# PERSPECTIVES

cph1 strains make pseudohyphae like the wild type.

How do these varied observations on the control of cell morphology in Candida fit together? The most striking observations suggest that the yeast form is the result of negative control of the pseudohyphal growth state. The results of Braun and Johnson, implicating TUP1, are particularly provocative because TUP1 in Saccharomyces cerevisiae functions as a global repressor, turning off genes in response to major alterations in the state of the cells. There Tup1p protein (together with Ssn6p, the product of the SSN6 gene) controls such processes as cell type (a or  $\alpha$ ) and glucose repression. Tup1p is targeted to a particular site by DNA binding proteins like Alpha2p and Mcm1p (6, 7); it then acts to block transcription at the adjacent promoter, probably by altering the chromatin structure. TUP1-mediated repression is alleviated when the targeting DNA binding protein is no longer present. Interestingly, deletion of TUP1 in S. cerevisiae depresses formation of pseudohyphae.

We are still early in the genetic analysis of the polymorphic character of this fungus, but a working model that incorporates almost all the data can be constructed (see figure). A likely explanation for the Braun and Johnson results is that in the absence of the TUP1 gene product, the genes for pseudohypha formation are constitutively on. These genes may include a repressor for the yeast genes. The basic feature of the model is that the process is controlled by a series of negative and positive transcription factors, which ultimately affect the TUP-mediated regulation of pseudohyphae and yeast cell formation. In this model, Efg1p, the product of the gene studied by Stoldt et al. (3), would function as a repressor of the TUP complex (perhaps by repressing the synthesis of the putative DNA targeting protein). Then overexpression of Efg1p would block the formation of the TUP complex and allow expression of the pseudohyphal genes. RBF1 would serve as a repressor of EFG1; a deletion of RBF1 would increase the synthesis of Efg1p and lead once more to a decrease in the amount of TUP repression. Chp1p would activate the synthesis of Efg1p; when CPH1 is deleted, little Efg1p is made, and synthesis of the TUPcomplex continues at a high level. Thus, no pseudohyphae can be made. One piece of evidence not accounted for by this model is the pseudohyphal growth of cells in which the amount of Efg1p is reduced to a low level. Because EFG1 is an essential gene, this phenotype may reflect some other function of Efg1p. The model in the figure makes several predictions: Overexpression of Efg1p, like deletion of TUP1, should be epistatic to CPH1 deletion, but if Cph1p is necessary for the synthesis of Efg1p, deletion of CPH1 should itself be epistatic to deletion of RBF1. In other words, the RBF1-CPH1 double-deletion strain should not form pseudohyphae. Finally, overexpression of Tup1p might prevent formation of pseudohyphae under most or all conditions. Given the early state of our analysis of this complex regulatory circuit, many other models can no doubt be devised to suit the data, but the one presented here, although highly speculative, has the advantage of being easily testable.

There are several other implications of this group of papers. Most important, perhaps, is that C. albicans should not be thought of as a yeast that can assume various forms. The yeast form exists only if the general repressor TUP1 is active. A strong case can be made that there is no "default" form for this organism. A second important implication is that hypha formation seems to occur by means of a pathway separate from that for pseudohyphae; none of the deletions or constructs that affect pseudohyphae affect hyphae in similar ways. It also seems likely that alterations in chromatin structure are an important mode of control of morphogenesis in Candida; the sort of silencing that has been well studied in S. cerevisiae may occur in pseudohyphal genes. Finally, neither deletion of RBF1 nor overexpression of Efg1p completely blocks formation of yeast cells; hence, neither of these alterations causes as profound a change in the cell morphology as the TUP1 deletion, reinforcing the view that the ultimate control of pseudohypha formation is mediated by a TUP complex.

The role of polymorphism in the pathogenesis of C. *albicans* is far from proven, but the capability seems intuitively important for an organism that is found in a variety of different niches in the body. The elucidation of the regulatory cascade in the papers by Braun and Johnson and others is likely to be of great value in understanding the ways in which this organism causes disease, a topic about which we know little for any pathogenic fungus.

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## Online archaeology www.ncl.ac.uk/~nktg/wintro/

Extensive digging is not necessary to find useful archaeology information at this site, created by K. Greene of the University of Newcastle upon Tyne as an electronic companion to his text "Archaeology—An Introduction." The Web companion is a hyperlinked outline of the book with summaries of chapter contents, supplemented by a thorough set of Web links to other archaeology sites at the end of each section.

## Biocatalysis and biodegradation dragon.labmed.umn.edu/~lynda/ index.html

Understanding biocatalytic reaction pathways is important for both pollution control and applications in biotechnology. The University of Minnesota Biocatalysis/Biodegradation Database aids learning about these reactions and provides extensive links to Web resources on microbial enzymecatalyzed chemistry. Each chemical pathway is described in text and is accompanied by a graphical represention of the reaction. There is also a graduate level course on biocatalysis offered over the Internet by the site's creators L. Ellis and L. Wackett.

## Neuro net

#### www.neuroguide.com

"Neurosciences on the Internet" is part Web guide and part electronic journal created by N. Busis of Shadyside Hospital, Pittsburgh. The site has a large set of pointers to Web, gopher, mail list, and ftp resources of interest to neuroscientists, including links to image repositories and software collections. In addition, the Web site publishes original contributions vetted by peer reviewers on topics ranging from amyotrophic lateral sclerosis to viral infections in the central nervous system.

Edited by David Voss

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