

# References

- Chodera, J. D. A Simple Method for Automated Equilibration Detection in Molecular Simulations. *Journal of Chemical Theory and Computation* 2016, 12 (4), 1799–1805. <https://doi.org/10.1021/acs.jctc.5b00784>.

# Some other software

- MDTraj: <http://mdtraj.org/1.9.3/index.html>
- ProDy: [http://prody.csb.pitt.edu/tutorials/trajectory\\_analysis/trajectory.html](http://prody.csb.pitt.edu/tutorials/trajectory_analysis/trajectory.html)