

**folding@home EXTRA CREDIT:**

You may be interested to check out the “folding@home” website: <http://folding.stanford.edu/>. This is a distributed computing project in which people throughout the world download and run protein folding software to form a large supercomputer. You can help this project by running some of the software yourself. To do so, go to <http://folding.stanford.edu/English/Download> and download the program. When you first run the program, it will ask you to create a username of your choosing and join a team by typing in the team number. Pick any user name and then join our team, “Caltech Bi1 2009,” by typing in the team number **164672**. The program will run in the background. To get extra credit, you must finish at least one WU (working unit) and contribute it to the Bi1 2009 team – more would be great and may get you a few extra points! To get credit, you must also email your folding@home username to Damien (soghoian@caltech.edu).

**folding@home extra credit will be calculated at the end of the term.**

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