Jagdeep Kaur

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Microbial Mannanases: An Overview of Production and Applications. Critical Reviews in Biotechnology, 2007, 27, 197-216.	5.1	361
2	Strategies for optimization of heterologous protein expression in E. coli: Roadblocks and reinforcements. International Journal of Biological Macromolecules, 2018, 106, 803-822.	3.6	245
3	Structure-Activity Determinants in Antifungal Plant Defensins MsDef1 and MtDef4 with Different Modes of Action against Fusarium graminearum. PLoS ONE, 2011, 6, e18550.	1.1	159
4	Stimulatory effect of phosphate-solubilizing fungal strains (Aspergillus awamori and Penicillium) Tj ETQq0 0 0 rgB 718-727.	T /Overloc 4.2	k 10 Tf 50 6 141
5	Structural and Functional Studies of a Phosphatidic Acid-Binding Antifungal Plant Defensin MtDef4: Identification of an RGFRRR Motif Governing Fungal Cell Entry. PLoS ONE, 2013, 8, e82485.	1.1	120
6	Silencing of <i>ABCC13</i> transporter in wheat reveals its involvement in grain development, phytic acid accumulation and lateral root formation. Journal of Experimental Botany, 2016, 67, 4379-4389.	2.4	100
7	An essential oil and its major constituent isointermedeol induce apoptosis by increased expression of mitochondrial cytochrome c and apical death receptors in human leukaemia HL-60 cells. Chemico-Biological Interactions, 2008, 171, 332-347.	1.7	95
8	Purification, characterization and thermostability of lipase from a thermophilic Bacillus sp. J33. , 2000, 206, 91-96.		93
9	Biodegradation of malathion by Brevibacillus sp. strain KB2 and Bacillus cereus strain PU. World Journal of Microbiology and Biotechnology, 2012, 28, 1133-1141.	1.7	79
10	Immobilization, stability and esterification studies of a lipase from a Bacillus sp Biotechnology and Applied Biochemistry, 2002, 36, 7.	1.4	77
11	Peanuts that keep aflatoxin at bay: a threshold that matters. Plant Biotechnology Journal, 2018, 16, 1024-1033.	4.1	71
12	Lipid hydrolizing enzymes in virulence: <i>Mycobacterium tuberculosis</i> as a model system. Critical Reviews in Microbiology, 2010, 36, 259-269.	2.7	70
13	Differential expression of structural genes for the late phase of phytic acid biosynthesis in developing seeds of wheat (Triticum aestivum L.). Plant Science, 2014, 224, 74-85.	1.7	68
14	De Novo Transcriptome Sequencing Reveals Important Molecular Networks and Metabolic Pathways of the Plant, Chlorophytum borivilianum. PLoS ONE, 2013, 8, e83336.	1.1	65
15	Title is missing!. Biotechnology Letters, 1998, 20, 997-1000.	1.1	65
16	An improved method for single step purification of metagenomic DNA. Molecular Biotechnology, 2007, 36, 61-63.	1.3	60
17	Sphingolipid C-9 Methyltransferases Are Important for Growth and Virulence but Not for Sensitivity to Antifungal Plant Defensins in <i>Fusarium graminearum </i> . Eukaryotic Cell, 2009, 8, 217-229.	3.4	59
18	Microbial degradation of an organophosphate pesticide, malathion. Critical Reviews in Microbiology, 2014, 40, 146-154	2.7	53

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19	Genome-wide identification and expression characterization of ABCC-MRP transporters in hexaploid wheat. Frontiers in Plant Science, 2015, 6, 488.	1.7	50
20	Microbial remediation of explosive waste. Critical Reviews in Microbiology, 2012, 38, 152-167.	2.7	49
21	Can plant defensins be used to engineer durable commercially useful fungal resistance in crop plants?. Fungal Biology Reviews, 2011, 25, 128-135.	1.9	46
