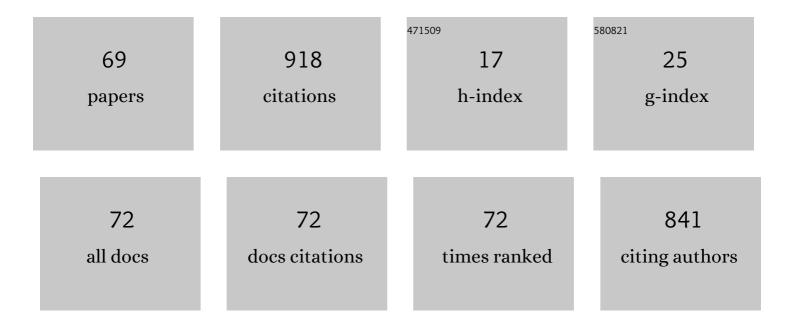
## Bharat Bhusan Patnaik

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Current knowledge of immune priming in invertebrates, emphasizing studies on Tenebrio molitor. Developmental and Comparative Immunology, 2022, 127, 104284.	2.3	11
2	Molecular cloning, sequence characterization, and expression analysis of C-type lectin (CTL) and ER-Golgi intermediate compartment 53-kDa protein (ERGIC-53) homologs from the freshwater prawn, Macrobrachium rosenbergii. Aquaculture International, 2022, 30, 1011-1035.	2.2	2
3	Characterization of <scp><i>Haemaphysalis longicornis</i></scp> microbiome collected from different regions of Korean peninsula. Entomological Research, 2022, 52, 271-280.	1.1	1
4	Gene expression analysis of inflammation-related genes in macrophages treated with α-(1Â→Â3, 1Â→Â6)-D-gluc extracted from Streptococcus mutans. International Journal of Biological Macromolecules, 2021, 166, 45-53.	can 7.5	5
5	Transcriptome analysis of <i>Macrobrachium rosenbergii</i> hepatopancreas in response to <i>Vibrio harveyi</i> infection. Aquaculture Research, 2021, 52, 1855-1875.	1.8	3
6	Deep sequencing and phylogenetic analysis of severe fever with thrombocytopenia syndrome virus from the tick, Haemaphysalis longicornis , in Korea. Entomological Research, 2021, 51, 3-11.	1.1	1
7	Identification, <i>in silico</i> characterization, and expression analysis of <scp><i>Tenebrio molitor</i></scp> Cecropinâ€2. Entomological Research, 2021, 51, 74-82.	1.1	11
8	Autophagy in Tenebrio molitor Immunity: Conserved Antimicrobial Functions in Insect Defenses. Frontiers in Immunology, 2021, 12, 667664.	4.8	16
9	TmSpz-like Plays a Fundamental Role in Response to E. coli but Not S. aureus or C. albican Infection in Tenebrio molitor via Regulation of Antimicrobial Peptide Production. International Journal of Molecular Sciences, 2021, 22, 10888.	4.1	16
10	Tenebrio molitor SpÃæle 1b Is Required to Confer Antibacterial Defense Against Gram-Negative Bacteria by Regulation of Antimicrobial Peptides. Frontiers in Physiology, 2021, 12, 758859.	2.8	9
11	TmIKKε Is Required to Confer Protection Against Gram-Negative Bacteria, E. coli by the Regulation of Antimicrobial Peptide Production in the Tenebrio molitor Fat Body. Frontiers in Physiology, 2021, 12, 758862.	2.8	8
12	Transcriptome studies of the floodwater mosquito, <scp><i>Aedes vexans</i></scp> (Diptera:) Tj ETQq0 0 0 rgBT Research, 2020, 50, 563-574.	/Overlock 1.1	10 Tf 50 30 5
13	Tickâ€borne viruses: Current trends in largeâ€scale viral surveillance. Entomological Research, 2020, 50, 379-392.	1.1	3
14	Bacterial but not fungal challenge upâ€regulates the transcription of <i>Coleoptericin</i> genes in <scp><i>Tenebrio molitor</i></scp> . Entomological Research, 2020, 50, 440-449.	1.1	14
15	IKKÎ <sup>3</sup> /NEMO Is Required to Confer Antimicrobial Innate Immune Responses in the Yellow Mealworm, Tenebrio Molitor. International Journal of Molecular Sciences, 2020, 21, 6734.	4.1	12
16	<i>In silico</i> identification and expression analyses of <i>Defensin</i> genes in the mealworm beetle <scp><i>Tenebrio molitor</i></scp> . Entomological Research, 2020, 50, 575-585.	1.1	12
17	TmSpz4 Plays an Important Role in Regulating the Production of Antimicrobial Peptides in Response to Escherichia coli and Candida albicans Infections. International Journal of Molecular Sciences, 2020, 21, 1878.	4.1	19
