

**SCHOOL:**  
**COMPUTER SIMULATIONS OF BIOLOGICAL MEMBRANES**  
**&**  
**FREE ENERGY CALCULATIONS OF BIOMOLECULAR SYSTEMS**

**DATE:** August 13-17, 2018

**VENUE:** University of Los Andes, Bogotá, Colombia

**WEB:** <https://mptg-cbp.github.io/teaching/mem-fe-2018/>

**ORGANIZER:**  
 Max Planck Tandem Group in  
 Computational Biophysics



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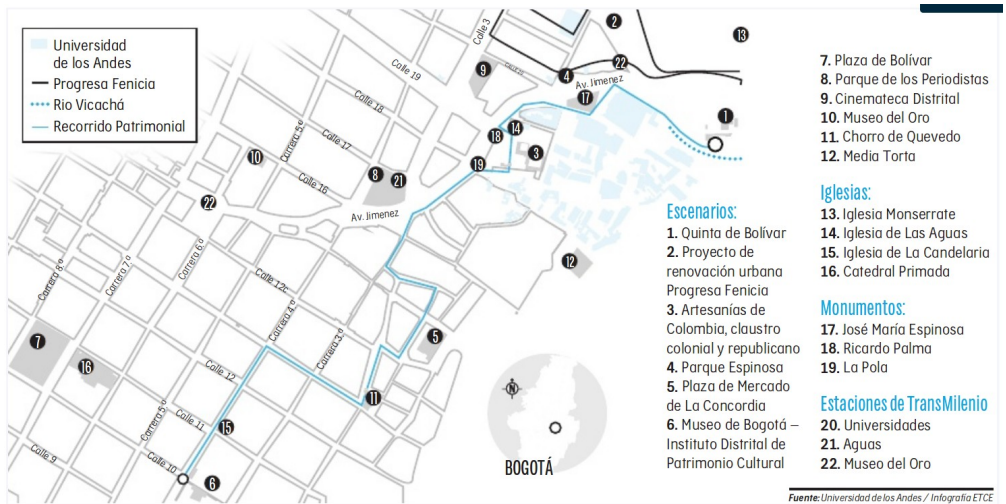
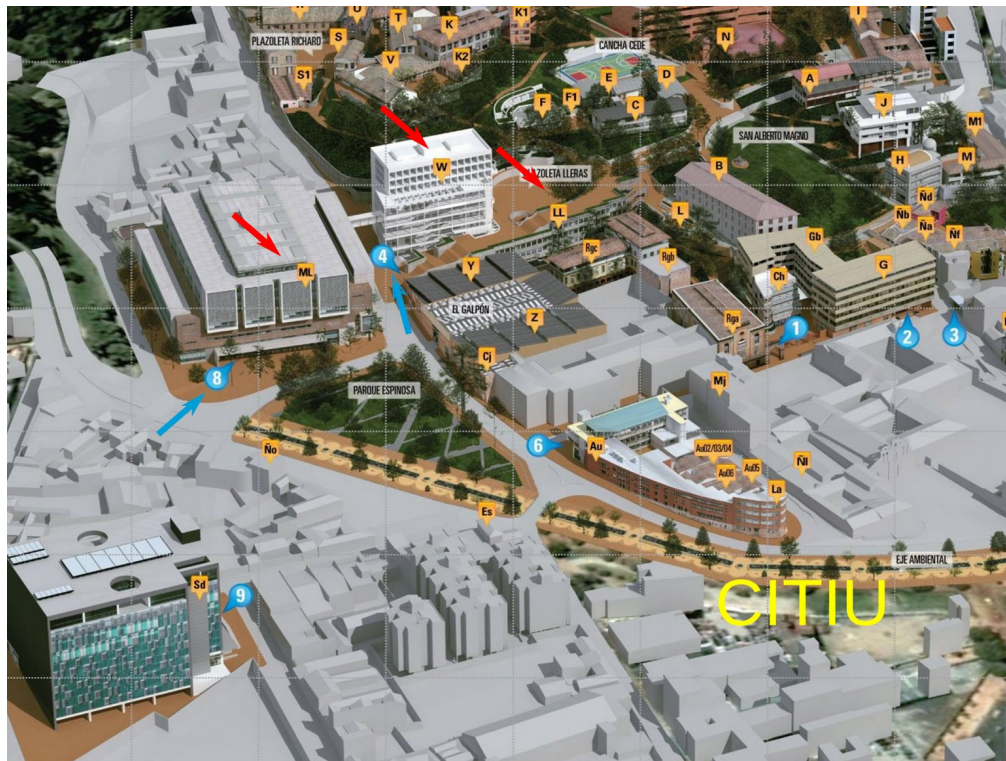
MAPS

