George A. PANTELOPULOS

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USA Nationality

Single with no children

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Graduate Student

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Education 2015-Current

Expected Graduation: 2020/9/25

2013-2015 B.S. OF CHEMISTRY. Temple University, Philadelphia, USA.

Graduation: 2015/5/8

A.S. OF Science. Community College of Philadelphia, Philadelphia, USA. 2011-2013

Ph.D of Chemistry. Boston University, USA.

Graduation: 2013/5/4

Honors

2019 LICHTIN AWARD FOR EXCELLENCE IN RESEARCH.

RIKEN SHORT-TERM INTERNATIONAL PROGRAM ASSOCIATE. 2018

2017 NSF GROW/JSPS Fellow.

2015-Current NSF GRADUATE RESEARCH FELLOWSHIP.

PECO SCHOLAR, PECO. 2015

2015 FEYNMAN MEMORIAL SCHOLAR, Temple University. 2014 DIAMOND SCHOLAR SUMMER FELLOW, Temple University.

2013 NSF RESEARCH EXPERIENCE FOR UNDERGRADUATES, Boston University.

Service and Teaching

2018-2019 TEACHING FELLOW, BOSTON UNIVERSITY CH225 - MATHEMATICAL ME-

THODS FOR CHEMISTS.

2016-2018 GRADUATE FELLOWSHIPS MENTORSHIP AND WORKSHOPS, BOSTON UNIVER-

SITY CHEMISTRY DEPARTMENT.

2015-2016 WRITING MENTOR, BOSTON UNIVERSITY CHEMICAL WRITING PROGRAM.

2015-2016 HIGH SCHOOL SCIENCE OUTREACH INSTRUCTOR, BOSTON UNIVERSITY WO-

MEN IN CHEMISTRY.

2014-2015 SCIENCE INSTRUCTOR, TEENSHARP.

Presentations

2020	"Exploring APP-C99 complexes in lipid bilayers and their role in Alzheimer's disease," ACS Annual Meeting, Philadelphia (Virtual Poster).
2019	"Phase separations in Lipid membranes, what they have to do with proteins, and recent developments in MD-based modeling," Boston University Physical Chemistry Lecture Series (Talk).
2019	"Probing the roles of membrane and cholesterol on $A\beta$ biogenesis and toxicity," ACS Annual Meeting, Orlando (Talk).
2018	"Understanding an Intrinsically Disordered Membrane Protein in Alzheimer's Disease via Molecular Simulation," RIKEN Summer School (Poster); Computational Biophysics at the Molecular and Meso Scales, ICISE Vietnam (Talk).
2017	"Structure of APP-C99 1-99 and Implications for its Role in Amy- Loidogenesis," <i>Kyoto University (Talk)</i> .
2017	"Role of Cholesterol in Ternary Lipid Membrane Phase Separation observed via coarse-grained simulations," American Theoretical Chemistry Conference (Poster); Japanese Biophysical Society (Poster).
2017	"Tempering in OpenMM and GENESIS," Temple University Current Trends in Molecular Dynamics Software Design Workshop (Poster).
2017	"Critical investigation of finite size and cholesterol effects in li- pid domain formation," RIKEN, Wako-shi (Talk); Boston University Che- mistry Student Graduate Seminar Series; Nagoya University IGER Seminar Series (Talk).
2016	"Exploring phase separation and domain formation in lipid bilayers through molecular simulation," Boston University Graduate Research Symposium; ACS National Meeting, Philadelphia (Poster).
2015	"Examining the conformational dynamics of the N-terminal region of mdm2 using markov state model approaches," ACS National Meeting, Boston (Poster).
2015	"MICROSECOND SIMULATIONS OF MDM2 AND ITS COMPLEX WITH P53 YIELD INSIGHT INTO FORCE FIELD ACCURACY AND CONFORMATIONAL DYNAMICS," Temple University Undergraduate Research Forum and Creative Works Symposium (Talk).
2014-2015	"MICROSECOND SIMULATIONS OF MDM2 AND ITS COMPLEX WITH P53 YIELD INSIGHT INTO FORCE FIELD ACCURACY AND CONFORMATIONAL DYNAMICS," ACS Philadelphia Younger Chemists Committee Conference (Poster); Emory STEM Research Symposium (Poster).
2013-2014	"Molecular Dynamics Simulations of the P53-MDM2 Binding Interface," ACS Philadelphia Younger Chemists Committee Conference (Poster); ACS 248th National Meeting (Poster).

