

ICM

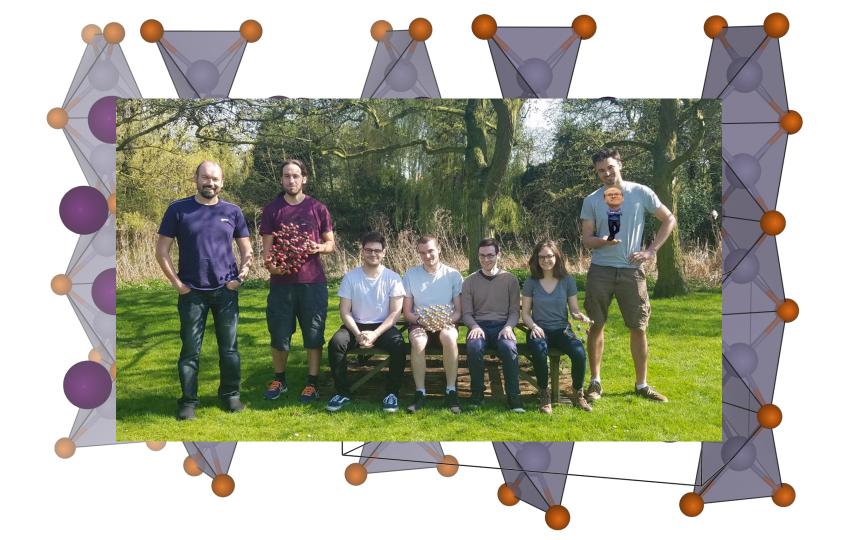




{Databases & Crystal Structure Prediction} in the Andrew Morris Group

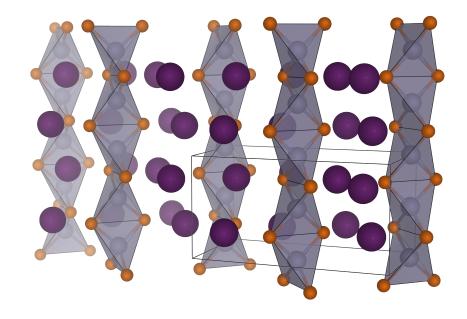
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http://ml-evs.science



What flavour of crystal structure prediction?

- Ab initio random structure searching (AIRSS) provides an unbiased way of sampling configuration space.
- Supplemented by data mining, prototyping, evolutionary algorithms, etc.
- Focus on alloying/conversion anodes for batteries, where dense sampling of metastable phases is important.
- A more niche focus: encapsulated nanowires
- ~100 DFT relaxations per stoichiometry,
 ~100 stoichiometries per phase diagram.



Spec sheet

- ~850k geometry optimisations targeted towards alloying anodes for Li and beyond-Li batteries
- Native CASTEP 18, 17, 16, 9, 8... 7... 6
- Messy, nonuniform historical data
- Big phase diagrams: some as large as 100k calculations.
- Should fit CLI workflow of group, and be usable for new members
- Improved data mining capabilities
- Selectively open data, but either open to all or closed; no auth

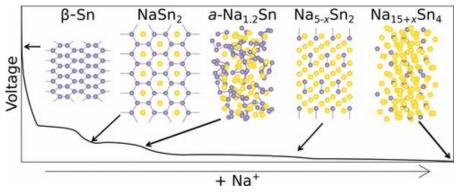
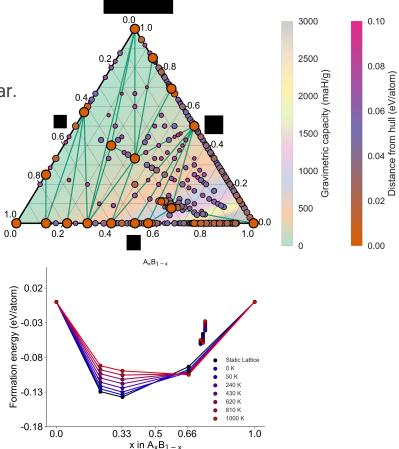


Figure 1: Na-Sn voltage curve adapted from Ref [1].

