### **SCHOOL:**

## COMPUTER SIMULATIONS OF BIOLOGICAL MEMBRANES

## FREE ENERGY CALCULATIONS OF BIOMOLECULAR SYSTEMS

**VENUE:** University of Los Andes, Bogotá, Colombia **DATE:** August 13-17, 2018

## PROGRAM OVERVIEW

|               | Monday (13.08)             | Tuesday (14.08)               | Wed. (15.08)       | Thursday (16.08)               | Friday (17.08)                         |
|---------------|----------------------------|-------------------------------|--------------------|--------------------------------|--|
| 8:00 – 9:00   | Registration               |                               |                    |                                |  |
| 9:00 – 9:30   | Welcome                    |                               |                    | Gian Pietro Miscione           |  |
| 9:30 - 10:30  | Bert de Groot              | Thomas Walz<br>(webinar) ML-C | Chad Leidy         | Gian Pietro Miscione           | Bert de Groot<br>ML-C                  |
|               |                            |                               |                    | Contributed talk 4             | WIE C                                  |
| 10:30 – 11:00 | Coffee break               |                               |                    |                                |  |
| 11:00 – 12:00 | Helgi Ingólfson            | Gerhard Hummer<br>ML-C        | Contributed talk 2 | Esteban Vohringer-<br>Martínez | Vytas Gapsys                           |
|               |                            |                               | Contributed talk 3 |                                | ML-C                                   |
| 12:00 – 14:00 | Lunch                      |                               | Lun                |                                | nch                                    |
| 14:00 – 15:00 | Gerhard Hummer             | Bert de Groot<br>ML-C         |                    | Andrés González                | Hands on<br>ML-C                       |
| 15:00 – 15:30 | Vytas Gapsys<br>ML-C       | Helgi Ingólfson               | Free program       | Coffee break                   |  |
| 15:30 – 16:00 | Contributed talk 1<br>ML-C | ML-C                          |                    | Contributed talk 5             |  |
| 16:00 – 16:30 | Coffee break               |                               |                    | Contributed talk 6             | Closing remarks &<br>departure<br>ML-C |
| 16:30 – 17:00 | Heading to computer room   |                               |                    | Heading to computer room       |  |
| 17:00 – 19:00 | Hands on                   | Hands on<br>ML-C              |                    | Poster session                 |  |

# PROGRAM IN DETAIL

| PART 1: COMPUTER SIMULATIONS OF BIOLOGICAL MEMBRANES |          |   |   |  |
|--|----------|---|---|--|
| Monday, Augi   | ıst 13   |   |   |  |
| Time   | Room     |   |   |  |
| 8:00 – 9:00  |          | Registration  |   |  |
| 9:00 - 9:30  |          | Welcome   |   |  |
| 9:30 – 10:30   |          | Bert de Groot<br>(MPIBPC, Göttingen)                            | The molecular dynamics of solute permeation through biological membranes and membrane proteins                  |  |
| 10:30 - 11:00  |          | Coffee break  |   |  |
| 11:00 – 12:00  |          | Helgi Ingólfson<br>(LLNL, Livermore)                            | Coarse-grained simulations: introduction and application to membranes   |  |
| 12:00 – 14:00  |          | Lunch   |   |  |
| 14:00 – 15:00  |          | Gerhard Hummer<br>(MPIB, Frankfurt)                             | tba   |  |
| 15:00 – 15:30  |          | Vytautas Gapsys (MPIBPC,<br>Göttingen)                          | tba   |  |
| 15:30 – 16:00  |          | Contributed talk 1:<br>Florian Franz<br>(IWR, Heidelberg)       | Stability of Biological Membranes upon Mechanical Indentation   |  |
| 16:00 – 16:10  |          | Photo   |   |  |
| 16:10 – 16:40  |          | Coffee break  |   |  |
| 16:40 – 17:00  |          | Heading to computer room  |   |  |
| 17:00 – 19:00  |          | Hands on: all-atom MD simulations of biological membranes       |   |  |
| 19:00 –  |          | Invited speakers dinner   |   |  |
| Tuesday, Augu  | ust 14   |   |   |  |
| 9:30 – 10:30   |          | Webinar, Thomas Walz<br>(Rockefeller, New York)                 | Electron Microscopy Approaches to Studying Lipid-Protein<br>Interactions  |  |
| 10:30 – 11:00  |          | Coffee break  |   |  |
| 11:00 – 12:00  |          | Gerhard Hummer (MPIB,<br>Frankfurt)                             | <i>tba</i>  |  |
| 12:00 – 14:00  |          | Lunch   |   |  |
| 14:00 – 15:00  |          | Bert de Groot (MPIBPC,<br>Göttingen)                            | Permeation, selectivity and gating of ion channels studied by computational electrophysiology                   |  |
| 15:00 – 16:00  |          | Helgi Ingolfson<br>(LLNL, Livermore)                            | Capturing biological membranes in silico at different compositional complexity                                  |  |
| 16:00 - 16:40  |          | Coffee break  |   |  |
| 16:40 – 17:00  |          | Heading to computer room  |   |  |
| 17:00 – 19:00  |          | Hands on: coarse-grained MD simulations of biological membranes |   |  |
| Wednesday, Au  | ugust 15 |   |   |  |
| 9:30 – 10:30   |          | Chad Leidy<br>(UANDES, Bogotá)                                  | Integrating Computational and Experimental Approaches to InvestigateAntimicrobial Peptide Activity in Membranes |  |
| 10:30 - 11:00  |          | Coffee break  |   |  |
| 11:00 – 11:30  |          | Contributed talk 2:   | Exponential consensus ranking improves outcome in docking and   |  |

