

# 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](#)

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- [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](#)  
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- [Julien's page](#)  
Sep 11, 2013 • updated by Anonymous • [view change](#)
- [Glide Docking Tutorial: Docking on an RNA receptor](#)  
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- [ligand-charge.png](#)  
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- [labels.png](#)  
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