Brian Hendrich

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50 8,495 32 65 g-index

65 9,355 14 5.71 ext. papers ext. citations avg, IF L-index

| # | Paper | IF | Citations |
|----|--|-------|-----------|
| 50 | A mouse Mecp2-null mutation causes neurological symptoms that mimic Rett syndrome. <i>Nature Genetics</i> , 2001 , 27, 322-6 | 36.3 | 1223 |
| 49 | Identification and characterization of a family of mammalian methyl-CpG binding proteins. <i>Molecular and Cellular Biology</i> , 1998 , 18, 6538-47 | 4.8 | 1079 |
| 48 | Dynamic reprogramming of DNA methylation in the early mouse embryo. <i>Developmental Biology</i> , 2002 , 241, 172-82 | 3.1 | 984 |
| 47 | MBD2 is a transcriptional repressor belonging to the MeCP1 histone deacetylase complex. <i>Nature Genetics</i> , 1999 , 23, 58-61 | 36.3 | 692 |
| 46 | The thymine glycosylase MBD4 can bind to the product of deamination at methylated CpG sites. <i>Nature</i> , 1999 , 401, 301-4 | 50.4 | 539 |
| 45 | 3D structures of individual mammalian genomes studied by single-cell Hi-C. <i>Nature</i> , 2017 , 544, 59-64 | 50.4 | 485 |
| 44 | The p120 catenin partner Kaiso is a DNA methylation-dependent transcriptional repressor. <i>Genes and Development</i> , 2001 , 15, 1613-8 | 12.6 | 359 |
| 43 | The methyl-CpG binding domain and the evolving role of DNA methylation in animals. <i>Trends in Genetics</i> , 2003 , 19, 269-77 | 8.5 | 323 |
| 42 | The NuRD component Mbd3 is required for pluripotency of embryonic stem cells. <i>Nature Cell Biology</i> , 2006 , 8, 285-92 | 23.4 | 299 |
| 41 | Enhanced CpG mutability and tumorigenesis in MBD4-deficient mice. <i>Science</i> , 2002 , 297, 403-5 | 33.3 | 266 |
| 40 | NuRD-mediated deacetylation of H3K27 facilitates recruitment of Polycomb Repressive Complex 2 to direct gene repression. <i>EMBO Journal</i> , 2012 , 31, 593-605 | 13 | 185 |
| 39 | NuRD suppresses pluripotency gene expression to promote transcriptional heterogeneity and lineage commitment. <i>Cell Stem Cell</i> , 2012 , 10, 583-94 | 18 | 168 |
| 38 | Deficiency of Mbd2 suppresses intestinal tumorigenesis. <i>Nature Genetics</i> , 2003 , 34, 145-7 | 36.3 | 150 |
| 37 | Kaiso-deficient mice show resistance to intestinal cancer. <i>Molecular and Cellular Biology</i> , 2006 , 26, 199- | 20,88 | 136 |
| 36 | Mbd3, a component of the NuRD co-repressor complex, is required for development of pluripotent cells. <i>Development (Cambridge)</i> , 2007 , 134, 1123-32 | 6.6 | 132 |
| 35 | MBD3/NuRD facilitates induction of pluripotency in a context-dependent manner. <i>Cell Stem Cell</i> , 2014 , 15, 102-10 | 18 | 125 |
| 34 | Somatic frameshift mutations in the MBD4 gene of sporadic colon cancers with mismatch repair deficiency. <i>Oncogene</i> , 1999 , 18, 8044-7 | 9.2 | 121 |

