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Corrigendum: White-nose syndrome without borders: *Pseudogymnoascus destructans* infection tolerated in Europe and Palearctic Asia but not in North America

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This Article contains typographical errors.

In the Results section under subheading ‘Quantitative comparison of WNS on bats’

“The fungal load on qPCR-positive bats ranged from 0.21 pg to 3.41 µg across the surface of the left wing (Supplementary Fig. S1, see Table 1 for sample sizes)”.

should read:

“The fungal load on qPCR-positive bats ranged from 0.21 fg to 3.41 ng across the surface of the left wing (Supplementary Fig. S1, see Table 1 for sample sizes)”.

“The fungal load from UV-negative individuals (median = 3.78×10^{-5} µg) overlapped that from UV-positive individuals (median = 7.46×10^{-4} µg; Fig. 8)”.

should read:

“The fungal load from UV-negative individuals (median = 3.78×10^{-5} ng) overlapped that from UV-positive individuals (median = 7.46×10^{-4} ng; Fig. 8)”.

In the upper panel of Figure 4, the y-axis ‘Fungal load per cm² (log₁₀(ng))’ was incorrectly given as ‘Fungal load per cm² (log₁₀(µg))’

In the upper panel of Figure 5, the y-axis ‘*P. destructans* load per cm² (log₁₀(ng))’ was incorrectly given as ‘*P. destructans* load per cm² (log₁₀(µg))’

In Figure 6, the x-axis ‘*P. destructans* load per cm² (log₁₀(ng))’ was incorrectly given as ‘*P. destructans* load per cm² (log₁₀(µg))’

In Figure 8, the x-axis ‘Fungal load per cm² (log₁₀(ng))’ was incorrectly given as ‘Fungal load per cm² (log₁₀(µg))’

In the legend of Figure 8,

“Bats were identified as positive (grey; $n = 255$) and negative (orange; $n = 151$) using UV trans-illumination”.

should read:

“Bats were identified as positive (orange; $n = 255$) and negative (grey; $n = 151$) using UV trans-illumination”.

The correct Figures 4, 5, 6 and 8 appear below as Figures 1, 2, 3 and 4 respectively.

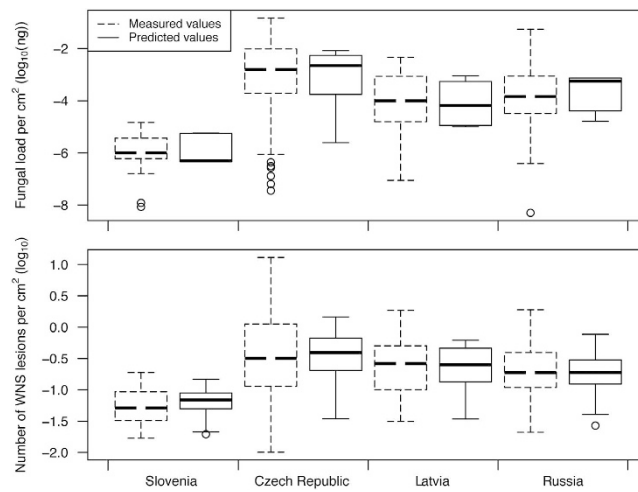


Figure 1.

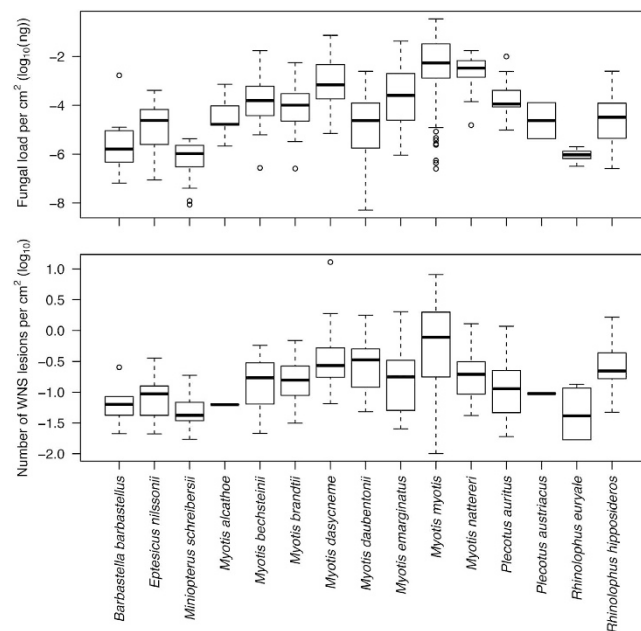


Figure 2.

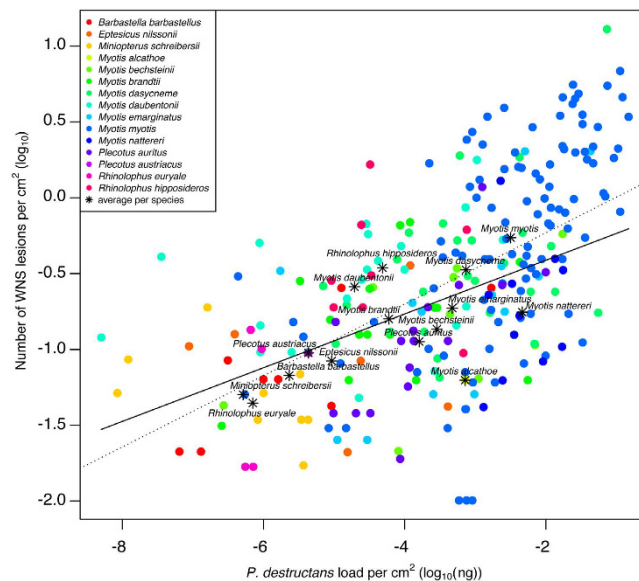


Figure 3.

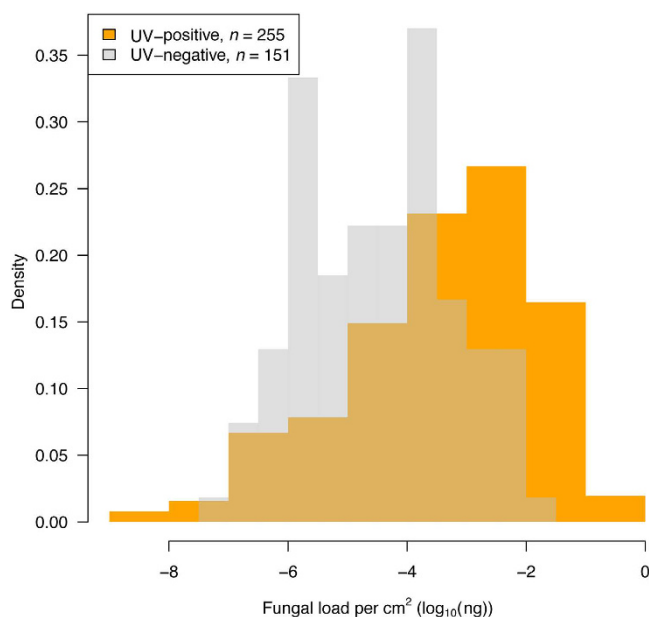


Figure 4.



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