





Correction

Correction: Díaz-García et al. *Candida* Genotyping of Blood Culture Isolates from Patients Admitted to 16 Hospitals in Madrid: Genotype Spreading during the COVID-19 Pandemic Driven by Fluconazole-Resistant *C. parapsilosis*. *J. Fungi* 2022, 8, 1228

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Error in Figure 2:

In the original publication [1], a format error in Figure 2 was detected. A correction has been made to Figure 2 in the Results section, Section 3.1. The legend to the figure was truncated, and the missing parts made the interpretation of the whole figure difficult. Figure 2 should be changed to:

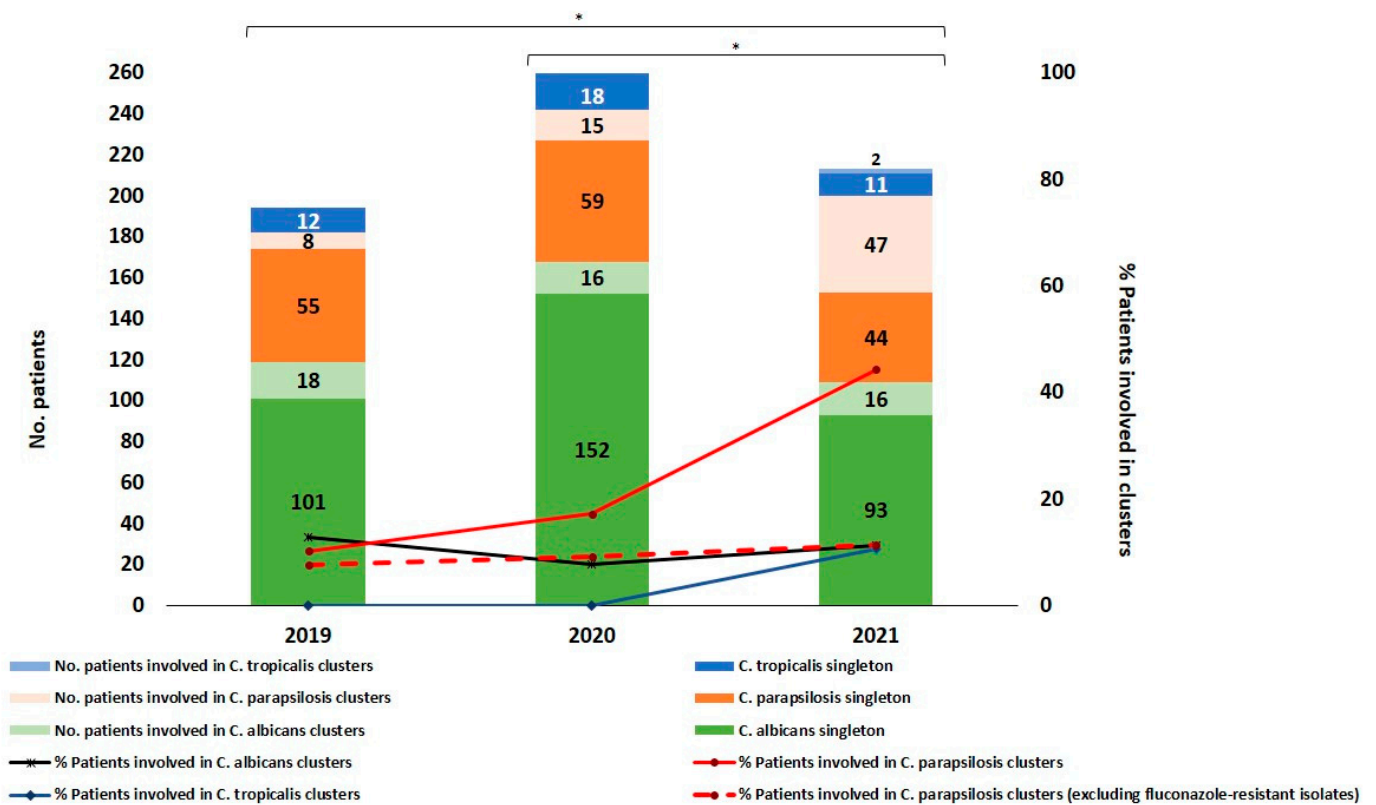


Figure 2. Numbers (and percentage) of patients involved in singleton and intra-hospital clusters per species over the study period. * Differences reaching statistical significance ($p < 0.05$).

The authors apologize for any inconvenience caused and state that the scientific conclusions are unaffected. This correction was approved by the Academic Editor. The original publication has also been updated.

Reference

1. Díaz-García, J.; Gómez, A.; Machado, M.; Alcalá, L.; Reigadas, E.; Sánchez-Carrillo, C.; Pérez-Ayala, A.; de la Pedrosa, E.G.-G.; González-Romo, F.; Cuétara, M.S.; et al. *Candida* Genotyping of Blood Culture Isolates from Patients Admitted to 16 Hospitals in Madrid: Genotype Spreading during the COVID-19 Pandemic Driven by Fluconazole-Resistant *C. parapsilosis*. *J. Fungi* **2022**, *8*, 1228. [CrossRef] [PubMed]

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