

University of North Bengal

Rajarammohunpur, Siliguri-734013 India

Department of Botany



Dr. Arnab Sen

MSc, PhD, FIAAT, FLS, FNABS **PROFESSOR**

- Fellow of the National Academy of Biological Sciences (FNABS)
- Fellow of the Linnean Society, London, UK (FLS)
- Fellow of Indian Association of Angiosperm Taxonomy (FIAAT)

Members of Learned Societies:

- Society for General Microbiology, UK (2014-2016, 2019)
- American Society for Microbiology (ASM), USA (2007-2008)
- International Society for Computational Biology (ISCB), USA
- Int. Association of Computer Science and Information Technology (Singapore)
- Association of Microbiologists of India (AMI)
- Society for Plant Biochemistry and Biotechnology
- Indian Association for Angiosperm taxonomy (IAAT)
- National Academy of Biological Sciences
- Asian Phytopathological Society

Contact Addresses:

Contact No. +91-9434307487; 7864979539

Mailing Address

Department of Botany, University of North Bengal, PO-NBU, Siliguri-734013, Dist-Darjeeling, West Bengal, India.

Coordinator, Bioinformatics Facility Centre, University of North Bengal

In-charge, Biswa Bangla Genome Centre

IEmail: arnab.nbu@gmail.com; arnab.sen@nbu.ac.in

Subject Specialization (Areas of Research Interest): Nano-technology, Cytogenetics, Meta-genomoics, Bio-informatics, Drug designing and Network pharmacology

No. of Ph.D. students:	(a) Supervised: 21		(b): On-going: <mark>08</mark>		
No. of M.Phil. students:	(a) Supervise	(a) Supervised: N/A		-going: N/A	
No. of Publications:	Total: 195	(a) Journals: 1	87	(b) Books: 02	(c): Chapters: 6.
Impact Factor: 255.79 (2023	3, as per Thomso	n Reuters)			
Citation Index: For latest cit	ation index CLIC	CK HERE			

Country Visited (for Academic purposes): Bangladesh, USA, Canada, Portugal, France, Algeria, Singapore, UK, Tunisia etc. **University Education:**

- PhD from North Eastern Hill University in "Electron Microscopy and Molecular Biology of Frankia" in the year 1997.
- MSc in Botany with specialization in Cytogenetics from Center for Life Sciences Department of Botany, North Bengal University securing **First class second position** in the University in the year 1990.
- Graduation: BSc (Honors) in 1988.

Awards & Recognition:

- **DBT CREST** Award 2011-2012
- DBT Overseas Award 2006-2007
- DST Fast Track Young Scientist Award, 2001
- Principal Investigator, Frankia Genome Project (US Deptt. of Energy sponsored)
- University Silver Medal for securing 1st class second position in MSc.
- Honorary Chief Advisor, Sikkim State Biotech Hub 2012
- Member of the Advisory Board: Bioinformatics Centre, Sikkim Central University, Sikkim.
- Visiting Faculty: Rajiv Gandhi University, 2004
- Visiting Faculty: Vidyasagar University
- Visiting Faculty: Sikkim Central University 2012-2017
- Visiting Faculty: Post Graduate Diploma in Tea Management, NBU

Special Honors & Achievements:

- Establishment of Biswa-Bangla Genome Center at North Bengal University
- Establishment of Bioinformatics Facility at North Bengal University
- Delivered **Constance Ledward Rollins Distinguished Lecture** at Department of Microbiology, University of New Hampshire, USA in the year 2007.
- Member of the PhD defense board, University of New Hampshire, USA
- Coordinator and Member Secretary: DBT-Bioinformatics Facility
- Academic Coordinator: Certificate course in plant tissue culture 2006
- Associate co-Coordinator: Medicinal plants workshop programs 2006
- Reviewer of scientific papers of various National and International Journals.

Editorship of Journals:

- Editor: Indian Journal of Microbiology (a Springer publication) (2006-2011)
- Editor: NBU Journal of Plant Sciences (2010-2012; 2017-2019)
- Guest Editor: Indian Journal of Biotechnology (a CSIR publication)

Fellowships/Scholarships availed

- DBT Overseas associateship to Univ. of New Hampshire, USA, 2005-06 (one year)
- **CREST Fellowship** to University of Lyon, France (6 months)
- Fellowship for a period of 3.5 years, sponsored by IFCPAR (Indo-French Center for the Promotion of Advanced Research) ay NEHU, Shillong
- Fellow of the Linnean Society, London, UK (FLS)
- Research associate-ship in a Indo-US Project on Wheat Biotechnology.
- Fellow of Indian Association of Angiosperm Taxonomy (FIAAT)
- Research associateship in a Transgenic Brassica Project at IARI, New Delhi
- Research associateship in a Wheat Biotechnology Network Project at Meerut University, Meerut

Membership of Learned Societies

- Society for General Microbiology, UK (2014-2016, 2019)
- American Society for Microbiology (ASM), USA (2007-2008)
- International Society for Computational Biology (ISCB), USA
- *Life Member:* International Association of Computer Science and Information Technology (Singapore)
- Life Member: Association of Microbiologists of India (AMI)
- Life Member: National Academy of Biological Sciences (Chennai, India)
- Life Member: Society for Plant Biochemistry and Biotechnology
- Life Member: Indian Association for Angiosperm taxonomy (IAAT)

Member of the Statutory Bodies

- Member of the Court, University of North Bengal
- Member of PhD Defense Board, University of New Hampshire (2007, 2012), USA
- External Member of the Executive Committee of Bioinformatics Centre, Vidyasagar University
- Member of Board of Research Studies (BRS), Department of Zoology and, Department of Botany, CBPBU
- BRS member, Department of Botany, Microbiology, Raiganj University
- BRS member, Department of Biotechnology, North-Eastern Hill University, Shillong
- BRS member, Assam University, Silchar
- Member of the Syllabus Committee, Botany, Diamond Harbour Women's University
- Member of PhD Defense Board, Gauhati University
- Member of the Teacher Recruitment Board, Bodoland University
- Member of the Executive Committee, Bioinformatics Centre, Sikkim Central University
- Member of the Executive Committee, Bioinformatics Centre, TM Bhagalpur University
- Member of PhD Defense Board, Dibrugarh University
- Member of PhD Defense Board, Banaras Hindu University
- Member of PhD Defense Board, BN Mandal University, Madhepura
- Member of PhD Defense Board, Magadha University, Bodhgaya
- Member of PhD Defense Board, Rajsahi University, Bangladesh
- Member of PhD Defense Board, Visva Bharati University, Shantiniketan
- Member of PhD Defense Board, Calcutta University, Kolkata
- Member of PhD Defense Board, Osmania University, Hydrabad
- Member of PhD Defense Board, USTM, Meghalaya

AREAS OF RESEARCH INTEREST

Nanobiotechnology:

Our group is continuously working in phytochemistry and phytomedicines. These types of studies may prove to be helpful in the medicinal field. Phytomedicines derived from herbs and shrubs that possess therapeutic and curative properties have been utilized globally since time immemorial. The North-Eastern region of India is a **"biodiversity**"



hub" (Kar et al., 2016). The entire region is rich in flora and fauna and contains various edible underutilized juicy fruit-bearing plants like Elaeagnus pyriformis, Myrica nagi, Rubas plicatus, etc. Nutritionally, underutilized fruits contain an enormous amount of macro elements such as nitrogen, phosphorus, potassium, calcium, magnesium, and sodium and contain micro-elements which include Iron, Zinc, Copper, and Manganese which promise to be of imtherapeutic significance. mense Native people in Northeast India

use various types of fruits for various purposes, often grown wild and sold in local markets (<u>Kar et al., 2019</u>). *Baccaurea ramiflora*, a popular fruit in East and Northeast India, is known for its ethnomedicinal and high nutritive values. The seeds of *B. ramiflora* are effective against indigestion, constipation, and snake bites, making it an important part of the Rath Yatra in Odisha, India.

Plant and fruit phytocompounds are essential for the **green synthesis of silver nanoparticles**, which are increasingly used as anti-microbial agents. AgNPs, synthesized from secondary metabolites, have antioxidant efficacy and antimicrobial activity against pathogenic microorganisms. Their in-silico molecular docking analysis suggests the potential for cancer combat by manipulating apoptosis (Banerjee et al., 2022) . Underutilized fruit *Phyllanthus acidus is used* for ethnomedicinal purposes. Our recent study compared the efficacy of raw fruit juice and nanoparticles on gentamicin-induced nephrotoxicity in a mouse model. Results showed that high doses of *P. acidus* nanoparticles significantly improved creatinine and urea levels, enhanced antioxidant enzymes, and restored glomerulus structure. The study suggests that PANH is more ameliorative than high doses of *P. acidus* fruit juice in treating kidney injury.

Our study investigated the potential of beta-sitosterol conjugated silver nanoparticles (BSAgNPs) to ameliorate carbon tetracycline-induced liver injury in Swiss albino mice. The results showed that BSAgNPs significantly increased liver

tissue SOD activity, and catalase activity, and reduced nuclear factor erythroid-2-related factor 2 (Nrf2) levels. The antifibrotic effect of BSAgNPs may promote the lowering of chronic inflammation, oxidative stress, and collagen deposition, suggesting that **nanoparticle-mediated drug delivery** of beta-sitosterol may have therapeutic promise against liver complications (Kar et al., 2022a).

India experiences abundant rainfall during monsoon, but groundwater supplies irrigation water, causing As contamination in agricultural fields. Nanoparticle-mediated plant toxicity amelioration offers promising results in recent advancements. Arsenic (As), a toxic metalloid, is a widespread issue in the Earth's crust and is primarily caused by anthropogenic factors such as excessive use of groundwater, pesticides, mining activities, and sewage sludge. India, Bangladesh, and China are more prone to arsenic contamination due to the high As content in their bedrock. Blackgram, a vital pulse crop, is a major producer of black gram, which is high in protein, carbohydrates, and fiber content. We have used Zinc Oxide Nanoparticles (ZnONPs to address heavy

metal and metalloid toxicity in plants, with being a popular solution. **ZnONPs have been used as nano fertilizers** to combat zinc deficiency in agricultural soils, reducing As-accumulation and enhancing growth and photosynthesis in rice seedlings. However, the feasibility of applying ZnONPs to alleviate As-induced stress in black gram has not been documented. The study found that ZnONPs priming altered metabolic pathways decreased ROS generation, and increased osmoregulatory, acting as an alternative source of Zn2+, an important micronutrient. These findings have significant potential for improving the overall health of black gram grown in arsenic-contaminated agricultural fields (Banerjee et al., 2023).

Heavy metals in agricultural soils are a concern for human health and global food security. Nanoparticles have the potential to reduce plant absorption and bioavailability, but environmental toxicity and ecosystem impact remain. Future research should focus on biodegradable and biocompatible NPs, safety, dependability, effectiveness, and long-term impacts.

