Chiranjib Chakraborty

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

176 papers

4,367 citations

36 h-index 60 g-index

198 ext. papers

5,710 ext. citations

5.3 avg, IF

6.45 L-index

| # | Paper | IF | Citations |
|-----|---|------------------|-----------|
| 176 | Bioengineering of Novel Non-Replicating mRNA (NRM) and Self-Amplifying mRNA (SAM) Vaccine Candidates Against SARS-CoV-2 Using Immunoinformatics Approach <i>Molecular Biotechnology</i> , 2022 , 1 | 3 | 3 |
| 175 | COMMENTARYOmicron (B.1.1.529) variant of SARS-CoV-2- Concerns, challenges and recent updates <i>Journal of Medical Virology</i> , 2022 , | 19.7 | 26 |
| 174 | A Detailed Overview of Immune Escape, Antibody Escape, Partial Vaccine Escape of SARS-CoV-2 and Their Emerging Variants With Escape Mutations <i>Frontiers in Immunology</i> , 2022 , 13, 801522 | 8.4 | 11 |
| 173 | TN strain proteome mediated therapeutic target mapping and multi-epitopic peptide-based vaccine development for Mycobacterium leprae <i>Infection, Genetics and Evolution</i> , 2022 , 99, 105245 | 4.5 | 2 |
| 172 | A Paradigm Shift in the Combination Changes of SARS-CoV-2 Variants and Increased Spread of Delta Variant (B.1.617.2) across the World 2022 , 13, 927 | | 3 |
| 171 | Evaluation and Designing of Epitopic-Peptide Vaccine Against Using M-Polyprotein Target Sequences. <i>International Journal of Peptide Research and Therapeutics</i> , 2022 , 28, 5 | 2.1 | |
| 170 | Omicron variant (B.1.1.529) of SARS-CoV-2: understanding mutations in the genome, S-glycoprotein, and antibody-binding regions <i>GeroScience</i> , 2022 , | 8.9 | 7 |
| 169 | Hybrid immunity against COVID-19 in different countries with a special emphasis on the Indian scenario during the Omicron period <i>International Immunopharmacology</i> , 2022 , 108, 108766 | 5.8 | 2 |
| 168 | The recombinant variants of SARS-CoV-2: concerns continues amid COVID-19 pandemic <i>Journal of Medical Virology</i> , 2022 , | 19.7 | 4 |
| 167 | Comparative genomics, evolutionary epidemiology, and RBD-hACE2 receptor binding pattern in B.1.1.7 (alpha) and B.1.617.2 (delta) related to their pandemic response in UK and India <i>Infection, Genetics and Evolution</i> , 2022 , 105282 | 4.5 | 1 |
| 166 | Recombinant SARS-CoV-2 variants XD, XE, and XF: The emergence of recombinant variants requires an urgent call for research - Correspondence <i>International Journal of Surgery</i> , 2022 , 106670 | 7.5 | 3 |
| 165 | Emerging cases of acute hepatitis of unknown origin in children amid the ongoing COVID-19 pandemic: Needs attention © Correspondence. <i>International Journal of Surgery</i> , 2022 , 102, 106682 | 7.5 | 0 |
| 164 | Challenges of Long Non Coding RNAs in Human Disease Diagnosis and Therapies: Bio-Computational Approaches. <i>Studies in Big Data</i> , 2022 , 121-131 | 0.9 | |
| 163 | Understanding Gene Expression and Transcriptome Profiling of COVID-19: An Initiative Towards the Mapping of Protective Immunity Genes Against SARS-CoV-2 Infection <i>Frontiers in Immunology</i> , 2021 , 12, 724936 | 8.4 | 3 |
| 162 | A Next-Generation Vaccine Candidate Using Alternative Epitopes to Protect against Wuhan and All Significant Mutant Variants of SARS-CoV-2: An Immunoinformatics Approach 2021 , 12, 2173-2195 | | 8 |
| 161 | D614G mutation and SARS-CoV-2: impact on S-protein structure, function, infectivity, and immunity. <i>Applied Microbiology and Biotechnology</i> , 2021 , 105, 9035-9045 | 5.7 | 10 |
| 160 | The Drug Repurposing for COVID-19 Clinical Trials Provide Very Effective Therapeutic Combinations: Lessons Learned From Major Clinical Studies. <i>Frontiers in Pharmacology</i> , 2021 , 12, 70420 | 5 ^{5.6} | 13 |

