Zhengyuan Zhai

List of Publications by Year in descending order

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623734 526287 34 761 14 27 citations g-index h-index papers 36 36 36 926 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Transcriptome analysis revealed growth phase-associated changes of a centenarian-originated probiotic Bifidobacterium animalis subsp. lactis A6. BMC Microbiology, 2022, 22, 61.	3.3	2
2	<scp>L</scp> -Malic Acid Protects <i>Lacticaseibacillus paracasei</i> L9 from Glycodeoxycholic Acid Stress via the Malolactic Enzyme Pathway. Journal of Agricultural and Food Chemistry, 2022, 70, 9007-9016.	5.2	4
3	Homologous Over-Expression of Chain Length Determination Protein EpsC Increases the Molecular Weight of Exopolysaccharide in Streptococcus thermophilus 05-34. Frontiers in Microbiology, 2021, 12, 696222.	3.5	4
4	Lactobacillus rhamnosus GG Derived Extracellular Vesicles Modulate Gut Microbiota and Attenuate Inflammatory in DSS-Induced Colitis Mice. Nutrients, 2021, 13, 3319.	4.1	54
5	Establishment and Evaluation of a Loop-Mediated Isothermal Amplification Assay for Rapid Detection of Pseudomonas fluorescens in Raw Milk. Frontiers in Microbiology, 2021, 12, 810511.	3.5	3
6	Global transcriptomic analysis of Lactobacillus plantarum CAUH2 in response to hydrogen peroxide stress. Food Microbiology, 2020, 87, 103389.	4.2	31
7	A Novel Major Pilin Subunit Protein FimM Is Involved in Adhesion of Bifidobacterium longum BBMN68 to Intestinal Epithelial Cells. Frontiers in Microbiology, 2020, 11, 590435.	3.5	10
8	Expression, purification, and characterization of phospholipase B1 from Candida albicans in Escherichia coli. 3 Biotech, 2020, 10, 538.	2.2	1
9	Combined transcriptomic and proteomic analysis of the response to bile stress in a centenarian-originated probiotic Lactobacillus salivarius Ren. Food Research International, 2020, 137, 109331.	6.2	10
10	The N-terminal domain of rhamnosyltransferase EpsF influences exopolysaccharide chain length determination in <i>Streptococcus thermophilus</i> 05-34. PeerJ, 2020, 8, e8524.	2.0	6
11	Synergy between Probiotic <i>Lactobacillus casei</i> and Milk to Maintain Barrier Integrity of Intestinal Epithelial Cells. Journal of Agricultural and Food Chemistry, 2019, 67, 1955-1962.	5.2	20
12	Complete genome sequencing of Lactobacillus plantarum CAUH2 reveals a novel plasmid pCAUH203 associated with oxidative stress tolerance. 3 Biotech, 2019, 9, 116.	2.2	8
13	Low-fat yogurt alleviates the pro-inflammatory cytokine IL- \hat{I}^2 -induced intestinal epithelial barrier dysfunction. Journal of Dairy Science, 2019, 102, 976-984.	3.4	14
14	The MarR Family Regulator BmrR Is Involved in Bile Tolerance of <i>Bifidobacterium longum</i> BBMN68 via Controlling the Expression of an ABC Transporter. Applied and Environmental Microbiology, 2019, 85, .	3.1	6
15	<i>Lactobacillus plantarum</i> bacteriocin is associated with intestinal and systemic improvements in diet-induced obese mice and maintains epithelial barrier integrity <i>in vitro</i> . Gut Microbes, 2019, 10, 382-397.	9.8	90
16	Global Transcriptomic Analysis and Function Identification of Malolactic Enzyme Pathway of Lactobacillus paracasei L9 in Response to Bile Stress. Frontiers in Microbiology, 2018, 9, 1978.	3.5	27
17	Characterization of a Cryptic Rolling-Circle Replication Plasmid pMK8 from Enterococcus durans 1–8. Current Microbiology, 2018, 75, 1198-1205.	2.2	5
18	Characterization of a Rolling-Circle Replication Plasmid pM411 from Lactobacillus plantarum 1–3. Current Microbiology, 2016, 73, 820-826.	2,2	12

#	Article	IF	CITATIONS
19	Complete genome sequence of Lactobacillus helveticus CAUH18, a potential probiotic strain originated from koumiss. Journal of Biotechnology, 2016, 224, 18-19.	3.8	17
20	Functional role of pyruvate kinase from Lactobacillus bulgaricus in acid tolerance and identification of its transcription factor by bacterial one-hybrid. Scientific Reports, 2015, 5, 17024.	3.3	20
21	Functional role of <i>oppA</i> encoding an oligopeptide-binding protein from <i>Lactobacillus salivarius</i> Ren in bile tolerance. Journal of Industrial Microbiology and Biotechnology, 2015, 42, 1167-1174.	3.0	43
22	Proteomic characterization of the acid tolerance response in <i><scp>L</scp>actobacillus delbrueckii</i> subsp. <i>bulgaricus</i> â€ <scp>CAUH1</scp> and functional identification of a novel acid stressâ€related transcriptional regulator <scp>L</scp> db0677. Environmental Microbiology, 2014, 1524-1537.	3.8	61
23	Characterization of the cryptic plasmid pWCZ from Lactobacillus paracasei WCZ isolated from silage. Annals of Microbiology, 2014, 64, 809-814.	2.6	0
24	Integrated Transcriptomic and Proteomic Analysis of the Bile Stress Response in a Centenarian-originated Probiotic Bifidobacterium longum BBMN68. Molecular and Cellular Proteomics, 2014, 13, 2558-2572.	3.8	76
25	Coexpression of the Superoxide Dismutase and the Catalase Provides Remarkable Oxidative Stress Resistance in Lactobacillus rhamnosus. Journal of Agricultural and Food Chemistry, 2011, 59, 3851-3856.	5.2	42
26	A novel vector for lactic acid bacteria that uses a bile salt hydrolase gene as a potential food-grade selection marker. Journal of Biotechnology, 2011, 152, 49-53.	3.8	14
27	Characterization of a rolling-circle replication plasmid pXY3 from Lactobacillus plantarum XY3. Plasmid, 2010, 64, 36-40.	1.4	11
28	Identification of the bacterial biodiversity in koumiss by denaturing gradient gel electrophoresis and species-specific polymerase chain reaction. Journal of Dairy Science, 2010, 93, 1926-1933.	3.4	44
29	Characterization of a novel rolling-circle replication plasmid pYSI8 from Lactobacillus sakei YSI8. Plasmid, 2009, 62, 30-34.	1.4	13
30	Functional analysis of the plasmid pM4 replicon from Lactobacillus plantarum M4: Determination of the minimal replicon and functionality identification of the putative sso. Plasmid, 2009, 62, 166-171.	1.4	6
31	Characterization of a Rolling-Circle Replication Plasmid pLR1 from Lactobacillus plantarum LR1. Current Microbiology, 2009, 58, 106-110.	2.2	11
32	Characterization and application of an anti-Listeria bacteriocin produced by Pediococcus pentosaceus 05-10 isolated from Sichuan Pickle, a traditionally fermented vegetable product from China. Food Control, 2009, 20, 1030-1035.	5.5	63
33	Characterization of a cryptic plasmid pM4 from <i>Lactobacillus plantarum</i> M4. FEMS Microbiology Letters, 2008, 285, 183-187.	1.8	20
34	Expression, purification and characterization of pectin methylesterase inhibitor from kiwi fruit in Escherichia coli. Protein Expression and Purification, 2008, 60, 221-224.	1.3	13