Fengtang Yang

List of Publications by Year
in descending order

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142 papers	19,261 citations	38742 50 h-index	12597 132 g-index
153 all docs	153 docs citations	153 times ranked	29184 citing authors

#	Article	IF	Citations
1	Integrated genome and transcriptome analyses reveal the mechanism of genome instability in ataxia with oculomotor apraxia 2. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	16
2	Evidence for multi-copy Mega-NUMTsin the human genome. Nucleic Acids Research, 2021, 49, 1517-1531.	14.5	42
3	Analysis of multiple chromosomal rearrangements in the genome of Willisornis vidua using BAC-FISH and chromosome painting on a supposed conserved karyotype. Bmc Ecology and Evolution, 2021, 21, 34.	1.6	7
4	Werner Helicase Is a Synthetic-Lethal Vulnerability in Mismatch Repairâ€“Deficient Colorectal Cancer Refractory to Targeted Therapies, Chemotherapy, and Immunotherapy. Cancer Discovery, 2021, 11, 1923-1937.	9.4	48
5	Optogenetic modeling of human neuromuscular circuits in Duchenne muscular dystrophy with CRISPR and pharmacological corrections. Science Advances, 2021, 7, eabi8787.	10.3	14
6	Chromosomal painting of the sandpiper (Actitis macularius) detects several fissions for the Scolopacidae family (Charadriiformes). Bmc Ecology and Evolution, 2021, 21, 8.	1.6	8
7	Combinatorial CRISPR screen identifies fitness effects of gene paralogues. Nature Communications, 2021, 12, 1302.	12.8	59
8	Analysis pipelines for cancer genome sequencing in mice. Nature Protocols, 2020, 15, 266-315.	12.0	25
9	Population Structure, Stratification, and Introgression of Human Structural Variation. Cell, 2020, 182, 189-199.e15.	28.9	79
10	Structural variation of the malaria-associated human glycophorin A-B-E region. BMC Genomics, 2020, 21, 446.	2.8	7
11	Expanded potential stem cell media as a tool to study human developmental hematopoiesis in vitro. Experimental Hematology, 2019, 76, 1-12.e5.	0.4	9
12	Birth, expansion, and death of VCY-containing palindromes on the human Y chromosome. Genome Biology, 2019, 20, 207.	8.8	8
13	Establishment of porcine and human expanded potential stem cells. Nature Cell Biology, 2019, 21, 687-699.	10.3	261
14	Derivation and maintenance of mouse haploid embryonic stem cells. Nature Protocols, 2019, 14, 1991-2014.	12.0	12
15	Evolutionary and functional analysis of RBMY1 gene copy number variation on the human Y chromosome. Human Molecular Genetics, 2019, 28, 2785-2798.	2.9	9
16	Functional linkage of gene fusions to cancer cell fitness assessed by pharmacological and CRISPR-Cas9 screening. Nature Communications, 2019, 10, 2198.	12.8	92
17	PiggyBac transposon tools for recessive screening identify B-cell lymphoma drivers in mice. Nature Communications, 2019, 10, 1415.	12.8	37
18	Structural rearrangements generate cell-specific, gene-independent CRISPR-Cas9 loss of fitness effects. Genome Biology, 2019, 20, 27.	8.8	35

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19	A new patientâ€derived iPSC model for dystroglycanopathies validates a compound that increases glycosylation of Î±â€dystroglycan. EMBO Reports, 2019, 20, e47967.	4.5	18
20	Chromosome-level genome assembly for giant panda provides novel insights into Carnivora chromosome evolution. Genome Biology, 2019, 20, 267.	8.8	31
21	ATM orchestrates the DNA-damage response to counter toxic non-homologous end-joining at broken replication forks. Nature Communications, 2019, 10, 87.	12.8	133
22	Human blood vessel organoids as aÂmodel ofÂdiabetic vasculopathy. Nature, 2019, 565, 505-510.	27.8	500
23	The Origins and Vulnerabilities of Two Transmissible Cancers in Tasmanian Devils. Cancer Cell, 2018, 33, 607-619.e15.	16.8	88
24	Evolutionary routes and KRAS dosage define pancreatic cancer phenotypes. Nature, 2018, 554, 62-68.	27.8	328
