

# Richard J Gibbons

## List of Publications by Year in descending order

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108  
papers

10,283  
citations

43973

48  
h-index

35952

97  
g-index

113  
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113  
docs citations

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times ranked

11337  
citing authors

#	ARTICLE	IF	CITATIONS
1	The chromatin remodeller ATRX facilitates diverse nuclear processes, in a stochastic manner, in both heterochromatin and euchromatin. <i>Nature Communications</i> , 2022, 13, .	5.8	20
2	Genetic and functional insights into CDA-I prevalence and pathogenesis. <i>Journal of Medical Genetics</i> , 2021, 58, 185-195.	1.5	9
3	A gain-of-function single nucleotide variant creates a new promoter which acts as an orientation-dependent enhancer-blocker. <i>Nature Communications</i> , 2021, 12, 3806.	5.8	18
4	An evolutionarily ancient mechanism for regulation of hemoglobin expression in vertebrate red cells. <i>Blood</i> , 2020, 136, 269-278.	0.6	16
5	Synergistic silencing of $\hat{1}\pm$ -globin and induction of $\hat{1}^3$ -globin by histone deacetylase inhibitor, vorinostat as a potential therapy for $\hat{1}^2$ -thalassaemia. <i>Scientific Reports</i> , 2019, 9, 11649.	1.6	21
6	ATRX Contributes to MeCP2-Mediated Pericentric Heterochromatin Organization during Neural Differentiation. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5371.	1.8	19
7	The macroH2A1.2 histone variant links ATRX loss to alternative telomere lengthening. <i>Nature Structural and Molecular Biology</i> , 2019, 26, 213-219.	3.6	36
8	Persistence of skewed X-chromosome inactivation in pre-B acute lymphoblastic leukemia of a female ATRX mutation carrier. <i>Blood Advances</i> , 2019, 3, 2627-2631.	2.5	2
9	The Loss of ATRX Increases Susceptibility to Pancreatic Injury and Oncogenic KRAS in Female But Not Male Mice. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2019, 7, 93-113.	2.3	14
10	Histone Lysine Methylases and Demethylases in the Landscape of Human Developmental Disorders. <i>American Journal of Human Genetics</i> , 2018, 102, 175-187.	2.6	204
11	Phenotypic and molecular characterization of a serum-free miniature erythroid differentiation system suitable for high-throughput screening and single-cell assays. <i>Experimental Hematology</i> , 2018, 60, 10-20.	0.2	6
12	2 new cases of pontocerebellar hypoplasia type 10 identified by whole exome sequencing in a Turkish family. <i>European Journal of Medical Genetics</i> , 2018, 61, 273-279.	0.7	14
13	Mutant IDH1 Promotes Glioma Formation In Vivo. <i>Cell Reports</i> , 2018, 23, 1553-1564.	2.9	91
14	Anaemia among females in child-bearing age: Relative contributions, effects and interactions of $\hat{1}\pm$ - and $\hat{1}^2$ -thalassaemia. <i>PLoS ONE</i> , 2018, 13, e0206928.	1.1	17
15	Robust CRISPR/Cas9 Genome Editing of the HUDEP-2 Erythroid Precursor Line Using Plasmids and Single-Stranded Oligonucleotide Donors. <i>Methods and Protocols</i> , 2018, 1, 28.	0.9	17
16	How to Tackle Challenging ChIP-Seq, with Long-Range Cross-Linking, Using ATRX as an Example. <i>Methods in Molecular Biology</i> , 2018, 1832, 105-130.	0.4	7
17	Selective silencing of $\hat{1}\pm$ -globin by the histone demethylase inhibitor IOX1: a potentially new pathway for treatment of $\hat{1}^2$ -thalassaemia. <i>Haematologica</i> , 2017, 102, e80-e84.	1.7	33
18	The chromatin remodelling factor $\langle\text{sc}\rangle\text{ATRX}\langle/\text{sc}\rangle$ suppresses R-loops in transcribed telomeric repeats. <i>EMBO Reports</i> , 2017, 18, 914-928.	2.0	99

