## Federico E Rey

## List of Publications by Citations

Source: https://exaly.com/author-pdf/2509422/federico-e-rey-publications-by-citations.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

59	15,966	34	71
papers	citations	h-index	g-index
71	19,736 ext. citations	13.8	6.15
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
59	Human gut microbiome viewed across age and geography. <i>Nature</i> , <b>2012</b> , 486, 222-7	50.4	4616
58	Gut microbiota from twins discordant for obesity modulate metabolism in mice. <i>Science</i> , <b>2013</b> , 341, 124	41323134	2251
57	The effect of diet on the human gut microbiome: a metagenomic analysis in humanized gnotobiotic mice. <i>Science Translational Medicine</i> , <b>2009</b> , 1, 6ra14	17.5	1977
56	Effects of the gut microbiota on host adiposity are modulated by the short-chain fatty-acid binding G protein-coupled receptor, Gpr41. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 16767-72	11.5	1070
55	Gut microbiome alterations in Alzheimerঙ disease. <i>Scientific Reports</i> , <b>2017</b> , 7, 13537	4.9	712
54	Olfactory receptor responding to gut microbiota-derived signals plays a role in renin secretion and blood pressure regulation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2013</b> , 110, 4410-5	11.5	640
53	Characterizing a model human gut microbiota composed of members of its two dominant bacterial phyla. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2009</b> , 106, 5859-	-6 <sup>4</sup> 1.5	478
52	Predicting a human gut microbiotald response to diet in gnotobiotic mice. Science, 2011, 333, 101-4	33.3	391
51	Intestinal microbiota composition modulates choline bioavailability from diet and accumulation of the proatherogenic metabolite trimethylamine-N-oxide. <i>MBio</i> , <b>2015</b> , 6, e02481	7.8	389
50	Diet-Microbiota Interactions Mediate Global Epigenetic Programming in Multiple Host Tissues. <i>Molecular Cell</i> , <b>2016</b> , 64, 982-992	17.6	280
49	Metabolic niche of a prominent sulfate-reducing human gut bacterium. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2013</b> , 110, 13582-7	11.5	239
48	Bacteria from diverse habitats colonize and compete in the mouse gut. Cell, 2014, 159, 253-66	56.2	226
47	Dissecting the in vivo metabolic potential of two human gut acetogens. <i>Journal of Biological Chemistry</i> , <b>2010</b> , 285, 22082-90	5.4	225
46	Interactions between Roseburia intestinalis and diet modulate atherogenesis in a murine model. <i>Nature Microbiology</i> , <b>2018</b> , 3, 1461-1471	26.6	170
45	The gut microbiota-derived metabolite trimethylamine N-oxide is elevated in Alzheimer <b>u</b> disease. <i>Alzheimers Research and Therapy</i> , <b>2018</b> , 10, 124	9	156
44	A Cardiovascular Disease-Linked Gut Microbial Metabolite Acts via Adrenergic Receptors. <i>Cell</i> , <b>2020</b> , 180, 862-877.e22	56.2	146
43	The reactive adventitia: fibroblast oxidase in vascular function. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , <b>2002</b> , 22, 1962-71	9.4	140

## (2019-2000)

42	Vascular effects following homozygous disruption of p47(phox): An essential component of NADPH oxidase. <i>Circulation</i> , <b>2000</b> , 101, 1234-6	16.7	140
41	Functional genomic analysis of three nitrogenase isozymes in the photosynthetic bacterium Rhodopseudomonas palustris. <i>Journal of Bacteriology</i> , <b>2005</b> , 187, 7784-94	3.5	126
40	Redirection of metabolism for biological hydrogen production. <i>Applied and Environmental Microbiology</i> , <b>2007</b> , 73, 1665-71	4.8	121
39	Fatty acid synthase modulates intestinal barrier function through palmitoylation of mucin 2. <i>Cell Host and Microbe</i> , <b>2012</b> , 11, 140-52	23.4	103
38	Metabolic, Epigenetic, and Transgenerational Effects of Gut Bacterial Choline Consumption. <i>Cell Host and Microbe</i> , <b>2017</b> , 22, 279-290.e7	23.4	100
37	Perivascular superoxide anion contributes to impairment of endothelium-dependent relaxation: role of gp91(phox). <i>Circulation</i> , <b>2002</b> , 106, 2497-502	16.7	99
36	Trimethylamine N-Oxide Binds and Activates PERK to Promote Metabolic Dysfunction. <i>Cell Metabolism</i> , <b>2019</b> , 30, 1141-1151.e5	24.6	98
35	Host Genotype and Gut Microbiome Modulate Insulin Secretion and Diet-Induced Metabolic Phenotypes. <i>Cell Reports</i> , <b>2017</b> , 18, 1739-1750	10.6	91
34	Creating and characterizing communities of human gut microbes in gnotobiotic mice. <i>ISME Journal</i> , <b>2010</b> , 4, 1094-8	11.9	91
33	Regulation of uptake hydrogenase and effects of hydrogen utilization on gene expression in Rhodopseudomonas palustris. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 6143-52	3.5	90
32	Loss of Gut Microbiota Alters Immune System Composition and Cripples Postinfarction Cardiac Repair. <i>Circulation</i> , <b>2019</b> , 139, 647-659	16.7	85
31	Close social relationships correlate with human gut microbiota composition. <i>Scientific Reports</i> , <b>2019</b> , 9, 703	4.9	81
30	Untargeted metabolomics identifies trimethyllysine, a TMAO-producing nutrient precursor, as a predictor of incident cardiovascular disease risk. <i>JCI Insight</i> , <b>2018</b> , 3,	9.9	78
29	Microbial Transplantation With Human Gut Commensals Containing CutC Is Sufficient to Transmit Enhanced Platelet Reactivity and Thrombosis Potential. <i>Circulation Research</i> , <b>2018</b> , 123, 1164-1176	15.7	68
28	A common antimicrobial additive increases colonic inflammation and colitis-associated colon tumorigenesis in mice. <i>Science Translational Medicine</i> , <b>2018</b> , 10,	17.5	62
27	Hydrogen production by photoreactive nanoporous latex coatings of nongrowing Rhodopseudomonas palustris CGA009. <i>Biotechnology Progress</i> , <b>2007</b> , 23, 124-30	2.8	55
26	FixK, a global regulator of microaerobic growth, controls photosynthesis in Rhodopseudomonas palustris. <i>Molecular Microbiology</i> , <b>2010</b> , 75, 1007-20	4.1	40
25	Genetic determinants of gut microbiota composition and bile acid profiles in mice. <i>PLoS Genetics</i> , <b>2019</b> , 15, e1008073	6	32

