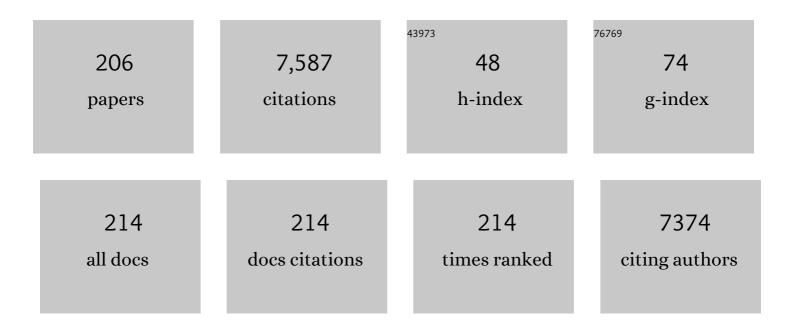
Sanjay Kumar

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Chicken Gut Microbiota: Importance and Detection Technology. Frontiers in Veterinary Science, 2018, 5, 254.	0.9	274
2	Assessment of genetic diversity and plant growth promoting attributes of psychrotolerant bacteria allied with wheat (Triticum aestivum) from the northern hills zone of India. Annals of Microbiology, 2015, 65, 1885-1899.	1.1	245
3	Molecular diversity and multifarious plant growth promoting attributes of Bacilli associated with wheat (<i>Triticum aestivum</i> L.) rhizosphere from six diverse agroâ€ecological zones of India. Journal of Basic Microbiology, 2016, 56, 44-58.	1.8	229
4	An RNA isolation system for plant tissues rich in secondary metabolites. BMC Research Notes, 2011, 4, 85.	0.6	214
5	Expression of SOD and APX genes positively regulates secondary cell wall biosynthesis and promotes plant growth and yield in Arabidopsis under salt stress. Plant Molecular Biology, 2015, 87, 615-631.	2.0	183
6	Phenylalanine ammonia-lyase (PAL) and cinnamate 4-hydroxylase (C4H) and catechins (flavan-3-ols) accumulation in tea. Functional and Integrative Genomics, 2009, 9, 125-134.	1.4	161
7	Protein complex directs hemoglobin-to-hemozoin formation in <i>Plasmodium falciparum</i> . Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 5392-5397.	3.3	143
8	Magnesium ammonium phosphate formation, recovery and its application as valuable resources: a review. Journal of Chemical Technology and Biotechnology, 2013, 88, 181-189.	1.6	141
9	Appraisal of diversity and functional attributes of thermotolerant wheat associated bacteria from the peninsular zone of India. Saudi Journal of Biological Sciences, 2019, 26, 1882-1895.	1.8	134
10	Probiotics and Atopic Dermatitis: An Overview. Frontiers in Microbiology, 2016, 7, 507.	1.5	128
11	De novo sequencing and characterization of Picrorhiza kurrooa transcriptome at two temperatures showed major transcriptome adjustments. BMC Genomics, 2012, 13, 126.	1.2	124
12	New aspects and strategies for methane mitigation from ruminants. Applied Microbiology and Biotechnology, 2014, 98, 31-44.	1.7	120
13	Purification and characterization of glutaminase-free l-asparaginase from Pectobacterium carotovorum MTCC 1428. Bioresource Technology, 2011, 102, 2077-2082.	4.8	113
14	Associative patterns among anaerobic fungi, methanogenic archaea, and bacterial communities in response to changes in diet and age in the rumen of dairy cows. Frontiers in Microbiology, 2015, 6, 781.	1.5	112
15	Effect of antibiotic withdrawal in feed on chicken gut microbial dynamics, immunity, growth performance and prevalence of foodborne pathogens. PLoS ONE, 2018, 13, e0192450.	1.1	112
16	Fuzzy time series forecasting method based on hesitant fuzzy sets. Expert Systems With Applications, 2016, 64, 557-568.	4.4	99
17	26S rRNA-based internal control gene primer pair for reverse transcription-polymerase chain reaction-based quantitative expression studies in diverse plant species. Analytical Biochemistry, 2004, 335, 330-333.	1.1	94
18	A comprehensive analysis of fifteen genes of steviol glycosides biosynthesis pathway in Stevia rebaudiana (Bertoni). Gene, 2012, 492, 276-284.	1.0	94

#	Article	IF	CITATIONS
19	Metagenomic assessment of the functional potential of the rumen microbiome in Holstein dairy cows. Anaerobe, 2016, 38, 50-60.	1.0	93
20	INTUITIONISTIC FUZZY SETS BASED METHOD FOR FUZZY TIME SERIES FORECASTING. Cybernetics and Systems, 2012, 43, 34-47.	1.6	90
