• PERSONAL DETAILS:-

1. Name: ARUNDHATI BANERJEE

2. Father's Name: NILOTPAL BANERJEE

3. Date of Birth: 23rd June, 1992



4. Temporary Address: A-72/104, SHUKHOBRISHTI COMPLEX, SHAPOORJI PALONJI HOUSING COMPLEX, ACTION AREA III, NEWTOWN, KOLKATA – 700135, WEST BENGAL, INDIA

5. Permanent Address: A-26/9, BATAYAN COOPERATIVE, GROUP HOUSING COMPLEX, SECTOR 2A, BIDHANNAGAR, DURGAPUR-713212, WEST BENGAL, INDIA

6. Contact:

E-mail: <u>arundhati.92star@gmail.com</u>, <u>arundhati.bt92star@gmail.com</u> Phone numbers: 8145245649 & 9474533102

• ACADEMIC DETAILS:-

| Qualification | Institution | Year of Passing | Board | % of Marks |
|--------------------------------------|--|-----------------|--|---------------------------------|
| PhD Department of Biochemistry | University of Kalyani (under CSIR-SRF Direct) | 1st March 2022 | | Coursework : 88% |
| M.Tech (Biotechnology) | National Institute of Technology Durgapur | 2016 | National Institute of Technology, Durgapur | CGPA: 8.5 (points out of 10) |
| B.Tech (Biotechnology) | West Bengal University of Technology (now MAKAUT) | 2014 | West Bengal University of Technology | DGPA: 8.9 (points out of 10) |
| Class XII-CBSE | Hem Sheela Model School, Durgapur | 2010 | CBSE-XII | 84% |
| Class X - ICSE | Carmel Convent High School, Durgapur | 2008 | CISCE | 88% |

Title of the PhD Thesis: *In silico* Analysis and Interactions among Associated Proteins for AIDS and Leukemia: An Insight to Explore the Therapeutic Aspects of Gp130

Curriculum Highlights: Recombinant DNA Technology, Bioreactor Designing and Analysis, Bioinformatics, Molecular Modeling and Drug Designing, Immunology, Human Genomics, Biochemistry, Proteomics, Plant and Animal Biotechnology, Food Technology, Molecular Biology and Genetics.

• PROFESSIONAL QUALIFICATION

- Guest Faculty in Department of Biotechnology, Nutrition Science, Medical Lab Technology and Physiotherapy at Durgapur Paramedical College (affiliated under MAKAUT (formerly West Bengal University of Technology) and WBUHS) - 16th September 2022 – present
- 2. Worked as a Faculty (Assistant Professor) in Durgapur Institute of Science and Technology for B.Sc and M.Sc (Department of Biotechnology and Biochemistry), under KAZI NAZRUL UNIVERSITY, January 2017 to June 2017
- 3. As a Faculty (Lecturer) at Erudite (2014-2015)

• PROFESSIONAL RESPONSIBILITY

- Conducting semester examinations
- 0 Working as Examiner & Scrutinizer for Biochemistry in Kazi Nazrul University, 2017
- Conducting meetings through conference calls
- Presenting slides in meetings
- Guiding Bachelor students for projects
- o Guided THREE Dissertations from Bachelors Level
- Presently, examiner under the affiliation of MAKAUT

• **<u>PUBLICATIONS</u>**

| Publications in: | National | International |
|------------------------|----------|--|
| Research Papers | - | 26 published (details attached below) h-index as per SCOPUS: 5 Cumulative Impact Factor: 54.76 Citations so far: 79 |
| Book Chapters | - | 9 |
| Conference Proceedings | - | 16 |

• <u>AS A REVIEWER</u>

| Articles Reviewed (Nos.) | 60 (Elsevier, Springer, Nature and Taylor & Francis) |
|--------------------------|--|
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PARTICIPATION

| Participation in: | Numbers |
|-------------------------------|-------------------------------|
| Seminars | 5 |
| Workshops | 3 |
| Conference Paper Presentation | 8 (International Conferences) |

• **<u>GUIDED UG DISSERTATIONS:</u>** THREE IN NUMBER

• **RESEARCH TRAINING UNDERGONE**

| | Numbers |
|---------------------|---------|
| Laboratory Training | 3 |
| Computer Training | 4 |

