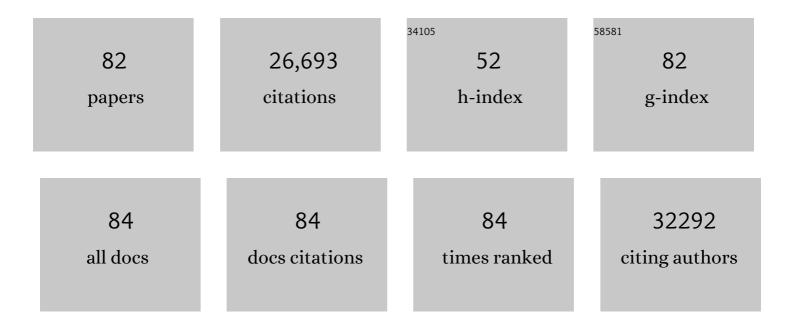
## Patricia Lepage

List of Publications by Year in descending order

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DATRICIA LERACE

#	Article	lF	CITATIONS
1	A human gut microbial gene catalogue established by metagenomic sequencing. Nature, 2010, 464, 59-65.	27.8	9,342
2	Anticancer immunotherapy by CTLA-4 blockade relies on the gut microbiota. Science, 2015, 350, 1079-1084.	12.6	2,539
3	The Intestinal Microbiota Modulates the Anticancer Immune Effects of Cyclophosphamide. Science, 2013, 342, 971-976.	12.6	1,580
4	Dietary intervention impact on gut microbial gene richness. Nature, 2013, 500, 585-588.	27.8	1,485
5	Baseline gut microbiota predicts clinical response and colitis in metastatic melanoma patients treated with ipilimumab. Annals of Oncology, 2017, 28, 1368-1379.	1.2	908
6	Resistance Mechanisms to Immune-Checkpoint Blockade in Cancer: Tumor-Intrinsic and -Extrinsic Factors. Immunity, 2016, 44, 1255-1269.	14.3	797
7	Intestinal microbiota determines development of non-alcoholic fatty liver disease in mice. Gut, 2013, 62, 1787-1794.	12.1	777
8	Enterococcus hirae and Barnesiella intestinihominis Facilitate Cyclophosphamide-Induced Therapeutic Immunomodulatory Effects. Immunity, 2016, 45, 931-943.	14.3	645
9	High-fat diet alters gut microbiota physiology in mice. ISME Journal, 2014, 8, 295-308.	9.8	583
10	Twin Study Indicates Loss of Interaction Between Microbiota and Mucosa of Patients With Ulcerative Colitis. Gastroenterology, 2011, 141, 227-236.	1.3	518
11	Intestinal microbiota contributes to individual susceptibility to alcoholic liver disease. Gut, 2016, 65, 830-839.	12.1	429
12	Specificities of the fecal microbiota in inflammatory bowel disease. Inflammatory Bowel Diseases, 2006, 12, 106-111.	1.9	373
13	Earlyâ€life establishment of the swine gut microbiome and impact on host phenotypes. Environmental Microbiology Reports, 2015, 7, 554-569.	2.4	320
14	Phylogenetic network analysis applied to pig gut microbiota identifies an ecosystem structure linked with growth traits. ISME Journal, 2016, 10, 2973-2977.	9.8	308
15	A metagenomic insight into our gut's microbiome. Gut, 2013, 62, 146-158.	12.1	302
16	Involvement of tissue bacteria in the onset of diabetes in humans: evidence for a concept. Diabetologia, 2011, 54, 3055-3061.	6.3	283
17	Resilience of the Dominant Human Fecal Microbiota upon Short-Course Antibiotic Challenge. Journal of Clinical Microbiology, 2005, 43, 5588-5592.	3.9	281
18	Systemic short chain fatty acids limit antitumor effect of CTLA-4 blockade in hosts with cancer. Nature Communications, 2020, 11, 2168.	12.8	231