21	Intuitionistic Fuzzy Time Series: An Approach for Handling Nondeterminism in Time Series Forecasting. IEEE Transactions on Fuzzy Systems, 2016, 24, 1270-1281.	6.5	86
22	Light and temperature regulated terpene biosynthesis: hepatoprotective monoterpene picroside accumulation in Picrorhiza kurrooa. Functional and Integrative Genomics, 2010, 10, 393-404.	1.4	85
23	Bacterial diversity associated with feeding dry forage at different dietary concentrations in the rumen contents of Mehshana buffalo (Bubalus bubalis) using 16S pyrotags. Anaerobe, 2014, 25, 31-41.	1.0	84
24	The basic and applied aspects of superoxide dismutase. Journal of Molecular Catalysis B: Enzymatic, 2011, 68, 129-138.	1.8	80
25	Engineering a thermo-stable superoxide dismutase functional at sub-zero to >50°C, which also tolerates autoclaving. Scientific Reports, 2012, 2, 387.	1.6	79
26	An early gene of the flavonoid pathway, flavanone 3-hydroxylase, exhibits a positive relationship with the concentration of catechins in tea (Camellia sinensis). Tree Physiology, 2008, 28, 1349-1356.	1.4	78
27	p-Coumarate:CoA ligase as a key gene in the yield of catechins in tea [Camellia sinensis (L.) O. Kuntze]. Functional and Integrative Genomics, 2009, 9, 271-275.	1.4	76
28	Superoxide dismutase: an industrial perspective. Critical Reviews in Biotechnology, 2011, 31, 65-76.	5.1	76
29	Factors affecting rumen methanogens and methane mitigation strategies. World Journal of Microbiology and Biotechnology, 2009, 25, 1557-1566.	1.7	75
30	Purification and partial characterization of a low temperature responsive Mn-SOD from tea (Camellia) Tj ETQq	0 0 0 _{rg} BT /(Dverlock 10 T
31	Overexpression of Camellia sinensis Thaumatin-Like Protein, CsTLP in Potato Confers Enhanced Resistance to Macrophomina phaseolina and Phytophthora infestans Infection. Molecular Biotechnology, 2013, 54, 609-622.	1.3	71
32	A new anaerobic fungus (Oontomyces anksri gen. nov., sp. nov.) from the digestive tract of the Indian camel (Camelus dromedarius). Fungal Biology, 2015, 119, 731-737.	1.1	71
33	Bacterial Diversity Dynamics Associated with Different Diets and Different Primer Pairs in the Rumen of Kankrej Cattle. PLoS ONE, 2014, 9, e111710.	1.1	71
34	Partitions based computational method for high-order fuzzy time series forecasting. Expert Systems With Applications, 2012, 39, 12158-12164.	4.4	65
35	Transgenic Potato Plants Overexpressing SOD and APX Exhibit Enhanced Lignification and Starch Biosynthesis with Improved Salt Stress Tolerance. Plant Molecular Biology Reporter, 2017, 35, 504-518.	1.0	65
36	Molecular regulation of catechins biosynthesis in tea [Camellia sinensis (L.) O. Kuntze]. Gene, 2012, 495, 205-210.	1.0	63

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37	D1/D2 Domain of Large-Subunit Ribosomal DNA for Differentiation of Orpinomyces spp. Applied and Environmental Microbiology, 2011, 77, 6722-6725.	1.4	62
38	Methanobrevibacter boviskoreani sp. nov., isolated from the rumen of Korean native cattle. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 4196-4201.	0.8	61
39	Growth-Inhibitory Effect of Heparin on Babesia Parasites. Antimicrobial Agents and Chemotherapy, 2004, 48, 236-241.	1.4	57
40	Recycle of electrolytically dissolved struvite as an alternative to enhance phosphate and nitrogen recovery from swine wastewater. Journal of Hazardous Materials, 2011, 195, 175-181.	6.5	53
41	RNA-seq-mediated transcriptome analysis of actively growing and winter dormant shoots identifies non-deciduous habit of evergreen tree tea during winters. Scientific Reports, 2014, 4, 5932.	1.6	52
