

# Polina L Perelman

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2865511/publications.pdf>

Version: 2024-02-01

70  
papers

4,933  
citations

196777

29  
h-index

111975

67  
g-index

72  
all docs

72  
docs citations

72  
times ranked

7352  
citing authors

#	ARTICLE	IF	CITATIONS
1	Evolution of Tandemly Arranged Repetitive DNAs in Three Species of Cyprinoidei with Different Ploidy Levels. <i>Cytogenetic and Genome Research</i> , 2021, 161, 32-42.	0.6	3
2	Draft de novo Genome Assembly of the Elusive Jaguarundi, <i>Puma yagouaroundi</i> . <i>Journal of Heredity</i> , 2021, 112, 540-548.	1.0	5
3	Repetitive Sequence Distribution on <i>Saguinus</i> , <i>Leontocebus</i> and <i>Leontopithecus</i> Tamarins (Platyrrhine). <i>Tj ETQq1 1,0,784314 rgBT /O</i>	1.3	4
4	Massive LINE-1 retrotransposon enrichment in tamarins of the Cebidae family (Platyrrhini, Primates) and its significance for genome evolution. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2021, 59, 2553-2561.	0.6	3
5	Karyotype Evolution in 10 Pinniped Species: Variability of Heterochromatin versus High Conservatism of Euchromatin as Revealed by Comparative Molecular Cytogenetics. <i>Genes</i> , 2020, 11, 1485.	1.0	8
6	A draft genome assembly of spotted hyena, <i>Crocuta crocuta</i> . <i>Scientific Data</i> , 2020, 7, 126.	2.4	6
7	Evolution of the Human Chromosome 13 Synteny: Evolutionary Rearrangements, Plasticity, Human Disease Genes and Cancer Breakpoints. <i>Genes</i> , 2020, 11, 383.	1.0	10
8	Uso de la micromatriz de alta densidad de bovino para la construcción de un mapa físico de polimorfismos de nucleótido simple en alpacas ( <i>Vicugna pacos</i> ). <i>Revista De Investigaciones Veterinarias Del Peru</i> , 2020, 31, e18725.	0.0	0
9	Chromosome-Level Alpaca Reference Genome VicPac3.1 Improves Genomic Insight Into the Biology of New World Camelids. <i>Frontiers in Genetics</i> , 2019, 10, 586.	1.1	19
10	Improving Illumina assemblies with Hi-C and long reads: An example with the North African dromedary. <i>Molecular Ecology Resources</i> , 2019, 19, 1015-1026.	2.2	67
11	Evolution of gene regulation in ruminants differs between evolutionary breakpoint regions and homologous synteny blocks. <i>Genome Research</i> , 2019, 29, 576-589.	2.4	39
12	Comparative Chromosome Mapping of Musk Ox and the X Chromosome among Some Bovidae Species. <i>Genes</i> , 2019, 10, 857.	1.0	8
13	Construction of two whole genome radiation hybrid panels for dromedary ( <i>Camelus dromedarius</i> ): 5000RAD and 15000RAD. <i>Scientific Reports</i> , 2018, 8, 1982.	1.6	20
14	Multiple intrasyntenic rearrangements and rapid speciation in voles. <i>Scientific Reports</i> , 2018, 8, 14980.	1.6	11
15	A combined banding method that allows the reliable identification of chromosomes as well as differentiation of AT- and GC-rich heterochromatin. <i>Chromosome Research</i> , 2018, 26, 307-315.	1.0	19
16	Red fox genome assembly identifies genomic regions associated with tame and aggressive behaviours. <i>Nature Ecology and Evolution</i> , 2018, 2, 1479-1491.	3.4	113
17	The Case of X and Y Localization of Nucleolus Organizer Regions (NORs) in <i>Tragulus javanicus</i> (Cetartiodactyla, Mammalia). <i>Genes</i> , 2018, 9, 312.	1.0	7
18	Sequencing of Supernumerary Chromosomes of Red Fox and Raccoon Dog Confirms a Non-Random Gene Acquisition by B Chromosomes. <i>Genes</i> , 2018, 9, 405.	1.0	22