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19	Biodiversity of the Mucosa-Associated Microbiota Is Stable Along the Distal Digestive Tract in Healthy Individuals and Patients With Ibd. Inflammatory Bowel Diseases, 2005, 11, 473-480.	1.9	220
20	Dysbiosis in inflammatory bowel disease: a role for bacteriophages?. Gut, 2008, 57, 424-425.	12.1	206
21	Microbial ecology perturbation in human IgA deficiency. Science Translational Medicine, 2018, 10, .	12.4	206
22	Enterocolitis due to immune checkpoint inhibitors: a systematic review. Gut, 2018, 67, 2056-2067.	12.1	179
23	Replication of Obesity and Associated Signaling Pathways Through Transfer of Microbiota From Obese-Prone Rats. Diabetes, 2014, 63, 1624-1636.	0.6	171
24	Intestinal microbiota in metabolic diseases. Gut Microbes, 2014, 5, 544-551.	9.8	170
25	The gut microbiota drives the impact of bile acids and fat source in diet on mouse metabolism. Microbiome, 2018, 6, 134.	11.1	169
26	Intestinal TM7 bacterial phylogenies in active inflammatory bowel disease. Journal of Medical Microbiology, 2008, 57, 1569-1576.	1.8	164
27	Intestinal inhibition of Atg7 prevents tumour initiation through a microbiome-influenced immune response and suppresses tumour growth. Nature Cell Biology, 2015, 17, 1062-1073.	10.3	154
28	Review article: the role of bacteria in onset and perpetuation of inflammatory bowel disease. Alimentary Pharmacology and Therapeutics, 2006, 24, 11-18.	3.7	150
29	Epidemiology of inflammatory bowel disease in a German twin cohort: Results of a nationwide study. Inflammatory Bowel Diseases, 2008, 14, 968-976.	1.9	137
30	Temperature Gradient Gel Electrophoresis of Fecal 16S rRNA Reveals Active Escherichia coli in the Microbiota of Patients with Ulcerative Colitis. Journal of Clinical Microbiology, 2006, 44, 3172-3177.	3.9	131
31	Isoflavones and Functional Foods Alter the Dominant Intestinal Microbiota in Postmenopausal Women. Journal of Nutrition, 2005, 135, 2786-2792.	2.9	129
32	Gut bacteria are critical for optimal muscle function: a potential link with glucose homeostasis. American Journal of Physiology - Endocrinology and Metabolism, 2019, 317, E158-E171.	3.5	126
33	Transcriptional activity of the dominant gut mucosal microbiota in chronic inflammatory bowel disease patients. Journal of Medical Microbiology, 2010, 59, 1114-1122.	1.8	121
34	Clostridium difficile Colonization in Early Infancy Is Accompanied by Changes in Intestinal Microbiota Composition. Journal of Clinical Microbiology, 2011, 49, 858-865.	3.9	120
35	Chemotherapy-induced ileal crypt apoptosis and the ileal microbiome shape immunosurveillance and prognosis of proximal colon cancer. Nature Medicine, 2020, 26, 919-931.	30.7	118
36	Colon Immune-Related Adverse Events: Anti-CTLA-4 and Anti-PD-1 Blockade Induce Distinct Immunopathological Entities. Journal of Crohn's and Colitis, 2017, 11, 1238-1246.	1.3	116

