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Adaptation to environmental pH in *Candida albicans* and its relation to pathogenesis

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In the section “Rim101p processing in *C. albicans*” 2nd paragraph, the sentence “However, several issues need to be resolved, including the fact that PacC is localized to the nucleus at acidic pH and in the presence of upstream pathway mutants.” should have read “However, several issues need to be resolved, including the fact that PacC is not excluded from the nucleus at acidic pH and in presence of upstream pathway mutants.”

Also, these two references were omitted from the manuscript.

The reference Diez E et al. (2002) Activation of the Aspergillus PacC zinc finger transcription factor requires two proteolytic steps. *Embo J* 21:1350–1359 was omitted from the section “Rim101p processing in *C. albicans*”

first paragraph. “In *A. nidulans*, processing occurs in two steps, the first which gives rise to an ~500 residue product similar to that seen in *S. cerevisiae* and a second processing event that occurs distal to the zinc finger domain and yields an ~250 residues active product (Mingot et al. 1999; Espeso et al. 2000; Diez et al. 2002).”

The reference Espeso EA, Arst HN, Jr. (2000) On the mechanism by which alkaline pH prevents expression of an acid-expressed gene. *Mol Cell Biol* 20:3355–3363 was omitted from the section “Rim101p, activator or repressor?” first paragraph. “*A. nidulans* PacC binds to the consensus sequence GCCAAG with high affinity and GCCAGG with lower affinity and it can act as an activator and a repressor at these sites (Tilburn et al. 1995; Espeso and Penalva 1996; Espeso et al 1997; Espeso and Arst 2000).”

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