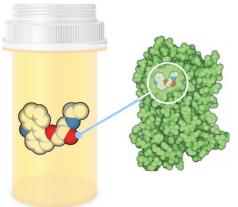
### Jessica Maat

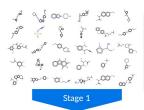
## University of California, Irvine

**Data-Driven Force Field Development for Drug Discovery** 



Openforcefield.org





#### Parse inputs & parameter usage

Parse input molecule & expand tautomeric and isomeric state.

Determine the bond, angle and orsion parameters from .offxml that the input molecule utilizes.



#### Molecule Selection

Using randomization, select a nolecule from each cluster that is representative of the chemistry.

Generate QM data in QCA





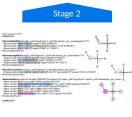






#### **Starting Datasets**

- 1. Roche set
- 2. Coverage set 3. eMolecules Discrepancy
- 4. Pfizer Discrepancy set
- 5. Bayer set



#### **Determine** fingerprints & Cluster

Calculate the MACCS key for each

Cluster the molecule based off the MACCS key using DBSCAN.











# Jessica Maat University of California, Irvine

In my research I deal with large quantum chemistry training datasets for force field optimization, the following topics would benefit my research:

- GPU Programming
- Big data processing
- Machine learning
- Workflow management

