Andreas Houben

List of Publications by Year
in descending order

241 papers	11,989 citations	28242 55 h-index	39638 94 g-index
261 all docs	261 docs citations	261 times ranked	8747 citing authors

#	Article	IF	Citations
1	A chromosome conformation capture ordered sequence of the barley genome. Nature, 2017, 544, 427-433.	13.7	1,365
2	DNA methylation controls histone H3 lysine 9 methylation and heterochromatin assembly in Arabidopsis. EMBO Journal, 2002, 21, 6549-6559.	3.5	439
3	Chromosomal histone modification patterns â€“ from conservation to diversity. Trends in Plant Science, 2006, 11, 199-208.	4.3	338
4	Loss of centromeric histone H3 (CENH3) from centromeres precedes uniparental chromosome elimination in interspecific barley hybrids. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, E498-505.	3.3	260
5	B chromosomes in plants: escapees from the A chromosome genome?. Trends in Plant Science, 2003, 8, 417-423.	4.3	204
6	Stable barley chromosomes without centromeric repeats. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 9842-9847.	3.3	199
7	Methylation of histone H3 in euchromatin of plant chromosomes depends on basic nuclear DNA content. Plant Journal, 2003, 33, 967-973.	2.8	186
8	Uniparental Chromosome Elimination at Mitosis and Interphase in Wheat and Pearl Millet Crosses Involves Micronucleus Formation, Progressive Heterochromatinization, and DNA Fragmentation. Plant Cell, 2005, 17, 2431-2438.	3.1	185
9	Selfish supernumerary chromosome reveals its origin as a mosaic of host genome and organellar sequences. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 13343-13346.	3.3	173
10	Genetics of sex determination in flowering plants. Genesis, 1994, 15, 214-230.	3.3	142
11	Chromosome-scale genome assembly provides insights into rye biology, evolution and agronomic potential. Nature Genetics, 2021, 53, 564-573.	9.4	138
12	Evolution and biology of supernumerary B chromosomes. Cellular and Molecular Life Sciences, 2014, 71, 467-478.	2.4	136
13	Identification and Dynamics of Two Classes of Aurora-Like Kinases in Arabidopsis and Other Plants. Plant Cell, 2005, 17, 836-848.	3.1	135
14	The transcript elongation factor FACT affects Arabidopsis vegetative and reproductive development and genetically interacts with HUB1/2. Plant Journal, 2010, 61, 686-697.	2.8	134
15	Construction of a map-based reference genome sequence for barley, Hordeum vulgare L.. Scientific Data, 2017, 4, 170044.	2.4	130
16	A Century of B Chromosomes in Plants: So What?. Annals of Botany, 2008, 101, 767-775.	1.4	126
17	Point mutation impairs centromeric CENH3 loading and induces haploid plants. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 11211-11216.	3.3	126
18	A wholeâ€genome snapshot of 454 sequences exposes the composition of the barley genome and provides evidence for parallel evolution of genome size in wheat and barley. Plant Journal, 2009, 59, 712-722.	2.8	125

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19	The chromosomal distribution of phosphorylated histone H3 differs between plants and animals at meiosis. Chromosoma, 2000, 109, 308-317.	1.0	119
20	The cell cycle dependent phosphorylation of histone H3 is correlated with the condensation of plant mitotic chromosomes. Plant Journal, 1999, 18, 675-679.	2.8	116
21	Liveâ€cell <scp>CRISPR</scp> imaging in plants reveals dynamic telomere movements. Plant Journal, 2017, 91, 565-573.	2.8	114
22	Plant Elongator regulates auxin-related genes during RNA polymerase II transcription elongation. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 1678-1683.	3.3	112
23	Phosphorylation of histone H3 in plantsâ€"A dynamic affair. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2007, 1769, 308-315.	2.4	110
24	CENH3 interacts with the centromeric retrotransposon cereba and GC-rich satellites and locates to centromeric substructures in barley. Chromosoma, 2007, 116, 275-283.	1.0	107
