Lipids of *Candida albicans*: subcellular distribution and biosynthesis

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Lipids constituted around 5% of the dry weight in *Candida albicans* 3153, while sterols and phospholipids accounted for 1.2% and 1.1% respectively. Phospholipids were mainly localized in the microsomal fraction; phosphatidylserine (PS), phosphatidylcholine (PC), phosphatidylethanolamine (PE) and phosphatidylinositol (PI) were the major phospholipids. Incorporation studies with [14C]acetate and [32P]orthophosphoric acid demonstrated that PS was synthesized at the highest rate followed by PC, PE and PI. There was little difference in either the content or the rate of biosynthesis of PC, PE and PI. Incorporation of labelled serine, ethanolamine and choline revealed serine to be a precursor for PC, PE and PS, ethanolamine for PC and PE, and choline for PC biosynthesis only.

Introduction

Lipids of micro-organisms are important not only due to their interaction with antimicrobial drugs (Fryberg et al., 1975; Yamaguchi, 1977), but also because of their response to changes in the growth medium. For *Candida albicans*, there are reports on the lipid composition of the plasma membrane and cell wall (Marriott, 1975), variation in lipid composition associated with dimorphism (Ghannoum et al., 1986), and the effect of antifungal agents on lipids (Singh et al., 1979). Total lipid content ranges from 7% to 32% in various species of *Candida* (Kaneko et al., 1976). Phosphatidylcholine (PC), phosphatidylethanolamine (PE), phosphatidylserine (PS) and phosphatidylinositol (PI) represent the major phospholipids, while triglycerides, free sterols, esterified sterols and free fatty acids are the major neutral lipids (Henry et al., 1981; Kaneko et al., 1976). Although *C. albicans* lipids have been studied in detail, information on lipid biosynthesis is meagre and variation in lipid content among different strains has not been studied. The study of lipid composition of different subcellular organelles and their synthesis should help to determine their functional capabilities, as phospholipids are the structural components of biomembranes. Therefore, we have examined the composition and biosynthesis of lipids in *C. albicans* using various precursors.

Methods

**Materials.** [1-3H]Ethanolamine (specific activity 8.8 mCi mmol⁻¹; 325·6 MBq mmol⁻¹) was obtained from Amersham. [methyl-14C]Choline chloride (specific activity 5·mCi mmol⁻¹; 185 MBq mmol⁻¹), [U-14C]serine (specific activity 173·mCi mmol⁻¹; 6401 MBq mmol⁻¹), [1-14C]acetate (specific activity 60·3 mCi mmol⁻¹; 2231 MBq mmol⁻¹) and [32P]orthophosphoric acid (carrier-free) were obtained from BARC, Bombay, India.

**Organism and growth conditions.** *C. albicans* 3153, obtained from the Mycological Reference Laboratory, Colindale, London, UK, was maintained on Sabouraud's dextrose broth and cultivated in a defined medium containing 0·3% KH2PO4, 0·3% (NH4)2SO4, 0·025% MgSO4, 0·025% CaCl2, 2H2O, 0·001% biotin and 0·5% glucose as carbon source at 37 °C on a gyrotatory shaker (200 r.p.m.). Cells were harvested in the stationary phase (after 24 h) for lipid analysis and in the exponential phase (after 16 h) for studies of lipid biosynthesis.

**Subcellular fractionation of cells.** Cells were harvested, washed and sonicated in 0·01 M-Tris/HCl pH 8·0 containing 250 mM-sucrose in a Branson Sonifier for 30 min in ice at 60 W. The homogenate was then centrifuged at 5000 g for 20 min to remove cell debris. Cell fractionation and characterization was done according to the procedure of Chavant et al. (1980) except that mitochondria and microsomes were isolated by centrifugation at 15000 g and 105000 g respectively.

**Labelling of lipids.** Cells were washed with 0.85% NaCl and resuspended in medium (for 32P-incorporation experiments, phosphorus-free medium was used) under sterile conditions. Cells were preincubated at 37 °C for 30 min with shaking. Precursors (10 μCi [14C]acetate, 4 mCi [32P]orthophosphate, 6 μCi [14C]serine, 6 μCi [3H]ethanolamine or 6 μCi [14C]choline chloride per g of cells) were added to cells suspended in 50 ml medium and at different time intervals 3 ml samples from each culture were transferred into tubes containing 0·5 ml 1 M-KCN (Kilburn et al., 1981). The tubes were centrifuged at 2700 g for 15 min. The cell pellet was recovered and lipids extracted. Radioactivity was counted in Packard Tricarb scintillation counter using a toluene-based scintillation fluid.

