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List of Publications by Year in descending order

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279798 330143 1,477 48 23 37 citations h-index g-index papers 49 49 49 2448 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Cultivated Tomato (Solanum lycopersicum L.) Suffered a Severe Cytoplasmic Bottleneck during Domestication: Implications from Chloroplast Genomes. Plants, 2020, 9, 1443.	3.5	11
2	Peruvian chicha: A Focus on the Microbial Populations of This Ancient Maize-Based Fermented Beverage. Microorganisms, 2020, 8, 93.	3.6	21
3	Adaptive responses along a depth and a latitudinal gradient in the endemic seagrass Posidonia oceanica. Heredity, 2019, 122, 233-243.	2.6	18
4	Genetic and functional characterization of the bacterial community on fruit of three raspberry (Rubus idaeus) cultivars. Journal of Berry Research, 2019, 9, 227-247.	1.4	11
5	Plant Microbiome and Its Link to Plant Health: Host Species, Organs and Pseudomonas syringae pv. actinidiae Infection Shaping Bacterial Phyllosphere Communities of Kiwifruit Plants. Frontiers in Plant Science, 2018, 9, 1563.	3.6	51
6	Comparative transcriptome analysis of the interaction between Actinidia chinensis var. chinensis and Pseudomonas syringae pv. actinidiae in absence and presence of acibenzolar-S-methyl. BMC Genomics, 2018, 19, 585.	2.8	33
7	Biofilm Formation on Stainless Steel by Streptococcus thermophilus UC8547 in Milk Environments Is Mediated by the Proteinase PrtS. Applied and Environmental Microbiology, 2017, 83, .	3.1	24
8	Transcriptome Analysis of the Melon-Fusarium oxysporum f. sp. melonis Race 1.2 Pathosystem in Susceptible and Resistant Plants. Frontiers in Plant Science, 2017, 8, 362.	3.6	43
9	Survey on the phage resistance mechanisms displayed by a dairy Lactobacillus helveticus strain. Food Microbiology, 2017, 66, 110-116.	4.2	22
10	Comparative transcriptome profiling of resistant and susceptible rice genotypes in response to the seedborne pathogen Fusarium fujikuroi. BMC Genomics, 2016, 17, 608.	2.8	99
11	Draft Genome Sequence of <i>Clostridium sporogenes</i> Strain UC9000 Isolated from Raw Milk. Genome Announcements, 2016, 4, .	0.8	O
12	Transcriptome characterisation and simple sequence repeat marker discovery in the seagrass Posidonia oceanica. Scientific Data, 2016, 3, 160115.	5.3	13
13	Draft Genome Sequence of Pediococcus parvulus 2.6, a Probiotic \hat{l}^2 -Glucan Producer Strain. Genome Announcements, 2016, 4, .	0.8	6
14	Viable But Not Culturable (VBNC) state of Brettanomyces bruxellensis in wine: New insights on molecular basis of VBNC behaviour using a transcriptomic approach. Food Microbiology, 2016, 59, 196-204.	4.2	76
15	Draft Genome Sequence of Lactobacillus collinoides CUPV237, an Exopolysaccharide and Riboflavin Producer Isolated from Cider. Genome Announcements, 2016, 4, .	0.8	3
16	Draft genome sequence and chemical profiling of Fusarium langsethiae, an emerging producer of type A trichothecenes. International Journal of Food Microbiology, 2016, 221, 29-36.	4.7	27
17	Integrate genome-based assessment of safety for probiotic strains: Bacillus coagulans GBI-30, 6086 as a case study. Applied Microbiology and Biotechnology, 2016, 100, 4595-4605.	3.6	76
18	Physical Mapping of Bread Wheat Chromosome 5A: An Integrated Approach. Plant Genome, 2015, 8, eplantgenome2015.03.0011.	2.8	11