25	Repeat associated mechanisms of genome evolution and function revealed by the Mus caroli and Mus pahari genomes. Genome Research, 2018, 28, 448-459.	5.5	99
26	Copy number variation arising from gene conversion on the human Y chromosome. Human Genetics, 2018, 137, 73-83.	3.8	9
27	Sixteen diverse laboratory mouse reference genomes define strain-specific haplotypes and novel functional loci. Nature Genetics, 2018, 50, 1574-1583.	21.4	169
28	The Malaria-Protective Human Glycophorin Structural Variant DUP4 Shows Somatic Mosaicism and Association with Hemoglobin Levels. American Journal of Human Genetics, 2018, 103, 769-776.	6.2	21
29	Chromosome assembly of large and complex genomes using multiple references. Genome Research, 2018, 28, 1720-1732.	5.5	94
30	Multicolor Fluorescence In Situ Hybridization (FISH) Approaches for Simultaneous Analysis of the Entire Human Genome. Current Protocols in Human Genetics, 2018, 99, e70.	3.5	4
31	Shieldin complex promotes DNA end-joining and counters homologous recombination in BRCA1-null cells. Nature Cell Biology, 2018, 20, 954-965.	10.3	291
32	Chromosome painting in Glyphorynchus spirurus (Vieillot, 1819) detects a new fission in Passeriformes. PLoS ONE, 2018, 13, e0202040.	2.5	6
33	Recurrent Rearrangements of Human Amylase Genes Create Multiple Independent CNV Series. Human Mutation, 2017, 38, 532-539.	2.5	29
34	Establishment of mouse expanded potential stem cells. Nature, 2017, 550, 393-397.	27.8	223
35	Revealing hidden complexities of genomic rearrangements generated with Cas9. Scientific Reports, 2017, 7, 12867.	3.3	45
36	A reversible haploid mouse embryonic stem cell biobank resource for functional genomics. Nature, 2017, 550, 114-118.	27.8	58

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37	Reply to Gatesy and Springer: Claims of homology errors and zombie lineages do not compromise the dating of placental diversification. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E9433-E9434.	7.1	37
38	Karyotype Evolution and Phylogenetic Relationships of <i>Cricetulus sokolovi</i> Orlov et Malygin 1988 (Cricetidae, Rodentia) Inferred from Chromosomal Painting and Molecular Data. Cytogenetic and Genome Research, 2017, 152, 65-72.	1.1	10
39	Genomic evidence reveals a radiation of placental mammals uninterrupted by the KPg boundary. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E7282-E7290.	7.1	119
40	Animal Probes and ZOO-FISH. Springer Protocols, 2017, , 395-415.	0.3	3
41	Generation and Characterisation of a Pax8-CreERT2 Transgenic Line and a Slc22a6-CreERT2 Knock-In Line for Inducible and Specific Genetic Manipulation of Renal Tubular Epithelial Cells. PLoS ONE, 2016, 11, e0148055.	2.5	11
42	Genomic Alteration in Head and Neck Squamous Cell Carcinoma (HNSCC) Cell Lines Inferred from Karyotyping, Molecular Cytogenetics, and Array Comparative Genomic Hybridization. PLoS ONE, 2016, 11, e0160901.	2.5	15
43	Rapid Karyotype Evolution in Lasiopodomys Involved at Least Two Autosome â€“ Sex Chromosome Translocations. PLoS ONE, 2016, 11, e0167653.	2.5	19
44	Multiplexed pancreatic genome engineering and cancer induction by transfection-based CRISPR/Cas9 delivery in mice. Nature Communications, 2016, 7, 10770.	12.8	145
45	Formin Is Associated with Left-Right Asymmetry in the Pond Snail and the Frog. Current Biology, 2016, 26, 654-660.	3.9	135
46	Punctuated bursts in human male demography inferred from 1,244 worldwide Y-chromosome sequences. Nature Genetics, 2016, 48, 593-599.	21.4	273
47	Chromosome engineering in zygotes with <scp>CRISPR</scp>/<scp>C</scp>as9. Genesis, 2016, 54, 78-85.	1.6	78
48	Chromosomal phylogeny of Vampyressine bats (Chiroptera, Phyllostomidae) with description of two new sex chromosome systems. BMC Evolutionary Biology, 2016, 16, 119.	3.2	20