<u>AWARDS/CERTIFICATES</u>

- Awarded Senior Research Fellowship (Direct) by CSIR (Council of Scientific and Industrial Research), India – 2018 (for two years and one extension)
- o Awarded Senior Research Fellow by ICMR (Indian Council of Medical Research), India
- Have conducted workshop classes (theory and practical) in 21 days Summer Training Programme at **Bioinformatics Infrastructure Facility (BIF)**, University of Kalyani
- Have been invited as a <u>Keynote Speaker</u> in The ESMED General Assembly in August , 2022
- o Attended a Seminar on "Networking Drug Discovery and Pharm Innovation" on 6th February, 2012.
- Attended Workshops on Spoken Tutorial Project on "<u>Python</u>" and "<u>Linux</u>" developed by IIT Bombay, an initiative of NATIONAL MISSION on Education through Information and Communication Technology (**NMEICT**), MHRD GOVT OF INDIA.

- Awarded Merit Certificates for securing 1st and 2nd Rank in 4 year and 3rd year, respectively in B. Tech
- Merit certificates for *PowerPoint Presentation* (stood first in competitions).
- 0 Merit Certificate in *National Science Olympiad* (NSO), for rank 8 in School, 11 in City.

• <u>COMMUNICATION SKILLS</u>

| Languages | Speak | Read | Write |
|-----------|-------|--------------|-------|
| English | | \checkmark | |
| Bengali | | \checkmark | |
| Hindi | | \checkmark | |

• COMPUTER PROFICIENCY:

- C-Language.
- Perl Programming.
- MODELLER9.14 Software packages
- Discovery Studio packages from Accelyrs.
- o Competent user of basic MS Office packages like MS Word, MS Excel, MS Power Point.
- Strength: Determination, Organization, Communication, Task Prioritization and Team Leader
- Weakness: Being too honest, selflessly helpful, Too attached, Inability to say "no".
- **Skills/Attitude:** Deep Critical Thinking, Investigate and develop new strategies, Eager to solve problems, motivated, efficient in team work, committed to a work, efficient in managing time for a particular work.
- About Yourself :

• <u>Special Interest in</u>: Bioinformatics, Computational Structural Biology and *In silico* Studies, Drug Discovery and Molecular Modelling, Pharmacogenetics, Genomics, rDNA technology, Bioreactor Designing and Analysis.

• Have rendered <u>Theory and Hands on Demonstration Sessions</u> in 21 Days Summer Training Programme 2019 at Bioinformatics Infrastructure Facility (BIF), University of Kalyani

Skilled in preparing presentations, doing team work and have secured awards for presentations.
Have sent CV upon invitation for the <u>Editorial Board Membership</u> for International Peer-Review Journals.

 \circ Skilled in preparing presentations, doing team work and have secured awards for presentations. \circ Confident in research work documentation and authoring technical reports.

• Have performed as an *examiner and scrutinizer for university semester examinations*.

• **REFERENCES** (any 2)

| Name: Dr. N.B Hui | Name: Dr. Apurba Dey |
|---|---|
| Designation: Professor | Designation: Senior Professor |
| Company Name: National Institute of Technology, | Company Name: National Institute of Technology, |
| Durgapur | Durgapur |
| Contact No.: 9434788117, | Contact No.: 9434788098 |
| Email: nirmal.hui@me.nitdgp.ac.in | Email: apurba.dey@bt.nitdgp.ac.in |

Arundhati Banerjee

Date: 21st November, 2023

[ARUNDHATI BANERJEE]

LIST OF PUBLICATIONS:

