



## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** International Journal of Biological Macromolecules

**IF:** 8.7

**Title:** Emerging roles of micropeptides in calcium cycling dysregulation and atrial fibrillation: A review

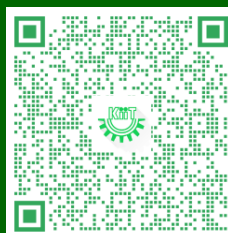
**Author:** Pani P.; Swalsingh G.; Sadayappan S.; Bal N.C.

**Details:** Volume 370, July 2026

**Abstract:** Atrial fibrillation (AF) is the most common sustained cardiac arrhythmia globally, marked by chaotic electrical impulses and irregular atrial contraction. At the cellular level, defective calcium ( $\text{Ca}^{2+}$ ) handling plays a central role in AF pathogenesis, with the sarco/endoplasmic reticulum  $\text{Ca}^{2+}$ -ATPase (SERCA) pump being a critical determinant of intracellular  $\text{Ca}^{2+}$  reuptake and myocardial relaxation. Recent research has uncovered a class of small transmembrane micropeptides, phospholamban (PLB), sarcolipin (SLN), dwarf open reading frame (DWORF), myoregulin (MLN), endoregulin (ELN) and another-regulin (ALN), that directly modulate SERCA activity. Interestingly, these micropeptides exhibit chamber-specific expression and diverse regulatory mechanisms, functioning as inhibitors, uncouplers or facilitators of SERCA activity and are increasingly linked to atrial arrhythmogenesis. We explore evidence from genetically modified animal models and patient-derived data to elucidate how dysregulation of these peptides, particularly SLN and PLB, contributes to abnormal EC-coupling like  $\text{Ca}^{2+}$  cycling, delayed afterdepolarizations, oxidative stress and atrial remodeling. We also examine the influence of systemic metabolic regulators, such as thyroid hormones, catecholamines, dexamethasone and various exercise paradigms on micropeptide expression and function, offering insight into their intersection with AF progression. By dissecting the spatial, temporal and metabolic regulation of SERCA micropeptides, this review aims to offer current understanding about chamber-specific modulation of  $\text{Ca}^{2+}$  homeostasis and use these insights towards treatment of AF.



**URL:** <https://www.sciencedirect.com/science/article/abs/pii/S0141813026028503?via%3Dihub>





# SCHOLARLY PUBLICATIONS

## School of Biotechnology

### KIIT Deemed to be University

**Journal Name:** Materials Today Advances

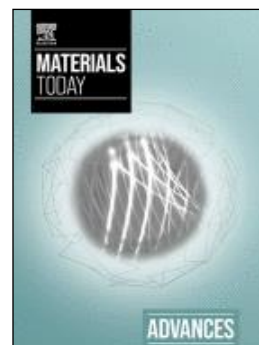
**IF:** 8.0

**Title:** Zein nanoparticles as emerging moderators of the gut microbiome: Mechanisms, biomedical potentials and translational challenges

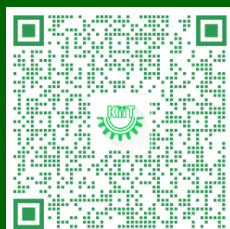
**Author:** Dutta J.; Parashar R.; Mishra S.; Kumari S.; Kujawska M.; Kaushik N.K.; Ghosh A.; Verma S.K.

**Details:** Volume 30, June 2026

**Abstract:** Zein, a hydrophobic prolamin protein derived from maize, has emerged as a sustainable material for oral nanocarrier design due to its biodegradability, biocompatibility, and spontaneous self-assembly into stable NPs. This review critically examines the synthesis strategies, physicochemical behavior, and biological performance of zein NPs (ZNPs) with particular emphasis on gastrointestinal delivery and microbiome-associated outcomes. Current evidence indicates that ZNPs act primarily as protective and transport matrices: their intrinsic properties govern gastric resistance, mucus interaction, and controlled intestinal release, thereby improving stability and bioavailability of encapsulated compounds. The review insights into the microbiome pathway within a carrier-enabled exposure framework to correlate the attributing causality by ZNPs in the frame of the reported alterations in microbial composition, metabolite production, and host physiology arising from delivered bioactive (e.g., drugs, polyphenols, or probiotics). Moreover, the study details the advances in fabrication approaches, including desolvation, electrospraying, flash nanoprecipitation, and microfluidic micromixing, which have improved particle uniformity and scalability, supporting applications in pharmaceutical and nutraceutical delivery. Future progress will require integrating empty-carrier controls, longitudinal microbiome analyses, harmonized production standards, and nano-specific regulatory evaluation frameworks. Establishing these criteria will enable rational design of zein-based systems as controlled oral delivery platforms rather than assuming intrinsic microbiome-modulating activity.



