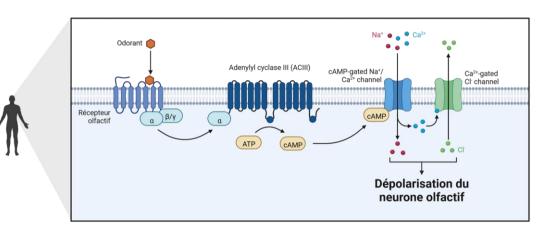


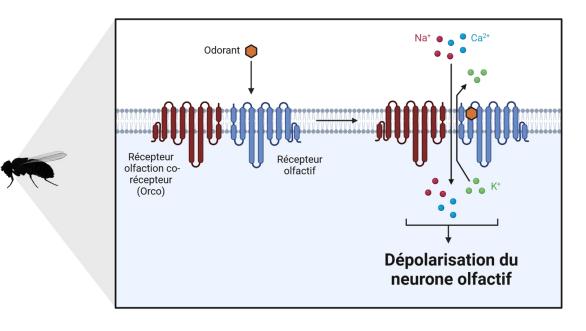
Numerical models decipher odorant binding and receptor activation at the molecular level

UNIVERSITÉ CÔTE D'AZUR ChEmoSim

O Dr. Jérémie Topin

Nature created two olfaction systems





Massive yet grossly underestimated global costs of invasive insects

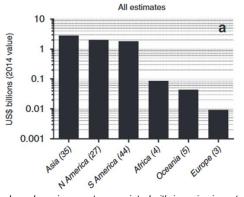


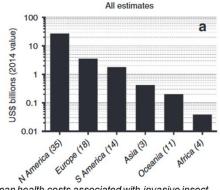
Dr. J. Pacalon







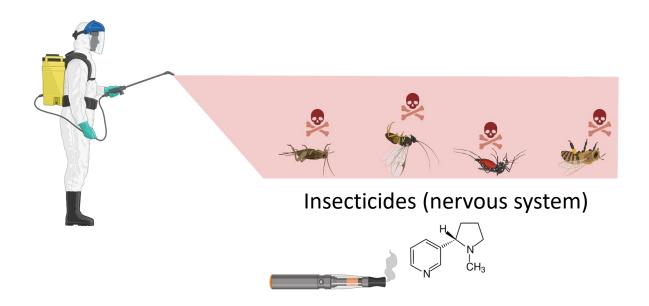




Targeting Insect Olfaction



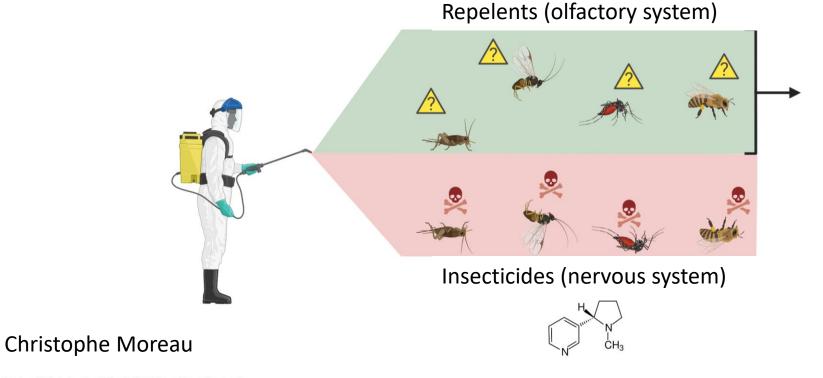
Dr. J. Pacalon