Publications

- 20. "MolPainter and MolSolvator: tools for drawing and solvating complex planar molecular systems of arbitrary composition and molecule placement," G.A. Pantelopulos*, A. Liberatore, *In Preparation*, (2020)
- 19. "Evaluation and applicability of prominent coarse-granined models of cholesterol in molecular dynamics simulation." G.A. Pantelopulos*, A. Bandara, J.E. Straub, *In Preparation*, (2020)
- 18 "Characterization of the intrinsically disordered monomer and homodimer of APP-C99 1-99" G.A. Pantelopulos*, J. Hutchinson, D. Matsuoka, C. Sanders, Y. Sugita, J.E. Straub, *In Preparation*, (2020) 17. "*De novo* all-atom molecular dynamics simulation of amyloid precursor protein in phase separating ternary lipid mixture." G.A. Pantelopulos*, A. Bandara, J.E. Straub, *In Preparation*, (2020)
- 15. "Cholesterol effects amyloid precursor protein via the liquid ordered phase as a nonbinding solvent." G.A. Pantelopulos*, A. Panahi, J.E. Straub, *In Preparation*, (2020)
- 14. "Direct Observation of Cholesterol Dimers and Tetramers in Lipid Bilayers," M.R. Elkins, A. Bandara, G.A. Pantelopulos, J.E. Straub, M. Hong, Submitted (2020)
- 13. "Bicelles Rich in both Sphingolipids and Cholesterol and Their Use in Studies of Membrane Proteins," J. Hutchinson, K. Shih, H. Scheidt, S. Fantin, G.A. Pantelopulos, H. Harrington, K. Mittendorf, S. Qian, R. Stein, S. Collier, M. Chambers, J. Kastaras, M. Voehler, B. Ruotolo, D. Huster, R. McFeeters, J. Straub, M. Neih, C. Sanders J. Am. Chem. Soc., Accepted (2020)
- 12. "Exploring the impact of line tension on the spatial localization of protein in phase-separated lipid bilayers," A. Bandara, A. Panahi, G.A. Pantelopulos, T. Nagai, J. E. Straub, *J. Phys. Chem. B*, 20, 204702 (2019)
- 11. "Aerosol-OT Surfactant Forms Stable Reverse Micelles in Aploar Solvent in the Absence of Water," R. Urano, G.A. Pantelopulos, J. E. Straub, J. Chem. Phys. 14, 149 (2019).
- 10. "Structural Role of Cholesterol in Complex Lipid Bilayer Phases Observed via MD Simulation," G.A. Pantelopulos*, J. E. Straub, Biophys. J. 115, 2167-2178 (2018).
- 9. "Characterization of dynamics and mechanism in the self-assembly of AOT reverse micelles," R. Urano,
 G.A. Pantelopulos*, S. Song, J. E. Straub, J. Chem. Phys. 14, 144901 (2018).
- 8. "Structure of APP-C99₁₋₉₉ and Implications for Role of Extra-Membrane Domains in Function and Oligomerization," G.A. Pantelopulos, J. E. Straub, D. Thirumalai, Y. Sugita, *Biochim. Biophys. Acta Biomembranes* 1860, 1698-1708 (2018).
- 7. "Critical size dependence of domain formation observed in coarse-grained simulations of bilayers composed of ternary lipid mixtures," G.A. Pantelopulos*, T. Nagai, A. Bandara, A. Panahi, J. E. Straub, J. Chem. Phys. 147 095101 (2017).
- **6.** "Bridging microscopic and macroscopic mechanisms of p53-MDM2 binding using molecular simulations and kinetic network models," G. Zhou, **G.A. Pantelopulos**, S. Mukherjee, V. Voelz, *Biophys. J.* **113**, 785-793 (2017).
- 5. "Exploring the structure and stability of cholesterol dimer formation in multicomponent lipid bilayers,"
- A. Bandara, A. Panahi, **G.A. Pantelopulos**, J.E. Straub *J. Comp. Chem.* **38**, 1479-1488 (2016).
- 4. "Specific binding of cholesterol to C99 domain of Amyloid Precursor Protein depends critically on charge state of protein," A. Panahi, A. Bandara, G.A. Pantelopulos, L. Dominguez and J.E. Straub, J. Phys. Chem. Lett. 7, 3535-3541 (2016).
- **3.** "On the use of mass scaling for stable and efficient simulated tempering with molecular dynamics," T. Nagai, **G.A. Pantelopulos**, T. Takahashi, J.E. Straub *J. Comp. Chem.* **37** 2017-2028 (2016).
- 2. "Markov models of the apo-MDM2 lid region reveal diffuse yet two-state binding dynamics and receptor poses for computational docking," S. Mukherjee, G.A. Pantelopulos*, V.A. Voelz Sci. Rep. 6 31631 (2016).
- 1. "Microsecond simulations of mdm2 and its complex with p53 yield insight into force field accuracy and conformational dynamics," **G.A. Pantelopulos***, S. Mukherjee, V.A. Voelz, *Proteins* **83** 1665-1676 (2015).