**Abbreviations:** CL, cardiolipin; LPC, lysophosphatidylcholine; PC, phosphatidylcholine; PE, phosphatidylethanolamine; PI, phosphatidylinositol; PS, phosphatidylserine.
Quantification of lipids. Lipids were extracted by the method of Folch et al. (1957). Phospholipids were quantified by the method of Marinetti (1962), and sterols by the method of Zlatkis et al. (1953), which is specific for ergosterol/cholesterol. Lipids of the subcellular fractions were extracted by the method of Bligh & Dyer (1959). Individual phospholipids were separated by one-dimensional thin-layer chromatography on silica gel H plates in chloroform/methanol/7 m-NH$_3$ (65:25:4, by vol.). Fatty acid methyl esters of phospholipids (separated from the neutral lipids by thin-layer chromatography) were prepared by transesterification with methanol in the presence of thionyl chloride (Khuller et al., 1981) and were resolved in an AILNucogas chromatogram fitted with a DEGS column at 180 °C using nitrogen as a carrier gas (flow rate 40 ml min$^{-1}$). Fatty acid methyl esters were identified by comparison of their retention times with those of authentic standards. Fatty acids were quantified by triangulation.

Results and Discussion

Lipid composition

The total lipid content of C. albicans 3153 was 5.15 ± 0.27% of the dry weight. Phospholipids and sterols represented 1.11 ± 0.11 and 1.22 ± 0.13% of the dry weight, respectively, and these components were present in equimolar ratio (Table 1). Hitchcock et al. (1986) reported the total lipid content of this strain to be 0.85%, and the ratio of phospholipids to sterols to be 3:1. These variations in results are presumably due to differences in growth medium or stage of growth. In earlier studies with different strains of C. albicans, total lipid content of C. albicans ATCC 10231 was found to vary from 0.3 to 6.3% of the dry weight, and that of C. albicans 1 Ha 582 was 13.9% of the dry weight (Weete, 1980). The group and fatty acid composition of lipids from 62 species and strains of Candida was investigated by Malkhas'ian et al. (1982) but no strict correlation between the composition of lipids and yeast species was found.

Our analysis of individual phospholipids of C. albicans 3153 revealed the presence of PC, PE, PS, PI and cardiolipin (CL) of which PS constituted the greatest amount and CL the least. Hitchcock et al. (1986) and Ghannoum et al. (1986) have also reported PC, PE, PS and PI to be major phospholipids of C. albicans, although in their case PC was present in the largest quantity followed by PE, PS and PI. Since the lipid composition of plasma membranes of C. albicans has been determined earlier (Marriott, 1975), it was of interest to study the lipid composition of two other metabolically active organelles, mitochondria and microsomes, of which little is known. Phospholipid analysis of subcellular fractions revealed that the greatest amounts were present in the microsomal fraction [30.11 ± 5.05 mg (g dry wt)$^{-1}$ (mean ± SD, n = 4)]. The mitochondrial fraction contained a considerable amount of phospholipids [12.26 ± 1.91 mg (g dry wt)$^{-1}$] while the cytosolic fraction contained only negligible amounts [0.12 ± 0.04 mg (g dry wt)$^{-1}$].

Fatty acid analysis of the phospholipids of whole cells of C. albicans revealed that C$_{16}$:0 (24%), C$_{18}$:1 (15%), C$_{18}$:2 (32%) and C$_{18}$:0 (8.6%) were the main fatty acids while C$_{16}$:0, C$_{12}$:0, C$_{14}$:0, C$_{14}$:1, C$_{16}$:1 and C$_{16}$:2 were minor ones (each less than 5%). These results are in agreement with those of Pierce et al. (1978) and Hitchcock et al. (1986), who reported C$_{16}$:0, C$_{16}$:1, C$_{18}$:1 and C$_{18}$:2 as major fatty acids. Differences between results of the present study and earlier ones (Ghannoum et al., 1986) are probably due to differences in growth phase and composition of the medium. In comparison to whole cells, subcellular fractions contained larger amounts of C$_{16}$:0 (60% in mitochondria and microsomes and 35% in cytosol), while the levels of C$_{18}$:2 were significantly lower (32% in whole cells to traces in various fractions) with marginal changes in C$_{18}$:0. There was little change in C$_{18}$:1, while C$_{16}$:1 increased considerably. Since relative percentage was measured and not the absolute amount, abundance of one fatty acid would decrease the percentage of other fatty acids. The mitochondrial and microsomal fractions had almost the same fatty acid composition while cytosolic fractions contained two extra fatty acids (C$_{14}$:0, C$_{14}$:1) as major components (20% and 14%) in addition to C$_{16}$:0, C$_{16}$:1 and C$_{18}$:1 present in mitochondria and microsomes.