(2021-2021)

| 159 | Emerging mutations in the SARS-CoV-2 variants and their role in antibody escape to small molecule-based therapeutic resistance <i>Current Opinion in Pharmacology</i> , 2021 , 62, 64-73 | 5.1 | 7 |
|-----|---|------|----|
| 158 | Understanding the molecular evolution of tiger diversity through DNA barcoding marker ND4 and NADH dehydrogenase complex using computational biology. <i>Genes and Genomics</i> , 2021 , 43, 759-773 | 2.1 | O |
| 157 | SARS-CoV-2 and other human coronaviruses: Mapping of protease recognition sites, antigenic variation of spike protein and their grouping through molecular phylogenetics. <i>Infection, Genetics and Evolution</i> , 2021 , 89, 104729 | 4.5 | 1 |
| 156 | Immunoinformatics Approach for the Identification and Characterization of T Cell and B Cell Epitopes towards the Peptide-Based Vaccine against SARS-CoV-2. <i>Archives of Medical Research</i> , 2021 , 52, 362-370 | 6.6 | 10 |
| 155 | SARS-CoV-2 Brazil variants in Latin America: More serious research urgently needed on public health and vaccine protection. <i>Annals of Medicine and Surgery</i> , 2021 , 66, 102428 | 2 | 12 |
| 154 | Asian-Origin Approved COVID-19 Vaccines and Current Status of COVID-19 Vaccination Program in Asia: A Critical Analysis. <i>Vaccines</i> , 2021 , 9, | 5.3 | 7 |
| 153 | Determination of k-mer density in a DNA sequence and subsequent cluster formation algorithm based on the application of electronic filter. <i>Scientific Reports</i> , 2021 , 11, 13701 | 4.9 | 2 |
| 152 | Lessons Learned from Cutting-Edge Immunoinformatics on Next-Generation COVID-19 Vaccine Research. <i>International Journal of Peptide Research and Therapeutics</i> , 2021 , 27, 1-9 | 2.1 | 3 |
| 151 | Therapeutic advances of miRNAs: A preclinical and clinical update. <i>Journal of Advanced Research</i> , 2021 , 28, 127-138 | 13 | 86 |
| 150 | Response to: Status of Remdesivir: Not Yet Beyond Question!. <i>Archives of Medical Research</i> , 2021 , 52, 104-106 | 6.6 | 5 |
| 149 | A Novel Multi-Epitopic Peptide Vaccine Candidate Against: In-Silico Identification, Design, Cloning and Validation Through Molecular Dynamics. <i>International Journal of Peptide Research and Therapeutics</i> , 2021 , 27, 1-18 | 2.1 | 12 |
| 148 | SARS-CoV-2 protein drug targets landscape: a potential pharmacological insight view for the new drug development. <i>Expert Review of Clinical Pharmacology</i> , 2021 , 14, 225-238 | 3.8 | 10 |
| 147 | CRISPR-Cas9: A Preclinical and Clinical Perspective for the Treatment of Human Diseases. <i>Molecular Therapy</i> , 2021 , 29, 571-586 | 11.7 | 37 |
| 146 | From COVID-19 to Cancer mRNA Vaccines: Moving From Bench to Clinic in the Vaccine Landscape. <i>Frontiers in Immunology</i> , 2021 , 12, 679344 | 8.4 | 23 |
| 145 | Strategies for transdermal drug delivery against bone disorders: A preclinical and clinical update. Journal of Controlled Release, 2021 , 336, 375-395 | 11.7 | 2 |
| 144 | Evolution, Mode of Transmission, and Mutational Landscape of Newly Emerging SARS-CoV-2 Variants. <i>MBio</i> , 2021 , 12, e0114021 | 7.8 | 21 |
| 143 | The current second wave and COVID-19 vaccination status in India. <i>Brain, Behavior, and Immunity</i> , 2021 , 96, 1-4 | 16.6 | 18 |
| 142 | Recent research progress on circular RNAs: Biogenesis, properties, functions, and therapeutic potential. <i>Molecular Therapy - Nucleic Acids</i> , 2021 , 25, 355-371 | 10.7 | 3 |

