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

Supporting Information, S4 Table

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S4 Table. Summary of read/sequence counts in the MicroCOPD study.

Data	Number of samples	Read/sequence count	Mean read/sequence count
Raw data			
NCS	189	667,033	3,529
OW	193	22,933,493	118,826
BAL	193	20,348,685	105,434
Post ITSxpress			
NCS	184	529,519	2,878
OW	193	11,907,308	61,696
BAL	193	5,790,865	30,004
Post DADA2			
NCS	161	472,574	2,935
OW	193	11,236,458	58,220
BAL	193	5,317,673	27,553
Post LULU			
NCS	157	410,205	2,613
OW	193	10,881,332	56,380
BAL	192	4,300,589	22,399
Without low abundance and contaminants			
NCS	122	73,658	604
OW	192	9,490,666	49,431
BAL	185	1,663,488	8,992

NCS: Negative control sample, OW: oral wash, BAL: bronchoalveolar lavage, DADA2:

Divisive Amplicon Denoising Algorithm version 2.