**URL:** <https://www.sciencedirect.com/science/article/pii/S2590049826001530?via%3Dihub>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Biomedicine and Pharmacotherapy

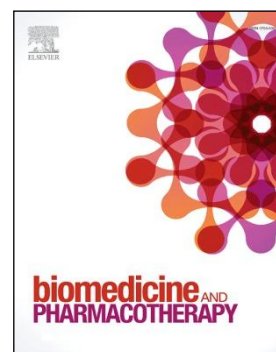
**IF:** 7.5

**Title:** Corrigendum to “The posterity of Zebrafish in paradigm of in vivo molecular toxicological profiling” (Biomedicine & Pharmacotherapy, (2024), 171, C, (116160), (S0753332224000416), 10.1016/j.biopha.2024.116160)

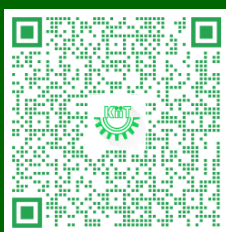
**Author:** Verma S.K.; Nandi A.; Sinha A.; Patel P.; Mohanty S.; Jha E.; Jena S.; Kumari P.; Ghosh A.; Jerman I.; Chouhan R.S.; Dutt A.; Samal S.K.; Mishra Y.K.; Varma R.S.; Panda P.K.; Kaushik N.K.; Singh D.; Suar M.

**Details:** Volume 199, June 2026

**Abstract:** This Corrigendum relates to the above article. In this article, the authors have included references [4, 6, and 59] that require correction due to contextual inappropriateness: [4] M. Harper Book of the month: MMR: science and fiction, J. R. Soc. Med., 97 (2004), pp. 552–553, 10.1177/014107680409701115 [6] C. Yu, R. Chen, J.J. Li, J.J. Li, M. Drahansky, M. Paridah, A. Moradbak, A. Mohamed, H. Abdulwahab taiwo Owolabi, FolaLi, M. Asniza, S.H. Abdul Khalid, T. Sharma, N. Dohare, M. Kumari, U.K. Singh, A.B. Khan, M.S. Borse, R. Patel, A. Paez, A. Howe, D. Goldschmidt, C. Corporation, J. Coates, F. Reading, We are IntechOpen, the world ' s leading publisher of Open Access books Built by scientists, for scientists TOP 1%, Intech, 13 (2012), 10.1016/j.colsurfa.2011.12.014 [59] E. Duffey Global biodiversity: status of the earth's living resources, Biol. Conserv., 66 (1993), p. 145, 10.1016/0006–3207(93)90147-s The authors confirm that these references were inadvertently introduced in the article. The updated references for [4, 6 and 59] should be as below: [4] J. Payne-James, Clinical Toxicology: Principles & Mechanisms, J. R. Soc. Med. 97 (2004) 554–555. <https://doi.org/10.1177/014107680409701117>. [6] R. Nagel, K. Isberner, Testing of chemicals with fish — a critical evaluation of tests with special regard to zebrafish, Fish Ecotoxicol. (1998) 337–352. [https://doi.org/10.1007/978–3-0348–8853-0\\_11](https://doi.org/10.1007/978–3-0348–8853-0_11). [59] R. Turner. Acute toxicity: The determination of LD50. Screening Methods In Pharmacology.



**URL:** <https://www.sciencedirect.com/science/article/pii/S0753332226005536?via%3Dihub>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Journal of Environmental Chemical Engineering

**IF:** 7.5

**Title:** Nano-biohybrid photocatalysts for sustainable wastewater treatment: Recent advances and future perspectives

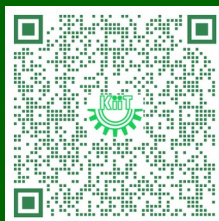
**Author:** Kumar M.; Sahoo P.C.; Pillai S.; Raut R.; Parida K.

**Details:** Volume-14, Issue-5, October 2026

**Abstract:** Currently, the rising discharge of industrial, agricultural and domestic effluents has severely contaminated the water bodies with a wide variety of organic and inorganic pollutants such as dyes, medicines, heavy metals and persistent organic compounds. Traditional wastewater treatment processes are often characterized by low efficiency, being costly to operate, secondary contamination, and weak elimination of the refractory pollutants. In that regard, it is possible to note that nano-biohybrid photocatalysts are becoming a prospective and viable alternative to advanced wastewater treatment. The mini-review will include a detailed account of the latest breakthroughs in designing, manufacturing and applications of nano- biohybrid photocatalysts with an accent on their prospects to alleviate the existing environmental issues. In detail, the synthesis strategies of nano-biohybrid systems (including incorporation of nanoparticles with enzymes, microbial cells or biomolecules), the importance of surface engineering, heterojunction formation and defect modulation in enhancing photocatalytic performance are reviewed. The mechanistic understanding of the pollutant degradation is established, where emphasis is made on the reactive oxygen species (ROS) formation, the efficiency of charge separation, and synergistic bio-nano interactions which enable the selective and quick degradation of a broad spectrum of pollutants. Recent studies have established that nano- biohybrid photocatalysts are more effective in degrading efficiencies, broad-spectrum pollutant removal, stability in operation, reuse and less energy is required as compared to traditional photocatalysts. These systems are special in that they incorporate biological specificity and catalytic activity with high surface area, tunable electronic structure, and light absorption characteristics of nanomaterials that lead to environmentally friendly, energy efficient, and long-term wastewater treatment. Lastly, the review mentions the current obstacles like material, scale-up, cost-effectiveness, and regulatory issues, and future trends including the creation of multifunctional, self-regenerative and scalable nano-biohybrid photocatalysts.