#	ARTICLE	IF	CITATIONS
19	Naked mole rat cells display more efficient excision repair than mouse cells. <i>Aging</i> , 2018, 10, 1454-1473.	1.4	38
20	Centromere repositioning explains fundamental number variability in the New World monkey genus <i>Saimiri</i> . <i>Chromosoma</i> , 2017, 126, 519-529.	1.0	12
21	Intrachromosomal Rearrangements in Rodents from the Perspective of Comparative Region-Specific Painting. <i>Genes</i> , 2017, 8, 215.	1.0	12
22	X Chromosome Evolution in Cetartiodactyla. <i>Genes</i> , 2017, 8, 216.	1.0	24
23	The Ancestral Carnivore Karyotype As Substantiated by Comparative Chromosome Painting of Three Pinnipeds, the Walrus, the Steller Sea Lion and the Baikal Seal ( <i>Pinnipedia</i> , <i>Carnivora</i> ). <i>PLoS ONE</i> , 2016, 11, e0147647.	1.1	15
24	Rapid Karyotype Evolution in <i>Lasiopodomys</i> Involved at Least Two Autosome " Sex Chromosome Translocations. <i>PLoS ONE</i> , 2016, 11, e0167653.	1.1	19
25	Comparative Chromosome Map and Heterochromatin Features of the Gray Whale Karyotype ( <i>Cetacea</i> ). <i>Cytogenetic and Genome Research</i> , 2016, 148, 25-34.	0.6	13
26	A First Generation Comparative Chromosome Map between Guinea Pig ( <i>Cavia porcellus</i> ) and Humans. <i>PLoS ONE</i> , 2015, 10, e0127937.	1.1	14
27	Genomic legacy of the African cheetah, <i>Acinonyx jubatus</i> . <i>Genome Biology</i> , 2015, 16, 277.	3.8	167
28	Platinum coat color in red fox ( <i>Vulpes vulpes</i> ) is caused by a mutation in an autosomal copy of <i>KIT</i> . <i>Animal Genetics</i> , 2015, 46, 190-199.	0.6	13
29	Whole-genome analyses resolve early branches in the tree of life of modern birds. <i>Science</i> , 2014, 346, 1320-1331.	6.0	1,583
30	A Comprehensive Whole-Genome Integrated Cytogenetic Map for the Alpaca ( <i>Lama</i> ). <i>Chromosome Research</i> , 2014, 22, 107-119.	0.8	59
31	Development and Application of Camelid Molecular Cytogenetic Tools. <i>Journal of Heredity</i> , 2014, 105, 952-963.	1.0	36
32	A New Case of an Inherited Reciprocal Translocation in Cattle: rcp(13;26)(q24;q11). <i>Cytogenetic and Genome Research</i> , 2014, 144, 208-211.	0.6	2
33	Comparative chromosome painting of pronghorn ( <i>Antilocapra americana</i> ) and saola ( <i>Pseudoryx</i> ). <i>Chromosome Research</i> , 2014, 22, 107-119.	0.784314	7
34	Molecular cytogenetic insights to the phylogenetic affinities of the giraffe ( <i>Giraffa camelopardalis</i> ) and pronghorn ( <i>Antilocapra americana</i> ). <i>Chromosome Research</i> , 2013, 21, 447-460.	1.0	17
35	Transcription of a protein-coding gene on B chromosomes of the Siberian roe deer ( <i>Capreolus</i> ). <i>Chromosome Research</i> , 2013, 21, 447-460.	1.7	58
36	Tissue sampling methods and standards for vertebrate genomics. <i>GigaScience</i> , 2012, 1, 8.	3.3	51

#	ARTICLE	IF	CITATIONS
37	Chromosome Painting of the Pygmy Tree Shrew Shows that No Derived Cytogenetic Traits Link Primates and Scandentia. <i>Cytogenetic and Genome Research</i> , 2012, 136, 175-179.	0.6	11
38	Comparative Chromosome Painting in Carnivora and Pholidota. <i>Cytogenetic and Genome Research</i> , 2012, 137, 174-193.	0.6	18
39	Chromosomal evolution in Rodentia. <i>Heredity</i> , 2012, 108, 4-16.	1.2	70
40	Chromosomal rearrangements and karyotype evolution in carnivores revealed by chromosome painting. <i>Heredity</i> , 2012, 108, 17-27.	1.2	45
41	A Molecular Phylogeny of Living Primates. <i>PLoS Genetics</i> , 2011, 7, e1001342.	1.5	1,130
42	Karyotype Evolution of Eulipotyphla (Insectivora): The Genome Homology of Seven Sorex Species Revealed by Comparative Chromosome Painting and Banding Data. <i>Cytogenetic and Genome Research</i> , 2011, 135, 51-64.	0.6	23
43	Comparative cytogenetics of main Laurasiatheria taxa. <i>Russian Journal of Genetics</i> , 2010, 46, 1132-1137.	0.2	1
44	Comparative cytogenetics of rodents. <i>Russian Journal of Genetics</i> , 2010, 46, 1138-1142.	0.2	4
45	Cross-species chromosome painting in Cetartiodactyla: Reconstructing the karyotype evolution in key phylogenetic lineages. <i>Chromosome Research</i> , 2009, 17, 419-436.	1.0	45
46	Tracking genome organization in rodents by Zoo-FISH. <i>Chromosome Research</i> , 2008, 16, 261-274.	1.0	29
47	Multidirectional cross-species painting illuminates the history of karyotypic evolution in Perissodactyla. <i>Chromosome Research</i> , 2008, 16, 89-107.	1.0	68
48	Phylogenomics of the dog and fox family (Canidae, Carnivora) revealed by chromosome painting. <i>Chromosome Research</i> , 2008, 16, 129-143.	1.0	58
49	Chromosome painting shows that skunks (Mephitidae, Carnivora) have highly rearranged karyotypes. <i>Chromosome Research</i> , 2008, 16, 1215-1231.	1.0	16
50	The Ancestral Carnivore Karyotype (2n = 38) Lives Today in Ringtails. <i>Journal of Heredity</i> , 2008, 99, 241-253.	1.0	16
51	Karyotype evolution and phylogenetic relationships of hamsters (Cricetidae, Muroidea, Rodentia) inferred from chromosomal painting and banding comparison. <i>Chromosome Research</i> , 2007, 15, 283-97.	1.0	52
52	Chromosomal evolution of Arvicolinae (Cricetidae, Rodentia). I. The genome homology of tundra vole, field vole, mouse and golden hamster revealed by comparative chromosome painting. <i>Chromosome Research</i> , 2007, 15, 447-456.	1.0	49
53	Chromosomal evolution of Arvicolinae (Cricetidae, Rodentia). II. The genome homology of two mole voles (genus <i>Ellobius</i> ), the field vole and golden hamster revealed by comparative chromosome painting. <i>Chromosome Research</i> , 2007, 15, 891-897.	1.0	57
54	Novel genes identified by manual annotation and microarray expression analysis in the pancreas. <i>Genomics</i> , 2006, 88, 752-761.	1.3	6