Targeting Insect Olfaction

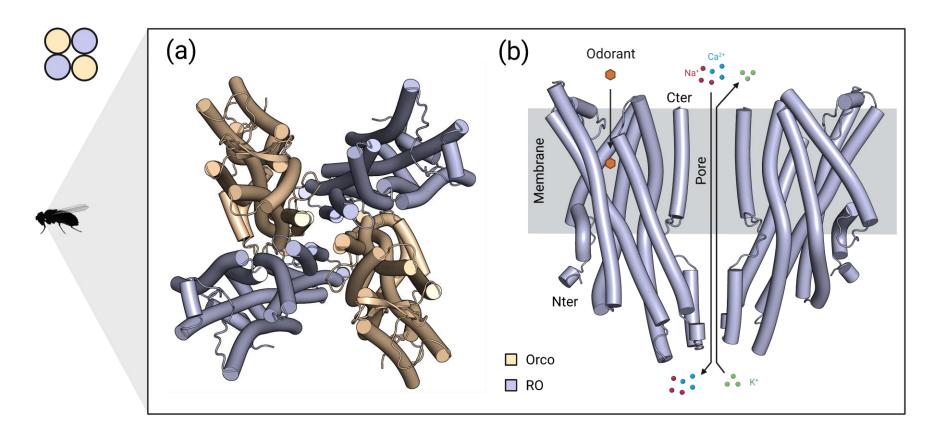


Dr. J. Pacalon



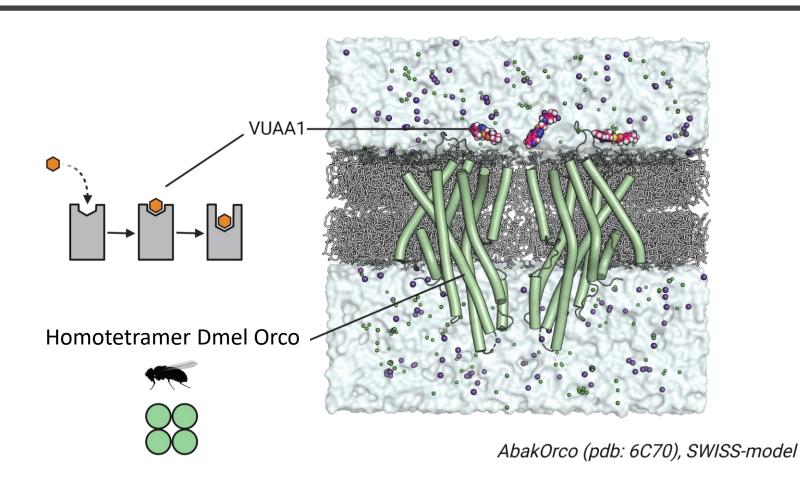


Quaternary structure of Insect OR

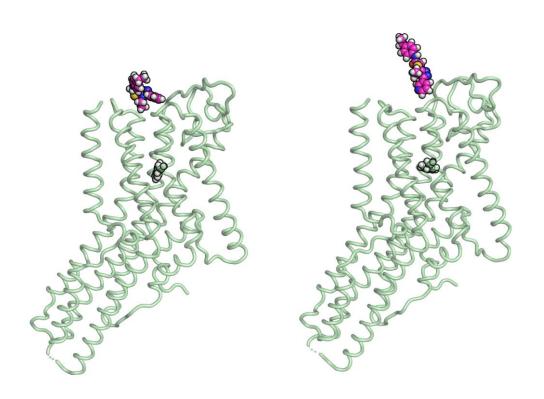


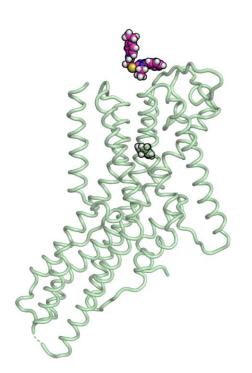
CAN WE IDENTIFY THE BINDING POCKET OF A COMPLEX MOLECULAR ASSEMBLY?

Binding Simulation

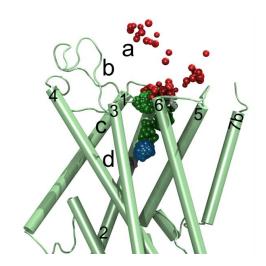


Binding Simulation / Results

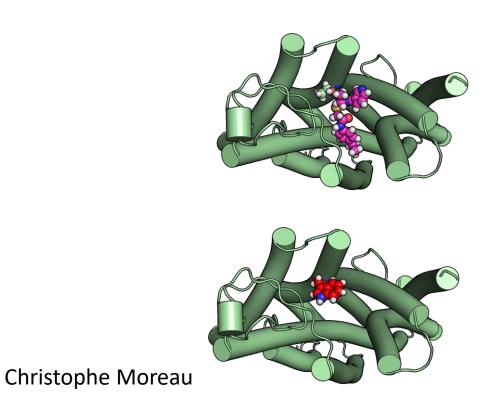


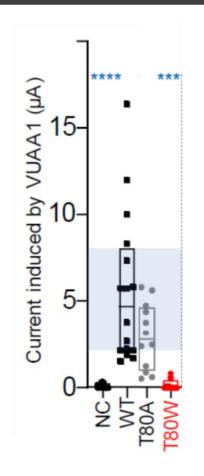


22 independent simulations (31 μs) 288 000 atoms (ff14SB/lipid14/gaff2)