**URL:** <https://www.sciencedirect.com/science/article/abs/pii/S2213343726026175>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Journal of Agriculture and Food Research

**IF:** 7.2

**Title:** Functional properties, health benefits, and biotechnological applications of the oyster mushroom (*Pleurotus ostreatus*): A comprehensive review

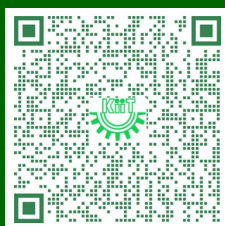
**Author:** Mukherjee K.; Panda S.K.; Samanta I.; Goswami L.; Maity P.J.; Love S.K.; Sasikumar R.; Chowdhary G.; T S.K.; Jaiswal A.K.

**Details:** Volume-29, July 2026

**Abstract:** *Pleurotus ostreatus*, commonly known as the “multi-purpose mushroom,” is an edible fungus distinguished by its unique convergence of nutritional, medicinal, and environmental attributes. In recent years, it has emerged as one of the most extensively studied and widely cultivated mushroom species, owing to its broad substrate adaptability and low input requirements. This review consolidates current knowledge of its nutritional composition and phytochemical profile, emphasising a diverse array of bioactive compounds with antimicrobial, anticancer, antioxidant, antidiabetic, anti-arthritis, and antiviral activities. The species is increasingly recognised as a functional food and a sustainable reservoir of bioactive molecules with significant potential for pharmaceutical applications. Emerging evidence also suggests its role in promoting mental health and overall well-being. The review further explores the interface between *P. ostreatus* and nanobiotechnology, as well as its potential applications. *P. ostreatus* secretes a spectrum of oxidative and ligninolytic enzymes, including lignin peroxidase, manganese peroxidase, laccases, and cellulases, enabling the degradation of pesticides, recalcitrant hydrocarbons, complex lignin polymers, and industrial pollutants. The multifaceted significance of *P. ostreatus* across functional foods, nutraceuticals, nanotechnology, therapeutics, and bioremediation positions it as an underexploited yet powerful platform for sustainable development, with strong potential to contribute to achieving the United Nations Sustainable Development Goals (2, 3, and 17).



**URL:** <https://www.sciencedirect.com/science/article/pii/S2666154326004734>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** LWT

**IF:** 7.1

**Title:** Development and characterisation of *Prunus nepalensis* fruit-aquafaba probiotic powder by vacuum freeze drying

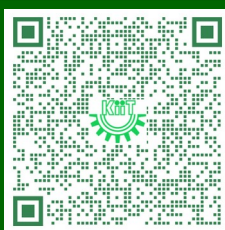
**Author:** Nongmaithem R.; Senthilkumar K.; Selva Kumar T.; Kaviarasu G.; Biswas S.; Panda S.K.; Rajarathinam R.; Sasikumar R.; Jaiswal A.K.

**Details:** Volume-253, 1<sup>st</sup> August, 2026

**Abstract:** Usage of probiotics in plant-based matrices has increased due to the addition of functional foods to the human diet. This work subjected to produce powdered probiotic sohiong juice to add value to a fruit rich in bioactives. *Lactobacillus plantarum* subsp. *lactis* (MCC 2974) inoculated into sohiong juice with different concentrations of aquafaba as a cryoprotectant and freeze dried. The freeze dried samples were characterized for the viable cell count, physicochemical properties, storage properties, and morphological characterisation. A formulation (F3) that contains probiotic juice and aquafaba at a 1:0.3 (v/v) ratio exhibits improved physicochemical properties, including lower hygroscopicity ( $13.12 \pm 0.39\%$ ), indicating lower moisture absorption. This is followed by a wettability measurement of  $38.15 \pm 0.36$  s, which indicates the time taken to sink the powder sample in water; it also retained colour parameters. The aquafaba mixed formulation shows increasing loose bulk density and reduction in porosity. Hausner Ratio and Carr's Index value prove that the formulated F3 sample passes the scale of flowability as per United States Pharmacopeia standards. Field Emission Scanning Electron Microscopy (FESEM) images confirmed proper particle formation of probiotics, followed by Fourier Transform Infrared (FTIR) Spectroscopy, which proved that the aquafaba formulation doesn't lose functional groups that present in the probiotic juice. Thermogravimetric Analysis (TGA) (30% residue at 800 °C) and Differential Scanning Calorimetry (DSC) (18.995 J/g) analysis results showed that aquafaba improved the thermal stability of sohiong fruit juice. It shows the potential of combining underutilized fruits with plant-based stabilizers to develop sustainable, health-promoting nutraceuticals.