18	TmRelish is required for regulating the antimicrobial responses to Escherichia coli and Staphylococcus aureus in Tenebrio molitor. Scientific Reports, 2020, 10, 4258.	3.3	25

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19	TmSpz6 Is Essential for Regulating the Immune Response to Escherichia coli and Staphylococcus aureus Infection in Tenebrio molitor. Insects, 2020, 11, 105.	2.2	24
20	Aedes albopictus Autophagy-Related Gene 8 (AaAtg8) Is Required to Confer Anti-Bacterial Gut Immunity. International Journal of Molecular Sciences, 2020, 21, 2944.	4.1	8
21	Lectin-Like Activity of Hemocyanin in Freshwater Prawn, Macrobrachium rosenbergii. Protein Journal, 2020, 39, 358-365.	1.6	8
22	Regulation of the expression of nine antimicrobial peptide genes by TmIMD confers resistance against Gram-negative bacteria. Scientific Reports, 2019, 9, 10138.	3.3	28
23	Molecular cloning and characterization of SOCS2 from the mealworm beetle Tenebrio molitor. Entomological Research, 2019, 49, 313-322.	1.1	1
24	TmDorX2 positively regulates antimicrobial peptides in Tenebrio molitor gut, fat body, and hemocytes in response to bacterial and fungal infection. Scientific Reports, 2019, 9, 16878.	3.3	33
25	Molecular Cloning and Expression Analysis of Three Suppressors of Cytokine Signaling Genes (SOCS5,) Tj ETQq1	1 0.78431 2.2	L4 rgBT /Ove 1P
26	TmToll-7 Plays a Crucial Role in Innate Immune Responses Against Gram-Negative Bacteria by Regulating 5 AMP Genes in Tenebrio molitor. Frontiers in Immunology, 2019, 10, 310.	4.8	26
27	Transcriptome analysis of air-breathing land slug, Incilaria fruhstorferi reveals functional insights into growth, immunity, and reproduction. BMC Genomics, 2019, 20, 154.	2.8	9
28	<i>In silico</i> identification, characterization and expression analysis of <i>attacin</i> gene family in response to bacterial and fungal pathogens in <scp><i>Tenebrio molitor</i></scp> . Entomological Research, 2018, 48, 45-54.	1.1	19
29	RNA Sequencing, <i>De novo</i> assembly, functional annotation and SSR analysis of the endangered diving beetle <scp><i>Cybister chinensis</i></scp> (= <scp><i>Cybister japonicus</i></scp> ) using the Illumina platform. Entomological Research, 2018, 48, 60-72.	1.1	3
30	Transcriptome analysis of the threatened snail Ellobium chinense reveals candidate genes for adaptation and identifies SSRs for conservation genetics. Genes and Genomics, 2018, 40, 333-347.	1.4	6
31	Molecular Cloning and Effects of Tm14-3-3ζ-Silencing on Larval Survivability Against E. coli and C. albicans in Tenebrio molitor. Genes, 2018, 9, 330.	2.4	5
32	TmCactin plays an important role in Gram-negative and -positive bacterial infection by regulating expression of 7 AMP genes in Tenebrio molitor. Scientific Reports, 2017, 7, 46459.	3.3	34
33	Sequencing and de novo assembly of visceral mass transcriptome of the critically endangered land snail Satsuma myomphala: Annotation and SSR discovery. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2017, 21, 77-89.	1.0	10
34	De novo Transcriptome Generation and Annotation for Two Korean Endemic Land Snails, Aegista chejuensis and Aegista quelpartensis, Using Illumina Paired-End Sequencing Technology. International Journal of Molecular Sciences, 2016, 17, 379.	4.1	7
35	Transcriptome Profile of the Asian Giant Hornet ( <i>Vespa mandarinia</i> ) Using Illumina HiSeq 4000 Sequencing: <i>De Novo</i> Assembly, Functional Annotation, and Discovery of SSR Markers. International Journal of Genomics, 2016, 2016, 1-15.	1.6	24