In Vitro evaluation Entry



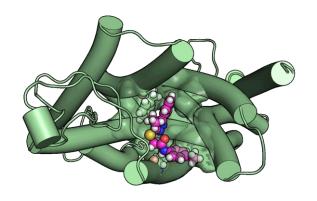


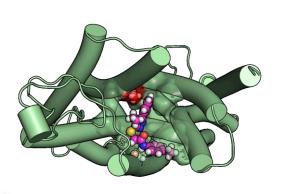


A molecular vision of life

In Vitro evaluation Binding Site

Current induced by VUAA1 (μΑ)

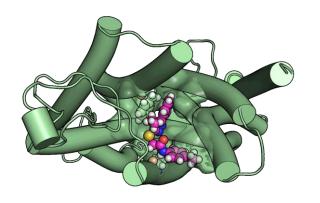


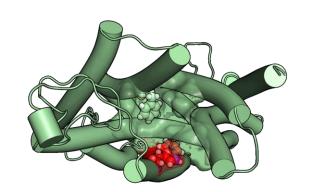


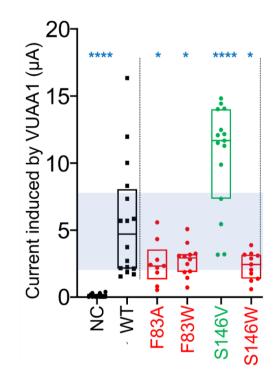


Christophe Moreau

In Vitro evaluation Binding Site



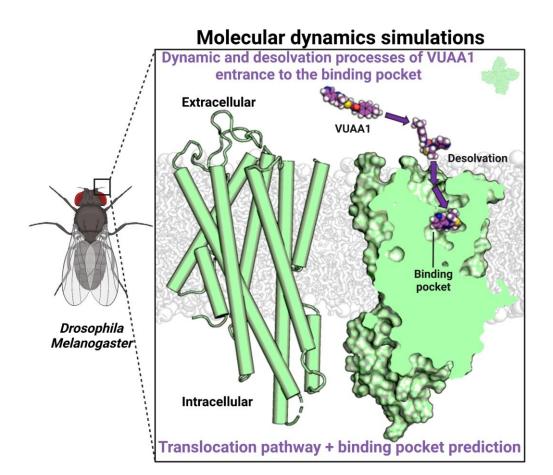


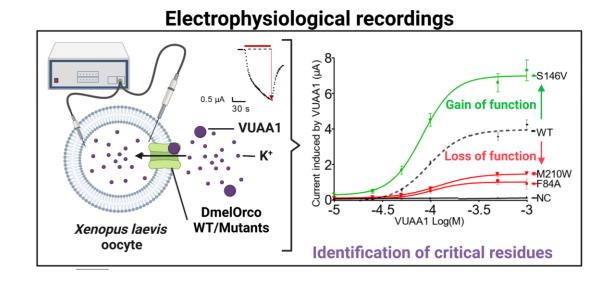




Christophe Moreau

Translocation pathway in Orco



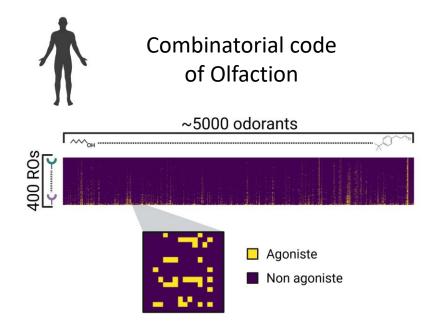


CAN WE PREDICT ACTIVATION OF A PROTEIN BY A MOLECULE WITHOUT A KNOWN PROTEIN STRUCTURE?

Cracking the Combinatorial Code of Olfaction



M. Lalis & M. Hladiš



The majority of Odorants-ORs pairs remains unknown

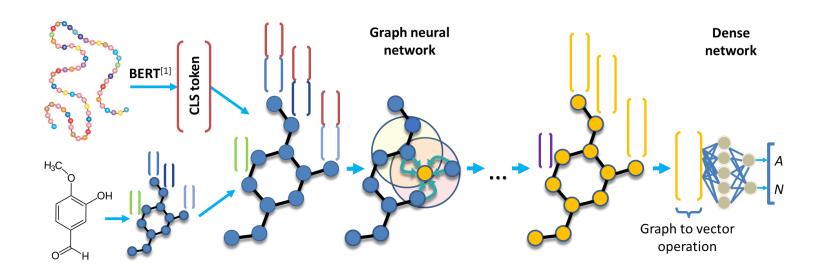
No experimental structure for all ORs



Cracking the Combinatorial Code of Olfaction



M. Lalis & M. Hladiš



Matching receptor to odorant with protein language and graph neural networks

Matej Hladiš, Maxence Lalis, Sebastien Fiorucci, Jérémie Topin