(a) For Refereed Journal Papers

| Sl. No. | Author(s) | Title of paper | Name of the Journal | Impact Factor | Volume | Year | Pages |
|------------|---|---|--|------------------|---------|------|---------------|
| 1 | Shreya Bhattacharya, Arundhati Banerjee , Sujay Ray | Development of new vaccine target against SARS-CoV2 using envelope (E) protein: An evolutionary, molecular modeling and docking based study | International Journal of Biological Macromolecules | 8.025 | 172 | 2021 | pp: 74- 81 |
| 2 | Arundhati Banerjee , Sujay Ray | Molecular level biodegradation of phenol and its derivatives through dmp operon of <i>Pseudomonas</i> <i>putida</i> : A biomolecular modelling and docking analysis | Journal of Environmental Sciences | 6.796 | 36 | 2016 | 144- 151 |
| 3 | Arundhati Banerjee , Sujay Ray | Molecular modeling, mutational analysis and conformational switching in IL27: An <i>in silico</i> structural insight towards AIDS research | Gene | 3.913 | 610 | 2017 | 72-78 |
| 4 | Arundhati Banerjee , Sujay Ray | Structural insight with mutational impact on tyrosinase and PKC- β interaction from Homo sapiens: Molecular modeling and docking studies for melanogenesis, albinism and increased risk for melanoma | Gene | 3.913 | 610 | 2017 | 99-109 |
| 5 | Arundhati Banerjee , Sujay Ray | Mutations and Interactions in Human ERα and bZIP Proteins: An <i>In silico</i> Approach for Cell Signaling in Breast Oncology | Gene | 3.913 | 592 (1) | 2016 | 90-102 |

| | A 11 (* | 0, 1, 11, 1 | | - | | r | 110 |
|----|--|---|---|-------|---------|------|---------------|
| 6 | Arundhati Banerjee , Sujay Ray | Stability in Interactions with Human Beta- Catenin and Helix- to- Coil Transitions upon Mutations in Human SOX17 Protein: A Computational Molecular Outlook for Histogenesis and Organogenesis | Gene | 3.913 | 576 (1) | 2015 | 118- 126 |
| 7 | Sujay Ray, Abishek Basnet, Shreya Bhattacharya, Arundhati Banerjee , Koustav Biswas | A comprehensive analysis of NAC gene family in <i>Oryza sativa</i> <i>japonica</i> : a structural and functional genomics approach | Journal of Biomolecular Structure & Dynamics | 5.235 | 41 (3) | 2023 | 1-15 |
| 8 | Arundhati Banerjee , Sujay Ray | Molecular interactions and mutational impact upon rhodopsin (G90→D90) for hindering dark adaptation of eye: A comparative structural level outlook for signaling mechanism in night blindness | Mutation Research/Fundamental and Molecular Mechanisms of Mutagenesis | 3.15 | 814 | 2019 | 7-14 |
| 9 | Arundhati Banerjee, Rakhi Dasgupta, Sujay Ray | Residual Participation and Thermodynamic Stability Due to Molecular Interactions in IL11, IL11Rα and Gp130 from <i>Homo sapiens</i> : An In Silico Outlook for IL11 as a Therapeutic Remedy | International Journal of Peptide Research and Therapeutics | 2.191 | 26 | 2020 | 2009– 2020 |
| 10 | Shreya Bhattacharya, Pragati Prasad Sah, Arundhati Banerjee , SujayRay | Structural impact due to PPQEE deletion in multiple cancer associated protein - Integrin αV: An In silico exploration | Biosystems | 1.957 | 198 | 2020 | 104216 |

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|----|--|--|---|-------|--------|------|--------------|
| 11 | Shreya Bhattacharya, Shreyeshi Dhar, Arundhati Banerjee , Sujay Ray | Structural, Functional, and Evolutionary Analysis of Late Embryogenesis Abundant Proteins (LEA) in Triticum Aestivum: A Detailed Molecular Level Biochemistry Using in Siliag | Computational Biology and Chemistry | 3.737 | 82 | 2019 | 9-24 |
| | | Using in Silico Approach | | | | | |
| 12 | Shreya Bhattacharya, Arundhati Banerjee , Pragati P. Sah, Chittobrata Mal, Sujay Ray | Mutations and Functional Analysis of 14-3-3 Stress Response Protein from Triticum Aestivum: An Evolutionary Analysis through in Silico Structural Biochemistry Approach | Computational Biology and Chemistry | 3.737 | 77 | 2018 | 343– 353 |
| 13 | Pragati Sah, Shreya Bhattacharya, Arundhati Banerjee , Sujay Ray | Identification of novel therapeutic target and epitopes through proteome mining from essential hypothetical proteins in Salmonella strains: An In silico approach towards antivirulence therapy and vaccine development. | Infection, Genetics and Evolution | 4.393 | 83 | 2020 | 104315 |
| 14 | Shreya Bhattacharya, Shreyeshi Dhar, Arundhati Banerjee and Sujay Ray | Detailed Molecular Biochemistry for Novel Therapeutic Design against Nipah and Hendra Virus: A Systematic Review | Current Molecular Pharmacology | 3.855 | 13 (2) | 2020 | 108 - 125 |
| 15 | Arundhati Banerjee , Rakhi Dasgupta and Sujay Ray | A Computational Study to Prevent HIV Invasion by Bovine LF in Mucosal- Layer via Blocking of DC- SIGN_GP120 Interaction | Current Proteomics | 0.642 | 17 (5) | 2020 | 413 - 424 |