[1] P. K. Allan, et al. J. Am. Chem. Soc. 138 (2016)

matador

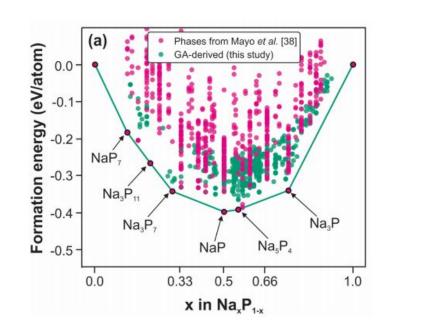
- Open-source, Python 3.6, bus factor increasing by 1 per year.
- MongoDB backend.
- Fundamental sub-commands:
 - **import** data to local database, on-the-fly
 - query with powerful macros and DFT parameter matching
 - hull construct phase diagrams and make pretty plots,
 + ability to track changes over time (as searches progress)
 - swap leveraging OQMD, materials project and ICSD for structural prototypes
 - **run** high-throughput relaxations, electronic, vibrational calculations with CASTEP or QE.
- Somewhere between fully-automated luxury DFT and handle-turning monkey.

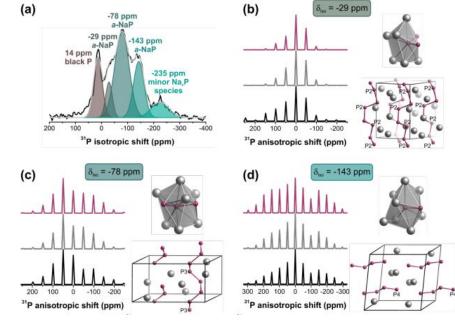


Python API for "Science"

- Structure analysis and descriptors:
 - Voronoi for local structure, others planned (SOAP, BP symmetry functions)
 - Crystal graphs for global structure
- Screening/fitting against experimental NMR, PDF, XRD, electrochemistry
- WIP: Finite-temperature phase diagrams and non-equilibrium electrochemistry
- WIP: Training of interatomic potentials
 - Overall ~850e3 geometry optimisations => ~50e6 "snapshots" of forces and energies
 - At least 10,000 per phase diagram; can learn on-the-fly
- Genetic algorithm running from database: separate open source code called ilustrado.

Case study: P helices in disordered NaP





[2] (De)Sodiation via Helical Phosphorus Intermediates in High Capacity Anodes for Sodium-ion Batteries. L. E. Marbella, M. L. Evans, M. Groh, J. Nelson, A. J. Morris, C. P. Grey, JACS (In print) (2018)

OPTiMaDe in matador 1.0

- Query selected OPTiMaDe databases from Python, using matador's own query structure.
 - Will implement two-way conversion between filter languages.
 - Will typically only query relaxations, preferably with quality tolerance.
- Host (at least) our own published data with OPTiMaDe REST-API (both internally and externally).
- **Provenance tracking**: we re-relax an OQMD prototype based on an ICSD phase?
 - Currently we would track both under provenance: {oqmd_id: 1234, icsd_id: 5678},
 - List of DOIs enough? How many allowed per entry?
- **Discoverability**: we have many results on few systems
 - Access via e.g. http://matador.science/optimade or http://optimade.org/matador, should lead to same interface (c.f. Skyscanner comment)
 - Explorability? (e.g. wow, where else can I fly with this company?)
- **Calculation quality/type**: phase stability and interdependence of structures & calculations.
- Fuzzier searches: discussed in part yesterday,
 - \circ ~ elements HAVE ALL ['[III], '[V]'], n_elements IS 2 ~
 - Give me all A_2BO_3 ?

Thanks for listening!



Engineering and Physical Sciences Research Council



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matador's (ab)users In the AJM group

Computing:

- Tier-1: ARCHER
- Tier-2: CSD3, HPC Midlands +
- CFN Cluster Brookhaven National Labs