

Philip A Gregory

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

47
papers

8,569
citations

29
h-index

49
g-index

49
ext. papers

9,534
ext. citations

8.7
avg, IF

5.47
L-index

#	Paper	IF	Citations
47	The miR-200 family and miR-205 regulate epithelial to mesenchymal transition by targeting ZEB1 and SIP1. <i>Nature Cell Biology</i> , 2008 , 10, 593-601	23.4	3060
46	The RNA binding protein quaking regulates formation of circRNAs. <i>Cell</i> , 2015 , 160, 1125-34	56.2	1206
45	A double-negative feedback loop between ZEB1-SIP1 and the microRNA-200 family regulates epithelial-mesenchymal transition. <i>Cancer Research</i> , 2008 , 68, 7846-54	10.1	852
44	An autocrine TGF-beta/ZEB/miR-200 signaling network regulates establishment and maintenance of epithelial-mesenchymal transition. <i>Molecular Biology of the Cell</i> , 2011 , 22, 1686-98	3.5	425
43	MicroRNAs as regulators of epithelial-mesenchymal transition. <i>Cell Cycle</i> , 2008 , 7, 3112-8	4.7	417
42	Contextual extracellular cues promote tumor cell EMT and metastasis by regulating miR-200 family expression. <i>Genes and Development</i> , 2009 , 23, 2140-51	12.6	376
41	Glucuronidation and the UDP-glucuronosyltransferases in health and disease. <i>Drug Metabolism and Disposition</i> , 2004 , 32, 281-90	4	209
40	The Notch ligand Jagged2 promotes lung adenocarcinoma metastasis through a miR-200-dependent pathway in mice. <i>Journal of Clinical Investigation</i> , 2011 , 121, 1373-85	15.9	154
39	Epigenetic modulation of the miR-200 family is associated with transition to a breast cancer stem-cell-like state. <i>Journal of Cell Science</i> , 2013 , 126, 2256-66	5.3	150
38	Mutant p53 drives invasion in breast tumors through up-regulation of miR-155. <i>Oncogene</i> , 2013 , 32, 2992-3000	12.4	124
37	ZEB1 drives prometastatic actin cytoskeletal remodeling by downregulating miR-34a expression. <i>Journal of Clinical Investigation</i> , 2012 , 122, 3170-83	15.9	119
36	Genome-wide identification of miR-200 targets reveals a regulatory network controlling cell invasion. <i>EMBO Journal</i> , 2014 , 33, 2040-56	13	112
35	MiRNA-205 modulates cellular invasion and migration via regulating zinc finger E-box binding homeobox 2 expression in esophageal squamous cell carcinoma cells. <i>Journal of Translational Medicine</i> , 2011 , 9, 30	8.5	111
34	Regulation of UDP glucuronosyltransferase genes. <i>Current Drug Metabolism</i> , 2003 , 4, 249-57	3.5	104
33	The role of microRNAs in metastasis and epithelial-mesenchymal transition. <i>Cellular and Molecular Life Sciences</i> , 2009 , 66, 1682-99	10.3	102
32	MiR-200 can repress breast cancer metastasis through ZEB1-independent but moesin-dependent pathways. <i>Oncogene</i> , 2014 , 33, 4077-88	9.2	95
31	Regulation of UDP glucuronosyltransferases in the gastrointestinal tract. <i>Toxicology and Applied Pharmacology</i> , 2004 , 199, 354-63	4.6	95

