

**CURRICULUM VITAE****Dr. Pradipta Bandyopadhyay**

Professor, Jawaharlal Nehru University  
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**ACADEMIC/INDUSTRIAL POSITIONS**

Nov 2014-present Professor, Jawaharlal Nehru University  
 Jan-Mar, 2020 Visiting Scientist, Stony Brook University  
 May-Oct, 2019 Visiting Scientist, University of Oklahoma  
 Aug 2007-2014 Associate Professor, Jawaharlal Nehru University  
 Aug-Oct 2008 Visiting Associate Professor, University of California, San Francisco  
 2004-2007 Assistant Professor, Indian Institute of Technology, Guwahati, India  
 2003 Post-Doctoral Fellow, Lawrence Berkeley National Laboratory  
 Sep-Dec, 2002 Scientist at ARQULE INC., Redwood City, California

**EDUCATION/TRAINING**

| Institution                                     | Degree   | Year(s)   | Field                                     |
|---|----------|-----------|---|
| Indian Institute of Technology, Kanpur          | M.Sc.    | 1994      | Chemistry                                 |
| Graduate University for Advanced Studies, Japan | Ph.D.    | 1999      | Theoretical and Computational Chemistry   |
| Iowa State University                           | Post-Doc | 1999-2001 | Theoretical and Computational Chemistry   |
| University of California, San Francisco         | Post-Doc | 2001-2002 | Computational Biochemistry and Biophysics |

**RESEARCH ACHIEVEMENTS:****Advancement of several computer simulation algorithms/methods:**

- Published several papers on a nested Monte Carlo technique that can increase the speed of computer simulation by orders of magnitude for both quantum mechanical/molecular mechanical (QM/MM) and classical systems
- Developed a new method to explore the energy surface of complex molecules known as Monte Carlo Temperature Basin Paving (MCTBP)
- In collaboration with Prof. Ken Dill, developing a fast and accurate solvation model for MD simulation of biomolecules to significantly increase the applicability of MD simulation

**State-of-the-art applications to biological systems:**

- Diffusion of proteins in a full *E. coli* cell
- Accurate binding free energy calculations for designing the active sites of calcium-binding proteins
- Innovative calculations of protein-protein binding affinities in solutions with high salt concentrations

**FELLOWSHIPS, HONORS AND AWARDS:**

- Distinguished Lecture award from the Chemical Society of Japan, 2013

- Finalist for the Swarnajayanti Award in chemistry (An award to recognize scientists below 40 years old in India), 2009
- Indo-US research fellowship to conduct research at the University of California, San Francisco, 2008.
- Monbusho (Japanese Government) scholarship for foreign graduate students, 1996-1999
- Best Masters project award in chemistry from the Indian Institute of Technology, Kanpur, 1994

#### **CURRENT GRANTS:**

- Toward manipulation of the binding affinity of a calcium ( $\text{Ca}^{2+}$ ) ion to calcium-binding proteins using computational approaches with experimental validations; with Co-PI Prof. S. Gourinath; from the Department of Biotechnology of India.
- Path towards an ultrafast model of solvation of molecules and its connection to complex processes in biology; with Prof. T. Urbic of University of Lubjijana, Slovenia; Indo-Slovenia Bilateral Grant
- Development of an analytical model of solvation in water using statistical mechanics; Mathematical Research Impact Centric Support (MATRICS), India

#### **PERSONNEL TRAINED:**

- 7 PhD students awarded degrees; • 5 current Ph.D. students: • 2 Post-Doctoral Fellows; • 15 Master's Students

#### **SELECTED INVITED LECTURES AND MEETING SESSIONS CHAIRED:**

|          |   |
|----------|---|
| Aug 2023 | WATER MEETING, Stony Brook University                 |
| Feb 2023 | APACC, Vietnam  |
| 2022     | Theoretical Chemistry Symposium, IISER Kolkata, India |
| 2019     | University of Texas, Austin                           |
| 2019     | PNNL, USA   |
| 2017     | Rare Event Simulation, Agra, India                    |
| 2017     | Energy Landscape meeting, Goa,                        |
| 2010     | Telluride Meeting on Energy Landscapes,               |

#### **TEACHING EXPERIENCE:**

##### **Multiple Undergraduate, Masters and Doctorate level courses in the last 19 years including:**

- Biomolecular simulation, • Computational Structural Biology, • Basic Thermodynamic and Statistical Mechanics, • Statistical Mechanics for Biomolecules, • Advances in Physical Science, • Basic Quantum Chemistry

#### **COURSES DESIGNED:**

- Biomolecular simulation, • Statistical Mechanics for Biomolecules

#### **MAJOR ADMINISTRATIVE COMMITTEES**

- Chair, The Graduate Advisory Committee
- Spearheaded the introduction of a regular Ph.D. course in our school in 2009
- Faculty recruitment committee

#### **OTHER PROFESSIONAL SERVICE:**

- Organized 2 International conferences in 2011 and 2013; • Ph.D. Thesis reviewer of top Indian institutes and universities such Indian Institute of Science, and different IITs. • Peer reviewer for J Chem

Phys, JPhysChem, JBiolStrdyn, Biophys J, ACS Omega, ChemPhysLett; • Selection Committee member for the faculty member recruitment in another university.

