

The pulmonary mycobiome - a study of subjects with and without chronic obstructive pulmonary disease

Supporting Information, S6 Fig

Einar M. H. Martinsen^{1*}, Tomas M. L. Eagan^{1,2}, Elise O. Leiten¹, Ingvild Haaland¹, Gunnar R. Husebø^{1,2}, Kristel S. Knudsen², Christine Drengenes^{1,2}, Walter Sanseverino³, Andreu Paytuví-Gallart³, and Rune Nielsen^{1,2}

¹Department of Clinical Science, University of Bergen, Bergen, Norway

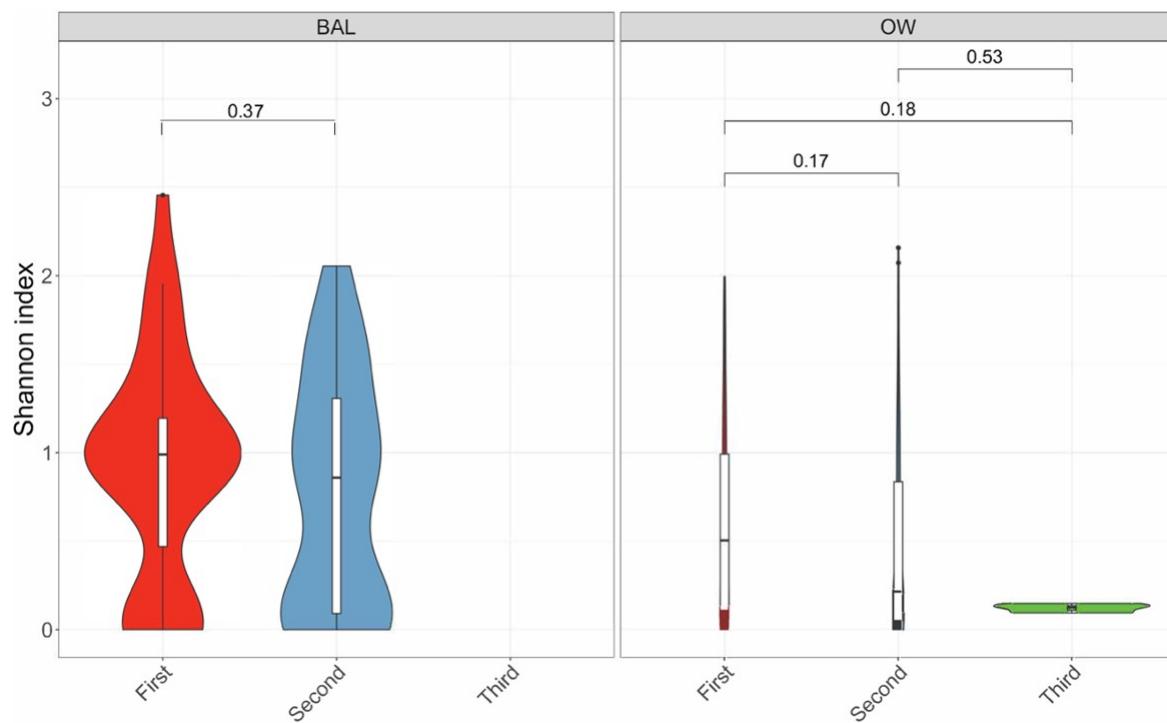
²Department of Thoracic Medicine, Haukeland University Hospital, Bergen, Norway

³Sequentia Biotech SL, Barcelona, Spain

* Corresponding author

E-mail: einar.martinsen@uib.no

S6 Fig. Alpha diversity plots and comparisons between sequencing runs.



BAL: bronchoalveolar lavage, OW: oral wash. Alpha diversity was evaluated using Shannon index. Statistical differences in alpha diversity were tested using Kruskal-Wallis.