

Franca Pelliccia

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

392
citations

759233

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docs citations

22
times ranked

567
citing authors

#	ARTICLE	IF	CITATIONS
1	Chromosome length and DNA loop size during early embryonic development of <i>Xenopus laevis</i> . <i>Chromosoma</i> , 1993, 102, 478-483.	2.2	45
2	Influence of family history of dementia in the development and progression of late-onset Alzheimer's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 250-256.	1.7	45
3	Organization, nucleotide sequence, and chromosomal mapping of a tandemly repeated unit containing the four core histone genes and a 5S rRNA gene in an isopod crustacean species. <i>Genome</i> , 2000, 43, 341-345.	2.0	36
4	5S ribosomal and U1 small nuclear RNA genes: A new linkage type in the genome of a crustacean that has three different tandemly repeated units containing 5S ribosomal DNA sequences. <i>Genome</i> , 2001, 44, 331-335.	2.0	33
5	Characterization of FRA7B, a human common fragile site mapped at the 7p chromosome terminal region. <i>Cancer Genetics and Cytogenetics</i> , 2010, 202, 47-52.	1.0	31
6	5S ribosomal and U1 small nuclear RNA genes: A new linkage type in the genome of a crustacean that has three different tandemly repeated units containing 5S ribosomal DNA sequences. <i>Genome</i> , 2001, 44, 331-335.	2.0	28
7	Molecular characterization of the human common fragile site FRA1H. <i>Genes Chromosomes and Cancer</i> , 2007, 46, 487-493.	2.8	24
8	Identification and characterization of U1 small nuclear RNA genes from two crustacean isopod species. <i>Chromosome Research</i> , 2003, 11, 365-373.	2.2	23
9	Breakages at common fragile sites set boundaries of amplified regions in two leukemia cell lines K562 and K562-1. Molecular characterization of FRA2H and localization of a new CFS FRA2S. <i>Cancer Letters</i> , 2010, 299, 37-44.	7.2	21
10	Clonal inheritance of rRNA gene activity: Cytological evidence in human cells. <i>Chromosoma</i> , 1981, 84, 345-351.	2.2	19
11	Impaired Replication Timing Promotes Tissue-Specific Expression of Common Fragile Sites. <i>Genes</i> , 2020, 11, 326.	2.4	16
12	Analysis of the Association Between TERC and TERT Genetic Variation and Leukocyte Telomere Length and Human Lifespan: A Follow-Up Study. <i>Genes</i> , 2019, 10, 82.	2.4	14
13	rRNA polymerase I transcription is modulated by spatial learning in different brain regions. <i>Journal of Neurochemistry</i> , 2016, 136, 706-716.	3.9	13
14	Study of the relationships between common fragile sites, chromosome breakages and sister chromatid exchanges. <i>Mutagenesis</i> , 1995, 10, 257-260.	2.6	11
15	Sex chromosome differentiation revealed by genomic in-situ hybridization. <i>Chromosome Research</i> , 2000, 8, 459-464.	2.2	8
16	DNA methylation, histone H3 methylation, and histone H4 acetylation in the genome of a crustacean. <i>Genome</i> , 2006, 49, 87-90.	2.0	6
17	Biallelic deletion and loss of expression analysis of genes at FRA2G common fragile site in tumor-derived cell lines. <i>Cancer Genetics and Cytogenetics</i> , 2005, 161, 181-186.	1.0	5
18	Transcriptional profiling of genes at the human common fragile site FRA1H in tumor-derived cell lines. <i>Cancer Genetics and Cytogenetics</i> , 2007, 178, 144-150.	1.0	5

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19	The importance of molecular cytogenetic analysis prior to using cell lines in research: The case of the KG-1a leukemia cell line. <i>Oncology Letters</i> , 2012, 4, 237-240.	1.8	4
20	Characterization of Chromosomal Instability in Glioblastoma. <i>Frontiers in Genetics</i> , 2021, 12, 810793.	2.3	3
21	Cytogenetic mapping of five YAC clones to human chromosome region 2q31 -> q32.1 in relation to the FRA2G common fragile site. <i>Genetica</i> , 2002, 115, 269-272.	1.1	2
22	Effects of Potassium Cyanide on Silver Stainability of Specific Cell Structures. <i>Biotechnic & Histochemistry</i> , 1982, 57, 259-263.	0.4	0