

Andrea Quagliariello

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

1,308
citations

430442

18
h-index

377514

34
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41
all docs

41
docs citations

41
times ranked

2483
citing authors

#	ARTICLE	IF	CITATIONS
1	Gut Microbiota Markers in Obese Adolescent and Adult Patients: Age-Dependent Differential Patterns. <i>Frontiers in Microbiology</i> , 2018, 9, 1210.	1.5	139
2	Gut metabolomics profiling of non-small cell lung cancer (NSCLC) patients under immunotherapy treatment. <i>Journal of Translational Medicine</i> , 2020, 18, 49.	1.8	114
3	Gut microbiota profile in children affected by atopic dermatitis and evaluation of intestinal persistence of a probiotic mixture. <i>Scientific Reports</i> , 2019, 9, 4996.	1.6	107
4	Effect of <i>Bifidobacterium breve</i> on the Intestinal Microbiota of Coeliac Children on a Gluten Free Diet: A Pilot Study. <i>Nutrients</i> , 2016, 8, 660.	1.7	106
5	Autism, Gastrointestinal Symptoms and Modulation of Gut Microbiota by Nutritional Interventions. <i>Nutrients</i> , 2019, 11, 2812.	1.7	102
6	Gut Microbiota Profiling and Gut-Brain Crosstalk in Children Affected by Pediatric Acute-Onset Neuropsychiatric Syndrome and Pediatric Autoimmune Neuropsychiatric Disorders Associated With Streptococcal Infections. <i>Frontiers in Microbiology</i> , 2018, 9, 675.	1.5	88
7	Evaluation of the effects of intrapartum antibiotic prophylaxis on newborn intestinal microbiota using a sequencing approach targeted to multi hypervariable 16S rDNA regions. <i>Applied Microbiology and Biotechnology</i> , 2016, 100, 5537-5546.	1.7	84
8	Clinical intervention using <i>Bifidobacterium</i> strains in celiac disease children reveals novel microbial modulators of TNF- α and short-chain fatty acids. <i>Clinical Nutrition</i> , 2019, 38, 1373-1381.	2.3	79
9	Ancient and recent admixture layers in Sicily and Southern Italy trace multiple migration routes along the Mediterranean. <i>Scientific Reports</i> , 2017, 7, 1984.	1.6	52
10	Complex interplay between neutral and adaptive evolution shaped differential genomic background and disease susceptibility along the Italian peninsula. <i>Scientific Reports</i> , 2016, 6, 32513.	1.6	41
11	Daily Consumption of Orange Juice from <i>Citrus sinensis</i> L. Osbeck cv. Cara Cara and cv. Bahia Differently Affects Gut Microbiota Profiling as Unveiled by an Integrated Meta-Omics Approach. <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 1381-1391.	2.4	39
12	Genetic signature of differential sensitivity to stevioside in the Italian population. <i>Genes and Nutrition</i> , 2014, 9, 401.	1.2	33
13	Fecal microbiota signatures of insulin resistance, inflammation, and metabolic syndrome in youth with obesity: a pilot study. <i>Acta Diabetologica</i> , 2021, 58, 1009-1022.	1.2	32
14	Identification of new biomarkers of bronchopulmonary dysplasia using metabolomics. <i>Metabolomics</i> , 2019, 15, 20.	1.4	31
15	Distinct gut microbiota profile in antiretroviral therapy-treated perinatally HIV-infected patients associated with cardiac and inflammatory biomarkers. <i>Aids</i> , 2019, 33, 1001-1011.	1.0	31
16	A Metagenomic and in Silico Functional Prediction of Gut Microbiota Profiles May Concur in Discovering New Cystic Fibrosis Patient-Targeted Probiotics. <i>Nutrients</i> , 2017, 9, 1342.	1.7	24
17	Towards a disease-associated common trait of gut microbiota dysbiosis: The pivotal role of <i>Akkermansia muciniphila</i> . <i>Digestive and Liver Disease</i> , 2020, 52, 1002-1010.	0.4	23
18	Mutation Rates and Discriminating Power for 13 Rapidly-Mutating Y-STRs between Related and Unrelated Individuals. <i>PLoS ONE</i> , 2016, 11, e0165678.	1.1	22