22	Specific domains of plant defensins differentially disrupt colony initiation, cell fusion and calcium homeostasis in <scp><i>N</i></scp> <i>eurospora crassa</i> . Molecular Microbiology, 2014, 92, 1357-1374.	1.2	46
23	Engineering of Bacillus lipase by directed evolution for enhanced thermal stability: effect of isoleucine to threonine mutation at protein surface. Molecular Biology Reports, 2011, 38, 2919-2926.	1.0	45
24	Differential Expression of RDC1/CXCR7 in the Human Placenta. Journal of Clinical Immunology, 2009, 29, 379-386.	2.0	44
25	Lipoxygenase in Caragana jubata responds to low temperature, abscisic acid, methyl jasmonate and salicylic acid. Gene, 2011, 483, 49-53.	1.0	42
26	Transgenic maize plants expressing the Totivirus antifungal protein, KP4, are highly resistant to corn smut. Plant Biotechnology Journal, 2011, 9, 857-864.	4.1	40
27	Immobilization and stability studies of a lipase from thermophilic Bacillus sp: The effect of process parameters on immobilization of enzyme. Electronic Journal of Biotechnology, 2006, 9, 0-0.	1.2	39
28	Engineering of a metagenome derived lipase toward thermal tolerance: Effect of asparagine to lysine mutation on the protein surface. Gene, 2012, 491, 264-271.	1.0	39
29	Cloning, expression and characterization of a metagenome derived thermoactive/thermostable pectinase. Molecular Biology Reports, 2012, 39, 8353-8361.	1.0	39
30	p16INK4a and p15INK4b gene promoter methylation in cervical cancer patients. Oncology Letters, 2012, 3, 1331-1335.	0.8	38
31	Promoter hypermethylation of p73 and p53 genes in cervical cancer patients among north Indian population. Molecular Biology Reports, 2012, 39, 9145-9157.	1.0	38
32	Characterization of LipN (Rv2970c) of <i>Mycobacterium Tuberculosis</i> H37Rv and its Probable Role in Xenobiotic Degradation. Journal of Cellular Biochemistry, 2016, 117, 390-401.	1.2	38
33	Expression of apoplast-targeted plant defensin MtDef4.2 confers resistance to leaf rust pathogen Puccinia triticina but does not affect mycorrhizal symbiosis in transgenic wheat. Transgenic Research, 2017, 26, 37-49.	1.3	38
34	Low-pH-induced apoptosis: role of endoplasmic reticulum stress-induced calcium permeability and mitochondria-dependent signaling. Cell Stress and Chaperones, 2015, 20, 431-440.	1.2	37
35	Studies on lipolytic isoenzymes from a thermophilic Bacillus sp.: Production, purification and biochemical characterization. Enzyme and Microbial Technology, 2007, 40, 881-887.	1.6	35
36	Role of Oxidative Stress and Apoptosis in the Placental Pathology of Plasmodium berghei Infected Mice. PLoS ONE, 2012, 7, e32694.	1.1	35

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37	Molecular principles behind pyrazinamide resistance due to mutations in panD gene in Mycobacterium tuberculosis. Gene, 2016, 581, 31-42.	1.0	34
38	A thermostable lipolytic enzyme from a thermophilic Bacillus sp.: Purification and characterization. Molecular and Cellular Biochemistry, 2006, 290, 17-22.	1.4	32
39	Characterization of a thermostable lipase showing loss of secondary structure at ambient temperature. Molecular Biology Reports, 2012, 39, 2795-2804.	1.0	32
40	De Novo Transcriptome Sequencing and Analysis for Venturia inaequalis, the Devastating Apple Scab Pathogen. PLoS ONE, 2013, 8, e53937.	1.1	32
41	Production of Biodiesel From Used Mustard Oil and Its Performance Analysis in Internal Combustion Engine. Journal of Energy Resources Technology, Transactions of the ASME, 2010, 132, .	1.4	31
42	Characterization of Squalene synthase Gene from Chlorophytum borivilianum (Sant. and Fernand.). Molecular Biotechnology, 2013, 54, 944-953.	1.3	31