25	CRISPRâ€"Cas9-mediated induction of heritable chromosomal translocations in Arabidopsis. Nature Plants, 2020, 6, 638-645.	4.7	104
26	DNA and proteins of plant centromeres. Current Opinion in Plant Biology, 2003, 6, 554-560.	3.5	99
27	<i>Arabidopsis</i>Î± Aurora Kinases Function in Formative Cell Division Plane Orientation. Plant Cell, 2011, 23, 4013-4024.	3.1	97
28	Holocentromeres in <i>Rhynchospora</i> are associated with genome-wide centromere-specific repeat arrays interspersed among euchromatin. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 13633-13638.	3.3	96
29	Haploidization via Chromosome Elimination: Means and Mechanisms. Annual Review of Plant Biology, 2016, 67, 421-438.	8.6	95
30	Extrachromosomal circular DNA derived from tandemly repeated genomic sequences in plants. Plant Journal, 2008, 53, 1027-1034.	2.8	92
31	State-of-the-art and novel developments of in vivo haploid technologies. Theoretical and Applied Genetics, 2019, 132, 593-605.	1.8	91
32	Highly effective cell synchronization in plant roots by hydroxyurea and amiprophos-methyl or colchicine. Genome, 1993, 36, 387-390.	0.9	89
33	Isolation and characterization of X chromosome-derived DNA sequences from a dioecious plant Melandrium album. Chromosome Research, 1997, 5, 57-6.	1.0	89
34	Uniparental loss of ribosomal DNA in the allotetraploid grassZingeria trichopoda(2n= 8). Genome, 2003, 46, 156-163.	0.9	87
35	Arabidopsis Chromatin-Associated HMGA and HMGB Use Different Nuclear Targeting Signals and Display Highly Dynamic Localization within the Nucleus. Plant Cell, 2006, 18, 2904-2918.	3.1	86
36	The holocentric species <i><scp>L</scp>uzula elegans</i> shows interplay between centromere and largeâ€scale genome organization. Plant Journal, 2013, 73, 555-565.	2.8	86

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37	Molecular cytogenetic characterisation of the terminal heterochromatic segment of the B-chromosome of rye (Secale cereale). Chromosoma, 1996, 105, 97-103.	1.0	84
38	European maize genomes highlight intraspecies variation in repeat and gene content. Nature Genetics, 2020, 52, 950-957.	9.4	84
39	The chromatin remodelling complex FACT associates with actively transcribed regions of the Arabidopsis genome. Plant Journal, 2004, 40, 660-671.	2.8	82
40	Changing local recombination patterns in Arabidopsis by CRISPR/Cas mediated chromosome engineering. Nature Communications, 2020, 11, 4418.	5.8	82
41	The temporal and spatial pattern of histone H3 phosphorylation at serine 28 and serine 10 is similar in plants but differs between mono- and polycentric chromosomes. Cytogenetic and Genome Research, 2003, 101, 172-176.	0.6	81
42	Refined examination of plant metaphase chromosome structure at different levels made feasible by new isolation methods. Chromosoma, 1993, 102, 96-101.	1.0	79
43	Nondisjunction in Favor of a Chromosome: The Mechanism of Rye B Chromosome Drive during Pollen Mitosis. Plant Cell, 2012, 24, 4124-4134.	3.1	77
44	Alternative meiotic chromatid segregation in the holocentric plant Luzula elegans. Nature Communications, 2014, 5, 4979.	5.8	77
45	Transcriptionally Active Heterochromatin in Rye B Chromosomes. Plant Cell, 2007, 19, 1738-1749.	3.1	75
46	Highâ€copy sequences reveal distinct evolution of the rye B chromosome. New Phytologist, 2013, 199, 550-558.	3.5	75
47	Formation and Expression of Pseudogenes on the B Chromosome of Rye. Plant Cell, 2013, 25, 2536-2544.	3.1	74
48	Microdissection and microcloning of the barley (Hordeum vulgare L.) chromosome 1HS. Theoretical and Applied Genetics, 1993, 86, 629-636.	1.8	73
49	Chromosome ?painting? in plants ? a feasible technique?. Chromosoma, 1996, 104, 315-320.	1.0	72
50	Molecular Evidence for Transcription of Genes on a B Chromosome in Crepis capillaris. Genetics, 2005, 171, 269-278.	1.2	72