36	Transcriptomic Analysis of the Endangered Neritid Species Clithon retropictus: De Novo Assembly, Functional Annotation, and Marker Discovery. Genes, 2016, 7, 35.	2.4	13

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37	The Silencing of a 14-3-3É› Homolog in Tenebrio molitor Leads to Increased Antimicrobial Activity in Hemocyte and Reduces Larval Survivability. Genes, 2016, 7, 53.	2.4	5
38	Transcriptome Analysis of the Tadpole Shrimp (Triops longicaudatus) by Illumina Paired-End Sequencing: Assembly, Annotation, and Marker Discovery. Genes, 2016, 7, 114.	2.4	15
39	Transcriptome sequencing and de novo characterization of Korean endemic land snail, Koreanohadra kurodana for functional transcripts and SSR markers. Molecular Genetics and Genomics, 2016, 291, 1999-2014.	2.1	14
40	<scp>RNA</scp> sequencing, <i>de novo</i> assembly, and functional annotation of an endangered <scp>N</scp> ymphalid butterfly, <scp><i>F</i></scp> <i>abriciana nerippe</i> â€ <scp>F</scp> elder, 1862. Entomological Research, 2016, 46, 148-161.	1.1	7
41	Mutan: A mixed linkage α-[(1,3)- and (1,6)]-d-glucan from Streptococcus mutans, that induces osteoclast differentiation and promotes alveolar bone loss. Carbohydrate Polymers, 2016, 137, 561-569.	10.2	15
42	Sequencing, De Novo Assembly, and Annotation of the Transcriptome of the Endangered Freshwater Pearl Bivalve, Cristaria plicata, Provides Novel Insights into Functional Genes and Marker Discovery. PLoS ONE, 2016, 11, e0148622.	2.5	61
43	Transcriptome Characterization for Non-Model Endangered Lycaenids, Protantigius superans and Spindasis takanosis, Using Illumina HiSeq 2500 Sequencing. International Journal of Molecular Sciences, 2015, 16, 29948-29970.	4.1	13
44	DEPLETION OF AUTOPHAGYâ€RELATED GENES ATG3 AND ATG5 IN <i>Tenebrio molitor</i> LEADS TO DECREASED SURVIVABILITY AGAINST AN INTRACELLULAR PATHOGEN, <i>Listeria monocytogenes</i> . Archives of Insect Biochemistry and Physiology, 2015, 88, 85-99.	1.5	18
45	Cloning, expression analysis, and RNA interference study of a HORMA domain containing autophagy-related gene 13 (ATG13) from the coleopteran beetle, Tenebrio molitor. Frontiers in Physiology, 2015, 6, 180.	2.8	12
46	Molecular cloning and characterization of autophagy-related gene TmATG8 in Listeria-invaded hemocytes of Tenebrio molitor. Developmental and Comparative Immunology, 2015, 51, 88-98.	2.3	24
47	Silencing of apolipophorinâ€ <scp>III</scp> causes abnormal adult morphological phenotype and susceptibility to <i><scp>L</scp>isteria monocytogenes</i> infection in <i><scp>T</scp>enebrio molitor</i> . Entomological Research, 2015, 45, 116-121.	1.1	5
48	Brazilin isolated from Caesalpinia sappan L. inhibits rheumatoid arthritis activity in a type-II collagen induced arthritis mouse model. BMC Complementary and Alternative Medicine, 2015, 15, 124.	3.7	32
49	Genomic organization, sequence characterization and expression analysis of Tenebrio molitor apolipophorin-III in response to an intracellular pathogen, Listeria monocytogenes. Gene, 2014, 534, 204-217.	2.2	17
50	Characterization of chitinaseâ€producing <i><scp>S</scp>erratia</i> and <i><scp>B</scp>acillus</i> strains isolated from insects. Entomological Research, 2014, 44, 109-120.	1.1	6
51	Gene structure, cDNA characterization and RNAi-based functional analysis of a myeloid differentiation factor 88 homolog in Tenebrio molitor larvae exposed to Staphylococcus aureus infection. Developmental and Comparative Immunology, 2014, 46, 208-221.	2.3	25
