NAME		Dr. Puniti Mathur		
DESIGNATION			Professor	
EMAIL ID		pmathur@amity.edu	(a)	
CONTACT NUMBER			0120-4392204	
RESEARCH INTERESTS			 Design, synthesis, bioactivity, and conformational studies of non-coded amino acids containing peptides as potential activators of glucokinase-Implications in Type 2 diabetes. Computer-aided drug design: Homology modelling, pharmacophore development, virtual screening, docking and molecular dynamics studies 	
	AL QUALIFICATION ege / University	NS:	Degree	Year
Tvanic of Conege / Oniversity			Degree	1001
All India Institute of Medical Sciences, New Delhi			Ph.D	2003
IIT Roorkee			MSc. Biotechnology	1997
Title of Ph.D. t	<u> </u>	onformati	onal Studies on Bioactive P	eptides containing α , β -
	E (in chronological ord			
Designation	Type of post held (teaching/ research)	Name of	the Institute	Year (From – To)
Research Assistant	Research	International Center for Genetic Engineering and Biotechnology, New Delhi 2003-2004		
Assistant	Industry research	Dabur Pharma Research 2004-2		2004-2008
Manager	T1-1		ion, Sahibabad, Ghaziabad	2000 M 2017
Assistant	Teaching and	Amity Institute of Biotechnology,		2008-May 2017
Professor Associate	research Teaching and		Iniversity, Noida or Computational Biology	June 2017-November
Professor	research	and Bioi of Biotec	nformatics, Amity Institute chnology, Amity	2019
Professor	Teaching and research	University, Noida Centre for Computational Biology and Bioinformatics, Amity Institute of Biotechnology, Amity University, Noida December 2019-till date		
No. OF PhD STUDENTS		Awarded		
				·

Ongoing: 3

SUPERVISED

RESEARCH PROJECTS as PI (completed)	DBT Sponsored Design, synthesis and biological activity of non-coded amino acids containing peptides as potential activators of hepatic glucokinase: Implications in Type 2 diabetes therapy for Rs. 56.41 lakh: 2017-21	
PATENTS (filed/granted)	 A modified peptide as glucokinase activator for treatment of Type 2 diabetes and method thereof, Patent granted (2023) No.462368 Novel peptides containing unnatural amino acids as activators of human glucokinase - Filed on May 31, (1) 2022, Application no. 202211031250 	
Databases created	(1) www.glucokinasedb.in (2) www.pepengine.in	
	1. Das AP, Mathur P , Agarwal SM (2024) Machine Learning, Molecular Docking, and Dynamics-Based Computational Identification of Potential Inhibitors against Lung Cancer <i>ACS Omega</i> 9,4, 4528-4539 DOI: 10.1021/acsomega.3c07338 (IF: 4.1)	
	2. Das AP, Nandekar P, Mathur P , Agarwal SM (2023) A systematic pipeline of protein structure selection for computer-aided drug discovery: A case study on T790M/L858R mutant EGFR structures <i>Protein Science</i> 32(10): https://doi.org/10.1002/pro.4740 (IF: 8.0)	
	3. Yadav S, Bharti S, Mathur P. (2023) GlucoKinaseDB: A comprehensive, curated resource of glucokinase modulators for clinical and molecular research <i>Computational Biology and Chemistry</i> https://doi.org/10.1016/j.compbiolchem.2023.107818 (IF: 3.73)	
PUBLICATIONS	4. Yadav Siddharth, Ahamad, Shahzaib, Gupta, Dinesh, Mathur Puniti (2023) Lead optimization, pharmacophore development and scaffold design of Protein kinase CK2 inhibitors as potential COVID-19 therapeutics <i>Journal of Biomolecular Structure and Dynamics</i> 41(5):1811-1827 doi: 10.1080/0739	
	1102.2021.2024449 (IF: 5.23) 5. Buddham R, Chauhan S, Narad P, Mathur P (2022) Reconstruction and Exploratory Analysis of mTORC1 Signaling Pathway and Its Applications to Various Diseases Using Network-Based Approach. <i>J. Microbiol. Biotechnol.</i> 32 (3):365-377. https://doi.org/10.4014/jmb.2108.08007 (IF: 3.277)	
	6. Siddharth Yadav, Samuel Bharti, Priyansh Srivastava, Puniti Mathur (2022) PepEngine: A manually curated structural database of peptides containing α, β- dehydrophenylalanine (ΔPhe) and α-aminoisobutyric acid (Aib). <i>International Journal of Peptide Research and Therapeuti</i>	
	cs 28, 57 doi: 10.1007/s10989-022-10362-9 (IF: 2.19) 7. Richa Buddham, Siddharth Yadav, Priyanka Narad, Puniti Mathur (2022) Network based identification of Potential Key Genes Associated with Alzheimer disease and Type 2 Diabetes using mTOR signalling. <i>Research Journal of Biotechnology</i> 17	