42	Differential display-mediated identification of three drought-responsive expressed sequence tags in tea [Camellia sinensis (L.) O. Kuntze]. Journal of Biosciences, 2005, 30, 231-235.	0.5	51
43	An Improved Protocol for the Isolation of RNA from Roots of Tea (Camellia sinensis (L.) O. Kuntze). Molecular Biotechnology, 2012, 52, 82-88.	1.3	51
44	PROBABILISTIC AND INTUITIONISTIC FUZZY SETS–BASED METHOD FOR FUZZY TIME SERIES FORECASTING. Cybernetics and Systems, 2014, 45, 349-361.	1.6	51
45	Molecular dissection of transcriptional reprogramming of steviol glycosides synthesis in leaf tissue during developmental phase transitions in Stevia rebaudiana Bert. Scientific Reports, 2017, 7, 11835.	1.6	51
46	Simulation of salt and water movement and estimation of water productivity of rice crop irrigated with saline water. Paddy and Water Environment, 2010, 8, 333-346.	1.0	50
47	Tea (Camellia sinensis) clones with shorter periods of winter dormancy exhibit lower accumulation of reactive oxygen species. Tree Physiology, 2007, 27, 1253-1259.	1.4	49
48	Expression of 3-hydroxy-3-methylglutaryl-CoA reductase, p-hydroxybenzoate-m-geranyltransferase and genes of phenylpropanoid pathway exhibits positive correlation with shikonins content in arnebia [Arnebia euchroma (Royle) Johnston]. BMC Molecular Biology, 2010, 11, 88.	3.0	49
49	Tea (Camellia sinensis (L.) O. Kuntze) clone with lower period of winter dormancy exhibits lesser cellular damage in response to low temperature. Plant Physiology and Biochemistry, 2005, 43, 383-388.	2.8	47
50	Microbial profiles, in vitro gas production and dry matter digestibility based on various ratios of roughage to concentrate. Annals of Microbiology, 2013, 63, 541-545.	1,1	47
51	Role of live microbial feed supplements with reference to anaerobic fungi in ruminant productivity: A review. Journal of Integrative Agriculture, 2015, 14, 550-560.	1.7	46
52	<i>In-vivo</i> Therapeutic Efficacy Trial with Artemisinin Derivative, Buparvaquone and Imidocarb Dipropionate against <i>Babesia equi</i> Infection in Donkeys. Journal of Veterinary Medical Science, 2003, 65, 1171-1177.	0.3	44
53	Dysbiosis of the Fecal Microbiota in Cattle Infected with Mycobacterium avium subsp. paratuberculosis. PLoS ONE, 2016, 11, e0160353.	1.1	44
54	Characterization of dihydroflavonol 4-reductase cDNA in tea [Camellia sinensis (L.) O. Kuntze]. Plant Biotechnology Reports, 2009, 3, 95-101.	0.9	43

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55	Differential display mediated cloning of anthocyanidin reductase gene from tea (Camellia sinensis) and its relationship with the concentration of epicatechins. Tree Physiology, 2009, 29, 837-846.	1.4	43
56	Lipoxygenase in Caragana jubata responds to low temperature, abscisic acid, methyl jasmonate and salicylic acid. Gene, 2011, 483, 49-53.	1.0	42
57	Particle swarm optimization and intuitionistic fuzzy set-based novel method for fuzzy time series forecasting. Granular Computing, 2022, 7, 285-303.	4.4	40
58	The Promoter of AtUSP Is Co-regulated by Phytohormones and Abiotic Stresses in Arabidopsis thaliana. Frontiers in Plant Science, 2016, 7, 1957.	1.7	39
59	Complete genome sequence of Pseudomonas frederiksbergensis ERDD5:01 revealed genetic bases for survivability at high altitude ecosystem and bioprospection potential. Genomics, 2019, 111, 492-499.	1.3	38
60	Effect of altitude on the primary products of photosynthesis and the associated enzymes in barley and wheat. Photosynthesis Research, 2006, 88, 63-71.	1.6	37
