ADAPTIVE LOWERING OF THE LIPID CLUSTERING TEMPERATURE WITHIN TETRAHYMENA MEMBRANES

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1. Introduction

Variations in temperature can effect dramatic changes in structural and functional features of biomembranes. For instance, freeze-etch electron microscopy has recently revealed a lateral separation of membrane components directly within the membrane cores of the endoplasmic reticulum in cells of the unicellular eukaryote Tetrahymena upon lowering these cells' environmental temperature from that optimal for growth at 28°C to below 18-16°C [1]. At this temperature range, a discontinuity is also detected in the activity of the endoplasmic reticulum marker enzyme, glucose 6-phosphatase, which is bound to the smooth microsomal membranes isolated from these 28° C-Tetrahymena cells [1]. Different physical methods such as electron spin resonance, calorimetry, fluorescence probing, and proton nuclear magnetic resonance have indicated that these thermotropic membrane changes are very probably induced by a clustering of 'rigid' liquid crystalline membrane lipids ([1] cf. also [2]) rather than by a true liquid causes such membrane changes in other cells, e.g. [3-6]. The present paper provides evidence that Tetrahymena cells grown at 18°C have adaptively shifted the temperature of the lipid clustering within endoplasmic reticulum membranes down to $\sim 12^{\circ}$ C by replacing the more saturated with more unsaturated membrane phospholipid species.

2. Materials and methods

Cell Static cultures of the ciliate protozoan

Tetrahymena pyriformis GL were axenically grown at 18°C in the logarithmic growth phase (~20 000 cells/ml) in a medium consisting of 0.75% proteose peptone, 0.75% yeast extract, 1.5% glucose, 0.1 mM ferric citrate, 0.05 mM CaCl₂, and 1 mM MgSO₄.

Freeze-etch electron microscopy. In three experiments, a *Tetrahymena* culture was divided into several aliquots, which were equilibrated for 10 min at 18°C, 15°C, 11.5°C, 8°C, and 5°C. At these temperatures, the cells were fixed with 2% glutaraldehyde buffered with 0.05 N Na-cacodylate (pH 7.2). After 10 min, the cells were twice washed with the cacodylate buffer and stepwise glycerinated up to 25% for 2 hr, before they were frozen on cardboard disks in Freon 22 cooled by liquid nitrogen. Fracturing, etching (1 min at–100°C) and replication were carried out in a Balzers freeze-etch device model BA 260M. The replicas were studied in a Siemens Elmiskop Ia.

Membrane isolation. The light smooth microsomal fraction containing only smooth-surfaced membrane vesicles was isolated on step sucrose gradients as described recently [7].

Glucose 6-phosphatase. This enzyme was determined at V_{max} in the membrane vesicles fraction as a function of temperature as described previously [1].

Lipid analysis. Total lipids were extracted from the membrane vesicles with chloroform/methanol (2:1) at 0-4°C according to Folch et al. [8]. Polar and neutral lipids were separated, identified and quantified by two- and one-dimensional thin-layer chromatography, respectively, as described recently. [7]. The fatty acids of the total lipids, polar lipids and neutral lipids were saponified, methylated and identified in a Varian gas chromatograph as described previously [7].

3. Results

3.1. Thermotropic membrane response

At cleavage, freeze-etch electron microscopy exposes two faces of the hydrophobic core of the endoplasmic reticulum membranes in Tetrahymena [9,10]. At 18°C and 15°C, the PF-faces (i.e. the faces bordering the cytoplasm) are studded with numerous, uniformly distributed particles, ~ 7.5 nm in diameter (fig.1a). The opposite EF-faces (i.e. the faces oriented to the cisternal space) show only some particles and numerous depressions (fig.1a). At 11.5°C, however,

about 40% of both fracture faces reveal particle-depleted areas (cf. fig.1b). These can be regularly seen in *Tetrahymena* cells equilibrated at 8° C. At 5° C, these smooth areas have largely expanded and can reach diameters of up to $1 \mu m$ (fig.1b).

The temperature range of the emergence of pricicle-depleted areas is the same as that ($\sim 12^{\circ}$ C), where the activity of the glucose 6-phosphatase associated with the smooth microsomal membrane vesicles shows a discontinuity (fig.2). Above and below this discontinuity the activation energies lie in the range of ~ 6 Kcal/mol and ~ 10 Kcal/mol.





Fig.1. Fracture faces of endoplasmic reticulum membranes of *Tetrahymena*; shadowing direction from below. (a) At 18° C, numerous particles distribute uniformly on the pF-faces, while the EF-faces reveal numerous depressions. (b) A large smooth area can be seen on a PF-face at 5° C. Note that particles in the particle-bearing areas are not aggregated. Magnification \times 70 000.

