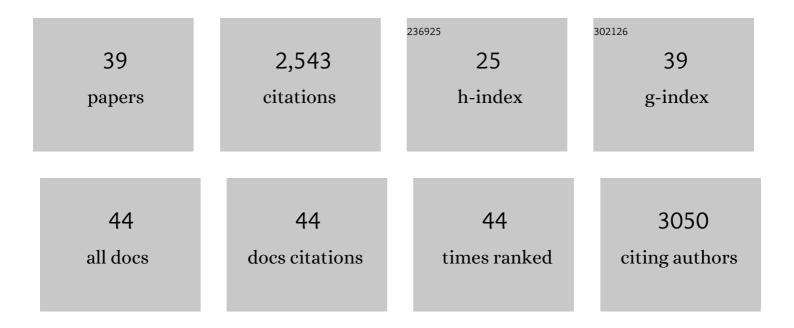
Humaira Gowher

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Mechanism of Stimulation of Catalytic Activity of Dnmt3A and Dnmt3B DNA-(cytosine-C5)-methyltransferases by Dnmt3L. Journal of Biological Chemistry, 2005, 280, 13341-13348.	3.4	250
2	Dnmt3a and Dnmt1 functionally cooperate during <i>de novo</i> methylation of DNA. FEBS Journal, 2002, 269, 4981-4984.	0.2	221
3	Enzymatic properties of recombinant Dnmt3a DNA methyltransferase from mouse: the enzyme modifies DNA in a non-processive manner and also methylates non-CpA sites. Journal of Molecular Biology, 2001, 309, 1201-1208.	4.2	217
4	Molecular Enzymology of the Catalytic Domains of the Dnmt3a and Dnmt3b DNA Methyltransferases. Journal of Biological Chemistry, 2002, 277, 20409-20414.	3.4	177
5	Chromatin Targeting of de Novo DNA Methyltransferases by the PWWP Domain. Journal of Biological Chemistry, 2004, 279, 25447-25454.	3.4	176
6	Mammalian DNA methyltransferases: new discoveries and open questions. Biochemical Society Transactions, 2018, 46, 1191-1202.	3.4	122
7	Chromatin domains, insulators, and the regulation of gene expression. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2012, 1819, 644-651.	1.9	115
8	DNMT3L facilitates DNA methylation partly by maintaining DNMT3A stability in mouse embryonic stem cells. Nucleic Acids Research, 2019, 47, 152-167.	14.5	99
9	The Escherichia coli Dam DNA Methyltransferase Modifies DNA in a Highly Processive Reaction. Journal of Molecular Biology, 2002, 319, 1085-1096.	4.2	95
10	VEZF1 Elements Mediate Protection from DNA Methylation. PLoS Genetics, 2010, 6, e1000804.	3.5	91
11	Mechanism of inhibition of DNA methyltransferases by cytidine analogs in cancer therapy. Cancer Biology and Therapy, 2004, 3, 1062-1068.	3.4	85
12	Mutational Analysis of the Catalytic Domain of the Murine Dnmt3a DNA-(cytosine) Tj ETQq0 0 0 rgBT /Overlock	10 Tf 50 30 4.2	02 ₈₃ Td (C5)-n
13	De Novo Methylation of Nucleosomal DNA by the Mammalian Dnmt1 and Dnmt3A DNA Methyltransferases. Biochemistry, 2005, 44, 9899-9904.	2.5	78
14	An epigenetic switch regulates <i>de novo</i> DNA methylation at a subset of pluripotency gene enhancers during embryonic stem cell differentiation. Nucleic Acids Research, 2016, 44, 7605-7617.	14.5	70
15	Catalytic Mechanism of DNA-(cytosine-C5)-methyltransferases Revisited: Covalent Intermediate Formation is not Essential for Methyl Group Transfer by the Murine Dnmt3a Enzyme. Journal of	4.2	69

	Molecular Biology, 2003, 329, 675-684.		
16	Molecular enzymology of the Eco RV DNA-(adenine-N6)-methyltransferase: kinetics of DNA binding and bending, kinetic mechanism and linear diffusion of the enzyme on DNA. Journal of Molecular Biology, 2000, 303, 93-110.	4.2	65
17	Extra-coding RNAs regulate neuronal DNA methylation dynamics. Nature Communications, 2016, 7, 12091.	12.8	57
18	Mutations in DNA methyltransferase DNMT3B in ICF syndrome affect its regulation by DNMT3L. Human	2.9	52

18 ^y Molecular Genetics, 2006, 15, 1375-1385.

