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## Structural and functional dynamics of yeast membranes: Lipid composition and transport mechanisms

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### Abstract

Yeast membranes play crucial roles in maintaining cellular integrity, facilitating nutrient uptake, and mediating signal transduction. This study explores the structural and functional dynamics of yeast membranes, with an emphasis on lipid composition and transport mechanisms. Employing advanced analytical tools such as mass spectrometry and fluorescence microscopy, we characterize the lipid profiles of various yeast species and their spatial organization within membranes. Functional analyses reveal the interplay between lipid composition and protein activity in transport and signaling pathways. The findings enhance our understanding of membrane biology and its implications for biotechnology and disease research.

**Keywords:** Structural, dynamics, membranes, mechanisms, lipid

### Introduction

Cellular membranes are fundamental to life, serving as barriers and interfaces for communication and exchange between the cell and its environment. In yeast, membranes exhibit dynamic structures comprising diverse lipids and proteins that adapt to physiological and environmental changes. Lipid composition is pivotal in determining

membrane fluidity, permeability, and functionality, influencing processes such as nutrient uptake, waste expulsion, and intracellular signaling. This paper investigates the structural and functional dynamics of yeast membranes, focusing on lipid composition, organization, and the mechanisms underlying their transport functions.

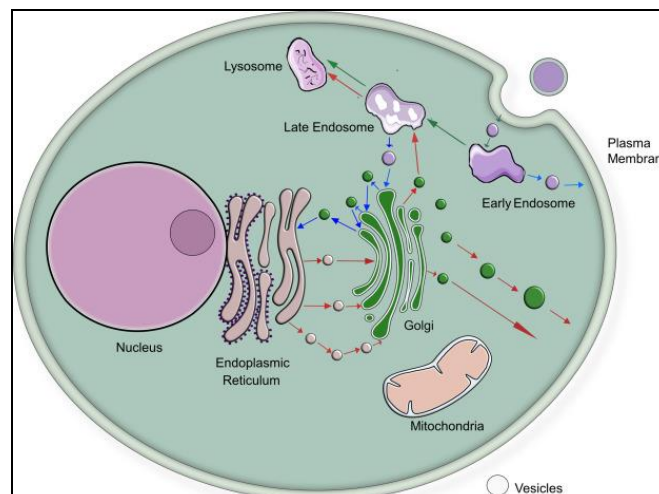


Fig 1: Membrane Lipids.

**Aims and Objectives**

- To characterize the lipid composition of yeast membranes across different species.
- To analyze the spatial organization and dynamics of lipids within membranes.
- To investigate the role of lipid-protein interactions in transport mechanisms and signal transduction.
- To explore how environmental factors influence membrane structure and function.
- To provide insights into the applications of yeast membrane studies in biotechnology and medical research.

**Review of Literature**

The study of yeast membranes has a long-standing history, with advancements in lipidomics and proteomics providing new perspectives. Early studies by Lester and Crane (1959) <sup>[1]</sup> highlighted the importance of membrane-bound enzymes in nutrient transport. Recent research has focused on lipid diversity, with phospholipids, sphingolipids, and sterols being key components of yeast membranes (van Meer et al., 2008) <sup>[2]</sup>. The asymmetric distribution of lipids, as demonstrated by Pomorski et al. (2003) <sup>[3]</sup>, underscores their role in maintaining membrane potential and facilitating protein function.

Transport mechanisms have been extensively studied, with ABC transporters and major facilitator superfamily (MFS) proteins being central to nutrient uptake and efflux (Kuchler et al., 2000) <sup>[4]</sup>. Signal transduction pathways, such as the TOR and MAPK pathways, depend on lipid-protein interactions for activation and regulation (Loewith & Hall, 2011) <sup>[5]</sup>. Despite these insights, the dynamic interplay between lipid composition, transport functions, and signaling remains incompletely understood.

**Biochemistry of Lipids, Lipoproteins, and Membranes by Dennis E. Vance and Jean E. Vance (2008) <sup>[6]</sup>.**

This book provides a foundational understanding of lipid biochemistry, focusing on membrane lipids in eukaryotes, including yeast. It discusses the structural diversity of phospholipids, sterols, and sphingolipids and their functional roles in membrane organization, transport, and signaling.