**URL:** <https://www.sciencedirect.com/science/article/pii/S0023643826006869>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** International Journal of Pharmaceutics

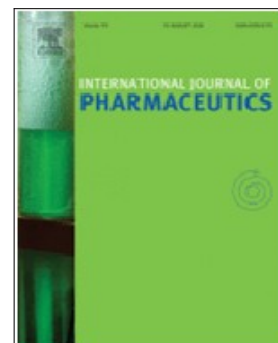
**IF:** 6.0

**Title:** Artificial intelligence and CRISPR-based approaches for targeted delivery of bacteriophages

**Author:** Pradhan R.R.; Pati S.; Samal S.K.

**Details:** Volume-701, 10<sup>th</sup> August, 2026

**Abstract:** The rapid emergence of Multidrug-Resistant (MDR) bacteria has increased interest in bacteriophage therapy as a promising alternative to conventional antibiotics. Bacteriophages are host-specific bacterial viruses that selectively infect and destroy pathogenic bacterial strains. Recent developments in artificial intelligence (AI) and Clustered Regularly Interspaced Short Palindromic Repeats (CRISPR)-based technologies offer innovative approaches to address challenges such as narrow host range, rapid immune clearance, phage instability, bacterial resistance, and biofilm penetration barriers. By integrating AI-driven structural modeling with CRISPR-mediated genome editing, these methods enable the targeted delivery of bacteriophages. This review focuses on next-generation approaches that combine AI-assisted phage identification, host prediction, and therapeutic optimization with CRISPR-based genome engineering for targeted phage delivery and improved safety. Overall, this review highlights the potential of AI- and CRISPR-assisted phage therapy for the treatment of MDR bacterial infections. It will also provide a systematic overview of bacteriophage biology, life cycle, and mechanisms of action, while focusing on the influence of phage morphology on therapeutic performance, recent advances, current clinical, preclinical studies, and future perspectives. Although phage therapy shows considerable potential against MDR bacterial infections, several challenges related to delivery, safety, and clinical translation remain. The integration of AI and CRISPR technologies has the potential to improve phage selection, targeting specificity, and therapeutic performance. However, continued research, clinical validation, and regulatory development will be essential for translating these advances into practical antimicrobial therapies.



**URL:** <https://www.sciencedirect.com/science/article/abs/pii/S0378517326006034>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Frontiers in Pharmacology

**IF:** 5.4

**Title:** Editorial: Emerging and reemerging neglected tropical diseases: their epidemiology, transmission, mitigation, and vaccines and chemotherapeutics advancements, volume II

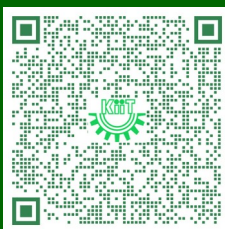
**Author:** Mohapatra, RK; Mishra, S; Kandi, V; Mohapatra, NP; Sirka, CS

**Details:** Volume-17, 5<sup>th</sup> June, 2026

**Abstract:** Neglected tropical diseases (NTDs) remain a major global health challenge that affects more than 1.5 billion people across nations. Diseases like lymphatic filariasis, leishmaniasis, schistosomiasis, and trachoma, to mention a few, impact impoverished communities that lack clean water, sanitation and healthcare disproportionately. The World Health Organisation (WHO) recognised 21 NTDs, which collectively cause over 200,000 deaths annually and contribute to millions of disability-adjusted life years (DALYs), outdoing the Human Immunodeficiency virus (HIV)/Acquired Immunodeficiency Syndrome (AIDS), tuberculosis and malaria burden. Termed as 'neglected', many of these diseases have been wiped out from most of the developed societies, while still persisting in the poor and most marginalised ones (Álvarez-Hernández et al., 2020). Despite the progress, including eliminating the guinea worm disease in most regions and reducing trachoma and yaws significantly, many NTDs still persist in Africa, Southeast Asia and Latin America (Srivastava et al., 2023; Ca et al., 2024; Mohapatra et al., 2024). Mass drug administration programmes have reached billions worldwide, yet institutional funding remains critically low, with only 0.6% of the global health development assistance directed toward NTDs. WHO's 2030 roadmap sets ambitious targets of reducing the number of people requiring NTD interventions by 90%, cutting DALYs by 75%, and achieving eradication of dracunculiasis and yaws (WHO, 2025). However, challenges of underreporting, weak surveillance, climate change and disruptions due to Coronavirus Disease 2019 (COVID-19) pandemic continue to hinder the progress.



