**STEPS TO SET UP NG-CHM PROJECTS IN ECLIPSE**

1. Install the latest Eclipse IDE and make sure it has GIT support installed.
2. Go to Menu: *File->Import…*
3. Select: *Projects from GIT* (under the GIT folder in list)
4. Select: *Clone URI*
5. Next Screen: Enter URI on next screen in URI field
   1. [*https://github.com/MD-Anderson-Bioinformatics/NG-CHM.git*](https://github.com/MD-Anderson-Bioinformatics/NG-CHM.git)
   2. [*https://github.com/MD-Anderson-Bioinformatics/NG-CHM\_Galaxy.git*](https://github.com/MD-Anderson-Bioinformatics/NG-CHM_Galaxy.git)
   3. [*https://github.com/MD-Anderson-Bioinformatics/NG-CHM\_GUI\_Builder.git*](https://github.com/MD-Anderson-Bioinformatics/NG-CHM_GUI_Builder.git)
   4. [*https://github.com/MD-Anderson-Bioinformatics/pathway-mapper.git*](https://github.com/MD-Anderson-Bioinformatics/pathway-mapper.git)
   5. [*https://github.com/MD-Anderson-Bioinformatics/ScatterPlotPlugin.git*](https://github.com/MD-Anderson-Bioinformatics/ScatterPlotPlugin.git)
6. The system should fill in host: github.com
7. Enter authentication to Git username/password …and press NEXT
8. Next screen: I use the fully selected branch set on this screen …and press NEXT
9. Next screen: I usually accept destination but put the code where you want it …and press NEXT
10. Next screen: Import as General Project …and press NEXT (note: if you have any issues starting a server in Tomcat. You can delete the project, repeat steps and Import as GIT project. Go with General Project first, though)
11. Press FINISH