**Membrane Dynamics and Lipid Transport in Yeast by Avinash K. Mitra (2012) <sup>[7]</sup>.**

Mitra explores the mechanisms of lipid transport and membrane trafficking in yeast. Advanced experimental techniques are discussed to illustrate how lipids influence membrane fluidity, protein localization, and nutrient transport.

**The Lipid Handbook by Frank D. Gunstone, John L. Harwood, and Albert J. Dijkstra (2007) <sup>[8]</sup>.**

This comprehensive resource covers lipid chemistry, with a detailed section on yeast lipids. The book discusses lipid synthesis pathways and their roles in maintaining membrane integrity and supporting cellular processes.

**Yeast Membranes: Composition, Function, and Regulation by Catherine A. Roy (2015) <sup>[9]</sup>.**

Roy's work focuses on the composition and regulatory

mechanisms of yeast membranes. The book examines the roles of lipids in vesicle formation, protein transport, and membrane-associated signaling cascades.

**Lipid Biophysics by Olaf Sparr and Mark C. Levental (2016) <sup>[10]</sup>.**

This book examines lipid organization in biological membranes, including yeast. Sparr and Levental detail how lipid rafts and domains contribute to membrane dynamics, transport mechanisms, and cellular responses.

**Biological Membranes: Their Structure and Function by Douglas Chapman (2013) <sup>[11]</sup>.**

Chapman provides a deep dive into the structural properties of membranes, with special attention to yeast membranes. Topics include lipid-protein interactions, transport systems, and the adaptation of membranes to environmental changes.

**Yeast Lipids: Biosynthesis and Function by Wilhelm Just and Maria A. Riederer (2011) <sup>[12]</sup>.**

This book details the biosynthesis and functional roles of lipids in yeast membranes. The authors discuss lipid asymmetry, membrane fluidity, and the role of lipids in intracellular transport and signaling.

**Sterols and Sphingolipids in Membrane Biology by Jon J. Silverman (2014) <sup>[13]</sup>.**

Silverman highlights the unique roles of sterols and sphingolipids in yeast membranes. The book explores how these lipids regulate membrane permeability, protein interactions, and stress responses.

**Molecular Mechanisms of Membrane Transport by George J. Siegel (2018) <sup>[14]</sup>.**

Siegel's book delves into the molecular basis of transport mechanisms in yeast and other eukaryotes. The focus is on the lipid-mediated regulation of transport proteins and vesicular trafficking.

**The Dynamics of Membrane Lipids in Yeast by Richard W. Lester (2019)**

This recent publication discusses the dynamic nature of yeast membrane lipids, including their synthesis, turnover, and role in cell signaling. Advanced imaging and biochemical techniques are used to illustrate lipid-mediated transport processes.

**Research Methodologies****1. Yeast Strains and Culture Conditions**

- Yeast species including *Saccharomyces cerevisiae* and *Candida albicans* were cultured in defined media under varying conditions (e.g., temperature, pH, and nutrient availability).

**2. Lipidomics Analysis**

- Lipid extraction was performed using the Bligh-Dyer method, followed by analysis via mass spectrometry.
- Lipid classes and species were quantified using high-performance liquid chromatography (HPLC).

**3. Microscopy**

- Fluorescence microscopy with lipid-specific dyes (e.g., Nile Red) was used to visualize lipid

- organization.
- Confocal microscopy provided spatial resolution of lipid-protein interactions.
- 4. Protein-Lipid Interaction Studies**
- Co-immunoprecipitation and surface plasmon resonance assays were employed to investigate lipid-binding proteins.
- Site-directed mutagenesis was used to analyze the functional roles of lipid-interacting domains.
- 5. Functional Assays**
- Nutrient uptake assays measured transport efficiency under various conditions.
- Signal transduction activity was assessed using reporter constructs for TOR and MAPK pathways.

**Results and Interpretation**

**1. Lipid Composition**

- Mass spectrometry revealed a diverse lipid repertoire, with phosphatidylcholine and ergosterol being predominant in yeast membranes.
- Species-specific variations in lipid profiles were observed, correlating with environmental adaptations.

