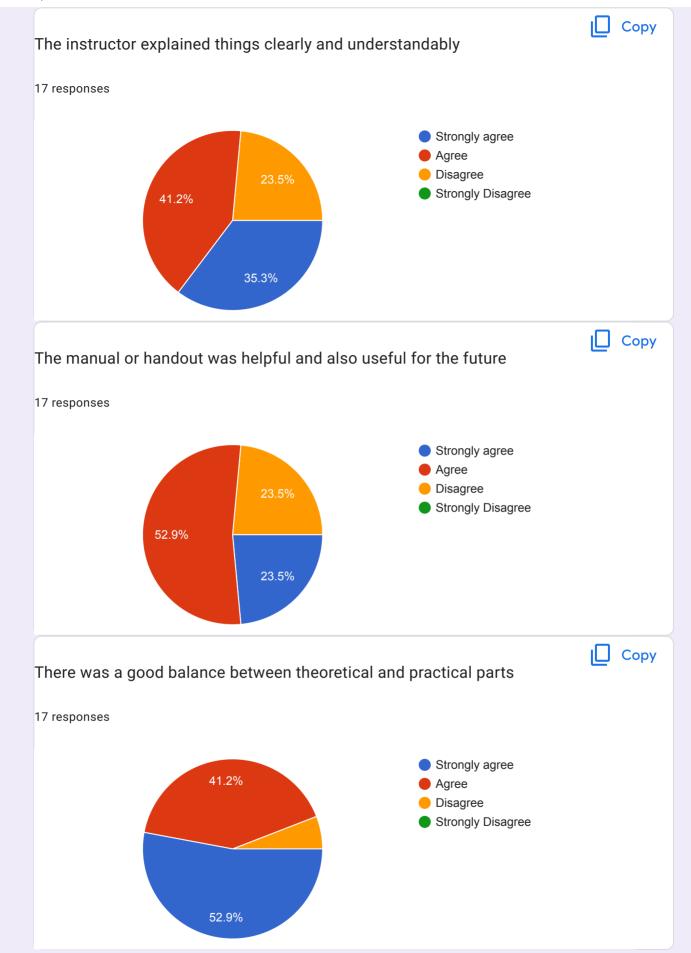


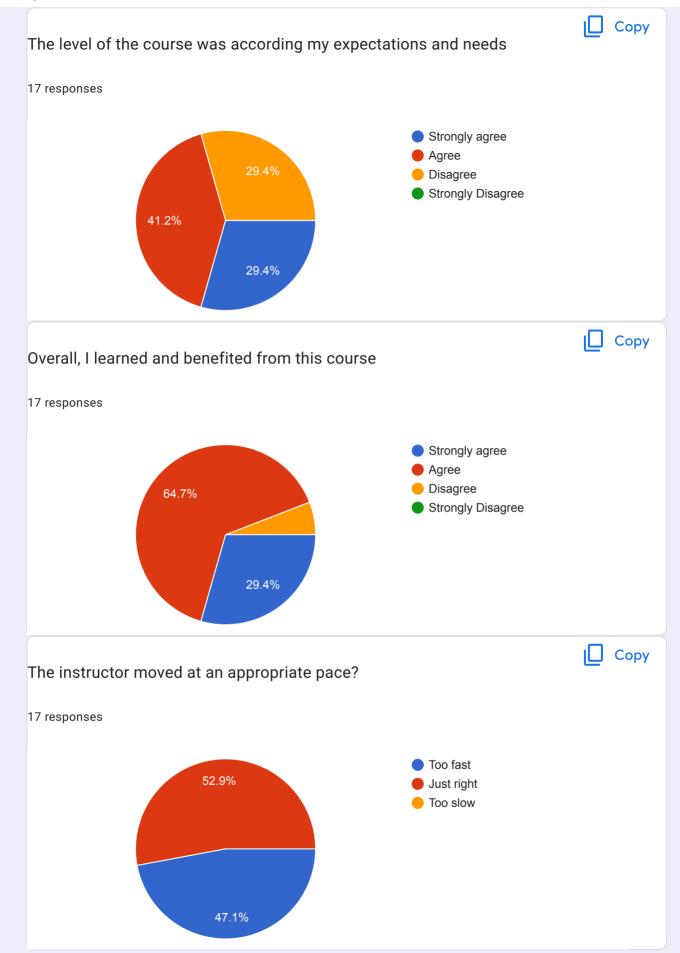


31/10/2022, 16:35 Evaluation of BIO609





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Why did you attend this course? 17 responses
NGS analysis
learn more about linux
to learn about bioinformatics
Too see if I missed something so far using Linux
Get familiar with BASH script
Relevant for Master
We often use Linux-based systems, this course is a good way to acquire familiarity with BASH scripting, so to be able to progress autonomously later.
To learn about running scripts from command line
Usefull for my PhD
To get a good intro to using Linux
To learn bioinformatics tools
It looks interesting.
I wanted to refresh and extend my command line basics
I was interested in the theory behind sequencing, I didnt expect that much bioinformatics
To learn the basics of linux coding
Because it was recommended for BIO610
To learn analysis on genomic data



What is your research field? 7 responses	
Evolutionary ecology	
evolutionary biology	
genomics	
Plant Biology	
Plant sciences	
Evolutionary Genetics	
Physics	
Ecology	
Microbiomics	
Neuroscience	
Evolutionary biology	
Plant pathology	
Plant biology	
Plant Sciences	
-	
Genomics	
Plant Science	



Would you recommend this course to others? Why?
17 responses
Yes, it's relevant
yes, its a good intro to linux for bioinformatics
yes
Sure, it covers the basics really nicely and I learned something even as an "intermediate" Linux user
Yes
Yes, it's a great and simple enough start into bash scripting
Yes, especially to those who have no knowledge whatsoever of Linux systems.
Yes but it's not paced well for a complete beginner
Yes, get a first overview to the LINUX language
Yes, if anyone wants an intro to Linux I would recommend it
yes, this is very important course to analysis sequence data
Maybe.
Only if they need to refresh command line. I do not think I could have followed conceptually if domebody had not explained it to me previously
Very useful
