

# Hackathon & Tutorials

Materials Available:

<https://github.com/openchemistry/avogadro-ugm2018>

# Current & Future APIs

- RPC: Your program => "visualize this file"
- C++ Plugins
- Python
  - File Formats
  - Input Generators
  - Workflows
  - Force Fields / Energies
  - Charge Models

# Plugin Directory / Ecosystem

- Register GitHub repo
- Search for plugin (author?)
  - "Splash" icon for plugin?
- Reviews / feedback on repo
  - How do we help file bugs on plugins?

# Generator Ideas

## Chemistry:

GAMESS  
Gaussian  
Mopac  
Orca  
Q-Chem  
NWChem

## Needed:

Turbomole  
Psi4  
???

## Solid-state:

VASP  
~~CP2K~~  
Abinit  
Quantum Espresso  
Crystal  
Wien2k  
(etc.)

## Molecular Dynamics:

Amber  
~~OpenMM~~  
Gromacs  
Tinker  
~~LAMMPS~~

- Add job types (TS, Saddle, IRC, NMR, TDDFT, *etc.*)
- Update methods ( $\omega$ B97X-D, SCAN, *etc.*)
- Update basis sets (def2-X, pc-X, *etc.*)
- Add solvation
- Syntax highlighting

# Workflow Ideas

## Avogadro

- get / modify selection (e.g., enlarge by  $x \text{ \AA}$ )
- create enantiomer
- “Random alloy” - Replace  $x\%$  atom X  $\rightarrow$  atom Y

## Scikit-nano <https://docs.scikit-nano.org>

- MWNT
- Graphene or TMDC multilayers?

## Pymatgen <https://github.com/materialsproject/pymatgen>

- create a slab
- nanoparticle with Wulff

## STK <https://lukasturcani.github.io/stk/docs/build/html/>

- polymers
- cages
- COF
  - hexagonal, honeycomb, kagome, square

# Workflow Ideas

**RDKit** <http://www.rdkit.org/docs/GettingStartedInPython.html>

- get ETKDG conformer

## ASE

- Generate MoS2 <https://wiki.fysik.dtu.dk/ase/ase/build/surface.html#ase.build.mx2>
- Generate TIP water box: [https://wiki.fysik.dtu.dk/ase/tutorials/tipnp\\_equil/tipnp\\_equil.html](https://wiki.fysik.dtu.dk/ase/tutorials/tipnp_equil/tipnp_equil.html)

**GenIce** - <https://github.com/vitroid/GenIce/>

- Generate ice unit cell

**PackMol** <https://github.com/choderalab/openmoltools/blob/master/openmoltools/packmol.py>

- Pack current molecule <http://timvdm.blogspot.com/2010/03/avogadros-packmol-plugin.html>

## Glycam(?)

- Sugar / oligosaccharide builder?

# Avogadro Demos

# Avogadro Demo: Small Molecule Building

- Draw Tool: Click & Drag
- Click on bond: Single, Double, Triple
- Change Element, Click on atom: Alchemy
- Auto-Adjust Hydrogens
- Manually Add H after drawing
- **Keyboard Shortcuts:**
  - Type element symbol to change
  - Type 1, 2, 3, 4 to change bond order
  - Control + 1, 2, 3 to change tools



# Avogadro Demo: Larger Molecules

- Copy/paste & Undo
- Insert Fragments
  - Select h-atom to "grow" fragment (v 1.1)
- Insert SMILES
  - Useful for polymers
- "Sculpting" using Auto-Optimization
- Adjusting angles, bond lengths, dihedrals
  - Bond-Centric Manipulate Tool
  - View  $\Rightarrow$  Properties ...

# Avogadro Demo: Peptides

- Custom Sequences
- Helix, Sheets, Custom Conformations
- Multiple Chains
- Visualization methods

# Avogadro Demo: Building Crystals

- New Interface in v1.1
- Still in development: feedback welcome
  - Build slabs and nanoparticles (soon)
- Edit cell parameters
- Build supercells (simple slabs)
- Edit Fractional Coordinates (v1.1)
- Detect / Set Space Group (v1.1)

# Avogadro Demo: Visualization Modes

- Standard Representations
  - Balls & Sticks, Labels, etc.
- Forces
- Hydrogen-Bonding
- Color Modes

# Avogadro Demo: Force Fields / Conformers

- Auto-Optimize Tool
- Extension / Menu Item
- Setting Constraints
  - Fixed / Frozen Atoms
  - Ignored Atoms
  - General Constraints (bonds, angles, ...)
- Finding & Generating Conformers

# Avogadro Demo: More Extensions

- Spectra Viewer (IR, UV/Vis, NMR)
  - Import experimental data
- Vibrations & Orbitals
- Molecular Surfaces
- External Packages
  - Gaussian, GAMESS, Q-Chem...
  - Abinit
  - Packmol, XtalOpt (3<sup>rd</sup> party)