CGRB Genotyping-by-Sequencing (GBS) Workshop

April 12-May5, 2016

You are receiving this survey because you are signed up to attend the GBS workshop, either as a formal registrant or as a sit-in attendee. If you believe you are receiving this message in error, please contact Adelaide Rhodes ([Adelaide.Rhodes@cgrb.oregonstate.edu](mailto:Adelaide.Rhodes@cgrb.oregonstate.edu)).

1. What is your experience with GBS data? Please select one of the following:  
   \_\_ I have no experience with GBS data.  
   \_\_ I don’t have experience yet, but anticipate working with GBS data.  
   \_\_ I have GBS data in hand, but don’t know what to do with it.  
   \_\_ I have GBS data, and someone else processed it for me.  
   \_\_ I have processed GBS data myself.
2. What is your familiarity with the linux command line?  
   \_\_ I have no experience.  
   \_\_ My only experience is an Intro to Command Line course at the CGRB with:  
    \_\_Adelaide Rhodes  
    \_\_Shawn O’Neil   
    \_\_Matthew Peterson (the Orientation Bootcamp)  
   \_\_I have experience and am comfortable on the command line.
3. Do you have a workshop account at the CGRB (you’ll need one)?  
   \_\_Yes  
   \_\_No
4. What type of laptop do you have? What is its hard drive capacity and RAM?
5. Is there a specific bioinformatic analysis pipeline you’re hoping to explore in the workshop? If yes, what is it?  
   \_\_No, I just want to learn as much as possible about what’s available  
   \_\_Yes, I want to learn about \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_.
6. Is there anything else you’d like us to know?