**Buildings** (red arrows): W, ML, and LL

**Entrances** (blue arrows):

Entrance 4. “Lleras” Address: Calle 19A N° 1 - 37 Este

Entrance 8. “Mario Laserna” Address: Cra. 1 Este N° 19A - 40



- 7. Plaza de Bolívar
- 8. Parque de los Periodistas
- 9. Cinemateca Distrital
- 10. Museo del Oro
- 11. Chorro de Quevedo
- 12. Media Torta

**Iglesias:**

- 13. Iglesia Monserrate
- 14. Iglesia de Las Aguas
- 15. Iglesia de La Candelaria
- 16. Catedral Primada

**Monumentos:**

- 17. José María Espinosa
- 18. Ricardo Palma
- 19. La Pola

**Estaciones de TransMilenio**

- 20. Universidades
- 21. Aguas
- 22. Museo del Oro

**Escenarios:**

- 1. Quinto de Bolívar
- 2. Proyecto de renovación urbana Progresía Fenicia
- 3. Artesanías de Colombia, claustro colonial y republicano
- 4. Parque Espinosa
- 5. Plaza de Mercado de La Concordia
- 6. Museo de Bogotá – Instituto Distrital de Patrimonio Cultural

## PROGRAM OVERVIEW (ROOM)

	PART 1: membranes			PART 2: free energy	
	Monday (13.08)	Tuesday (14.08)	Wed. (15.08)	Thursday (16.08)	Friday (17.08)
8:00 – 9:00	Registration W101				
9:00 – 9:30	Welcome W101				
9:30 - 10:30	Bert de Groot W101	Thomas Walz (webinar) MLC	Chad Leidy W101	CT4: Hugo J. Bohórquez / W101	Coffee break
10:30 – 11:00	Coffee break				Hands on LL002 W504
11:00 – 12:00	Helgi Ingólfsson W101	Gerhard Hummer MLC	CT2: Karen Palacio CT3: Sergio Mares-Samano W101	Esteban Vohringer-Martínez W101	
2:00 – 14:00	Lunch		Free program	Lunch	
14:00 – 15:00	Gerhard Hummer W101	Bert de Groot MLC		Andrés González W101	
15:00 – 16:00	Vytas Gapsys / Camilo Aponte / CT1: Florian Franz W101	Helgi Ingólfsson MLC		Coffee break	Vytas Gapsys MLC
				CT5: Angélica Sandoval CT6: Stefany Botero Rodríguez W101	
16:00 – 16:40	Coffee break				Closing remarks & departure MLC
16:40 – 19:00	Hands on LL001 LL002	Hands on LL001 LL002		Poster session Hall ML AB	

## PROGRAM IN DETAIL

### PART 1: COMPUTER SIMULATIONS OF BIOLOGICAL MEMBRANES

#### Monday, August 13

Time	Room		
8:00 – 9:00	W101	Registration	
9:00 – 9:30		Welcome	
9:30 – 10:30	W101	<b>Bert de Groot MPIBPC, Göttingen</b>	<i>The molecular dynamics of solute permeation through biological membranes and membrane proteins</i>
10:30 – 11:00		Coffee break	
11:00 – 12:00	W101	<b>Helgi Ingólfsson LLNL, Livermore</b>	<i>Coarse-grained simulations: introduction and application to membranes</i>
12:00 – 14:00		Lunch	
14:00 – 15:00	W101	<b>Gerhard Hummer MPIB, Frankfurt</b>	<i>Large finite-size effects in molecular dynamics simulations of membranes</i>
15:00 – 15:20	W101	<b>Vytautas Gapsys MPIBPC, Göttingen</b>	<i>Grid-based methods for the analysis of MD simulations of biological membranes I: <u>g_lomempro</u></i>
15:20 – 15:40	W101	<b>Camilo Aponte UANDES, Bogotá</b>	<i>Grid-based methods for the analysis of MD simulations of biological membranes II: <u>GROmars</u></i>
15:40 – 16:00	W101	<b>Contributed talk 1: Florian Franz IWR, Heidelberg</b>	<i>Stability of Biological Membranes upon Mechanical Indentation</i>
16:00 – 16:10		Photo	
16:10 – 16:30		Coffee break	
16:30 – 16:40		Heading to computer room	
16:40 – 19:00	LL001/ LL002	<b>Hands on: all-atom MD simulations of biological membranes Introductory (LL001) – Advanced (LL002)</b>	
19:00 –		Invited speakers dinner	

