



AutoIMD Controls



File

Settings

Help

1. Type in the atom selections for imd (top molecule):

indmolten: protein and resid 78 to 85

indfixed: within 8 of indmolten

2. Choose where and how to run the simulation:

Local ☐ using / 1 processors

3. Submit your job:

Submit

Connect

Pause

Finish

Discard

status: Waiting for user input