Research on Covid-19 and Post-Covid:

This group started working on SARS-CoV-2 since the emergence of the novel virus. The first study was conducted *in silico* using natural compounds of *Clerodendrum spp*. to find its potency against the disease (Kar et al., 2020). In the study they found potential drug candidates such as taraxerol that exhibited significant results against S (spike) protein, M^{pro} (main protease) and RdRp (RNA-dependent RNA polymerase) of the virus. It was found in another study carried out by this group that common cold drugs Dextromethorphan along with Prednisolone and Dexamethasone have potential against Covid-19 (Sarkar and Sen, 2020). Well known medicinal plants *Justicia adhatoda*, *Ocimum sanctum* and *Swertia chirata* were extensively studied by this group; in silico studies of compounds anisotine and amarogentin extracted from theses plants showed promising results against spike protein (crucial for host recognition and attachment) and M^{pro} or main protease enzyme (crucial for replication of viral genome) of SARS-CoV-2 (Kar et al., 2020).



Other bio-sources were also studied such as cyanobacteria; cyanobacterial proteins cyanovirin-N, scytovirin and phycocyanin showed promising results in preventing the spike protein, the papain-like protease (PL^{pro}) and the main protease (M^{pro}) of SARS-CoV-2 that helps in the attachment and replication of the virus (<u>Naidoo et al., 2021</u>). Work has also been done on different mutants of the SARS-CoV-2 virus that emerged over the past few years. *In silico* study based on molecular docking and molecular dynamics simulation was carried out to detect inhibitory potential of 605 phytocompounds against spike protein of SARS-CoV-2 wild type as well as mutant variants B.1.1.7 (Alpha), B.1.351 (Beta), P.1 (Gamma), B.1.617.2 (Delta), and B.1.1.529 (Omicron) (Kar et al., 2022).

Some symptoms were found to be persistent in the survivors of Covid-19 disease, known as Post-Covid symptoms or Long-haul Covid-19. This group has studied effectiveness of Cannabidiol (CBD) and Cannabivarin (CVN) obtained from *Cannabis sativa* plant, using molecular docking and simulation, that revealed these compounds' ability to down-regulate post-Covid related central nervous system (CNS) proteins (Sarkar et al., 2021). This can be beneficial in post -covid symptoms treatment strategy. Another *in silico* study carried out by this group suggests that the natural phyto-compounds derived from *Citrus macroptera* could be employed in Post-covid inflammation complexities (Lala et al., 2022). Moreover, the emergence of Mucormycosis as a syndemic or synergistic epidemic in the post-covid phase, caused many casualties. The group explored the possible role of traditional antifungal plants, Neem (*Azadiracta in-dica*) and Turmeric (*Curcuma longa*), in management of Covid-Associated Mucormycosis (Datta et al., 2022). The results suggested a triple-front attack on the fungal pathogens through necrosis inhibition, iron chelation and immunoboosting.

Biological network Analysis:

We work on determination of the constituent compounds present in medicinally important plants. These phytocompounds are bioactive agents that hold much potential. On consumption or application of plant extracts, these compounds entre our body and work through interaction with different enzymes and proteins. To understand these interactions we took the aid of Network pharmacology. It is an emerging subject of systematic drug research in the era of artificial intelligence and big data. Human body is controlled by a network of Proteins and metabolic pathways. Plants, through their phytocompounds, have the ability to control multiple points of these networks and thus, provide a better management of diseases. The less adverse effects, affordability, and easy accessibility highlight their potential in traditional remedies. Thus, identifying the target proteins of medicinal plants has become an important aspect in treatment of numerous diseases. In our lab we have worked on the plants Neem and Turmeric, both of which have



well known antifungal potential. The plants were repurposed against Covid associated Mucormycosis (CAM) or black fungus (Datta et al., 2022). We took a step-by-step approach for identifying the target proteins and ligands associated with Mucormycosis treatment. Functional network analysis and Molecular docking approaches were applied to validate our findings. Quercetin derived from both Neem and Turmeric was found to be one of the main phytocompounds working against Mucormycosis. Along with that, Caffeic acid, Curcumin, Kaempferol, Tetrahydrocurcumin and Myricetin also play a pivotal role in fighting against Black-Fungus. A thorough analysis of our result suggested a triple-

front attack on the fungal pathogens and the approaches are necrosis inhibition, iron chelation and immuno-boosting. Moreover, we have worked on the potential of an anthocyanin rich tea variety, Purple tea, through network pharmacology and have been able to explore it's medicinal properties against lifestyle diseases. We determined it's constituent phytocompounds through GCMS. Our study showed the mechanism of action of Purple tea against diabetes, coronary heart disease, obesity, cancer, etc. Presently we are working on the in silico exploration of the potential of the



plant *Phyllanthus acidus* against Monkey pox virus since the leaves of this plant have a history of being used in treatment of chicken pox and small pox in northeast India. We are also working on the efficacy of Tupistra nutans against diabetes through in silico and in vivo studies. The plant is consumed as vegetable and is also traditionally used in treatment of diabetes in Sikkim and in the hilly region of West Bengal. Another traditionally significant plant, Tacca integrefolia is also being explored for scientific validation of its potential in regulation of menstrual disorders in females through insilico and invitro cell culture studies.

Microbial Genomics:

This lab has pioneered *in-silico* studies on diazotrophic and clinical Actinobacteria since the year 2000. The actinobacteria *Frankia* and its host plant interactions were studied extensively and have collaborations with labs of the University of New Hampshire, USA, University of Lyon, France, University of Tunisia. Niche adaptation, biosynthetic energy cost, and resource partitioning in *Frankia* genomes were studied and the works were published in reputed journals of international acclaim (<u>Sarkar et al, 2017</u>, <u>Sen et al, 2019</u>). The insights into the actinobacteria genomes by the *In-silico* approach have overhauled taxonomy and are recognized by Bergey's Manual (<u>Sen et al, 2014</u>). This group orchestrated several whole-genome sequencing of Actinobacteria *-Frankia, Streptomyces*, N2-fixing *Rhizobium*, etc. and these are available in NCBI database (<u>Rai et al, 2017</u>; <u>Gtari et al, 2017</u>). The lab has to its credit a couple of databases, on Mycobacterium secretome-MycoSec to facilitate Global Health. Another aspect is the study of hostmicrobial interaction through reverse ecology analysis, proteome and meta barcoding studies (<u>Sen et al, 2022</u>, <u>Gueddou et al, 2017</u>). Recently, this group is dealing with the bacterial reverse transcriptase, the nitrogen fixing operon of *Frankia*, Surfacteome of bacteria and so on. Prof. Sen has recently coined a term 'Surfactome' representing all the proteins present in the bacterial surface and has studied the *Frankia* surfactome to a deeper extent. This group is aiming to study about the surfactome and RTs of several other microbial group. Along with that, they have also ventured in the development of the epitope-based vaccine against the Monkey Pox Virus.

Ethno pharmacology and Nano biology:

The lab is actively involved in the elucidation of various phyto-compound from indigenous medicinal plants, underutilized fruits, etc. of the northeast region. Biogenic synthesis of nanoparticles using the extract and their antioxidant, antimicrobial, and anti-cancer properties are studied to tap their efficacy as potential drugs for the future. Bioactive metabolites are also studied both in *in vitro* (molecular docking) and *in vivo* (murine models). Another arena of research is the empowerment of the indigenous communities by promoting their traditional foods and beverages.

Micro-propagation and molecular marker studies:

Micropropagation is the rapid vegetative propagation of plants under in vitro conditions Micropropagation of economically important crops such as tea, turmeric, etc. and barcoding of clones of tea using molecular markers is also another area of interest for this group.

Ethnic fermented beverages consumed by different tribes in North Bengal:

Consumption of alcoholic or fermented beverages had a cultural connotation, since Indus Valley Civilization. The process of preparing ethnic brews has been developed by several tribal communities in India. Fermentation not only increases the shelf life of food and beverages but also influences the texture of the food, nutrition uptake, and aroma. Every tribal community has its native brew. The substrate selection and starter culture preparation differs for every community depending on the tropo geographical region of their habitat. North Bengal is home to different tribal communities such as Toto, Rabha, Orao/Santal, Bodo, Lepcha, Rabha and others. The 'Toto' tribe from West Bengal, India is the primitive mongoloid Indo-Bhutanese isolated endemic group and their livelihood is mainly based on forest



produce 'Eu'(/ju:/), is the most popular millet-based ethnic brew of this tribe and this drink is available only in this small enclave called Toto para. The microbial diversity present in the brew was mainly constituted of probiotic and fermenting bacteria. The nutrient profiling of Eu estimated the amount of protein, carbohydrate, free amino acid, and free fatty acid. The nutritional substance present in the brew probably supports the growth of the probiotic microflora. Reverse ecology-based network analysis revealed considerable complementary interaction between humans and the good bacteria pre-

sent in the Eu. Scientific analysis revealed that Eu is a non-toxic, low-alcohol content brew that has plenty of beneficial probiotic bacteria with prebiotic substances. Eu is a probiotic health drink rather than a recreational drink. 'Haria' or Rice beer is an ethnic, alcoholic, refreshing stable food prepared from low-grade rice by the indigenous experts of rural areas in central and eastern India. Similar to Haria, another rice-based beer consumed by the Rabha tribe in India is known as 'Choko'/'chokot'. It is also nearly impossible to know when the tribe first started preparing this native brew. 'Choko'/'chokot' is a rice-based fermented brew where the starter culture ('Bakhor or phap') is made up of eleven different plants. The metagenomic analysis of the starter culture of chokot showed that several dominant bacterial genera out of this Bacillus, Arthrobacter, Lactobacillus, Ilyobacter, and Lactococcus are prevalent. Commercialization of this type of fermented ethnic non-industrial brew can be an alternative source of income for different tribal communities of North Bengal, India.

Tea Garden Weed Research:

Tea is regarded as one of the most consumed beverages throughout the world but its productivity is at risk because of various pests and infections. Oligonychus coffeae or Red spider mites are the most prevalent pest causing huge losses to the tea industry. The control measures using chemical pesticides have a negative impact on the quality of tea and make it unfit for human consumption. In this context, floral diversity can be explored to effectively control pests.Pests



have been managed using pesticides for as long as there has been agriculture. Pesticides have been used in ancient China, Egypt, Greece, and India from far before 2000 BC.

Pest control methods used by the Sumerians around 4,500 years ago included sulphur dusting. Surapala, a scientist in ancient India, investigated and recommended many plants and other goods with biocidal properties to treat plant illnesses. The well-known "panchamula" (five plant roots), which was widely used at the time, has antiviral, antibacterial, antifungal, and antifeeding properties. The use of synthetic chemical pesticides is either outlawed or regulated owing to the residual issue and consumer health concerns. Pesticide resistance affects at least 500 different insect and mite species. Using botanicals to manage pests is successful and safe pest management.

There are various types of pest that attack on the tea plant like tea mosquito bug, red slug, red spider mite, thrips, jassids etc. Out of them tea red spider mite and tea mosquito bug are more dominant pest in these region and causes huge

In last few years, the world experienced an unprecedented pandemic called COVID-19. The pandemic took 6,709,387 lives and 663,248,631 positive cases (as of 8:18pm CET, 10 January 2023 according to WHO) were reported world wide.

PhytoResp

It has been experienced that the people who had chronic respiratory problems were particularly vulnerable to COVID-19 pandemic. It was actually a syndemic where comorbid people suffered more.

One of the main causes of respiratory problems is viral infection and in such cases no effective medicines are available so far. Herbal drugs on the other hand showed promise in giving relief to respiratory ailments. As a result more and more researchs are tilting towards phytocompounds as a remedy to such problems.

