Antonio Manuel Martin Platero

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8532502/publications.pdf

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41 papers 3,746 citations

279798 23 h-index 276875 41 g-index

43 all docs 43 docs citations

43 times ranked

6304 citing authors

#	Article	IF	Citations
1	Beneficial Shifts in the Gut Bacterial Community of Gilthead Seabream (Sparus aurata) Juveniles Supplemented with Allium-Derived Compound Propyl Propane Thiosulfonate (PTSO). Animals, 2022, 12, 1821.	2.3	5
2	Allium-Based Phytobiotic Enhances Egg Production in Laying Hens through Microbial Composition Changes in Ileum and Cecum. Animals, 2021, 11, 448.	2.3	21
3	Allium Extract Implements Weaned Piglet's Productive Parameters by Modulating Distal Gut Microbiota. Antibiotics, 2021, 10, 269.	3.7	14
4	Enterocin Cross-Resistance Mediated by ABC Transport Systems. Microorganisms, 2021, 9, 1411.	3.6	5
5	Synergy of the Bacteriocin AS-48 and Antibiotics against Uropathogenic Enterococci. Antibiotics, 2020, 9, 567.	3.7	13
6	Egg Production in Poultry Farming Is Improved by Probiotic Bacteria. Frontiers in Microbiology, 2019, 10, 1042.	3.5	32
7	Multiple Genome Sequences of Lactobacillus pentosus Strains Isolated from Biofilms on the Skin of Fermented Green Table Olives. Microbiology Resource Announcements, 2019, 8, .	0.6	5
8	Bacterial density rather than diversity correlates with hatching success across different avian species. FEMS Microbiology Ecology, 2018, 94, .	2.7	21
9	High resolution time series reveals cohesive but short-lived communities in coastal plankton. Nature Communications, 2018, 9, 266.	12.8	122
10	Diversity and antimicrobial potential in sea anemone and holothurian microbiomes. PLoS ONE, 2018, 13, e0196178.	2.5	30
11	Draft Genome Sequences of Six Lactobacillus pentosus Strains Isolated from Brines of Traditionally Fermented Spanish-Style Green Table Olives. Genome Announcements, 2018, 6, .	0.8	7
12	A communal catalogue reveals Earth's multiscale microbial diversity. Nature, 2017, 551, 457-463.	27.8	1,942
13	BSocial: Deciphering Social Behaviors within Mixed Microbial Populations. Frontiers in Microbiology, 2017, 8, 919.	3.5	6
14	Nestedness of hoopoes' bacterial communities: symbionts from the uropygial gland to the eggshell. Biological Journal of the Linnean Society, 2016, 118, 763-773.	1.6	9
15	The Hoopoe's Uropygial Gland Hosts a Bacterial Community Influenced by the Living Conditions of the Bird. PLoS ONE, 2015, 10, e0139734.	2.5	29
16	Preening as a Vehicle for Key Bacteria in Hoopoes. Microbial Ecology, 2015, 70, 1024-1033.	2.8	19
17	Eggshell Bacterial Load Is Related to Antimicrobial Properties of Feathers Lining Barn Swallow Nests. Microbial Ecology, 2014, 67, 480-487.	2.8	25
18	Special structures of hoopoe eggshells enhance the adhesion of symbiontâ€carrying uropygial secretion that increase hatching success. Journal of Animal Ecology, 2014, 83, 1289-1301.	2.8	54

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19	Environmental Factors Shape the Community of Symbionts in the Hoopoe Uropygial Gland More than Genetic Factors. Applied and Environmental Microbiology, 2014, 80, 6714-6723.	3.1	25
20	Distribution-Based Clustering: Using Ecology To Refine the Operational Taxonomic Unit. Applied and Environmental Microbiology, 2013, 79, 6593-6603.	3.1	140
21	Avian life history traits influence eggshell bacterial loads: a comparative analysis. Ibis, 2012, 154, 725-737.	1.9	33
22	Antimicrobial Activity and Genetic Profile of Enteroccoci Isolated from Hoopoes Uropygial Gland. PLoS ONE, 2012, 7, e41843.	2.5	36
23	Horizontal transmission of streptococcus mutans in schoolchildren. Medicina Oral, Patologia Oral Y Cirugia Bucal, 2012, 17, e495-e500.	1.7	20
24	The evolution of size of the uropygial gland: mutualistic feather mites and uropygial secretion reduce bacterial loads of eggshells and hatching failures of European birds. Journal of Evolutionary Biology, 2012, 25, 1779-1791.	1.7	60
25	Cognitive skills and bacterial load: comparative evidence of costs of cognitive proficiency in birds. Die Naturwissenschaften, 2012, 99, 111-122.	1.6	19
26	DNA sampling from eggshell swabbing is widely applicable in wild bird populations as demonstrated in 23 species. Molecular Ecology Resources, 2011, 11, 481-493.	4.8	23
27	Typing of bacteriophages by randomly amplified polymorphic DNA (RAPD)-PCR to assess genetic diversity. FEMS Microbiology Letters, 2011, 322, 90-97.	1.8	49
28	Innate humoural immunity is related to eggshell bacterial load of European birds: a comparative analysis. Die Naturwissenschaften, 2011, 98, 807-813.	1.6	23
29	Comparative analysis of microbial DNA extraction protocols for groundwater samples. Analytical Biochemistry, 2011, 416, 240-242.	2.4	9
30	Chelex-based DNA isolation procedure for the identification of microbial communities of eggshell surfaces. Analytical Biochemistry, 2010, 397, 253-255.	2.4	20
31	Number and colour composition of nest lining feathers predict eggshell bacterial community in barn swallow nests: an experimental study. Functional Ecology, 2010, 24, 426-433.	3.6	77
32	Symbiotic bacteria living in the hoopoe's uropygial gland prevent feather degradation. Journal of Experimental Biology, 2009, 212, 3621-3626.	1.7	96
33	Polyphasic study of microbial communities of two Spanish farmhouse goats' milk cheeses from Sierra de Aracena. Food Microbiology, 2009, 26, 294-304.	4.2	68
34	Characterization and safety evaluation of enterococci isolated from Spanish goats' milk cheeses. International Journal of Food Microbiology, 2009, 132, 24-32.	4.7	155
35	Seasonal, sexual and developmental differences in hoopoe <i>Upupa epops</i> preen gland morphology and secretions: evidence for a role of bacteria. Journal of Avian Biology, 2009, 40, 191-205.	1.2	85
36	Symbiotic association between hoopoes and antibioticâ€producing bacteria that live in their uropygial gland. Functional Ecology, 2008, 22, 864-871.	3.6	108

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37	Biodiversity of the microbial community in a Spanish farmhouse cheese as revealed by culture-dependent and culture-independent methods. International Journal of Food Microbiology, 2008, 127, 200-208.	4.7	79
38	Polyphasic Approach to Bacterial Dynamics during the Ripening of Spanish Farmhouse Cheese, Using Culture-Dependent and -Independent Methods. Applied and Environmental Microbiology, 2008, 74, 5662-5673.	3.1	46
39	Fast, convenient, and economical method for isolating genomic DNA from lactic acid bacteria using a modification of the protein "salting-out―procedure. Analytical Biochemistry, 2007, 366, 102-104.	2.4	92
40	Characterisation of Escherichia coli isolated from raw milk cheeses. Annals of Microbiology, 2007, 57, 49-54.	2.6	12
41	Characterization of Antimicrobial Substances Produced by Enterococcus faecalis MRR 10-3, Isolated from the Uropygial Gland of the Hoopoe (Upupa epops). Applied and Environmental Microbiology, 2006, 72, 4245-4249.	3.1	112