## SELECTED PUBLICATIONS

For the full list see [Pradipta Bandyopadhyay - Google Scholar](#)

1. Srivastava, R.; Bandyopadhyay, P. Nested Monte Carlo Simulation of Ionic Systems with the Primitive Model Using Debye-Hückel (DH) Potential as an Importance Function and Optimizing the DH Potential with Kullback-Leibler Divergence Minimization. *J. Chem. Sci.* **2023**, *135* (2). <https://doi.org/10.1007/s12039-023-02167-0>.
2. Roy, U. C.; Bandyopadhyay, P. Correlation between Protein Conformations and Water Structure and Thermodynamics at High Pressure: A Molecular Dynamics Study of the Bovine Pancreatic Trypsin Inhibitor (BPTI) Protein. *J. Chem. Phys.* **2023**, *158* (9). <https://doi.org/10.1063/5.0124837>.
3. Yadav, A. K.; Bandyopadhyay, P.; Coutsiyas, E. A.; Dill, K. A. Crustwater: Modeling Hydrophobic Solvation. *J. Phys. Chem. B* **2022**, *126* (32), 6052–6062. <https://doi.org/10.1021/acs.jpcc.2c02695>.
4. Basit, A.; Yadav, A. K.; Bandyopadhyay, P. Calcium Ion Binding to the Mutants of Calmodulin: A Structure-Based Computational Predictive Model of Binding Affinity Using a Charge Scaling Approach in Molecular Dynamics Simulation. *J. Chem. Inf. Model.* **2022**, *62* (11), 2821–2834. <https://doi.org/10.1021/acs.jcim.2c00428>.
5. Srivastava, R.; Chattopadhyaya, M.; Bandyopadhyay, P. Calculation of Salt-Dependent Free Energy of Binding of  $\beta$ -Lactoglobulin Homodimer Formation and Mechanism of Dimer Formation Using Molecular Dynamics Simulation and Three-Dimensional Reference Interaction Site Model (3D-RISM): Diffuse Salt Ions and Non-Polar Interactions between the Monomers Favor the Dimer Formation. *Phys. Chem. Chem. Phys.* **2020**, *22* (4), 2142–2156. <https://doi.org/10.1039/c9cp05578a>.
6. Rakshit, A.; Bandyopadhyay, P.; Heindel, J. P.; Xantheas, S. S. Atlas of Putative Minima and Low-Lying Energy Networks of Water Clusters  $n = 3-25$ . *J. Chem. Phys.* **2019**, *151* (21). <https://doi.org/10.1063/1.5128378>.
7. Singh, P.; Sarkar, S. K.; Bandyopadhyay, P. Folding-Unfolding Transition in the Mini-Protein Villin Headpiece (HP35): An Equilibrium Study Using the Wang-Landau Algorithm. *Chem. Phys.* **2016**, *468*, 1–8. <https://doi.org/10.1016/j.chemphys.2016.01.005>.
8. Hasnain, S.; Bandyopadhyay, P. An Analytical Correlated Random Walk Model and Its Application to Understand Subdiffusion in Crowded Environment. *J. Chem. Phys.* **2015**, *143* (11). <https://doi.org/10.1063/1.4930275>.
9. Hasnain, S.; McClendon, C. L.; Hsu, M. T.; Jacobson, M. P.; Bandyopadhyay, P. A New Coarse-Grained Model for E. Coli Cytoplasm: Accurate Calculation of the Diffusion Coefficient of Proteins and Observation of Anomalous Diffusion. *PLoS One* **2014**, *9* (9). <https://doi.org/10.1371/journal.pone.0106466>.
10. Bandyopadhyay, P. Increasing the Efficiency of Monte Carlo Simulation with Sampling from an Approximate Potential. *Chem. Phys. Lett.* **2013**, *556*, 341–345. <https://doi.org/10.1016/j.cplett.2012.11.047>.
11. Bandyopadhyay, P.; Kuntz, I. D. Computational Investigation of Kinetics of Cross-Linking Reactions in Proteins: Importance in Structure Prediction. *Biopolymers* **2009**, *91* (1), 68–77. <https://doi.org/10.1002/bip.21083>.
12. Bandyopadhyay, P. Accelerating Quantum Mechanical/Molecular Mechanical Sampling Using Pure Molecular Mechanical Potential as an Importance Function: The Case of Effective Fragment Potential. *J. Chem. Phys.* **2005**, *122* (9). <https://doi.org/10.1063/1.1861890>.