

# AMBER in Utah Wiki

This is the beginnings of the AMBER @ Utah and Cheatham lab Wiki pages.

The intent is to provide a place for the group and/or local AMBER users to post scripts, information, and helpful hints for using AMBER and performing our research. Note that this information is (by default) accessible to all, however it can be restricted in various ways...

- [The AMBER home page \(www.ambermd.org\)](http://www.ambermd.org)
- [The AMBER tutorials](#)
- [Amber 11 AmberTools manual](#)
- [Amber 11 manual](#)

[current projects](#)

[ptraj scripts](#)

[tec3 to do list](#)

## Recently Updated

[Amber Installs](#)

Mar 04, 2016 • updated by Dan Roe • [view change](#)

[Amber Benchmarks](#)

Oct 15, 2015 • updated by Dan Roe • [view change](#)

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Oct 14, 2015 • updated by Rodrigo Galindo-Murillo • [view change](#)

[combined-clustering.in](#)

Jun 08, 2015 • created by Rodrigo Galindo-Murillo

[ptraj scripts](#)

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[DNA force field including e/z OL1 - chi OL4](#)

Apr 11, 2015 • updated by Rodrigo Galindo-Murillo • [view change](#)

[CPPTRAJ one-liners](#)

Apr 01, 2015 • updated by Rodrigo Galindo-Murillo • [view change](#)

[Artículos cómo](#)

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[Julien's page](#)

Sep 11, 2013 • updated by Anonymous • [view change](#)

[Glide Docking Tutorial: Docking on an RNA receptor](#)

Apr 12, 2013 • updated by Hamed Sadat Hayatshahi • [view change](#)

[ligand-charge.png](#)

Mar 21, 2013 • attached by Hamed Sadat Hayatshahi

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[atom-labels.png](#)

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Feb 10, 2013 • updated by Anonymous • [view change](#)

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