| 16 | Arundhati | Mutational, | Meta Gene | 0.201 | 21 | 2019 | 100591 |
|----|---|---|---|-------|-------|------|--------------|
| | Banerjee , Shreya Bhattacharya, Rakhi Dasgupta, Sujay Ray | Functional and Evolutionary Analysis of Interleukin-11 in <i>Homo sapiens:</i> A Detailed <i>In silico</i> Exploration for Platelet Recovery due to Chemotherapy Induced Thrombocytopenia. | | | | | |
| 17 | Shreyeshi Dhar, Shreya Bhattacharya, Arundhati Banerjee , Sujay Ray | Evolutionary, gene ontology and physiochemical relationships in LEA proteins of <i>Oryza</i> <i>sativa indica</i> : Detailed computational sequence-based insight. | Plant Gene | 0.427 | 21 | 2020 | 100218 |
| 18 | Shreya Bhattacharya, Puja Ghosh, Debasmita Banerjee, Arundhati Banerjee , Sujay Ray | In silico Drug Target Discovery Through Proteome Mining from M. tuberculosis: An Insight towards Antivirulent Therapy. | Combinatorial Chemistry & High Throughput Screening. | 1.714 | 23(3) | 2020 | 253- 268 |
| 19 | Shreya Bhattacharya, Pragati Prasad Sah, Arundhati Banerjee and Sujay Ray | Exploring Single Nucleotide Polymorphisms in ITGAV for Gastric, Pancreatic and Liver Malignancies: An Approach Towards the Discovery of Biomarker | Combinatorial Chemistry & High Throughput Screening. | 1.714 | 23(6) | 2021 | 860 - 873 |
| 20 | Arundhati Banerjee * and Rakhi Dasgupta | Molecular Modeling, Interacting Residues and other Structural Analyses for Human SOCS3, Gp130 and JAK Proteins: A Detailed Computational Approach for Proteins Involved in Feedback Inhibition | Current Biotechnology | NA | 10(3) | 2021 | 218 - 229 |

| | | | . | | 1 | | 221 |
|----|--|---|--|-------|-------|------|--------------|
| 21 | Sujay Ray and Arundhati Banerjee | Computational structural biology and modes of interaction between human annexin A6 with influenza A virus protein M2: a possible mechanism for reducing viral infection | International Journal of Bioinformatics Research and Applications | 0.184 | 14(4) | 2018 | 321- 336 |
| 22 | Arundhati Banerjee , Rakhi Dasgupta* and Sujay Ray* | Mutational Impact on the Interaction Between Human IL27 and gp130: In silico Approach for Defending HIV Infection | Current HIV Research | 0.802 | 15(5) | 2017 | 327 - 335 |
| 23 | Arundhati Banerjee , Sujay Ray* | Novel insight into mutational impacts and binding mechanism of human glutaminase and glutaminase- interacting protein: A bio-molecular modeling and docking analysis | Gene Reports | NA | 8 | 2017 | 49-60 |
| 24 | Sujay Ray and Arundhati Banerjee | Comparative Binding Mode and Residual Contribution from Lactoferrins (bLF and hLF) and HIV Gp120: An In silico Structural Perspective to Design Potent Peptide Inhibitor for HIV | Current Enzyme Inhibition | NA | 13(3) | 2017 | 226 - 234 |
| 25 | Arundhati Banerjee , Sujay Ray* | Structural Exploration and Conformational Transitions in MDM2 upon DHFR Interaction from Homo sapiens: A Computational Outlook for Malignancy via Epigenetic Disruption | Scientifica, | NA | 2016 | 2016 | 11 pages |
| 26 | Sujay Ray and Arundhati Banerjee | A Proteome-Level Computational Biology Insight on Sequence Based Aggregation Propensity Profile of | Current Biotechnology | NA | 4(1) | 2015 | 46 - 55 |

| Human Hydrolase Structural Enzymes | | | |
|---------------------------------------|--|--|--|
| | | | |

