and prediction. 2nd ed. Springer; 2009.

19. Koo H, Hakim JA, Morrow CD, Eipers PG, Davila A, Andersen DT, et al. Comparison of two bioinformatics tools used to characterize the microbial diversity and predictive functional attributes of microbial mats from lake obersee, antarctica. Journal of microbiological methods. Elsevier; 2017;140:15–22.

20. Dedysh SN, Ivanova AO. Abundance, diversity, and depth distribution of planctomycetes in acidic northern wetlands. Frontiers in microbiology. Frontiers; 2012;3:5.

21. McFarland KD. OTU vs. ASV. https://github.com/EDUCE-UBC/MICB425/blob/master/Module_03/Project1/20180307_intro_OTU_ASV.pdf; 2018.

22. McFarland KD. Major differences in alph diversity between q1 and q2. https://forum.qiime2.org/t/major-differences-in-alph-diversity-between-q1-and-q2/2419/3; 2018.