51	A Set of Cytogenetic Markers Allows the Precise Identification of All A-Genome Chromosomes in Diploid and Polyploid Wheat. Cytogenetic and Genome Research, 2015, 146, 71-79.	0.6	69
52	Genes on B chromosomes: Old questions revisited with new tools. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2015, 1849, 64-70.	0.9	68
53	B Chromosomes â€“ A Matter of Chromosome Drive. Frontiers in Plant Science, 2017, 08, 210.	1.7	68
54	Holocentric Chromosomes of <i>Luzula elegans</i> Are Characterized by a Longitudinal Centromere Groove, Chromosome Bending, and a Terminal Nucleolus Organizer Region. Cytogenetic and Genome Research, 2011, 134, 220-228.	0.6	65

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55	Colchicine-induced polyploidization depends on tubulin polymerization in c-metaphase cells. Protoplasma, 2006, 227, 147-153.	1.0	61
56	Centromeric and non-centromeric satellite DNA organisation differs in holocentric Rhynchospora species. Chromosoma, 2017, 126, 325-335.	1.0	59
57	Centromere location in <i>Arabidopsis</i> is unaltered by extreme divergence in CENH3 protein sequence. Genome Research, 2017, 27, 471-478.	2.4	58
58	Chromosome ?painting? in plants ? a feasible technique?. Chromosoma, 1996, 104, 315-320.	1.0	58
59	Elimination of chromosomes in <i>Hordeum vulgare</i> Ã— <i>H. bulbosum</i> crosses at mitosis and interphase involves micronucleus formation and progressive heterochromatinization. Cytogenetic and Genome Research, 2006, 114, 169-174.	0.6	56
60	The Expression Level of the Chromatin-Associated HMGB1 Protein Influences Growth, Stress Tolerance, and Transcriptome in Arabidopsis. Journal of Molecular Biology, 2008, 384, 9-21.	2.0	56
61	The plant-specific family of DNA-binding proteins containing three HMG-box domains interacts with mitotic and meiotic chromosomes. New Phytologist, 2011, 192, 577-589.	3.5	55
62	The transcript elongation factor SPT4/SPT5 is involved in auxin-related gene expression in <i>Arabidopsis</i>. Nucleic Acids Research, 2014, 42, 4332-4347.	6.5	54
63	Rye B chromosomes are weakly transcribed and might alter the transcriptional activity of A chromosome sequences. Chromosoma, 2009, 118, 607-616.	1.0	52
64	Novel phosphorylation of histone H3 at threonine 11 that temporally correlates with condensation of mitotic and meiotic chromosomes in plant cells. Cytogenetic and Genome Research, 2005, 109, 148-155.	0.6	51
65	Chromosomes Carrying Meiotic Avoidance Loci in Three Apomictic Eudicot <i>Hieracium</i> Subgenus <i>Pilosella</i> Species Share Structural Features with Two Monocot Apomicts Â Â. Plant Physiology, 2011, 157, 1327-1341.	2.3	51
66	Rye B chromosomes encode a functional Argonauteâ€like protein with <i>inÂvitro</i> slicer activities similar to its A chromosome paralog. New Phytologist, 2017, 213, 916-928.	3.5	51
67	Polymerase chain reaction mediated localization of RFLP clones to microisolated translocation chromosomes of barley. Genome, 1994, 37, 550-555.	0.9	50
68	Synteny between Brachypodium distachyon and Hordeum vulgare as revealed by FISH. Chromosome Research, 2010, 18, 841-850.	1.0	50
69	Anti-Phosphorylated Histone H2AThr120: A Universal Microscopic Marker for Centromeric Chromatin of Mono- and Holocentric Plant Species. Cytogenetic and Genome Research, 2014, 143, 150-156.	0.6	50
70	B-chromosome origin in the endemic New Zealand frog<i>Leiopelma</i><i>hochstetteri</i>through sex chromosome devolution. Genome, 1998, 41, 14-22.	0.9	48
71	Additive inheritance of histone modifications in <i>Arabidopsis thaliana</i> intraâ€specific hybrids. Plant Journal, 2011, 67, 691-700.	2.8	48
72	The ultrastructure of mono- and holocentric plant centromeres: an immunological investigation by structured illumination microscopy and scanning electron microscopy. Chromosoma, 2015, 124, 503-517.	1.0	48