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19	Draft Genome Sequence of Lactobacillus plantarum Lp90 Isolated from Wine. Genome Announcements, 2015, 3, .	0.8	17
20	Transcriptome changes associated with cold acclimation in leaves of olive tree (Olea europaea L.). Tree Genetics and Genomes, 2015, 11, 1.	1.6	31
21	Early transcriptional changes in Beta vulgaris in response to low temperature. Planta, 2015, 242, 187-201.	3.2	31
22	Draft Genome Sequence of Three Antibiotic-Resistant Leuconostoc mesenteroides Strains of Dairy Origin. Genome Announcements, 2015, 3, .	0.8	6
23	Integrated views in plant breeding: from the perspective of biotechnology. , 2015, , 467-486.		2
24	Genome Sequence of Oenococcus oeni OM27, the First Fully Assembled Genome of a Strain Isolated from an Italian Wine. Genome Announcements, 2014, 2, .	0.8	28
25	Genome Sequences of Five Oenococcus oeni Strains Isolated from Nero Di Troia Wine from the Same Terroir in Apulia, Southern Italy. Genome Announcements, 2014, 2, .	0.8	35
26	Draft Genome Sequence of Bacillus coagulans GBI-30, 6086, a Widely Used Spore-Forming Probiotic Strain. Genome Announcements, 2014, 2, .	0.8	16
27	De novo genome assembly of the soil-borne fungus and tomato pathogen Pyrenochaeta lycopersici. BMC Genomics, 2014, 15, 313.	2.8	39
28	Leptin gene haplotypes are associated with change in immunological and hematological variables in dairy cow during the peripartum period1. Journal of Animal Science, 2012, 90, 16-26.	0.5	11
29	A Survey of MicroRNA Length Variants Contributing to miRNome Complexity in Peach (Prunus Persica) Tj ETQq1 1	l <u>9.7</u> 8431	4 rgBT /Ove
30	Comparative Transcriptome Profiling of the Early Response to Magnaporthe oryzae in Durable Resistant vs Susceptible Rice (Oryza sativa L.) Genotypes. PLoS ONE, 2012, 7, e51609.	2.5	149
31	Association analyses of single nucleotide polymorphisms in the LEP and SCD1 genes on the fatty acid profile of muscle fat in Simmental bulls. Meat Science, 2011, 87, 344-348.	5 . 5	37
32	Diversity in the Response to Low Temperature in Representative Barley Genotypes Cultivated in Europe. Crop Science, 2011, 51, 2759-2779.	1.8	42
33	Characterization of a SNPs panel for meat traceability in six cattle breeds. Food Control, 2009, 20, 856-860.	5. 5	26
34	Exploring polymorphisms and effects on milk traits of the DGAT1, SCD1 and GHR genes in four cattle breeds. Livestock Science, 2009, 125, 74-79.	1.6	23
35	Fatty acid composition and sensory properties of Italian Simmental beef as affected by gene frequency of Montbéliarde origin. Meat Science, 2009, 83, 543-550.	5. 5	11
36	Breed differences in sheep milk fatty acid profiles: Opportunities for sustainable use of animal genetic resources. Small Ruminant Research, 2008, 78, 24-31.	1.2	44

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37	DNA Polymorphisms in River Buffalo Leptin Gene. Italian Journal of Animal Science, 2007, 6, 342-344.	1.9	9
38	Short Communication: Effect of Stearoyl-Coenzyme A Desaturase Polymorphism on Fatty Acid Composition of Milk. Journal of Dairy Science, 2007, 90, 3553-3558.	3.4	84
39	Meat molecular traceability: How to choose the best set of microsatellites? Meat Science, 2006, 72, 312-317.	5.5	38
40	Analysis of the genetic diversity between Gentile di Puglia, Sopravissana and Sarda sheep breeds using microsatellite markers. Italian Journal of Animal Science, 2006, 5, 73-78.	1.9	9
41	Single nucleotide polymorphism detection in promoter III of the acetyl-CoA carboxylase-alpha gene in sheep. Journal of Animal Breeding and Genetics, 2005, 122, 418-420.	2.0	13
42	Single nucleotide polymorphism detection in promoter I of the acetyl-CoA carboxylase-α gene in sheep. Small Ruminant Research, 2005, 59, 49-53.	1.2	6
43	Partial sequencing of Stearoyl-CoA desaturase gene in buffalo. Italian Journal of Animal Science, 2005, 4, 25-27.	1.9	3
44	A novel protein kinase C \hat{l} ±-dependent signal to ERK1/2 activated by \hat{l} ± \hat{V} \hat{l} 23 integrin in osteoclasts and in Chinese hamster ovary (CHO) cells. Journal of Cell Science, 2005, 118, 3263-3275.	2.0	60
45	Melanoma cells stimulate osteoclastogenesis, c-Src expression and osteoblast cytokines. European Journal of Cancer, 2001, 37, 629-640.	2.8	26
46	Gene silencing by S-adenosylmethionine in muscle differentiation. FEBS Letters, 2001, 508, 337-340.	2.8	31
47	Erythroid differentiation and regulatory gene expression are modulated by adenosine derivatives interfering with S-adenosylmethionine metabolic pathway. Journal of Cellular Biochemistry, 2001, 81, 401-412.	2.6	3
48	Dermorphin and Deltorphin Glycosylated Analogues:  Synthesis and Antinociceptive Activity after Systemic Administration. Journal of Medicinal Chemistry, 1999, 42, 400-404.	6.4	69