List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4294704/publications.pdf Version: 2024-02-01



IFSUS A FEDDO

#	Article	lF	CITATIONS
1	XAC4296 Is a Multifunctional and Exclusive Xanthomonadaceae Gene Containing a Fusion of Lytic Transglycosylase and Epimerase Domains. Microorganisms, 2022, 10, 1008.	3.6	0
2	Identification of novel mRNA isoforms associated with meat tenderness using RNA sequencing data in beef cattle. Meat Science, 2021, 173, 108378.	5.5	17
3	Riboswitch theo/metE as a Transcription Regulation Tool for Xanthomonas citri subsp. citri. Microorganisms, 2021, 9, 329.	3.6	2
4	Potential Bioinoculants for Sustainable Agriculture Prospected from Ferruginous Caves of the Iron Quadrangle/Brazil. Sustainability, 2021, 13, 9354.	3.2	2
5	Transcriptional changes involved in kumquat (Fortunella spp) defense response to Xanthomonas citri subsp. citri in early stages of infection. Physiological and Molecular Plant Pathology, 2021, 116, 101729.	2.5	4
6	Gene expression profiling and identification of hub genes in Nellore cattle with different marbling score levels. Genomics, 2020, 112, 873-879.	2.9	31
7	Transcriptome profiling of muscle in Nelore cattle phenotypically divergent for the ribeye muscle area. Genomics, 2020, 112, 1257-1263.	2.9	13
8	Bactericidal Effect of Entomopathogenic Bacterium Pseudomonas entomophila Against Xanthomonas citri Reduces Citrus Canker Disease Severity. Frontiers in Microbiology, 2020, 11, 1431.	3.5	12
9	CitrusKB: a comprehensive knowledge base for transcriptome and interactome of <i>Citrus</i> spp. infected by <i>Xanthomonas citri</i> subsp. <i>citri</i> at different infection stages. Database: the Journal of Biological Databases and Curation, 2020, 2020, .	3.0	2
10	mCherry fusions enable the subcellular localization of periplasmic and cytoplasmic proteins in Xanthomonas sp PLoS ONE, 2020, 15, e0236185.	2.5	4
11	Spliced genes in muscle from Nelore Cattle and their association with carcass and meat quality. Scientific Reports, 2020, 10, 14701.	3.3	21
12	A Genomic and Transcriptomic Overview of MATE, ABC, and MFS Transporters in Citrus sinensis Interaction with Xanthomonas citri subsp. citri. Plants, 2020, 9, 794.	3.5	9
13	Use of gene expression profile to identify potentially relevant transcripts to myofibrillar fragmentation index trait. Functional and Integrative Genomics, 2020, 20, 609-619.	3.5	10
14	Complete genome sequence and analysis of Alcaligenes faecalis strain Mc250, a new potential plant bioinoculant. PLoS ONE, 2020, 15, e0241546.	2.5	9
15	24 Structural variants affecting mRNAs isoforms splice sites associated with marbling in Nellore cattle. Journal of Animal Science, 2020, 98, 24-25.	0.5	0
16	Prediction of hub genes associated with intramuscular fat content in Nelore cattle. BMC Genomics, 2019, 20, 520.	2.8	24
17	Analyses of Seven New Genomes of Xanthomonas citri pv. aurantifolii Strains, Causative Agents of Citrus Canker B and C, Show a Reduced Repertoire of Pathogenicity-Related Genes. Frontiers in Microbiology, 2019, 10, 2361.	3.5	14
18	Gene expression analysis identifies hypothetical genes that may be critical during the infection process of Xanthomonas citri subsp. citri. Electronic Journal of Biotechnology, 2019, 42, 30-41.	2.2	2

#	Article	IF	CITATIONS
19	Origin and diversification of Xanthomonas citri subsp. citri pathotypes revealed by inclusive phylogenomic, dating, and biogeographic analyses. BMC Genomics, 2019, 20, 700.	2.8	33