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19	Editing an $\hat{1}$ -globin enhancer in primary human hematopoietic stem cells as a treatment for $\hat{2}$ -thalassemia. Nature Communications, 2017, 8, 424.	5.8	85
20	Understanding $\hat{1}$ -globin gene regulation and implications for the treatment of $\hat{2}$ -thalassemia. Annals of the New York Academy of Sciences, 2016, 1368, 16-24.	1.8	44
21	Genetic dissection of the $\hat{1}$ -globin super-enhancer in vivo. Nature Genetics, 2016, 48, 895-903.	9.4	308
22	Maintaining memory of silencing at imprinted differentially methylated regions. Cellular and Molecular Life Sciences, 2016, 73, 1871-1879.	2.4	18
23	Suppression of the alternative lengthening of telomere pathway by the chromatin remodelling factor ATRX. Nature Communications, 2015, 6, 7538.	5.8	219
24	ATRX Plays a Key Role in Maintaining Silencing at Interstitial Heterochromatic Loci and Imprinted Genes. Cell Reports, 2015, 11, 405-418.	2.9	152
25	$\hat{1}$ -Globin as a molecular target in the treatment of $\hat{2}$ -thalassemia. Blood, 2015, 125, 3694-3701.	0.6	102
26	The Fanconi Anemia Pathway Maintains Genome Stability by Coordinating Replication and Transcription. Molecular Cell, 2015, 60, 351-361.	4.5	283
27	ATRX Dysfunction Induces Replication Defects in Primary Mouse Cells. PLoS ONE, 2014, 9, e92915.	1.1	84
28	Mutations in Kr $\hat{1}$ ppel-like factor 1 cause transfusion-dependent hemolytic anemia and persistence of embryonic globin gene expression. Blood, 2014, 123, 1586-1595.	0.6	76
29	Analysis of hundreds of cis-regulatory landscapes at high resolution in a single, high-throughput experiment. Nature Genetics, 2014, 46, 205-212.	9.4	417
30	Variant ATRX Syndrome with Dysfunction of ATRX and MAGT1 Genes. Human Mutation, 2014, 35, 58-62.	1.1	7
31	JAK2V617F promotes replication fork stalling with disease-restricted impairment of the intra-S checkpoint response. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 15190-15195.	3.3	35
32	The chromatin remodeller ATRX: a repeat offender in human disease. Trends in Biochemical Sciences, 2013, 38, 461-466.	3.7	103
33	ATRX and the replication of structured DNA. Current Opinion in Genetics and Development, 2013, 23, 289-294.	1.5	40
34	Analysis of Sequence Variation Underlying Tissue-specific Transcription Factor Binding and Gene Expression. Human Mutation, 2013, 34, 1140-1148.	1.1	10
35	High-resolution analysis of cis-acting regulatory networks at the $\hat{1}$ -globin locus. Philosophical Transactions of the Royal Society B: Biological Sciences, 2013, 368, 20120361.	1.8	12
36	GENE SILENCING. , 2013, , 143-184.		0

