

# Monica Di Paola

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3508272/publications.pdf>

Version: 2024-02-01

29  
papers

6,021  
citations

471509

17  
h-index

501196

28  
g-index

29  
all docs

29  
docs citations

29  
times ranked

10505  
citing authors

#	ARTICLE	IF	CITATIONS
1	Impact of diet in shaping gut microbiota revealed by a comparative study in children from Europe and rural Africa. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 14691-14696.	7.1	4,561
2	Role of social wasps in <i>Saccharomyces cerevisiae</i> ecology and evolution. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 13398-13403.	7.1	259
3	Diet, Environments, and Gut Microbiota. A Preliminary Investigation in Children Living in Rural and Urban Burkina Faso and Italy. Frontiers in Microbiology, 2017, 8, 1979.	3.5	222
4	Characterization of cervico-vaginal microbiota in women developing persistent high-risk Human Papillomavirus infection. Scientific Reports, 2017, 7, 10200.	3.3	188
5	Age and Gender Affect the Composition of Fungal Population of the Human Gastrointestinal Tract. Frontiers in Microbiology, 2016, 7, 1227.	3.5	170
6	Phenotype and Disease Course of Early-onset Pediatric Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2014, 20, 597-605.	1.9	119
7	Alteration of Fecal Microbiota Profiles in Juvenile Idiopathic Arthritis. Associations with HLA-B27 Allele and Disease Status. Frontiers in Microbiology, 2016, 7, 1703.	3.5	65
8	Oleoylethanolamide treatment affects gut microbiota composition and the expression of intestinal cytokines in Peyer's patches of mice. Scientific Reports, 2018, 8, 14881.	3.3	39
9	Gut microbiota in children and altered profiles in juvenile idiopathic arthritis. Journal of Autoimmunity, 2019, 98, 1-12.	6.5	39
10	Genomics approach to the analysis of bacterial communities dynamics in Hirschsprung's disease-associated enterocolitis: a pilot study. Pediatric Surgery International, 2010, 26, 465-471.	1.4	36
11	Gut microbiota profiles and characterization of cultivable fungal isolates in IBS patients. Applied Microbiology and Biotechnology, 2021, 105, 3277-3288.	3.6	31
12	Population genomics reveals evolution and variation of <i>Saccharomyces cerevisiae</i> in the human and insects gut. Environmental Microbiology, 2019, 21, 50-71.	3.8	30
13	Early melanoma invasivity correlates with gut fungal and bacterial profiles. British Journal of Dermatology, 2022, 186, 106-116.	1.5	26
14	Nod2 Deficiency in mice is Associated with Microbiota Variation Favouring the Expansion of mucosal CD4+ LAP+ Regulatory Cells. Scientific Reports, 2018, 8, 14241.	3.3	25
15	IL-13 mRNA Tissue Content Identifies Two Subsets of Adult Ulcerative Colitis Patients With Different Clinical and Mucosa-Associated Microbiota Profiles. Journal of Crohn's and Colitis, 2020, 14, 369-380.	1.3	25
16	Comparative immunophenotyping of <i>Saccharomyces cerevisiae</i> and <i>Candida</i> spp. strains from Crohn's disease patients and their interactions with the gut microbiome. Journal of Translational Autoimmunity, 2020, 3, 100036.	4.0	24
17	Genomic and Phenotypic Variation in Morphogenetic Networks of Two <i>Candida albicans</i> Isolates Subtends Their Different Pathogenic Potential. Frontiers in Immunology, 2018, 8, 1997.	4.8	23
18	<i>Saccharomyces cerevisiae</i> Induces Immune Enhancing and Shapes Gut Microbiota in Social Wasps. Frontiers in Microbiology, 2019, 10, 2320.	3.5	22

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19	A Metagenomics Study on Hirschsprung's Disease Associated Enterocolitis: Biodiversity and Gut Microbial Homeostasis Depend on Resection Length and Patient's Clinical History. <i>Frontiers in Pediatrics</i> , 2019, 7, 326.	1.9	19
20	<i>Saccharomyces cerevisiae</i> "Insects Association: Impacts, Biogeography, and Extent. <i>Frontiers in Microbiology</i> , 2020, 11, 1629.	3.5	18
21	Changing Dietary Habits: The Impact of Urbanization and Rising Socio-Economic Status in Families from Burkina Faso in Sub-Saharan Africa. <i>Nutrients</i> , 2022, 14, 1782.	4.1	16
22	Active microbial ecosystem in Iron-Age tombs of the Etruscan civilization. <i>Environmental Microbiology</i> , 2021, 23, 3957-3969.	3.8	15
23	Identification of Vaginal Microbial Communities Associated with Extreme Cervical Shortening in Pregnant Women. <i>Journal of Clinical Medicine</i> , 2020, 9, 3621.	2.4	12
24	Study on a Fermented Whole Wheat: Phenolic Content, Activity on PTP1B Enzyme and In Vitro Prebiotic Properties. <i>Molecules</i> , 2019, 24, 1120.	3.8	11
25	Intestinal microbiota profiles in a genetic model of colon tumorigenesis correlates with colon cancer biomarkers. <i>Scientific Reports</i> , 2022, 12, 1432.	3.3	9
26	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
27	Applications of Wild Isolates of <i>Saccharomyces</i> Yeast for Industrial Fermentation: The Gut of Social Insects as Niche for Yeast Hybrids™ Production. <i>Frontiers in Microbiology</i> , 2020, 11, 578425.	3.5	5
28	Resistance to Arsenite and Arsenate in <i>Saccharomyces cerevisiae</i> Arises through the Subtelomeric Expansion of a Cluster of Yeast Genes. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 8119.	2.6	5
29	P419 Long-term outcome of ulcerative colitis in pediatric patients who achieved mucosal and histological healing: a real-life referral center experience. <i>Journal of Crohn's and Colitis</i> , 2022, 16, i407-i408.	1.3	0