|                     | Karen Palacio Rodríguez<br>(UDEA, Medellín)                       | receptor ensemble docking  |  |  |  |  |
|---------------------|---|--|--|--|--|--|
| 11:30 – 12:00       | Contributed talk 3:<br>Sergio Mares-Samano<br>(UNAM, México City) | Molecular dynamics studies of the solvation effects of water and trifluoroethanol on the peptides pandinin-2 (pin2) and pin2-GVG |  |  |  |  |
| 12:00 –             | free program  |  |  |  |  |  |
| PART 2) FREE EN     | ERGY CALCULATIONS OF  | BIOLOGICAL SYSTEMS   |  |  |  |  |
| Thursday, August 16 | Thursday, August 16   |  |  |  |  |  |
| 9:00 – 10:00        | Gian Pietro Miscione<br>(UANDES, Bogotá)                          | Mechanism of L-Tryptophan Prenylation, Catalyzed by FgaPT2: a Computational Study  |  |  |  |  |
| 10:00 – 10:30       | Contributed talk 4:<br>Hugo J. Bohórquez<br>(FIDIC, Bogotá)       | The frozen molecule approach in peptide-protein binding energies   |  |  |  |  |
| 10:30 – 11:00       | Coffee break  |  |  |  |  |  |
| 11:00 – 12:00       | Esteban Vohringer-<br>Martínez (CONCEPCIÓN,<br>Concepción)        | Force field improvement with ab-initio methods   |  |  |  |  |
| 12:00 – 14:00       | Lunch   |  |  |  |  |  |
| 14:00 – 15:00       | Andrés González<br>(UANDES, Bogotá)                               | Applications of product design departing from free energy calculations   |  |  |  |  |
| 15:00 – 15:30       | Coffee break  |  |  |  |  |  |
| 15:30 – 16:00       | Contributed talk 5:<br>Angélica Sandoval<br>(UANDES, Bogotá)      | Specific at one side unspecific at the other: the interaction of blood proteins with extracellular DNA                           |  |  |  |  |
| 16:00 – 16:30       | Contributed talk 6:<br>Stefany Botero Rodríguez<br>(UNAL, Bogotá) | Computational Tracing of Protein and ncRNA epitopes inside<br>Dengue Virus   |  |  |  |  |
| 16:30 – 17:00       | Heading to poster room  |  |  |  |  |  |
| 17:00 – 19:00       | Poster session (with wine and finger food)                        |  |  |  |  |  |
| Friday, August 17   |   |  |  |  |  |  |
| 9:30 – 10:30        | Bert de Groot<br>(MPIBPC, Göttingen)                              | Introduction to alchemical free energy calculations  |  |  |  |  |
| 10:30 – 11:00       | Coffee break  |  |  |  |  |  |
| 11:00 – 12:00       | Vytas Gapsys<br>(MPIBPC, Göttingen)                               | tba  |  |  |  |  |
| 12:00 – 14:00       | Lunch   |  |  |  |  |  |
| 14:00 – 16:00       | Hands on: Free energy calculations                                |  |  |  |  |  |
| 16:00 – 17:00       | Closing remarks and departure                                     |  |  |  |  |  |