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37	Enteric Delivery of Regenerating Family Member 3 alpha AltersÂthe Intestinal Microbiota and Controls Inflammation inÂMice WithÂColitis. Gastroenterology, 2018, 154, 1009-1023.e14.	1.3	107
38	Structural robustness of the gut mucosal microbiota is associated with Crohn's disease remission after surgery. Gut, 2016, 65, 954-962.	12.1	106
39	Gut flora and inflammatory bowel disease. Alimentary Pharmacology and Therapeutics, 2004, 20, 18-23.	3.7	102
40	Multi-hit early life adversity affects gut microbiota, brain and behavior in a sex-dependent manner. Brain, Behavior, and Immunity, 2019, 80, 179-192.	4.1	102
41	Gut microbiome and anticancer immune response: really hot Sh*t!. Cell Death and Differentiation, 2015, 22, 199-214.	11.2	100
42	Synergistic convergence of microbiota-specific systemic IgG and secretory IgA. Journal of Allergy and Clinical Immunology, 2019, 143, 1575-1585.e4.	2.9	86
43	Hemidesmosome integrity protects the colon against colitis and colorectal cancer. Gut, 2017, 66, 1748-1760.	12.1	84
44	Mucosal Healing and Bacterial Composition in Response to Enteral Nutrition Vs Steroid-based Induction Therapy—A Randomised Prospective Clinical Trial in Children With Crohn's Disease. Journal of Crohn's and Colitis, 2019, 13, 846-855.	1.3	82
45	Rapid analysis of bile acids in different biological matrices using LC-ESI-MS/MS for the investigation of bile acid transformation by mammalian gut bacteria. Analytical and Bioanalytical Chemistry, 2017, 409, 1231-1245.	3.7	81
46	Altered gut microbiota composition in immune-impaired <i>Nod2</i> <sup>â^'/â^'</sup> mice. Gut, 2012, 61, 634-635.	12.1	80
47	Bacterial protein signals are associated with Crohn's disease. Gut, 2014, 63, 1566-1577.	12.1	80
48	Search for Localized Dysbiosis in Crohn's Disease Ulcerations by Temporal Temperature Gradient Gel Electrophoresis of 16S rRNA. Journal of Clinical Microbiology, 2005, 43, 4654-4658.	3.9	76
49	Nutritional strategies and gut microbiota composition as risk factors for necrotizing enterocolitis in very-preterm infants. American Journal of Clinical Nutrition, 2017, 106, 821-830.	4.7	71
50	The Human Gut Microbiome and Its Dysfunctions. Digestive Diseases, 2013, 31, 278-285.	1.9	65
51	Modulation of the microbiota by oral antibiotics treats immunoglobulin A nephropathy in humanized mice. Nephrology Dialysis Transplantation, 2019, 34, 1135-1144.	0.7	59
52	TREM-1 Inhibition Restores Impaired Autophagy Activity and Reduces Colitis in Mice. Journal of Crohn's and Colitis, 2018, 12, 230-244.	1.3	55
53	Patchy distribution of mucosal lesions in ileal Crohn's disease is not linked to differences in the dominant mucosa-associated bacteria. Inflammatory Bowel Diseases, 2007, 13, 684-692.	1.9	54
54	Cellular and Physiological Effects of Probiotics and Prebiotics. Mini-Reviews in Medicinal Chemistry, 2004, 4, 889-896.	2.4	47