- (5): 38-50 doi: 10.25303/1705rjbt38050
- 8. Siddharth Yadav, **Puniti Mathur** (2020) Orthosteric and allosteric modulation of human kinases: A mechanistic view. *Frontiers in Bioscience, Landmark*, 25, 8,1462-1487 doi: 10.2741/4864] (IF: 2.34)
- 9. Dua D, Nagoorvali D, Chauhan MS, Palta P, **Mathur P**, Singh MK (2019) Calcium ionophore enhanced developmental competence and apoptotic dynamics of goat parthenogenetic embryos produced in vitro. *In Vitro Cellular & Developmental Biology Animal* Mar;55(3):159-168. doi: 10.1007/s11626-019-00322-x. (IF: 1.447)
- 10. Uzma Khanam, Balwant Kishan Malik, **Puniti Mathur** Bhawna Rathi (2019) Human Caveolin-1 a potent inhibitor for prostate cancer therapy: a computational approach *International Journal of Computational Biology and Drug Design* 12(3): 203-218 (IF:0.7)
- 11. Supriya Srivastava, Seneha Santoshi, Balwant Kishan Malik, **Puniti Mathur** (2017) Molecular modeling and molecular dynamics studies of SPECT protein of Plasmodium falciparum and in silico screening of lead compounds *International Journal of Pharmaceutical Sciences and Research* 8(12):5077-87 https://doi.org/10.13040/IJPSR.0975-8232.8(12).5077-87
- 12. Rajesh Pal, Gauri Misra and **Puniti Mathur** (2017) In Silico Screening Of Small Molecule Modulators Of Zika Virus Proteins *Proceedings of the 7th International Conference Confluence 2017 on Cloud Computing, Data Science and Engineering* 7943179, pp. 381-386 (IEEE) doi: 10.1109/CONFLUENCE.2017.7943179
- 13. Supriya Srivastava, **Puniti Mathur** (2016) Homology Modeling and Docking Studies of Pfmc2TM Maurer's cleft two transmembrane protein –A Potential Drug Target in Malaria. International Journal of Control Theory and Applications 9(3):219-225
- 14. Uzma Khanam, Balwant Kishan Malik, Puniti Mathur, Bhawna Rathi (2016) Identification of novel inhibitors for mitogen-activated protein kinase kinase 4 by virtual screening and molecular dynamics simulation techniques. International Journal of Pharmacy and Pharmaceutical Sciences 8(7): 262-268 (IF:0.55)
- 15. Madhvi Gupta Datta, **Puniti Mathur**, V.S. Chauhan (2011) De novo design, synthesis and solution conformational study of two didehydroundecapeptides: effect of nature and number of amino acids interspersed between ΔPhe residues. Journal of Peptide Science 17(12): 783-790 (IF:1.95)
- 16. Gupta Madhavi, Bagaria Ashima, Mishra Aseem, **Mathur Puniti**, Basu Atanu, Ramakumar S, Chauhan VS (2007) Self-assembly of a dipeptide containing conformationally restricted dehydrophenylalanine residue to form ordered nanotubes. Advanced Materials 19(6): 858-861 (IF: 13.87)
- 17. **Puniti Mathur**, N.R. Jagannathan, V.S. Chauhan (2007) α, β Dehydrophenylalanine containing cecropin-melittin hybrid peptides: conformation and activity. J. Peptide Science 13(4): 253-262. (IF: 1.95)

	 Puniti Mathur, U.A. Ramagopal, S. Ramakumar, N. R. Jagannathan, V.S. Chauhan (2006) Stabilization of unusual structures in peptides using α, β – dehydrophenylalanine: Crystal and solution structures of Boc-Pro-ΔPhe-Val-ΔPheAla-OMe and Boc-Pro-ΔPhe-Gly-ΔPhe-Ala-OMe. Biopolymers (Peptide Science) 84 (3): 298-309 (IF:2.57) Puniti Mathur, S. Ramakumar, V. S. Chauhan (2004) Peptide Design using α, β - dehydroamino acids: From β- turns to helical Hairpins. Biopolymers (Peptide Science) 76(2): 150-161 (IF: 2.57) Padyana A. K., Ramakumar S., Mathur P., Jagannathan N. R., Chauhan V. S.(2003) Role of a two-residue spacer in an α, β-didehydrophenylalanine containing hexapeptide: Crystal and solution structure of Boc-Val-ΔPhe-Leu-Ala- ΔPhe-Ala-OMe. J. Peptide Science 9: 54-63. (IF: 1.95) Ramagopal U. A., Ramakumar S., Mathur P., Joshi, R. M., Chauhan V. S. (2002) Dehydrophenylalanine zippers: strong helix-helix clamping through a network of weak interactions. Protein Engineering 15: 331-335. (IF:3.02) 	
	22. Mathur P., Srivatsun S., Joshi R.M., Jagannathan N. R., Chauhan V. S. (1999) Dehydrophenylalanine containing analogs of Tritrpticin show increased biological activity. <i>J. BioSciences</i> . 24: 42. (IF: 1.88)	
AWARDS & HONOURS/ DISTINCTIONS	 50% Bursary awarded by Wellcome Trust, to attend advanced training on small molecule drug discovery at Wellcome Genome campus, Cambridge, UK (June 2018) Senior Research Fellowship Council of Scientific and Industrial Research (CSIR) Government of India (2001-2003) Qualified GATE (1997) 2 years (1995-1997) Scholarship from Department of Biotechnology Government of India, during MSc. Biotechnology 	
MEMBERSHIP with Professional/ Academic bodies	 Life member of Indian Biophysical Society Life member of Indian Peptide Society Member of American Peptide Society Member of American Chemical Society 	