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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Gut Microbiome Modification through Dietary Intervention in Patients with Colorectal Cancer: Protocol for a Prospective, Interventional, Controlled, Randomized Clinical Trial in Patients with Scheduled Surgical Intervention for CRC. Journal of Clinical Medicine, 2022, 11, 3613.	1.0	0
2	Dietary modulation of gut microbiota in patients with colorectal cancer undergoing surgery: A review. International Journal of Surgery, 2022, 104, 106751.	1.1	2
3	A New Paradigm in the Relationship between Melatonin and Breast Cancer: Gut Microbiota Identified as a Potential Regulatory Agent. Cancers, 2021, 13, 3141.	1.7	12
4	Gut and Endometrial Microbiome Dysbiosis: A New Emergent Risk Factor for Endometrial Cancer. Journal of Personalized Medicine, 2021, 11, 659.	1.1	17
5	Relationships of Gut Microbiota Composition, Short-Chain Fatty Acids and Polyamines with the Pathological Response to Neoadjuvant Radiochemotherapy in Colorectal Cancer Patients. International Journal of Molecular Sciences, 2021, 22, 9549.	1.8	13
6	Genome Profiling of H3k4me3 Histone Modification in Human Adipose Tissue during Obesity and Insulin Resistance. Biomedicines, 2021, 9, 1363.	1.4	4
7	Connection between the Gut Microbiome, Systemic Inflammation, Gut Permeability and FOXP3 Expression in Patients with Primary Sj¶gren's Syndrome. International Journal of Molecular Sciences, 2020, 21, 8733.	1.8	36
8	Microbial Signature in Adipose Tissue of Crohn's Disease Patients. Journal of Clinical Medicine, 2020, 9, 2448.	1.0	15
9	Breast and Gut Microbiota Action Mechanisms in Breast Cancer Pathogenesis and Treatment. Cancers, 2020, 12, 2465.	1.7	90
10	Gut Microbiota-Mediated Inflammation and Gut Permeability in Patients with Obesity and Colorectal Cancer. International Journal of Molecular Sciences, 2020, 21, 6782.	1.8	63
11	Measurement of Serum Testosterone in Nondiabetic Young Obese Men: Comparison of Direct Immunoassay to Liquid Chromatography-Tandem Mass Spectrometry. Biomolecules, 2020, 10, 1697.	1.8	4
12	The Role of the Gut Microbiome in Colorectal Cancer Development and Therapy Response. Cancers, 2020, 12, 1406.	1.7	185
13	Epigenetic regulation of white adipose tissue in the onset of obesity and metabolic diseases. Obesity Reviews, 2020, 21, e13054.	3.1	8
14	Detection of TP53 and PIK3CA Mutations in Circulating Tumor DNA Using Next-Generation Sequencing in the Screening Process for Early Breast Cancer Diagnosis. Journal of Clinical Medicine, 2019, 8, 1183.	1.0	38
15	Crossâ€Sectional, Primary Care–Based Study of the Prevalence of Hypoandrogenemia in Nondiabetic Young Men with Obesity. Obesity, 2019, 27, 1584-1590.	1.5	16
16	Effects of SHBG rs1799941 Polymorphism on Free Testosterone Levels and Hypogonadism Risk in Young Non-Diabetic Obese Males. Journal of Clinical Medicine, 2019, 8, 1136.	1.0	5
17	Human adipose tissue H3K4me3 histone mark in adipogenic, lipid metabolism and inflammatory genes is positively associated with BMI and HOMA-IR. PLoS ONE, 2019, 14, e0215083.	1.1	33
18	Dietary and Gut Microbiota Polyamines in Obesity- and Age-Related Diseases. Frontiers in Nutrition, 2019, 6, 24.	1.6	133

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19	Gender-Related Differences on Polyamine Metabolome in Liquid Biopsies by a Simple and Sensitive Two-Step Liquid-Liquid Extraction and LC-MS/MS. Biomolecules, 2019, 9, 779.	1.8	10
20	Altered Adipose Tissue DNA Methylation Status in Metabolic Syndrome: Relationships Between Global DNA Methylation and Specific Methylation at Adipogenic, Lipid Metabolism and Inflammatory Candidate Genes and Metabolic Variables. Journal of Clinical Medicine, 2019, 8, 87.	1.0	67
21	Elevated circulating levels of succinate in human obesity are linked to specific gut microbiota. ISME Journal, 2018, 12, 1642-1657.	4.4	260
22	Adipose Tissue LPL Methylation is Associated with Triglyceride Concentrations in the Metabolic Syndrome. Clinical Chemistry, 2018, 64, 210-218.	1.5	30
23	Gut Microbiota Differs in Composition and Functionality Between Children With Type 1 Diabetes and MODY2 and Healthy Control Subjects: A Case-Control Study. Diabetes Care, 2018, 41, 2385-2395.	4.3	176
24	Complement Factor C3 Methylation and mRNA Expression Is Associated to BMI and Insulin Resistance in Obesity. Genes, 2018, 9, 410.	1.0	13
25	Chromatin immunoprecipitation improvements for the processing of small frozen pieces of adipose tissue. PLoS ONE, 2018, 13, e0192314.	1.1	6
26	Neonatal Androgen Exposure Causes Persistent Gut Microbiota Dysbiosis Related to Metabolic Disease in Adult Female Rats. Endocrinology, 2016, 157, 4888-4898.	1.4	76
27	Insulin resistance is associated with specific gut microbiota in appendix samples from morbidly obese patients. American Journal of Translational Research (discontinued), 2016, 8, 5672-5684.	0.0	60
28	Metabolomic insights into the intricate gut microbial–host interaction in the development of obesity and type 2 diabetes. Frontiers in Microbiology, 2015, 6, 1151.	1.5	108
29	Benefits of polyphenols on gut microbiota and implications in human health. Journal of Nutritional Biochemistry, 2013, 24, 1415-1422.	1.9	1,146
30	Adipose Tissue Gene Expression of Factors Related to Lipid Processing in Obesity. PLoS ONE, 2011, 6, e24783.	1.1	94
31	Rapid Differential Diagnosis between Extrapulmonary Tuberculosis and Focal Complications of Brucellosis Using a Multiplex Real-Time PCR Assay. PLoS ONE, 2009, 4, e4526.	1.1	28
32	Rapid Diagnosis of Brucella Epididymo-Orchitis by Real-Time Polymerase Chain Reaction Assay in Urine Samples. Journal of Urology, 2006, 176, 2290-2293.	0.2	31