It is, in this context, we at North Bengal University decided to create a database which will provide valuable and effective information on the plants and their constituent compounds which are essentially effective against respiratory disorders. In the present version of the database, we have provided 329 medicinal plants traditionally known to cure respiratory disorders in Darjeeling region along with the parts used.

Also we have taken a step to evaluate the efficiency of the phytocompounds through in silico methods.





PhytoResp HOME BROWSE STATISTICS PHY crop lose on tea industry in every year. The objective of our work is to acquire information on the pest control status of Dooars and Terai concerning botanical formulations used against Tea pests and select new plant species effective for pest control.

PHYTORESP-a Database

In last few years, the world experienced an unprecedented pandemic called COVID-19. The pandemic took 6,709,387 lives and 663,248,631 positive cases

(as of 8:18pm CET, 19 January 2023 according to WHO) were reported world wide. It has been experienced that the people who had chronic respiratory problems were particularly vulnerable to COVID-19 pandemic. It was actually a syndemic where comorbid people suffered more.

One of the main causes of respiratory problems is viral infection and in such cases no effective medicines are available so far. Herbal drugs on the other hand showed promise in giving relief to respiratory ailments. As a result more and more researches are tilting towards phytocompounds as a remedy to such problems.

It is, in this context, we at North Bengal University decided to create a database, named PhytoResp which will provide valuable and effective information on the plants and their constituent compounds which are essentially effective

Technology generation:

• Novel techniques for isolation of *Frankia* in pure culture and generation of single spore cultures of *Frankia*.

Research projects (recently carried out/ongoing):

- Establishment of Biswa-Bangla Genome Centre
- Indo-Tunis Project on "Phytoremediation of Heavy Metal-Contaminated Soils in Tunisia and India using Frankia-Alnus symbiosis". 2017-2019 (Two Years)
- A Bioinformatics perspective to impede the tubercle bacillus (TB) devastation in West Bengal (Three Years)
- Frankia Genome Project (Sequencing and analysis of FIVE Frankia genomes, Sponsored by US Department of Energy, Govt. of United States of America) (Principal Investigator)
- Development of novel methodology for understanding the mechanism of nitrogen fixation through codon usage analysis
 related genes and protein structure modeling of relevant gene product (protein) and its implication DBT, Govt. of India
 sponsored (Co-PI)
- Establishment of Bioinformatics Facility. Sponsored by DBT, New Delhi. (Principal coordinator)
- Production of Transgenic tea for fungal resistance. Sponsored by DST, New Delhi (completed) (Principal Investigator)
- Detection of Best alder-Frankia symbiosis using molecular markers. Sponsored by UGC, New Delhi (Principal Investigator)
- Rehabilitation of the Tissue Culture Laboratory. Sponsored by Department of Horticulture, Government of West Bengal, India. (Principal Investigator)
- Survey, conservation, development of propagation techniques and popularization of cultivation of medicinal plants in Terai, Dooars, Hills of North Bengal and Sikkim. NMPB, Government of India sponsored. (Co-Principal Investigator)

PhD Supervised/Supervising

- Molecular characterization of Frankia and alder-Frankia symbiosis in Eastern India- Balwinder Singh Bajwa (Awarded)
- Study of genetic diversity and standardization of genetic transformation in Camellia sinensis (L.) O. Kuntze- Manprit Gill (Awarded)
- Studies of codon usage, prteome analysis and evolution of nitrogen fixing genes in some microorganisms- A bioinformatic approach- Saubashya Sur (Awarded)
- Genomics of some pathogenic food bacteria and molecular modeling of their important oxins and their interactions- Uttam Kumar Mondal (Awarded)
- Genetic diversity of Frankia associated with Hippophae L. in Lachen valley of North Sikkim. Bharat Basistha (Awarded)
- Diversity of Frankia associated with Alnus nepalensis and Casuarina equisetifolia in West Bengal. Debadin Bose. (Awarded)
- In silico characterization of some nitrogenase proteins found in symbiotic diazotrops and cyanobacteria -Subarna Thakur (Awarded)
- Study of genetic diversity and micropropagation of bamboos growing in North Bengal- Arvind Kumar Goyal (Awarded)
- Diversity study of Rhizobium from North Bengal and Sikkim through Biochemical and Molecular characterization and exploration of their role as biofertilizer- Ritu Rai (Awarded)
- Diversity and micropropagation of Canna from West Bengal and Orissa- Tanmayee Mishra (Awarded)
- Micropropagation, diversity study and detection of antioxidants in some medicinal Zingibers- Malay Bhattacharya (Awarded)
- Medicinal and molecular documentation of some members of Mimosaceae and their micro-symbionts- Manas Ranjan Saha (Awarded)

- Riddles of human intestinal microflora- a bioinformatics perspective.- Ayan Roy (Awarded)
- Characterization and diversity of selected Actinorhizal haemoglobin genes and proteins with reference to Alnus-Frankia symbiosis- Sanghati Bhattacharya (Awarded)
- Relationship of Multiple Tannase Sequences and Structures through Bio-informatics- Amrita Banerjee (Awarded)
- Studies of molecular diversity and chemical properties of selected medicinal members under the genus Clerodendrum L.-Pallab Kar (Awarded)
- Comparative and Evolutionary Genomics of Selected Actinobacteria with Special Reference to Pathogenicity- Shilpee Pal (Awarded)
- Bioinformatic characterization of selected Actinobacteria with a focus on biotopic diversity- Indrani Sarkar (Awarded).
- Molecular detection and diversity analysis of Bipolaris sorokiniana in sorghum and induction of systemic resistance by bioinoculants- Priyanka Bhattacharya (Awarded)
- Medicinal and molecular profiling of selected tea varieties of Darjeeling and Dooars- Reha Labar (Awarded).
- Medicinal and molecular profiling of selected members of Araliaceae- Arnab Chakraborty (Registered).
- Biochemical, molecular and genomic studies of selected Streptomyces isolates from tea garden soils of North Bengal- Saroja Chhetrri (Registered).
- Medicinal and molecular profiling of selected Rutaceous members with a focus on Citrus L.- Mousikha Lala (Registered).
- Ethnomedicinal and molecular profiling of the fermented beverages used by different tribes in North Bengal, India Soumita Bhattacharjee
- STUDIES ON THE SELECTED TEA GARDEN WEEDS FOR THE PROMOTION OF TEA PLANT HEALTH -Sandipan Ghosh
- Selected underutilized fruits of north east India for overall health improvement of humankind-SWARNENDRA BANERJEE

Selected Post-Graduate Dissertations:

- Isolation and characterization of the Streptomyces sp from the rhizospheric soil of Areca catechu of plants of jaigaon, west Bengal -Abisekh Karki (2023)
- Ayurveda plants against respiratory disease and their anti-inflammatory efficacy- Ananya Banik (2023)
- Phytoresp: a curated database of plants against respiratory disease- Anushka Das (2023)
- In vitro antioxidant and antimicrobial activities of Elaeagnus pyriformis (silverbeery) fruit extract and its effective role against testicular damage, oxidative stress linked to male infertility.- Usashi Shome (2023)
- Improvement of Pisum sativum (I) yield under arsenic stress potential role of the zinc oxide nanoparticles in abiotic stress management- *Rajarshi Sarkar* (2023)
- In vitro and in silico evaluation of medicinal plants used in starter culture (ranu dabai) of fermented rice beer by the inhabitants of tea garden of terai region of west Bengal- *Prasant Guptta* (2022)
- Selected native plants of sub-Himalayan West Bengal effective against neurological disorders as revealed by network pharmacology- *Thering Dhendup Lepcha* (2022)
- Isolation, identification, and characterization of the root nodule associating endophytic diazotrophs from the leguminous plants growing near the tea garden area of the University of North Bengal- *Anipa Saha* (2022)
- Effect of Arsenic contaminated soil in vigna moong plants and its amelioration with Zinc oxide nanoparticle priming Jarzis Islam (2022)
- Mitigation of Arsenic toxicity in early growth phase of vigna moong with the help of zinc oxide nanoparticles priming-*Sourik Mondal* (2022)

- In silico screening of chemical compounds of Citrus in different disease-related targets- Anup Barman (2021)
- Fruit juice of some Actinorhizal plants (Myrica nagi, M. esculanta, Eleagnus pyriformis) may help in reducing nephrotoxicity in kidney- *Babita Sah* (2021)
- Anti-inflammatory, anti-cancerous, anti-oxidant, anti-fungal, anti-microbial, and CNS stimulant activity evaluation through molecular docking studies of phytochemicals found in 5-AV2 and 1-ASSAM cultivars of Camellia sinensis - *Reshmi Sarki* (2021)
- In silico investigations of chemical constituents of Clerodendrum species in the anti-cancer, anti-inflammatory, anti-diabetic, hepatoprotective, anti-oxidant, and anti-bacterial drug targets- *Sangita Bose* (2021)
- Synthesis of silver nanoparticles using Ficus rumphii leaf extract and their antioxidant activity-Sagarika Ray (2021)
- Comparative study on antioxidant property of distilled alcohol made from Rhododendron arboreum sm. and Zingiber officinale consumed by ethnic groups of Darjeeling Hills- *Anjana Sharma* (2020)
- Biogenic synthesis of silver nanoparticles using Clerodendrum inerme leaf extract and antioxidant activity- *Moumita Barman* (2020)
- Optimization and characterization of silver nanoparticle synthesized using green approach A review *Pinki Sutradhar* (2020)
- Characterization and application of green synthesis of metallic sliver nanoparticles (AgNps) A review- Debarati Saha (2020)
- Bioinformatics perspective of Atopobium sp.- Dipayan Pal (2019)
- Production and biochemical characterization of Averrhoe carambola (Kamranga) wine- an underutilized fruit and beverage-Mampi Mondal (2019)
- Antioxidant and medicinal profiling of three members of Chlorophyceae- Deep Deb (2019)
- Isolation and characterization of Streptomyces strains Namita Das (2019)
- Antioxidant Profiling of Schefflera arboricola Uttam Roy (2018)
- Antioxidant Profiling of Trevesia palmate- Rahul Kumar Prasad (2018)
- Isolation and characterization of Actinobacterial strains from Tea Garden of NBU Campus Sandipan Ghosh (2018)
- Comparative antioxidant activity of four different cultivars of Ziziphus sp.- Sourav Paul (2018)
- Small RNAs of some selected species of Actinobacteria in light of Pseudoknot, free energy, and target protein classes: a bioinformatics approach- *Md Sohaib* (2017)
- Codon usage pattern of carbohydrate-active enzymes (CAZy) in Actinobacteria: an in silico approach- *Priyashree Bhadra* (2017)
- Investigation of codon usage pattern on CAZyme of Actinobacteria- Frankia, Leifsonia, and Streptomyces genome *Riya Das* (2017)
- Small RNAs of some selected species of Actinobacteria in light of Pseudoknot, free energy, and target protein classes: a bioinformatics approach- Satyaki Biswas (2017)
- Molecular diversity of 18 tea [Camellia sinensis (L.) Kuntze] varieties in North Bengal with RAPD and ISSR markers Suman Seth (2016)
- In silico drug targeting against tubercle Bacillus from India- Dwipanwita Adhikari (2016)
- Antioxidant profiling of Marsilea minuta Linn.- Alokejyoti Biswas (2016)
- High profile medicinal properties screening of selected home and country-made wines Mousikha Lala (2015)
- In-silico study on heat shock proteins of selected Actinobacteria Md. Azmalul Alam (2015)