61	Responses to winter dormancy, temperature, and plant hormones share gene networks. Functional and Integrative Genomics, 2011, 11, 659-664.	1.4	37
62	Complete genome sequence of Arthrobacter sp. ERGS1:01, a putative novel bacterium with prospective cold active industrial enzymes, isolated from East Rathong glacier in India. Journal of Biotechnology, 2015, 214, 139-140.	1.9	37
63	Fuzzy Time Series Model Based on Intuitionistic Fuzzy Sets for Empirical Research in Stock Market. International Journal of Applied Evolutionary Computation, 2012, 3, 71-84.	0.7	36
64	Transcriptome sequencing of rhizome tissue of Sinopodophyllum hexandrum at two temperatures. BMC Genomics, 2014, 15, 871.	1.2	35
65	Arnebia euchroma, a Plant Species of Cold Desert in the Himalayas, Harbors Beneficial Cultivable Endophytes in Roots and Leaves. Frontiers in Microbiology, 2021, 12, 696667.	1.5	35
66	Global Transcriptional Insights of Pollen-Pistil Interactions Commencing Self-Incompatibility and Fertilization in Tea [Camellia sinensis (L.) O. Kuntze]. International Journal of Molecular Sciences, 2019, 20, 539.	1.8	34
67	A Broad Temperature Active Lipase Purified From a Psychrotrophic Bacterium of Sikkim Himalaya With Potential Application in Detergent Formulation. Frontiers in Bioengineering and Biotechnology, 2020, 8, 642.	2.0	34
68	Genome Sequence of Oscillibacter ruminantium Strain GH1, Isolated from Rumen of Korean Native Cattle. Journal of Bacteriology, 2012, 194, 6362-6362.	1.0	33
69	Changes in methane emission, rumen fermentation in response to diet and microbial interactions. Research in Veterinary Science, 2013, 94, 263-268.	0.9	33
70	Development of an antibody-ELISA for seroprevalence of Trypanosoma evansi in equids of North and North-western regions of India. Veterinary Parasitology, 2013, 196, 251-257.	0.7	32
71	Metagenomic Evidence of the Prevalence and Distribution Patterns of Antimicrobial Resistance Genes in Dairy Agroecosystems. Foodborne Pathogens and Disease, 2016, 13, 296-302.	0.8	32
72	Strategies for high-altitude adaptation revealed from high-quality draft genome of non-violacein producing Janthinobacterium lividum ERGS5:01. Standards in Genomic Sciences, 2018, 13, 11.	1.5	32

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73	Ectopic expression of SOD and APX genes in Arabidopsis alters metabolic pools and genes related to secondary cell wall cellulose biosynthesis and improve salt tolerance. Molecular Biology Reports, 2019, 46, 1985-2002.	1.0	32
74	Transcriptional profiling of contrasting genotypes revealed key candidates and nucleotide variations for drought dissection in Camellia sinensis (L.) O. Kuntze. Scientific Reports, 2019, 9, 7487.	1.6	32
75	A perspective on Theileria equi infections in donkeys. Japanese Journal of Veterinary Research, 2009, 56, 171-80.	0.7	32
76	Over-expression of superoxide dismutase exhibits lignification of vascular structures in Arabidopsis thaliana. Journal of Plant Physiology, 2010, 167, 757-760.	1.6	31
77	Induction of a new isozyme of superoxide dismutase at low temperature in Potentilla astrisanguinea Lodd. variety argyrophylla (Wall. ex. Lehm) Griers. Journal of Plant Physiology, 2001, 158, 1093-1097.	1.6	30
78	Isolation and characterization of methanogens from rumen of Murrah buffalo. Annals of Microbiology, 2012, 62, 345-350.	1.1	30
79	A COMPUTATIONAL METHOD FOR FUZZY TIME SERIES FORECASTING BASED ON DIFFERENCE PARAMETERS. International Journal of Modeling, Simulation, and Scientific Computing, 2013, 04, 1250023.	0.9	30