#### Tuesday, August 14

9:30 – 10:30	MLC	<b>Webinar, Thomas Walz Rockefeller, New York</b>	<i>Electron Microscopy Approaches to Studying Lipid-Protein Interactions</i>
10:30 –		Coffee break	

11:00			
11:00 – 12:00	MLC	<b>Gerhard Hummer MPIB, Frankfurt</b>	<i>Molecular dynamics simulations of proteins that sense the composition and state of lipid membranes</i>
12:00 – 14:00		Lunch	
14:00 – 15:00	MLC	<b>Bert de Groot MPIBPC, Göttingen</b>	<i>Permeation, selectivity and gating of ion channels studied by computational electrophysiology</i>
15:00 – 16:00	MLC	<b>Helgi Ingolfson LLNL, Livermore</b>	<i>Capturing biological membranes in silico at different compositional complexity</i>
16:00 – 16:30		Coffee break	
16:30 – 16:40		Heading to computer room	
16:40 – 19:00	LL001/ LL002	<b>Hands on: coarse-grained MD simulations of biological membranes Introductory (LL001) – Advanced (LL002)</b>	
Wednesday, August 15			
9:30 – 10:30	W101	<b>Chad Leidy UANDES, Bogotá</b>	<i>Integrating Computational and Experimental Approaches to Investigate Antimicrobial Peptide Activity in Membranes</i>
10:30 – 11:00		Coffee break	
11:00 – 11:30	W101	<b>Contributed talk 2: Karen Palacio Rodríguez UDEA, Medellín</b>	<i>Exponential consensus ranking improves outcome in docking and receptor ensemble docking</i>
11:30 – 12:00	W101	<b>Contributed talk 3: Sergio Mares-Samano UNAM, México City</b>	<i>Molecular dynamics studies of the solvation effects of water and trifluoroethanol on the peptides pandinin-2 (pin2) and pin2-GVG</i>
12:00 –		free program	
PART 2) FREE ENERGY CALCULATIONS OF BIOLOGICAL SYSTEMS			
Thursday, August 16			
9:00 – 10:00	W101	<b>Gian Pietro Miscione UANDES, Bogotá</b>	<i>Mechanism of L-Tryptophan Prenylation, Catalyzed by FgaPT2: a Computational Study</i>
10:00 – 10:30	W101	<b>Contributed talk 4: Hugo J. Bohórquez FIDIC, Bogotá</b>	<i>The frozen molecule approach in peptide-protein binding energies</i>
10:30 – 11:00		Coffee break	
11:00 – 12:00	W101	<b>Esteban Vohringer-Martínez</b>	<i>Force field improvement with ab-initio methods</i>



		<b>UCONCEPCIÓN, Concepción</b>	
12:00 – 14:00		Lunch	
14:00 – 15:00	W101	<b>Andrés González UANDES, Bogotá</b>	<i>Applications of product design departing from free energy calculations</i>
15:00 – 15:30		Coffee break	
15:30 – 16:00	W101	<b>Contributed talk 5: Angélica Sandoval UANDES, Bogotá</b>	<i>Specific at one side unspecific at the other: the interaction of blood proteins with extracellular DNA</i>
16:00 – 16:30	W101	<b>Contributed talk 6: Stefany Botero Rodríguez UNAL, Bogotá</b>	<i>Computational Tracing of Protein and ncRNA epitopes inside Dengue Virus</i>
16:30 – 16:40		Heading to poster room	
16:40 – 19:00	Hall ML AB	<b>Poster session (with wine and finger food)</b>	
Friday, August 17			
9:00 – 10:00	MLC	<b>Bert de Groot MPIBPC, Göttingen</b>	<i>Introduction to alchemical free energy calculations</i>
10:00 – 10:45		Coffee break	
10:45 – 12:45	LL002 W504	<b>Hands on: Free energy calculations Introductory (LL002) – Advanced (W504)</b>	
12:45 – 15:00		Lunch	
15:00 – 16:00	MLC	<b>Vytas Gapsys MPIBPC, Göttingen</b>	<i>Application of the alchemical free energy calculations to proteins, nucleic acids and ligands</i>
16:00 – 16:30	MLC	Closing remarks and departure	