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19	16S Metagenomics Reveals Dysbiosis of Nasal Core Microbiota in Children With Chronic Nasal Inflammation: Role of Adenoid Hypertrophy and Allergic Rhinitis. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 458.	1.8	21
20	Fecal Microbiota Transplant in Two Ulcerative Colitis Pediatric Cases: Gut Microbiota and Clinical Course Correlations. <i>Microorganisms</i> , 2020, 8, 1486.	1.6	18
21	Shared language, diverging genetic histories: high-resolution analysis of Y-chromosome variability in Calabrian and Sicilian Arbereshe. <i>European Journal of Human Genetics</i> , 2016, 24, 600-606.	1.4	16
22	Active microbial ecosystem in Iron Age tombs of the Etruscan civilization. <i>Environmental Microbiology</i> , 2021, 23, 3957-3969.	1.8	15
23	Gut Microbiota Profile in Children with IgE-Mediated Cow's Milk Allergy and Cow's Milk Sensitization and Probiotic Intestinal Persistence Evaluation. <i>International Journal of Molecular Sciences</i> , 2021, 22, 1649.	1.8	15
24	An omic approach to congenital diaphragmatic hernia: a pilot study of genomic, microRNA, and metabolomic profiling. <i>Journal of Perinatology</i> , 2020, 40, 952-961.	0.9	13
25	Anti-tumor necrosis factor α therapy associates to type 17 helper T lymphocytes immunological shift and significant microbial changes in dextran sodium sulphate colitis. <i>World Journal of Gastroenterology</i> , 2019, 25, 1465-1477.	1.4	11
26	Positive selection of lactase persistence among people of Southern Arabia. <i>American Journal of Physical Anthropology</i> , 2016, 161, 676-684.	2.1	9
27	Multiple selective events at the PRDM16 functional pathway shaped adaptation of western European populations to different climate conditions. <i>Journal of Anthropological Sciences</i> , 2017, 95, 235-247.	0.4	7
28	Gut microbiota composition in Himalayan and Andean populations and its relationship with diet, lifestyle and adaptation to the high-altitude environment. <i>Journal of Anthropological Sciences</i> , 2019, 96, 189-208.	0.4	7
29	Faecal microbiota transplantation in <i>Clostridioides difficile</i> infection: real-life experience from an academic Italian hospital. <i>Therapeutic Advances in Gastroenterology</i> , 2020, 13, 175628482093431.	1.4	6
30	Combined metagenomic and archaeobotanical analyses on human dental calculus: A cross-section of lifestyle conditions in a Copper Age population of central Italy. <i>Quaternary International</i> , 2023, 653-654, 69-81.	0.7	6
31	Ancient pathogen-driven adaptation triggers increased susceptibility to non-celiac wheat sensitivity in present-day European populations. <i>Genes and Nutrition</i> , 2016, 11, 15.	1.2	5
32	Massive parallel sequencing of human whole mitochondrial genomes with Ion Torrent technology: an optimized workflow for Anthropological and Population Genetics studies. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017, 28, 843-850.	0.7	4
33	The impact of intestinal microbiota on weight loss in Parkinson's disease patients: a pilot study. <i>Future Microbiology</i> , 2020, 15, 1393-1404.	1.0	4
34	Effects of a Synbiotic Formula on Functional Bowel Disorders and Gut Microbiota Profile during Long-Term Home Enteral Nutrition (LTHEN): A Pilot Study. <i>Nutrients</i> , 2021, 13, 87.	1.7	3
35	Editorial: The Role of Microbiota in the Onset and Development of Intestine and Liver Diseases and Cancer: Molecular and Cell Mechanisms. <i>Frontiers in Cell and Developmental Biology</i> , 2022, 10, 852188.	1.8	1
36	OC.05.5 ALTERATION IN THE ABUNDANCE OF AKKERMANSIA MUCINIPHILA IS ASSOCIATED TO GASTROINTESTINAL AND EXTRA-INTESTINAL DISEASES: TOWARDS THE IDENTIFICATION OF SPECIFIC MICROBIAL SIGNATURES OF DISEASE. <i>Digestive and Liver Disease</i> , 2019, 51, e90.	0.4	0

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37	OC.04.5 IL-33/ST2 LEVELS AND GUT MICROBIOTA CHARACTERIZATION CAN PREDICT MUCOSAL RESPONSE TO ANTI-TNF THERAPY IN ULCERATIVE COLITIS. <i>Digestive and Liver Disease</i> , 2019, 51, e87-e88.	0.4	0
38	P497 IL-33/ST2 levels and gut microbiota characterisation can predict mucosal response to anti-TNF therapy in ulcerative colitis. <i>Journal of Crohn's and Colitis</i> , 2019, 13, S360-S361.	0.6	0
39	Beneficial Commensal Bacteria Promote Drosophila Growth by Down-Regulating the Expression of Peptidoglycan Recognition Proteins. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0