43	Subcellular targeting of an evolutionarily conserved plant defensin <scp>M</scp> t <scp>D</scp> ef4.2 determines the outcome of plant–pathogen interaction in transgenic <scp>A</scp> rabidopsis. Molecular Plant Pathology, 2012, 13, 1032-1046.	2.0	29
44	Characterization of a novel esterase Rv1497 of Mycobacterium tuberculosis H37Rv demonstrating β-lactamase activity. Enzyme and Microbial Technology, 2016, 82, 180-190.	1.6	29
45	Characterization of an acid inducible lipase Rv3203 from Mycobacterium tuberculosis H37Rv. Molecular Biology Reports, 2014, 41, 285-296.	1.0	28
46	Infergen Stimulated Macrophages Restrict Mycobacterium tuberculosis Growth by Autophagy and Release of Nitric Oxide. Scientific Reports, 2016, 6, 39492.	1.6	28
47	Modulation of Trehalose Dimycolate and Immune System by Rv0774c Protein Enhanced the Intracellular Survival of Mycobacterium smegmatis in Human Macrophages Cell Line. Frontiers in Cellular and Infection Microbiology, 2017, 7, 289.	1.8	28
48	Molecular Characterization of Oxidative Stress-Inducible LipD of Mycobacterium tuberculosis H37Rv. Current Microbiology, 2014, 68, 387-396.	1.0	27
49	Development of genic SSR marker resource from RNA-Seq data in Dendrocalamus latiflorus. Journal of Plant Biochemistry and Biotechnology, 2016, 25, 179-190.	0.9	27
50	Characterization and molecular modelling of an engineered organic solvent tolerant, thermostable lipase with enhanced enzyme activity. Journal of Molecular Catalysis B: Enzymatic, 2013, 97, 243-251.	1.8	26
51	Functional characterization of hypothetical proteins of <i>Mycobacterium tuberculosis</i> with possible esterase/lipase signature: a cumulative <i>in silico</i> and <i>in vitro</i> approach. Journal of Biomolecular Structure and Dynamics, 2017, 35, 1226-1243.	2.0	26
52	De novo Transcriptome Analysis Revealed Genes Involved in Flavonoid and Vitamin C Biosynthesis in Phyllanthus emblica (L.). Frontiers in Plant Science, 2016, 7, 1610.	1.7	24
53	Multifaceted role of lipids inMycobacterium leprae. Future Microbiology, 2017, 12, 315-335.	1.0	24
54	Biochemical Analysis of a Native and Proteolytic Fragment of a High-Molecular-Weight Thermostable Lipase from a Mesophilic Bacillus sp Protein Expression and Purification, 2002, 24, 71-75.	0.6	23

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55	Transformation of malathion by Lysinibacillus sp. isolated from soil. Biotechnology Letters, 2012, 34, 863-867.	1.1	23
56	Primer Based Approach for PCR Amplification of High GC Content Gene: <i>Mycobacterium</i> Gene as a Model. Molecular Biology International, 2014, 2014, 1-7.	1.7	23
57	Dynamics of fluoroquinolones induced resistance in DNA gyrase of <i>Mycobacterium tuberculosis</i> . Journal of Biomolecular Structure and Dynamics, 2018, 36, 362-375.	2.0	23
58	A thermostable glucoamylase from a thermophilic Bacillus sp.: characterization and thermostability. Journal of Industrial Microbiology and Biotechnology, 2004, 31, 540-543.	1.4	22
59	RNA-Seq mediated root transcriptome analysis of Chlorophytum borivilianum for identification of genes involved in saponin biosynthesis. Functional and Integrative Genomics, 2016, 16, 37-55.	1.4	22
60	mesT, a unique epoxide hydrolase, is essential for optimal growth of <i>Mycobacterium tuberculosis</i> in the presence of styrene oxide. Future Microbiology, 2017, 12, 527-546.	1.0	22
61	Morbid Sequences Suggest Molecular Mimicry between Microbial Peptides and Self-Antigens: A Possibility of Inciting Autoimmunity. Frontiers in Microbiology, 2017, 8, 1938.	1.5	22