#	ARTICLE	IF	CITATIONS
55	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. <i>Chromosome Research</i> , 2006, 14, 283-296.	1.0	58
56	Reciprocal chromosome painting between three laboratory rodent species. <i>Mammalian Genome</i> , 2006, 17, 1183-1192.	1.0	35
57	The proto-oncogene C-KIT maps to canid B-chromosomes. <i>Chromosome Research</i> , 2005, 13, 113-122.	1.0	72
58	Karyotypic conservatism in the suborder Feliformia (Order Carnivora). <i>Cytogenetic and Genome Research</i> , 2005, 108, 348-354.	0.6	31
59	Comparative chromosome painting defines the karyotypic relationships among the domestic dog, Chinese raccoon dog and Japanese raccoon dog. <i>Chromosome Research</i> , 2003, 11, 735-740.	1.0	38
60	Reciprocal chromosome painting among human, aardvark, and elephant (superorder Afrotheria) reveals the likely eutherian ancestral karyotype. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 1062-1066.	3.3	164
61	Assignment of the bovine attractin (ATRN) gene to chromosome 13q21â†’q22 by in situ hybridization. <i>Cytogenetic and Genome Research</i> , 2003, 103, 204K-204K.	0.6	2
62	Comparative molecular cytogenetic studies in the order Carnivora: mapping chromosomal rearrangements onto the phylogenetic tree. <i>Cytogenetic and Genome Research</i> , 2002, 96, 137-145.	0.6	64
63	Complex structure of B-chromosomes in two mammalian species: <i>Apodemus peninsulae</i> (Rodentia) and <i>Nyctereutes procyonoides</i> (Carnivora). <i>Chromosome Research</i> , 2002, 10, 109-116.	1.0	58
64	Comparative Chromosome Painting. <i>Russian Journal of Genetics</i> , 2002, 38, 869-876.	0.2	1
65	Karyotype relationships between distantly related marsupials from South America and Australia. <i>Chromosome Research</i> , 2001, 9, 301-308.	1.0	46
66	Phylogenetic implications of the 38 putative ancestral chromosome segments for four canid species. <i>Cytogenetic and Genome Research</i> , 2001, 92, 243-247.	0.6	81
67	Dog chromosome-specific paints reveal evolutionary inter- and intrachromosomal rearrangements in the American mink and human. <i>Cytogenetic and Genome Research</i> , 2000, 90, 275-278.	0.6	51
68	Comparative chromosome analysis in three <i>Sorex</i> species: <i>S. raddei</i> , <i>S. minutus</i> and <i>S. caecutiens</i> . <i>Acta Theriologica</i> , 2000, 45, 119-130.	1.1	15
69	Comparative Cytogenetics of Some Species of <i>Crocidura</i> (Insectivora) with 2n=40.. <i>Cytologia</i> , 1999, 64, 293-299.	0.2	4
70	Localization of the genes for major ribosomal RNA on chromosomes of the house musk shrew, <i>Suncus murinus</i> , at meiotic and mitotic cells by fluorescence in situ hybridization and silver staining.. <i>Genes and Genetic Systems</i> , 1997, 72, 215-218.	0.2	9