49	Karyotypic Evolution in Malagasy Flying Foxes (Pteropodidae, Chiroptera) and Their Hipposiderid Relatives as Determined by Comparative Chromosome Painting. Cytogenetic and Genome Research, 2016, 148, 185-198.	1.1	4
50	The pig X and Y Chromosomes: structure, sequence, and evolution. Genome Research, 2016, 26, 130-139.	5.5	69
51	Genome-wide comparative chromosome maps of Arvicola amphibius, Dicrostonyx torquatus, and Myodes rutilus. Chromosome Research, 2016, 24, 145-159.	2.2	9
52	Multidirectional chromosome painting substantiates the occurrence of extensive genomic reshuffling within Accipitriformes. BMC Evolutionary Biology, 2015, 15, 205.	3.2	19
53	A First Generation Comparative Chromosome Map between Guinea Pig (Cavia porcellus) and Humans. PLoS ONE, 2015, 10, e0127937.	2.5	14
54	Integration of molecular cytogenetics, dated molecular phylogeny, and model-based predictions to understand the extreme chromosome reorganization in the Neotropical genus Tonatia (Chiroptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5		20

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55	Frequent somatic transfer of mitochondrial DNA into the nuclear genome of human cancer cells. Genome Research, 2015, 25, 814-824.	5.5	69
56	CRISPR/Cas9 somatic multiplex-mutagenesis for high-throughput functional cancer genomics in mice. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 13982-13987.	7.1	172
57	Obesity, starch digestion and amylase: association between copy number variants at human salivary (AMY1) and pancreatic (AMY2) amylase genes. Human Molecular Genetics, 2015, 24, 3472-3480.	2.9	105
58	Expansion of the HSFY gene family in pig lineages. BMC Genomics, 2015, 16, 442.	2.8	10
59	Successful Generation of Human Induced Pluripotent Stem Cell Lines from Blood Samples Held at Room Temperature for up to 48Âhr. Stem Cell Reports, 2015, 5, 660-671.	4.8	51
60	Evolution of the rapidly mutating human salivary agglutinin gene (DMBT1) and population subsistence strategy. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 5105-5110.	7.1	35
61	Phylogenetic Reconstruction by Cross-Species Chromosome Painting and G-Banding in Four Species of Phyllostomini Tribe (Chiroptera, Phyllostomidae) in the Brazilian Amazon: An Independent Evidence for Monophyly. PLoS ONE, 2015, 10, e0122845.	2.5	15
62	Quantitative Genetics of CTCF Binding Reveal Local Sequence Effects and Different Modes of X-Chromosome Association. PLoS Genetics, 2014, 10, e1004798.	3.5	55
63	Sequence of a Complete Chicken BG Haplotype Shows Dynamic Expansion and Contraction of Two Gene Lineages with Particular Expression Patterns. PLoS Genetics, 2014, 10, e1004417.	3.5	31
64	Haptoglobin (HP) and Haptoglobin-related protein (HPR) copy number variation, natural selection, and trypanosomiasis. Human Genetics, 2014, 133, 69-83.	3.8	72
65	Transmissible Dog Cancer Genome Reveals the Origin and History of an Ancient Cell Lineage. Science, 2014, 343, 437-440.	12.6	144
66	Two new cytotypes reinforce that Micronycteris hirsuta Peters, 1869 does not represent a monotypic taxon. BMC Genetics, 2013, 14, 119.	2.7	14
67	Genetic Basis of Y-Linked Hearing Impairment. American Journal of Human Genetics, 2013, 92, 301-306.	6.2	25
68	The zebrafish reference genome sequence and its relationship to the human genome. Nature, 2013, 496, 498-503.	27.8	3,708
69	Massively Parallel Sequencing Reveals the Complex Structure of an Irradiated Human Chromosome on a Mouse Background in the Tc1 Model of Down Syndrome. PLoS ONE, 2013, 8, e60482.	2.5	93
70	Fine Mapping of the Pond Snail Left-Right Asymmetry (Chirality) Locus Using RAD-Seq and Fibre-FISH. PLoS ONE, 2013, 8, e71067.	2.5	26
71	Disruption of Mouse Cenpj, a Regulator of Centriole Biogenesis, Phenocopies Seckel Syndrome. PLoS Genetics, 2012, 8, e1003022.	3.5	84
72	A comprehensive molecular cytogenetic analysis of chromosome rearrangements in gibbons. Genome Research, 2012, 22, 2520-2528.	5.5	32

