# Nicholas Leioatts

Department of Biochemistry and Biophysics University of Rochester Medical Center Box 712, 601 Elmwood Ave. Rochester, NY 14642 Phone: (772) 633-1719

Email: Nicholas\_Leioatts@urmc.rochester.edu

#### **Research Interests**

- Structure and dynamics of allosteric macromolecules.
- Computational methods, especially those that allow modeling at various scales of time and resolution.
- Analysis methods for directly comparing computational results to experimental observables.

#### **Education**

Ph.D. Biophysics, University of Rochester Medical Center, (**expected 2014**). M.S. Biophysics, University of Rochester Medical Center, 2012. B.S. Physics, Florida Institute of Technology, 2009.

## **Research Experience**

#### Graduate Research

• University of Rochester Medical Center

**Focus:** *Understanding allostery in GPCRs using computer simulation at multiple resolutions* Advisor: Dr. Alan Grossfield

### REU/Internships

• Amertron Inc., Spring 2007 - Spring 2008.

Manufacturing Engineer (Designing/Testing RF connectors)

• Fermi National Accelerator Lab, Summer 2006.

Performed monte carlo simulations of particle accelerator beamline

#### Undergraduate Research

• Detector Physics, Summer 2005 - Spring 2009.

Muon Tomography (Designed & built scintillation and gaseous-based particle detectors)

## **Teaching**

- Mentor for research students: Hanna Shebert (2012), Pooja Suresh (2013).
- GRE Quantitative Preparatory Class, Instructor, Fall 2010 & 2012.
- General Biochemistry (Ph.D.), Teaching Assistant, Fall 2010.

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### Honors, Awards, & Fellowships

- Elon H. Hooker fellowship, 2013-2014.
- William F. Neuman Award, 2013.
- Training Grant, NIH T32 GM068411, 2011-2012.
- Biochemistry Department Retreat Poster Presentation Award, 2012.
- Departmental Seminar Award, 2011 & 2012.
- Departmental Travel Award, 2011.

#### **Professional Activities and Service**

- Associate Chair for Gordon Research Seminar in Computational Chemistry, 2014.
- Host for annual "T32 student-invited speaker" seminar, 2013.
- Reviewer
  - Chem. Phys. Lipids, 2013.
  - J. Chem. Theor. & Comp., 2012-2013.
- Departmental service:
  - Organizer Biophysics Program Retreat, 2010 & 2011.

#### **Conference and Seminar Presentations**

- Retinal Changes Conformation During the Early Stages of Rhodopsin Activation.
- Understanding the Rhodopsin Activation Mechanism with Molecular Dynamics.
  57th Annual Biophysical Society Meeting. Philadelphia, PA, 2-6 February 2013.
- Unraveling Allostery with Simulations of Rhodopsin and Opsin.
  Computational Chemistry Gordon Research Conference. Mount Snow, VT, 22-27 July 2012.
- Elucidating Elastic Network Model Robustness by Parameterization with Molecular Dynamics. **56th Annual Biophysical Society Meeting.** San Diego, CA, 25-29 February 2012.
- Validating and Improving Elastic Network Models with Microsecond Scale Molecular Dynamics. **55th Annual Biophysical Society Meeting.** Baltimore, MD, 5-9 March 2011.

#### **Publications**

- 1. **Leioatts, N.**, Suresh, P., and Grossfield, A., *Tracking the Activation of Two Model GPCRs using Structure-Based Dynamics*, (in preparation, submission planned February 2014)
- 2. **Leioatts, N.**, Mertz, B., Martínez-Mayorga, K., Romo, T.D., Pitman, M.C., Feller, S.E., Grossfield, A., and Brown, M.F., *Retinal dynamics explained by a concerted transition early in rhodopsin activation*, Biochemistry, 2013, DOI: 10.1021/bi4013947
- 3. Seckler, J.M., Leioatts, N., Mao, H., and Grossfield, A., *The interplay of structure and dynamics in the function of HIV-1 Reverse Transcriptase*, Proteins: Struc. Func. Bioinf., 2013, DOI: 10.1002/prot.24325
- 4. **Leioatts, N.**, and Grossfield, A., *Molecular Dynamics Simulations of Membranes and Membrane Proteins* in "Molecular Modeling at the Atomic Scale", ed. Ruhong Zhou, CRC Press (in press, June 2014)
- 5. **Leioatts, N.**, Romo, T.D., and Grossfield, A., *Elastic network models are robust to variations in formalism*, J. Chem. Theor. Comput., 2012, DOI:10.1021/ct3000316

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