• Altitudinal diversity study of Alnus nepalensis D.DON with RAPD markers - Mamta Kumari Chhetri (2015)

Publications:

GenBank submission (Genome Projects):

- Lucas,S., Han,J., Lapidus,A., Cheng,J.-F., Goodwin,L., Pitluck,S., Peters,L., Mikhailova,N., Teshima,H., Detter,J.C., Han,C., Tapia,R., Land,M., Hauser,L., Kyrpides,N., Ivanova,N., Pagani,I., Beauchemin,N., Sen,A., Gtari,M., Wall,L., Tisa,L. and Woyke,T. Improved High-Quality Draft sequence of Frankia sp. QA3. http://www.ncbi.nlm.nih.gov/nuccore/CM001489.
- Lucas, S., Han, J., Lapidus, A., Cheng, J.-F., Goodwin, L., Pitluck, S., Peters, L., Mikhailova, N., Teshima, H., Detter, J.C., Han, C., Tapia, R., Land, M., Hauser, L., Kyrpides, N., Ivanova, N., Pagani, I., Beauchemin, N., Sen, A., Gtari, M., Wall, L., Tisa, L. and Woyke, T. The draft genome of Frankia sp. CN31. http://www.ncbi.nlm.nih.gov/nuccore/NZ_AGJN00000000.1
- Lucas, S,... Sen, A, et al. (2011). Frankia sp. Eul1c chromosome, complete genome. http://www.ncbi.nlm.nih.gov/nuccore/ NC_014666.1
- Lucas, S,Sen, A, Fernandez, http://www.ncbi.nlm.nih.gov/nuccore/NZ_AGJN00000000.1 M and Tisa,L (2011). Frankia sp. EUN1f ctg00163, whole genome shotgun sequence. http://www.ncbi.nlm.nih.gov/nuccore/NZ_ADGX01000396.1
- Rai,R., Swanson,E., Sarkar,I., Sen,A., Lama,D., Abebe-Akele,F.,Thomas,K., Morris,K., Simpson,S. and Tisa,L.S., Permanent Draft Genome Sequence of the French Bean Symbiont Rhizobium sp. Strain RSm-3 Isolated from the Eastern Himalayan Region of India. https://www.ncbi.nlm.nih.gov/nuccore/MAWZ00000000
- Chhettri,S., Sevigny,J.L., Sen,A., Ennis,N. and Tisa,L., Genome sequences of Streptomyces and Rhizobium isolates collected from root and soil. https://www.ncbi.nlm.nih.gov/nuccore/RZYA00000000.1
- Chhettri,S., Sevigny,J.L., Sen,A., Ennis,N. and Tisa,L., Genome sequences of Streptomyces and Rhizobium isolates collected from root and soil. https://www.ncbi.nlm.nih.gov/nuccore/RZXZ01000001
- Yamashita,T., Arnab,Sen., Sevigny,J.L., Ennis,N., Saroja,C., Yamashita,T., Thomas,W.K., Thomas,D., Alam,M., Alroobi,R.M., Atkinson,E.C., Baer,N., Bieser,K., Blouin,N., Brogan,L.J., Chen,J., Edgington,N.P., George,O.L., Heda,G.D., Howerton,A., Luek,J., Mazzer,P., Miller,K., Moore,D.P., Page,S.T., Roe,J.L., Shuman,K.E., Townsend,K. and Lou,T., Genome sequencing of Pseudomonas sp. SDr-06. https://www.ncbi.nlm.nih.gov/nuccore/QOLA00000000
- Sen,G., Sen,A., Chhettri,S., Sarkar,I. and Bhattacharya,M. Whole genome sequencing of Streptomyces sp. MAG02. https://www.ncbi.nlm.nih.gov/nuccore/jaignw000000000.1
- Sen,G. and Sen,A. Whole genome sequencing of Streptomyces actuosus VRA1 https://www.ncbi.nlm.nih.gov/nuccore/ NZ_JAFFZS000000000.1
- Sen,G., Sen,A., Chhettri,S., Sarkar,I. and Bhattacharya,M. Whole genome sequencing of Streptomyces sp. DHA010. https:// www.ncbi.nlm.nih.gov/nuccore/jaignv000000000.1
- G Sen, A Sen, I Sarkar, Bhattacharya, M. and Chettri, S. Whole genome sequencing of Streptomyces species from India. https://www.ncbi.nlm.nih.gov/nuccore/NZ_JAFVLN010000042.1

GenBank submission (Partial sequences):

Bamboo	: 26
Canna	: 08
Turmeric	: 02
Frankia	: 22
Tea (Camellia)	: 40
Clerodendrum	: 29
Streptocaulon sylvestre	: 03
Food sample (Eu)	: 01
Yeast sample	: 03

Database created:

- Bamboo-infoline: a database for North Bengal Bamboos (www.bamboodb.ind.in)
- MycoSec: a database for Mycobacterium Secretome analysis (www.bicnbu.in/mycosec)

Books:

- Biology of Useful Plants and Microbes by Arnab Sen (ed), Narosa Publication House, New Delhi.
- Medicinal plants by AP Das, MU Alam and Arnab Sen et al.

Research Papers & Book Chapters

Total Impact Factor: 255.79 as of October. 2023 (Thomson Reuters)

2023

- Chhettri S, J Sevigny, C Pesce, I Sarkar, T Kelley, I Nouioui, G Sen, LS Tisa, A Sen (2023). Whole genome sequencing of *Streptomyces antnestii* sp. nov. with a potency to become an industrial strain. *J Genomics* (*accepted*).
- Banerjee S, J Islam, S Mondal, A Saha, B Saha, A Sen (2023). Proactive attenuation of arsenic-stress by nano-priming: Zinc Oxide Nanoparticles in Vigna mungo (L.) Hepper trigger antioxidant defense response and reduce root-shoot arsenic translocation. J *Hazardous Materials*. https://doi.org/10.1016/j.jhazmat.2023.130735 IF :13.6
- Bhattacharjee, S., Sharma, A., Lala, M., Gupta, P., Ghosh, C., & Sen, A. (2023). Efficacy of 'Harchur raksi', a traditional fermented beverage of high altitude Darjeeling hills as an anti-inflammatory and antiviral remedy. Vegetos, 1-9. https://doi.org/10.1007/s42535-023-00681-7
- Bhattacharjee, S., Lala, M., Gupta, P., Ghosh, C., & Sen, A. (2023). In vitro, in silico and chemical profiling of Timbur Raksi, a lesser-known ethnic fermented beverage from Darjeeling Himalaya. Res. Jr. Agril Sci, 14(2), 381-384.
- Bhattacharjee, S, C Ghosh, A Sen, M Lala (2023). Characterization of *Firmania colorata* (Roxb.) leaf extract and its silver nanoparticles reveal their Antioxidative, Anti-microbial, and Anti-inflammatory properties. *Int. Nano Lett.* https://doi.org/10.1007/s40089-023-00392-6
- <u>Bhattacharjee, S., Sarkar, I., Sen, G., Ghosh, C., & Sen, A. (2023</u>). Biochemical and Metagenomic sketching of microbial populations in the starter culture of 'Chokot', a rice-based fermented liquor of Rabha Tribe in North Bengal, India. *Ecological Genetics and Genomics*, 29, 100193. <u>https://doi.org/10.1016/j.egq.2023.100193</u>
- Datta S, R Labar, I Sarkar, A Sen (2023) GC-MS profiling and *in silico* polypharmacology establish antioxidant rich Purple tea as a major functional beverage against various lifestyle diseases. *Lett. Appl. NanoBio Sc.* <u>https://doi.org/10.33263/LIANBS120.000</u>.
- Datta, S., Sarkar, I., Ghosh, S., Goswami, S., Sen, G. and Sen, A., (2023). The faster evolution of signal peptide genes of Frankia in surfactome may be attributed to their cardinal role in symbiotic association. Symbiosis, pp.1-8. <u>https://doi.org/10.1007/s13199-023-00935-7</u> IF: 2.5
- <u>Ghosh S, Sen G, Sarkar I, Ghosh C, Sen A</u>. (2023). Molecular docking studies to validate the efficacy of the indigenous Bioformulation of North East India against the reproductive proteins of the pest, Tetranychus urticae. **Res. Jr. Agril. Sci**. 14(5): 1259-1264.
- <u>S Ghosh, S Sengupta, G Sen, C Ghosh, A Sen*</u>, (2023) Acaricidal and Ovicidal effects of two common weeds of tea gardens of North Bengal, India against *Oligonychus coffeae*. Journal of research im weed science, 6(2), 40-49. <u>http://dx.doi.org/10.26655/JRWEEDSCI.2023.6.5</u>

- <u>Ghosh, Sandipan; Sarkar, Indrani; Sen, Arnab</u> (2022). Persepectives on Extremophilic Actinobacteria-A Review. NBU Journal of Plant Sciences, Vol. 14 (2022), pp. 1-7
- Das, Anushka; Banik, Ananya; Datta, Sutapa; Sen, Arnab (2022). PhytoResp : A Database for Medicinal Plants of Darjeeling, Against Respiratory Ailments, Vol. 14 (2022), pp. 61-67
- Lungphi P, S Bhattacharjee, R Labar, A V Singh, C Ghosh, AP Das, A Sen (2022). 'Bitter the Better' A comprehensive evaluation of pharmacological compounds in the bitter tea Phalap-khah of the Tangsa tribe in Arunachal Pradesh (India). J. Botanical Soc. Ben. 76(2) : 62-69.
- Sarkar I, G Sen, MA Ali, SM Almutairi, J Lee, A Sen (2022). Detection of the peptidyl epitope for vaccine development against MPV. Journal of King Saud University-Science. doi: 10.1016/j.jksus.2022.102458. IF: 3.829
- Fernandes I, OS Paulo, I Marques, I Sarkar, A Sen, I Graça, K Pawlowski, JC. Ramalho, AI. Ribeiro-Barros (2022). Salt stress tolerance in Casuarina glauca: insights from the branchlets transcriptome. Plants DOI: 10.3390/plants11212942 IF*:4.658
- <u>Banerjee, S., Kar, P., Islam, R., Naidoo, D., Roy, A., Sarkar, I., Sen, G., Saha, T., Yasmin, H. and Sen, A., (2022).</u> Synthesis of silver nanoparticles from secondary metabolites of star gooseberry fruit (*Phyllanthus acidus*) and their nephroprotective efficiency. South African Journal of Botany, 151, pp.385-395. DOI: 10.1016/j.sajb.2022.10.021 IF: 3.11
- <u>Kar P, Banerjee S,Sen A, Naidoo D, Roy A, Choi YE. (2022).</u> β-sitosterol conjugated silver nanoparticle-mediated amelioration of CCl4induced liver injury in swiss albino mice. J King Saud Univ-Sc. https://doi.org/10.1016/j.jksus.2022.102113 IF: 3.829
- Sarkar I, G Sen, S Bhattacharyya, M Gtari, A Sen (2022). Inter-cluster competition and resource partitioning may govern the ecology of Frankia. Archives of Microbiology. DOI: 10.21203/rs.3.rs-1183956/v1 IF: 2.8