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73	Isolation of homozygous mutant mouse embryonic stem cells using a dual selection system. Nucleic Acids Research, 2012, 40, e21-e21.	14.5	21
74	Generation of diversity by somatic mutation in the <scp>C</scp>amelus dromedarius<scp>T</scp>â€cell receptor gamma variable domains. European Journal of Immunology, 2012, 42, 3416-3428.	2.9	27
75	Estimation of rearrangement phylogeny for cancer genomes. Genome Research, 2012, 22, 346-361.	5.5	108
76	Genome Sequencing and Analysis of the Tasmanian Devil and Its Transmissible Cancer. Cell, 2012, 148, 780-791.	28.9	300
77	Massive Genomic Rearrangement Acquired in a Single Catastrophic Event during Cancer Development. Cell, 2011, 144, 27-40.	28.9	2,020
78	ATMIN Is Required for Maintenance of Genomic Stability and Suppression of B Cell Lymphoma. Cancer Cell, 2011, 19, 587-600.	16.8	33
79	High-resolution chromosome painting reveals the first genetic signature for the chiropteran suborder Pteropodiformes (Mammalia: Chiroptera). Chromosome Research, 2011, 19, 507-519.	2.2	18
80	Chromosome painting in Tragulidae facilitates the reconstruction of Ruminantia ancestral karyotype. Chromosome Research, 2011, 19, 531-539.	2.2	25
81	Comparative cytogenetic mapping of Sox2 and Sox14 in cichlid fishes and inferences on the genomic organization of both genes in vertebrates. Chromosome Research, 2011, 19, 657-667.	2.2	14
82	Chromosomal evolution of Arvicolinae (Cricetidae, Rodentia). III. Karyotype relationships of ten Microtus species. Chromosome Research, 2010, 18, 459-471.	2.2	37
83	Cross-species chromosome painting in bats from Madagascar: the contribution of Myzopodidae to revealing ancestral syntenies in Chiroptera. Chromosome Research, 2010, 18, 635-653.	2.2	11
84	Signatures of mutation and selection in the cancer genome. Nature, 2010, 463, 893-898.	27.8	661
85	Avian comparative genomics: reciprocal chromosome painting between domestic chicken (Gallus) Tj ETQq1 1 0.784314 rgBT /Overlock diploid number. Chromosome Research, 2009, 17, 99-113.	2.2	58
86	Chromosomal rearrangements underlying karyotype differences between Chinese pangolin (Manis) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 Research, 2009, 17, 321-329.	2.2	12
87	Cross-species chromosome painting in Cetartiodactyla: Reconstructing the karyotype evolution in key phylogenetic lineages. Chromosome Research, 2009, 17, 419-436.	2.2	45
88	Animal Probes and ZOO-FISH. , 2009, , 323-346.		20
89	Tracking genome organization in rodents by Zoo-FISH. Chromosome Research, 2008, 16, 261-274.	2.2	29
90	Multidirectional cross-species painting illuminates the history of karyotypic evolution in Perissodactyla. Chromosome Research, 2008, 16, 89-107.	2.2	68

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91	Phylogenomics of the dog and fox family (Canidae, Carnivora) revealed by chromosome painting. Chromosome Research, 2008, 16, 129-143.	2.2	58
92	Comparative cytogenetics of bats (Chiroptera): The prevalence of Robertsonian translocations limits the power of chromosomal characters in resolving interfamily phylogenetic relationships. Chromosome Research, 2008, 16, 155-170.	2.2	40
93	Foreword. Chromosome Research, 2008, 16, 1-4.	2.2	11
94	Flying lemurs â€“ The 'flying tree shrews'? Molecular cytogenetic evidence for a Scandentia-Dermoptera sister clade. BMC Biology, 2008, 6, 18.	3.8	44
95	Adaptive Evolution of UGT2B17 Copy-Number Variation. American Journal of Human Genetics, 2008, 83, 337-346.	6.2	137
96	Neo-sex chromosomes in the black muntjac recapitulate incipient evolution of mammalian sex chromosomes. Genome Biology, 2008, 9, R98.	9.6	36
97	Copy number variation and evolution in humans and chimpanzees. Genome Research, 2008, 18, 1698-1710.	5.5	215