62	Engineering lipases for temperature adaptation: Structure function correlation. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2019, 1867, 140261.	1.1	22
63	Discovery and Utilization of EST-SSR Marker Resource for Genetic Diversity and Population Structure Analyses of a Subtropical Bamboo, Dendrocalamus hamiltonii. Biochemical Genetics, 2019, 57, 652-672.	0.8	22
64	Rv0518, a nutritive stress inducible GDSL lipase of Mycobacterium tuberculosis, enhanced intracellular survival of bacteria by cell wall modulation. International Journal of Biological Macromolecules, 2019, 135, 180-195.	3.6	21
65	Enantiomeric separation of pharmaceutically important drug intermediates using a Metagenomic lipase and optimization of its large scale production. International Journal of Biological Macromolecules, 2017, 95, 995-1003.	3.6	20
66	Characterization of ML0314c of Mycobacterium leprae and deciphering its role in the immune response in leprosy patients. Gene, 2018, 643, 26-34.	1.0	20
67	Characterization and evolution of a metagenome-derived lipase towards enhanced enzyme activity and thermostability. Molecular and Cellular Biochemistry, 2013, 373, 149-159.	1.4	19
68	Rv0774c, an iron stress inducible, extracellular esterase is involved in immune-suppression associated with altered cytokine and TLR2 expression. International Journal of Medical Microbiology, 2017, 307, 126-138.	1.5	19
69	Elucidating genes involved in sesquiterpenoid and flavonoid biosynthetic pathways in Saussurea lappa by de novo leaf transcriptome analysis. Genomics, 2019, 111, 1474-1482.	1.3	19
70	smRNAome profiling to identify conserved and novel microRNAs in Stevia rebaudiana Bertoni. BMC Plant Biology, 2012, 12, 197.	1.6	18
71	Combinatorial reshaping of a lipase structure for thermostability: Additive role of surface stabilizing single point mutations. Biochemical and Biophysical Research Communications, 2014, 447, 626-632.	1.0	18
72	Alanine mutation of the catalytic sites of Pantothenate Synthetase causes distinct conformational changes in the ATP binding region. Scientific Reports, 2018, 8, 903.	1.6	18

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73	Rv2037c, a stress induced conserved hypothetical protein of Mycobacterium tuberculosis, is a phospholipase: Role in cell wall modulation and intracellular survival. International Journal of Biological Macromolecules, 2020, 153, 817-835.	3.6	18
74	Differentially expressed transcripts from leaf and root tissue of Chlorophytum borivilianum: A plant with high medicinal value. Gene, 2012, 511, 79-87.	1.0	17
75	Chemopreventive activity of lantadenes on two-stage carcinogenesis model in Swiss albino mice: AP-1 (c-jun), NFI°B (p65) and P53 expression by ELISA and immunohistochemical localization. Molecular and Cellular Biochemistry, 2008, 314, 1-8.	1.4	16
76	A novel parthenin analog exhibits anti-cancer activity: Activation of apoptotic signaling events through robust NO formation in human leukemia HL-60 cells. Chemico-Biological Interactions, 2011, 193, 204-215.	1.7	16
77	Rv1288, a Two Domain, Cell Wall Anchored, Nutrient Stress Inducible Carboxyl-Esterase of Mycobacterium tuberculosis, Modulates Cell Wall Lipid. Frontiers in Cellular and Infection Microbiology, 2018, 8, 421.	1.8	16
78	Analysis of mutations leading to para-aminosalicylic acid resistance in Mycobacterium tuberculosis. Scientific Reports, 2019, 9, 13617.	1.6	16