- Kar P, Saleh-E-In MM, Sen A, Naidoo D, Roy A, Choi YE. (2022) Computational profiling of natural compounds as promising inhibitors against the spike proteins of SARS-CoV-2 wild type and the variants of concern, viral cell-entry process, and cytokine storm in COVID-19. *J Cell Biochem.* https://doi.org/10.1002/jcb.30243. IF: 2.6
- Lala M, S Bhattacharya, A Sen, I Sarkar (2022). In-silico studies on wild orange (*Citrus macroptera* Mont.) compounds against COVID-19 proinflammation targets. Journal of Biomolecular Structure and Dynamics, DOI:10.1080/07391102.2022.2051744. IF: 4.4
- Datta S, I Sarkar, G Sen and A Sen. (2022). Neem and Turmeric in the management of Covid associated Mucormycosis (CAM) derived through network pharmacology. *Journal of Biomolecular Structure and Dynamics*, http://dx.doi.org/10.1080/07391102.2022.2048077. IF: 4.4
- Sen G, I Sarkar, S Chetrri, P Kar, A Roy, A Sen, M Bhattacharya (2022). Rhizospheric soil metabarcoding analysis of Alnus nepalensis from Darjeeling hills reveals the abundance of nitrogen-fixing symbiotic microbes. J Forest Res. https:// doi.org/10.1080/13416979.2022.2037813 IF: 1.5
- Sarkar I, A Sen (2022). Niche adaptation of *Frankia* do not drastically influence their metabolic profiling. *J Forest Res.* http://dx.doi.org/10.1080/13416979.2022.2028707. IF: 1.5

2021

- Sarkar I, P Kar, G Sen, S. Chhetri, M Bhattacharya, S Bhattacharyya, A Sen (2021). Metagenomic outlooks of microbial dynamics influenced by organic manure in tea garden soils of North Bengal, India. Archives of Microbiology. DOI: 10.1007/s00203-021-02635-6. IF: 2.8
- <u>Gueddou A, I Sarker, A Sen, F Ghodhbane-Gtari, D R Benson, J Armengaud, M Gtari (2021)</u>. Effect of actinorhizal root exudates on the proteomes of *Frankia soli* NRRL B-16219, a strain colonizing the root tissues of its actinorhizal host via intercellular pathway. *Research in Microbiology*. DOI: https://doi.org/10.1016/j.resmic.2021.103900 IF: 2.6
- Nandi S, P Upadhyay, A Roy, A Dasgupta, A Sen, A Adhikary, K Acharya. (2021). A natural derivative from ethnomedicinal mushroom potentiates apoptosis, autophagy and attenuates cell migration, via fine tuning the Akt signaling in human Lung Adenocarcinoma Cells (A549). Environmental Toxicology https://doi.org/10.1002/tox.23377. IF: 4.5
- Banerjee S, S Islam, A Chatterjee, A Sen and P Kar. (2021). Synthesis of silver nanoparticles using underutilized fruit Baccaurea ramiflora (Latka) juice and its biological and cytotoxic efficacy against MCF-7 and MDA-MB 231 cancer cell lines. South African J. Bot. https:// doi.org/10.1016/j.sajb.2021.09.016. IF: 3.111
- Bhattacharjee S, I Sarkar, P Kar, A Sen (2021). Biochemical and microbial profiling establish "Eu" (a traditional fermented beverage of Toto people) as a probiotic health drink. *J Ethnic Food* https://doi.org/10.1186/s42779-021-00093-5
- Naidoo D, P Kar, A Roy, T Mutanda, J Bwapwa, **A Sen**, A Anandraj (2021). Structural insight into the binding of cyanovirin-N with the spike glycoprotein, Mpro and PLpro of SARS-CoV-2: protein-protein interactions, dynamics simulations and free energy calculations. *Molecules* https://doi.org/10.3390/molecules26175114. IF: 4.927.
- Banerjee S, Kar P, Sarkar I, Chhetri A, Mishra DK, Dutta A, Kumar A, Sinha B, Sen A. (2021). Structural elucidation and chemical characterization of underutilized fruit silverberry (*Elaeagnus pyriformis*) silver nanoparticles playing a dual role as anti-cancer agent by promoting apoptosis and inhibiting ABC transporters. *South African J. Bot.* https://doi.org/10.1016/j.sajb.2021.06.029. IF: 3.111.
- <u>Kar P, Banerjee S, Chhetri A, Sen A (2021)</u>. Synthesis, physicochemical characterization and biological activity of synthesized Silver and Rajat Bhasma nanoparticles using medicinally important Clerodendrum inerme. *Journal of Phytology*. 13: 64-71. DOI:https:// doi.org/10.25081/jp.2021.v13.7026.

- Kar P, V Kumar, B Vellingiri, A Sen, N Jaishee, A Anandraj, H Malhotra, S Bhattacharyya, S Mukhopadhyay, M Kinoshita, V Govindasamy, A Roy, D Naidoo, MD Subramaniam (2020) Anisotine and amarogentin as promising inhibitory candidates against SARS-CoV-2. Journal of Biomolecular Structure and Dynamics,. doi: 10.1080/07391102.2020.1860133 IF: 4.4
- Bhattacharya S, A Sen and M Bhattacharya (2020). Ecology and population genetics studies of *Alnus Nepalensis* in subHimalayan West Bengal and Sikkim. *Int. J Lead Res Pub* 1(3) 1-9
- Sarkar I and A Sen (2020). In silico screening predicts common cold drug Dextromethorphan along with Prednisolone and Dexamethasone can be effective against novel Coronavirus disease (Covid-19). Journal of Bimolecular Structure and Dynamics. DOI: 10.1080/07391102.2020.1850528. IF: 4.4
- Kar P, Dutta S, Chakraborty AK, Bhattacharya M and Sen A (2020). Leaf extract of ethnomedicinally important Bharangi (Clerodendrum

serratum) may improve neuromodulatory activity in mice model. Indian Journal of Traditional Knowledge, . IF: 0.8

- Labar R, Kar P, Biswas P, Sen A and Bhattacharya M (2020). Evolution of matK gene among the elite tea clones (*Camellia sinensis*) revealed by nucleotide substitution within the consensus region. *Journal of Applied Biology and Biotechnology*.DOI: <u>10.7324/JABB.2021.9105</u>
- Kar P, Mishra D, Roy A, Chakraborty AK, Sinha B and Sen A (2020). Elucidation of phytomedicinal efficacies of *Clerodendrum inerme* (L.) Gaertn. (Wild Jasmine). South African Journal of Botany, IF: 3.111
- <u>Kar P, Sharma NJ, Singh B, Sen A and Roy A</u> (2020). Natural compounds from Clerodendrum spp. as possible therapeutic candidates against SARS-CoV-2: an in silico investigation. Journal of Biomolecular Structure and Dynamics, https:// doi.org/10.1080/07391102.2020.1780947. IF: 4.4
- Lala, M, D Modak, S Paul, I Sarkar, A Dutta, A Kumar, S Bhattacharya and A Sen. 2020. Potent bioactive methanolic extract of wild orange (Citrus macroptera Mont.) shows antioxidative, anti-inflammatory, and antimicrobial properties in in- vitro, in vivo, and *in silico* studies. Bulletin of the National Research Centre.
- Misra AK and A Sen (2020) Frankia the endo-micro-symbiont of Hippophae sp. in "Seabucthron", Sharma PC (ed.) Springer pub.
- Sarkar I, G Sen and **A Sen** (2020) Methods for whole-genome analysis of Actinobacteria through Bioinformatics approaches. In Methods in Actinobacteriology, Springer Protocols Handbooks (Springer Protocols), Springer Nature, New York.

2019

- Labar R, Sen A, Bhattacharya M., 2019. Effect of solvents on qualitative and quantitative phytochemical constituent profiles of fresh leaves of TV26. NBU Journal of Plant Sciences, 11: 115-123.
- Labar R, Sarkar I, Sen A, Bhattacharya M., 2019. Effect of solvent with varying polarities on phytochemical extraction from mature tea leaves and its evaluation using biochemical, antimicrobial and in-silico approaches. International Research Journal of Pharmacy. DOI: 10.7897/2230-8407.1008247.
- Kar P, Chakraborty AK, Dutta S, Bhattacharya M, Chaudhuri TK and Sen A (2019) Fruit juice of silverberry (Elaeagnus) and bayberry (Myrica) may help in combating against kidney dysfunctions. Clinical Phytoscience, 5: 1-9.
- Kar P, Chakraborty AK, Bhattacharya M, Mishra T and Sen A (2019) Micropropagation, genetic fidelity assessment and phytochemical studies of Clerodendrum thomsoniae Balf. f. with special reference to its anti-stress properties. Research in Plant Biology, 9: 9-15.
- <u>Chakraborty AK, Sarkar I and Sen A. (2019).</u> Herbal medicine meets bioinformatics for remedy of Tuberculosis by Mycobacterium tuberculosis RGTB423. International Journal of Data Mining and Bioinformatics. DOI: 10.1504/IJDMB.2019.100619. IF: 0.339

- Dey P, Saha MR,...Sen A...Chaudhuri T (2018) Oleander stem and root standardized extracts mitigates acute hyperglycaemia by limiting systemic oxidative stress response in diabetic mice. Advances in Pharmacological Sciences, 2019: 1-12.
- Paul, S, Modak, D, Chakraborty, AK, Sen, A, & Bhattacharjee, S (2018). In vivo and in silico approaches to investigate the toxicological and analgesic properties of unprocessed Aloe vera gel in experimental rat models. Archives of Biological Sciences. https://doi.org/10.2298/ ABS180524036P . IF: 0.856
- <u>Kar P, Dutta S, Chakraborty AK, Roy A, Sen S, Kumar A, Lee J, Chaudhuri TK, and Sen A (2018).</u> The antioxidant rich active principles of *Clerodendrum* sp. controls haloalkane xenobiotic induced hepatic damage in murine model. *Saudi Journal of Biological Sciences*, 26: 1539-1547. Impact Factor: 3.111
- Sen, A., Tisa, L. S., Gtari, M., & Sarkar, I. (2018). Contrasted evolutionary constraints on carbohydrate active enzymes (CAZymes) in selected Frankia strains. Antonie van Leeuwenhoek, 1-11. IF: 2.674
- Gtari, M., Nouioui, I., Sarkar, I., Ghodhbane-Gtari, F., Tisa, L.S., Sen, A. and Klenk, H.P., An update on the taxonomy of the genus Frankia Brunchorst, 1886, 174 AL. Antonie van Leeuwenhoek, pp.1-17. IF: 2.674
- Sarkar, I., Gtari, M., Tisa, L.S. and Sen, A., (2018). A novel phylogenetic tree based on the presence of protein domains in selected actinobacteria. Antonie van Leeuwenhoek, pp.1-7. IF: 2.674
- Mishra DK, UK Singha, A Das, S Dutta, P Kar, A Chakraborty, A Sen, and B Sinha (2018). DNA binding, Amelioration of Oxidative stress and

molecular docking study of Zn(II) metal complex of a new Schiff base ligand. J Coord Chem. https:// doi.org/10.1080/00958972.2018.1476687. IF: 1.9

