

Supplementary Table 6

Counts of responses to pre- and post- questions for University of Cincinnati students about annotation and bioinformatics.

Note: Not all students completed the pre- and post-test resulting on only 22 students that could be assessed.

No.	Question	Pre-test	Post-test
1	What are some major difficulties associated with genome sequencing and assembly?	11/22	22/22
2	What information is utilized to predict genes in a newly assembled genome?	6/22	17/22
3	How does a sequenced genome increase tools available for research on an organism?	8/22	20/22