98	Construction, Characterization, and Chromosomal Mapping of a Fosmid Library of the White-Cheeked Gibbon (Nomascus leucogenys). Genomics, Proteomics and Bioinformatics, 2007, 5, 207-215.	6.9	3
99	Flow analysis and sorting of microchromosomes (<3 Mb). Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2007, 71A, 410-413.	1.5	10
100	Definition of the zebrafish genome using flow cytometry and cytogenetic mapping. BMC Genomics, 2007, 8, 195.	2.8	43
101	Paired-End Mapping Reveals Extensive Structural Variation in the Human Genome. Science, 2007, 318, 420-426.	12.6	1,003
102	Karyotypic evolution and phylogenetic relationships in the order Chiroptera as revealed by G-banding comparison and chromosome painting. Chromosome Research, 2007, 15, 257-67.	2.2	35
103	Karyotype evolution and phylogenetic relationships of hamsters (Cricetidae, Muroidea, Rodentia) inferred from chromosomal painting and banding comparison. Chromosome Research, 2007, 15, 283-97.	2.2	52
104	Chromosomal evolution of Arvicolinae (Cricetidae, Rodentia). I. The genome homology of tundra vole, field vole, mouse and golden hamster revealed by comparative chromosome painting. Chromosome Research, 2007, 15, 447-456.	2.2	49
105	Cross-species chromosome painting among camel, cattle, pig and human: further insights into the putative Cetartiodactyla ancestral karyotype. Chromosome Research, 2007, 15, 499-514.	2.2	110
106	Karyotype evolution in Rhinolophus bats (Rhinolophidae, Chiroptera) illuminated by cross-species chromosome painting and G-banding comparison. Chromosome Research, 2007, 15, 835-848.	2.2	42
107	Chromosomal evolution of Arvicolinae (Cricetidae, Rodentia). II. The genome homology of two mole voles (genus Ellobius), the field vole and golden hamster revealed by comparative chromosome painting. Chromosome Research, 2007, 15, 891-897.	2.2	57
108	High-density comparative BAC mapping in the black muntjac (Muntiacus crinifrons): Molecular cytogenetic dissection of the origin of MCR 1p+4 in the X1X2Y1Y2Y3 sex chromosome system. Genomics, 2006, 87, 608-615.	2.9	17

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109	Global variation in copy number in the human genome. Nature, 2006, 444, 444-454.	27.8	3,831
110	Phylogenomics of several deer species revealed by comparative chromosome painting with Chinese muntjac paints. Genetica, 2006, 127, 25-33.	1.1	36
111	Cross-species chromosome painting unveils cytogenetic signatures for the Eulipotyphla and evidence for the polyphyly of Insectivora. Chromosome Research, 2006, 14, 151-159.	2.2	41
112	Chromosome localization of microsatellite markers in the shrews of the Sorex araneus group. Chromosome Research, 2006, 14, 253-262.	2.2	23
113	Comparative genome maps of the pangolin, hedgehog, sloth, anteater and human revealed by cross-species chromosome painting: further insight into the ancestral karyotype and genome evolution of eutherian mammals. Chromosome Research, 2006, 14, 283-296.	2.2	58
114	Comparative genomic analysis links karyotypic evolution with genomic evolution in the Indian Muntjac (Muntiacus muntjak vaginalis). Chromosoma, 2006, 115, 427-436.	2.2	6
115	Reciprocal chromosome painting between three laboratory rodent species. Mammalian Genome, 2006, 17, 1183-1192.	2.2	35
116	Chromosome painting between human and lorisiform prosimians: Evidence for the HSA 7/16 synteny in the primate ancestral karyotype. American Journal of Physical Anthropology, 2006, 129, 250-259.	2.1	29
117	Are molecular cytogenetics and bioinformatics suggesting diverging models of ancestral mammalian genomes?. Genome Research, 2006, 16, 306-310.	5.5	73
118	Phylogenomics of the genus Mus (Rodentia; Muridae): extensive genome repatterning is not restricted to the house mouse. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 2925-2934.	2.6	58