- Dutta S, AK Chakraborty, P Dey, P Kar, P Guha, S Sen, A Kumar, A Sen, TK Chaudhuri (2018) Amelioration of CCl₄ induced liver injury in swiss albino mice by antioxidant rich leaf extract of *Croton bonplandianus* Baill. *PLOS One* DOI: 10.1371/journal.pone.0196411. IF: 3.752
- Dutta S, A Chakraborty, P Kar, A Sen and TK Chaudhury (2018) Stimulation of Murine Immune Response by *Clerodendrum infortunatum*. *Pharmacogn Mag* DOI: 10.4103/pm.pm_549_17 . IF: 0.7
- Saha MS, S Pal, I Sarkar, A Roy, PKD Mohapatra and A Sen (2018). Comparative genomics of Mycobacterium reveals evolutionary trends of *M. avium* complex. *Genomics* doi: 10.1016/j.ygeno.2018.02.019. IF: 4.4
- Roy A, Sen A, Chakraborty S and Sarkar I (2018). Comprehensive profiling of functional attributes, virulence potential and evolutionary dynamics in mycobacterial secretomes. *World J Microbiol Biotechnol.* 34:5 https://doi.org/10.1007/s11274-017-2388-1. IF: 4.1
- Saha MR, P Kar, A Sen (2018). Assessment of phytochemical, antioxidant and genetic diversities among selected medicinal plant species of Mimosoidae (Mimosaceae). Ind. J Trad. Knowled. 17: 132-140 IF: 0.8
- Saha MR, Dey P, Sarkar I, Sarker D, Haldar B, Chaudhuri TK, Sen A.(2018) Acacia nilotica leaf improves insulin resistance and hyperglycemia associated acute hepatic injury and nephropathy by improving systemic antioxidant status in diabetic mice. J Ethnopharmacology. 210:275–286 IF: 5.4

2017

- Sarkar I, Tisa LS, Gtari M and Sen A (2017). Biosynthetic energy cost of potentially highly expressed proteins vary with niche in selected actinobacteria. J Basic Microbiol. DOI: 10.1002/jobm.201700350. IF: 3.1
- Dutta S, P Dey, MRSaha, I Sarkar, R Sarkar, JA Mardi, J Barman, A Sen, TK Chaudhuria (2017). Differential interaction with O₂ and N₂ freeradicals, phytochemical fingerprinting and molecular docking reveals potent antioxidant activities of three major recreational foods of the Indian subcontinent. *J Functional Foods* 39:112–122. IF: 5.6
- Pal S, I Sarkar, A Roy. PKDM Patra, KC Mondal and A Sen (2017). Comparative evolutionary genomics of *Corynebacterium* with special reference to codon and amino acid usage diversities. *Genetica* (DOI:10.1007/s10709-017-9986-6). IF: 1.5
- Goyal AK, Middha S, Usha T, Sen A (2017). Analysis of toxic, antidiabetic and antioxidant potential of Bambusa balcooa Roxb. leaf extracts in alloxan-induced diabetic rats. *3 Biotech* 7:120, DOI:10.1007/s13205-017-0776-8 IF: 2.893
- Gueddou A, Swansonb E, Ktari A, Nouioui I, Hezbri K, Ghodhbane-Gtari F, Simpson S, Morris K, Thomas W. k, Sen A, Gtaria M, Tisa L. (2017).Permanent Draft Genome Sequences of Three Frankia sp. Strains That Are Atypical, Noninfective, Ineffective Isolates. *GenomeA* vol. 5 no. 15 e00174-17
- Rai, R., Swanson, E., Sarkar, I., Lama, D., Abebe-Aleke, F., Simpson, S., ... & Sen, A. (2017). Permanent Draft Genome Sequence of the French Bean Symbiont *Rhizobium* sp. Strain RSm-3 Isolated from the Eastern Himalayan Region of India. *GenomeA*, 5(15), e00175-17.
- Ktari, A., Gueddou, A., Nouioui, I., Miotello, G., Sarkar, I., Ghodhbane-Gtari, F., ...**Sen A** & Gtari, M. (2017). Host plant compatibility shapes the proteogenome of *Frankia coriariae*. *Frontiers in Microbiology*, 8. DOI: 10.3389/fmicb.2017.00720 IF: 5.2

- <u>Ghodhbane-Gtari F, N Beauchemin, M Louati, I Nouioui, A Ktari, K Hezbri, A Gueddou, A Chen, M Huntemann, N Ivanova, N Kyrpides, V Markowitz, K Mavrommatis, I Pagani, A Sen, L Wall, T Woyke, M Gtari, and L Tisa (2016)</u> Permanent improved high quality draft genome sequence of Nocardia casuarinae strain BMG51109, an endophyte of actinorhizal root nodules of *Casuarina glauca. GenomeA* vol. 4 no. 4 e00799-16
- <u>Ghodhbane-Gtari F, N Beauchemin, A Geddou, K Hezbri, A Ktari, M Louati, I Nouioui, A Chen, M Huntemann, N Ivanova, N Kyrpides, V</u> <u>Markowitz, K Mavrommatis, I Pagani, **A Sen**, L Wall, T Woyke, M Gtari, and L Tisa (2016)</u> Permanent Draft Genome sequence of Nocardia sp. BMG111209, an actinobacterium isolated from nodules of Casuarina glauca. *GenomeA* vol. 4 no. 4 e00770-16.
- Goyal AK and A Sen (2016). In vitro regeneration of bamboos, the Green gold: An overview. Ind. J Biotechnol. 15:9-16. IF: 0.413
- Thakur S and Sen A (2016) Comparative analysis of metabolic machinery of *Frankia* along with other selected actinobacteria. *Symbiosis*. 70: 59-68. DOI 10.1007/s13199-016-0410-2. IF: 2.5

- Kar P, Dey P, Misra AK, Tapas Kumar Chaudhuri TK and Sen A (2016). Phytometabolomic fingerprinting of selected actinorhizal fruits popularly consumed in North-East India. *Symbiosis*. 70: 159-168. DOI: 10.1007/s13199-016-0415-x IF: 2.5
- Sarkar I, Normand P, Tisa S. L, Gtari M, Bothra A and Sen A (2016). Characterization of PAS domains in Frankia and selected Actinobacteria and their possible interaction with other co-domains for environmental adaptation. *Symbiosis*. 70: 69-78. IF: 2.5
- Bose D, Sarkar I, Labar R, Oshone R, Ghazal R, Morris K, Abebe-Akele F, Thomas K W, Tisa S. L, Sen A (2016). Comparative genomics of Prauserella sp. Am3, an actinobacterium isolated from root nodules of Alnus nepalensis in India. *Symbiosis*. 70:49-58. DOI 10.1007/s13199-016-0401-3. IF: 2.5
- Tisa S. L., Oshone R, Sarkar I, Ktari A, Sen A, Gtari M (2016). Genomic approaches toward understanding the actinorhizal symbiosis: an update on the status of the *Frankia* genomes. *Symbiosis*. 70: 5-16. DOI 10.1007/s13199-016-0390-2 IF: 2.5
- Saha MR, Dey P, Begum S, De B, Chaudhuri TK, Sarker DD, Das AP, Sen A (2016). Effect of Acacia catechu (L.f.) Willd. on Oxidative Stress with Possible Implications in Alleviating Selected Cognitive Disorders. PLoS One. 2016 Mar 7;11(3):e0150574. doi:10.1371/ journal.pone.0150574. eCollection 2016. IF: 3.752
- Sghaier H, Hezbri K, Ghodhbane-Gtari F, Pujic P, Sen A, Daffonchio D, Boudabous A, Tisa LS, Klenk HP, Armengaud J, Normand P, Gtari M (2016). Stone-dwelling actinobacteria *Blastococcus saxobsidens, Modestobacter marinus* and *Geodermatophilus obscurus* proteogenomes. *ISME J*. doi: 10.1038/ismej.2015.108. **IF: 11.217**
- Saha MR, P Dey, TK Chaudhuri, DD Sarker, AK Goyal and A Sen (2016). Assessment of haemolytic cytotoxic and free radical scavenging activities of an underutilized fruit, *Baccaurea sapida* (Roxb.) Muell. Arg. *Ind. J. Expl. Biol.* 54:115-125.

2015

- Rai, R., & Sen, A. (2015). Biochemical Characterization of French Bean Associated Rhizobia found in North Bengal and Sikkim. Journal of Academia and Industrial Research (JAIR), 4(1), 10-18.
- Tisa, L. S., Beauchemin, N., Cantor, M. N., Furnholm, T., Ghodhbane-Gtari, F., Goodwin, L., ... & Kyrpides, N. (2015). Draft genome sequence of Frankia sp. strain DC12, an atypical, noninfective, ineffective isolate from *Datisca cannabina*. *GenomeA*, 3(4), e00889-15.
- Labar R, Sen A (2015). Efficacy of anthocyanin in production of remedial tea. NBU Journal of Plant Sciences.9:18-32
- Banerjee A, DK Sahoo, H Thatoi, BR Pati, KC Mondal, A Sen, PK Das Mohapatra (2015). Structural Characterization and Active Site Prediction of Bacterial Keratinase through Molecular Docking. J Bioinform 1:67-82
- <u>Roy A, Mukhopadhyay S, Sarkar I, Sen A (2015)</u>. Comparative investigation of the various determinants that influence the codon and amino acid usage patterns in the genus *Bifidobacterium*. World J Microbiol Biotechnol. 31(6):959-81. doi: 10.1007/s11274-015-1850-1. IF: 3.312
- Mishra T, AK Goyal and A Sen (2015). An overview on the in vitro regeneration of Canna. *Int. J. Fund. Appl. Sc.* Int. J. Fund. Appl. Sci. 4(2) 39-49.
- Mishra T, AK Goyal, A Sen (2015) Somatic embryogenesis and genetic fidelity study of micropropagated medicinal species, *Canna indica*. *Horticulturae* (in`.
- Mishra T, AK Goyal, M Bhattacharya, P Kar and **A Sen**. (2015). Polyethylene glycol mediated protoplast fusion of medicinally important *Canna*. *Research in Plant Biology*. 5(1): 20-24. IF:2.331
- Saha MR, R Rai, P Kar, A Sen and DD Sarker (2015) Ethnobotany, traditional knowledge and socioeconomic importance of native drink among the Oraon tribe of Malda district in India. J Intercult Ethnopharmacol. 4: 34–39 DOI: 10.5455/jice.20141202060743.
- Goyal AK, S Pradhan, BC Basistha and A Sen (2015). Micropropagation and assessment of genetic fidelity of Dendrocalamus strictus (Roxb.) Nees using RAPD and ISSR markers. *3Biotech*. 5:473-482. DOI: 10.1007/s13205-014-0244-7 IF: 2.893
- Goyal AK and A Sen (2015). Phylogenetic relationships among 29 accessions of bamboos encountered in North Bengal, India based on RAPD and ISSR markers. Ind J Biotechnol. 14:495-503. IF: 0.324

2014

Dey P, MR Saha, SR Chowdhuri, A Sen, MP Sarkar and TK Chaudhuri (2014). Assessment of anti-diabetic activity of an ethnopharmacological

plant Nerium indicum through alloxan induced diabetes in mice. J Ethnopharma. 161:128-37. doi: 10.1016/j.jep.2014.12.012. IF: 5.4

Sen A, V Daubin, D Abrouk, I Gifford, AM Berry, and P Normand (2014) The phylogeny of actinobacteria revisited in the light of complete genomes, the orders Frankiales and Micrococcales should be split into coherent entities. Proposal of Frankiales ord. nov., Geodermatophilales ord. nov., Acidothermales ord. nov. and Nakamurellales ord. nov. Int J Syst Evol Microbiol. 64:3821-32. doi: 10.1099/ijs.0.063966-0. IF: 2.166

Mishra T and A Sen (2014). Antioxidant Potential of Canna: an overview. NBU J Pl Sc.