| 141 | Designing an effective therapeutic siRNA to silence RdRp gene of SARS-CoV-2. <i>Infection, Genetics and Evolution</i> , 2021 , 93, 104951 | 4.5 | 8 |
|-----|---|---------------|------------------|
| 140 | Therapeutics development for Ebola virus disease: A recent scenario. <i>Current Opinion in Pharmacology</i> , 2021 , 60, 208-215 | 5.1 | 2 |
| 139 | All Nations Must Prioritize the COVID-19 Vaccination Program for Elderly Adults Urgently 2021 , 12, 688 | -690 | 6 |
| 138 | Ongoing Clinical Trials of Vaccines to Fight against COVID-19 Pandemic. <i>Immune Network</i> , 2021 , 21, e5 | 6.1 | 12 |
| 137 | COVID-19 vaccine: Challenges in developing countries and India's initiatives. <i>Infezioni in Medicina</i> , 2021 , 29, 165-166 | 3.6 | 3 |
| 136 | Evaluation of molecular interaction, physicochemical parameters and conserved pattern of SARS-CoV-2 Spike RBD and hACE2: in silico and molecular dynamics approach. <i>European Review for Medical and Pharmacological Sciences</i> , 2021 , 25, 1708-1723 | 2.9 | 4 |
| 135 | India's cost-effective COVID-19 vaccine development initiatives. <i>Vaccine</i> , 2020 , 38, 7883-7884 | 4.1 | 21 |
| 134 | Single-cell sequencing of miRNAs: A modified technology. <i>Cell Biology International</i> , 2020 , 44, 1773-178 | 04.5 | 5 |
| 133 | Consider TLR5 for new therapeutic development against COVID-19. <i>Journal of Medical Virology</i> , 2020 , 92, 2314-2315 | 19.7 | 40 |
| 132 | Extensive Partnership, Collaboration, and Teamwork is Required to Stop the COVID-19 Outbreak. <i>Archives of Medical Research</i> , 2020 , 51, 728-730 | 6.6 | 38 |
| 131 | COVID-19: Consider IL-6 receptor antagonist for the therapy of cytokine storm syndrome in SARS-CoV-2 infected patients. <i>Journal of Medical Virology</i> , 2020 , 92, 2260-2262 | 19.7 | 47 |
| 130 | Tocilizumab: A Therapeutic Option for the Treatment of Cytokine Storm Syndrome in COVID-19. <i>Archives of Medical Research</i> , 2020 , 51, 595-597 | 6.6 | 60 |
| 129 | Insight into Evolution and Conservation Patterns of B1-Subfamily Members of GPCR. <i>International Journal of Peptide Research and Therapeutics</i> , 2020 , 26, 1-13 | 2.1 | О |
| 128 | Interaction between miRNAs and signaling cascades of Wnt pathway in chronic lymphocytic leukemia. <i>Journal of Cellular Biochemistry</i> , 2020 , 121, 4654-4666 | 4.7 | 1 |
| 127 | Development of epitope-based peptide vaccine against novel coronavirus 2019 (SARS-COV-2): Immunoinformatics approach. <i>Journal of Medical Virology</i> , 2020 , 92, 618-631 | 19.7 | 237 |
| 126 | Comparative Analysis and Molecular Evolution of Class I PI3K Regulatory Subunit p85Reveal the Structural Similarity Between nSH2 and cSH2 Domains. <i>International Journal of Peptide Research and Therapeutics</i> , 2020 , 26, 2555-2569 | 2.1 | |
| 125 | Identification and Design of a Next-Generation Multi Epitopes Bases Peptide Vaccine Candidate Against Prostate Cancer: An In Silico Approach. <i>Cell Biochemistry and Biophysics</i> , 2020 , 78, 495-509 | 3.2 | 8 |
| 124 | SARS-CoV-2 causing pneumonia-associated respiratory disorder (COVID-19): diagnostic and proposed therapeutic options. <i>European Review for Medical and Pharmacological Sciences</i> , 2020 , 24, 401 | 6-40 <u>2</u> | 6 ¹³⁰ |