79	Insights into controlling role of substitution mutation, E315G on thermostability of a lipase cloned from metagenome of hot spring soil. 3 Biotech, 2014, 4, 189-196.	1.1	15
80	A βâ€mannanase from <i>Paenibacillus</i> sp.: Optimization of production and its possible prebiotic potential. Biotechnology and Applied Biochemistry, 2016, 63, 669-678.	1.4	15
81	Point Mutation Ile137-Met Near Surface Conferred Psychrophilic Behaviour and Improved Catalytic Efficiency to Bacillus Lipase of 1.4 Subfamily. Applied Biochemistry and Biotechnology, 2016, 178, 753-765.	1.4	15
82	Cell Wall Associated Factors of Mycobacterium tuberculosis as Major Virulence Determinants: Current Perspectives in Drugs Discovery and Design. Current Drug Targets, 2017, 18, 1904-1918.	1.0	15
83	Structural and functional insights into thermostable and organic solvent stable variant Pro247-Ser of Bacillus lipase. International Journal of Biological Macromolecules, 2018, 108, 845-852.	3.6	15
84	Streptomycin sulphate loaded solid lipid nanoparticles show enhanced uptake in macrophage, lower MIC in Mycobacterium and improved oral bioavailability. European Journal of Pharmaceutics and Biopharmaceutics, 2021, 160, 100-124.	2.0	15
85	Plasmodium berghei: Influence of infection on the oxidant and antioxidants levels in pregnant BALB/c mice. Experimental Parasitology, 2012, 131, 215-222.	0.5	14
86	CXCL12–CXCR7 Signaling Activates ERK and Akt Pathways in Human Choriocarcinoma Cells. Cell Communication and Adhesion, 2014, 21, 221-228.	1.0	14
87	Bioremediation of malathion in soil by mixed Bacillus culture. Advances in Bioscience and Biotechnology (Print), 2013, 04, 674-678.	0.3	14
88	Role of N-Terminal Domain of Streptokinase in Protein Transport. Biochemical and Biophysical Research Communications, 1996, 227, 303-310.	1.0	13
89	Point mutation Gln121-Arg increased temperature optima of Bacillus lipase (1.4 subfamily) by fifteen degrees. International Journal of Biological Macromolecules, 2016, 88, 507-514.	3.6	13
90	Characterization of an extracellular protein, Rv1076 from M. tuberculosis with a potential role in humoral response. International Journal of Biological Macromolecules, 2017, 101, 621-629.	3.6	13

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91	2,4,6-Trinitrophenol degradation by Bacillus cereus isolated from a firing range. Biotechnology Letters, 2011, 33, 2411-2415.	1.1	12
92	The virally encoded killer proteins from Ustilago maydis. Fungal Biology Reviews, 2013, 26, 166-173.	1.9	12
93	Thirty-degree shift in optimum temperature of a thermophilic lipase by a single-point mutation: effect of serine to threonine mutation on structural flexibility. Molecular and Cellular Biochemistry, 2017, 430, 21-30.	1.4	12
94	Drug targeted virtual screening and molecular dynamics of LipU protein of <i>Mycobacterium tuberculosis</i> and <i>Mycobacterium leprae</i> . Journal of Biomolecular Structure and Dynamics, 2019, 37, 1254-1269.	2.0	12
95	TLR Agonist Augments Prophylactic Potential of Acid Inducible Antigen Rv3203 against Mycobacterium tuberculosis H37Rv in Experimental Animals. PLoS ONE, 2016, 11, e0152240.	1.1	12
96	Disruption of N terminus long range non covalent interactions shifted temp.opt 25°C to cold: Evolution of point mutant Bacillus lipase by error prone PCR. Gene, 2016, 576, 237-243.	1.0	11
97	Impact of novel N-aryl substituted piperamide on NF-kappa B translocation as a potent anti-neuroinflammatory agent. Biomedicine and Pharmacotherapy, 2020, 127, 110199.	2.5	11
