

Mariano Rocchi

List of Publications by Year in descending order

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Version: 2024-02-01

28
papers

3,571
citations

394421

19
h-index

477307

29
g-index

31
all docs

31
docs citations

31
times ranked

4938
citing authors

#	ARTICLE	IF	CITATIONS
1	Evolutionary and Biomedical Insights from the Rhesus Macaque Genome. <i>Science</i> , 2007, 316, 222-234.	12.6	1,283
2	Comparative and demographic analysis of orang-utan genomes. <i>Nature</i> , 2011, 469, 529-533.	27.8	541
3	Gibbon genome and the fast karyotype evolution of small apes. <i>Nature</i> , 2014, 513, 195-201.	27.8	320
4	Gene amplification as double minutes or homogeneously staining regions in solid tumors: Origin and structure. <i>Genome Research</i> , 2010, 20, 1198-1206.	5.5	194
5	Heterozygous Submicroscopic Inversions Involving Olfactory Receptor Gene Clusters Mediate the Recurrent t(4;8)(p16;p23) Translocation. <i>American Journal of Human Genetics</i> , 2002, 71, 276-285.	6.2	185
6	Evolutionary Formation of New Centromeres in Macaque. <i>Science</i> , 2007, 316, 243-246.	12.6	136
7	Recurrent Sites for New Centromere Seeding. <i>Genome Research</i> , 2004, 14, 1696-1703.	5.5	135
8	Centromere Repositioning. <i>Genome Research</i> , 1999, 9, 1184-1188.	5.5	124
9	The genome of the vervet (<i>Chlorocebus aethiops sabaues</i>). <i>Genome Research</i> , 2015, 25, 1921-1933.	5.5	114
10	Evolutionary movement of centromeres in horse, donkey, and zebra. <i>Genomics</i> , 2006, 87, 777-782.	2.9	100
11	Genomic organization and evolution of double minutes/homogeneously staining regions with MYC amplification in human cancer. <i>Nucleic Acids Research</i> , 2014, 42, 9131-9145.	14.5	91
12	Independent centromere formation in a capricious, gene-free domain of chromosome 13q21 in Old World monkeys and pigs. <i>Genome Biology</i> , 2006, 7, R91.	9.6	60
13	Centromere sliding on a mammalian chromosome. <i>Chromosoma</i> , 2015, 124, 277-287.	2.2	49
14	Evolutionary history of chromosome 10 in primates. <i>Chromosoma</i> , 2002, 111, 267-272.	2.2	36
15	Evolutionary descent of a human chromosome 6 neocentromere: A jump back to 17 million years ago. <i>Genome Research</i> , 2009, 19, 778-784.	5.5	34
16	Epigenetic origin of evolutionary novel centromeres. <i>Scientific Reports</i> , 2017, 7, 41980.	3.3	30
17	Evolutionary history of chromosome 11 featuring four distinct centromere repositioning events in Catarrhini. <i>Genomics</i> , 2007, 90, 35-43.	2.9	28
18	Great ape Y Chromosome and mitochondrial DNA phylogenies reflect subspecies structure and patterns of mating and dispersal. <i>Genome Research</i> , 2016, 26, 427-439.	5.5	27

#	ARTICLE	IF	CITATIONS
19	Evolutionary History of Chromosome 20. <i>Molecular Biology and Evolution</i> , 2004, 22, 360-366.	8.9	21
20	The Hidden Genomic and Transcriptomic Plasticity of Giant Marker Chromosomes in Cancer. <i>Genetics</i> , 2018, 208, 951-961.	2.9	13
21	22q11.2 Low Copy Repeats Expanded in the Human Lineage. <i>Frontiers in Genetics</i> , 2021, 12, 706641.	2.3	11
22	A satellite-like sequence, representing a "clone gap" in the human genome, was likely involved in the seeding of a novel centromere in macaque. <i>Chromosoma</i> , 2009, 118, 269-277.	2.2	9
23	Ring chromosomes, breakpoint clusters, and neocentromeres in sarcomas. <i>Genes Chromosomes and Cancer</i> , 2015, 54, 156-167.	2.8	9
24	The 14/15 association as a paradigmatic example of tracing karyotype evolution in New World monkeys. <i>Chromosoma</i> , 2016, 125, 747-756.	2.2	8
25	Rapid emergence of independent "chromosomal lineages" in silvered-leaf monkey triggered by Y/autosome translocation. <i>Scientific Reports</i> , 2018, 8, 3250.	3.3	5
26	Fluorescence In Situ Hybridization Probe Preparation. <i>Methods in Molecular Biology</i> , 2017, 1541, 91-100.	0.9	3
27	Molecular characterization of an anaphoid supernumerary marker chromosome derived from 18q22.1qter in prenatal diagnosis: a case report. <i>Molecular Cytogenetics</i> , 2014, 7, 69.	0.9	1
28	Eight million years of maintained heterozygosity in chromosome homologs of cercopithecine monkeys. <i>Chromosoma</i> , 2020, 129, 57-67.	2.2	1