**URL:** <https://www.frontiersin.org/journals/pharmacology/articles/10.3389/fphar.2026.1864825/full>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Materials Chemistry and Physics

**IF:** 5.2

**Title:** Intraoral biocompatible mucoadhesive buccal patch imbued with bromelain for targeting oral fibrosis: Evaluation of functional activity, cytotoxicity, and safety

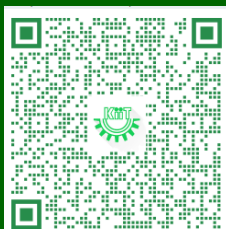
**Author:** Kokkanti R.R.; Bajoria A.A.; Parida N.; Viswanathan S.; Pritam P.; Kumar N.S.; Kushwaha G.S.; Panda A.K.; Veeraraghavan G.; Patnaik S.

**Details:** Volume 361, August 2026

**Abstract:** A bromelain-loaded mucoadhesive buccal patch was developed for enzymatic therapy in oral submucous fibrosis (OSF), a progressive fibrotic disorder characterized by excessive collagen deposition. The patch was analytically characterized for content, release kinetics, and enzymatic stability. HPLC and LC-MS/MS confirmed bromelain content and integrity, while in vitro release studies demonstrated near-complete release within 90 min in phosphate-buffered saline and artificial saliva. Zymography revealed 76.67 U/mL enzymatic activity at 2 h, even after 6 months of storage, confirming long-term stability. LC-MS peptide mapping validated collagen degradation. Ex vivo studies showed prolonged mucoadhesion (~137 min), supporting effective mucosal retention. HPLC confirmed drug purity. Biological assays including MTT, colony formation, wound healing, and live/dead staining indicated selective cytotoxicity and apoptosis in KB-3-1 oral cancer cells, with high viability of normal fibroblasts. Acute toxicity studies in rats revealed no systemic toxicity or organ damage. Together, these findings establish a robust analytical and biological profile for the patch, supporting its potential as a non-invasive, enzyme-based therapy for OSF. Future studies will focus on dose optimization, long-term in vivo efficacy, and translational potential.



**URL:** <https://www.sciencedirect.com/science/article/pii/S0254058426006814?via%3Dihub>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** ACS Omega

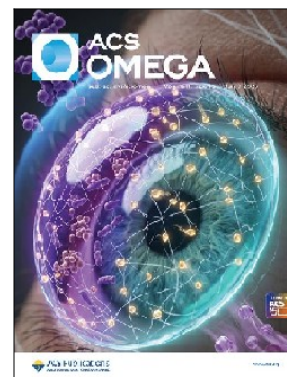
**IF:** 5.2

**Title:** Designing a Whole-Cell Biosensor for Detection of Toxic Metals Using Intein Splicing Inhibition of Mycobacterium tuberculosis SufB Protein

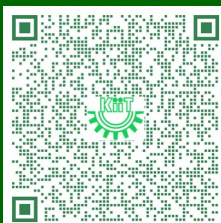
**Author:** Mehra A.; Nanda A.; Nayak S.

**Details:** Volume-11, Issue-26, 7<sup>th</sup> July, 2026

**Abstract:** Heavy metal contamination, driven by anthropogenic disruption of natural geochemical cycles, has led to widespread bioaccumulation, posing a serious global health threat. Addressing this crisis demands the development of simple and sensitive strategies capable of monitoring metal pollution in the environment. Leveraging nongenetically modified native organisms, this research offers a promising low-cost, eco-friendly screening approach for detecting heavy metals in contaminated samples. Current study uses post-translational splicing of the Mycobacterium tuberculosis (Mtb) SufB precursor protein to detect metals by linking metal-induced splicing inhibition to viability loss of native mycobacterial cells. Toxic metal ions like  $\text{Cd}^{2+}$  and  $\text{Hg}^{2+}$  blocked splicing activity of Mtb SufB precursor protein over a concentration range of 25  $\mu\text{M}$ –2 mM, while  $\text{Pb}^{2+}$  and  $\text{Cr}^{3+}$  failed to do so. An innovative biosensor platform was designed to detect metals by a simple Alamar Blue assay using attenuated Mtb H37Ra as indicator cells. Qualitative metal detection was assessed via colorimetric variation relating to mycobacterial viability, while concurrent spectral absorbance measurement enabled metal ion quantification. Loss of H37Ra cell viability by metal ions over the 25  $\mu\text{M}$ –2 mM concentration range highlighted the sensitivity of the designed biosensor, while the addition of metal-specific chelators reversed the effect. Multiplexing ability was evaluated by including known splicing inhibitors like  $\text{Cu}^{2+}$ ,  $\text{Zn}^{2+}$ , and  $\text{Pt}^{4+}$  over various concentration ranges alongside  $\text{Cd}^{2+}$  and  $\text{Hg}^{2+}$  in a simple 96-well plate format. The designed intein-based biosensor offers a user-friendly platform, readily standardizable for high-throughput detection using native organisms harboring metal-sensing precursor proteins. As a proof-of-concept, this study demonstrates the applicability of intein-based biosensing for initial heavy metal screening in environmental and industrial effluents, serving as a rapid and accessible tool prior to targeted advanced analysis.