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37	An interspecies analysis reveals a key role for unmethylated CpG dinucleotides in vertebrate Polycomb complex recruitment. <i>EMBO Journal</i> , 2012, 31, 317-329.	3.5	173
38	Î±-Thalassemia, Mental Retardation, and Myelodysplastic Syndrome. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2012, 2, a011759-a011759.	2.9	29
39	Intragenic Enhancers Act as Alternative Promoters. <i>Molecular Cell</i> , 2012, 45, 447-458.	4.5	237
40	How genetically heterogeneous is Kabuki syndrome?: MLL2 testing in 116 patients, review and analyses of mutation and phenotypic spectrum. <i>European Journal of Human Genetics</i> , 2012, 20, 381-388.	1.4	142
41	Nprl3 is required for normal development of the cardiovascular system. <i>Mammalian Genome</i> , 2012, 23, 404-415.	1.0	38
42	Combinatorial readout of histone H3 modifications specifies localization of ATRX to heterochromatin. <i>Nature Structural and Molecular Biology</i> , 2011, 18, 777-782.	3.6	187
43	Systematic documentation and analysis of human genetic variation in hemoglobinopathies using the microattribution approach. <i>Nature Genetics</i> , 2011, 43, 295-301.	9.4	142
44	Generation of bivalent chromatin domains during cell fate decisions. <i>Epigenetics and Chromatin</i> , 2011, 4, 9.	1.8	54
45	Functional significance of mutations in the Snf2 domain of ATRX. <i>Human Molecular Genetics</i> , 2011, 20, 2603-2610.	1.4	46
46	ATRX: Taming tandem repeats. <i>Cell Cycle</i> , 2010, 9, 4605-4606.	1.3	4
47	Distinct Factors Control Histone Variant H3.3 Localization at Specific Genomic Regions. <i>Cell</i> , 2010, 140, 678-691.	13.5	1,069
48	ATR-X Syndrome Protein Targets Tandem Repeats and Influences Allele-Specific Expression in a Size-Dependent Manner. <i>Cell</i> , 2010, 143, 367-378.	13.5	365
49	The Molecular Basis of Î±-Thalassemia: A Model for Understanding Human Molecular Genetics. <i>Hematology/Oncology Clinics of North America</i> , 2010, 24, 1033-1054.	0.9	36
50	Adventitious changes in long-range gene expression caused by polymorphic structural variation and promoter competition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 21771-21776.	3.3	77
51	Research Highlights. <i>Epigenomics</i> , 2009, 1, 231-234.	1.0	0
52	Altered visual function and interneuron survival in Atrx knockout mice: inference for the human syndrome. <i>Human Molecular Genetics</i> , 2009, 18, 966-977.	1.4	34
53	The role of Xâ€œinactivation in the gender bias of patients with acquired Î±-thalassaemia and myelodysplastic syndrome (ATMDS). <i>British Journal of Haematology</i> , 2009, 144, 538-545.	1.2	9
54	X-linked Î± thalassaemia/mental retardation syndrome: a case with gonadal dysgenesis, caused by a novel mutation in ATRX gene. <i>Clinical Dysmorphology</i> , 2009, 18, 168-171.	0.1	10

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55	Mutations in the chromatin-associated protein ATRX. <i>Human Mutation</i> , 2008, 29, 796-802.	1.1	155
56	Neuronal Death Resulting from Targeted Disruption of the Snf2 Protein ATRX Is Mediated by p53. <i>Journal of Neuroscience</i> , 2008, 28, 12570-12580.	1.7	61
57	Epigenetics and its Genetic Syndromes. , 2008, , 155-174.		0
58	Analysis of DNA Methylation at the Human Alpha Globin Cluster during Hematopoiesis.. <i>Blood</i> , 2008, 112, 1861-1861.	0.6	0
59	Structural consequences of disease-causing mutations in the ATRX-DNMT3-DNMT3L (ADD) domain of the chromatin-associated protein ATRX. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 11939-11944.	3.3	138
60	Defining the Cause of Skewed X-Chromosome Inactivation in X-Linked Mental Retardation by Use of a Mouse Model. <i>American Journal of Human Genetics</i> , 2007, 80, 1138-1149.	2.6	32
61	Tissue-specific histone modification and transcription factor binding in $\hat{\pm}$ globin gene expression. <i>Blood</i> , 2007, 110, 4503-4510.	0.6	69
62	Using Genomics to Study How Chromatin Influences Gene Expression. <i>Annual Review of Genomics and Human Genetics</i> , 2007, 8, 299-325.	2.5	33
63	A Regulatory SNP Causes a Human Genetic Disease by Creating a New Transcriptional Promoter. <i>Science</i> , 2006, 312, 1215-1217.	6.0	254
64	Alpha thalassaemia-mental retardation, X linked. <i>Orphanet Journal of Rare Diseases</i> , 2006, 1, 15.	1.2	130
65	A novel mutation in the last exon of ATRX in a patient with alpha-thalassemia myelodysplastic syndrome. <i>European Journal of Haematology</i> , 2006, 76, 432-435.	1.1	10
66	5-Azacytidine treatment of the patient with ATMDS. <i>European Journal of Haematology</i> , 2006, 76, 453-453.	1.1	0
67	Gastrointestinal phenotype of ATR-X syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 1172-1176.	0.7	24
68	Loss of Atrx Affects Trophoblast Development and the Pattern of X-Inactivation in Extraembryonic Tissues. <i>PLoS Genetics</i> , 2006, 2, e58.	1.5	140
69	Two Novel Somatic Mutations of the ATRX Gene in Female Patients with Acquired Alpha Thalassemia of Myelodysplastic Syndrome (ATMDS).. <i>Blood</i> , 2006, 108, 1765-1765.	0.6	2
70	Acquired $\hat{\pm}$ -thalassemia in association with myelodysplastic syndrome and other hematologic malignancies. <i>Blood</i> , 2005, 105, 443-452.	0.6	95
71	Understanding $\hat{\pm}$ -Globin Gene Regulation: Aiming to Improve the Management of Thalassemia. <i>Annals of the New York Academy of Sciences</i> , 2005, 1054, 92-102.	1.8	47
72	Somatic point mutations in RUNX1/CBFA2/AML1 are common in high-risk myelodysplastic syndrome, but not in myelofibrosis with myeloid metaplasia. <i>European Journal of Haematology</i> , 2005, 74, 47-53.	1.1	62

