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List of Articles by Year in descending order

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h-index

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citing authors

#	ARTICLE	IF	CITATIONS
1	PXR-dependant dysregulation of glucose metabolism induced by chronic exposure to NOAEL-level pesticide cocktail in mice. <i>Environment International</i> , 2025, 202, 109697.	10.2	1
2	More than a mutagenic Aflatoxin B1 precursor: The multiple cellular targets of Versicolorin A revealed by global gene expression analysis. <i>Environmental Pollution</i> , 2024, 363, 125138.	7.7	1
3	Effect of <i>Streptomyces roseolus</i> Cell-Free Supernatants on the Fungal Development, Transcriptome, and Aflatoxin B1 Production of <i>Aspergillus flavus</i> . <i>Toxins</i> , 2023, 15, 428.	3.8	5
4	CAR Protects Females from Diet-Induced Steatosis and Associated Metabolic Disorders. <i>Cells</i> , 2023, 12, 2218.	4.7	4
5	Integrative study of diet-induced mouse models of NAFLD identifies PPAR α as a sexually dimorphic drug target. <i>Gut</i> , 2022, 71, 807-821.	16.8	75
6	Intestinal toxicity of the new type A trichothecenes, NX and 3ANX. <i>Chemosphere</i> , 2022, 288, 132415.	8.2	30
7	Tissular Genomic Responses to Oral FB1 Exposure in Pigs. <i>Toxins</i> , 2022, 14, 83.	3.8	8
8	The pregnane X receptor drives sexually dimorphic hepatic changes in lipid and xenobiotic metabolism in response to gut microbiota in mice. <i>Microbiome</i> , 2021, 9, .	11.5	27
9	Comparative sensitivity of proliferative and differentiated intestinal epithelial cells to the food contaminant, deoxynivalenol. <i>Environmental Pollution</i> , 2021, 277, 116818.	7.7	28
10	Statistical Integration of Omics Data Increases Biological Knowledge Extracted from Metabolomics Data: Application to Intestinal Exposure to the Mycotoxin Deoxynivalenol. <i>Metabolites</i> , 2021, 11, 407.	3.4	9
11	Chronic exposure to Cytolethal Distending Toxin (CDT) promotes a cGAS-dependent type I interferon response. <i>Cellular and Molecular Life Sciences</i> , 2021, 78, 6319-6335.	5.5	18
12	The Solvent Dimethyl Sulfoxide Affects Physiology, Transcriptome and Secondary Metabolism of <i>Aspergillus flavus</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 1055.	3.5	12
13	The <i>brlA</i> Gene Deletion Reveals That Patulin Biosynthesis Is Not Related to Conidiation in <i>Penicillium expansum</i> . <i>International Journal of Molecular Sciences</i> , 2020, 21, 6660.	4.4	18
14	Versicolorin A, a precursor in aflatoxins biosynthesis, is a food contaminant toxic for human intestinal cells. <i>Environment International</i> , 2020, 137, 105568.	10.2	27
15	Hepatocyte-specific deletion of Ppar α promotes NAFLD in the context of obesity. <i>Scientific Reports</i> , 2020, 10, .	3.4	135
16	Gene Expression Profiling Reveals that PXR Activation Inhibits Hepatic PPAR α Activity and Decreases FGF21 Secretion in Male C57Bl6/J Mice. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3767.	4.4	17
17	Haem iron reshapes colonic luminal environment: impact on mucosal homeostasis and microbiome through aldehyde formation. <i>Microbiome</i> , 2019, 7, .	11.5	55
18	Dimorphic metabolic and endocrine disorders in mice lacking the constitutive androstane receptor. <i>Scientific Reports</i> , 2019, 9, .	3.4	13

