Federico E Rey

List of Publications by Year in Descending Order

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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 ext. papers	19,736 ext. citations	13.8 avg, IF	6.15 L-index

#	Paper	IF	Citations
59	Vocal fold mucus layer: Comparison of histological protocols for visualization in mice <i>Laryngoscope Investigative Otolaryngology</i> , 2022 , 7, 444-453	2.8	1
58	The human gut microbiota contributes to type-2 diabetes non-resolution 5-years after Roux-en-Y gastric bypass <i>Gut Microbes</i> , 2022 , 14, 2050635	8.8	1
57	Gut microbiome variation modulates the effects of dietary fiber on host metabolism. <i>Microbiome</i> , 2021 , 9, 117	16.6	14
56	Extraction optimization for combined metabolomics, peptidomics, and proteomics analysis of gut microbiota samples. <i>Journal of Mass Spectrometry</i> , 2021 , 56, e4625	2.2	3
55	Gut microbes impact stroke severity via the trimethylamine N-oxide pathway. <i>Cell Host and Microbe</i> , 2021 , 29, 1199-1208.e5	23.4	17
54	Aronia berry polyphenols have matrix-dependent effects on the gut microbiota. <i>Food Chemistry</i> , 2021 , 359, 129831	8.5	9
53	A Cardiovascular Disease-Linked Gut Microbial Metabolite Acts via Adrenergic Receptors. <i>Cell</i> , 2020 , 180, 862-877.e22	56.2	146
52	Gut-derived Flavonifractor species variants are differentially enriched during in vitro incubation with quercetin. <i>PLoS ONE</i> , 2020 , 15, e0227724	3.7	2
51	Selective Bacterial Colonization of the Murine Larynx in a Gnotobiotic Model. <i>Frontiers in Microbiology</i> , 2020 , 11, 594617	5.7	2
50	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. <i>Analytical Chemistry</i> , 2020 , 92, 14021-14030	7.8	6
49	Effects of Smoking and Smoking Cessation on the Intestinal Microbiota. <i>Journal of Clinical Medicine</i> , 2020 , 9,	5.1	10
48	Differential Catabolism of an Anthocyanin-Rich Elderberry Extract by Three Gut Microbiota Bacterial Species. <i>Journal of Agricultural and Food Chemistry</i> , 2020 , 68, 1837-1843	5.7	9
47	Genetic determinants of gut microbiota composition and bile acid profiles in mice. <i>PLoS Genetics</i> , 2019 , 15, e1008073	6	32
46	Trimethylamine N-Oxide Binds and Activates PERK to Promote Metabolic Dysfunction. <i>Cell Metabolism</i> , 2019 , 30, 1141-1151.e5	24.6	98
45	Close social relationships correlate with human gut microbiota composition. <i>Scientific Reports</i> , 2019 , 9, 703	4.9	81
44	Starch Utilization Promotes Quercetin Degradation and Butyrate Production by. <i>Frontiers in Microbiology</i> , 2019 , 10, 1145	5.7	17
43	Autometa: automated extraction of microbial genomes from individual shotgun metagenomes. <i>Nucleic Acids Research</i> , 2019 , 47, e57	20.1	27

(2017-2019)

42	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> , 2019 , 116, 7990-7999	11.5	24
41	The emerging role of gut microbial metabolism on cardiovascular disease. <i>Current Opinion in Microbiology</i> , 2019 , 50, 64-70	7.9	19
40	Loss of Gut Microbiota Alters Immune System Composition and Cripples Postinfarction Cardiac Repair. <i>Circulation</i> , 2019 , 139, 647-659	16.7	85
39	Fecal Aliquot Straw Technique (FAST) allows for easy and reproducible subsampling: assessing interpersonal variation in trimethylamine-N-oxide (TMAO) accumulation. <i>Microbiome</i> , 2018 , 6, 91	16.6	8
38	Is maternal microbial metabolism an early-life determinant of health?. Lab Animal, 2018, 47, 239-243	0.4	7
37	Sexual dimorphism of cardiometabolic dysfunction: Gut microbiome in the play?. <i>Molecular Metabolism</i> , 2018 , 15, 70-81	8.8	28
36	Untargeted metabolomics identifies trimethyllysine, a TMAO-producing nutrient precursor, as a predictor of incident cardiovascular disease risk. <i>JCI Insight</i> , 2018 , 3,	9.9	78
35	Gut Microbial and Metabolic Responses to Salmonella enterica Serovar Typhimurium and Candida albicans. <i>MBio</i> , 2018 , 9,	7.8	21
34	The gut microbiota-derived metabolite trimethylamine N-oxide is elevated in Alzheimer u disease. <i>Alzheimers Research and Therapy</i> , 2018 , 10, 124	9	156
33	Interactions between Roseburia intestinalis and diet modulate atherogenesis in a murine model. <i>Nature Microbiology</i> , 2018 , 3, 1461-1471	26.6	170
32	Social and population health science approaches to understand the human microbiome. <i>Nature Human Behaviour</i> , 2018 , 2, 808-815	12.8	25
31	Microbial Transplantation With Human Gut Commensals Containing CutC Is Sufficient to Transmit Enhanced Platelet Reactivity and Thrombosis Potential. <i>Circulation Research</i> , 2018 , 123, 1164-1176	15.7	68
30	A common antimicrobial additive increases colonic inflammation and colitis-associated colon tumorigenesis in mice. <i>Science Translational Medicine</i> , 2018 , 10,	17.5	62
29	Host Genotype and Gut Microbiome Modulate Insulin Secretion and Diet-Induced Metabolic Phenotypes. <i>Cell Reports</i> , 2017 , 18, 1739-1750	10.6	91
28	Chemical signaling between gut microbiota and host chromatin: What is your gut really saying?. <i>Journal of Biological Chemistry</i> , 2017 , 292, 8582-8593	5.4	27
27	Gut microbiome alterations in Alzheimer disease. Scientific Reports, 2017, 7, 13537	4.9	712
26	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> , 2017 , 73, 124-133	4.6	10
25	Metabolic, Epigenetic, and Transgenerational Effects of Gut Bacterial Choline Consumption. <i>Cell Host and Microbe</i> , 2017 , 22, 279-290.e7	23.4	100