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73	Asplenia in ATR-X syndrome: A second report. American Journal of Medical Genetics, Part A, 2005, 139A, 37-39.	0.7	15
74	Histone modifying and chromatin remodelling enzymes in cancer and dysplastic syndromes. Human Molecular Genetics, 2005, 14, R85-R92.	1.4	141
75	The chromatin-remodeling protein ATRX is critical for neuronal survival during corticogenesis. Journal of Clinical Investigation, 2005, 115, 258-267.	3.9	169
76	The chromatin-remodeling protein ATRX is critical for neuronal survival during corticogenesis. Journal of Clinical Investigation, 2005, 115, 258-267.	3.9	119
77	Attenuation of an amino-terminal premature stop codon mutation in the ATRX gene by an alternative mode of translational initiation. Journal of Medical Genetics, 2004, 41, 951-956.	1.5	39
78	Acquired somatic ATRX mutations in myelodysplastic syndrome associated with $\hat{\alpha}$ thalassemia (ATMDS) convey a more severe hematologic phenotype than germline ATRX mutations. Blood, 2004, 103, 2019-2026.	0.6	84
79	A conserved truncated isoform of the ATR-X syndrome protein lacking the SWI/SNF-homology domain. Gene, 2004, 326, 23-34.	1.0	53
80	Deletion of the $\hat{\alpha}$ -globin gene cluster as a cause of acquired $\hat{\alpha}$ -thalassemia in myelodysplastic syndrome. Blood, 2004, 103, 1518-1520.	0.6	34
81	A Novel Splicing Mutation in the Gene Encoding the Chromatin-Associated Factor ATRX Associated with Acquired Hemoglobin H Disease in Myelodysplastic Syndrome (ATMDS).. Blood, 2004, 104, 3606-3606.	0.6	3
82	Identification of acquired somatic mutations in the gene encoding chromatin-remodeling factor ATRX in the $\hat{\alpha}$ -thalassemia myelodysplasia syndrome (ATMDS). Nature Genetics, 2003, 34, 446-449.	9.4	132
83	The ATRX syndrome protein forms a chromatin-remodeling complex with Daxx and localizes in promyelocytic leukemia nuclear bodies. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 10635-10640.	3.3	322
84	102. Cloning and characterisation of tammar ATRX and ATRY. Reproduction, Fertility and Development, 2003, 15, 102.	0.1	0
85	Mutations in the general transcription factor TFIIF result in beta-thalassaemia in individuals with trichothiodystrophy. Human Molecular Genetics, 2001, 10, 2797-2802.	1.4	115
86	Monosomy for the most telomeric, gene-rich region of the short arm of human chromosome 16 causes minimal phenotypic effects. European Journal of Human Genetics, 2001, 9, 217-225.	1.4	47
87	Molecular-clinical spectrum of the ATR-X syndrome. American Journal of Medical Genetics Part A, 2000, 97, 204-212.	2.4	208
88	A nonsense mutation of the ATRX gene causing mild mental retardation and epilepsy. Annals of Neurology, 2000, 47, 117-121.	2.8	72
89	Mutations in ATRX, encoding a SWI/SNF-like protein, cause diverse changes in the pattern of DNA methylation. Nature Genetics, 2000, 24, 368-371.	9.4	476
90	Localization of a putative transcriptional regulator (ATRX) at pericentromeric heterochromatin and the short arms of acrocentric chromosomes. Proceedings of the National Academy of Sciences of the United States of America, 1999, 96, 13983-13988.	3.3	233