80	Complete genome sequence of Arthrobacter alpinus ERGS4:06, a yellow pigmented bacterium tolerant to cold and radiations isolated from Sikkim Himalaya. Journal of Biotechnology, 2016, 220, 86-87.	1.9	30
81	Diverse culturable bacterial communities with cellulolytic potential revealed from pristine habitat in Indian trans-Himalaya. Canadian Journal of Microbiology, 2018, 64, 798-808.	0.8	30
82	Bioplastic reservoir of diverse bacterial communities revealed along altitude gradient of Pangi-Chamba trans-Himalayan region. FEMS Microbiology Letters, 2018, 365, .	0.7	30
83	Differential expression of Histone H3 gene in tea (Camellia sinensis (L.) O. Kuntze) suggests its role in growing tissue. Molecular Biology Reports, 2009, 36, 537-542.	1.0	29
84	A shared response of thaumatin like protein, chitinase, and late embryogenesis abundant protein3 to environmental stresses in tea [Camellia sinensis (L.) O. Kuntze]. Functional and Integrative Genomics, 2012, 12, 565-571.	1.4	29
85	Cloning and expression analysis of ten genes associated with picrosides biosynthesis in Picrorhiza kurrooa. Gene, 2013, 515, 320-328.	1.0	29
86	Spatial distribution, risk factors and haemato-biochemical alterations associated with Theileria equi infected equids of Punjab (India) diagnosed by indirect ELISA and nested PCR. Acta Tropica, 2016, 155, 104-112.	0.9	29
87	Copper, zinc superoxide dismutase of Curcuma aromatica is a kinetically stable protein. Process Biochemistry, 2014, 49, 1288-1296.	1.8	28
88	Bioprocess for co-production of polyhydroxybutyrate and violacein using Himalayan bacterium Iodobacter sp. PCH194. Bioresource Technology, 2021, 319, 124235.	4.8	28
89	Early low-temperature responsive mitogen activated protein kinases RaMPK1 and RaMPK2 from Rheum australe D. Don respond differentially to diverse stresses. Molecular Biology Reports, 2010, 37, 933-938.	1.0	27
90	Characterization of gene expression of QM from Caragana jubata, a plant species that grows under extreme cold. Molecular Biology Reports, 2010, 37, 1003-1010.	1.0	27

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91	Biotechnological interventions for harnessing podophyllotoxin from plant and fungal species: current status, challenges, and opportunities for its commercialization. Critical Reviews in Biotechnology, 2017, 37, 739-753.	5.1	27
92	Network design for groundwater level monitoring in Upper Bari Doab Canal tract, Punjab, India. Irrigation and Drainage, 2005, 54, 431-442.	0.8	26
93	Structure of a superoxide dismutase and implications for copper-ion chelation. Acta Crystallographica Section D: Biological Crystallography, 2008, 64, 892-901.	2.5	26
94	Expression of Babesia equi EMA-1 and EMA-2 during merozoite developmental stages in erythrocyte and their interaction with erythrocytic membrane skeleton. Molecular and Biochemical Parasitology, 2004, 133, 221-227.	0.5	25
95	Passive transfer of Theileria equi antibodies to neonate foals of immune tolerant mares. Veterinary Parasitology, 2008, 151, 80-85.	0.7	25
96	A fuzzy time series forecasting method induced by intuitionistic fuzzy sets. International Journal of Modeling, Simulation, and Scientific Computing, 2015, 06, 1550041.	0.9	25
97	Cloning and differential expression of QM like protein homologue from tea [Camellia sinensis (L.) O. Kuntze]. Molecular Biology Reports, 2009, 36, 921-927.	1.0	24