20	Serratia liquefaciens FG3 isolated from a metallophyte plant sheds light on the evolution and mechanisms of adaptive traits in extreme environments. Scientific Reports, 2019, 9, 18006.	3.3	10
21	Gene Expression in the Salivary Gland of Rhipicephalus (Boophilus) microplus Fed on Tick-Susceptible and Tick-Resistant Hosts. Frontiers in Cellular and Infection Microbiology, 2019, 9, 477.	3.9	12
22	Detection and identification of <i>Xanthomonas</i> pathotypes associated with citrus diseases using comparative genomics and multiplex PCR. PeerJ, 2019, 7, e7676.	2.0	10
23	Comparative proteomic analysis of <i>Xanthomonas citri</i> ssp. <i>citri</i> periplasmic proteins reveals changes in cellular envelope metabolism during <i>in vitro</i> pathogenicity induction. Molecular Plant Pathology, 2018, 19, 143-157.	4.2	11
24	Biotechnological potential of plant growth-promoting bacteria from the roots and rhizospheres of endemic plants in ironstone vegetation in southeastern Brazil. World Journal of Microbiology and Biotechnology, 2018, 34, 156.	3.6	15
25	Transposons and pathogenicity in <i>Xanthomonas</i> : acquisition of murein lytic transglycosylases by Tn <i>Xax1</i> enhances <i>Xanthomonas citri</i> subsp. <i>citri</i> 306 virulence and fitness. PeerJ, 2018, 6, e6111.	2.0	8
26	Proteomics-based identification of differentially abundant proteins reveals adaptation mechanisms of Xanthomonas citri subsp. citri during Citrus sinensis infection. BMC Microbiology, 2017, 17, 155.	3.3	18
27	Differences in global gene expression in muscle tissue of Nellore cattle with divergent meat tenderness. BMC Genomics, 2017, 18, 945.	2.8	32
28	Quorum Sensing, Its Role in Virulence and Symptomatology in Bacterial Citrus Canker. , 2017, , .		5
29	Identification of New Genes Related to Virulence of <i>Xanthomonas axonopodis</i> Pv. <i>Citri</i> during Citrus Host Interactions. Advances in Microbiology, 2017, 07, 22-46.	0.6	3
30	Unravelling potential virulence factor candidates in <i>Xanthomonas citri</i> . subsp. <i>citri</i> by secretome analysis. PeerJ, 2016, 4, e1734.	2.0	35
31	Proline accumulation in sugarcane roots subjected to drought conditions. CientÃfica, 2016, 44, 592.	0.2	5
32	A TALE of Transposition: Tn <i>3</i> -Like Transposons Play a Major Role in the Spread of Pathogenicity Determinants of Xanthomonas citri and Other Xanthomonads. MBio, 2015, 6, e02505-14.	4.1	43
33	Chemotactic signal transduction and phosphate metabolism as adaptive strategies during citrus canker induction by Xanthomonas citri. Functional and Integrative Genomics, 2015, 15, 197-210.	3.5	39
34	Expression of genes related to mitochondrial function in Nellore cattle divergently ranked on residual feed intake. Molecular Biology Reports, 2015, 42, 559-565.	2.3	20
35	Type IV Secretion System Is Not Involved in Infection Process in Citrus. International Journal of Microbiology, 2014, 2014, 1-9.	2.3	16
36	Comparative proteomic analysis reveals that T3SS, Tfp, and xanthan gum are key factors in initial stages of Citrus sinensis infection by Xanthomonas citri subsp. citri. Functional and Integrative Genomics, 2014, 14, 205-217.	3.5	30

#	Article	IF	CITATIONS
37	Study of whole genome linkage disequilibrium in Nellore cattle. BMC Genomics, 2013, 14, 305.	2.8	106
38	The Genome of Anopheles darlingi , the main neotropical malaria vector. Nucleic Acids Research, 2013, 41, 7387-7400.	14.5	102