24	Sexual dimorphism of cardiometabolic dysfunction: Gut microbiome in the play?. <i>Molecular Metabolism</i> , <b>2018</b> , 15, 70-81	8.8	28
23	Chemical signaling between gut microbiota and host chromatin: What is your gut really saying?. <i>Journal of Biological Chemistry</i> , <b>2017</b> , 292, 8582-8593	5.4	27
22	Autometa: automated extraction of microbial genomes from individual shotgun metagenomes. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, e57	20.1	27
21	Social and population health science approaches to understand the human microbiome. <i>Nature Human Behaviour</i> , <b>2018</b> , 2, 808-815	12.8	25
20	Critical symbiont signals drive both local and systemic changes in diel and developmental host gene expression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2019</b> , 116, 7990-7999	11.5	24
19	Gut Microbial and Metabolic Responses to Salmonella enterica Serovar Typhimurium and Candida albicans. <i>MBio</i> , <b>2018</b> , 9,	7.8	21
18	The emerging role of gut microbial metabolism on cardiovascular disease. <i>Current Opinion in Microbiology</i> , <b>2019</b> , 50, 64-70	7.9	19
17	Starch Utilization Promotes Quercetin Degradation and Butyrate Production by. <i>Frontiers in Microbiology</i> , <b>2019</b> , 10, 1145	5.7	17
16	Gut microbes impact stroke severity via the trimethylamine N-oxide pathway. <i>Cell Host and Microbe</i> , <b>2021</b> , 29, 1199-1208.e5	23.4	17
15	Gut microbiome variation modulates the effects of dietary fiber on host metabolism. <i>Microbiome</i> , <b>2021</b> , 9, 117	16.6	14
14	The Influence of Social Conditions Across the Life Course on the Human Gut Microbiota: A Pilot Project With the Wisconsin Longitudinal Study. <i>Journals of Gerontology - Series B Psychological Sciences and Social Sciences</i> , <b>2017</b> , 73, 124-133	4.6	10
13	Effects of Smoking and Smoking Cessation on the Intestinal Microbiota. <i>Journal of Clinical Medicine</i> , <b>2020</b> , 9,	5.1	10
12	Differential Catabolism of an Anthocyanin-Rich Elderberry Extract by Three Gut Microbiota Bacterial Species. <i>Journal of Agricultural and Food Chemistry</i> , <b>2020</b> , 68, 1837-1843	5.7	9
11	Aronia berry polyphenols have matrix-dependent effects on the gut microbiota. <i>Food Chemistry</i> , <b>2021</b> , 359, 129831	8.5	9
10	Fecal Aliquot Straw Technique (FAST) allows for easy and reproducible subsampling: assessing interpersonal variation in trimethylamine-N-oxide (TMAO) accumulation. <i>Microbiome</i> , <b>2018</b> , 6, 91	16.6	8
9	Is maternal microbial metabolism an early-life determinant of health?. Lab Animal, 2018, 47, 239-243	0.4	7
8	The Plot Thickens: Diet Microbe Interactions May Modulate Thrombosis Risk. <i>Cell Metabolism</i> , <b>2016</b> , 23, 573-5	24.6	7
7	Integrated Label-Free and 10-Plex DiLeu Isobaric Tag Quantitative Methods for Profiling Changes in the Mouse Hypothalamic Neuropeptidome and Proteome: Assessment of the Impact of the Gut Microbiome. Analytical Chemistry 2020, 92, 14021-14030	7.8	6

## LIST OF PUBLICATIONS

6	Extraction optimization for combined metabolomics, peptidomics, and proteomics analysis of gut microbiota samples. <i>Journal of Mass Spectrometry</i> , <b>2021</b> , 56, e4625	2.2	3
5	Gut-derived Flavonifractor species variants are differentially enriched during in vitro incubation with quercetin. <i>PLoS ONE</i> , <b>2020</b> , 15, e0227724	3.7	2
4	Social relationships, social isolation, and the human gut microbiota		2
3	Selective Bacterial Colonization of the Murine Larynx in a Gnotobiotic Model. <i>Frontiers in Microbiology</i> , <b>2020</b> , 11, 594617	5.7	2
2	Vocal fold mucus layer: Comparison of histological protocols for visualization in mice <i>Laryngoscope Investigative Otolaryngology</i> , <b>2022</b> , 7, 444-453	2.8	1
1	The human gut microbiota contributes to type-2 diabetes non-resolution 5-years after Roux-en-Y gastric bypass <i>Gut Microbes</i> , <b>2022</b> , 14, 2050635	8.8	1