

Supplementary information

**Evolution of the human pathogenic lifestyle
in fungi**

In the format provided by the
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Figure S1. Human pathogenicity has repeatedly evolved in fungi. Genera and lineages harboring major and emerging fungal pathogens (see Table 1) are shown in red and non-pathogenic taxa are shown in black. The fungal tree of life based on a phylogenomic analysis of 1,644 species and 290 genes from Li et al. (2021). A genome-scale phylogeny of the kingdom Fungi. *Curr. Biol.* 31: 1653–1665.