(2018-2020)

| 123 | The 2019 novel coronavirus disease (COVID-19) pandemic: A zoonotic prospective. <i>Asian Pacific Journal of Tropical Medicine</i> , 2020 , 13, 242 | 2.1 | 52 |
|-----|---|--------------------|----|
| 122 | Application of Internet Assistance Computation for Disease Prediction and Bio-modeling: Modern Trends in Medical Science. <i>Intelligent Systems Reference Library</i> , 2020 , 327-346 | 0.8 | 1 |
| 121 | Probable Molecular Mechanism of Remdesivir for the Treatment of COVID-19: Need to Know More. <i>Archives of Medical Research</i> , 2020 , 51, 585-586 | 6.6 | 85 |
| 120 | MicroRNAs: Possible Regulatory Molecular Switch Controlling the BBB Microenvironment. <i>Molecular Therapy - Nucleic Acids</i> , 2020 , 19, 933-936 | 10.7 | 3 |
| 119 | Immunoinformatics approach to understand molecular interaction between multi-epitopic regions of SARS-CoV-2 spike-protein with TLR4/MD-2 complex. <i>Infection, Genetics and Evolution</i> , 2020 , 85, 1045 | 8 1 7.5 | 36 |
| 118 | A SARS-CoV-2 vaccine candidate: cloning and validation. <i>Informatics in Medicine Unlocked</i> , 2020 , 20, 100 | 3 9 .4 | 40 |
| 117 | Repurposing Drugs, Ongoing Vaccine, and New Therapeutic Development Initiatives Against COVID-19. <i>Frontiers in Pharmacology</i> , 2020 , 11, 1258 | 5.6 | 61 |
| 116 | Fibroblast-Like-Synoviocytes Mediate Secretion of Pro-Inflammatory Cytokines via ERK and JNK MAPKs in Ti-Particle-Induced Osteolysis. <i>Materials</i> , 2020 , 13, | 3.5 | 4 |
| 115 | The Interplay among miRNAs, Major Cytokines, and Cancer-Related Inflammation. <i>Molecular Therapy - Nucleic Acids</i> , 2020 , 20, 606-620 | 10.7 | 33 |
| 114 | Computer aided novel antigenic epitopes selection from the outer membrane protein sequences of Aeromonas hydrophila and its analyses. <i>Infection, Genetics and Evolution</i> , 2020 , 82, 104320 | 4.5 | 9 |
| 113 | Diabetes and COVID-19: a major challenge in pandemic period?. <i>European Review for Medical and Pharmacological Sciences</i> , 2020 , 24, 11409-11420 | 2.9 | 4 |
| 112 | Advances in nanocarriers enabled brain targeted drug delivery across blood brain barrier. <i>International Journal of Pharmaceutics</i> , 2019 , 559, 360-372 | 6.5 | 83 |
| 111 | Understanding the molecular interaction of human argonaute-2 and miR-20a complex: A molecular dynamics approach. <i>Journal of Cellular Biochemistry</i> , 2019 , 120, 19915-19924 | 4.7 | 6 |
| 110 | Influence of single nucleotide polymorphisms (SNPs) in genetic susceptibility towards periprosthetic osteolysis. <i>Genes and Genomics</i> , 2019 , 41, 1113-1125 | 2.1 | 3 |
| 109 | Ebola virus disease: Recent advancs in diagnostics and therapeutics. <i>Asian Pacific Journal of Tropical Medicine</i> , 2019 , 12, 385 | 2.1 | 3 |
| 108 | Computational and modeling approaches to understand the impact of the Fabry's disease causing mutation (D92Y) on the interaction with pharmacological chaperone 1-deoxygalactonojirimycin (DGJ). <i>Advances in Protein Chemistry and Structural Biology</i> , 2019 , 114, 341-407 | 5.3 | 9 |
| 107 | The novel strategies for next-generation cancer treatment: miRNA combined with chemotherapeutic agents for the treatment of cancer. <i>Oncotarget</i> , 2018 , 9, 10164-10174 | 3.3 | 53 |
| 106 | Anesthetic Molecule Interaction of Noble Gases with Proteins and Lipids and their Effect: A Review. <i>Current Drug Delivery</i> , 2018 , 15, 1381-1392 | 3.2 | 3 |