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19	The GMO90+ Project: Absence of Evidence for Biologically Meaningful Effects of Genetically Modified Maize-based Diets on Wistar Rats After 6-Months Feeding Comparative Trial. <i>Toxicological Sciences</i> , 2019, 168, 315-338.	3.8	14
20	Insights into the role of hepatocyte PPAR α activity in response to fasting. <i>Molecular and Cellular Endocrinology</i> , 2018, 471, 75-88.	3.4	63
21	Metabolic Effects of a Chronic Dietary Exposure to a Low-Dose Pesticide Cocktail in Mice: Sexual Dimorphism and Role of the Constitutive Androstane Receptor. <i>Environmental Health Perspectives</i> , 2018, 126, .	8.3	126
22	<i>Saccharomyces cerevisiae</i> Boulardii Reduces the Deoxynivalenol-Induced Alteration of the Intestinal Transcriptome. <i>Toxins</i> , 2018, 10, 199.	3.8	31
23	The protective role of liver X receptor (LXR) during fumonisin B1-induced hepatotoxicity. <i>Archives of Toxicology</i> , 2018, 93, 505-517.	5.8	40
24	The sunflower genome provides insights into oil metabolism, flowering and Asterid evolution. <i>Nature</i> , 2017, 546, 148-152.	38.0	749
25	A Specific ChREBP and PPAR α Cross-Talk Is Required for the Glucose-Mediated FGF21 Response. <i>Cell Reports</i> , 2017, 21, 403-416.	6.3	123
26	Identification of Signaling Pathways Targeted by the Food Contaminant FB1: Transcriptome and Kinome Analysis of Samples from Pig Liver and Intestine. <i>Molecular Nutrition and Food Research</i> , 2017, 61, .	4.1	38
27	Intestinal toxicity of the type B trichothecene mycotoxin fusarenon-X: whole transcriptome profiling reveals new signaling pathways. <i>Scientific Reports</i> , 2017, 7, .	3.4	34
28	Comparing the intestinal transcriptome of Meishan and Large White piglets during late fetal development reveals genes involved in glucose and lipid metabolism and immunity as valuable clues of intestinal maturity. <i>BMC Genomics</i> , 2017, 18, .	3.3	16
29	Time course study of the response to LPS targeting the pig immune gene networks. <i>BMC Genomics</i> , 2017, 18, .	3.3	17
30	Deciphering the Anti-Aflatoxinogenic Properties of Eugenol Using a Large-Scale q-PCR Approach. <i>Toxins</i> , 2016, 8, 123.	3.8	56
31	Patulin is a cultivar-dependent aggressiveness factor favouring the colonization of apples by <i>Aspergillus expansum</i> . <i>Molecular Plant Pathology</i> , 2016, 17, 220-230.	5.1	102
32	Liver PPAR α is crucial for whole-body fatty acid homeostasis and is protective against NAFLD. <i>Gut</i> , 2016, 65, 1202-1214.	16.8	676
33	Foxo3 Transcription Factor Drives Pathogenic T H 1 Differentiation by Inducing the Expression of Eomes. <i>Immunity</i> , 2016, 45, 774-787.	22.6	65
34	Microbial biotransformation of DON: molecular basis for reduced toxicity. <i>Scientific Reports</i> , 2016, 6, .	3.4	150
35	Hepatic circadian clock oscillators and nuclear receptors integrate microbiome-derived signals. <i>Scientific Reports</i> , 2016, 6, .	3.4	112
36	Patulin transformation products and last intermediates in its biosynthetic pathway, E- and Z-ascladiol, are not toxic to human cells. <i>Archives of Toxicology</i> , 2016, 91, 2455-2467.	5.8	92

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37	Exploring transcriptomic diversity in muscle revealed that cellular signaling pathways mainly differentiate five Western porcine breeds. <i>BMC Genomics</i> , 2015, 16, .	3.3	11
38	Time course of the response to ACTH in pig: biological and transcriptomic study. <i>BMC Genomics</i> , 2015, 16, .	3.3	25
39	Pattern recognition receptors in the gut: analysis of their expression along the intestinal tract and the crypt/villus axis. <i>Physiological Reports</i> , 2015, 3, e12225.	1.6	49
40	Intestinal toxicity of the masked mycotoxin deoxynivalenol-3- β -D-glucoside. <i>Archives of Toxicology</i> , 2015, 90, 2037-2046.	5.8	115
41	Muscle transcriptomic investigation of late fetal development identifies candidate genes for piglet maturity. <i>BMC Genomics</i> , 2014, 15, .	3.3	35
42	Src is activated by the nuclear receptor peroxisome proliferator-activated receptor β in ultraviolet radiation-induced skin cancer. <i>EMBO Molecular Medicine</i> , 2014, 6, 80-98.	7.1	59
43	Sequencing, physical organization and kinetic expression of the patulin biosynthetic gene cluster from <i>Penicillium expansum</i> . <i>International Journal of Food Microbiology</i> , 2014, 189, 51-60.	4.9	106
44	Combined linkage and association mapping of flowering time in Sunflower (<i>Helianthus annuus</i> L.). <i>Theoretical and Applied Genetics</i> , 2013, 126, 1337-1356.	3.7	53
45	Consensus mapping of major resistance genes and independent QTL for quantitative resistance to sunflower downy mildew. <i>Theoretical and Applied Genetics</i> , 2012, 125, 909-920.	3.7	50
46	Genetic analysis of phytosterol content in sunflower seeds. <i>Theoretical and Applied Genetics</i> , 2012, 125, 1589-1601.	3.7	48
47	Transcriptomic data of piglet blood compartments with 3° mRNA sequencing. <i>Data in Brief</i> , 0, 66, 112679.	1.1	0