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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Negative environmental impacts of antibiotic-contaminated effluents from pharmaceutical industries. Water Research, 2017, 126, 79-87.	5.3	240
2	De Novo Assembly and Functional Annotation of the Olive (Olea europaea) Transcriptome. DNA Research, 2013, 20, 93-108.	1.5	84
3	Pollution from azithromycin-manufacturing promotes macrolide-resistance gene propagation and induces spatial and seasonal bacterial community shifts in receiving river sediments. Environment International, 2019, 123, 501-511.	4.8	74
4	Antibiotic-manufacturing sites are hot-spots for the release and spread of antibiotic resistance genes and mobile genetic elements in receiving aquatic environments. Environment International, 2019, 130, 104735.	4.8	63
5	Usefulness of a New Large Set of High Throughput EST-SNP Markers as a Tool for Olive Germplasm Collection Management. Frontiers in Plant Science, 2018, 9, 1320.	1.7	57
6	Role of metabolism during viral infections, and crosstalk with the innate immune system. Intractable and Rare Diseases Research, 2016, 5, 90-96.	0.3	52
7	Fever as an important resource for infectious diseases research. Intractable and Rare Diseases Research, 2016, 5, 97-102.	0.3	45
8	Functional Repertoire of Antibiotic Resistance Genes in Antibiotic Manufacturing Effluents and Receiving Freshwater Sediments. Frontiers in Microbiology, 2017, 8, 2675.	1.5	40
9	Development of EST-derived SSR Markers with Long-core Repeat in Olive and Their Use for Paternity Testing. Journal of the American Society for Horticultural Science, 2013, 138, 290-296.	0.5	38
10	Transcriptomic Analysis Using Olive Varieties and Breeding Progenies Identifies Candidate Genes Involved in Plant Architecture. Frontiers in Plant Science, 2016, 7, 240.	1.7	25
11	Effects of industrial effluents containing moderate levels of antibiotic mixtures on the abundance of antibiotic resistance genes and bacterial community composition in exposed creek sediments. Science of the Total Environment, 2020, 706, 136001.	3.9	24
12	Small RNAs as Fundamental Players in the Transference of Information During Bacterial Infectious Diseases. Frontiers in Molecular Biosciences, 2020, 7, 101.	1.6	22
13	Genetic Analysis of the Individual Contribution to Virulence of the Type III Effector Inventory of Pseudomonas syringae pv. phaseolicola. PLoS ONE, 2012, 7, e35871.	1.1	21
14	Effect of Roux-en-Y gastric bypass-induced weight loss on the transcriptomic profiling of subcutaneous adipose tissue. Surgery for Obesity and Related Diseases, 2016, 12, 257-263.	1.0	21
15	Small RNAs in cell-to-cell communications during bacterial infection. FEMS Microbiology Letters, 2018, 365, .	0.7	14
16	Molecular Phylogeny Reveals the Past Transoceanic Voyages of Drywood Termites (Isoptera,) Tj ETQq0 0 0 rgBT /	Oyerlock I	10 ₁₂ 50 142

17	Oesophageal squamous cell carcinoma (ESCC): Advances through omics technologies, towards ESCC salivaomics. Drug Discoveries and Therapeutics, 2015, 9, 247-257.	0.6	11
18	Characterization of macrolide resistance in bacteria isolated from macrolide-polluted and unpolluted river sediments and clinical sources in Croatia. Science of the Total Environment, 2020, 749, 142357.	3.9	10

#	Article	IF	CITATIONS
19	The changes in the transcriptomic profiling of subcutaneous adipose tissue after bariatric surgery depend on the insulin resistance state. Surgery for Obesity and Related Diseases, 2018, 14, 1182-1191.	1.0	9
20	Identification of an olive (Olea europaea L.) core collection with a new set of SSR markers. Genetic Resources and Crop Evolution, 2021, 68, 117-133.	0.8	9
21	Comment on: Improvements in humoral immune function and glucolipid metabolism after laparoscopic sleeve gastrectomy in patients with obesity. Surgery for Obesity and Related Diseases, 2019, 15, 1463-1464.	1.0	0