



ALAGAPPA UNIVERSITY

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DEPARTMENT OF BIOINFORMATICS

(UGC- INNO, DST-FIST-PURSE, DBT-BIC and RUSA 2.0 Sponsored Department)
KARAIKUDI-630 004, TAMIL NADU, INDIA.

2nd International Conference cum Workshop on "RECENT TRENDS IN STRUCTURAL BIOINFORMATICS AND COMPUTER AIDED DRUG DESIGN" (ICSBCADD-2022) November 21st – 25th, 2022

S.No.	Title of the Abstract	Talk / Posters
1.	Mapping the protein binding site of the (pro)renin receptor using <i>in silico</i> 3D structural analysis Akio Ebihara Faculty of Applied Biological Sciences, Gifu University, Tokai National Higher Education and Research System, 1-1 Yanagido, Gifu 501-1193, Japan	Plenary Lecture
2.	<i>In-silico</i> approaches to understand antibiotic resistance in clinically important bacterial pathogens Sravan Kumar Miryala, Sudha Ramaiah and Anand Anbarasu* Medical & Biological computing laboratory, School of Biosciences & Technology, Vellore Institute of Technology, Vellore-632014, India	Plenary Lecture
3.	The current status of computational screening of chemical libraries for druggability prediction : Accuracy and Speed N. Arul Murugan Department of Computational Biology, Indraprastha Institute of Information Technology, New Delhi 110020	Plenary Lecture
4.	Proteomics studies for biomarker discovery in head and neck squamous cell carcinoma treated with Radiotherapy Lipi Das ^{a,b} , Ashok K Varma ^{a,b,*} a. Advanced Centre of Treatment, Research and Education in Cancer, Kharghar, Navi Mumbai. b. Homi Bhabha National Institute, Anushaktinagar, Mumbai - 400 094	Plenary Lecture
5.	Structural biology using synchrotron radiation and cryo-electron microscopy Atsushi Nakagawa Institute for Protein Research, Osaka University 3-2 Yamadaoka, Suita, Osaka 565-0861 Japan	Plenary Lecture
6.	Genomic biomarkers for early detection of complex diseases Basant K. Tiwary Department of Bioinformatics, School of Life Sciences, Pondicherry University,	Plenary Lecture

	Pondicherry-605014	
7.	Translating bioinformatics for patient care in hospitals Chandra Verma Atomistic Simulations and Design in Biology, Biomolecular modelling and Design Function, Bioinformatics Institute, A* STAR, Singapore-138671	Plenary Lecture
8.	Structural Biology at NSRRC and recent applications Chun-Jung Chen* Life Science Group, Scientific Research Division, National Synchrotron Radiation Research Center, Hsinchu 30076, Taiwan	Plenary Lecture
9.	Modern Machine Learning Methods for Drug Design U. Deva Priyakumar Center for Computational Natural Sciences and Bioinformatics, International Institute of Information Technology, Hyderabad 500 032	Plenary Lecture
10.	Structure-based virtual screening for identification of novel greatwall kinase inhibitors A. Dinakara Rao, Anbumani Velmurugan Ilavarasi, Tulsi, Saswati Sarita Mohanty and Katike Umamahesh. Department of Bioinformatics, School of Life Sciences, Pondicherry University, Puducherry	Plenary Lecture
11.	QSAR Modeling approach in Virtual Screening of Active Compounds Feroz Khan Computational Biology Unit, CSIR-Central Institute of Medicinal & Aromatic Plants, P.O.-CIMAP, Lucknow-226015 (U.P.), INDIA	Plenary Lecture
12.	The Success story of PAK1-RUNX3 Ganesh Venkatraman *and Suresh K Rayala Department of Human Genetics, Sri Ramachandra Institute of Higher Education and Research, Chennai-600 116	Plenary Lecture
13.	Plant Diseases in Guyana -Past, Present and Control Measures Gomathinayagam Subramanian and Rekha Gomathinayagam University of Guyana Berbice Campus, Corentyne, Berbice, Guyana, South America	Plenary Lecture
14.	Targeting pyruvate dehydrogenase kinase signalling in the development of effective cancer therapy: A combined computational and experimental approach Md. Imtaiyaz Hassan Centre for Interdisciplinary Research in Basic Sciences, Jamia Millia Islamia, Jamia Nagar, New Delhi 10025, INDIA	Plenary Lecture
15.	Dhanvantari-2022: Genome to Drug in silico - A country path today; A highway tomorrow B. Jayaram Department of Chemistry, Kusuma School of Biological Sciences & Supercomputing Facility for Bioinformatics & Computational Biology, Indian Institute of Technology, Hauz Khas, New Delhi-110016, India	Plenary Lecture
16.	Biomedical Text Mining and Applications in Drug Discovery	Plenary Lecture

