## Swapna Joshi

List of Publications by Year in descending order

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759233 752698 37 476 12 20 h-index citations g-index papers 37 37 37 798 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A long noncoding RNA signature for ulcerative colitis identifies IFNG-AS1 as an enhancer of inflammation. American Journal of Physiology - Renal Physiology, 2016, 311, G446-G457.	3.4	99
2	Association of cathepsin B gene polymorphisms with tropical calcific pancreatitis. Gut, 2006, 55, 1270-1275.	12.1	75
3	Lysine methyltransferase 2D regulates pancreatic carcinogenesis through metabolic reprogramming. Gut, 2019, 68, 1271-1286.	12.1	53
4	Transcriptomic and CRISPR/Cas9 technologies reveal FOXA2 as a tumor suppressor gene in pancreatic cancer. American Journal of Physiology - Renal Physiology, 2016, 310, G1124-G1137.	3.4	46
5	The effect of sex and irritable bowel syndrome on HPA axis response and peripheral glucocorticoid receptor expression. Psychoneuroendocrinology, 2016, 69, 67-76.	2.7	43
6	The Colonic Mucosal MicroRNAs, MicroRNA-219a-5p, and MicroRNA-338-3p Are Downregulated in Irritable Bowel Syndrome and Are Associated With Barrier Function and MAPK Signaling. Gastroenterology, 2021, 160, 2409-2422.e19.	1.3	26
7	Identification of Circulating MicroRNA Signatures in Crohnʽs Disease Using the Nanostring nCounter Technology. Inflammatory Bowel Diseases, 2016, 22, 2063-2069.	1.9	24
8	Epigenetic Mechanisms in Irritable Bowel Syndrome. Frontiers in Psychiatry, 2020, 11, 805.	2.6	23
9	Effect of Exclusion Diets on Symptom Severity and the Gut Microbiota in Patients With Irritable Bowel Syndrome. Clinical Gastroenterology and Hepatology, 2022, 20, e465-e483.	4.4	20
10	Sigmoid colon mucosal gene expression supports alterations of neuronal signaling in irritable bowel syndrome with constipation. American Journal of Physiology - Renal Physiology, 2018, 315, G140-G157.	3 <b>.</b> 4	18
11	MKAD-21 Suppresses the Oncogenic Activity of the miR-21/PPP2R2A/ERK Molecular Network in Bladder Cancer. Molecular Cancer Therapeutics, 2018, 17, 1430-1440.	4.1	17
12	Comprehensive screening for бreg1l̂±ï¿½'½'g gene rules out association with tropical calcific pancreatitis. World Journal of Gastroenterology, 2007, 13, 5938.	3.3	15
13	1090 - Epigenetic Changes in Blood Cells and Colonic Mucosa are Associated with Irritable Bowel Syndrome (IBS). Gastroenterology, 2018, 154, S-214.	1.3	4
14	The visceral sensitivity index: A novel tool for measuring Glâ€symptomâ€specific anxiety in inflammatory bowel disease. Neurogastroenterology and Motility, 2022, 34, e14384.	3.0	4
15	Dysregulation of the Long-Noncoding RNA, Ghrlos, in Irritable Bowel Syndrome. Gastroenterology, 2017, 152, S722.	1.3	3
16	Guanylate Cyclase-C Receptor and Ligand Expression in Colonic Mucosa in Chronic Constipation. American Journal of Gastroenterology, 2014, 109, S540.	0.4	2
17	Colonic Mucosal Microbiome is Associated with Mucosal Microrna Expression in Irritable Bowel Syndrome. Gastroenterology, 2017, 152, S40-S41.	1.3	1
18	Expression Profiling of Sigmoid Biopsies in Irritable Bowel Syndrome vs Healthy Controls. Gastroenterology, 2017, 152, S722.	1.3	1

