Lila E Mullany

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7629733/publications.pdf

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36	1,371	23	36
papers	citations	h-index	g-index
36	36	36	2515
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	The p53-signaling pathway and colorectal cancer: Interactions between downstream p53 target genes and miRNAs. Genomics, 2019, 111, 762-771.	2.9	80
2	The functional role of miRNAs in colorectal cancer: insights from a large population-based study. Cancer Biology and Medicine, 2019, 16, 211.	3.0	8
3	The MAPK-Signaling Pathway in Colorectal Cancer: Dysregulated Genes and Their Association With MicroRNAs. Cancer Informatics, 2018, 17, 117693511876652.	1.9	45
4	Dysregulated genes and miRNAs in the apoptosis pathway in colorectal cancer patients. Apoptosis: an International Journal on Programmed Cell Death, 2018, 23, 237-250.	4.9	73
5	The PI3K/AKT signaling pathway: Associations of miRNAs with dysregulated gene expression in colorectal cancer. Molecular Carcinogenesis, 2018, 57, 243-261.	2.7	83
6	The NF- $\hat{l}^{\circ}B$ signalling pathway in colorectal cancer: associations between dysregulated gene and miRNA expression. Journal of Cancer Research and Clinical Oncology, 2018, 144, 269-283.	2.5	71
7	MicroRNAâ€transcription factor interactions and their combined effect on target gene expression in colon cancer cases. Genes Chromosomes and Cancer, 2018, 57, 192-202.	2.8	42
8	The TGF \hat{l}^2 -signaling pathway and colorectal cancer: associations between dysregulated genes and miRNAs. Journal of Translational Medicine, 2018, 16, 191.	4.4	35
9	miRNA involvement in cell cycle regulation in colorectal cancer cases. Genes and Cancer, 2018, 9, 53-65.	1.9	29
10	MicroRNA-messenger RNA interactions involving JAK-STAT signaling genes in colorectal cancer. Genes and Cancer, 2018, 9, 232-246.	1.9	6
11	Expression of Wnt-signaling pathway genes and their associations with miRNAs in colorectal cancer. Oncotarget, 2018, 9, 6075-6085.	1.8	17
12	Diet and lifestyle factors associated with miRNA expression in colorectal tissue. Pharmacogenomics and Personalized Medicine, 2017, Volume10, 1-16.	0.7	28
13	Infrequently expressed miRNAs in colorectal cancer tissue and tumor molecular phenotype. Modern Pathology, 2017, 30, 1152-1169.	5.5	17
14	Alterations in microRNA expression associated with alcohol consumption in rectal cancer subjects. Cancer Causes and Control, 2017, 28, 545-555.	1.8	7
15	The miRNA landscape of colorectal polyps. Genes Chromosomes and Cancer, 2017, 56, 347-353.	2.8	8
16	Transcription factorâ€microRNA associations and their impact on colorectal cancer survival. Molecular Carcinogenesis, 2017, 56, 2512-2526.	2.7	13
17	The coâ€regulatory networks of tumor suppressor genes, oncogenes, and miRNAs in colorectal cancer. Genes Chromosomes and Cancer, 2017, 56, 769-787.	2.8	67
18	Single nucleotide polymorphisms within MicroRNAs, MicroRNA targets, and MicroRNA biogenesis genes and their impact on colorectal cancer survival. Genes Chromosomes and Cancer, 2017, 56, 285-295.	2.8	21

#	Article	IF	Citations
19	An Assessment of Database-Validated microRNA Target Genes in Normal Colonic Mucosa: Implications for Pathway Analysis. Cancer Informatics, 2017, 16, 117693511771640.	1.9	9
20	Genetic variants in the TGF \hat{l}^2 -signaling pathway influence expression of miRNAs in colon and rectal normal mucosa and tumor tissue. Oncotarget, 2017, 8, 16765-16783.	1.8	25
21	Infrequently expressed miRNAs influence survival after diagnosis with colorectal cancer. Oncotarget, 2017, 8, 83845-83859.	1.8	28
22	Dietary intake alters gene expression in colon tissue. Pharmacogenetics and Genomics, 2016, 26, 294-306.	1.5	18
23	Colorectal tumor molecular phenotype and miRNA: expression profiles and prognosis. Modern Pathology, 2016, 29, 915-927.	5.5	41
24	Expression Profiles of miRNA Subsets Distinguish Human Colorectal Carcinoma and Normal Colonic Mucosa. Clinical and Translational Gastroenterology, 2016, 7, e152.	2.5	82
25	Association of cigarette smoking and microRNA expression in rectal cancer: Insight into tumor phenotype. Cancer Epidemiology, 2016, 45, 98-107.	1.9	36
26	Impact of polymorphisms in microRNA biogenesis genes on colon cancer risk and microRNA expression levels: a population-based, case-control study. BMC Medical Genomics, 2016, 9, 21.	1.5	33
27	MicroRNA profiles in colorectal carcinomas, adenomas and normal colonic mucosa: variations in miRNA expression and disease progression. Carcinogenesis, 2016, 37, 245-261.	2.8	107
28	MicroRNA Seed Region Length Impact on Target Messenger RNA Expression and Survival in Colorectal Cancer. PLoS ONE, 2016, 11, e0154177.	2.5	52
29	Telomere Length, TERT, and miRNA Expression. PLoS ONE, 2016, 11, e0162077.	2.5	14
30	Site-specific associations between miRNA expression and survival in colorectal cancer cases. Oncotarget, 2016, 7, 60193-60205.	1.8	41
31	Gene expression in colon cancer: A focus on tumor site and molecular phenotype. Genes Chromosomes and Cancer, 2015, 54, 527-541.	2.8	39
32	Differential Gene Expression in Colon Tissue Associated With Diet, Lifestyle, and Related Oxidative Stress. PLoS ONE, 2015, 10, e0134406.	2.5	26
33	SNP Regulation of microRNA Expression and Subsequent Colon Cancer Risk. PLoS ONE, 2015, 10, e0143894.	2.5	25
34	Effectiveness and Usability of Bioinformatics Tools to Analyze Pathways Associated with miRNA Expression. Cancer Informatics, 2015, 14, CIN.S32716.	1.9	8
35	Improved survival among colon cancer patients with increased differentially expressed pathways. BMC Medicine, 2015, 13, 75.	5.5	18
36	An evaluation and replication of mi <scp>RNA</scp> s with disease stage and colorectal cancerâ€specific mortality. International Journal of Cancer, 2015, 137, 428-438.	5.1	119