	Dr. N. Jeyakumar, Dept. of Bioinformatics, School of Life Sciences, Bharathiar University, Coimbatore 641 046	
17.	Listeria Monocytogenes High Temperature Requirement A (HtrA) Protease and Its Interaction with Extracellular Matrix Molecules Karthe Ponnuraj Centre of Advanced Study in Crystallography and Biophysics, University of Madras, Guindy Campus, Chennai 600025, India	Plenary Lecture
18.	Designing and development of hybrid conjugates of commercial drugs and bioactive herbals a possible approach for novel efficacious nonresistant therapeutic agents Krishna Misra Honorary Professor, Indian Institute of Information Technology, Allahabad	Plenary Lecture
19.	Computational Explorations to Overcome the Anti-Microbial Resistance in Bacteria Lakshmi P.T.V Department of Bioinformatics, School of Life Sciences, Pondicherry University, Puducherry-605014	Plenary Lecture
20.	Structural intermediates in the assembly landscape of the human mitochondrial anion channel Radhakrishnan Mahalakshmi* Molecular Biophysics Laboratory, Department of Biological Sciences, Indian Institute of Science Education and Research, Bhopal, India.	Plenary Lecture
21.	Binding affinity of protein-carbohydrate complexes: database development, analysis and prediction M. Michael Gromiha Department of Biotechnology, Indian Institute of Technology Madras Chennai 600036, INDIA	Plenary Lecture
22.	Dissecting Hubs and Bottlenecks in a Protein-Protein Interaction Network Prof H. A. Nagarajaram Department of Systems and Computational Biology, School of Life Sciences University of Hyderabad	Plenary Lecture
23.	Exopolysaccharides and their functions in disease: The story in an oral bacterium N. Ramasubbu, C. Cugini and E. Shimizu Oral Biology, Rutgers School of Dental Medicine, Newark, NJ 07101	Plenary Lecture
24.	Structural characterization and engineering of a plasticizer-degrading enzyme Pravindra Kumar* and Jai Krishna Mahto Department of Biosciences and Bioengineering, IIT Roorkee, Roorkee, India.	Plenary Lecture
25.	Design, identification and evaluation of anti-cancer agents against Aurora kinases	Plenary Lecture

	<p>Punit Kaur¹, Deepali Gupta¹, Mukesh Kumar¹, Mandeep Singh¹, A.S. Ethayathulla¹, Andrey E. Shchekotikhin² and Uddipan Das¹</p> <p>¹Department of Biophysics, All India Institute of Medical Sciences, New Delhi, Delhi 110029, India and ²Gause Institute of New Antibiotics, Moscow, 11 B. Pirogovskaya Street, Moscow 119021, Russia</p>	
26.	<p>Computer aided drug design for developing potential inhibitors of oncogene PELP1 for treating breast cancer Ratna K Vadlamudi</p> <p>Professor, UT Health San Antonio, USA</p>	Plenary Lecture
27.	<p>Role of sulfur-mediated chalcogen bonds in the structure and substrate specificity of proteins Saikrishnan Kayarat*, Vishal Adhav</p> <p>Department of Biology, Indian Institute of Science Education and Research, Pune – 411008</p>	Plenary Lecture
28.	<p>SciWhyLab tools for integrating machine learning, transcriptomics and lead generation Anuja Jain, Shruti Gupta, Ajay Kumar Verma, Ajay Arya, Manisha Kalsan, Dana Mary Varghese and Shandar Ahmad*</p> <p>School of Computational and Integrative Sciences, Jawaharlal Nehru University, New Delhi</p>	Plenary Lecture
29.	<p>Inhibition Mechanism of Human Kidney Type Glutaminase to Arrest Cancer Cell Proliferation J. Sivaraman</p> <p>Department of Biological Sciences, National University of Singapore, Singapore 117546</p>	Plenary Lecture
30.	<p>PELP1 – a mediator for inflammation induced tumorigenesis Suresh K Rayala*and Ganesh Venkatraman [R1]</p> <p>Department of Biotechnology, IIT Madras, Chennai-600 032</p>	Plenary Lecture
31.	<p>Exploring the medicinally important secondary metabolites in fenugreek through transcriptome studies R. Sowdhamini</p> <p>National Centre for Biological Sciences, Tata Institute of Fundamental Research, Bellary Road, Bangalore-560 065.</p>	Plenary Lecture
32.	<p>Structure based understanding of the innate immunity proteins and their exploitation for therapeutic applications T.P. Singh</p> <p>Department of Biophysics, All India Institute of Medical Sciences, New Delhi</p>	Plenary Lecture
33.	<p>Potential Drug Resistance in SARS-CoV-2: Importance of Mutational Surveillance in Pandemics Timir Tripathi^{1,2}</p> <p>¹Regional Director's Office, Indira Gandhi National Open University (IGNOU), Regional Centre Kohima, Kohima- 797001, Nagaland, India</p>	Plenary Lecture