39	Evidence for Reductive Genome Evolution and Lateral Acquisition of Virulence Functions in Two Corynebacterium pseudotuberculosis Strains. PLoS ONE, 2011, 6, e18551.	2.5	75
40	Selection and validation of reference genes for gene expression studies by reverse transcription quantitative PCR in Xanthomonas citri subsp. citri during infection of Citrus sinensis. Biotechnology Letters, 2011, 33, 1177-1184.	2.2	31
41	Novel insights into the genomic basis of citrus canker based on the genome sequences of two strains of Xanthomonas fuscans subsp. aurantifolii. BMC Genomics, 2010, 11, 238.	2.8	102
42	Development and validation of a Xanthomonas axonopodis pv. citri DNA microarray platform (XACarray) generated from the shotgun libraries previously used in the sequencing of this bacterial genome. BMC Research Notes, 2010, 3, 150.	1.4	3
43	Proteome of the phytopathogen Xanthomonas citri subsp. citri: a global expression profile. Proteome Science, 2010, 8, 55.	1.7	28
44	Sensitivity evaluation of a single-step PCR assay using Ehrlichia canis p28 gene as a target and its application in diagnosis of canine ehrlichiosis. Brazilian Journal of Veterinary Parasitology, 2010, 19, 75-79.	0.7	16
45	New genes of Xanthomonas citri subsp. citri involved in pathogenesis and adaptation revealed by a transposon-based mutant library. BMC Microbiology, 2009, 9, 12.	3.3	67
46	The Brazilian experience of sugarcane ethanol industry. In Vitro Cellular and Developmental Biology - Plant, 2009, 45, 372-381.	2.1	87
47	Cellular and molecular characterization of an embryonic cell line (BME26) from the tick Rhipicephalus (Boophilus) microplus. Insect Biochemistry and Molecular Biology, 2008, 38, 568-580.	2.7	57
48	Structure and genetic relationships between Brazilian naturalized and exotic purebred goat domestic goat (Capra hircus) breeds based on microsatellites. Genetics and Molecular Biology, 2007, 30, 356-363.	1.3	17
49	Genes diferencialmente expressos em cana-de-açúcar inoculada com Xanthomonas albilineans, o agente causal da escaldadura da folha. Summa Phytopathologica, 2006, 32, 328-338.	0.1	4
50	The Complete Nucleotide Sequence and Genomic Organization of Citrus Leprosis Associated Virus, Cytoplasmatic type (CiLV-C). Virus Genes, 2006, 32, 289-298.	1.6	90
51	Identification and Genomic Characterization of a New Virus (Tymoviridae Family) Associated with Citrus Sudden Death Disease. Journal of Virology, 2005, 79, 3028-3037.	3.4	76
52	Brain and hepatic Hsp70 protein levels in heat-acclimated broiler chickens during heat stress. Brazilian Journal of Poultry Science, 2004, 6, 201-206.	0.7	20
53	Hepatic mRNA expression and plasma levels of insulin-like growth factor-I (IGF-I) in broiler chickens selected for different growth rates. Genetics and Molecular Biology, 2004, 27, 39-44.	1.3	12
54	Expression of heat shock protein in broiler embryo tissues after acute cold or heat stress. Molecular Reproduction and Development, 2004, 67, 172-177.	2.0	46

#	Article	IF	CITATIONS
55	The zymogen-enteropeptidase system: A practical approach to study the regulation of enzyme activity by proteolytic cleavage. Biochemistry and Molecular Biology Education, 2004, 32, 45-48.	1.2	4
56	COMPARATIVE GENOMICS ANALYSES OF CITRUS-ASSOCIATED BACTERIA. Annual Review of Phytopathology, 2004, 42, 163-184.	7.8	57
57	Conditions affecting production of functional muscle recombinant α-tropomyosin in Saccharomyces cerevisiae. Protein Expression and Purification, 2003, 30, 105-111.	1.3	4