98	Alpha-tubulin (CsTUA) up-regulated during winter dormancy is a low temperature inducible gene in tea [Camellia sinensis (L.) O. Kuntze]. Molecular Biology Reports, 2012, 39, 3485-3490.	1.0	24
99	Improved callus induction, shoot regeneration, and salt stress tolerance in Arabidopsis overexpressing superoxide dismutase from Potentilla atrosanguinea. Protoplasma, 2015, 252, 41-51.	1.0	24
100	Microbial polyhydroxyalkanoates from extreme niches: Bioprospection status, opportunities and challenges. International Journal of Biological Macromolecules, 2020, 147, 1255-1267.	3.6	24
101	2C-methyl- d- erythritol 2,4-cyclodiphosphate synthase from Stevia rebaudiana Bertoni is a functional gene. Molecular Biology Reports, 2012, 39, 10971-10978.	1.0	22
102	Multiplex PCR for detection of Trypanosoma evansi and Theileria equi in equids of Punjab, India. Veterinary Parasitology, 2015, 211, 293-299.	0.7	22
103	Transcriptome analysis of stem wood of Nothapodytes nimmoniana (Graham) Mabb. identifies genes associated with biosynthesis of camptothecin, an anti-carcinogenic molecule. Journal of Biosciences, 2016, 41, 119-131.	0.5	22
104	Standardisation and comparison of serial dilution and single dilution enzyme linked immunosorbent assay (ELISA) using different antigenic preparations of the Babesia (Theileria) equi parasite. Veterinary Research, 2003, 34, 71-83.	1.1	22
105	Effect of a bioactive product SEL001 from Lactobacillus sakei probio65 on gut microbiota and its anti-colitis effects in a TNBS-induced colitis mouse model. Saudi Journal of Biological Sciences, 2020, 27, 261-270.	1.8	21
106	Elevated CO2 and temperature influence key proteins and metabolites associated with photosynthesis, antioxidant and carbon metabolism in Picrorhiza kurroa. Journal of Proteomics, 2020, 219, 103755.	1.2	21
107	Genome Sequence of <i>Methanobrevibacter</i> sp. Strain JH1, Isolated from Rumen of Korean Native Cattle. Genome Announcements, 2013, 1, .	0.8	20
108	Braving the attitude of altitude: Caragana jubata at work in cold desert of Himalaya. Scientific Reports, 2013, 3, 1022.	1.6	19

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109	Use of Intuitionistic Fuzzy Time Series in Forecasting Enrollments to an Academic Institution. Advances in Intelligent Systems and Computing, 2016, , 843-852.	0.5	19
110	Fuzzy time series forecasting based on hesitant fuzzy sets, particle swarm optimization and support vector machine-based hybrid method. Granular Computing, 2022, 7, 861-879.	4.4	19
111	Infectivity rate and transmission potential of Hyalomma anatolicum anatolicum ticks for Babesia equi infection. Veterinary Parasitology, 2007, 144, 338-343.	0.7	18
112	CsNAM-like protein encodes a nuclear localized protein and responds to varied cues in tea [Camellia sinensis (L.) O. Kuntze]. Gene, 2012, 502, 69-74.	1.0	18
113	Dehydrin2 is a stress-inducible, whereas Dehydrin1 is constitutively expressed but up-regulated gene under varied cues in tea [Camellia sinensis (L.) O. Kuntze]. Molecular Biology Reports, 2013, 40, 3859-3863.	1.0	18
114	The GATA and SORLIP motifs in the 3-hydroxy-3-methylglutaryl-CoA reductase promoter of Picrorhiza kurrooa for the control of light-mediated expression. Functional and Integrative Genomics, 2014, 14, 191-203.	1.4	18
115	Genome Assembly of Chryseobacterium polytrichastri ERMR1:04, a Psychrotolerant Bacterium with Cold Active Proteases, Isolated from East Rathong Glacier in India. Genome Announcements, 2015, 3, .	0.8	18
116	Effect of feeding different dietary levels of energy and protein on growth performance and immune status of Vanaraja chicken in the tropic. Veterinary World, 2016, 9, 893-899.	0.7	18