	² Molecular and Structural Biophysics Laboratory, Department of Biochemistry, North-Eastern Hill University, Shillong- 793022, Meghalaya, India.		
34.	Molecular docking and Molecular dynamics simulations on various Marine Bioactive compounds against Neuro Degenerative Disorders Devadasan Velmurugan* and Anantha Krishnan Dhanabalan Office of the Director, Research and Development Cell, AMET University, Kanathur 603112, Chennai.	Plenary Lecture	
35.	Investigating enhancer reprogramming and 3D organisation in p53 deficient/mutant colorectal cancer cells Vijayalakshmi Mahadevan Institute of Bioinformatics and Applied Biotechnology (IBAB), Bangalore, India	Plenary Lecture	
36.	Gene expression and interaction biology reveals toll-like receptor signaling pathway as a major trigger in COPD pathogenesis Pavithra Dhamodharan ^a , Mohanapriya Arumugam ^{a*} ^a Department of Biotechnology, School of Biosciences and Technology, Vellore Institute of Technology	Plenary Lecture	
37.	Design of Peptidomimetic Ligands for the Inhibition of BACE1 of Alzheimer's Disease: Computational and Chemical Biology Approach K. Muruga Poopathi Raja ^{1,2*} 1. Biophysical and Chemical Biology Laboratory, Department of Physical Chemistry, School of Chemistry, Madurai Kamaraj University, Palkalai Nagar, Madurai-625021. 2. Biophysical and Chemical Biology Laboratory, Department of Chemistry, School of Physical Sciences, Central University of Kerala, Periya, Kasaragod District, Kerala - 671 320.	Plenary Lecture	
38.	Charge-Charge Repulsion: An instance from <i>Candida albicans</i> Pyruvate kinase Lukkani Laxman Kumar, Ayaluru Murali* Department of Bioinformatics, Pondicherry University, Pondicherry - 605014	Poster	Accepted
39.	An <i>in silico</i> Approach to Investigate the Interaction of Fe³⁺ ion with Ftr1p of Mucormycosis Causing <i>Rhizopus sp.</i> Ahana Roy Choudhury and Ayaluru Murali* Department of Bioinformatics, School of Life Sciences, Pondicherry University, Puducherry- 605014	Poster	Accepted
40.	<i>In silico</i> Analysis of Red Seaweed <i>Gelideilla acerosa</i> Kirithiga. A ¹ , Anantha Krishnan. D ² , Velmurugan. D ² and Ramakritinan. C.M ^{1*} ¹ Department of Marine Biotechnology, AMET Deemed to be University, Kanathur - 603112, Chennai, Tamil Nadu, India ² Office of the Director, Research and Development Cell, AMET Deemed to be University, Kanathur - 603112, Chennai, Tamil Nadu, India	Poster	Accepted
41.	A Machine Learning Application for Bioactivity Prediction of the Target Replicase Polyprotein 1ab of SARS-CoV-2 Rajesh Das, Amouda Venkatesan*	Poster	Accepted

	Department of Bioinformatics, Pondicherry University, Kalapet, Puducherry-605014		
42.	<p>Screening of key genes involving in hormone signalling in a phytopathosystem (<i>Piper nigrum</i> - <i>Phytophthora capsici</i>) using transcript assembly</p> <p>B.Nagarathnam¹,Mohamed Shafi¹,Chidambareswaren Mahadevan², Manjula Sakuntala², R.Sowdhamini¹</p> <p>¹ National Centre for Biological Sciences, (TIFR),GKVK Campus,Bangalore-560065. India ²Plant Disease Biology and Biotechnology, Rajiv Gandhi Centre for Biotechnology, Thycaud P.O, Thiruvananthapuram – 695014, Kerala.</p>	Poster	Accepted
43.	<p>Big data analysis of <i>Rauwolfia serpentine</i> reveals the potency of indole alkaloids using <i>De-novo</i> Transcriptome studies</p> <p>Hithesh Kumar¹, Anand Anbarasu^{1,2} and Sudha Ramaiah^{3*}</p> <p>¹School of Bio-Sciences and Technology (SBST), Vellore Institute of Technology, Vellore, 632014 ²Department of Biotechnology, SBST, Vellore Institute of Technology, Vellore ³Department of Bio-sciences, SBST, Vellore Institute of Technology, Vellore</p>	Poster	Accepted
44.	<p><i>In-silico</i> investigation of curcumin-indole-3-propionic acid diester Against the enzymes involved in Diabetes-associated cognitive decline</p> <p>Sidhambaram Jayanthi, Loganathan Chitra, Sakayanathan Penislusshiyam and Thayumanavan Palvannan*</p> <p>Department of Biochemistry, Periyar University, Salem-636011*</p>	Poster	Accepted
45.	<p>Proteoglycan signaling FN1 as a potential therapeutic target for pancreatic adenocarcinoma: A systems biology approach</p> <p>Gayathri Ashok, Anand Anbarasu and Sudha Ramaiah*</p> <p>Medical and Biological Computing Laboratory, Department of Biosciences, School of Bio Sciences and Technology, Vellore Institute of Technology, Vellore, 632014</p>	Poster	Accepted
46.	<p>Novel antimicrobial peptide mutants can potentially inhibit β-lactamase enzymes in Gram-negative clinically important bacterial pathogens</p> <p>Soumya Basu, Sudha Ramaiah, Anand Anbarasu*</p> <p>Medical and Biological Computing Laboratory, School of Bio-Sciences and Technology (SBST), Vellore Institute of Technology, Vellore, India</p>	Poster	Accepted
47.	<p>Novel curcumin derivative can inhibit MurC ligase to obstruct peptidoglycan biosynthesis in multi-drug resistant <i>Salmonella Typhi</i>: Evidence from Molecular Docking and Dynamics Simulations</p> <p>Reetika Debroy, Anand Anbarasu and Sudha Ramaiah*</p> <p>Medical and Biological Computing Laboratory, School of Bio-Sciences and Technology, Vellore Institute of Technology (VIT), Vellore-632014, Tamil Nadu, India</p>	Poster	Accepted
48.	<p>A Novel Ursodeoxycholic Acid Derivative can be a potential inhibitor for therapeutic target Parkin for Parkinson's Disease: Evidence from gene interaction network analysis, structure evaluation and molecular dynamics</p>	Poster	Accepted