## POSTERS

1	<b>Diego Gomez Hernandez</b> <b>UDISTRITAL,</b> <b>Bogotá</b>	<i>In silico evaluation of the potential affinity of anandamide analogues for CB1 receptor.</i>
2	<b>Carlos Pinilla</b> <b>UNORTE,</b> <b>Barranquilla</b>	<i>FeS alloys at conditions of the Earth Interior</i>
3	<b>Juan David Orjuela</b> <b>UANDES, Bogotá</b>	<i>Effect of the lipid-protein interaction on the assembly of the platelet Glycoprotein-IB</i>
4	<b>Markus Kurth</b> <b>Heidelberg U.</b> <b>Heidelberg</b>	<i>Probing the lipid environment of the Metabotropic glutamate receptor 2</i>
5	<b>Laura Pinto</b> <b>Herrera</b> <b>UANDES, Bogotá</b>	<i>Effect of particulate matter on the lung surfactant at a high altitude city: Review</i>
6	<b>Johann Bedoya</b> <b>Cardona</b> <b>UNAL, Medellín</b>	<i>Applicability of some force field parameters of biomolecules to model the surface tension of hexadecane</i>
7	<b>Yaritza Fortaleché</b> <b>Rodríguez</b> <b>UNAL, Bogotá</b>	<i>Agonist, antagonist or super-activator effect of different ligands in the receptor-ligand interaction of the TraR protein</i>
8	<b>Paula Giraldo</b> <b>Hincapié</b> <b>UNAL, Medellín</b>	<i>Understanding insecticide peptides: Modeling peptide-membrane interactions with Molecular Dynamics</i>
9	<b>Jessica Martinez</b> <b>UAN, Bogotá</b>	<i>Does Using Virtual Sites in Proteins Affect the Protein Fold?</i>
10	<b>Juan Zarate</b> <b>Moreno</b> <b>SENA, Bogotá</b>	<i>Kinetic models for chitosan membranes on biological systems, a computational approach.</i>
11	<b>Cesar Quintana</b> <b>Cataño</b> <b>UANDES, Bogotá</b>	<i>Force spectroscopy of T4 bacteriophage adhesion during infection.</i>
12	<b>Santiago Agudelo</b> <b>Gómez</b> <b>UDEA, Medellín</b>	<i>Preferred sites of glycosaminoglycans in the envelope protein of the Zika virus</i>
13	<b>Alberto Mario</b> <b>Castillo</b> <b>UBJTL, Bogotá</b>	<i>Spring Network Model of Trypanosoma Cruzi bloodstream trypomastigote</i>
14	<b>Gesivaldo Santos</b> <b>State U. Bahia</b>	<i>Autism Spectro Disorder &amp; Alzheimer Disease: different conditions linked by mTOR pathway</i>
15	<b>Helman Amaya</b> <b>Espinosa</b> <b>UANDES, Bogotá</b>	<i>Cooperative dynamics of self-interacting Biopolymers in blood elucidated by Coarse-Grained Brownian Dynamics Simulations</i>

16	<b>Valeria Mejía Restrepo</b> UANDES, Bogotá	<i>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.</i>
17	<b>Andrés Cifuentes López</b> UDISTRITAL, Bogotá	<i>In silico evaluation of the potential affinity of anandamide analogues for CB1 receptor</i>
18	<b>Jhoan Ortiz Giron</b> UDEA, Medellín	<i>A novel methodology for cryo-EM map validation</i>