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73	Utility of DNA amplified by degenerate oligonucleotide-primed PCR (DOP-PCR) from the total genome and defined chromosomal regions of field bean. Molecular Genetics and Genomics, 1994, 243, 173-177.	2.4	46
74	Molecular-cytogenetic analysis of Aegilops triuncialis and identification of its chromosomes in the background of wheat. Molecular Cytogenetics, 2014, 7, 91.	0.4	46
75	Sequencing of Single Pollen Nuclei Reveals Meiotic Recombination Events at Megabase Resolution and Circumvents Segregation Distortion Caused by Postmeiotic Processes. Frontiers in Plant Science, 2017, 8, 1620.	1.7	46
76	B-chromosome origin in the endemic New Zealand frog <i>Leiopelma</i> <i>hochstetteri</i> through sex chromosome devolution. Genome, 1998, 41, 14-22.	0.9	45
77	Aurora1 phosphorylation activity on histone H3 and its crossâ€talk with other postâ€translational histone modifications in Arabidopsis. Plant Journal, 2009, 59, 221-230.	2.8	44
78	Induction of telomereâ€mediated chromosomal truncation and stability of truncated chromosomes in <i>Arabidopsis thaliana</i>. Plant Journal, 2011, 68, 28-39.	2.8	44
79	CENH3 distribution and differential chromatin modifications during pollen development in rye (Secale) Tj ETQq1 1 0.784314 rgBT /Overl	1.0	44
80	Holokinetic centromeres and efficient telomere healing enable rapid karyotype evolution. Chromosoma, 2015, 124, 519-528.	1.0	44
81	The differential loading of two barley CENH3 variants into distinct centromeric substructures is cell type- and development-specific. Chromosome Research, 2015, 23, 277-284.	1.0	44
82	Analysis of transposable elements and organellar <scp>DNA</scp> in male and female genomes of a species with a huge Y chromosome reveals distinct Y centromeres. Plant Journal, 2016, 88, 387-396.	2.8	44
83	Differential immunostaining of plant chromosomes by antibodies recognizing acetylated histone H4 variants. Chromosome Research, 1996, 4, 191-194.	1.0	42
84	Alterations in the distribution of histone H3 phosphorylation in mitotic plant chromosomes in response to cold treatment and the protein phosphatase inhibitor cantharidin. Chromosome Research, 2002, 10, 467-476.	1.0	42
85	Super-Resolution Microscopy Reveals Diversity of Plant Centromere Architecture. International Journal of Molecular Sciences, 2020, 21, 3488.	1.8	42
86	The evolution of the hexaploid grass Zingeria kochii (Mez) Tzvel. (2n=12) was accompanied by complex hybridization and uniparental loss of ribosomal DNA. Molecular Phylogenetics and Evolution, 2010, 56, 146-155.	1.2	41
87	Telomere-mediated truncation of barley chromosomes. Chromosoma, 2012, 121, 181-190.	1.0	41
88	Histone H4 acetylation in plant heterochromatin is altered during the cell cycle. Chromosoma, 1997, 106, 193-197.	1.0	40
89	The genomic complexity of micro B chromosomes of Brachycome dichromosomatica. Chromosoma, 2001, 110, 451-459.	1.0	40
90	Distribution patterns of phosphorylated Thr 3 and Thr 32 of histone H3 in plant mitosis and meiosis. Cytogenetic and Genome Research, 2008, 122, 73-79.	0.6	39

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91	Ribosomal RNA genes specific to the B chromosomes in <i>Brachycome dichromosomatica</i> are not transcribed in leaf tissue. Genome, 1997, 40, 674-681.	0.9	38
92	A monophyletic origin of the B chromosomes ofBrachycome dichromosomatica (Asteraceae). Plant Systematics and Evolution, 1999, 219, 127-135.	0.3	38
93	Intraspecific hybrids of Arabidopsis thaliana revealed no gross alterations in endopolyploidy, DNA methylation, histone modifications and transcript levels. Theoretical and Applied Genetics, 2010, 120, 215-226.	1.8	38
94	Sources of Stem Rust Resistance in Wheat-Alien Introgression Lines. Plant Disease, 2016, 100, 1101-1109.	0.7	38
95	B Chromosomes of Aegilops speltoides Are Enriched in Organelle Genome-Derived Sequences. PLoS ONE, 2014, 9, e90214.	1.1	38
96	Mapping nonrecombining regions in barley using multicolor FISH. Chromosome Research, 2013, 21, 739-751.	1.0	37