	P. Priyamvada, Suvitha Anbarasu, Anand Anbarasu, Sudha Ramaiah* Medical and Biological Computing Laboratory, School of Bio-Sciences and Technology, Vellore Institute of Technology, Vellore-632014, Tamil Nadu.		
49.	Two-component EvgS protein subdues both antimicrobial resistance and virulence in <i>Shigella flexneri</i> 2a str. 301: Evidence from Gene Interaction Network, Differential Gene Expression and Molecular Dynamics Simulation Aniket Naha, Anand Anbarasu, Sudha Ramaiah* Medical and Biological Computing Laboratory, School of Bio-Sciences and Technology, Vellore Institute of Technology, Vellore-632014, Tamil Nadu, India	Poster	Accepted
50.	Computational approach for the recognition of small molecule inhibitors for Toll-like receptor Shailya Verma and Ramanathan Sowdhamini* National Centre for Biological Sciences, TIFR, Bangalore -560065, India	Poster	Accepted
51.	Structural modelling of odorant receptors from <i>Aedes aegypti</i> and search for natural repellents Vikas Tiwari and R. Sowdhamini* National Centre for Biological Sciences, Tata Institute of Fundamental Research, Bangalore 560065, India	Poster	Accepted
52.	Integrative network analysis interweaves the missing links in cardiomyopathy diseasome Pankaj Kumar Chauhan & R. Sowdhamini* National Centre for Biological Sciences (TIFR), Bangalore, IN 560065	Poster	Accepted
53.	Structure analysis of HKT1;5 to understand the sodium transport affinity in wild rice species K. Mohamed Shafi¹, Gayatri Venkataraman² and Ramanathan Sowdhamini¹ ¹ National Centre for Biological Sciences (Tata Institute of Fundamental Research), Bangalore 560065 ² Plant Molecular Biology Laboratory, M.S. Swaminathan Research Foundation (MSSRF), Chennai 600113	Poster	Accepted
54.	A multi-epitope vaccine design against <i>Mycobacterium tuberculosis</i>: A subtractive proteomics and reverse vaccinology based immunoinformatics Approach Subhashree subhasmita Nayak, Ramadas Krishna* Department of Bioinformatics, Pondicherry University, Pondicherry-605014	Poster	Accepted
55.	The alteration of structural network by transient association between proteins Vasam Manjveekar Prabantu, Himani Tandon, Sandhya Sankaran, Ramanathan Sowdhamini, Narayanaswamy Srinivasan Molecular Biophysics Unit, Indian Institute of Science, Bangalore - 560012	Poster	Accepted
56.	In-Silico Identification of Hub Genes and their Regulatory Interaction Networks involved in Retinoic Acid Signaling Pathway of α-Synuclein in Parkinsons Diseases	Poster	Accepted

