Andrei Prodan

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Faecal microbiota transplantation halts progression of human new-onset type 1 diabetes in a randomised controlled trial. Gut, 2021, 70, 92-105.	6.1	161
2	Fecal Microbiota Transplantation from Overweight or Obese Donors in Cachectic Patients with Advanced Gastroesophageal Cancer: A Randomized, Double-blind, Placebo-Controlled, Phase II Study. Clinical Cancer Research, 2021, 27, 3784-3792.	3.2	30
3	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. Gut, 2021, , gutinl-2020-323297.	6.1	16
4	Donor metabolic characteristics drive effects of faecal microbiota transplantation on recipient insulin sensitivity, energy expenditure and intestinal transit time. Gut, 2020, 69, 502-512.	6.1	188
5	Untargeted accurate identification of highly pathogenic bacteria directly from blood culture flasks. International Journal of Medical Microbiology, 2020, 310, 151376.	1.5	12
6	Gut Microbiota in Hypertension and Atherosclerosis: A Review. Nutrients, 2020, 12, 2982.	1.7	183
7	Infusion of donor feces affects the gut–brain axis in humans with metabolic syndrome. Molecular Metabolism, 2020, 42, 101076.	3.0	50
8	Donor Fecal Microbiota Transplantation Alters Gut Microbiota and Metabolites in Obese Individuals With Steatohepatitis. Hepatology Communications, 2020, 4, 1578-1590.	2.0	71
9	Plasma Metabolites Related to Peripheral and Hepatic Insulin Sensitivity Are Not Directly Linked to Gut Microbiota Composition. Nutrients, 2020, 12, 2308.	1.7	6
10	Associations between gut microbiota, faecal short-chain fatty acids, and blood pressure across ethnic groups: the HELIUS study. European Heart Journal, 2020, 41, 4259-4267.	1.0	124
11	A salivary metabolite signature that reflects gingival host-microbe interactions: instability predicts gingivitis susceptibility. Scientific Reports, 2020, 10, 3008.	1.6	2
12	Comparing bioinformatic pipelines for microbial 16S rRNA amplicon sequencing. PLoS ONE, 2020, 15, e0227434.	1.1	282
13	Treatment with Anaerobutyricum soehngenii: a pilot study of safety and dose–response effects on glucose metabolism in human subjects with metabolic syndrome. Npj Biofilms and Microbiomes, 2020, 6, 16.	2.9	53
14	Intestinal and tumor microbiome analysis combined with metabolomics of the anti-PD-L1 phase II PERFECT trial for resectable esophageal adenocarcinoma Journal of Clinical Oncology, 2020, 38, 4556-4556.	0.8	1
15	The effect of having Christmas dinner with in-laws on gut microbiota composition. Human Microbiome Journal, 2019, 13, 100058.	3.8	0
16	Rapid diagnosis of lung infections. Nature Biotechnology, 2019, 37, 725-726.	9.4	3
17	Does disease start in the mouth, the gut or both?. ELife, 2019, 8, .	2.8	11
18	Effect of Vegan Fecal Microbiota Transplantation on Carnitine―and Cholineâ€Derived Trimethylamineâ€Nâ€Oxide Production and Vascular Inflammation in Patients With Metabolic Syndrome. Journal of the American Heart Association, 2018, 7, .	1.6	164

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19	Domain intelligible models. Methods, 2018, 149, 69-73.	1.9	4
20	Depicting the composition of gut microbiota in a population with varied ethnic origins but shared geography. Nature Medicine, 2018, 24, 1526-1531.	15.2	436
21	On the ecosystemic network of saliva in healthy young adults. ISME Journal, 2017, 11, 1218-1231.	4.4	132
22	Effect of experimental gingivitis induction and erythritol on the salivary metabolome and functional biochemistry of systemically healthy young adults. Metabolomics, 2016, 12, 1.	1.4	6
23	A Study of the Variation in the Salivary Peptide Profiles of Young Healthy Adults Acquired Using MALDI-TOF MS. PLoS ONE, 2016, 11, e0156707.	1.1	8
24	Interindividual variation, correlations, and sex-related differences in the salivary biochemistry of young healthy adults. European Journal of Oral Sciences, 2015, 123, 149-157.	0.7	83