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91	Germline and gonosomal mosaicism in the ATR-X syndrome. <i>European Journal of Human Genetics</i> , 1999, 7, 933-936.	1.4	26
92	Brachydactyly Type B: Linkage to Chromosome 9q22 and Evidence for Genetic Heterogeneity. <i>American Journal of Human Genetics</i> , 1999, 64, 578-585.	2.6	24
93	Comparison of the human and murine ATRX gene identifies highly conserved, functionally important domains. <i>Mammalian Genome</i> , 1998, 9, 400-403.	1.0	64
94	Mutations in transcriptional regulator ATRX establish the functional significance of a PHD-like domain. <i>Nature Genetics</i> , 1997, 17, 146-148.	9.4	196
95	The genetic basis for mental retardation. <i>QJM - Monthly Journal of the Association of Physicians</i> , 1996, 89, 169-176.	0.2	36
96	The $\hat{\alpha}$ -Thalassemia/Mental Retardation Syndromes. <i>Medicine (United States)</i> , 1996, 75, 45-52.	0.4	24
97	ATRX encodes a novel member of the SNF2 family of proteins: mutations point to a common mechanism underlying the ATR-X syndrome. <i>Human Molecular Genetics</i> , 1996, 5, 1899-1907.	1.4	217
98	Lumping Juberg-Marsidi syndrome and X-linked $\hat{\alpha}$ -thalassemia/mental retardation syndrome?. <i>American Journal of Medical Genetics Part A</i> , 1995, 55, 300-301.	2.4	19
99	X-linked $\hat{\alpha}$ -thalassemia/mental retardation (ATR-X) syndrome: A new kindred with severe genital anomalies and mild hematologic expression. <i>American Journal of Medical Genetics Part A</i> , 1995, 55, 302-306.	2.4	50
100	Mutations in a putative global transcriptional regulator cause X-linked mental retardation with $\hat{\alpha}$ -thalassemia (ATR-X syndrome). <i>Cell</i> , 1995, 80, 837-845.	13.5	583
101	Alpha thalassaemia mental retardation (ATR-X): an atypical family.. <i>Archives of Disease in Childhood</i> , 1994, 70, 439-440.	1.0	13
102	Absence of the XIST gene from late-replicating isodicentric X chromosomes in leukaemia. <i>Human Molecular Genetics</i> , 1994, 3, 1053-1059.	1.4	72
103	A newly defined X linked mental retardation syndrome associated with alpha thalassaemia.. <i>Journal of Medical Genetics</i> , 1991, 28, 729-733.	1.5	62
104	X linked alpha thalassaemia/mental retardation: spectrum of clinical features in three related males.. <i>Journal of Medical Genetics</i> , 1991, 28, 738-741.	1.5	31
105	The non-deletion type of alpha thalassaemia/mental retardation: a recognisable dysmorphic syndrome with X linked inheritance.. <i>Journal of Medical Genetics</i> , 1991, 28, 724-724.	1.5	19
106	The non-deletion alpha thalassaemia/mental retardation syndrome: further support for X linkage.. <i>Journal of Medical Genetics</i> , 1991, 28, 742-745.	1.5	31
107	Unusual Types of $\hat{\alpha}$ Thalassemia. , 0, , 296-320.		2
108	ATR-X Syndrome. , 0, , 309-334.		6