#	Article	IF	CITATIONS
19	H3K4me3 Affects Glucose Metabolism and Lipid Content in Pancreatic Cancer. Gastroenterology, 2017, 152, S115.	1.3	1
20	Tu1895 – Colonic Mucosal Microbiome Correlates with Dietary Intake in IBS Patients and Healthy Controls. Gastroenterology, 2019, 156, S-1164.	1.3	1
21	129 Identification of a LncRNA Signature in Ulcerative Colitis: IFNG-AS1 Is a CD4+ T-Cell LncRNA Associated With IBD SNP Loci. Gastroenterology, 2016, 150, S31.	1.3	O
22	Tu1794 Functional Pathways Associated With Differential Colonic Mucosal Expression of microRNA and mRNA in Irritable Bowel Syndrome. Gastroenterology, 2016, 150, S949.	1.3	0
23	338 Interplay Between DNA Methylation and KMT2D Histone Methyltransferase Regulates Pancreatic Cellular Growth Through a Glucose Metabolic Shift. Gastroenterology, 2016, 150, S80.	1.3	O
24	Sex-Dependent Alterations of Colonic Epithelial Permeability in Irritable Bowel Syndrome. Gastroenterology, 2017, 152, S723.	1.3	0
25	AODTH-010â€A mirna-epigenetic network in pancreatic cancer. , 2017, , .		O
26	Sa1598 - Dna Methylation Age Acceleration is Associated with Decreased Resilience in Irritable Bowel Syndrome. Gastroenterology, 2018, 154, S-325.	1.3	0
27	573 - Gene Expression Network Analysis of the Gut-Brain Axis Supports an Association Between Alpha-Synuclein and Markers of Enteric Glial Cells. Gastroenterology, 2018, 154, S-117-S-118.	1.3	O
28	Su1610 – Dysregulation of Parkinson's Disease Related Genes in Ulcerative Colitis and Murine Experimental Colitis. Gastroenterology, 2019, 156, S-582-S-583.	1.3	0
29	Tu1894 – Colonic Mucosal Microbiome is Associated with Bowel Habit Subtypes in Irritable Bowel Syndrome (IBS) Patients. Gastroenterology, 2019, 156, S-1163-S-1164.	1.3	0
30	Sa1874 – The Visceral Sensitivity Index: A Novel Tool for Measuring Gi-Symptom Specific Anxiety in Inflammatory Bowel Disease (IBD). Gastroenterology, 2019, 156, S-437-S-438.	1.3	0
31	Su1613 – The Association of Visceral Adiposity with Irritable Bowel Syndrome, Symptom Severity, and the Hypothalamic-Pituitaryadrenal Axis Response. Gastroenterology, 2019, 156, S-584.	1.3	0
32	Mo1956 ASSOCIATION OF FECAL MICROBIOME WITH RESILIENCE IN IRRITABLE BOWEL SYNDROME PATIENTS COMPARED TO HEALTHY CONTROLS. Gastroenterology, 2020, 158, S-992.	1.3	0
33	718 EFFECT OF EXCLUSION DIETS ON SYMPTOM SEVERITY AND GUT MICROBIOTA IN PATIENTS WITH IRRITABLE BOWEL SYNDROME (IBS). Gastroenterology, 2020, 158, S-151.	1.3	O
34	Mo1569 IDENTIFICATION OF COLONIC MUCOSAL MICRORNAS ALTERED IN IRRITABLE BOWEL SYNDROME AND THEIR ROLES IN INTESTINAL BARRIER FUNCTION Gastroenterology, 2020, 158, S-899.	1.3	0
35	647 FECAL MICRORNAS ARE ASSOCIATED WITH TGF-BETA AND GABA SIGNALING IN C. DIFFICILE INFECTION. Gastroenterology, 2020, 158, S-140.	1.3	O
36	Abstract 4528: Chromatin regulation by ING3 leads to tumor suppressive effects in pancreatic cancer through distinct signaling pathways. , 2016, , .		0

#	Article	lF	CITATIONS
37	Use Of Weighted Gene Coexpression Network Analysis To Identify Connectivity Between Gut And Brain Gene Expression. FASEB Journal, 2022, 36, .	0.5	O