## **POSTERS**

| PU3 | TERS  |   |  |
|-----|---|---|--|
| 1   | Diego Gomez Hernandez<br>(UDISTRITAL, Bogotá)   | In silico evaluation of the potential afinity of anandamide analogues for CB1 receptor.   |  |
| 2   | Carlos Pinilla<br>UNORTE, Barranquilla          | FeS all0ys at conditions of the Earth Interior  |  |
| 3   | Juan David Orjuela<br>UANDES, Bogotá            | Effect of the lipid-protein interaction on the assembly of the platelet Glycoprotein-IB   |  |
| 4   | Markus Kurth<br>Heidelberg U. Heidelberg        | Probing the lipid environment of the Metabotropic glutamate receptor 2  |  |
| 5   | Laura Pinto Herrera<br>UANDES, Bogotá           | Effect of particulate matter on the lung surfactant at a high altitude city: Review   |  |
| 6   | Johann Bedoya Cardona<br>UNAL, Medellín         | Applicability of some force field parameters of biomolecules to model the surface tension of hexadecane   |  |
| 7   | Yaritza Fortaleché<br>Rodríguez<br>UNAL, Bogotá | Agonist, antagonist or super-activator effect of different ligands in the receptor-ligand interaction of the TraR protein   |  |
| 8   | Paula Giraldo Hincapie<br>UNAL, Medellín        | Understanding insecticide peptides: Modeling peptide-membrane interactions with Molecular Dynamics  |  |
| 9   | Jessica Martinez<br>UAN, Bogotá                 | Does Using Virtual Sites in Proteins Affect the Protein Fold?   |  |
| 10  | Juan Zarate Moreno<br>SENA, Bogotá              | Kinetic models for chitosan membranes on biological systems, a computational approach.  |  |
| 11  | Cesar Quintana Cataño<br>UANDES, Bogotá         | Force spectroscopy of T4 bacteriophage adhesion during infection.   |  |
| 12  | Santiago Agudelo Gómez<br>UDEA, Medellín        | Preferred sites of glycosaminoglycans in the envelope protein of the Zika virus   |  |
| 13  | Alberto Mario Castillo<br>UBJTL, Bogotá         | Spring Network Model of Trypanosoma Cruzi bloodstream trypomastigote  |  |
| 14  | Gesivaldo Santos<br>State U. Bahia              | Autism Spectro Disorder & Alzheimer Disease: different conditions linked by mTOR pathway  |  |
| 15  | Helman Amaya Espinosa<br>UANDES, Bogotá         | Cooperative dynamics of self-interacting Biopolymers in blood elucidated by Coarse-<br>Grained Brownian Dynamics Simulations  |  |
| 16  | Valeria Mejía Restrepo<br>UANDES, Bogotá        | The effect of G1324A and G1324S mutations on the complex formed by the von Willebrand factor A1 domain and the glycoprotein Iba-platelet receptor, studied through molecular dynamics simulations and free energy calculations. |  |
| 17  | Andrés Cifuentes López<br>UDISTRITAL, Bogotá    | In silico evaluation of the potential afinity of anandamide analogues for CB1 receptor  |  |
| 18  | Jhoan Ortiz Giron<br>UDEA, Medellín             | A novel methodology for cryo-EM map validation  |  |
| 19  | Adrian Orjuela<br>UNAL, Bogotá                  | tba   |  |
|     |   |   |  |