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List of Publications by Year in descending order

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1040056 1058476 15 554 9 14 citations g-index h-index papers 16 16 16 827 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Probiotic roles of <i>Lactobacillus &lt; /i&gt; sp. in swine: insights from gut microbiota. Journal of Applied Microbiology, 2017, 122, 554-567.</i>	3.1	181
2	<i>In vitro</i> evaluation of the mucin-adhesion ability and probiotic potential of <i>Lactobacillus mucosae</i> LM1. Journal of Applied Microbiology, 2014, 117, 485-497.	3.1	109
3	Molecular cloning, characterization and comparison of bile salt hydrolases from <i>Lactobacillus johnsonii </i> PF01. Journal of Applied Microbiology, 2013, 114, 121-133.	3.1	61
4	Comparative genomic analysis of Lactobacillus mucosae LM1 identifies potential niche-specific genes and pathways for gastrointestinal adaptation. Genomics, 2019, 111, 24-33.	2.9	47
5	Genome Sequence of Lactobacillus mucosae LM1, Isolated from Piglet Feces. Journal of Bacteriology, 2012, 194, 4766-4766.	2.2	35
6	Quantitative Proteogenomics and the Reconstruction of the Metabolic Pathway in Lactobacillus mucosae LM1. Korean Journal for Food Science of Animal Resources, 2015, 35, 692-702.	1.5	29
7	Carbohydrate-binding specificities of potential probiotic Lactobacillus strains in porcine jejunal (IPEC-J2) cells and porcine mucin. Journal of Microbiology, 2016, 54, 510-519.	2.8	26
8	Complete genome analysis of Lactobacillus fermentum SK152 from kimchi reveals genes associated with its antimicrobial activity. FEMS Microbiology Letters, 2017, 364, .	1.8	20
9	Proteomic View of the Crosstalk between Lactobacillus mucosae and Intestinal Epithelial Cells in Co-culture Revealed by Q Exactive-Based Quantitative Proteomics. Frontiers in Microbiology, 2017, 8, 2459.	3.5	19
10	Comparative exoproteome analyses of Lactobacillus spp. reveals species- and strain-specific proteins involved in their extracellular interaction and probiotic potential. LWT - Food Science and Technology, 2018, 93, 420-426.	5.2	9
11	Exoproteome Perspective on the Bile Stress Response of Lactobacillus johnsonii. Proteomes, 2021, 9, 10.	3.5	8
12	Exploring the Bile Stress Response of Lactobacillus mucosae LM1 through Exoproteome Analysis. Molecules, 2021, 26, 5695.	3.8	5
13	Hacking Commensal Bacteria to Consolidate the Adaptive Mucosal Immune Response in the Gut–Lung Axis: Future Possibilities for SARS-CoV-2 Protection. BioTech, 2022, 11, 3.	2.6	4
14	A molecular systems analysis of HOX PPI networks in hematopoiesis and leukemogenesis. , 2011, , .		0
15	Application of a systematic exoproteogenomic profiling workflow on lactobacillus mucosae LM1., 2015, , .		O