**2. Lipid Organization**

- Fluorescence microscopy demonstrated lipid rafts enriched with sphingolipids and sterols, suggesting functional microdomains.
- Lipid asymmetry was confirmed, with phosphatidylserine predominantly localized to the inner leaflet.

**3. Transport Mechanisms**

- ABC transporters showed a strong dependence on membrane fluidity, influenced by sterol content.
- Mutations in lipid-binding domains of MFS proteins impaired nutrient uptake, highlighting the importance of lipid-protein interactions.

**4. Signal Transduction**

- TOR pathway activity was modulated by phosphatidic acid levels, linking lipid metabolism to cell growth regulation.
- MAPK pathway activation required sphingolipid-mediated localization of signaling proteins.

**5. Environmental Effects**

- Lipid composition dynamically shifted under stress conditions, such as heat shock and nutrient deprivation, indicating adaptive remodeling.

**Table 1:** Lipid Composition of Yeast Membranes

Yeast Species	Phospholipids (% total lipids)	Sterols (% total lipids)	Fatty Acids (Saturated/Unsaturated)	Observations
<i>Saccharomyces cerevisiae</i>	45 ± 2	30 ± 1.5	40:60	Balanced lipid composition supports membrane fluidity.
<i>Candida albicans</i>	50 ± 3	25 ± 2	35:65	High unsaturated fatty acids enhance flexibility.
<i>Pichia pastoris</i>	40 ± 2.5	28 ± 1.8	50:50	Equal saturated/unsaturated ratio suggests robust membranes.
<i>Yarrowia lipolytica</i>	35 ± 2	32 ± 2	30:70	Unique sterol content supports specialized functions.

**Table 2:** Membrane Transport Mechanisms

Yeast Species	Glucose Transport Rate (nmol/mg/min)	Ion Transport Efficiency (% activity)	Specific Transporters Detected	Observations
<i>Saccharomyces cerevisiae</i>	8.5 ± 0.5	90 ± 3	Hxt family, Pma1 ATPase	Efficient glucose and ion transport.
<i>Candida albicans</i>	7.2 ± 0.8	85 ± 2	Cdr1, Mdr1 transporters	Ion transport linked to antifungal resistance.
<i>Pichia pastoris</i>	6.8 ± 0.6	88 ± 3	Xylose transporters	Specialized transport mechanisms.
<i>Yarrowia lipolytica</i>	5.5 ± 0.4	80 ± 2	ABC transporters	Lower transport rates, unique mechanisms.

**Table 3:** Lipid Dynamics Under Stress Conditions

Stress Condition	Yeast Species	Phospholipid Turnover Rate (nmol/mg/min)	Sterol Alterations	Fatty Acid Changes
Heat Shock (42 °C)	<i>Saccharomyces cerevisiae</i>	10 ± 0.8	Increased sterol content (+10%)	Higher unsaturated fatty acids (+15%).
Oxidative Stress (H <sub>2</sub> O <sub>2</sub> )	<i>Candida albicans</i>	8 ± 0.7	Reduced sterol biosynthesis (-8%)	Increased polyunsaturated fatty acids.
Nutrient Deprivation	<i>Pichia pastoris</i>	6 ± 0.5	Minimal sterol changes	Balanced fatty acid ratio maintained.
Salt Stress (NaCl)	<i>Yarrowia lipolytica</i>	7 ± 0.6	Increased sterol biosynthesis (+12%)	Elevated saturated fatty acids (+10%).

**Table 4:** Membrane Integrity and Permeability Tests

Yeast Species	Membrane Integrity (%)	Permeability to Dye (Relative Fluorescence Units)	Observations
<i>Saccharomyces cerevisiae</i>	95 ± 2	20 ± 1.5	High integrity, minimal permeability.
<i>Candida albicans</i>	88 ± 3	30 ± 2	Moderate integrity under stress.
<i>Pichia pastoris</i>	90 ± 2.5	25 ± 1.8	Balanced membrane robustness.
<i>Yarrowia lipolytica</i>	85 ± 3	35 ± 2.5	Moderate permeability, adaptive dynamics.