98	Inhibition of NOTCH signaling pathway chemosensitizes HCC CD133+ cells to vincristine and 5-fluorouracil through upregulation of BBC3. Biochemical and Biophysical Research Communications, 2020, 525, 941-947.	1.0	11
99	Point mutation Arg153-His at surface of Bacillus lipase contributing towards increased thermostability and ester synthesis: insight into molecular network. Molecular and Cellular Biochemistry, 2018, 443, 159-168.	1.4	10
100	The immunosuppressive effects of a novel recombinant LipQ (Rv2485c) protein of Mycobacterium tuberculosis on human macrophage cell lines. Microbial Pathogenesis, 2017, 107, 361-367.	1.3	9
101	Gene expression analysis for selection and validation of suitable housekeeping gene(s) in cadmium exposed pigeonpea plants inoculated with arbuscular mycorrhizae. Plant Physiology and Biochemistry, 2021, 162, 592-602.	2.8	9
102	Antifungal Plant Defensins: Structure-Activity Relationships, Modes of Action, and Biotech Applications. ACS Symposium Series, 2012, , 317-336.	0.5	8
103	Rv0646c, an esterase from M. tuberculosis, up-regulates the host immune response in THP-1 macrophages cells. Molecular and Cellular Biochemistry, 2018, 447, 189-202.	1.4	8
104	Impact of novel <i>N</i> -aryl piperamide NO donors on NF-κB translocation in neuroinflammation: rational drug-designing synthesis and biological evaluation. Innate Immunity, 2018, 24, 24-39.	1.1	8
105	A comparative analysis of methylation status of tumor suppressor genes in paired biopsy and serum samples from cervical cancer patients among north indian population. Russian Journal of Genetics, 2016, 52, 226-230.	0.2	6
106	Double Mutants in DNA Gyrase Lead to Ofloxacin Resistance in <i>Mycobacterium tuberculosis</i> . Journal of Cellular Biochemistry, 2017, 118, 2950-2957.	1.2	6
107	Multidomain truncated hemoglobins: New members of the globin family exhibiting tandem repeats of globin units and domain fusion. IUBMB Life, 2017, 69, 479-488.	1.5	6
108	mbtJ: an iron stress-induced acetyl hydrolase/esterase of <i>Mycobacterium tuberculosis</i> helps bacteria to survive during iron stress. Future Microbiology, 2018, 13, 547-564.	1.0	6

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109	Rv2223c, an acid inducible carboxyl-esterase of <i>Mycobacterium tuberculosis</i> enhanced the growth and survival of <i>Mycobacterium smegmatis</i> . Future Microbiology, 2019, 14, 1397-1415.	1.0	6
110	Novel missense mutations in gidB gene associated with streptomycin resistance in <i>Mycobacterium tuberculosis:</i> insights from molecular dynamics. Journal of Biomolecular Structure and Dynamics, 2019, 37, 20-35.	2.0	6
111	Molecular Dynamics Assisted Mechanistic Insight of Val430-Ala Mutation of Rv1592c Protein in Isoniazid Resistant Mycobacterium Tuberculosis. Current Computer-Aided Drug Design, 2021, 17, 95-106.	0.8	6
112	Risk of Late-Onset Alzheimer's Disease by Plasma Cholesterol: Rational <i>In Silico</i> Drug Investigation of Pyrrole-Based HMG-CoA Reductase Inhibitors. Assay and Drug Development Technologies, 2017, 15, 342-351.	0.6	5
113	The critical role of piperamide derivative D4 in the regulation of inflammatory response by the microglia and astrocytic glial cells. Biomedicine and Pharmacotherapy, 2020, 132, 110895.	2.5	5
114	The lipolytic activity of LipJ, a stress-induced enzyme, is regulated by its C-terminal adenylate cyclase domain. Future Microbiology, 2021, 16, 487-507.	1.0	5
115	Expression of heat shock protein 90, 70, 60 and 25 in the placenta of Plasmodium berghei infected BALB/c mice. Asian Pacific Journal of Tropical Disease, 2014, 4, S442-S444.	0.5	4