97	The H3 histone chaperone NASP^{SIM3} escorts CenH3 in Arabidopsis. Plant Journal, 2020, 101, 71-86.	2.8	37
98	Mitotic Spindle Attachment to the Holocentric Chromosomes of Cuscuta europaea Does Not Correlate With the Distribution of CENH3 Chromatin. Frontiers in Plant Science, 2020, 10, 1799.	1.7	37
99	Differences of histone H4 acetylation and replication timing between A and B chromosomes of Brachycome dichromosomatica. Chromosome Research, 1997, 5, 233-237.	1.0	36
100	Immunogold labeling of chromosomes for scanning electron microscopy: a closer look at phosphorylated histone H3 in mitotic metaphase chromosomes of Hordeum vulgare. Chromosome Research, 2003, 11, 585-596.	1.0	36
101	How Next-Generation Sequencing Has Aided Our Understanding of the Sequence Composition and Origin of B Chromosomes. Genes, 2017, 8, 294.	1.0	36
102	A repetitive DNA sequence common to the different B chromosomes of the genus. Chromosoma, 1997, 106, 513.	1.0	36
103	The pericentromeric heterochromatin of the grass Zingeria biebersteiniana (2n = 4) is composed of Zbcen1-type tandem repeats that are intermingled with accumulated dispersedly organized sequences. Genome, 2001, 44, 955-961.	0.9	35
104	Characterization of Eu- and Heterochromatin of <i>Citrus</i> with a Focus on the Condensation Behavior of 45S rDNA Chromatin. Cytogenetic and Genome Research, 2011, 134, 72-82.	0.6	35
105	Measuring Meiotic Crossovers via Multi-Locus Genotyping of Single Pollen Grains in Barley. PLoS ONE, 2015, 10, e0137677.	1.1	34
106	Barley doubled-haploid production by uniparental chromosome elimination. Plant Cell, Tissue and Organ Culture, 2011, 104, 321-327.	1.2	33
107	Mitotic lifecycle of chromosomal 3x<scp>HMG</scp>â€box proteins and the role of their Nâ€terminal domain in the association with r<scp>DNA</scp> loci and proteolysis. New Phytologist, 2015, 208, 1067-1077.	3.5	33
108	Altered expression of Aurora kinases in Arabidopsis results in aneuâ€•and polyploidization. Plant Journal, 2014, 80, 449-461.	2.8	32

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109	Cytogenetic mapping with centromeric bacterial artificial chromosomes contigs shows that this recombinationâ€poor region comprises more than half of barley chromosome 3<scp>H</scp>. Plant Journal, 2015, 84, 385-394.	2.8	32
110	Restructuring of Holocentric Centromeres During Meiosis in the Plant <i>Rhynchospora pubera</i>. Genetics, 2016, 204, 555-568.	1.2	32
111	<scp>RNA</scp>â€guided endonuclease â€“ <i>inÂsitu</i> labelling (<scp>RGEN</scp>â€<scp>ISL</scp>): a fast <scp>CRISPR</scp>/Cas9â€based method to label genomic sequences in various species. New Phytologist, 2019, 222, 1652-1661.	3.5	32
112	A repetitive DNA sequence common to the different B chromosomes of the genus Brachycome. Chromosoma, 1997, 106, 513-519.	1.0	31
113	Engineered Plant Minichromosomes: A Bottom-Up Success?. Plant Cell, 2008, 20, 8-10.	3.1	31
114	Epigenetic Histone Marks of Extended Meta-Polycentric Centromeres of Lathyrus and Pisum Chromosomes. Frontiers in Plant Science, 2016, 7, 234.	1.7	31
115	Chromatin Ring Formation at Plant Centromeres. Frontiers in Plant Science, 2016, 7, 28.	1.7	30
116	Supernumerary B chromosomes of Aegilops speltoides undergo precise elimination in roots early in embryo development. Nature Communications, 2020, 11, 2764.	5.8	30
117	Immunostaining and interphase arrangement of field bean kinetochores. Chromosome Research, 1995, 3, 27-31.	1.0	29
118	Biology and Evolution of B Chromosomes. , 2013, , 149-165.		29
119	Localization of vicilin genes via polymerase chain reaction on microisolated field bean chromosomes. Plant Journal, 1993, 3, 883-886.	2.8	28
120	AtHaspin phosphorylates histone H3 at threonine 3 during mitosis and contributes to embryonic patterning in Arabidopsis. Plant Journal, 2011, 68, 443-454.	2.8	28