30	Mechanisms of vitamin D metabolite repression of IgE-dependent mast cell activation. <i>Journal of Allergy and Clinical Immunology</i> , 2014 , 133, 1356-64, 1364.e1-14	11.5	79
29	MicroRNA-194 Promotes Prostate Cancer Metastasis by Inhibiting SOCS2. <i>Cancer Research</i> , 2017 , 77, 1021-1034	10.1	74
28	A ZEB1-miR-375-YAP1 pathway regulates epithelial plasticity in prostate cancer. <i>Oncogene</i> , 2017 , 36, 24-34	9.2	73
27	The microRNA-200 family regulates epithelial to mesenchymal transition. <i>Scientific World Journal, The</i> , 2008 , 8, 901-4	2.2	63
26	Coordinate regulation of the human UDP-glucuronosyltransferase 1A8, 1A9, and 1A10 genes by hepatocyte nuclear factor 1alpha and the caudal-related homeodomain protein 2. <i>Molecular Pharmacology</i> , 2004 , 65, 953-63	4.3	63
25	Combinatorial Targeting by MicroRNAs Co-ordinates Post-transcriptional Control of EMT. <i>Cell Systems</i> , 2018 , 7, 77-91.e7	10.6	52
24	Specificity protein 1 (Sp1) maintains basal epithelial expression of the miR-200 family: implications for epithelial-mesenchymal transition. <i>Journal of Biological Chemistry</i> , 2014 , 289, 11194-11205	5.4	49
23	Regulation of vascular leak and recovery from ischemic injury by general and VE-cadherin-restricted miRNA antagonists of miR-27. <i>Blood</i> , 2013 , 122, 2911-9	2.2	48
22	Neuropilin-1 is upregulated in the adaptive response of prostate tumors to androgen-targeted therapies and is prognostic of metastatic progression and patient mortality. <i>Oncogene</i> , 2017 , 36, 3417-3427	8.27	47
21	miR-200/375 control epithelial plasticity-associated alternative splicing by repressing the RNA-binding protein Quaking. <i>EMBO Journal</i> , 2018 , 37,	13	46
20	Polymorphic variations in the expression of the chemical detoxifying UDP glucuronosyltransferases. <i>Toxicology and Applied Pharmacology</i> , 2005 , 207, 77-83	4.6	33
19	A novel polymorphism in a forkhead box A1 (FOXA1) binding site of the human UDP glucuronosyltransferase 2B17 gene modulates promoter activity and is associated with altered levels of circulating androstane-3 β -7 β -diol glucuronide. <i>Molecular Pharmacology</i> , 2010 , 78, 714-22	4.3	29
18	Tissue specific differences in the regulation of the UDP glucuronosyltransferase 2B17 gene promoter. <i>Pharmacogenetics and Genomics</i> , 2000 , 10, 809-20		29
17	The homeodomain Pbx2-Prep1 complex modulates hepatocyte nuclear factor 1alpha-mediated activation of the UDP-glucuronosyltransferase 2B17 gene. <i>Molecular Pharmacology</i> , 2002 , 62, 154-61	4.3	27
16	Cloning and characterization of the human UDP-glucuronosyltransferase 1A8, 1A9, and 1A10 gene promoters: differential regulation through an interior-like region. <i>Journal of Biological Chemistry</i> , 2003 , 278, 36107-14	5.4	25
15	Chromatinized protein kinase C directly regulates inducible genes in epithelial to mesenchymal transition and breast cancer stem cells. <i>Molecular and Cellular Biology</i> , 2014 , 34, 2961-80	4.8	24
14	The caudal-related homeodomain protein Cdx2 and hepatocyte nuclear factor 1alpha cooperatively regulate the UDP-glucuronosyltransferase 2B7 gene promoter. <i>Pharmacogenetics and Genomics</i> , 2006 , 16, 527-36	1.9	19
13	Epithelial plasticity in prostate cancer: principles and clinical perspectives. <i>Trends in Molecular Medicine</i> , 2014 , 20, 643-51	11.5	18

12	Identification and characterization of functional hepatocyte nuclear factor 1-binding sites in UDP-glucuronosyltransferase genes. <i>Methods in Enzymology</i> , 2005 , 400, 22-46	1.7	14
11	Regulation of splicing and circularisation of RNA in epithelial mesenchymal plasticity. <i>Seminars in Cell and Developmental Biology</i> , 2018 , 75, 50-60	7.5	12
10	Post-transcriptional Gene Regulation by MicroRNA-194 Promotes Neuroendocrine Transdifferentiation in Prostate Cancer. <i>Cell Reports</i> , 2021 , 34, 108585	10.6	10
9	MicroRNA-143-3p targets pyruvate carboxylase expression and controls proliferation and migration of MDA-MB-231 cells. <i>Archives of Biochemistry and Biophysics</i> , 2019 , 677, 108169	4.1	8
8	Insufficiently complex unique-molecular identifiers (UMIs) distort small RNA sequencing. <i>Scientific Reports</i> , 2020 , 10, 14593	4.9	5
7	Extensive transcriptional responses are co-ordinated by microRNAs as revealed by Exon-Intron Split Analysis (EISA). <i>Nucleic Acids Research</i> , 2019 , 47, 8606-8619	20.1	3
6	The miR-200-Quaking axis functions in tumour angiogenesis. <i>Oncogene</i> , 2019 , 38, 6767-6769	9.2	3
5	Polymorphisms in the mitochondrial ribosome recycling factor EF-G2mt/MEF2 compromise cell respiratory function and increase atorvastatin toxicity. <i>PLoS Genetics</i> , 2012 , 8, e1002755	6	3
4	Selective Microfluidic Capture and Detection of Prostate Cancer Cells from Urine without Digital Rectal Examination. <i>Cancers</i> , 2021 , 13,	6.6	1
3	Post-transcriptional control of EMT is coordinated through combinatorial targeting by multiple microRNAs		1
2	Neuropilin-1 is over-expressed in claudin-low breast cancer and promotes tumor progression through acquisition of stem cell characteristics and RAS/MAPK pathway activation.. <i>Breast Cancer Research</i> , 2022 , 24, 8	8.3	0
1	The Quaking RNA-binding proteins as regulators of cell differentiation.. <i>Wiley Interdisciplinary Reviews RNA</i> , 2022 , e1724	9.3	0