
Corrigendum: Whole-genome analyses reveal gene content differences between nontypeable *Haemophilus influenzae* isolates from chronic obstructive pulmonary disease compared to other clinical phenotypes

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In the published version of Supplementary Dataset 1 associated with this article and hosted on figshare at <https://doi.org/10.6084/m9.figshare.12545957.v2>, there was an error in the listing of the NTHi isolate on line 551. 'RHH-3' should have read 'RHH-15'.

This has now been corrected in the updated Supplementary Dataset which can be found on figshare at: <https://figshare.com/s/8d95658f8bda7eed27c4>.

The authors regret any inconvenience caused.

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