	Kajal Abrol, Amutha Ramaswamy* Department of Bioinformatics, School of Life Sciences, Pondicherry University, Kalapet, Puducherry-605014		
57.	Computational Analysis To Explore Possible Binding Between SIRT6 and PPARγ Parvathy J, Himani Tandon, N Srinivasan, N Ravi Sundaresan Interdisciplinary Mathematical sciences Initiative (IMI), IISc Bangalore, CV Raman Road, 560012	Poster	Accepted
58.	Pseudokinases repurpose flexibility signatures associated with the protein kinase fold for noncatalytic roles Seemadri Subhadarshini, Anindita Paul, Narayanaswamy Srinivasan Molecular Biophysics Unit, Indian Institute of Science, Bengaluru – 560012	Poster	Accepted
59.	Microsecond molecular dynamics simulations probing the conformational landscape of LIMK1 Hemavathy Nagarajan ^a , Vetrivel Umashankar ^b , Jeyaraman Jeyakanthan ^{a*} ^a Structural Biology and Bio-Computing Lab, Department of Bioinformatics, Science Block, Alagappa University, Karaikudi, 630 003, Tamil Nadu, India, ^b Department of Bioinformatics, ICMR-National Institute of Traditional Medicine, Nehru Nagar, Belagavi, Karnataka-590 010, India	Poster	Accepted
60.	Exploring the candidate genes encompassing the medicinal value of Athalakkai (<i>Momordica cymbalaria</i>) through transcriptomics approach Heyram Krishnakumar and Jeyaraman Jeyakanthan * Structural Biology and Bio-Computing Lab, Department of Bioinformatics, Science Block, Alagappa University, Karaikudi, 630 003, Tamil Nadu, India.	Poster	Accepted
61.	Comparative Studies on Bioactive Compounds of <i>Sphaeranthus amaranthoides</i> Against Parkinson's Disease: An <i>In-Silico</i> Approach Balasubramanian Sivaprakasam, Soundar Rajan Kulanthaivel, Priyanka Sappanimuthu, Joseph Sahayarayan Jesudass ^{1*} and Ramasamy Vidhyavathi Department of Bioinformatics, Alagappa University, Karaikudi, Tamilnadu, India-630004	Poster	Accepted
62.	Identification of Potent Lead Molecule against Serine/threonine-protein kinase B-raf receptor from <i>Endostemon viscosus</i> by using Computational approach Soundar Rajan Kulanthaivel, Balasubramanian Sivaprakasam, Priyanka Sappanimuthu, Jesudass Joseph Sahayarayan* and Ramasamy Vidhyavathi Department of Bioinformatics, Alagappa University, Karaikudi, Tamilnadu, India-630004	Poster	Accepted
63.	Consensus Secondary Structure Prediction (CSSP-2.0) Madhumathi Sanjeevi ^a , Ajitha Mohan ^b , Dhanalakshmi Ramachandran ^b , Jeyakanthan Jeyaraman ^{a*} and Kanagaraj Sekar ^{b*} ^a Structural Biology and BioComputing Laboratory, Department of Bioinformatics, Alagappa University, Karaikudi - 630 004, India and	Poster	Accepted

	^b Department of Computational and Data Sciences, Indian Institute of Science, Bangalore - 560012, India		
64.	Molecular Crosstalk between Sars-cov2 and Alzheimer's disease using Multi-Omics Datasets: A System Biology Approaches Premkumar T, Sajitha Lulu S*	Poster	Accepted
	Department of Biotechnology, School of Bio Sciences and Technology Vellore Institute of Technology, Vellore – 632014, Tamil Nadu		
65.	Role of NF-KB Pathway Genes In Rheumatoid Arthritis: A Network Biology Approach Devi Soorya Narayana S, Vino S*	Poster	Accepted
	School of Bio Sciences and Technology, Vellore Institute of Technology, Vellore, 632014		
66.	Evaluating Cell Surface Protein Liv1 as A Therapeutic Target In Triple-Negative Breast Cancer Patients Roshni S	Poster	Accepted
	Department of Human Genetics, Sri Ramachandra Institute of Higher Education & Research, Chennai.		
67.	Purification, Crystallization and Preliminary X-ray Diffraction Analysis of Succinyl-diaminopimelate desuccinylase from <i>Wolbachia</i> endosymbiont of <i>Brugia malayi</i> P. Saritha ^a , J. Jeyakanthan ^{a*} and Chun-Jung Chen ^{b*}	Poster	Accepted
	^a Department of Bioinformatics, Alagappa University, Science Block, Karaikudi – 630004 ^b Life Science Group, Scientific Research Division, National Synchrotron Radiation Research Center, No. 101號, Hsin Ann Rd, East District, Hsinchu, 300, Taiwan.		
68.	Identification of potential plant-based inhibitor against Scrub typhus through Computational approaches Sangavi P ¹ , Langeswaran K ^{2*}	Poster	Accepted
	¹ Department of Bioinformatics, Alagappa University, Karaikudi, Tamil Nadu, India ^{2*} Department of Biotechnology, Alagappa University, Karaikudi, Tamil Nadu, India		
69.	Identification of unique biomarkers of non-alcoholic and alcoholic based Hepatocellular carcinoma Annadurai Priyadharshini and Arnold Emerson Issac*	Poster	Accepted
	Department of Biotechnology, School of Bio science and Technology, Vellore Institute of Technology, Katpadi, Vellore -632014.		
70.	Development of novel amphiphilic drug-drug conjugates through the <i>in silico</i> approach against breast cancer S. Muthumanickam 1 , M.Manikandan 2 , P. Boomi 1* , J.Jeyakanthan 1 , M. Karunakaran	Poster	Accepted
	¹ Department of Bioinformatics, Alagappa University, Karaikudi, Tamil Nadu, India ² Department of Industrial Chemistry, Alagappa University, Karaikudi, Tamil Nadu, India ³ Department of Physics, Alagappa Govt. Arts College, Karaikudi, Tamil Nadu, India		
71.	<i>In silico</i> molecular docking and Molecular Dynamic Simulation of designed twin drug against cancer	Poster	Accepted