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19	From Phenylthiazoles to Phenylpyrazoles: Broadening the Antibacterial Spectrum toward Carbapenem-Resistant Bacteria. Journal of Medicinal Chemistry, 2019, 62, 7998-8010.	6.4	41
20	Vezf1 regulates genomic DNA methylation through its effects on expression of DNA methyltransferase Dnmt3b. Genes and Development, 2008, 22, 2075-2084.	5.9	38
21	Phosphorylation of Serine-515 Activates the Mammalian Maintenance Methyltransferase Dnmt1. Epigenetics, 2007, 2, 155-160.	2.7	36
22	Vezf1 protein binding sites genome-wide are associated with pausing of elongating RNA polymerase II. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2370-2375.	7.1	35
23	Integrative genomic analysis of pediatric T-cell lymphoblastic lymphoma reveals candidates of clinical significance. Blood, 2021, 137, 2347-2359.	1.4	31
24	DNA fromAspergillus flavuscontains 5-methylcytosine. FEMS Microbiology Letters, 2001, 205, 151-155.	1.8	30
25	Avidin plate assay system for enzymatic characterization of a histone lysine methyltransferase. Analytical Biochemistry, 2005, 342, 287-291.	2.4	28
26	The transcription factor Vezf1 represses the expression of the antiangiogenic factor Cited2 in endothelial cells. Journal of Biological Chemistry, 2018, 293, 11109-11118.	3.4	26
27	Lipophilic efficient phenylthiazoles with potent undecaprenyl pyrophosphatase inhibitory activity. European Journal of Medicinal Chemistry, 2019, 175, 49-62.	5.5	24
28	Dnmt3b Methylates DNA by a Noncooperative Mechanism, and Its Activity Is Unaffected by Manipulations at the Predicted Dimer Interface. Biochemistry, 2018, 57, 4312-4324.	2.5	23
29	Effect of Disease-Associated Germline Mutations on Structure Function Relationship of DNA Methyltransferases. Genes, 2019, 10, 369.	2.4	23
30	The acute myeloid leukemia variant DNMT3A Arg882His is a DNMT3B-like enzyme. Nucleic Acids Research, 2020, 48, 3761-3775.	14.5	18
31	Oct4-Mediated Inhibition of Lsd1 Activity Promotes the Active and Primed State of Pluripotency Enhancers. Cell Reports, 2020, 30, 1478-1490.e6.	6.4	17
32	A refined DNA methylation detection method using MspJI coupled quantitative PCR. Analytical Biochemistry, 2017, 533, 1-9.	2.4	12
33	Development of Biphenylthiazoles Exhibiting Improved Pharmacokinetics and Potent Activity Against Intracellular <i>Staphylococcus aureus</i> . ACS Infectious Diseases, 2020, 6, 2887-2900.	3.8	11
34	Editorial—Role of DNA Methyltransferases in the Epigenome. Genes, 2019, 10, 574.	2.4	8
35	Misregulation of the expression and activity of DNA methyltransferases in cancer. NAR Cancer, 2021, 3, zcab045.	3.1	8
36	Characterization of Small Molecules Inhibiting the Pro-Angiogenic Activity of the Zinc Finger Transcription Factor Vezf1. Molecules, 2018, 23, 1615.	3.8	3

#	Article	IF	CITATIONS
37	Simplified MethylRAD Sequencing to Detect Changes in DNA Methylation at Enhancer Elements in Differentiating Embryonic Stem Cells. Epigenomes, 2020, 4, 24.	1.8	3
38	RyR2/IRBIT regulates insulin gene transcript, insulin content, and secretion in the insulinoma cell line INS-1. Scientific Reports, 2022, 12, 7713.	3.3	3
39	Bgp1/Vezf1 couples RNA Pol II activity and maintenance of genomic methylation. FASEB Journal, 2011, 25, lb43.	0.5	Ο