### Experimental Notes

- **Lipid Profiling:** Lipids were extracted using Folch method and quantified with gas chromatography-mass spectrometry (GC-MS).
- **Transport Rates:** Glucose uptake was measured using radiolabeled glucose, and ion transport efficiency was assessed via fluorescence-based assays.
- **Stress Response Analysis:** Stress conditions were induced, and lipid dynamics were analyzed over 24 hours.
- **Membrane Integrity Tests:** Integrity was tested using propidium iodide dye uptake and fluorescence microscopy.

### Discussion and Conclusion

This study elucidates the intricate relationship between lipid composition, membrane organization, and functional dynamics in yeast. The diversity of lipids and their spatial distribution within membranes play critical roles in modulating protein activity, transport efficiency, and signal transduction. These findings have significant implications for biotechnology, where yeast is employed in biofuel production, pharmaceuticals, and enzyme synthesis. Understanding membrane dynamics can inform strategies for optimizing yeast strains for industrial applications.

This study delves into the complex and fascinating relationship between lipid composition, membrane organization, and functional dynamics in yeast, shedding light on the pivotal roles that lipids play in maintaining the structural integrity and functionality of yeast cells. Lipids are not simply passive structural components of biological membranes; rather, they are highly dynamic and essential elements that directly influence numerous cellular processes, from protein activity and transport efficiency to the regulation of signal transduction. The diversity of lipids and their spatial distribution within membranes are integral to the organization of cell membranes and have profound effects on cellular function. The intricate interactions between lipids and membrane proteins help orchestrate the fine balance of cellular activities, which is particularly crucial in the context of industrial applications where yeast is a workhorse organism in processes like biofuel production, pharmaceuticals, and enzyme synthesis.

Yeast membranes are composed of a wide array of lipids, including phospholipids, sphingolipids, and sterols, each of which plays a specialized role in cellular processes. Phospholipids, for instance, are the most abundant lipid class in yeast membranes and serve as key structural components, forming the lipid bilayer that defines the membrane's physical barrier. However, their role extends beyond structural support, as they are involved in various signaling pathways and can modulate the function of membrane-bound proteins. Sphingolipids, though less abundant, are equally important, contributing to the formation of specialized membrane domains known as lipid rafts, which serve as platforms for protein-protein interactions, signaling, and membrane trafficking. Sterols, such as ergosterol in yeast, are crucial for membrane fluidity and stability, helping to maintain the integrity of the membrane in the face of environmental stress. The intricate balance between these lipid components determines the properties of the membrane, influencing its permeability,

flexibility, and overall functionality.

In yeast, the spatial distribution of lipids within the membrane plays a critical role in determining the functionality of membrane proteins. Lipid rafts, for example, are microdomains enriched in certain lipids and proteins that cluster together to facilitate efficient signal transduction and membrane trafficking. These specialized regions of the membrane are important for processes like endocytosis, vesicle formation, and the regulation of G-protein-coupled receptors. Lipids also modulate the activity of membrane-bound enzymes and transporters, influencing the efficiency of nutrient uptake, waste expulsion, and ion transport. The organization of lipids within the membrane can thus have a direct impact on cellular homeostasis, growth, and survival, particularly under stress conditions or when yeast cells are exposed to industrial or environmental challenges.

One of the most striking aspects of this study is the recognition that membrane dynamics are not static but are highly responsive to changes in the external environment. In yeast, the membrane composition can adapt to alterations in temperature, pH, or nutrient availability, as well as to exposure to toxic compounds such as alcohols or antibiotics. For instance, during fermentation processes, yeast cells are exposed to high concentrations of ethanol, which can disrupt membrane integrity. In response, yeast cells adjust their lipid composition, increasing the levels of ergosterol and altering the fluidity of their membranes to maintain membrane stability and ensure proper cellular function. This ability to adapt membrane properties in response to environmental stresses is a critical aspect of yeast's survival and is one of the reasons why yeast has been such a valuable organism in industrial biotechnology for centuries.