| 105 | Rising Strengths Hong Kong SAR in Bioinformatics. <i>Interdisciplinary Sciences, Computational Life Sciences</i> , 2017 , 9, 224-236 | 3.5 | 1 |
|-----|---|---------|-----------------|
| 104 | The crucial role and regulations of miRNAs in zebrafish development. <i>Protoplasma</i> , 2017 , 254, 17-31 | 3.4 | 25 |
| 103 | Influence of V54M mutation in giant muscle protein titin: a computational screening and molecular dynamics approach. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017 , 35, 917-928 | 3.6 | 41 |
| 102 | Suppression of osteogenic activity by regulation of WNT and BMP signaling during titanium particle induced osteolysis. <i>Journal of Biomedical Materials Research - Part A</i> , 2017 , 105, 912-926 | 5.4 | 17 |
| 101 | Therapeutic miRNA and siRNA: Moving from Bench to Clinic as Next Generation Medicine. <i>Molecular Therapy - Nucleic Acids</i> , 2017 , 8, 132-143 | 10.7 | 464 |
| 100 | The Smart Programmable CRISPR Technology: A Next Generation Genome Editing Tool for Investigators. <i>Current Drug Targets</i> , 2017 , 18, 1653-1663 | 3 | 7 |
| 99 | The Molecular Concept of Atheromatous Plaques. <i>Current Drug Targets</i> , 2017 , 18, 1250-1258 | 3 | 1 |
| 98 | Review of Prospects of Biological Fluid Biomarkers in Osteoarthritis. <i>International Journal of Molecular Sciences</i> , 2017 , 18, | 6.3 | 68 |
| 97 | Regulatory functional territory of PLK-1 and their substrates beyond mitosis. <i>Oncotarget</i> , 2017 , 8, 3794 | 1233790 | 62 ₇ |
| 96 | Micro-Environmental Signature of The Interactions between Druggable Target Protein, Dipeptidyl Peptidase-IV, and Anti-Diabetic Drugs. <i>Cell Journal</i> , 2017 , 19, 65-83 | 2.4 | 1 |
| 95 | miRNAs in Alzheimer Disease - A Therapeutic Perspective. <i>Current Alzheimer Research</i> , 2017 , 14, 1198-1 | 296 | 51 |
| 94 | Deciphering the impact of somatic mutations in exon 20 and exon 9 of PIK3CA gene in breast tumors among Indian women through molecular dynamics approach. <i>Journal of Biomolecular Structure and Dynamics</i> , 2016 , 34, 29-41 | 3.6 | 25 |
| 93 | India's Computational Biology Growth and Challenges. <i>Interdisciplinary Sciences, Computational Life Sciences</i> , 2016 , 8, 263-76 | 3.5 | 2 |
| 92 | Zebrafish: A complete animal model to enumerate the nanoparticle toxicity. <i>Journal of Nanobiotechnology</i> , 2016 , 14, 65 | 9.4 | 174 |
| 91 | Mechanism of artemisinin resistance for malaria PfATP6 L263 mutations and discovering potential antimalarials: An integrated computational approach. <i>Scientific Reports</i> , 2016 , 6, 30106 | 4.9 | 21 |
| 90 | DNA barcoding to fishes: current status and future directions. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016 , 27, 2744-52 | 1.3 | 21 |
| 89 | Virtual screening of the inhibitors targeting at the viral protein 40 of Ebola virus. <i>Infectious Diseases of Poverty</i> , 2016 , 5, 12 | 10.4 | 36 |
| 88 | Profiling cell-free and circulating miRNA: a clinical diagnostic tool for different cancers. <i>Tumor Biology</i> , 2016 , 37, 5705-14 | 2.9 | 47 |

