Franca Pelliccia

List of Publications by Year
in descending order

22	392	759233 12	752698 20
papers	citations	h-index	g-index

22	22	22	567
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Characterization of Chromosomal Instability in Glioblastoma. Frontiers in Genetics, 2021, 12, 810793.	2.3	3
2	Impaired Replication Timing Promotes Tissue-Specific Expression of Common Fragile Sites. Genes, 2020, 11, 326.	2.4	16
3	Analysis of the Association Between TERC and TERT Genetic Variation and Leukocyte Telomere Length and Human Lifespanâ€"A Follow-Up Study. Genes, 2019, 10, 82.	2.4	14
4	Influence of family history of dementia in the development and progression of lateâ€onset Alzheimer's disease. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 250-256.	1.7	45
5	<scp>RNA</scp> polymerase I transcription is modulated by spatial learning in different brain regions. Journal of Neurochemistry, 2016, 136, 706-716.	3.9	13
6	The importance of molecular cytogenetic analysis prior to using cell lines in research: The case of the KG-1a leukemia cell line. Oncology Letters, 2012, 4, 237-240.	1.8	4
7	Characterization of FRA7B, a human common fragile site mapped at the 7p chromosome terminal region. Cancer Genetics and Cytogenetics, 2010, 202, 47-52.	1.0	31
8	Breakages at common fragile sites set boundaries of amplified regions in two leukemia cell lines K562 â€" Molecular characterization of FRA2H and localization of a new CFS FRA2S. Cancer Letters, 2010, 299, 37-44.	7.2	21
9	Molecular characterization of the human common fragile site FRA1H. Genes Chromosomes and Cancer, 2007, 46, 487-493.	2.8	24
10	Transcriptional profiling of genes at the human common fragile site FRA1H in tumor-derived cell lines. Cancer Genetics and Cytogenetics, 2007, 178, 144-150.	1.0	5
11	DNA methylation, histone H3 methylation, and histone H4 acetylation in the genome of a crustacean. Genome, 2006, 49, 87-90.	2.0	6
12	Biallelic deletion and loss of expression analysis of genes at FRA2G common fragile site in tumor-derived cell lines. Cancer Genetics and Cytogenetics, 2005, 161, 181-186.	1.0	5
13	Identification and characterization of U1 small nuclear RNA genes from two crustacean isopod species. Chromosome Research, 2003, 11, 365-373.	2.2	23
14	Cytogenetic mapping of five YAC clones to human chromosome region 2q31 --> q32.1 in relation to the FRA2G common fragile site. Genetica, 2002, 115, 269-272.	1.1	2
15	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. Genome, 2001, 44, 331-335.	2.0	33
16	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. Genome, 2001, 44, 331-335.	2.0	28
17	Sex chromosome differentiation revealed by genomic in-situ hybridization. Chromosome Research, 2000, 8, 459-464.	2.2	8
18	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. Genome, 2000, 43, 341-345.	2.0	36

#	Article	IF	Citations
19	Study of the relationships between common fragile sites, chromosome breakages and sister chromatid exchanges. Mutagenesis, 1995, 10, 257-260.	2.6	11
20	Chromosome length and DNA loop size during early embryonic development of Xenopus laevis. Chromosoma, 1993, 102, 478-483.	2.2	45
21	Effects of Potassium Cyanide on Silver Stainability of Specific Cell Structures. Biotechnic & Histochemistry, 1982, 57, 259-263.	0.4	0
22	Clonal inheritance of rRNA gene activity: Cytological evidence in human cells. Chromosoma, 1981, 84, 345-351.	2.2	19