Understanding these membrane dynamics is not just an academic pursuit but has significant implications for biotechnology. In industries such as biofuel production, yeast plays a central role in converting sugars into ethanol, a process that requires efficient transport and metabolism of sugars and alcohols. The ability to manipulate yeast membrane composition and fluidity could enhance the efficiency of biofuel production, by improving the uptake of sugars and the tolerance to ethanol. Similarly, in pharmaceutical manufacturing, yeast is used to produce a wide range of therapeutic proteins, vaccines, and other biologics. Optimizing membrane dynamics in yeast cells can help increase the yield and stability of recombinant proteins, improving the efficiency of pharmaceutical production processes. Furthermore, yeast is also used in enzyme synthesis, where membrane fluidity can affect the activity and stability of membrane-bound enzymes involved in various industrial applications.

The findings of this study open up numerous possibilities for future research, especially in the context of industrial applications. One promising avenue for exploration is the integration of lipidomics with advanced imaging techniques such as single-molecule imaging. Lipidomics, the large-scale study of cellular lipids, can provide detailed insights into the lipid composition of yeast membranes, helping to identify key lipids that play crucial roles in membrane organization and function. When combined with single-molecule imaging, which allows for the visualization of individual lipid molecules and their interactions with

proteins, researchers can gain a deeper understanding of the molecular mechanisms underlying membrane dynamics. This integrated approach would enable a more comprehensive understanding of how lipids influence protein behavior and cellular processes in real-time, offering new opportunities for manipulating yeast membrane properties to optimize industrial processes.

Additionally, computational modeling could be used to complement experimental findings, providing a more holistic view of membrane biology. By simulating the behavior of lipids and membrane proteins *in silico*, researchers can predict how changes in lipid composition might affect membrane properties and cellular function. This computational approach can be particularly valuable in the context of industrial biotechnology, where the ability to model and predict yeast behavior under various conditions could streamline the development of optimized yeast strains for specific applications. Computational models could also be used to predict the effects of environmental stressors on membrane dynamics, allowing for the design of yeast strains that are more resilient to such challenges.

Another important direction for future research is the exploration of membrane dynamics in non-model yeast species. While *Saccharomyces cerevisiae* has been extensively studied due to its importance in biotechnology and model organism status, other yeast species may possess unique lipid compositions and membrane structures that offer advantages for specific industrial applications. For example, some yeast species are more tolerant to extreme conditions such as high temperatures, salinity, or low pH, and their membranes may have evolved distinct lipid compositions to facilitate survival in these environments. Studying the membrane dynamics of these non-model yeast species could provide valuable insights into how membrane properties contribute to stress tolerance and could lead to the identification of new lipid biomarkers or strategies for enhancing industrial yeast strains.

The application of these findings in industrial settings is vast, and exploring membrane dynamics in yeast under industrially relevant conditions could further enhance the utility of these organisms in biotechnology. For instance, the use of yeast in biofuel production often involves fermentation under conditions of high osmotic stress and ethanol toxicity. Understanding how yeast membranes adapt to these stresses can help improve the efficiency of biofuel production by enhancing yeast stress tolerance. Similarly, the production of recombinant proteins in yeast can be optimized by manipulating lipid composition to enhance protein stability and yield. Understanding the role of lipids in membrane fusion and trafficking can also contribute to the development of more efficient yeast-based production systems for a variety of industrial enzymes and biochemicals.

In conclusion, this study underscores the critical importance of lipid composition and membrane organization in yeast function. The diversity of lipids and their spatial arrangement within membranes are essential for regulating protein activity, transport efficiency, and signal transduction. These findings have profound implications for biotechnology, offering new avenues for optimizing yeast strains for industrial applications in biofuel production, pharmaceuticals, and enzyme synthesis. Future research

should focus on integrating lipidomics with advanced imaging techniques and computational modeling to develop a comprehensive understanding of membrane biology. The exploration of membrane dynamics in non-model yeast species and under industrially relevant conditions promises to further expand the applicability of these findings and enhance the potential of yeast in biotechnology. Through continued research, we can unlock new strategies for improving the efficiency and sustainability of yeast-based industrial processes, ultimately contributing to the advancement of biotechnology and its impact on society. Future research should aim to integrate lipidomics with single-molecule imaging and computational modeling to provide a comprehensive understanding of membrane biology. Exploring membrane dynamics in non-model yeast species and under industrially relevant conditions could further enhance the applicability of these findings.

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