121	Arabidopsis NSE4 Proteins Act in Somatic Nuclei and Meiosis to Ensure Plant Viability and Fertility. Frontiers in Plant Science, 2019, 10, 774.	1.7	28
122	Quantum dots--a versatile tool in plant science?. Journal of Nanobiotechnology, 2006, 4, 5.	4.2	27
123	Engineered Plant Minichromosomes: A Resurrection of B Chromosomes?. Plant Cell, 2007, 19, 2323-2327.	3.1	27
124	Characterization of Centromeric Histone H3 (CENH3) Variants in Cultivated and Wild Carrots (Daucus) Tj ETQq0 0 0 rgBT /Overlock 10 T	1.1	27
125	Nondisjunction and unequal spindle organization accompany the drive of <i>Aegilops speltoides</i> B chromosomes. New Phytologist, 2019, 223, 1340-1352.	3.5	26
126	The molecular organisation of a B chromosome tandem repeat sequence fromBrachycome dichromosomatica. Chromosoma, 1996, 105, 223-230.	1.0	25

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127	Tissue culture triggers chromosome alterations, amplification, and transposition of repeat sequences in Allium fistulosum. Genome, 2007, 50, 435-442.	0.9	25
128	A Fast Air-dry Dropping Chromosome Preparation Method Suitable for FISH in Plants. Journal of Visualized Experiments, 2015, , e53470.	0.2	25
129	Evolution of Plant B Chromosome Enriched Sequences. Genes, 2018, 9, 515.	1.0	25
130	Application and prospects of CRISPR/Cas9-based methods to trace defined genomic sequences in living and fixed plant cells. Chromosome Research, 2020, 28, 7-17.	1.0	25
131	Cloning and characterisation of polymorphic heterochromatic segments of Brachycome dichromosomatica. Chromosoma, 2000, 109, 206-213.	1.0	24
132	CRISPR/Cas9-Based RGEN-ISL Allows the Simultaneous and Specific Visualization of Proteins, DNA Repeats, and Sites of DNA Replication. Cytogenetic and Genome Research, 2019, 159, 48-53.	0.6	24
133	Prospects and limitations of expansion microscopy in chromatin ultrastructure determination. Chromosome Research, 2020, 28, 355-368.	1.0	24
134	Prospects of telomereâ€toâ€telomere assembly in barley: Analysis of sequence gaps in the MorexV3 reference genome. Plant Biotechnology Journal, 2022, 20, 1373-1386.	4.1	24
135	The acetylation patterns of histones H3 and H4 along Vicia faba chromosomes are different. Chromosome Research, 1998, 6, 59-63.	1.0	23
136	Kmasker - A Tool for in silico Prediction of Single-Copy FISH Probes for the Large-Genome SpeciesHordeum vulgare. Cytogenetic and Genome Research, 2014, 142, 66-78.	0.6	23
137	B chromosomes of B. dichromosomatica show a reduced level of euchromatic histone H3 methylation marks. Chromosome Research, 2007, 15, 215-222.	1.0	22
138	Current SEM techniques for deâ€•and reâ€construction of centromeres to determine 3D CENH3 distribution in barley mitotic chromosomes. Journal of Microscopy, 2012, 246, 96-106.	0.8	22
139	B chromosomes of rye are highly conserved and accompanied the development of early agriculture. Annals of Botany, 2013, 112, 527-534.	1.4	22
140	Conserved molecular structure of the centromeric histone CENH3 in Secale and its phylogenetic relationships. Scientific Reports, 2017, 7, 17628.	1.6	22
141	Ultrastructure and Dynamics of Synaptonemal Complex Components During Meiotic Pairing and Synapsis of Standard (A) and Accessory (B) Rye Chromosomes. Frontiers in Plant Science, 2019, 10, 773.	1.7	22
142	Evolution and function of B chromosome 45S rDNA sequences in Brachycome dichromosomatica. Genome, 2007, 50, 638-644.	0.9	21
143	Molecular cytogenetic characterisation of the terminal heterochromatic segment of the B-chromosome of rye (Secale cereale). Chromosoma, 1996, 105, 97-103.	1.0	21
144	Molecular-cytogenetic characterization of a higher plant centromere/kinetochore complex. Theoretical and Applied Genetics, 1996, 93, 477-484.	1.8	20

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145	<i>Plantago lagopus</i>B Chromosome Is Enriched in 5S rDNA-Derived Satellite DNA. Cytogenetic and Genome Research, 2016, 148, 68-73.	0.6	20
