Robert G Nichols

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Gut microbiota and intestinal FXR mediate the clinical benefits of metformin. Nature Medicine, 2018, 24, 1919-1929.	15.2	632
2	Intestinal farnesoid X receptor signaling promotes nonalcoholic fatty liver disease. Journal of Clinical Investigation, 2015, 125, 386-402.	3.9	517
3	Intermittent Fasting Promotes White Adipose Browning and Decreases Obesity by Shaping the GutÂMicrobiota. Cell Metabolism, 2017, 26, 672-685.e4.	7.2	427
4	Intestine farnesoid X receptor agonist and the gut microbiota activate Gâ€protein bile acid receptorâ€1 signaling to improve metabolism. Hepatology, 2018, 68, 1574-1588.	3.6	348
5	Persistent Organic Pollutants Modify Gut Microbiota–Host Metabolic Homeostasis in Mice Through Aryl Hydrocarbon Receptor Activation. Environmental Health Perspectives, 2015, 123, 679-688.	2.8	262
6	The microbiome modulating activity of bile acids. Gut Microbes, 2020, 11, 979-996.	4.3	124
7	Farnesoid X Receptor Signaling Shapes the Gut Microbiota and Controls Hepatic Lipid Metabolism. MSystems, 2016, 1, .	1.7	95
8	Berberine Directly Affects the Gut Microbiota to Promote Intestinal Farnesoid X Receptor Activation. Drug Metabolism and Disposition, 2019, 47, 86-93.	1.7	84
9	Metabolomics Reveals that Aryl Hydrocarbon Receptor Activation by Environmental Chemicals Induces Systemic Metabolic Dysfunction in Mice. Environmental Science & Technology, 2015, 49, 8067-8077.	4.6	80
10	The relationship between the gut microbiome and host gene expression: a review. Human Genetics, 2021, 140, 747-760.	1.8	78
11	Dietary broccoli impacts microbial community structure and attenuates chemically induced colitis in mice in an Ah receptor dependent manner. Journal of Functional Foods, 2017, 37, 685-698.	1.6	62
12	Vitamin A deficiency in mice alters host and gut microbial metabolism leading to altered energy homeostasis. Journal of Nutritional Biochemistry, 2018, 54, 28-34.	1.9	60
13	Metatranscriptomic Analysis of the Mouse Gut Microbiome Response to the Persistent Organic Pollutant 2,3,7,8-Tetrachlorodibenzofuran. Metabolites, 2020, 10, 1.	1.3	55
14	Expression of the aryl hydrocarbon receptor contributes to the establishment of intestinal microbial community structure in mice. Scientific Reports, 2016, 6, 33969.	1.6	54
15	Modulation of Colon Cancer by Nutmeg. Journal of Proteome Research, 2015, 14, 1937-1946.	1.8	44
16	Vitamin D Regulates the Microbiota to Control the Numbers of RORÎ ³ t/FoxP3+ Regulatory T Cells in the Colon. Frontiers in Immunology, 2019, 10, 1772.	2.2	44
17	Perfluorooctane sulfonate alters gut microbiota-host metabolic homeostasis in mice. Toxicology, 2020, 431, 152365.	2.0	43
18	The aryl hydrocarbon receptor as a moderator of host-microbiota communication. Current Opinion in Toxicology, 2017, 2, 30-35.	2.6	28

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#	Article	IF	CITATIONS
19	Prebiotic effects of white button mushroom (Agaricus bisporus) feeding on succinate and intestinal gluconeogenesis in C57BL/6 mice. Journal of Functional Foods, 2018, 45, 223-232.	1.6	28
20	Metabolic impact of persistent organic pollutants on gut microbiota. Gut Microbes, 2020, 12, 1848209.	4.3	22
21	A Quantitative HILIC–MS/MS Assay of the Metabolic Response of Huh-7 Cells Exposed to 2,3,7,8-Tetrachlorodibenzo-p-Dioxin. Metabolites, 2019, 9, 118.	1.3	12
22	The aryl hydrocarbon receptor activates ceramide biosynthesis in mice contributing to hepatic lipogenesis. Toxicology, 2021, 458, 152831.	2.0	12
23	Omics Approaches To Probe Microbiota and Drug Metabolism Interactions. Chemical Research in Toxicology, 2016, 29, 1987-1997.	1.7	7
24	Structural and Functional Analysis of the Gut Microbiome for Toxicologists. Current Protocols in Toxicology / Editorial Board, Mahin D Maines (editor-in-chief) [et Al], 2018, 78, e54.	1.1	6
25	Multiplatform Physiologic and Metabolic Phenotyping Reveals Microbial Toxicity. MSystems, 2018, 3, .	1.7	5