(2014-2016)

| 87 | PLK-1: Angel or devil for cell cycle progression. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2016 , 1865, 190-203 | 11.2 | 26 |
|----|--|------|-----|
| 86 | Formulation and Application of Biodegradable Nanoparticles Based Biopharmaceutical Delivery - An Efficient Delivery System. <i>Current Pharmaceutical Design</i> , 2016 , 22, 3020-33 | 3.3 | 5 |
| 85 | Therapeutic microRNA Delivery Strategies with Special Emphasis on Cancer Therapy and Tumorigenesis: Current Trends and Future Challenges. <i>Current Drug Metabolism</i> , 2016 , 17, 469-77 | 3.5 | 21 |
| 84 | Dynamics of Diabetes and Obesity: An Alarming Situation in the Developing Countries in Asia. <i>Mini-Reviews in Medicinal Chemistry</i> , 2016 , 16, 1258-1268 | 3.2 | 11 |
| 83 | Application of Bioactive Quercetin in Oncotherapy: From Nutrition to Nanomedicine. <i>Molecules</i> , 2016 , 21, E108 | 4.8 | 90 |
| 82 | miRNA-Regulated Key Components of Cytokine Signaling Pathways and Inflammation in Rheumatoid Arthritis. <i>Medicinal Research Reviews</i> , 2016 , 36, 425-39 | 14.4 | 36 |
| 81 | miRNA-regulated cancer stem cells: understanding the property and the role of miRNA in carcinogenesis. <i>Tumor Biology</i> , 2016 , 37, 13039-13048 | 2.9 | 52 |
| 80 | MicroRNAs mediated regulation of MAPK signaling pathways in chronic myeloid leukemia. <i>Oncotarget</i> , 2016 , 7, 42683-42697 | 3.3 | 45 |
| 79 | Can the chemotherapeutic agents perform anticancer activity through miRNA expression regulation? Proposing a new hypothesis [corrected]. <i>Protoplasma</i> , 2015 , 252, 1603-10 | 3.4 | 5 |
| 78 | India's budget reduction and AIDS initiatives. Lancet Infectious Diseases, The, 2015, 15, 636 | 25.5 | 2 |
| 77 | Exploring the Genomic Roadmap and Molecular Phylogenetics Associated with MODY Cascades Using Computational Biology. <i>Cell Biochemistry and Biophysics</i> , 2015 , 71, 1491-502 | 3.2 | 2 |
| 76 | Methoxy poly(ethylene glycol)-poly(lactide) nanoparticles encapsulating quercetin act as an effective anticancer agent by inducing apoptosis in breast cancer. <i>Pharmaceutical Research</i> , 2015 , 32, 723-35 | 4.5 | 42 |
| 75 | Analysing the Effect of Mutation on Protein Function and Discovering Potential Inhibitors of CDK4: Molecular Modelling and Dynamics Studies. <i>PLoS ONE</i> , 2015 , 10, e0133969 | 3.7 | 36 |
| 74 | DNA pattern recognition using canonical correlation algorithm. <i>Journal of Biosciences</i> , 2015 , 40, 709-19 | 2.3 | 5 |
| 73 | Nanoparticle based insulin delivery system: the next generation efficient therapy for Type 1 diabetes. <i>Journal of Nanobiotechnology</i> , 2015 , 13, 74 | 9.4 | 102 |
| 72 | Profiling of phosphatidylinositol 3-kinase (PI3K) proteins in insulin signaling pathway. <i>Applied Biochemistry and Biotechnology</i> , 2015 , 175, 3431-46 | 3.2 | 3 |
| 71 | Drug Metabolizing Enzymes in Type II Diabetes and their Pharmacogenetics During Therapy of Anti-Diabetes Drugs. <i>Current Drug Metabolism</i> , 2015 , 16, 864-76 | 3.5 | 1 |
| 70 | Structural signature of the G719S-T790M double mutation in the EGFR kinase domain and its response to inhibitors. <i>Scientific Reports</i> , 2014 , 4, 5868 | 4.9 | 32 |