146	Evolution, Composition and Regulation of Supernumerary B Chromosomes. Genes, 2019, 10, 161.	1.0	20
147	Unequal contribution of two paralogous CENH3 variants in cowpea centromere function. Communications Biology, 2020, 3, 775.	2.0	20
148	Karyotype analysis and physical mapping of 18S-5.8S-25S and 5S ribosomal RNA loci in species of genus Lens Miller (Fabaceae). Caryologia, 2002, 55, 121-128.	0.2	19
149	The distribution of epigenetic histone marks differs between the X and Y chromosomes in Silene latifolia. Planta, 2019, 250, 487-494.	1.6	19
150	The B chromosomes in <i>Brachycome</i>. Cytogenetic and Genome Research, 2004, 106, 199-209.	0.6	18
151	Distribution of Eu- and Heterochromatin in <i>Plantago</i><i>ovata</i>. Cytogenetic and Genome Research, 2009, 125, 235-240.	0.6	18
152	De novo generation of plant centromeres at tandem repeats. Chromosoma, 2013, 122, 233-241.	1.0	18
153	Cytomixis doesnâ€™t induce obvious changes in chromatin modifications and programmed cell death in tobacco male meiocytes. Frontiers in Plant Science, 2015, 6, 846.	1.7	18
154	Two combinatorial patterns of telomere histone marks in plants with canonical and nonâ€canonical telomere repeats. Plant Journal, 2020, 102, 678-687.	2.8	18
155	Aneuploids as a key for new molecular cloning strategies: development of DNA markers by microdissection using Triticum aestivum-Aegilops markgrafii chromosome addition line B. Euphytica, 1996, 89, 41-47.	0.6	17
156	Chromatin Alterations during Pollen Development inHordeum vulgare. Cytogenetic and Genome Research, 2013, 141, 50-57.	0.6	17
157	Collinearity of homoeologous group 3 chromosomes in the genus Hordeum and Secale cereale as revealed by 3H-derived FISH analysis. Chromosome Research, 2016, 24, 231-242.	1.0	17
158	Together But Different: The Subgenomes of the Bimodal Eleutherine Karyotypes Are Differentially Organized. Frontiers in Plant Science, 2019, 10, 1170.	1.7	17
159	Application of Aptamers Improves CRISPR-Based Live Imaging of Plant Telomeres. Frontiers in Plant Science, 2020, 11, 1254.	1.7	17
160	Functional Divergence of Microtubule-Associated TPX2 Family Members in Arabidopsis thaliana. International Journal of Molecular Sciences, 2020, 21, 2183.	1.8	17
161	Centromere-specific acetylation of histone H4 in barley detected through three-dimensional microscopy. Plant Molecular Biology, 2003, 51, 533-541.	2.0	16
162	Analysis of hybrid lethality in F1 wheat-rye hybrid embryos. Euphytica, 2008, 159, 367-375.	0.6	16

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163	In Vitro Phosphorylation of Histone H3 at Threonine 3 by Arabidopsis Haspin Is Strongly Influenced by Posttranslational Modifications of Adjacent Amino Acids. Molecular Plant, 2013, 6, 574-576.	3.9	16
164	TPX2 Protein of Arabidopsis Activates Aurora Kinase 1, But Not Aurora Kinase 3 In Vitro. Plant Molecular Biology Reporter, 2015, 33, 1988-1995.	1.0	16
165	The distribution of Î±-kleisin during meiosis in the holocentromeric plant Luzula elegans. Chromosome Research, 2016, 24, 393-405.	1.0	16
166	Application of Tris-HCl Allows the Specific Labeling of Regularly Prepared Chromosomes by CRISPR-FISH. Cytogenetic and Genome Research, 2020, 160, 156-165.	0.6	16
167	Characterization of the constriction with neocentric activity of 5RL chromosome in wheat. Chromosoma, 2002, 111, 228-235.	1.0	15
168	Characterisation of the low-chromosome number grass <i>Colpodium versicolor</i> (Stev.) Schmalh. (2n = 4) by molecular cytogenetics. Caryologia, 2005, 58, 241-245.	0.2	15
169	<i>In Situ</i> Hybridization to Plant Tissues and Chromosomes. , 2006, 326, 203-218.		15
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171	Fine mapping and chromosome walking towards the Ror1 locus in barley (Hordeum vulgare L.). Theoretical and Applied Genetics, 2013, 126, 2969-2982.	1.8	15
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