	<p>M.Manikandan¹, S. Muthumanickam², P. Boomi^{1,2*}, J. Jeyakanthan², H. Gurumalles Prabu¹</p> <p>¹Department of Industrial Chemistry, Alagappa University, Karaikudi, Tamil Nadu, India ²Department of Bioinformatics, Alagappa University, Karaikudi, Tamil Nadu, India</p>		
72.	<p>Bioinformatic Analysis for Drug Discovery Against Esophageal Cancer C. N. Rahul^a, Vinod Jangid^c, Kanagaraj Sekar^b, Natarajan Aiswarya^b, Aarthi Rashmi B^c & Jeyakanthan Jeyaraman^a</p> <p>^aStructural Biology and Biocomputing Lab, Department of Bioinformatics, Alagappa University, Karaikudi, 630004, India and ^bDepartment of Computational and Data Sciences, Indian Institute of Science, Bangalore 560012, India and ^cDepartment of Bioinformatics, Sri Krishna Arts and Science College, Coimbatore 641038, India</p>	Poster	Accepted
73.	<p>CAPS_protocol: A knowledge-driven approach to identify proteins of interest with an emphasis on biosynthetic pathways Adwait G. Joshi^a, K. Harini^a, Iyer Meenakshi^a, K. Mohamed Shafi^{a,b}, Shaik Naseer Pasha^a, Jarjapu Mahita^a, Radha Sivarajan Sajeevan^a, Snehal D. Karpe^a, Pritha Ghosh^a, Sathyanarayanan Nitish^{a,b}, A. Gandhimathi^a, Oommen K. Mathew^a, Subramanian Hari Prasanna^a, Manoharan Malini^a, Eshita Mutt^a, Mahantesha Naika^a, Nithin Ravooru^a, Rajas M. Rao^a, Prashant N. Shingate^a, Anshul Sukhwal^a, Margaret S. Sunitha^a, Atul K. Upadhyay^{a,c}, Rithvik S. Vinekar^a and Ramanathan Sowdhamini^{a,*}</p> <p>^a: National Centre for Biological Sciences (NCBS-TIFR), GKVK campus, Bellary road, Bangalore, 560065, Karnataka, India; ^b: The University of Trans-Disciplinary Health Sciences & Technology (TDU), Yelahanka, Bangalore, 560064, Karnataka, India; ^c: Department of Biotechnology, Thapar Institute of Engineering and Technology, Patiala, Punjab, 147004</p>	Poster	Accepted
74.	<p>Antimicrobial Resistance (AMR) Profiling and Phylogenetic Analysis of <i>Neisseria gonorrhoeae</i> Clinical Isolates from Kenya Juma M¹, Sankaradoss A², Ndombi R¹, Mwaura P¹, Damodar T², Nazir J², Pandit A², Khurana R², Masika M¹, Chirchir R¹, Gachie J¹, Krishna S^{2,3}, Sowdhamini R², Anzala O¹, Meenakshi IS^{1,*}</p> <p>¹KAVI Institute of Clinical Research, University of Nairobi, Nairobi, Kenya ² National Centre for Biological Sciences, Tata Institute of Fundamental Research (TIFR), Bengaluru, India ³ School of Interdisciplinary Life Sciences, Indian Institute of Technology Goa, Ponda, India</p>	Poster	Accepted
75.	<p>Identification of Critical Genes in Leukemia from Single-cell RNA Sequence Data: A Bioinformatics Approach Delsy Gnana Silvia D and Dr. Arnold Emerson^{1*}</p> <p>Department of Biotechnology, School of Bio Science and Technology, Vellore Institute of Technology (VIT). Katpadi, Vellore – 6320-14</p>	Poster	Accepted
76.	<p>Characterization of obligate and non-obligate classes of domain-domain interactions in multi-domain proteins Sidhanta Swayam Prakash Das, R. Sowdhamini and N. Srinivasan</p> <p>Molecular Biophysics Unit, Indian Institute of Science, Bangalore-560012</p>	Poster	Accepted