| 69 | Computational approaches and resources in single amino acid substitutions analysis toward clinical research. <i>Advances in Protein Chemistry and Structural Biology</i> , 2014 , 94, 365-423 | 5.3 | 19 |
|----------------------------|--|---------------|--|
| 68 | DNA barcoding to map the microbial communities: current advances and future directions. <i>Applied Microbiology and Biotechnology</i> , 2014 , 98, 3425-36 | 5.7 | 34 |
| 67 | Understanding the conservation patterns and molecular phylogenetics of human death receptors family through computational biology. <i>3 Biotech</i> , 2014 , 4, 177-187 | 2.8 | |
| 66 | Application of evolutionary based in silico methods to predict the impact of single amino acid substitutions in vitelliform macular dystrophy. <i>Advances in Protein Chemistry and Structural Biology</i> , 2014 , 94, 177-267 | 5.3 | 10 |
| 65 | Understanding the molecular dynamics of type-2 diabetes drug target DPP-4 and its interaction with Sitagliptin and inhibitor Diprotin-A. <i>Cell Biochemistry and Biophysics</i> , 2014 , 70, 907-22 | 3.2 | 12 |
| 64 | Influence of miRNA in insulin signaling pathway and insulin resistance: micro-molecules with a major role in type-2 diabetes. <i>Wiley Interdisciplinary Reviews RNA</i> , 2014 , 5, 697-712 | 9.3 | 142 |
| 63 | A novel zebrafish model to provide mechanistic insights into the inflammatory events in carrageenan-induced abdominal edema. <i>PLoS ONE</i> , 2014 , 9, e104414 | 3.7 | 22 |
| 62 | TNF/TNFR: drug target for autoimmune diseases and immune-mediated inflammatory diseases. <i>Frontiers in Bioscience - Landmark</i> , 2014 , 19, 1028-40 | 2.8 | 42 |
| 61 | Novel biomarker for prostate cancer diagnosis by MRS. Frontiers in Bioscience - Landmark, 2014, 19, 118 | 3622801 | 9 |
| | | | |
| 60 | Ebola eradication may need wider partnership. <i>Cmaj</i> , 2014 , 186, 1170 | 3.5 | Ο |
| 60 59 | Ebola eradication may need wider partnership. <i>Cmaj</i> , 2014 , 186, 1170 Effect of Wnt3a on keratinocytes utilizing in vitro and bioinformatics analysis. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 5472-95 | 3·5 6.3 | 0 |
| | Effect of Wnt3a on keratinocytes utilizing in vitro and bioinformatics analysis. <i>International Journal</i> | | |
| 59 | Effect of Wnt3a on keratinocytes utilizing in vitro and bioinformatics analysis. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 5472-95 Integrating in silico prediction methods, molecular docking, and molecular dynamics simulation to predict the impact of ALK missense mutations in structural perspective. <i>BioMed Research</i> | 6.3 | 0 |
| 59 58 | Effect of Wnt3a on keratinocytes utilizing in vitro and bioinformatics analysis. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 5472-95 Integrating in silico prediction methods, molecular docking, and molecular dynamics simulation to predict the impact of ALK missense mutations in structural perspective. <i>BioMed Research International</i> , 2014 , 2014, 895831 Computational biophysical, biochemical, and evolutionary signature of human R-spondin family proteins, the member of canonical Wnt/Etatenin signaling pathway. <i>BioMed Research International</i> , | 6.3 | 0 29 |
| 59 58 57 | Effect of Wnt3a on keratinocytes utilizing in vitro and bioinformatics analysis. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 5472-95 Integrating in silico prediction methods, molecular docking, and molecular dynamics simulation to predict the impact of ALK missense mutations in structural perspective. <i>BioMed Research International</i> , 2014 , 2014, 895831 Computational biophysical, biochemical, and evolutionary signature of human R-spondin family proteins, the member of canonical Wnt/Ecatenin signaling pathway. <i>BioMed Research International</i> , 2014 , 2014, 974316 Next generation delivery system for proteins and genes of therapeutic purpose: why and how?. | 6.3 | o 29 6 |
| 59 58 57 56 | Effect of Wnt3a on keratinocytes utilizing in vitro and bioinformatics analysis. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 5472-95 Integrating in silico prediction methods, molecular docking, and molecular dynamics simulation to predict the impact of ALK missense mutations in structural perspective. <i>BioMed Research International</i> , 2014 , 2014, 895831 Computational biophysical, biochemical, and evolutionary signature of human R-spondin family proteins, the member of canonical Wnt/Etatenin signaling pathway. <i>BioMed Research International</i> , 2014 , 2014, 974316 Next generation delivery system for proteins and genes of therapeutic purpose: why and how?. <i>BioMed Research International</i> , 2014 , 2014, 327950 Evolution- and structure-based computational strategy reveals the impact of deleterious missense | 6.3 | o 29 6 29 |
| 59 58 57 56 55 | Effect of Wnt3a on keratinocytes utilizing in vitro and bioinformatics analysis. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 5472-95 Integrating in silico prediction methods, molecular docking, and molecular dynamics simulation to predict the impact of ALK missense mutations in structural perspective. <i>BioMed Research International</i> , 2014 , 2014, 895831 Computational biophysical, biochemical, and evolutionary signature of human R-spondin family proteins, the member of canonical Wnt/Etatenin signaling pathway. <i>BioMed Research International</i> , 2014 , 2014, 974316 Next generation delivery system for proteins and genes of therapeutic purpose: why and how?. <i>BioMed Research International</i> , 2014 , 2014, 327950 Evolution- and structure-based computational strategy reveals the impact of deleterious missense mutations on MODY 2 (maturity-onset diabetes of the young, type 2). <i>Theranostics</i> , 2014 , 4, 366-85 Recent trends of polymer mediated liposomal gene delivery system. <i>BioMed Research International</i> , | 6.3 3 3 | 02962939 |