116	New Insight into Old <i>Bacillus</i> Lipase: Solvent Stable Mesophilic Lipase Demonstrating Enzyme Activity towards Cold. Journal of Molecular Microbiology and Biotechnology, 2015, 25, 340-348.	1.0	4
117	Gauging the trends of pseudogenes in plants. Critical Reviews in Biotechnology, 2021, 41, 1114-1129.	5.1	4
118	Mutation in Eth A protein of Mycobacterium tuberculosis conferred drug tolerance against enthinoamide in Mycobacterium smegmatis mc2155. Computational Biology and Chemistry, 2022, 98, 107677.	1.1	4
119	Genomic Insights into Omega-3 Polyunsaturated Fatty Acid Producing Shewanella sp. N2AIL from Fish Gut. Biology, 2022, 11, 632.	1.3	4
120	Cloning and Characterization of Promoter-Active DNA Sequences from Streptococcus equisimilis. Current Microbiology, 2007, 54, 48-53.	1.0	3
121	Differential expression of two members of Rv1922-LipD operon in Mycobacterium tuberculosis: Does rv1923 qualify for membership?. Pathogens and Disease, 2015, 73, .	0.8	3
122	Conserved cysteine variants of metagenomic derived polygalacturonase concurrently shift its optima at acidic pH and enhanced thermostability: structural and functional analysis. Journal of Biomolecular Structure and Dynamics, 2019, 37, 265-273.	2.0	3
123	Integration of VEKâ€30 peptide enhances fibrinolytic properties of staphylokinase. Biotechnology and Applied Biochemistry, 2021, 68, 213-220.	1.4	3
124	Correlation of over-expression of rv1900c with enhanced survival of M. smegmatis under stress conditions: Modulation of cell surface properties. Gene, 2021, 791, 145720.	1.0	3
125	Degradation of TNP, RDX, and CL-20 Explosives by Microbes. Environmental Science and Engineering, 2014, , 87-111.	0.1	3
126	In-Silico Characterization of a Hypothetical Protein, Rv1288 of Mycobacterium tuberculosis Containing an Esterase Signature and an Uncommon LytE Domain. Current Computer-Aided Drug Design, 2017, 13, 101-111.	0.8	3

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127	Comparative analysis of point mutations on protein COOH terminal near surface and its hydrophobic core provide insights on thermostability of Bacillus Lipase LipJ. Journal of Molecular Catalysis B: Enzymatic, 2016, 133, S482-S490.	1.8	2
128	Intrinsically Unstructured Carboxy Terminus of Bacillus Lipase is Essential for its Function. Protein and Peptide Letters, 2014, 21, 1265-1272.	0.4	2
129	Intrinsically unstructured carboxy terminus of Bacillus lipase is essential for its function. Protein and Peptide Letters, 2014, 21, 1265-72.	0.4	2
130	Environment dependent expression of mycobacterium hormone sensitive lipases: expression pattern under ex-vivo and individual in-vitro stress conditions in M. tuberculosis H37Ra. Molecular Biology Reports, 2022, 49, 4583-4593.	1.0	2
131	Antifungal Plant Defensins: Insights into Modes of Action and Prospects for Engineering Disease-Resistant Plants. , 2018, , 129-140.		1
132	Biomarkers of Cardiac Health and Disease. Critical Reviews in Biomedical Engineering, 2019, 47, 395-407.	0.5	1
133	Molecular characterization and immunogenic function of ML1899 (LipG) of Mycobacterium leprae. Journal of Medical Microbiology, 2019, 68, 1629-1640.	0.7	1
134	A Phagosomally Expressed Gene, rv0428c, of Mycobacterium tuberculosis Demonstrates Acetyl Transferase Activity and Plays a Protective Role Under Stress Conditions. Protein Journal, 2022, 41, 260-273.	0.7	1
135	Studies on Recombinant Lipase Production byÂE. Coli: Effect of Media And Bacterial Expression System	0.2	0