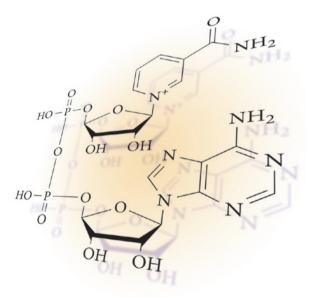
FUNGAL VIRULENCE

Salvageable research

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NAD⁺ is an important coenzyme that has essential roles in many metabolic pathways. Researchers from Brendan Cormack's laboratory are interested in the connections between this coenzyme and virulence in <u>Candida</u> <u>glabrata</u>. They have now successfully defined the NAD⁺-biosynthesis pathways in this opportunistic fungal pathogen, and the results are presented in a recent issue of <u>Molecular</u> <u>Microbiology</u>.

C. glabrata is an NAD⁺ auxotroph and must therefore synthesize NAD⁺ from the vitamin precursors that are present in the environment. Ma and colleagues began by investigating which precursors *C. glabrata* can salvage, and they found that, like



Saccharomyces cerevisiae, C. glabrata can use nicotinic acid (NA), nicotinamide (NAM) and nicotinamide riboside (NR) as sources of NAD+. In S. cerevisiae, which, unlike C. glabrata, can also synthesize NAD⁺ de novo, NA and NAM are both salvaged through the evolutionarily conserved Preiss-Handler pathway. Npt1 and Qns1, which catalyse the first and last steps, respectively, are essential enzymes in this pathway. The presence of a functional Preiss-Handler pathway in C. glabrata, and its involvement in the conversion of NA to NAD⁺, was confirmed by deletion of the C. glabrata orthologues of these key S. cerevisiae enzymes, as the authors found that, with NA as the sole NAD⁺ source, the *npt1* Δ and $qns1\Delta$ C. glabrata strains were unable to grow.

Ma and colleagues went on to investigate the pathways that *C. glabrata* uses to salvage the other two vitamin precursors, NAM and NR. In S. cerevisiae, Pnc1 converts NAM to NA, which can then enter the Preiss-Handler pathway for conversion to NAD+. The *npt1* Δ , *qns1* Δ and *pnc1* Δ strains of C. glabrata could not grow in media that contained NAM as the only source of NAD+, indicating that the same NAM-salvage pathway is used by C. glabrata. For NR, it was known that S. cerevisiae can salvage this precursor independently of the Preiss-Handler pathway using a

pathway that requires the NR kinase Nrk1. Ma *et al.* confirmed that *C. glabrata* can salvage NR using this same pathway. Surprisingly, however, it was also found that *C. glabrata nrk1* Δ strains grow as well as the parent strain if NR is the sole source of NAD⁺, indicating that, in addition to this Nrk1-dependent pathway, *C. glabrata* also possesses an Nrk1independent NR-salvage pathway.

The authors postulated that this alternative pathway might involve the combined action of the nicotinamidase Pnc1 and a nucleosidase to form NA, which can then be funnelled into the Preiss–Handler pathway. Using a range of deletion mutants, they were able to demonstrate that Urh1 and Pnp1 are the main nucleosidases that are involved, with a minor role for a third nucleosidase, Meu1. Finally, studies using a mouse model revealed that NR is the primary NAD⁺ source during *C. glabrata* disseminated infection *in vivo*.

Previous work in the Cormack laboratory has shown that the availability of NAD⁺ vitamin precursors is a key regulator of *C. glabrata* virulence. This detailed paper has now fully delineated the salvage pathways by which these precursors are converted to NAD⁺.

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