

Evgeni Levin

List of Publications by Year in descending order

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33
papers

2,985
citations

471509

17
h-index

454955

30
g-index

34
all docs

34
docs citations

34
times ranked

4655
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeted proteomics improves cardiovascular risk prediction in secondary prevention. <i>European Heart Journal</i> , 2022, 43, 1569-1577.	2.2	55
2	Fecal microbiota transplantation as tool to study the interrelation between microbiota composition and miRNA expression. <i>Microbiological Research</i> , 2022, 257, 126972.	5.3	5
3	Epigenetic Signatures Discriminate Patients With Primary Sclerosing Cholangitis and Ulcerative Colitis From Patients With Ulcerative Colitis. <i>Frontiers in Immunology</i> , 2022, 13, 840935.	4.8	4
4	Faecal microbiota transplantation halts progression of human new-onset type 1 diabetes in a randomised controlled trial. <i>Gut</i> , 2021, 70, 92-105.	12.1	161
5	Metabolomics in Severe Aortic Stenosis Reveals Intermediates of Nitric Oxide Synthesis as Most Distinctive Markers. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3569.	4.1	12
6	Analyzing Type 2 Diabetes Associations with the Gut Microbiome in Individuals from Two Ethnic Backgrounds Living in the Same Geographic Area. <i>Nutrients</i> , 2021, 13, 3289.	4.1	17
7	Duodenal <i>Anaerobutyricum soehngenii</i> infusion stimulates GLP-1 production, ameliorates glycaemic control and beneficially shapes the duodenal transcriptome in metabolic syndrome subjects: a randomised double-blind placebo-controlled cross-over study. <i>Gut</i> , 2021, , gutinl-2020-323297.	12.1	16
8	Distinct Metabolomic Signatures in Preclinical and Obstructive Hypertrophic Cardiomyopathy. <i>Cells</i> , 2021, 10, 2950.	4.1	5
9	Impact drugs targeting cardiometabolic risk on the gut microbiota. <i>Current Opinion in Lipidology</i> , 2021, 32, 38-54.	2.7	0
10	Effects of fecal microbiota transplant on DNA methylation in subjects with metabolic syndrome. <i>Gut Microbes</i> , 2021, 13, 1993513.	9.8	25
11	Donor metabolic characteristics drive effects of faecal microbiota transplantation on recipient insulin sensitivity, energy expenditure and intestinal transit time. <i>Gut</i> , 2020, 69, 502-512.	12.1	188
12	Oral butyrate does not affect innate immunity and islet autoimmunity in individuals with longstanding type 1 diabetes: a randomised controlled trial. <i>Diabetologia</i> , 2020, 63, 597-610.	6.3	60
13	Untargeted accurate identification of highly pathogenic bacteria directly from blood culture flasks. <i>International Journal of Medical Microbiology</i> , 2020, 310, 151376.	3.6	12
14	Infusion of donor feces affects the gut-brain axis in humans with metabolic syndrome. <i>Molecular Metabolism</i> , 2020, 42, 101076.	6.5	50
15	Donor Fecal Microbiota Transplantation Alters Gut Microbiota and Metabolites in Obese Individuals With Steatohepatitis. <i>Hepatology Communications</i> , 2020, 4, 1578-1590.	4.3	71
16	Differential DNA methylation in familial hypercholesterolemia. <i>EBioMedicine</i> , 2020, 61, 103079.	6.1	10
17	Improved cardiovascular risk prediction using targeted plasma proteomics in primary prevention. <i>European Heart Journal</i> , 2020, 41, 3998-4007.	2.2	68
18	Comparing bioinformatic pipelines for microbial 16S rRNA amplicon sequencing. <i>PLoS ONE</i> , 2020, 15, e0227434.	2.5	282

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19	Treatment with Anaerobutyricum soehngenii: a pilot study of safety and doseâ€‘response effects on glucose metabolism in human subjects with metabolic syndrome. Npj Biofilms and Microbiomes, 2020, 6, 16.	6.4	53
20	Manifold Mixing for Stacked Regularization. Communications in Computer and Information Science, 2020, , 444-452.	0.5	1
21	Predictive value of targeted proteomics for coronary plaque morphology in patients with suspected coronary artery disease. EBioMedicine, 2019, 39, 109-117.	6.1	42
22	Graph Space Embedding. , 2019, , .		3
23	Duodenal L cell density correlates with features of metabolic syndrome and plasma metabolites. Endocrine Connections, 2018, 7, 673-680.	1.9	4
24	Effect of Vegan Fecal Microbiota Transplantation on Carnitineâ€‘and Cholineâ€‘Derived Trimethylamineâ€‘Nâ€‘Oxide Production and Vascular Inflammation in Patients With Metabolic Syndrome. Journal of the American Heart Association, 2018, 7, .	3.7	164
25	Domain intelligible models. Methods, 2018, 149, 69-73.	3.8	4
26	Depicting the composition of gut microbiota in a population with varied ethnic origins but shared geography. Nature Medicine, 2018, 24, 1526-1531.	30.7	436
27	Intestinal Fungal Dysbiosis Is Associated With Visceral Hypersensitivity in Patients With Irritable Bowel Syndrome and Rats. Gastroenterology, 2017, 153, 1026-1039.	1.3	160
28	Improvement of Insulin Sensitivity after Lean Donor Feces in Metabolic Syndrome Is Driven by Baseline Intestinal Microbiota Composition. Cell Metabolism, 2017, 26, 611-619.e6.	16.2	689
29	Identification of microorganisms grown in blood culture flasks using liquid chromatographyâ€‘tandem mass spectrometry. Future Microbiology, 2017, 12, 1135-1145.	2.0	18
30	Unsupervised Multi-View Feature Selection for Tumor Subtype Identification. , 2017, , .		3
31	At least three phenotypes exist among periodontitis patients. Journal of Clinical Periodontology, 2017, 44, 1068-1076.	4.9	10
32	Distinct fecal and oral microbiota composition in human type 1 diabetes, an observational study. PLoS ONE, 2017, 12, e0188475.	2.5	163
33	Development of Upper Respiratory Tract Microbiota in Infancy is Affected by Mode of Delivery. EBioMedicine, 2016, 9, 336-345.	6.1	194