	National Centre for Biological Sciences, Bangalore		
77.	<p>Influence of ms 2 ct 6 A37: enlightening tRNA Lys ASL folding dynamics with codon AAA/AAG Ms. Ambika S. Dound*, Dr. Vishwambhar V. Bhandare and Prof. Kailas D. Sonawane</p> <p>Structural Bioinformatics Unit, Department of Biochemistry, Shivaji University, Vidyanagar, Kolhapur (MS), India- 416004</p>	Poster	Accepted
78.	<p>Structural Refinement and Validation of Modelled Streptomycin Binding Protein from <i>Staphylococcus aureus</i> Ms. Shweta D. Dhumale* and Ms. Minakshee D. Devadkar* and Dr. Kailas D. Sonawane</p> <p>Department of Microbiology, Shivaji University, Kolhapur 416004, India.</p>	Poster	Accepted
79.	<p>Homology modelling and molecular docking studies of penicillin binding protein Transpeptidase from <i>Salmonella enterica</i> subspecies Ms. Shivani J. Kharade*, Nitin M. Naik and Kailas D. Sonawane</p> <p>Shivaji University, Kolhapur (MS) 416004, India.</p>	Poster	Accepted
80.	<p>Structural insights to the novel PSEN1 mutations in Alzheimer disease Sanghamitra Uttam Chopade* and Vishwambhar V. Bhandare</p> <p>Department of Microbiology, Shivaji University, Kolhapur- 416 004, India</p>	Poster	Accepted
81.	<p>In silico studies of natural bioactives as a potential MAO-B and COMT inhibitors for the management of Parkinson's disease Pratibha Mali ^{a*}, Vishwambhar Bhandare ^b, Jyoti Jadhav ^a</p> <p>^a Department of Biochemistry, Shivaji University, Vidyanagar, Kolhapur, Maharashtra, India ^b Department of Microbiology, Shivaji University, Vidyanagar, Kolhapur, Maharashtra, India</p>	Poster	Accepted
82.	<p>Intermolecular interactions between Aβ peptides and human CathD in Alzhimers using molecular modelling Subodh Kamble* and Kailas Sonawane</p> <p>Structural Bioinformatics Unit, Department of Biochemistry, Shivaji University, Kolhapur, 416004, M.S., India</p>	Poster	Accepted
83.	<p>Structural destabilization of Luteinizing hormone beta (LHB) heterodimeric subunit by novel variant <i>p.Cys46Arg</i> Vishwambhar Bhandare ^{1*}, Anurag Lila ² and Ambarish Kunwar ³</p> <p>¹ Department of Microbiology, Shivaji University, Vidyanagar, Kolhapur, India - 416 004 ² Department of Endocrinology, Seth G S Medical College & KEM Hospital, Mumbai, India - 400 012 ³ Department of Biosciences and Bioengineering, IIT Bombay, Powai, Mumbai, India - 400 076</p>	Poster	Accepted
84.	<p>Computational search for potential COVID-19 drugs from FDA approved drugs and small molecules of natural origin identifies several anti-virals and plant products Abhishek Sharma, Vikas Tiwari and Ramanathan Sowdhamini*</p> <p>National Centre for Biological Sciences, TIFR, Bangalore -560065, India</p>	Poster	Accepted

85.	Comprehensive <i>in silico</i> analysis of structural and functional SNPs of the human CYP27B1 protein John Marshal Jayaraj and Karthikeyan Muthusamy*	Poster	Accepted
	Department of Bioinformatics, Alagappa University, Karaikudi – 630 004, Tamil Nadu, India		
86.	Hits Discovery on the NURR1 Receptor: Computational Approaches to Identify Agonist Compounds Thangam Nivethitha Saravanan, John Marshal Jayaraj, Swetha Subramanian and Karthikeyan Muthusamy*	Poster	Accepted
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	^a Department of Biotechnology, Dr. Umayal Ramanathan College for Women, Karaikudi-630003 ^b CSIR-Central Electrochemical Research Institute, Karaikudi – 630003 ^c Orbito Asia Diagnostics, Coimbatore – 641045 ^d Department of Bioinformatics, Alagappa University, Science Block, Karaikudi – 630004		
92.	<i>In silico</i> studies on protein based nanoparticles against colorectal cancer Muthulakshmi Muthumanickam ¹ , Rajaram Abhirami ² , Rameshthangam	Poster	Accepted

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102.	<p>Lawsonia inermis Flower Aqueous Extract Treatment Alleviates Diabetes associated cognitive Impairment Shahanaj. I & Natarajan. D</p> <p>Natural Drug Research Laboratory, Department of Biotechnology, Periyar University, Salem- 636011 TN, INDIA.</p>	Poster	Accepted
103.	<p>Identification of high affinity small molecule targeting 5-ALOX for the clinical treatment of pancreatic cancer Anushka Bhrdwaj¹, Anuraj Nayarisseri², Sanjeev Kumar Singh^{1*}</p> <p>¹ Department of Bioinformatics, Alagappa University, Science Block, Karaikudi – 630004, ² In silico Research Laboratory, Eminent Biosciences, Indore - 452010</p>	Poster	Accepted
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105.	<p>Structural Insights into the Molecular Design of NS2B/NS3 Phytochemical for the treatment of Dengue virus Khushboo Sharma¹, Anuraj Nayarisseri², Sanjeev Kumar Singh^{1,*}</p> <p>¹Department of Bioinformatics, Alagappa University, Science Block, Karaikudi-630004, ²In silico Research Laboratory, Eminent Biosciences, Indore-452010</p>	Poster	Accepted
106.	<p>Structure elucidation of phytochemicals in blocking Protein-DNA interactions of sporulation mechanism in <i>Bacillus Cereus</i> Rajaram Abhirami¹, Chandrabose Selvaraj¹, Sanjeev Kumar Singh^{1*}</p> <p>¹Computer Aided Drug Design and Molecular Modeling Lab, Department of Bioinformatics, Alagappa University, Karaikudi, India</p>	Poster	Accepted
107.	<p>Structural analysis of Gp46 protein for exploring the HTLV-1 inhibitors using virtual screening, Quantum Mechanical and Molecular Dynamics analysis Gurunathan Rubha Shri¹, Chandrabose Selvaraj¹, Sanjeev Kumar Singh^{1*}</p> <p>¹Computer Aided Drug Design and Molecular Modeling Lab, Department of Bioinformatics, Alagappa University, Karaikudi, India</p>	Poster	Accepted