(b) Papers publication in National/ International Conferences: 16 in number

| Sl. No. | Title of paper | Co- author(s), if | Name of the Conference | Date |
|------------|--|--------------------------------------|---|--------------------------------|
| 1. | Identification and targeting Orphan G-Protein Coupled Receptor (GPCR) in mosquito for application of herbal repellent | Krishnendu Adhikary | SASE-2023, BHU, Varanasi (Nesa India) | April 22- 23, 2023 |
| 2. | Molecular Computing and Structural Biology for Interactions in ER α and bZIP Proteins from <i>Homo sapiens</i> : An Insight into the Signal Transduction in Breast Cancer Metastasis | Sujay Ray | FICTA 2015, NIT Durgapur (Springer) | Nov 16, 2015 – Nov 18, 2015 |
| 3. | Structural and Bio- Molecular Interactions in Human Tenascin C and HIV: An <i>In silico</i> Approach to Avert AIDS, for Infants under the Exposure of HIV | Sujay Ray | ICRCICN, Kolkata (IEEE) 2015 | November 20- 22, 2015 |
| 4. | Interactions with Human CD4 Cells Leads to Helix to Coil Transition in HIVgp120 Aiding CCR5 Attachment and Viral Entry: An <i>In silico</i> Structural Biology Approach for AIDS | Sujay Ray | ICSNCS (LNEE) 2016 [JNU, New Delhi] | Feb 25, 2016 - Feb 27, 2016 |
| 5. | Coils to β -Sheets Transitions and Residual Interactions in Human SOX4 and Syntenin Protein: An In silico Insight into the Cytological Regulation | Sujay Ray | MicroCom, NIT Durgapur (IEEE) 2016 | |
| 6. | Bio-Molecular Interactions and Conformational Switches in Human ER β and SRC-1 Protein: A Molecular Docking and Simulation Approach to Suppress Tumor in Breast Cancer Malignancies | Sujay Ray | MicroCom, NIT Durgapur (IEEE) 2016 | |
| 7. | Computational Molecular Analysis of Human Rhodopsin, Transducin and Arrestin Interactions: An Insight into Signal Transduction for Ophthalmology | Tanushree Mukherjee, Sujay Ray | ICIC2 2016 (Springer) [University of Kalyani, Kalyani, | Feb 18, 2016 - Feb 19, 2016 |
| 8. | Structural and Molecular Insight into Human Glutaminase Protein Domain: An In silico Drug Targeting Approach | Sujay Ray, Apurba Dey | ABRMP 2016 (CBS), Kolkata | January 8-9, 2016 |
| 9. | Molecular Modeling and Computing for PDZ Protein (α-1 Syntrophin) from <i>Homo sapiens:</i> A Comparative Optimization Approach for Protein/Drug Affinity | Sujay Ray, Apurba Dey | ICADVC 2016 (Springer) NIT Durgapur | 25-27 February 2016 |
| 10. | Molecular and Protein Interaction Studies for Inhibiting Growth of Human Leukemic Cells: An <i>In silico</i> Structural Approach to Instigate Drug Discovery | Rakhi Dasgupta, Sujay Ray | Biospectrum, 2017 (Springer) | 25th to 26th August 2017 |
| 11. | Residual Exploration into Apoptosis of Leukemic Cells through Oncostatin M: A Computational Structural Oncologic Approach | Rakhi Dasgupta, Sujay Ray | ICCI 2018, Springer | 10-11 Dec,2018 |