24	Diet-Microbiota Interactions Mediate Global Epigenetic Programming in Multiple Host Tissues. <i>Molecular Cell</i> , 2016 , 64, 982-992	17.6	280
23	The Plot Thickens: Diet Microbe Interactions May Modulate Thrombosis Risk. <i>Cell Metabolism</i> , 2016 , 23, 573-5	24.6	7
22	Intestinal microbiota composition modulates choline bioavailability from diet and accumulation of the proatherogenic metabolite trimethylamine-N-oxide. <i>MBio</i> , 2015 , 6, e02481	7.8	389
21	Bacteria from diverse habitats colonize and compete in the mouse gut. <i>Cell</i> , 2014 , 159, 253-66	56.2	226
20	Gut microbiota from twins discordant for obesity modulate metabolism in mice. <i>Science</i> , 2013 , 341, 124	41 33 134	2251
19	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> , 2013 , 110, 4410-5	11.5	640
18	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> , 2013 , 110, 13582-7	11.5	239
17	Fatty acid synthase modulates intestinal barrier function through palmitoylation of mucin 2. <i>Cell Host and Microbe</i> , 2012 , 11, 140-52	23.4	103
16	Human gut microbiome viewed across age and geography. <i>Nature</i> , 2012 , 486, 222-7	50.4	4616
15	Predicting a human gut microbiotal response to diet in gnotobiotic mice. Science, 2011 , 333, 101-4	33.3	391
14	FixK, a global regulator of microaerobic growth, controls photosynthesis in Rhodopseudomonas palustris. <i>Molecular Microbiology</i> , 2010 , 75, 1007-20	4.1	40
13	Creating and characterizing communities of human gut microbes in gnotobiotic mice. <i>ISME Journal</i> , 2010 , 4, 1094-8	11.9	91
12	Dissecting the in vivo metabolic potential of two human gut acetogens. <i>Journal of Biological Chemistry</i> , 2010 , 285, 22082-90	5.4	225
11	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> , 2009 , 106, 5859-	·64 ^{1.5}	478
10	The effect of diet on the human gut microbiome: a metagenomic analysis in humanized gnotobiotic mice. <i>Science Translational Medicine</i> , 2009 , 1, 6ra14	17.5	1977
9	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> , 2008 , 105, 16767-72	11.5	1070
8	Redirection of metabolism for biological hydrogen production. <i>Applied and Environmental Microbiology</i> , 2007 , 73, 1665-71	4.8	121
7	Hydrogen production by photoreactive nanoporous latex coatings of nongrowing Rhodopseudomonas palustris CGA009. <i>Biotechnology Progress</i> , 2007 , 23, 124-30	2.8	55

LIST OF PUBLICATIONS

6	Regulation of uptake hydrogenase and effects of hydrogen utilization on gene expression in Rhodopseudomonas palustris. <i>Journal of Bacteriology</i> , 2006 , 188, 6143-52	3.5	90
5	Functional genomic analysis of three nitrogenase isozymes in the photosynthetic bacterium Rhodopseudomonas palustris. <i>Journal of Bacteriology</i> , 2005 , 187, 7784-94	3.5	126
4	Perivascular superoxide anion contributes to impairment of endothelium-dependent relaxation: role of gp91(phox). <i>Circulation</i> , 2002 , 106, 2497-502	16.7	99
3	The reactive adventitia: fibroblast oxidase in vascular function. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2002 , 22, 1962-71	9.4	140
2	Vascular effects following homozygous disruption of p47(phox): An essential component of NADPH oxidase. <i>Circulation</i> , 2000 , 101, 1234-6	16.7	140
1	Social relationships, social isolation, and the human gut microbiota		2