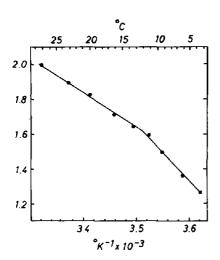


Fig. 2. Arrhenius plot of the glucose 6-phosphatase associated with the smooth microsomal membranes of *Tetrahymena*. Ordinate: Logarithm of glucose 6-phosphatase activity normalized to 27°C. Note the discontinuity at \sim 12°C!

3.2. Membrane composition

The smooth microsomal membranes of *Tetra-hymena* cells grown at 18°C exhibit the same ratios

of RNA/protein (0.09), phospholipid/protein (0.53) and neutral lipid/phosphotipid (0.17), as well as the same SDS-gelelectrophoretic protein pattern, as those isolated from cells grown at 28°C [7]. Differences between 28°C- and 18°C-Tetrahymena cells can be detected only in the pattern and fatty acid composition of the phospholipids (tables 1 and 2). In comparison to 28°C-cells, the smooth microsomes of Tetrahymena cells grown at 18°C contain 5% more phosphatidylethanolamine and its phosphonoanalogue glycerideaminoethylphosphonate, while ceramideaminoethylphosphonate, phosphatidylinositol and lysophosphatidylcholine have decreased by about the same amount (table 1). Since we have previously shown [7] that phosphatidylethanolamine and glycerideaminoethylphosphonate contain much more unsaturated acids than the other three phospholipid species, one would expect a higher degree of unsaturation in the polar lipid and total lipid fraction, respectively, of the smooth microsomes in 18°C-cells than in 28°C-cells. Indeed, this is found as can be seen from table 2.

Table 1
Phospholipid composition of the smooth microsomal membranes of *Tetrahymena* grown at 18°C^a and 28°C^b

Phospholipid	Mole % of total lipid phosphorus		
	18°C-cells	28°C-cells	
Phosphatidylethanolamine	31.8	30.0	
Glyceridearninoethylphosphonate	26.2	23.0	
Ceramideaminoethylphosphonate	12.9	14.2	
Phosphatidylinositol	2.8	3.9	
Lysophosphatidylcholine	2.2	4.8	
Phosphatidylcholine	18.2	18.1	
Monomethylaminoethylphosphonate	4.4	4.7	
Cardiolipin	1.2	1.3	

^a Average of three experiments.

b Data from Ronai and Wunderlich (1975).

Table 2
Major fatty acid composition of the total, polar and neutral lipid fractions of the smooth microsomal membranes of *Tetrahymena* grown at 18°C^a and 28°C^b (% of total fatty acids)

Fatty acids	Total lipids		Polar lipids		Neutral lipids	
	18°C-cells	28° C-cells	18°C-cells	28°C-cells	18°C-cells	28°C-cells
12:0	2.5	2.8	2.9	2.9	1.3	3.6
14:0 iso	10.3	10.7	8.5	10.3	4.4	11.2
14:0	2.0	3.2	3.2	3.4	2.1	4.4
15:0	1.6	1.2	1.5	1.6	2.9	3.7
16:0	10.4	11.9	6.6	12.0	2 4.6	12.8
16:1	13.0	10.4	12.5	8.9	13.2	10.2
16:2	1.4	1.3	2.2	1.2	1.4	2.5
17:0	2.1	2.3	3.9	1.4	4.0	1.8
18:0	2.9	4.7	2.6	3.2	8.5	9.6
18:1	7.2	8.0	6.0	8.1	9.3	7.4
18:2 △ 6,9	7.7	4.2	7.7	4.7	2.3	2.8
18:2 △ 9,12	15.9	17.5	18.4	19.3	13.2	15.8
18:3	23.2	21.6	24.5	23.1	12.3	14.0
Unsaturated	68.4	63.0	71.3	65.2	51.7	52.7

a Average of three experiments.

4. Discussion

Our data show that *Tetrahymena* cells grown at 18°C have shifted the lipid clustering temperature within the endoplasmic reticulum membranes down to $\sim 12^{\circ}\text{C}$. Interestingly, the ratio of lipid clustering temperature to cell growth temperature remains nearly the same in *Tetrahymena* cells grown at 28°C [1]. The lowering of the lipid clustering temperature is apparently achieved by replacing the more saturated phospholipid species by more unsaturated ones. This

supports the view that the poikilothermic eukaryote *Tetrahymena* is capable of adapting the fluidity of its membranes to long-time changes in environmental temperature [11,12], as it is also known in prokaryotes such as *Acholeplasma* [13,14], *Escherichia coli* [15,16] or *Bacillus* [17]. Unlike *Tetrahymena*, however, *E. coli* for instance lowers the liquid crystalline \Rightarrow crystalline phase transition of the membrane lipids upon lowering the cell growth temperature so that the difference between lipid phase transition temperature and growth temperature is kept almost constant [15].

b Data from Ronai and Wunderlich (1975).

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