| A Limelight on Human Gp130 and its Deleterious Mutations: A Computational Sequence Level Approach for Hepatocellular Carcinomas | ina Basu Ritika Nandi, Puja Ghosh, Sujay Ray | onal Conference on Computing and Communication Systems, I3CS | |
|--|---|--|---------------------------|
| Molecular Level Insight into the Interactions of SoxC and SoxD from Epsilonproteobacteria <i>Sulfurimonas denitrificans</i> : A Bio-Molecular Computational Approach | Angshuman Bagchi. | IC3T 2015, CMR Technical Campus, Hyderabad | 24th to 26th July 2015 |
| A Computational Structural Biology of SoxR and DNA: A Modelling and Interactive Discern to express the sox operon in <i>Pseudaminobacter salicylatoxidans</i> (KCT001) for global sulphur oxidation | Angshuman | ICICA 2014, NIT Durgapur (Springer) | 22-23 December 2014 |
| Study of the effect of pH on adsorption of phenol and catechol by the use of commercial activated carbon and assessment of toxicity reduction | Anuj Kumar, Tamal Mandal, Dalia Dasgupta | | December 27, 2013 |
| Isolation and characterization of phenol degrading bacteria from phenol contaminated industrial wastewater | Anuj Kumar, Tamal Mandal, Dalia Dasgupta. | | December 27 - 30, 2012 |

(c) Books/Monographs/Book chapters written:

| Sl. No. | Name of book/monograph/ Book chapters | Name of Co- author, if any | Year of Publicatio | Publisher with address |
|---------|--|---|-----------------------|--|
| 1 | <i>In silico</i> Perspective into Interactions and Mutations in Human TLR4 and Ebola Glycoprotein: An <i>In silico</i> Insight to Defend Ebola Virus Entry | Sujay Ray | 2016 | IGI Global, Pennsylvania, United States |
| 2 | Positive Regulation via DNA Interaction in Cellular Decisions for Arg to Ala Mutation in SOX11: An Optimized <i>In silico</i> Neuroinformatics Approach | | 2016 | IGI Global, Pennsylvania, United States |
| 3 | Molecular Computing and Residual Binding Mode in ERα and bZIP Proteins from Homo Sapiens: An Insight into the Signal Transduction in Breast Cancer Metastasis | Sujay Ray | 2016 | Advances in Intelligent Systems and Computing book series (AISC,volume 404) |
| 4 | A Limelight on Human Gp130 and Its Deleterious Mutations: A Computational Sequence Level Approach for Hepatocellular Carcinomas | Shreya Bhattacharya, Debina Basu, Ritika Nandi, Puja Ghosh, Sujay Ray | 2021 | Lecture Notes in Networks and Systems book series (LNNS,volume 170) |
| 5 | Molecular and Protein Interaction Studies for Inhibiting Growth of Human Leukemic Cells: An In Silico Structural Approach to Instigate Drug Discovery | Rakhi Dasgupta, Sujay Ray | 2020 | Biotechnological Applications in Human Health. Springer, Singapore. |
| 6 | Residual Exploration into Apoptosis of Leukemic Cells Through Oncostatin M: A Computational Structural Oncologic | Rakhi Dasgupta, Sujay Ray | 2020 | Advances in Intelligent Systems and Computing, vol 988. Springer, Singapore. |

| 7 | Computational Molecular Analysis of Human Rhodopsin, Transducin and Arrestin Interactions: An Insight into Signal Transduction for Ophthalmology | Tanushree Mukherjee, Sujay Ray | 2017 | Advances in Intelligent Systems and Computing, vol 458. Springer, Singapore. |
|---|---|--------------------------------------|------|--|
| 8 | Interactions with Human CD4 Protein Leads to Helix-to-Coil Transition in HIV- gp120 Aiding CCR5 Attachment and Viral Entry: An In Silico Structural Biology Approach for AIDS | Sujay Ray | 2016 | Lecture Notes in Electrical Engineering, vol 396. Springer, New Delhi. |
| 9 | Exploring the Human USP Gene Family and Its Association with Cancer: An In Silico Study | Sujay Ray | 2023 | Advances in Data Science and Computing Technologies pp 685– 694 Lecture Notes in Electrical Engineering book series (LNEE,volume 1056) |

(d) Number of UG and PG students guided THREE (UG) Dissertations