- Saha MR, P Kar, A Sen and DD Sarker (2014). Ethnobotany of Chanchal Block of Malda District of West Bengal (India): Plants used Local Healthcare. *Pleione*
- Saha MR, DD Sarker, P Kar, P Sengupta and A Sen (2014). Indigenous knowledge of plants in local healthcare management practices by tribal people of Malda district, India. J Intercult Ethnopharmacol. 3(4): 179-185.
- Ghosh S, UK Mondal, **A Sen** and AK Bothra (2014). Comparative analysis and codon usage study of nifD, nifK and nifH genes linked with free living nitrogen fixing bacteria. *Int. J. Integ. Biol.* 15:7-10
- <u>Kar, P, AK Goyal AP Das and A Sen (2014)</u>. Antioxidant and pharmaceutical potential of Clerodendrum L.: an overview. *Int. J. Green Pharma*.
 8: 210-216.
- Pal S, A Banerjee, PS Das, BR Pati, KC Mandal, B Chakraborty, A Sen and PKD Mohapatra (2014) Dengue virus type 4 evolution & genomics: a Bioinformatic approach. Ind J Biotechnol 13:330-336. IF: 0.324
- Banerjee A, S Pal, T Pal, KC Mondal, BR Pati, A Sen and PKD Mohapatra (2014) Characterization of Bacillus proteases through proteinprotein interaction: an in silico study of anthrax pathogenicity. *Hum. Trad. Med.* 4:e6.
- Goyal AK and A Sen (2014). Molecular Phylogeny of North Bengal bamboos inferred through PCR-RFLP based on the trnL-trnF region. *The Experiment Journal*. 19:1304-1315.
- Saha MR, DD Sarker and A Sen (2014). Ethnoveterinary practices among the tribal community of Malda district of West Bengal, India. Ind. J. Tradit. Knowl. 13: 359-367. IF: 1.091
- Pal A, UK Mandal, S Mukherjee, **A Sen** and AK Bothra (2014). The Implication of Codon Usage Design and Expression Level in Determining the Nature of Selection and Functionality amongst the Amino Acid Biosynthetic Pathway coding sequences of Arthrobacter sp. FB24. *Current Bioinformatics* 9:470-480. IF: 4.850

- Dey P, Saha MR and Sen A (2013). An overview of drug-induced hepatotoxicity. Asian J Pharma. Clinic. Res. 6(4): 1-4.
- Goyal AK, Middha SK and Sen A (2013). Bambusa vulgaris 'vittata' attenuates oxidative stress- an *in vitro* biochemical assay. *IJNPR* 4: 436-440.
- Dey P, Saha MR and Sen A (2013). Hepatotoxicity and the present herbal hepatoprotective scenario. Int. J. Green Pharm. 7: 265-273.
- Tisa LS, N Beauchemin, M Gtari, A Sen, and LG Wall (2013). What stories can the Frankia genomes tell us? J Biosciences 38:719–726. IF: 1.645
- Bhattacharya S, A Sen, S Thakur and LS Tisa (2013). Characterization of Hemoglobin from Actinorhizal Plants An in-silico approach. J Biosciences 38:777–787. IF: 2.795
- Goyal AK, T Mishra, M Bhattacharya, P Kar and A Sen (2013). Evaluation of phytochemical constituents and antioxidant activity of selected actinorhizal fruits growing in the sacred forests of North-East India. J Biosciences 38:797–803. IF: 2.795
- Thakur S, AK Bothra and A Sen (2013) Functional divergence outlines the evolution of novel protein function in NifH/BchL protein family. J Biosciences 38:733–740. IF: 2.795
- Sur S, S Saha, LS. Tisa, AK Bothra and A Sen (2013) Characterization of pseudogenes in members of the order Frankineae. J Biosciences 38:727–732. IF: 2.795
- Thakur S, Normand P, Daubin V, Tisa LS and Sen A (2013). Contrasted evolutionary constraints on secreted and non-secreted proteomes of selected Actinobacteria. BMC Genomics doi:10.1186/1471-2164-14-474. IF: 4.4
- Wall L, Sen A, ... Tisa LS. 2013. Draft Genome sequence of Frankia sp. strain BCU110501, a nitrogen-fixing actinobacterium isolated from

nodules of Discaria trinevis. GenomeA. 1: doi: 10.1128/genomeA.00503-13.

- Nouioui I, ... Sen A, ... Tisa LS. 2013. Draft Genome sequence of *Frankia* sp. strain BMG5.12, a nitrogen-fixing actinobacterium isolated from Tunisian soils." GenomeA. 1: doi: 10.1128/genomeA.00468-13.[23846272]
- Ray A, Bhattacharya S, Bothra AK and Sen A. (2013) A Database for Mycobacterium secretome analysis: 'MycoSec' to accelerate global health research. Omics: A Journal of Integrative Biology 17:502-509. IF: 3.374
- Bhattacharya M and Sen A (2013) In vitro regeneration of pathogen free Kaempferia galanga L. a rare medicinal plant. Res. Plant Biol. 3(3): 24-30, 2013
- Thakur S, AK Bothra and **A Sen** (2013). Exploring the genomes of symbiotic diazotrophs with relevance to biological nitrogen fixation. In R. Bandopadhyay and PB Kavi (ed). *Agriculture bioinformatics*. *Springer* publications Pp 235-257.
- <u>Sen A,Tisa LS. 2013.</u> Draft genome sequence of *Frankia* sp. strain QA3, a nitrogen-fixing actinobacterium isolated from the root nodule of *Alnus nitida*. Genome Announc. 1(2):e00103-13. doi:10.1128/genomeA.00103-13.
- Thakur S, AK Bothra and **A Sen** (2013). Insights into the nitrogenase protein an *in silico* approach. *In*: Biology of useful plants and microbes, A Sen (ed), Narosa Publishing House, New Delhi, India, pp 301-320.
- Goyal AK, P Kar and **A Sen** (2013). Advancement of bamboo taxonomy in the era of molecular biology: a review. *In*: Biology of useful plants and microbes, A Sen (ed), Narosa Publishing House, New Delhi, India, pp 197-208.
- <u>Ghodhbane-Gtari F, ... Sen A, ... Tisa LS. (2013).</u> Draft genome sequence of *Frankia* sp. strain CN3, an atypical, noninfective (Nod–) ineffective (Fix–) isolate from Coriaria nepalensis. GenomeA. 1(2):e00085-13. doi:10.1128/genomeA.00085-13.
- Goyal AK, T Misra and A Sen (2013) Antioxidant profiling of Latkan (Baccaurea ramiflora Lour.) wine. Ind. J Biotechnol. 12: 137-139. IF: 0.324
- Sur S, AK Bothra and A Sen (2013) Proteome analysis reveals the influence of isoelectric point and amino acid usages on the lifestyle of nitrogen fixing microorganisms. Ind J Biotechnol. 12: 88-97. IF: 0.324

2012

- Bose D and **A Sen** (2012). Resent advances in *Frankia* research. In Biology of plants and microbes, D Bose & S Roy (eds) Lavent Books, Kolkata. Pp 42-47.
- Thakur S, AK Bothra, S Sur and **A Sen** (2012) Molecular dynamics simulation of pr-1 protein from *Solanum tuberosum* provides an in-depth view of its structural features. In 'Microbial Resources for Crop Improvement'; BN Chakraborty and U Chakraborty (ed.); Satish Serial Publishing House Pp 173-181.
- Kundu, S, AK Bothra, LS Tisa and A Sen (2012). In silico analysis reveals the role of horizontally transferred genes (HGTs) in shaping the pathogenicity of Xanthomonas. Ind. J. Biotechnol. 11:404-411. IF: 0.324
- Goyal AK, PK Ghosh, AK Dubey and A Sen (2012). Inventorying bamboo biodiversity of North Bengal: A Case Study. Int. J. Fundam Appl Sci. 2012, http://bma.org.in/jfaas.apx 1(1), 5-8
- Saha, S, S Sur, A Bothra and A Sen (2012). A homology model for 16S rRNA tertiary structure of Frankia. NBU J Pl. Sc.6:89-94.

Mondal, U, A Sen and A Bothra (2012). Bioinformatics of pathogenic food bacteria. NBU J Pl. Sc.6:9-17.

- Thakur S, A Bothra and A Sen (2012) In silico studies of NifH protein structure and its post-translational modification in Bradyrhizobium sp. ors278. Int. J Pharm Bio Sc.
- Mishra T, Das AP and Sen A (2012). Phytochemical screening and in vitro antioxidant profiling of solvent fractions of Canna edulis Ker Gawler. Free Rad. and Antiox. 2:13-20.
- Sen A, S Thakur, A Bothra, S Sur and L Tisa (2012). Identification of TTA codon containing genes in *Frankia* and exploration of the role of tRNA in regulating these genes. Archives of Microbiology.194: 35-45. IF: 2.8

2011

Chakraborty D, UK Mondal, **A Sen** and AK Bothra (2011). Molecular dynamics simulation of hydrogenase isoenzyme formation protein HypC. IUP J Biotechnol. 5(3):56-64.

Sur S, AK Bothra and **A Sen** (2011). Bioinformatics of Nitrogen fixation – a review, In Recent studies in biodiversity and traditional knowledge in India. C Ghosh & AP Das (*eds.*) pp35-43.

Mishra T, AK Goyal, SK Middha and A Sen (2011) Antioxidative Properties of Canna edulis Ker Gawler. Ind. J Nat. Pro. Res. 2:315-321.

- Sen A and AK Misra (2011) Frankia and actinorhizal symbiosis. In: *Microorganisms in Sustainable Agriculture and Biotechnology* (ed. Satyanarayana T. Johri BN and Prakash A). Springer Science+Business Media B.V. DOI 10.1007/978-94-007-2214-9_7 pp113-126.
- <u>Goyal AK, BC Basistha</u>, **A Sen** and SK Middha (2011). Antioxidant profiling of *Hippophae salicifolia* D. Don growing in sacred forests of Sikkim, India. Functional Plant Biol. 38(9) 697-701. IF: 3.0
- Goyal AK, SK Middha and A Sen (2011). In vitro antioxidative profiling of different fractions of *Dendrocalamus strictus* (Roxb.) Nees Leaf Extract. Free Radicals and Antioxidants. 1:42-48
- Mondal UK, A Pal, A Sen and AK Bothra (2011). Bioinformatic study of pathogenicity related genes of three species of Helicobacter. Int. J Appl. Biotech. Biochem. 1(2): 193-200.
- Thakur S, Z Patra, M Biswal and **A Sen** (2011). Exploration of proteis with signal peptides and their respective genes in some actinomycetes: a bioinformatics approach. Int. J. Integrative Biol. 11(2): 97-102.
- Basistha BC and A Sen (2011). Sea buckthorn and its microsymbiont-a review. NBU J Pl. Sc.5:67-84.
- Mishra T, AK Goyal, P Mondal and A Sen (2011). Free radical scavenging activity of different ornamental and cultivars of Canna available in Eastern India. NBU J Pl. Sc.5:41-45.
- Mondal UK, A Sen and AK Bothra (2011). Characterization of pathogenic genes through condensed matrix method, case study through bacterial Zeta toxin. International journal of Genetic Engineering and Biotechnology. 2(1): 109-114.