**URL:** <https://pubs.acs.org/acsodf/article/11/26/38663/5167004/Designing-a-Whole-Cell-Biosensor-for-Detection-of>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Progress in Additive Manufacturing

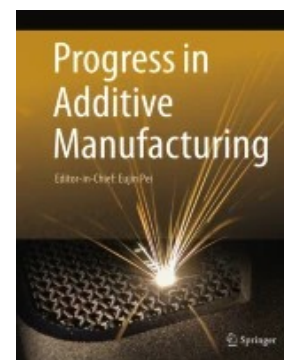
**IF:** 5.2

**Title:** Fabrication of cost-effective glass-bottom cell-culture device using fused deposition modeling: new avenue for bioimaging

**Author:** Swain S.; Shahul Hameed S.; Jain S.; Kumar S.S.; Pati F.; Unni H.N.; Suar M.; Giri L.

**Details:** 14<sup>th</sup> July, 2026

**Abstract:** Achieving high-resolution 4D imaging (XYZt) of larger areas is paramount for comprehensively characterizing engineered tissues and disease models. Yet the high cost and optical requirements of glass-bottom devices, which are essential for confocal microscopy, often hinder the use of such advanced imaging. This study presents an innovative method for crafting cost-effective microfluidic devices to overcome this challenge. Diverging from traditional soft lithography techniques, which require cleanroom facilities and costly materials, our approach harnesses fused deposition modeling (FDM) to fabricate glass-bottom polydimethylsiloxane (PDMS) devices that seamlessly incorporate a 0.17 mm glass coverslip optimized for laser scanning confocal microscopy (LSCM). Using glass-embedded acrylonitrile butadiene styrene (ABS) templates, we achieve precise fabrication of spiral channels in PDMS, thereby substantially reducing associated costs, including installation, infrastructure, and maintenance. The resultant device boasts numerous functionalities, facilitating diverse applications such as cell culture, reagent mixing, morphology monitoring, and on-chip immunoassays. Moreover, we showcase its versatility by demonstrating its efficacy for 4D calcium imaging with a resonance scanner in an LSCM, using HMC3 (human microglial) and MCF-7 (human breast cancer) cell lines. Beyond its cost-effectiveness, this biochip platform is suitable for applications such as toxicity analysis, drug screening, and real-time monitoring. This advancement promises new avenues for comprehensive bioimaging research, offering affordable and accessible solutions for studying complex biological systems. By democratizing access to high-resolution imaging, our method paves the way for a deeper understanding and characterization of diverse biological phenomena. Furthermore, its cost-effectiveness and ease of fabrication hold promise for adoption across various research fields, empowering researchers to further explore biological processes.



**URL:** <https://link.springer.com/article/10.1007/s40964-026-01837-y>





**SCHOLARLY PUBLICATIONS**  
**School of Biotechnology**  
**KIIT Deemed to be University**

**Journal Name:** Journal of Molecular Structure

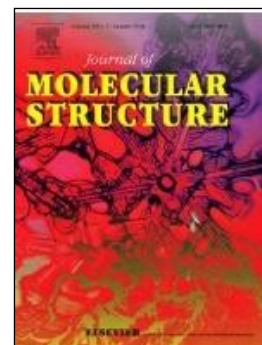
**IF:** 4.9

**Title:** Tetranuclear defective dicubanes of vanadium complexes: Synthesis, characterization and their potential biological activities

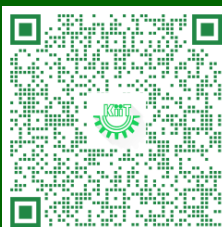
**Author:** Singh A.R.; Rajpurohit A.S.; Khongsai L.; Singh S.A.; Chingakham B.S.; Jaccob M.; Lonibala R.

**Details:** Volume 1373, October 2026

**Abstract:** Four tetranuclear V(V) complexes 1-4 with unique  $\{V_4O_8\}$  defective dicubane cores were synthesized through novel route under methanolic condition using bis(acetylacetonato) oxovanadium(IV) precursor along with 2,2'-bipyridine and substituted 2,2'-bipyridine ligands i.e. 2,2'-bipyridine in (1), 4,4'-dimethyl-2,2'-bipyridine (2), 5,5'-dimethyl-2,2'-bipyridine (3) and 4,4'-dimethoxy-2,2'-bipyridine (4). The complexes are characterized on the basis of CHN, UV-visible, IR and SC-XRD analysis data. Single-crystal X-ray diffraction analysis revealed the solid-state structures of all complexes (1-4) with distorted octahedral geometries for the four V(V) ions. The electronic nature of the complexes and their molecular surfaces were investigated using density functional theory (DFT) and Hirshfeld surface analysis. In vitro antidiabetic studies indicated that the complexes possess good antidiabetic activities against  $\alpha$ -glucosidase and in vitro studies of anticancer activities against A549 cells, complex 3 shows better activity with  $IC_{50} = 2.5 \pm 1.07 \mu M$ .