52	Isolation and Characterization of Chitinaseâ€Producing <i>Bacillus</i> and <i>Paenibacillus</i> Strains from Salted and Fermented Shrimp, <i>Acetes japonicus</i> . Journal of Food Science, 2014, 79, M665-74.	3.1	17
53	Mollusks Sequence Database: Version II. Korean Journal of Malacology, 2014, 30, 429-431.	0.1	6
54	<scp>E</scp> xpressed <scp>S</scp> equence <scp>T</scp> ags ( <scp>ESTs</scp> ) analysis of <i><scp>T</scp>enebrio molitor</i> larvae. Entomological Research, 2013, 43, 168-176.	1.1	5

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55	Identification and expression analysis of a novel R-type lectin from the coleopteran beetle, Tenebrio molitor. Journal of Invertebrate Pathology, 2013, 114, 226-229.	3.2	7
56	Molecular and immunohistochemical characterization of granulin gene encoded in Pieris rapae granulovirus genome. Journal of Invertebrate Pathology, 2013, 113, 7-17.	3.2	3
57	Molecular and immunohistochemical characterization of the chitinase gene from Pieris rapae granulovirus. Archives of Virology, 2013, 158, 1701-1718.	2.1	10
58	Molecular Cloning, Sequence Characterization and Expression Analysis of a CD63 Homologue from the Coleopteran Beetle, Tenebrio molitor. International Journal of Molecular Sciences, 2013, 14, 20744-20767.	4.1	16
59	Cloning, Characterization and Effect of TmPGRP-LE Gene Silencing on Survival of Tenebrio Molitor against Listeria monocytogenes Infection. International Journal of Molecular Sciences, 2013, 14, 22462-22482.	4.1	26
60	Expression analysis and immunohistochemical localization of putative tumor suppressor <scp>QM</scp> homologue from the cabbage butterfly, <i><scp>P</scp>ieris rapae</i> . Entomological Research, 2013, 43, 262-270.	1.1	1
61	Analysis of the Genome of a Korean Isolate of the Pieris rapae Granulovirus Enabled by Its Separation from Total Host Genomic DNA by Pulse-Field Electrophoresis. PLoS ONE, 2013, 8, e84183.	2.5	3
62	Reproductive Performance of Breeds and Hybrid of Silkworm, Bombyx mori L. with Special Reference to Egg Laying Rhythmicity. International Journal of Industrial Entomology, 2013, 26, 22-30.	0.1	0
63	Purification and characterization of tenecin 4, a new anti-Gram-negative bacterial peptide, from the beetle Tenebrio molitor. Developmental and Comparative Immunology, 2012, 36, 540-546.	2.3	65
64	Molecular Cloning and Characterization of Novel Morus alba Germin-Like Protein Gene Which Encodes for a Silkworm Gut Digestion-Resistant Antimicrobial Protein. PLoS ONE, 2012, 7, e50900.	2.5	7
65	lsozymic variations in specific and nonspecific esterase and its thermostability in silkworm, Bombyx mori L. Journal of Environmental Biology, 2012, 33, 837-42.	0.5	4
66	Induction of oxidative stress by non-lethal dose of mercury in rat liver: possible relationships between apoptosis and necrosis. Journal of Environmental Biology, 2010, 31, 413-6.	0.5	13
67	Molecular cloning and expression pattern of 14â€3â€3ζ from the malaria vector, <i>Anopheles sinensis</i> . Entomological Research, 2009, 39, 123-128.	1.1	4
68	Peptideâ€based polyclonal antibody against mosquito 14â€3â€3ζ recognizes 14â€3â€3 homolog from dipteran a lepidopteran insects. Entomological Research, 2009, 39, 129-134.	and 1.1	5
69	Current Status of Immune Deficiency Pathway in Tenebrio molitor Innate Immunity. Frontiers in Immunology, 0, 13, .	4.8	8