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55	<i>Nod2</i> Deficiency Leads to a Specific and Transmissible Mucosa-associated Microbial Dysbiosis Which Is Independent of the Mucosal Barrier Defect. Journal of Crohn's and Colitis, 2016, 10, 1428-1436.	1.3	45
56	Enhancing the clinical coverage and anticancer efficacy of immune checkpoint blockade through manipulation of the gut microbiota. OncoImmunology, 2017, 6, e1132137.	4.6	45
57	Extensive Expression Differences along Porcine Small Intestine Evidenced by Transcriptome Sequencing. PLoS ONE, 2014, 9, e88515.	2.5	44
58	Assessment of Neonatal Intensive Care Unit Practices and Preterm Newborn Gut Microbiota and 2-Year Neurodevelopmental Outcomes. JAMA Network Open, 2020, 3, e2018119.	5.9	44
59	Molecular comparison of dominant microbiota associated with injured versus healthy mucosa in ulcerative colitis. Gut, 2007, 56, 152-154.	12.1	40
60	A Guide for Ex Vivo Handling and Storage of Stool Samples Intended for Fecal Microbiota Transplantation. Scientific Reports, 2019, 9, 8897.	3.3	40
61	Harnessing the Intestinal Microbiome for Optimal Therapeutic Immunomodulation. Cancer Research, 2014, 74, 4217-4221.	0.9	39
62	Impact and consequences of intensive chemotherapy on intestinal barrier and microbiota in acute myeloid leukemia: the role of mucosal strengthening. Gut Microbes, 2020, 12, 1800897.	9.8	38
63	Host–microbe interactions in distal airways: relevance to chronic airway diseases. European Respiratory Review, 2015, 24, 78-91.	7.1	35
64	Microbiota Is Involved in Post-resection Adaptation in Humans with Short Bowel Syndrome. Frontiers in Physiology, 2017, 8, 224.	2.8	35
65	Late weaning is associated with increased microbial diversity and Faecalibacterium prausnitzii abundance in the fecal microbiota of piglets. Animal Microbiome, 2020, 2, 2.	3.8	31
66	The Family Coriobacteriaceae. , 2014, , 201-238.		31
67	Does Our Food (Environment) Change Our Gut Microbiome (â€~In-Vironment'): A Potential Role for Inflammatory Bowel Disease?. Digestive Diseases, 2012, 30, 33-39.	1.9	25
68	Dietary Protein Intake Level Modulates Mucosal Healing and Mucosa-Adherent Microbiota in Mouse Model of Colitis. Nutrients, 2019, 11, 514.	4.1	25
69	Respective Roles of Hematopoietic and Nonhematopoietic Nod2 on the Gut Microbiota and Mucosal Homeostasis. Inflammatory Bowel Diseases, 2016, 22, 763-773.	1.9	24
70	Association of germ-free mice with a simplified human intestinal microbiota results in a shortened intestine. Gut Microbes, 2014, 5, 176-182.	9.8	22
71	Inflammatory bowel disease and cancer response due to anti-CTLA-4: is it in the flora?. Seminars in Immunopathology, 2017, 39, 327-331.	6.1	22
72	Mucosal healing progression after acute colitis in mice. World Journal of Gastroenterology, 2019, 25, 3572-3589.	3.3	21

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73	Mining gut microbiome oligopeptides by functional metaproteome display. Scientific Reports, 2016, 6, 34337.	3.3	19
74	Effects of enteral polymeric diet on gut microbiota in children with Crohn's disease. Gut, 2017, 66, 194-195.	12.1	19
75	Metabolic Interplay between Gut Bacteria and Their Host. Frontiers of Hormone Research, 2014, 42, 73-82.	1.0	18
76	Why should we need the gut microbiota to respond to cancer therapies?. OncoImmunology, 2014, 3, e27574.	4.6	17
77	Perturbed Microbiota/Immune Homeostasis in Multiple Sclerosis. Neurology: Neuroimmunology and NeuroInflammation, 2021, 8, e997.	6.0	15
78	Immunome differences between porcine ileal and jejunal Peyer's patches revealed by global transcriptome sequencing of gut-associated lymphoid tissues. Scientific Reports, 2018, 8, 9077.	3.3	12
79	<i>Porphyromonas</i> , a potential predictive biomarker of <i>Pseudomonas aeruginosa</i> pulmonary infection in cystic fibrosis. BMJ Open Respiratory Research, 2019, 6, e000374.	3.0	12
80	Statement of Retraction. Replication of Obesity and Associated Signaling Pathways Through Transfer of Microbiota From Obese-Prone Rats. Diabetes 2014;63:1624–1636. DOI: 10.2337/db13-1526. Diabetes, 201 65, 1447-1447.	6,0.6	4
81	High engraftment capacity of frozen ready-to-use human fecal microbiota transplants assessed in germ-free mice. Scientific Reports, 2021, 11, 4365.	3.3	2
82	Gut Microbiota Diversity of Preterm Neonates Is Associated With Clostridioides Difficile Colonization. Frontiers in Cellular and Infection Microbiology, 0, 12, .	3.9	2