| 33 | Vestiges of a DNA methylation system in Drosophila melanogaster?. <i>Nature Genetics</i> , 1999 , 23, 389-90 | 36.3 | 111 |
|----|---|------|-----|
| 32 | Keeping things quiet: roles of NuRD and Sin3 co-repressor complexes during mammalian development. <i>International Journal of Biochemistry and Cell Biology</i> , 2009 , 41, 108-16 | 5.6 | 100 |
| 31 | c-Jun N-terminal phosphorylation antagonises recruitment of the Mbd3/NuRD repressor complex. <i>Nature</i> , 2011 , 469, 231-5 | 50.4 | 97 |
| 30 | Genomic structure and chromosomal mapping of the murine and human Mbd1, Mbd2, Mbd3, and Mbd4 genes. <i>Mammalian Genome</i> , 1999 , 10, 906-12 | 3.2 | 96 |
| 29 | Transcriptional repressors: multifaceted regulators of gene expression. <i>Development (Cambridge)</i> , 2013 , 140, 505-12 | 6.6 | 91 |
| 28 | The Nucleosome Remodeling and Deacetylation Complex Modulates Chromatin Structure at Sites of Active Transcription to Fine-Tune Gene Expression. <i>Molecular Cell</i> , 2018 , 71, 56-72.e4 | 17.6 | 70 |
| 27 | CHD4 in the DNA-damage response and cell cycle progression: not so NuRDy now. <i>Biochemical Society Transactions</i> , 2013 , 41, 777-82 | 5.1 | 67 |
| 26 | Mbd2 contributes to DNA methylation-directed repression of the Xist gene. <i>Molecular and Cellular Biology</i> , 2007 , 27, 3750-7 | 4.8 | 54 |
| 25 | The opposing transcriptional functions of Sin3a and c-Myc are required to maintain tissue homeostasis. <i>Nature Cell Biology</i> , 2011 , 13, 1395-405 | 23.4 | 49 |
| 24 | The methyl-CpG binding proteins Mecp2, Mbd2 and Kaiso are dispensable for mouse embryogenesis, but play a redundant function in neural differentiation. <i>PLoS ONE</i> , 2009 , 4, e4315 | 3.7 | 48 |
| 23 | Sin3a is essential for the genome integrity and viability of pluripotent cells. <i>Developmental Biology</i> , 2012 , 363, 62-73 | 3.1 | 40 |
| 22 | Methyl-CpG binding proteins and cancer: are MeCpGs more important than MBDs?. <i>Oncogene</i> , 2002 , 21, 5394-9 | 9.2 | 40 |
| 21 | Sall4 controls differentiation of pluripotent cells independently of the Nucleosome Remodelling and Deacetylation (NuRD) complex. <i>Development (Cambridge)</i> , 2016 , 143, 3074-84 | 6.6 | 40 |
| 20 | Constraint of gene expression by the chromatin remodelling protein CHD4 facilitates lineage specification. <i>Development (Cambridge)</i> , 2015 , 142, 2586-97 | 6.6 | 33 |
| 19 | The methyl binding domain 3/nucleosome remodelling and deacetylase complex regulates neural cell fate determination and terminal differentiation in the cerebral cortex. <i>Neural Development</i> , 2015 , 10, 13 | 3.9 | 32 |
| 18 | MeCP2 in neurons: closing in on the causes of Rett syndrome. <i>Human Molecular Genetics</i> , 2005 , 14 Spec No 1, R19-26 | 5.6 | 32 |
| 17 | A high-resolution map of transcriptional repression. <i>ELife</i> , 2017 , 6, | 8.9 | 31 |
| 16 | MeCP2 dependent heterochromatin reorganization during neural differentiation of a novel Mecp2-deficient embryonic stem cell reporter line. <i>PLoS ONE</i> , 2012 , 7, e47848 | 3.7 | 28 |

| 15 | The function of chromatin modifiers in lineage commitment and cell fate specification. <i>FEBS Journal</i> , 2015 , 282, 1692-702 | 5.7 | 27 |
|----|--|------|----|
| 14 | FRET-enhanced photostability allows improved single-molecule tracking of proteins and protein complexes in live mammalian cells. <i>Nature Communications</i> , 2018 , 9, 2520 | 17.4 | 23 |
| 13 | PWWP2A binds distinct chromatin moieties and interacts with an MTA1-specific core NuRD complex. <i>Nature Communications</i> , 2018 , 9, 4300 | 17.4 | 22 |
| 12 | NuRD-dependent DNA methylation prevents ES cells from accessing a trophectoderm fate. <i>Biology Open</i> , 2012 , 1, 341-52 | 2.2 | 18 |
| 11 | Methylation moves into medicine. <i>Current Biology</i> , 2000 , 10, R60-3 | 6.3 | 18 |
| 10 | The Nucleosome Remodelling and Deacetylation complex suppresses transcriptional noise during lineage commitment. <i>EMBO Journal</i> , 2019 , 38, | 13 | 16 |
| 9 | Mbd3/NuRD controls lymphoid cell fate and inhibits tumorigenesis by repressing a B cell transcriptional program. <i>Journal of Experimental Medicine</i> , 2017 , 214, 3085-3104 | 16.6 | 12 |
| 8 | Combining fluorescence imaging with Hi-C to study 3D genome architecture of the same single cell. <i>Nature Protocols</i> , 2018 , 13, 1034-1061 | 18.8 | 9 |
| 7 | Identification and characterization of a family of mammalian methyl CpG-binding proteins. <i>Genetical Research</i> , 1998 , 72, 59-72 | 1.1 | 7 |
| 6 | Live-cell 3D single-molecule tracking reveals how NuRD modulates enhancer dynamics | | 4 |
| 5 | Chromatin Remodelling Proteins and Cell Fate Decisions in Mammalian Preimplantation Development. <i>Advances in Anatomy, Embryology and Cell Biology</i> , 2018 , 229, 3-14 | 1.2 | 4 |
| 4 | Differential regulation of lineage commitment in human and mouse primed pluripotent stem cells by the nucleosome remodelling and deacetylation complex. <i>Stem Cell Research</i> , 2020 , 46, 101867 | 1.6 | 3 |
| 3 | Transcriptional control by Sall4 in blastocysts facilitates lineage commitment of inner cell mass cells | | 3 |
| 2 | The Nucleosome Remodelling and Deacetylation complex restricts Mediator access to enhancers to control transcription | | 1 |
| 1 | Subunit redundancy within the NuRD complex ensures fidelity of ES cell lineage commitment | | 1 |