58	Analysis and Functional Annotation of an Expressed Sequence Tag Collection for Tropical Crop Sugarcane. Genome Research, 2003, 13, 2725-2735.	5.5	254
59	Polymorphism analysis of the hsp70 stress gene in Broiler chickens (Gallus gallus) of different breeds. Genetics and Molecular Biology, 2003, 26, 275-281.	1.3	29
60	Carbohydrate metabolism of Xylella fastidiosa: Detection of glycolytic and pentose phosphate pathway enzymes and cloning and expression of the enolase gene. Genetics and Molecular Biology, 2003, 26, 203-211.	1.3	5
61	Performance and hormonal profile in broiler chickens fed with different energy levels during post restriction period. Pesquisa Agropecuaria Brasileira, 2003, 38, 697-702.	0.9	22
62	Respostas Fisiológicas Associadas à Termotolerância em Pintos de Corte de Duas Linhagens por Exposição a Altas Temperaturas. Revista Brasileira De Zootecnia, 2002, 31, 79-85.	0.8	8
63	Resistência à SÃndrome AscÃtica, Competência Homeotérmica e NÃveis de Hsp70 no Coração e Pulmão Frangos de Corte. Revista Brasileira De Zootecnia, 2002, 31, 1442-1450.	de.8	3
64	High-Level Production of Functional Muscle α-Tropomyosin in Pichia pastoris. Biochemical and Biophysical Research Communications, 2001, 284, 955-960.	2.1	9
65	Efeito do estresse térmico agudo sobre os nÃveis da proteÃna e RNA mensageiro da Hsp70, em fÃgado e cérebro de pintos de corte de diferentes linhagens. Revista Brasileira De Zootecnia, 2001, 30, 1506-1513.	0.8	3
66	Influence of Dietary Energy Level on Hepatic70-kDa Heat Shock Protein Expression in Broiler Chickens Submitted to Acute Heat Stress. Brazilian Journal of Poultry Science, 2000, 2, 259-266.	0.7	2
67	Sexagem de embriões bovinos fecundados in vitro pela técnica de PCR multiplex. Brazilian Journal of Veterinary Research and Animal Science, 2000, 37, 0-0.	0.2	1
68	Effect of heat stress or lipopolysaccharide (E. coli) injection on HSP70 levels in the liver and brain of adrenalectomized rats. Journal of Thermal Biology, 1999, 24, 429-432.	2.5	1
69	Digestibilidade e ganhos em peso de ovinos alimentados com diferentes fontes de volumosos e recebendo somatotropina recombinante bovina (rBST). Revista Brasileira De Zootecnia, 1999, 28, 1102-1108.	0.8	2
70	Concerted Action of the High Affinity Calcium Binding Sites in Skeletal Muscle Troponin C. Journal of Biological Chemistry, 1995, 270, 9770-9777.	3.4	38
71	Cloning and expression of chicken skeletal muscle troponin I in <i>Escherichia coli</i> : The role of rare codons on the expression level. Protein Science, 1993, 2, 1053-1056.	7.6	28
72	The complete sequence of a chicken-muscle cDNA encoding the acidic ribosomal protein P1. FEBS Journal, 1988, 177, 513-516.	0.2	7

#	Article	IF	CITATIONS
73	Cold-acclimation improves cold-tolerance of diabetic rats. Comparative Biochemistry and Physiology A, Comparative Physiology, 1987, 88, 425-429.	0.6	4
74	Calcium/calmodulin regulation of ATPase activity and endogenous phosphorylation of mammalian brain actomyosin. Biochemical and Biophysical Research Communications, 1987, 145, 1217-1224.	2.1	2
75	Effects of acute cold exposure on rectal temperature, blood glucose and plasma free fatty acids in alloxan-diabetic rats. Comparative Biochemistry and Physiology A, Comparative Physiology, 1986, 85, 63-65.	0.6	9
76	Isolation and purification of actomyosin ATPase from mammalian brain. Journal of Neuroscience Methods, 1986, 16, 47-58.	2.5	9