## Shamika Ketkar

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Somatic Dnmt3a inactivation leads to slow, canonical DNA methylation loss in murine hematopoietic cells. IScience, 2022, 25, 104004.	4.1	2
2	A novel, de novo intronic variant in <scp><i>POGZ</i></scp> causes <scp>White–Sutton</scp> syndrome. American Journal of Medical Genetics, Part A, 2022, 188, 2198-2203.	1.2	4
3	<scp><i>miRNAâ€34c</i></scp> Suppresses Osteosarcoma Progression In Vivo by Targeting Notch and <scp>E2F</scp> . JBMR Plus, 2022, 6, e10623.	2.7	8
4	Transcriptome-directed analysis for Mendelian disease diagnosis overcomes limitations of conventional genomic testing. Journal of Clinical Investigation, 2021, 131, .	8.2	87
5	Tumor suppressor function of <i>WT1</i> in acute promyelocytic leukemia. Haematologica, 2021, , .	3.5	4
6	<i>Dnmt3a</i> Inactivation Leads to Slow DNA Methylation Loss in Murine Hematopoietic Cells <i>In Vivo</i> . Blood, 2021, 138, 1087-1087.	1.4	0
7	The safety and clinical effects of administering a multiantigen-targeted T cell therapy to patients with multiple myeloma. Science Translational Medicine, 2020, 12, .	12.4	25
8	Remethylation of <i>Dnmt3a</i> <sup>â^'/â^'</sup> hematopoietic cells is associated with partial correction of gene dysregulation and reduced myeloid skewing. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 3123-3134.	7.1	27
9	Leukocyte-dependent effects of platelet-rich plasma on cartilage loss and thermal hyperalgesia in a mouse model of post-traumatic osteoarthritis. Osteoarthritis and Cartilage, 2020, 28, 1385-1393.	1.3	17
10	DNMT3AR882 Alters the Epigenome of Hematopoietic Cells. Blood, 2019, 134, 112-112.	1.4	0
11	Haploinsufficiency for DNA methyltransferase 3A predisposes hematopoietic cells to myeloid malignancies. Journal of Clinical Investigation, 2017, 127, 3657-3674.	8.2	80
12	PML-RARA requires DNA methyltransferase 3A to initiate acute promyelocytic leukemia. Journal of Clinical Investigation, 2015, 126, 85-98.	8.2	36
13	Identification of heart rate–associated loci and their effects on cardiac conduction and rhythm disorders. Nature Genetics, 2013, 45, 621-631.	21.4	282
14	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. Nature Genetics, 2013, 45, 690-696.	21.4	232
15	Genome-Wide Association and Functional Follow-Up Reveals New Loci for Kidney Function. PLoS Genetics, 2012, 8, e1002584.	3.5	166
16	Integration of genome-wide association studies with biological knowledge identifies six novel genes related to kidney function. Human Molecular Genetics, 2012, 21, 5329-5343.	2.9	64
17	Association of gene variants with lipid levels in response to fenofibrate is influenced by metabolic syndrome status. Atherosclerosis, 2011, 215, 435-439.	0.8	19
18	Hundreds of variants clustered in genomic loci and biological pathways affect human height. Nature, 2010, 467, 832-838.	27.8	1,789

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#	Article	IF	CITATIONS
19	New loci associated with kidney function and chronic kidney disease. Nature Genetics, 2010, 42, 376-384.	21.4	710
20	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. Nature Genetics, 2010, 42, 949-960.	21.4	836
21	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. Nature Genetics, 2010, 42, 937-948.	21.4	2,634
22	Sequence Variants of Toll-Like Receptor 4 and Susceptibility to Prostate Cancer. Cancer Research, 2005, 65, 11771-11778.	0.9	173