**URL:** <https://www.sciencedirect.com/science/article/abs/pii/S0022286026013293?via%3Dihub>





**SCHOLARLY PUBLICATIONS**  
**School of Biotechnology**  
**KIIT Deemed to be University**

**Journal Name:** Scientific Reports

**IF:** 4.9

**Title:** Structure-based virtual screening and molecular dynamics elucidate novel inhibitors of HIV-1 TAR RNA from enamine library

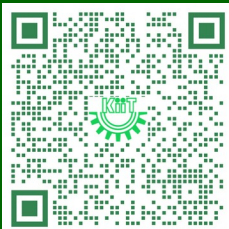
**Author:** Mundlia P.; Singh S.P.; Gartia J.; Ranawat P.; Singh G.; Barnwal R.P.

**Details:** Volume-16, Issue-1, December 2026

**Abstract:** The HIV-1 trans-activation response (TAR) RNA is a key regulatory element necessary for viral replication, presenting a high-value target for anti-HIV therapeutics. Recent advances in RNA-targeted drug discovery have leveraged large, diverse compound libraries, such as the Enamine collection, using structure-based virtual screening to uncover novel modulators of RNA structure and function. In this study, we employed an integrative in silico approach combining high-throughput virtual screening of the Enamine library against the central bulge region of HIV-1 TAR RNA with molecular dynamics (MD) simulations to assess binding stability and conformational adaptability of lead molecules. Virtual screening pinpointed multiple candidate ligands predicted to disrupt the critical Tat-TAR interaction interface, with subsequent MD simulations over 500-nanosecond timescales revealing ligand-induced stabilization of druggable RNA conformations and dynamic remodelling of key binding pockets. Notably, some top-scoring compounds featured strong and persistent contacts within the U23-C25 bulge and the adjacent major groove, recapitulating pharmacologically relevant mechanistic disruption. These findings reinforce the synergy of structure-based screening and MD for identifying and validating RNA-targeted small molecules and provide promising new scaffolds for further experimental testing as selective HIV-1 replication inhibitors.



**URL:** <https://www.nature.com/articles/s41598-026-50156-3>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Journal of the Science of Food and Agriculture

**IF:** 4.0

**Title:** Advantages of low-glycaemic-index carbohydrates in regulating glycaemic response of energy bar: A review

**Author:** Sasikumar, R; Gupta, A; Kumar, TS; Ganesan, K; Haorongbam, S; Rajarathinam, R; Janghu, S; Panda, SK; Jaiswal, AK

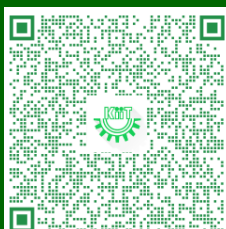
**Details:** June 2026

**Abstract:** Energy bars are widely consumed functional foods among athletes and physically active consumers who require convenient sources of sustained energy. Their carbohydrate composition, particularly the type and ratio of digestible sugars such as glucose, fructose and sucrose, strongly influences glycaemic index (GI), postprandial blood glucose response and subsequent metabolic outcomes. GI reflects the rise in blood glucose after food consumption relative to a reference carbohydrate - usually glucose. Low-GI carbohydrates are digested and absorbed more slowly, resulting in a moderated insulin response and potentially greater reliance on fat oxidation during exercise.. Evidence from formulation studies indicates that partial substitution of sucrose with inulin and FOS can reduce GI while maintaining energy density. For example, replacement of approximately 15% sucrose with inulin/FOS in papaya-based fruit bars reduced GI from about 65 to 54, while energy density remained within 3260-3350 kcal kg<sup>-1</sup>. Such formulations may produce lower postprandial glucose and insulin responses, support steadier blood glucose availability and promote substrate utilization patterns favourable for endurance performance. Standardizing sugar replacement strategies using ingredients such as inulin, FOS and selected sugar alcohols may support the development of energy bars with improved glycaemic control, sustained energy release and potential benefits for sports nutrition and metabolic health.



**URL:**

[https://scijournals.onlinelibrary.wiley.com/doi/epdf/10.1002/jsfa.70775?getft\\_integrator=clarivate&src=getftr&utm\\_source=clarivate](https://scijournals.onlinelibrary.wiley.com/doi/epdf/10.1002/jsfa.70775?getft_integrator=clarivate&src=getftr&utm_source=clarivate)





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Biochimie

**IF:** 3.4

**Title:** Sedentariness disrupts, while exercise restores, thermogenic and metabolic plasticity in inguinal adipose tissue of mice

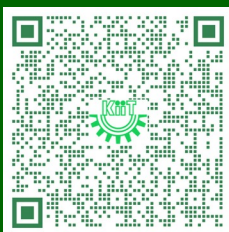
**Author:** Sahu B.; Senapati U.; Pani P.; Pani S.; Swalsingh G.; Pati B.; Tikoo O.; Giri S.; Kar B.P.; Trivedi R.; Bahadi C.K.; Nikhil K.; Kumar D.; Bal N.C.

