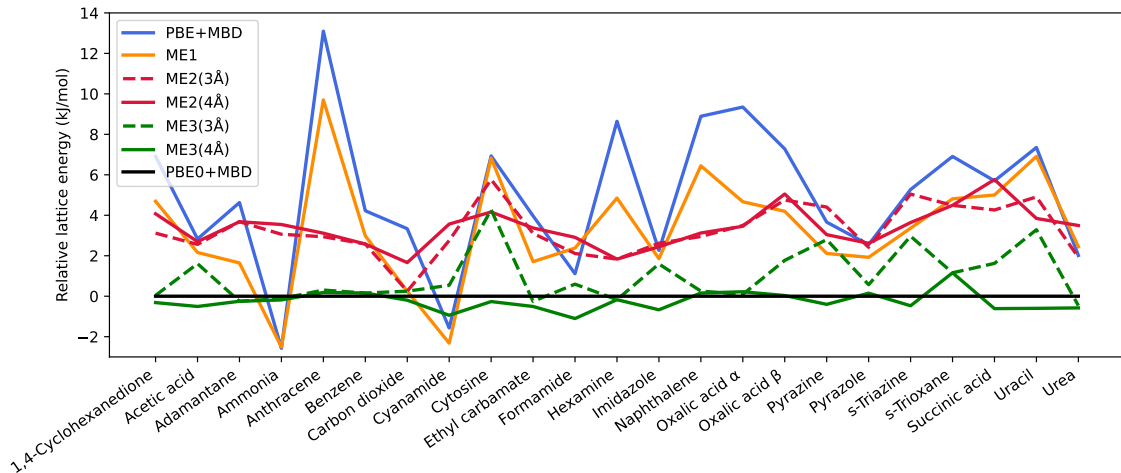
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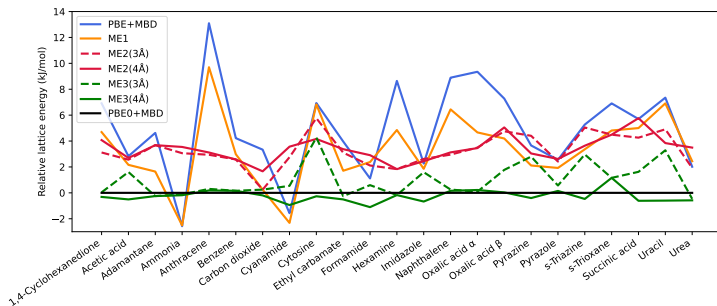
Calculated on top of PBE0+MBD-optimized structures (light settings)



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Errors of lattice energies in kJ/mol:

Method	ME	MAE	MAX
PBE+MBD	4.9	5.3	13.1
ME1	3.3	3.7	9.7
ME2(3Å)	3.3	3.3	5.8
ME2(4Å)	3.4	3.4	5.8
ME3(3Å)	1.0	1.1	4.3
ME3(4Å)	-0.2	0.4	1.2



Trimers needed for convergence to 1 kJ/mol

Embedding forces on atom a :

$$\mathbf{f}_{\text{per}}^{\text{high}}(a) \approx \mathbf{f}_{\text{per}}^{\text{low}}(a) + \Delta\mathbf{f}_i(a) + \sum_j \Delta\mathbf{f}_{ij}^{\text{int}}(a) + \sum_{j>k} \Delta\mathbf{f}_{ijk}^{\text{int}}(a)$$

$$\mathbf{f}_{ij}^{\text{int}}(a) = \mathbf{f}_{ij}(a) - \mathbf{f}_i(a)$$

$$\mathbf{f}_{ijk}^{\text{int}}(a) = \mathbf{f}_{ijk}(a) - \mathbf{f}_{ij}^{\text{int}}(a) - \mathbf{f}_{ik}^{\text{int}}(a) - \mathbf{f}_i(a)$$

Errors compared to PBE0+MBD results calculated at PBE+MBD-optimized structures:

Method	Forces (eV/Å)	
	MAE	MAX
PBE+MBD	0.226	1.422
ME1	0.023	0.216
ME2(3Å)	0.008	0.120
ME2(4Å)	0.006	0.055
ME3(3Å)	0.006	0.107
ME3(4Å)	0.004	0.043