Lipid biosynthesis

Lipid biosynthesis in C. albicans was studied with $[^{14}$C]$\text{acetate}$, a general precursor for all lipids, and $[^{32}$P]$\text{orthophosphoric acid}$, a specific precursor for

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* Values are means ± SD of five different batches.
† Values are means ± SD of eight different batches.
‡ UPL, unknown phospholipids.

Table 1. Lipid composition of whole cells of C. albicans 3153

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phospholipids. Incorporation of [14C]acetate into total lipids increased for 10 min, but it was only linear for 6 min (Fig. 1a); a similar pattern was exhibited for [14C]acetate incorporation into total phospholipids. Incorporation of [14C]acetate into individual phospholipids indicated that PS was synthesized at the highest rate while PI, PC and PE had almost the same rate of synthesis. These results differ from those of Ballman & Chaffin (1979), who found that the sequence of synthesis was PC > PE > PS. However, in that case lipid biosynthesis was studied during re-initiation of growth from the stationary phase and not in mid-exponential phase as in the present study. When [32P]orthophosphoric acid was used as a precursor, incorporation was linear for 6 min (Fig. 1b) and the rate of synthesis of PS was highest, as observed with [14C]acetate. Among the other phospholipids, PI was synthesized at a higher rate than PE and PC. Our results of lipid biosynthesis confirm the lipid composition of C. albicans (Table 1) and that PS was the major phospholipid while other phospholipids have lower, relatively similar, concentrations.

While studies with [14C]acetate and [32P]orthophosphoric acid as precursors give a generalized view of phospholipid biosynthesis, incorporation of various specific bases (choline, ethanolamine and serine) into lipid fractions was also examined to define the contribution of these bases to total PL biosynthesis. Of these three bases, maximum incorporation into total lipids and phospholipids was observed with [14C]choline chloride followed by [3H]ethanolamine and [14C]serine; incorporation of all three bases increased for 9 min (data not shown).

Incorporation of [14C]choline into individual phospholipids in 9 min (Table 2) showed around 87% incorporation into lysophosphatidylcholine (LPC) plus PC and negligible incorporation into the others. Direct incorporation of [14C]choline into PC indicated the cytidine pathway for PC synthesis, which has been shown in other fungi (Kasinathan & Khuller, 1983).

When [3H]ethanolamine was used as a precursor, maximum radioactivity (62%) was again observed in choline-containing lipids and only about 6% was detected in PE, while the rest was distributed in other phospholipids. Even after only 3 min, 54% of the radioactivity was present in the PC plus LPC fraction, whereas there was no difference in incorporation of radioactivity between 3 and 9 min into the PE fraction (data not shown). The rapidity of incorporation of labelled ethanolamine into PC and LPC suggests that PE is not the only substrate for methylation but that ethanolamine, phosphorylethanolamine and CDP-ethanolamine are also methylated using S-adenosylmethionine as the methyl donor, as suggested in mammalian systems (Salerno & Beler, 1973). However, this pathway needs further confirmation. Increased uptake into LPC as compared to other phospholipids may be due to degradation of PC to LPC by phospholipase A, which has been identified in this fungus (unpublished observations).

Studies with [14C]serine yielded results similar to the ethanolamine incorporation experiments. The incorporation into PS decreased from 36-9% after 3 min of pulse to 11-0% after 9 min while it increased in choline-containing lipids from 48-0% to 56-6%, indicating that PS is converted to PC. These results suggest that PS is decarboxylated by PS decarboxylase and then subsequently methylated to PC by methyltransferases. Earlier workers obtained similar results when they studied incorporation of [14C]serine in Saccharomyces cerevisiae in vivo and in vitro (Nikawa & Yamashita, 1983; Steiner & Lester, 1972).
This study has demonstrated that in C. albicans 3153, microsomes are the major reservoir of phospholipids represented by PS, PC, PE and PI. Choline was found to be a specific precursor for PC, whereas serine and ethanolamine could be used for the biosynthesis of other phospholipids.

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References