117	Enumeration of methanogens with a focus on fluorescence in situ hybridization. Die Naturwissenschaften, 2011, 98, 457-472.	0.6	17
118	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
119	Development of EMA-2 recombinant antigen based enzyme-linked immunosorbent assay for seroprevalence studies of Theileria equi infection in Indian equine population. Veterinary Parasitology, 2013, 198, 10-17.	0.7	17
120	Comparative genomics and physiological investigation supported safety, cold adaptation, efficient hydrolytic and plant growth-promoting potential of psychrotrophic Glutamicibacter arilaitensis LJH19, isolated from night-soil compost. BMC Genomics, 2021, 22, 307.	1.2	17
121	A comparison of rumen microbial profiles in dairy cows as retrieved by 454 Roche and Ion Torrent (PGM) sequencing platforms. PeerJ, 2016, 4, e1599.	0.9	17
122	Serodiagnosis of Babesia equi infection—a comparison of Dot-ELISA, complement fixation test and capillary tube agglutination test. Veterinary Parasitology, 1997, 69, 171-176.	0.7	16
123	Vaccination of donkeys against Babesia equi using killed merozoite immunogen. Veterinary Parasitology, 2002, 106, 19-33.	0.7	15
124	A stress-responsive late embryogenesis abundant protein 7 (CsLEA7) of tea [Camellia sinensis (L.) O. Kuntze] encodes for a chaperone that imparts tolerance to Escherichia coli against stresses. Molecular Biology Reports, 2014, 41, 7191-7200.	1.0	15
125	Impact estimation of animal trypanosomosis (surra) on livestock productivity in India using simulation model: Current and future perspective. Veterinary Parasitology: Regional Studies and Reports, 2017, 10, 1-12.	0.3	15
126	Intuitionistic Fuzzy Set-Based Computational Method for Financial Time Series Forecasting. Fuzzy Information and Engineering, 2018, 10, 307-323.	1.0	15

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127	Heterologous Expression of Key C and N Metabolic Enzymes Improves Re-assimilation of Photorespired CO ₂ and NH ₃ , and Growth. Plant Physiology, 2018, 177, 1396-1409.	2.3	15
128	A Computational Method of Forecasting Based on Intuitionistic Fuzzy Sets and Fuzzy Time Series. Advances in Intelligent and Soft Computing, 2012, , 993-1000.	0.2	15
129	Differences in the Activation State of Ribulose-1,5-bisphosphate Carboxylase/Oxygenase in Barley, Pea, and Wheat at Two Altitudes. Photosynthetica, 2004, 42, 303-305.	0.9	14
130	Modelling the effect of canal bed elevation on seepage and water table rise in a sand box filled with loamy soil. Irrigation Science, 2009, 27, 191-200.	1.3	14
131	Effect of elevated [CO 2] and temperature on growth, physiology and essential oil composition of Salvia sclarea L. in the western Himalayas. Journal of Applied Research on Medicinal and Aromatic Plants, 2017, 6, 22-30.	0.9	14
132	Comparative evaluation of lignocellulolytic activities of filamentous cultures of monocentric and polycentric anaerobic fungi. Anaerobe, 2018, 50, 76-79.	1.0	14
133	Genome-wide identification and expression profiling of genes encoding universal stress proteins (USP) identify multi-stress responsive USP genes in Arabidopsis thaliana. Plant Physiology Reports, 2019, 24, 434-445.	0.7	14
134	Optimized chromogenic dyes-based identification and quantitative evaluation of bacterial l-asparaginase with low/no glutaminase activity bioprospected from pristine niches in Indian trans-Himalaya. 3 Biotech, 2019, 9, 275.	1.1	14