- Bose D, B Bajwa, M Bajwa and **A Sen** (2010) Physiological characterization of *Frankia* strains isolated from sub-Himalayan West Bengal, India reveals two distinct groups. Biotech 3:6-15.
- <u>Goyal AK, SK Middha and A Sen (2010</u>). Evaluation of DPPH Radical Scavenging Activity, Total Phenols And Antioxidant Activities In Indian Wild Bambusa vulgaris 'Vittata' Methanolic Leaf Extract. Journal of Natural Pharmaceuticals 1:40-45.
- Goyal AK, SK Middha, U Talambedu, S Chatterjee, AK Bothra, MB Nagaveni and A Sen (2010). Bambooinfoline- a database for North Bengal Bamboo's. Bioinformation. 5(4):184-185.
- Goyal AK, A Sen, S Sur and AK Bothra (2010). Exploration of codon usage patterns in some Brucella genomes. IJPBS 1(4):B-239-252.
- Thakur S, AK Bothra, S Sur and A Sen (2010). Modeling and molecular dynamics simulation of PR-1 protein an integral part of plant defense. Int J of Biol and Chem Sci 4(4): 1251-1261.
- <u>Goyal, AK, K Ganguly, T Mishra and A Sen (2010)</u> In vitro multiplication of Curcuma longa Linn.–an important medicinal zingiber. NBU J Pl. Sc 4: 21-24.
- Ghosh MK, A Bothra and **A Sen** (2010) Bioinformatics of codon usage pattern in pathogenic proteobacteria *Burkholderia mallei*. NBU J Pl. Sc 4: 65-71.
- Sur S, A Bothra, TC Ghosh and A Sen (2010) Investigation of the molecular evolution of nitrogen fixation using nucleotide triplet based condensed matrix method. Int. J. Integrative Biol.10: 59-65
- Mondal UK, A Sen and A Bothra (2010) Homology modelling and molecular dynamics of CdtB of *Helicobacter hepaticus* ATCC51449. Int. J. Integrative Biol. 10: 35-40
- Sur S, A Bothra and A Sen (2010) Symbiotic nitrogen fixation-a bioinformatics perspective. Biotechnology DOI: 0000.19779-19779
- Basistha BC, NP Sharma, L Lepcha, ML Arrawatia and A Sen (2010). Ecology of Hippophae salicifolia d. Don of temperate and sub-alpine forests of north Sikkim Himalayas a case study. Symbiosis 50:87-95. IF: 2.5
- Sen A, S Sur, LS Tisa, AK Bothra, S Thakur and UK Mondal (2010) Homology modelling of the *Frankia* nitrogenase iron proteins. *Symbiosis* 50:37-44. IF: 2.5

- Mondal U, A Sen, and A Bothra (2009) Comparison in motional properties of *Staphylococcus aureus* exfoliative toxins A and B as revealed by their MD simulation. Int. J. Integrative Biol. 8:19-24.
- Sur S, G Sen, S Thakur, AK Bothra and A Sen (2009) In silico analysis of evolution in swine flu viral genomes through re-assortment by promulgation and mutation. *Biotechnol*. 8:434-441.
- Mondal U, **A Sen**, and A Bothra (2009) Molecular dynamics simulation receptor-binding C-terminal domain from *Clostridium difficile* Toxin A. *ICFAI J Biotech*. III: 22-37.
- Bhattacharya M, P Mondal and A Sen (2009). In vitro detection of antioxidants in different solvent fractions of Ginger (Zingiber officinale Rosc.). Ind J Plant Physiol. 14:23-27.
- Sur S, B Bajwa, M Bajwa, B Basistha, AK Bothra and A Sen (2009) Investigation of codon and amino-acid usages in a *Rhizobium* phage. *NBU J Pl. Sc.* 3: 49-51.
- Thakur S, S Sur, D Bose, AK Goyal, T Mishra, R Rai, M Bhattacharya, AK Bothra and A Sen (2009) Molecular modeling of a pathogenesis related protein from *Solanum tuberosum*. *NBU J Pl. Sc.* 3: 15-19.

2008

- Mondal UK, Das B, Ghosh TC, Sen A, and Bothra AK (2008). Nucleotide Triplet Based Molecular Phylogeny of class-I and classII Aminoacyl t-RNA. J. Biomol. Str. Dyna (JBSD) 26:321-328. IF: 4.4
- Mondal UK, S Sur, AK Bothra and A Sen (2008) Comparative analysis of codon usage patterns and identification of predicted highly expressed genes in five Salmonella genomes. Indian Journal of Medical Microbiology (26) 4: 313-321. IF: 1.347
- Sur S, AK Bothra, M Bajwa, LS Tisa & A Sen (2008). In Silico analysis of Chlorobium genomes divulge insights into the subsistence of the bacteria. Res. J. Microbiol. 3: 600-613.
- Sur S, M Bhattacharya, AK Bothra, LS Tisa and A Sen (2008). Bioinformatic analysis of codon usage patterns in a free living diazotroph, Azotobacter vinelandii. Biotechnology 7: 242-249.
- Sen A, S Sur, AK Bothra, DR Benson, P Normand, and LS Tisa. (2008) The implication of life style on codon usage patterns and predicted highly expressed genes for three *Frankia* Genomes. Antonie van Leeuwenhoek 93(4) 335-346. IF: 2.674

Chaudhury J and Sen A (2008). Anthrax and Bioterrorism. NBU J Plant Sc. 2:13-29

- Sur S, A Sen & AK Bothra (2007). Codon usage analysis of some archaeal (methanogenic) nitrogen fixing genes: relationship to gene expression and bias. *Bioinfo. Trends* 2: 47-60.
- Sen A, A Pal and D Bose (2007) Economic uses of Seabuckthorn (*Hippophae* L.) In Advances in Ethnobotany eds. AP Das and AK Pandey, BSMPS Dehra Dun, India, pp309-315.
- Bose D, S Sur, A Bothra and A Sen (2007) Study of the diversity of heavy metal resistance genes and their codon usage profiling. *ICFAI J Biotech* 2:49-59.
- Ghosh MK, S Sur and **A Sen** (2007) Nitrogen fixation in various microorganisms with special reference to *Frankia*. In Rhizosphere Biotechnology: Plant growth-Retrospect and prospect, ed. AK Roy, Scientific Publishers. pp 167-182.
- Bose D and A Sen (2007) Heavy metal induced pigment production by Frankia isolates from Mirik of Darjeeling hills. J. Hill Res. 20:36-38.
- Bajwa B, M Bajwa, S Sur, A Bothra, **A Sen** (2007) Molecular Systematics of *Frankia*-Actinorhizal Symbiosis. *In* Microbial Biotechnology, ed. R Saikia, New India Publishing Agency.
- Mondal U, S Sur, A Sen, and A Bothra (2007) Codon volatility: an efficient tool to detect the purity of genes. ICFAI J Biotech. 1: 45-49.
- Sur S, A Sen and AK Bothra. (2007) Mutational drift prevails over translational efficiency in *Frankia nif* operons. *Ind J Biotechnol* 6:321-328. IF: 0.324

2006

Bhattacharya M and A Sen (2006). Rapid in vitro multiplication of disease-free Zingiber officinale Rosc. Ind. J. Plant. Physiol. 11:379-384.

- Sur S, A Sen and AK Bothra (2006) Codon usage profiling and analysis of intergenic association of *Frankia* EulK1 *nif* genes. *Ind J Microbiol* 46:363-369. IF: 2.461
- Bose D and A Sen (2006) Isolation and heavy metal resistance pattern of *Frankia* from *Casuarina equisetifolia* nodules. *Ind J Microbiol* 46:9-12. IF: 2.461

2005

- Bose D and **A Sen** (2005) Heavy metal resistance pattern of *Frankia* isolated from sub-Himalayan West Bengal. In: Stress Biology (eds. U Chakraborty and BN Chakraborty) Narosa Publishing House, New Delhi. Pp 154-157.
- Sur S, A Pal, AK Bothra and **A Sen**. (2005) Moderate codon bias is attributed to translational selection for nitrogen fixing genes of *Bradyrhizobium japonicum* USDA 110. *Bioinformatics India*. 3:59-64.
- Bajwa M, A Sen and B Bajwa (2005) High level of polymorphism found in alder based *Frankia* of Darjeeling as revealed by PCR-RFLP. *Ind J Expl Biol*. 43:813-816. IF: 0.994

2002

Sen A and B Bajwa (2002) Taxonomy and evolution of actinorhizal plants and their microsymbionts. In Perspective of Plant Biodiversity. Ed. AP Das 417-436.

2001

Grover. A, Yashpal, A Sen, N Arora, P.B. Kirti and R.P. Sharma (2001) Removal of vacuolar targeting signal from classI vacular chitinase leads to its extracelluar secretation in transgenic tobacco. *J. Plant Biochem and Biotechnol*. 10:139-142. IF: 1.525

1998

- Sarma G., A. Sen, R. Varghese and A. K. Misra (1998). A novel technique for isolation of *Frankia* and generation of single spore cultures. *Canadian J. Microbiol*. 44: 490-492. IF: 3.226
- Yashpal, A Sen, J Venkateswari, P.B. Kirti, R.P. Sharma and A Grover (1998) Pyramiding of chitinase and glucanase genes for fungal resistance. *Ind. J. Expl. Biol.* 37:579-583. IF: 0.944
- Varshney R.K., P.C. Sharma, P.K. Gupta, H.S. Balyan, B. Ramesh, J.K. Roy, A. Kumar, and A. Sen (1998). Low level of polymorphism detected by SSR probes in bread wheat. *Plant Breeding* 117:182-184. IF: 2.536

1997

- Borthakur M., A. Sen and A. K. Misra (1997). Exogenous nitrogen sources do not drastically reduce nitrogenase activity in polymer entrapped *Frankia*, *Indian J. Exp. Biol.*, 35: 173-175. IF: 0.944
- Sen A., H.S. Balyan, P.C. Sharma, B. Ramesh, A. Kumar, J.K. Roy, R.K. Varshney and P.K. Gupta (1997). DNA amplification fingerprinting (DAF) as a new source of Molecular markers in bread wheat. *Wheat Information Service* 85:35-42.

1996

Borthakur M., A. Sen and A. K. Misra (1996). Immobilized *Frankia* spores remained viable on dry storage and on restoration to medium regenerated active colonies, *Plant and Soil*, 181: 227-231. IF: 3.299