**Details:** Volume-249, October 2026

**Abstract:** Inguinal WAT (IngWAT) quickly adapts its metabolism to body energy status variations, including uncoupling protein 1 (UCP1) expression. However, its long-term metabolic response to long-term sedentary and exercise training has not been fully characterized. This study investigated IngWAT's long-term response to sedentary (SED, restricted cage) vs. exercise (EXER, treadmill running with a defined protocol) conditions in mice. EXER mice showed increased expression of Sarcoendoplasmic Reticulum Calcium ATPase (SERCA)2b and UCP1, along with a reduction in adipocyte size, indicating activation of thermogenesis. Fascinatingly, exercise elevated the expression of CIDEA in tiny adipocytes and promoted a beiging effect in mice. Elevated OXPHOS complex proteins expression and NMR-based metabolite profiling in IngWAT revealed that exercise profoundly reprogrammed energy metabolism by strengthening phosphocreatine-creatine buffering, regulating TCA cycle metabolite cycling (citrate and succinate), and increasing ketone body formation (3-hydroxybutyrate and acetone). Elevated glycerol level, a higher glycerol-lactate ratio, and reduced carnitine in IngWAT of EXER mice reflected improved fatty acid uptake and  $\beta$ -oxidation, in contrast to the metabolic inefficiency observed during sedentariness. Moreover, increased vascularization was accomplished by upregulated expression of vascular endothelial growth factor 1a (VEGF1a), angiopoietins (ANG)1&2, supports elevated metabolic state in exercise. Conversely, SED mice showed larger adipocytes with uniform CIDEA expression, but a downregulation of thermogenic proteins. Interestingly, a significant downregulation of mitochondrial complex III was also observed, which indicates altered mitochondrial physiology in SED mice. Data presented here suggest that IngWAT is metabolically flexible and can undergo both structural and metabolic adaptations in accordance with physical activity level.



**URL:** <https://www.sciencedirect.com/science/article/abs/pii/S0300908426001513?via%3Dihub>





## SCHOLARLY PUBLICATIONS School of Biotechnology KIIT Deemed to be University

**Journal Name:** Chemical Biology & Drug Design

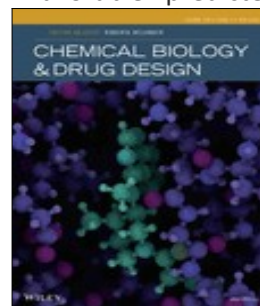
**IF:** 3.3

**Title:** Discovery of Potent Antimalarial Agents Targeting Plasmodium falciparum DNA Gyrase B by Integrating Computational and Experimental Approaches

**Author:** Naik, B; Sahu, W; Sethi, G; Prashar, C; Baldodiya, GM; Poswal, J; Mandal, CC; Hwang, JH; Panday, KC; Reddy, KS; Prusty, D

**Details:** Volume 108, Issue 1, 21<sup>st</sup> July, 2026

**Abstract:** The global spread of drug-resistant Plasmodium falciparum, particularly artemisinin-resistant strains, underscores the urgent need for novel antimalarial agents with distinct mechanisms of action and improved therapeutic potential. In this study, we employed an integrated in silico and in vitro strategy to identify compounds with inhibitory activity against P. falciparum apicoplast Gyrase B (PfGyrB). High-throughput virtual screening identified hit compounds with favorable predicted interactions against the target protein, which were subsequently evaluated using biochemical ATPase inhibition and parasite growth inhibition assays. UNC8153 and Fexofenadine hydrochloride demonstrated time-dependent antiplasmodial activity, with lower IC<sub>50</sub> values at 96 h than at 48 h under prolonged exposure conditions. UNC8153 exhibited greater antiplasmodial activity (~25-fold) than the reference compound novobiocin during the second intraerythrocytic cycle. Morphological analysis indicated impaired parasite development during the second intraerythrocytic cycle under prolonged exposure conditions. Importantly, UNC8153 retained inhibitory activity against the artemisinin-resistant C580Y strain and exhibited minimal cytotoxicity toward HEK-293 cells under the tested experimental conditions. Structural similarity analysis indicated that the identified compounds are chemically distinct from currently used antimalarial drugs, supporting their structural novelty. Collectively, these findings identify UNC8153 as a structurally distinct new antiplasmodial scaffold warranting further mechanistic, pharmacological, and in vivo evaluation.



**URL:** <https://onlinelibrary.wiley.com/doi/full/10.1111/cbdd.70364>