Not sure. I'm waiting for the 2 other days
Yes, because the starting point is basic commands and concepts



Additional comments or suggestions about this course.
17 responses
-
no
none
Provide the slides with the content in advance
A larger set of final exercises could be provided
If more time was allowed for the exercises I'd have learnt more
give the script or at least a list of commands before the lecture
It would be nice, if the Presentation would be awailable as a downloadable file. PDF conversion did not work in my case
nil
The slides on GitHub is difficult to use: hard to switch from pages far from each other, and it sometimes stuck.
The docker was new to me and an explanation to what it is and its benefits would have been nice
Maybe more explanations in the PDF so we can listen without taking notes and most of all, reproduce at home! screenshots of the Docker interface would be appreciated
The exercises are very well planned and appropiate. However, the pace of the explanation was too fast for me. Also I found that the cnical problems leading to log out/log in solution (which were very frequent for me) are highly disruptive to follow the course, as all the files created are lost and they are often needed afterwards.



General Comments about the PSC Graduate Program / the URPP Evolution in Action / the PhD Program in Evolutionary Biology
17 responses
-
/
none
no
Great
Nothing to add
I'm not in any of these programms
Very well organized
Nope.
No
N/A
NA



Other topics you would like to see offered.
17 responses
-
maybe a longer course for the same topic, a day is very short
none
Machine Learning
Maybe a more advanced module for BASH scripting
Nothing to add
The chosen topics were just right for what can bit squeezed into a 1day course
nil
Population genetics modeling.
Biochemical analytics
N/A
NA
If you would like to contribute to further improve the course, please leave your email.
2 responses
abhishek.meena@uzh.ch
-

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