135	Insight Into Dynamics of Gut Microbial Community of Broilers Fed With Fructooligosaccharides Supplemented Low Calcium and Phosphorus Diets. Frontiers in Veterinary Science, 2019, 6, 95.	0.9	14
136	Prickle morphogenesis in rose is coupled with secondary metabolite accumulation and governed by canonical MBW transcriptional complex. Plant Direct, 2021, 5, e00325.	0.8	13
137	An A20/AN1-zinc-finger domain containing protein gene in tea is differentially expressed during winter dormancy and in response to abiotic stress and plant growth regulators. Plant Gene, 2015, 1, 1-7.	1.4	12
138	Stress-Responsive cis-Regulatory Elements Underline Podophyllotoxin Biosynthesis and Better Performance of Sinopodophyllum hexandrum Under Water Deficit Conditions. Frontiers in Plant Science, 2021, 12, 751846.	1.7	12
139	Molecular evidence of Theileria equi infection in Hyalomma anatolicum ticks infested on sero-positive Indian horses. Acta Parasitologica, 2015, 60, 322-9.	0.4	11
140	Amino acids flanking the central core of Cu,Zn superoxide dismutase are important in retaining enzyme activity after autoclaving. Journal of Biomolecular Structure and Dynamics, 2016, 34, 475-485.	2.0	11
141	In vitro growth inhibitory efficacy of some target specific novel drug molecules against Theileria equi. Veterinary Parasitology, 2016, 217, 1-6.	0.7	11
142	Comparative transcriptome analysis of Rheum australe, an endangered medicinal herb, growing in its natural habitat and those grown in controlled growth chambers. Scientific Reports, 2021, 11, 3702.	1.6	11
143	Atypical human trypanosomosis: Potentially emerging disease with lack of understanding. Zoonoses and Public Health, 2022, 69, 259-276.	0.9	11
144	Comparative efficacy of different in vitro cultivation media for Trypanosoma evansi isolated from different mammalian hosts inhabiting different geographical areas of India. Journal of Parasitic Diseases, 2015, 39, 174-178.	0.4	10

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145	A parsimonious approach to delineating groundwater potential zones using geospatial modeling and multicriteria decision analysis techniques under limited data availability condition. Engineering Reports, 2019, 1, e12073.	0.9	10
146	The first draft genome of Picrorhiza kurrooa, an endangered medicinal herb from Himalayas. Scientific Reports, 2021, 11, 14944.	1.6	10
147	Managing reactive oxygen species—Some learnings from high altitude extremophytes. Environmental and Experimental Botany, 2021, 189, 104525.	2.0	10
148	Copper, zinc superoxide dismutase from Caragana jubata : A thermostable enzyme that functions under a broad pH and temperature window. Process Biochemistry, 2016, 51, 1434-1444.	1.8	9
149	Drug-induced reactive oxygen species–mediated inhibitory effect on growth of Trypanosoma evansi in axenic culture system. Parasitology Research, 2020, 119, 3481-3489.	0.6	9
150	Membrane localized thaumatin-like protein from tea (CsTLP) enhanced seed yield and the plant survival under drought stress in Arabidopsis thaliana. Plant Physiology and Biochemistry, 2021, 163, 36-44.	2.8	9
151	Timeâ€series ecoâ€metabolomics reveals extensive reshuffling in metabolome during transition from cold acclimation to deâ€acclimation in an alpine shrub. Physiologia Plantarum, 2021, 173, 1824-1840.	2.6	9
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153	Transplacental Transmission of Trypanosoma evansi From Experimentally Infected Donkey Mare to Neonatal Foal. Journal of Equine Veterinary Science, 2015, 35, 337-341.	0.4	8
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