119	Phylogenomic study of the subfamily Caprinae by cross-species chromosome painting with Chinese muntjac paints. Chromosome Research, 2005, 13, 389-399.	2.2	26
120	The proto-oncogene C-KIT maps to canid B-chromosomes. Chromosome Research, 2005, 13, 113-122.	2.2	72
121	Comparative cytogenetics of human chromosome 3q21.3 reveals a hot spot for ectopic recombination in hominoid evolution. Genomics, 2005, 85, 36-47.	2.9	19
122	Refined genome-wide comparative map of the domestic horse, donkey and human based on cross-species chromosome painting: insight into the occasional fertility of mules. Chromosome Research, 2004, 12, 65-76.	2.2	102
123	Chromosome evolution in bears: reconstructing phylogenetic relationships by cross-species chromosome painting. Chromosome Research, 2004, 12, 55-63.	2.2	33
124	Evolution of Genome Organizations of Squirrels (Sciuridae) Revealed by Cross-Species Chromosome Painting. Chromosome Research, 2004, 12, 317-335.	2.2	51
125	Karyotype of canine soft tissue sarcomas: a multi-colour, multi-species approach to canine chromosome painting. Chromosome Research, 2004, 12, 825-835.	2.2	13
126	Integrated Comparative Genome Maps and Their Implications for Karyotype Evolution of Carnivores. , 2004, , 215-224.		11

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127	Comparative chromosome painting defines the karyotypic relationships among the domestic dog, Chinese raccoon dog and Japanese raccoon dog. Chromosome Research, 2003, 11, 735-740.	2.2	38
128	Robertsonian translocation (8;14) in an infertile bitch (Canis familiaris). Journal of Applied Genetics, 2003, 44, 525-7.	1.9	3
129	The genome phylogeny of domestic cat, red panda and five mustelid species revealed by comparative chromosome painting and G-banding. Chromosome Research, 2002, 10, 209-222.	2.2	68
130	Cloning and chromosomal localization of human WIG-1/PAG608 and demonstration of amplification with increased expression in primary squamous cell carcinoma of the lung. Cancer Letters, 2001, 174, 179-187.	7.2	9
131	A classification efficiency test of spectral karyotyping and multiplex fluorescence in situ hybridization: Identification of chromosome homologies between Homo sapiens and Hylobates leucogenys. Genes Chromosomes and Cancer, 2001, 31, 65-74.	2.8	18
132	Cross-species color banding in ten cases of myeloid malignancies with complex karyotypes. Genes Chromosomes and Cancer, 2001, 30, 15-24.	2.8	18
133	Multiplex Fluorescence In Situ Hybridization and Cross Species Color Banding of a Case of Chronic Myeloid Leukemia in Blastic Crisis with a Complex Philadelphia Translocation. Cancer Genetics and Cytogenetics, 2000, 116, 105-110.	1.0	24
134	Reciprocal chromosome painting illuminates the history of genome evolution of the domestic cat, dog and human. Chromosome Research, 2000, 8, 393-404.	2.2	92
135	Use of Flow-Sorted Canine Chromosomes in the Assignment of Canine Linkage, Radiation Hybrid, and Syntenic Groups to Chromosomes: Refinement and Verification of the Comparative Chromosome Map for Dog and Human. Genomics, 2000, 69, 182-195.	2.9	56
136	Comparative chromosome painting. , 2000, , 259-265.		5
137	Mapping chromosomal homologies between humans and two langurs (Semnopithecus francoisi and S.) Tj ETQq1 1 0.784314 rgBT /Ov	2.2	40
138	Comparative Mapping Using Chromosome Sorting and Painting. ILAR Journal, 1998, 39, 68-76.	1.8	55
139	Comparative Chromosome Painting in Mammals: Human and the Indian Muntjac (Muntiacus muntjak) Tj ETQq1 1 0.784314 rgBT /Ov	2.9	90
140	Use of the Indian Muntjac Idiogram to Align Conserved Chromosomal Segments in Sheep and Human Genomes by Chromosome Painting. Genomics, 1997, 46, 143-147.	2.9	33
141	Chromosome painting without competitor DNA. Technical Tips Online, 1997, 2, 6-7.	0.2	9
142	Quenching autofluorescence in tissue immunofluorescence. Wellcome Open Research, 0, 2, 79.	1.8	14