



# “MODELLING LIPID MEMBRANE AND MEMBRANE PROTEIN PROCESSES”

**Dr. Juan José Nogueira**

MoBioChem

Department of Chemistry,  
Universidad Autónoma de Madrid,  
Calle Francisco Tomás y Valiente, 7, Madrid, Spain  
[juan.nogueira@uam.es](mailto:juan.nogueira@uam.es)

**September 24, 2021 (Friday)**  
**12pm (BRT time) – Google Meet**

**ORGANIZATION:**

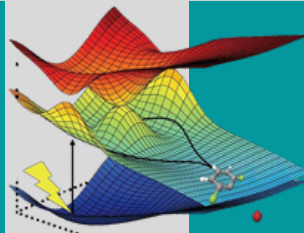
**Prof. Dr. Antonio Carlos Borin**

Instituto de Química, Universidade de São Paulo (USP), SP, Brazil

**INFORMATION AND REGISTRATION:**

**[ancborin@iq.usp.br](mailto:ancborin@iq.usp.br)**

**Registration:** send a message to [ancborin@iq.usp.br](mailto:ancborin@iq.usp.br)  
with the words “Juan José Nogueira – Virtual” on the “subject”  
**Deadline:** September 23, 2021 (Thursday), 06pm (BRT time)



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## **ABSTRACT**

### **Modelling Lipid Membrane and Membrane Protein Processes**

Juan J. Nogueira

Department of Chemistry, Universidad Autónoma de Madrid,  
Calle Francisco Tomás y Valiente, 7, Madrid, Spain  
[juan.nogueira@uam.es](mailto:juan.nogueira@uam.es)

Lipid membranes and membrane proteins are key biological assemblies involved in many relevant functions, including cell protection, molecular transport, and signal transduction. A deep understanding of these biological processes can be achieved by means of computer simulations. However, the simulation of membrane processes is a very complex task which requires the combination of different quantum and classical mechanical techniques within a dynamic framework. In this talk, I will discuss about the simulation of two different membrane processes recently published by our group: the passive transport of the anticancer drug cisplatin through a model lipid bilayer (1,2) and the binding of photoswitches to a human voltage-gated ion channel (3).

(1) Lorena Ruano, Gustavo Cárdenas, and Juan J. Nogueira. *ChemPhysChem*, 2021, 22, 1–12.

(2) Gustavo Cardenas, Álvaro Pérez-Barcia, Marcos Mandado, and Juan J. Nogueira. *PCCP*, 2021. Advanced manuscript. DOI: 10.1039/D1CP03382D.

(3) Vito F. Palmisano, Carlos Gómez-Rodellar, Hannah Pollak, Gustavo Cárdenas, Ben Corry, Shirin Faraji, and Juan J. Nogueira. *PCCP*, 2020, 23, 3552–3564.