

Anniina Keskitalo

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

862
citations

686830

13
h-index

752256

20
g-index

20
all docs

20
docs citations

20
times ranked

1720
citing authors

#	ARTICLE	IF	CITATIONS
1	Infant fecal microbiota composition and attention to emotional faces.. Emotion, 2022, 22, 1159-1170.	1.5	14
2	An Infancy-Onset 20-Year Dietary Counselling Intervention and Gut Microbiota Composition in Adulthood. Nutrients, 2022, 14, 2667.	1.7	2
3	Vascular Adhesion Protein 1 Mediates Gut Microbial Flagellin-Induced Inflammation, Leukocyte Infiltration, and Hepatic Steatosis. Sci, 2021, 3, 13.	1.8	3
4	Gut microbiota diversity but not composition is related to saliva cortisol stress response at the age of 2.5 months. Stress, 2021, 24, 551-560.	0.8	18
5	Maternal prenatal psychological distress and hair cortisol levels associate with infant fecal microbiota composition at 2.5 months of age. Psychoneuroendocrinology, 2020, 119, 104754.	1.3	40
6	Reply to the Letter to the Editor: Gut microbiota composition is associated with temperament traits in infants. Brain, Behavior, and Immunity, 2019, 81, 671-672.	2.0	1
7	Gut microbiota composition is associated with temperament traits in infants. Brain, Behavior, and Immunity, 2019, 80, 849-858.	2.0	91
8	Blocking Activin Receptor Ligands Is Not Sufficient to Rescue Cancer-Associated Gut Microbiotaâ€™s Role for Gut Microbial Flagellin in Colorectal Cancer and Cachexia?. Cancers, 2019, 11, 1799.	1.7	11
9	Chronic nonbacterial prostate inflammation in a rat model is associated with changes of gut microbiota that can be modified with a galactoglucomannanâ€™rich hemicellulose extract in the diet. BJU International, 2019, 123, 899-908.	1.3	29
10	Chlamydia trachomatis samples testing falsely negative in the Aptima Combo 2 test in Finland, 2019. Eurosurveillance, 2019, 24, .	3.9	25
11	Early fecal microbiota composition in children who later develop celiac disease and associated autoimmunity. Scandinavian Journal of Gastroenterology, 2018, 53, 403-409.	0.6	49
12	Six-Week Endurance Exercise Alters Gut Metagenome That Is not Reflected in Systemic Metabolism in Over-weight Women. Frontiers in Microbiology, 2018, 9, 2323.	1.5	145
13	Gut Microbiota Composition in Mid-Pregnancy Is Associated with Gestational Weight Gain but Not Prepregnancy Body Mass Index. Journal of Women's Health, 2018, 27, 1293-1301.	1.5	22
14	Enterobacter cloacae administration induces hepatic damage and subcutaneous fat accumulation in high-fat diet fed mice. PLoS ONE, 2018, 13, e0198262.	1.1	22
15	<i>Faecalibacterium prausnitzii</i> treatment improves hepatic health and reduces adipose tissue inflammation in high-fat fed mice. ISME Journal, 2017, 11, 1667-1679.	4.4	179
16	Gut Microbiota Analysis Results Are Highly Dependent on the 16S rRNA Gene Target Region, Whereas the Impact of DNA Extraction Is Minor. Journal of Biomolecular Techniques, 2017, 28, 19-30.	0.8	130
17	Evaluation of a multiplex real-time PCR kit AmplidiagÂ® Bacterial GE in the detection of bacterial pathogens from stool samples. Journal of Microbiological Methods, 2016, 128, 61-65.	0.7	9
18	Neuropeptide Y Overexpressing Female and Male Mice Show Divergent Metabolic but Not Gut Microbial Responses to Prenatal Metformin Exposure. PLoS ONE, 2016, 11, e0163805.	1.1	35

#	ARTICLE	IF	CITATIONS
19	The Microbiome Studies in Metabolic Diseases have Advanced but are Poorly Standardized and Lack a Mechanistic Perspective. <i>Journal of Diabetes & Metabolism</i> , 2015, 06, .	0.2	2
20	Fermentable fibres condition colon microbiota and promote diabetogenesis in NOD mice. <i>Diabetologia</i> , 2014, 57, 2183-2192.	2.9	35