J. Hoja, A. List, A. D. Boese, submitted, arXiv:2209.02687 (2022).

A. List, J. Hoja, A. D. Boese, in preparation.

Embedding stress tensor components:

$$\begin{aligned} \sigma_{pq}^{\text{high}} \approx \sigma_{pq}^{\text{low}} &- \frac{1}{V} \sum_i \sum_a n_i r_{i,p}(a) \Delta f_{i,q}(a) \\ &- \frac{1}{V} \sum_{i>j} \sum_a \frac{n_{ij}}{2} r_{ij,p}(a) \Delta f_{ij,q}^{\text{int}}(a) \\ &- \frac{1}{V} \sum_{i>j>k} \sum_a \frac{n_{ijk}}{3} r_{ijk,p}(a) \Delta f_{ijk,q}^{\text{int}}(a) \end{aligned}$$

Errors compared to PBE0+MBD results calculated at PBE+MBD-optimized structures:

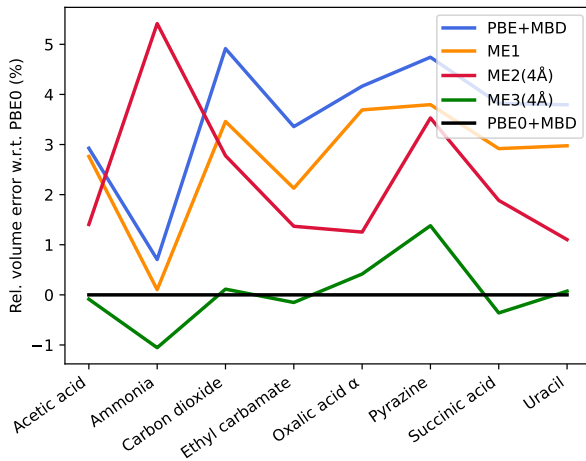
Method	Stress (eV/Å ³)	
	MAE	MAX
PBE+MBD	0.0190	0.0496
ME1	0.0009	0.0034
ME2(3Å)	0.0007	0.0039
ME2(4Å)	0.0007	0.0039
ME3(3Å)	0.0003	0.0032
ME3(4Å)	0.0002	0.0032

J. Hoja, A. List, A. D. Boese, submitted, arXiv:2209.02687 (2022).

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Errors of cell volumes in % compared to PBE0+MBD results (reduced X23 set):

Method	ME	MAE	MAX
PBE+MBD	3.55	3.55	4.91
ME1	2.73	2.73	3.80
ME2(4Å)	2.34	2.34	5.41
ME3(4Å)	0.04	0.45	1.38



J. Hoja, A. List, A. D. Boese, submitted, arXiv:2209.02687 (2022).

A. List, J. Hoja, A. D. Boese, in preparation.

- Finite displacements within sufficiently large super cells
- Number of multimers to calculate increases significantly

Multimers to calculate for Ammonia:

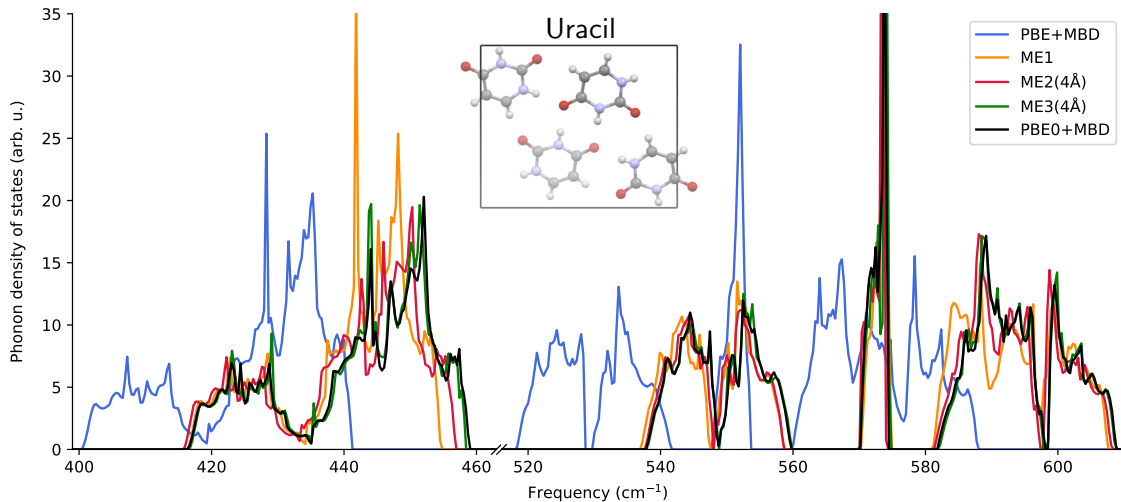
Multimer	SP	VIB
Monomers	1	8
Dimers (3Å)	2	96
Dimers (4Å)	3	144
Trimers (3Å)	7	192
Trimers (4Å)	11	480