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112.	<p>A Compendious study of Dengue NS1 protein and its interaction with Human RPL18 using Network Analysis and Drug Discovery approach Muthuraja Arun Pravin, Chandrabose Selvaraj and Sanjeev Kumar Singh*</p> <p>Computer Aided Drug Design and Molecular Modelling Lab, Department of Bioinformatics, Alagappa University, Karaikudi-630 004, Tamil Nadu, India</p>	Poster	Accepted
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115.	<p>Decoding the whole genome sequence of a clinical isolate of <i>Pseudomonas aeruginosa</i> Bhuvaneswari Narthanareeswaran and Jeyaraman Jeyakanthan*</p> <p>Structural Biology and Biocomputing Lab, Department of Bioinformatics, Alagappa University, Karaikudi – 630004, Tamil Nadu, India.</p>	Poster	Accepted
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117.	<p>Structure based design of anti-bacterial drugs using mammalian heme peroxidases as potential drug targets. V. Viswanathan ^{1,2}, J. Jeyakanthan ¹, P. Sharma ², S. Sharma ² and T. P. Singh ²</p> <p>¹ Department of Bioinformatics, Alagappa University, Karaikudi, Tamil Nadu. ² Department of Biophysics, All India Institute of Medical Sciences, New Delhi.</p>	Poster	Accepted
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119.	<p>Computational insights into the conformational dynamics of drug resistance mutation in the <i>pncA</i> gene of <i>Mycobacterium tuberculosis</i> Manikandan Jayaraman and Jeyakanthan Jeyaraman*</p> <p>Structural Biology and Biocomputing Lab, Department of Bioinformatics, Alagappa University, Karaikudi – 630004, Tamil Nadu, India.</p>	Poster	Accepted

120.	<p>Identification of potential inhibitor against Hepatocellular Carcinoma through <i>in silico</i> approach Srinithi R¹ , Nachammai KT² , Sangavi P¹ , Langeswaran K^{2*}</p> <p>¹Department of Bioinformatics, Science Campus, Alagappa University, Karaikudi, Tamil Nadu, India ²Department of Biotechnology, Science Campus, Alagappa University, Karaikudi, Tamil Nadu, India</p>	Poster	Accepted
121.	<p>Whole Genome Sequencing and annotation studies of <i>Nocardia farcinica</i> N.Shaslinah, Chitra jeyaraj pandian, Jemima D kingsley, and Jeyakanthan Jeyaraman*</p> <p>Structural Biology & Bio-Computing Lab, Department of Bioinformatics, Alagappa University, Karaikudi-630004, Tamil Nadu, India</p>	Poster	Accepted
122.	<p>Potent Inhibitors for Tetracycline Repressor Family of Transcription Regulator protein in Tetracycline-resistant strain of <i>Acinetobacter baumannii</i> Karthika Alagesan¹ , Chitra Jeyaraj Pandian² , Jemima Kingsley³ , Jeyaraman Jeyakanthan¹ *</p> <p>¹ Structural Biology and Bio-Computing Lab, Department of Bioinformatics, Alagappa University, Karaikudi 630004, Tamil Nadu, India. ² Department of Biotechnology, Dr. Umayal Ramanathan College for Women, Karaikudi 630003, Tamil Nadu, India.</p>	Poster	Accepted
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124.	<p>Unravelling the functional mode/mechanism of streptomycin adenylyltransferase from <i>Serratia marcescens</i> D. Prabhu¹, S. Rajamanikandan¹, J. Jeyakanthan², P. Ramasamy¹</p> <p>¹ R & D Wing, Sree Balaji Medical College and Hospital-BIHER, Chennai-600 044, India, ² Department of Bioinformatics, Alagappa University, Karaikudi-630 003, India E-mail:jjeyakanthan@alagappauniversity.ac.in</p>	Poster	Accepted

125.	<p>Computational screening of potent phytochemicals from medicinal plants as SARS-CoV-2 NSP1 inhibitors</p> <p>S. Rajamanikandan¹, D. Prabhu¹, J. Jeyakanthan², P. Ramasamy¹</p> <p>¹ R & D Wing, Sree Balaji Medical College and Hospital-BIHER, Chennai-600 044, India, ² Department of Bioinformatics, Alagappa University, Karaikudi-630 003, India E-mail:jjeyakanthan@alagappauniversity.ac.in</p>	Poster	Accepted
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