

Name: \_\_\_\_\_



## Worksheet 1: Interactive Molecular Graphics

course website -- <http://kinemage.biochem.duke.edu/teaching/bch258/>

MolProbit website -- [http://molprobit.biochem.duke.edu/...](http://molprobit.biochem.duke.edu/)

software website -- [http://kinemage.biochem.duke.edu/software/...](http://kinemage.biochem.duke.edu/software/)

### Your Contact Info

Please give us your full name:

your class, department and/or program:

your email address:

Do you have a laptop with wireless internet, that you could bring in for several of the class sessions?

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### Reading Assignment

Read (or review) chapter 1 in Branden & Tooze "Introduction to Protein Structure"

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### Graphics Assignment -

#### to be done before the next class, and this worksheet handed in then

The course will use kinemages, viewed with the KiNG display program, both for homework exercises and also as the medium for individual projects. The QuickStartKing.pdf file summarizes the basic operations. You will start out (below) directly on-line by uploading a kinemage file onto the MolProbit web site and viewing it there in KiNG (thus independent of hardware, OS, or installation - except for a browser, and Java). Later it will often be necessary to work off-line on your own computer, so the second half of this assignment leads you into trying that method.

### Viewing kinemages on-line

From the course web site (see above) download the files 3Dlit2a-exercise-KiNG.kin and Demo5\_4a-KiNG.kin to your computer. Go to the MolProbit site (see above, or navigate from the main kinemage site). If it says you need a current Java, follow its instructions. In the file upload section of the MolProbit main page, set file type to kinemage, browse to find the demo file, and upload it. Choose the "View in KiNG" option, which will start KiNG with this kinemage in the graphics window. Click the "Show text" button at lower right and go through the text following along in the graphics.

### Doing 3Dlit2a-exercise-KiNG.kin on-line

(see 3Dlit2a-exercise-KiNG.pdf) Please affirm (i.e. check off) that you did try:

Connect-the-atoms exercise \_\_\_\_\_ and Find-the-backbone-Hbonds exercise \_\_\_\_\_

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### Viewing Demo5\_4a-KiNG.kin on-line

Be sure to do kins 1, 2, 3, & 6; (4 & 5 are fun but optional).

In kinemage 3, which 6 sidechains make direct H-bonds to the biotin? \_\_\_\_\_  
\_\_\_\_\_

In kinemage 6, what is the distance between the N-cap sidechain O (red ball) and the backbone N(skyblue ball)? \_\_\_\_\_ Å

On the tools pulldown menu, what is the one choice under "Structural biology"?  
\_\_\_\_\_

### Installing KiNG stand-alone

Go to the software web page (see url above). Find and download the appropriate KiNG install package for your computer/operating system. Instructions are on that web page and in the document called QuickStartKiNG.pdf. Make a working folder for these class exercises (on MSWindows, it's advisable to put KiNG in that same folder). Launch KiNG and open the Demo5\_4a-KiNG.kin kinemage from the file menu.

Which operating system are you using (MacOSX, MSWindows, or Linux)? \_\_\_\_\_

Did the installation procedure work? \_\_\_\_\_

Did KiNG come up when launched? \_\_\_\_\_

Did the demo kinemage file open successfully, showing the carboxypeptidase active site in the black-background graphics window? \_\_\_\_\_

On the Tools pulldown menu, how many choices show now under "Structural biology"? \_\_\_\_\_

What is one of the new ones? \_\_\_\_\_

If the above steps did not work, please see one of us for help as soon as feasible.