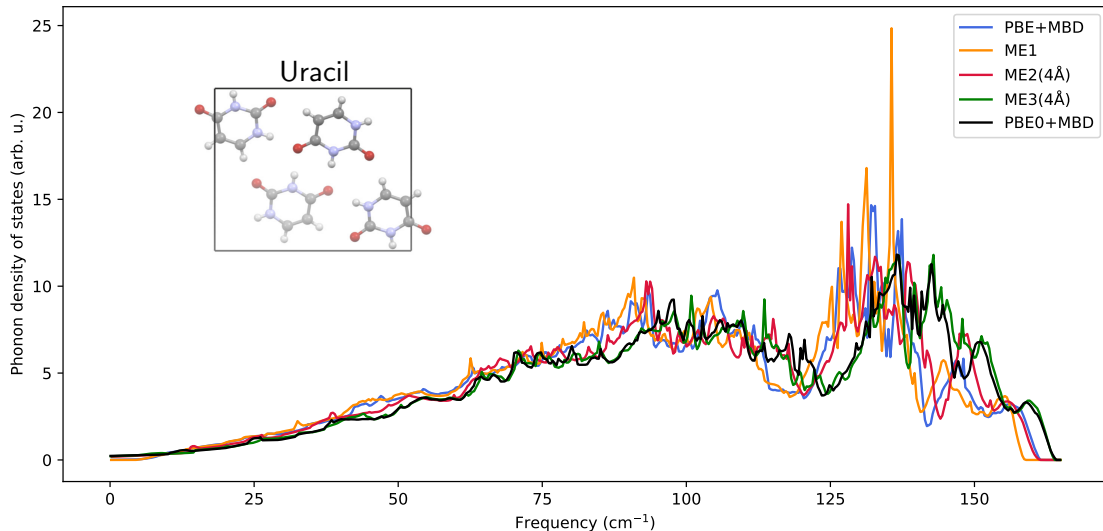
Errors compared to PBE0+MBD results:

Quantity	Method	ME	MAE
$\nu(\Gamma)$ (cm^{-1})	PBE+MBD	-48.2	49.0
	ME1	-1.2	4.2
	ME2(3Å)	-1.1	2.5
	ME2(4Å)	-1.0	2.7
F_{vib} (kJ/mol)	PBE+MBD	-10.9	10.9
	ME1	-1.2	1.2
	ME2(3Å)	-0.8	0.8
	ME2(4Å)	-0.8	0.8

Harmonic vibrational free energies (per molecule) at 300 K

J. Hoja, A. List, A. D. Boese, submitted, arXiv:2209.02687 (2022).





Relative timings (light basis set):

Method	Ammonia	Succinic acid
PBE+MBD	1.0	2.7
ME1	1.0	2.9
ME2(3Å)	1.2	7.9
ME2(4Å)	1.2	9.6
ME3(3Å)	2.0	16.4
ME3(4Å)	2.6	34.1
PBE0+MBD	3.4	15.3

calculated on 4 cores

Relative timings (tight basis set):

Method	Ammonia	Succinic acid
PBE+MBD	1.0	2.7
ME1	1.0	3.6
ME2(3Å)	1.2	24.8
ME2(4Å)	1.3	33.2
ME3(3Å)	2.8	64.1
ME3(4Å)	3.8	151.3
PBE0+MBD	93.3	258.6

calculated on 24 cores

- Python package available at <https://gitlab.com/membed/membed>
- Dependencies: ASE, Phonopy, Spglib
- Input files: geometry.in, membed.in
- Results of all individual calculations saved to an ASE database
- single points, lattice relaxations, phonon calculations

```
multimer_order      3
multimer_cutoff     4
atoms_per_molecule 3
identify_fragments  automatic
calculation_type    opt
f_max               0.005
symmetrize          True
calculator           aims

xc                  PBE          lo
xc                  PBE0         hi
many_body_dispersion ""
species_dir         /species_dir/light
aims_command        "srun aims.x"
k_grid              "4 4 4"         per
many_body_dispersion k_grid=5:5:5  per
```

- Multimer embedding scheme to approximate high-level methods
- Lattice energies can be converged to 1 kJ/mol with trimers
- Trimers needed for cell volumes within 1 % of PBE0 cell volumes
- Harmonic vib. free energies well described at monomer/dimer level



MSCA-IF VibMolCryst

Outlook:

Lattice optimizations/phonons with trimer interactions
Anharmonic vibrational effects
Multimer selection

Embedding code: MEmbed

<https://gitlab.com/membed/membed>



Boese Group at the University of Graz

Thank you for your attention!