(2012-2014)

| 51 | Evaluating protein-protein interaction (PPI) networks for diseases pathway, target discovery, and drug-design using 'in silico pharmacology'. <i>Current Protein and Peptide Science</i> , 2014 , 15, 561-71 | 2.8 | 13 |
|----|--|------|----|
| 50 | Does computational biology help us to understand the molecular phylogenetics and evolution of cluster of differentiation (CD) proteins?. <i>Protein Journal</i> , 2013 , 32, 143-54 | 3.9 | 1 |
| 49 | Predicting the impact of deleterious mutations in the protein kinase domain of FGFR2 in the context of function, structure, and pathogenesisa bioinformatics approach. <i>Applied Biochemistry and Biotechnology</i> , 2013 , 170, 1853-70 | 3.2 | 8 |
| 48 | Mapping the structural topology of IRS family cascades through computational biology. <i>Cell Biochemistry and Biophysics</i> , 2013 , 67, 1319-31 | 3.2 | 2 |
| 47 | Extrapolating the effect of deleterious nsSNPs in the binding adaptability of flavopiridol with CDK7 protein: a molecular dynamics approach. <i>Human Genomics</i> , 2013 , 7, 10 | 6.8 | 40 |
| 46 | In silico discrimination of nsSNPs in hTERT gene by means of local DNA sequence context and regularity. <i>Journal of Molecular Modeling</i> , 2013 , 19, 3517-27 | 2 | 6 |
| 45 | Computational analysis of C-reactive protein for assessment of molecular dynamics and interaction properties. <i>Cell Biochemistry and Biophysics</i> , 2013 , 67, 645-56 | 3.2 | 9 |
| 44 | Topology mapping of insulin-regulated glucose transporter GLUT4 using computational biology. <i>Cell Biochemistry and Biophysics</i> , 2013 , 67, 1261-74 | 3.2 | 5 |
| 43 | Sirtuins familyrecent development as a drug target for aging, metabolism, and age related diseases. <i>Current Drug Targets</i> , 2013 , 14, 666-75 | 3 | 15 |
| 42 | Crucial protein based drug targets and potential inhibitors for osteoporosis: new hope and possibilities. <i>Current Drug Targets</i> , 2013 , 14, 1707-13 | 3 | 8 |
| 41 | miRNAs in insulin resistance and diabetes-associated pancreatic cancer: the 'minute and miracle' molecule moving as a monitor in the 'genomic galaxy'. <i>Current Drug Targets</i> , 2013 , 14, 1110-7 | 3 | 58 |
| 40 | Nanoparticles as 'smart' pharmaceutical delivery. Frontiers in Bioscience - Landmark, 2013 , 18, 1030-50 | 2.8 | 22 |
| 39 | ATP-dependent fructose uptake system in Deinococcus radiodurans. <i>Applied Microbiology and Biotechnology</i> , 2012 , 93, 1241-8 | 5.7 | 4 |
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