To get started running bioinformatics software, you need to get an account on ITaP RCAC cluster computers. You may apply for an account by following these directions from ITaP.

After your ITaP/RCAC account has been created, you need to send email to the Bioinformatics Core Director. Describe the work you are doing and indicate you want access to the Bioinformatics queues in order to use the software you need.

Here is a list of the bioinformatics software currently installed.