Sandra Louzada

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8256364/publications.pdf

Version: 2024-02-01

22 papers 1,436 citations

687335 13 h-index 23 g-index

26 all docs

26 docs citations

times ranked

26

3863 citing authors

#	Article	IF	CITATIONS
1	Genomic Tackling of Human Satellite DNA: Breaking Barriers through Time. International Journal of Molecular Sciences, 2021, 22, 4707.	4.1	8
2	Optogenetic modeling of human neuromuscular circuits in Duchenne muscular dystrophy with CRISPR and pharmacological corrections. Science Advances, 2021, 7, eabi8787.	10.3	14
3	Decoding the Role of Satellite DNA in Genome Architecture and Plasticity—An Evolutionary and Clinical Affair. Genes, 2020, 11, 72.	2.4	47
4	Birth, expansion, and death of VCY-containing palindromes on the human Y chromosome. Genome Biology, 2019, 20, 207.	8.8	8
5	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
6	A new patientâ€derived iPSC model for dystroglycanopathies validates a compound that increases glycosylation of αâ€dystroglycan. EMBO Reports, 2019, 20, e47967.	4.5	18
7	Evolutionary routes and KRAS dosage define pancreatic cancer phenotypes. Nature, 2018, 554, 62-68.	27.8	328
8	Alcohol and endogenous aldehydes damage chromosomes and mutate stem cells. Nature, 2018, 553, 171-177.	27.8	284
9	Copy number variation arising from gene conversion on the human Y chromosome. Human Genetics, 2018, 137, 73-83.	3.8	9
10	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
11	Recurrent Rearrangements of Human Amylase Genes Create Multiple Independent CNV Series. Human Mutation, 2017, 38, 532-539.	2.5	29
12	Multiplexed pancreatic genome engineering and cancer induction by transfection-based CRISPR/Cas9 delivery in mice. Nature Communications, 2016, 7, 10770.	12.8	145
13	Punctuated bursts in human male demography inferred from 1,244 worldwide Y-chromosome sequences. Nature Genetics, 2016, 48, 593-599.	21.4	273
14	The pig X and Y Chromosomes: structure, sequence, and evolution. Genome Research, 2016, 26, 130-139.	5.5	69
15	A novel satellite DNA sequence in the Peromyscus genome (PMSat): Evolution via copy number fluctuation. Molecular Phylogenetics and Evolution, 2015, 92, 193-203.	2.7	10
16	A High-Resolution Comparative Chromosome Map of <i>Cricetus cricetus </i> and <i></i> Reveals the Involvement of Constitutive Heterochromatin in Breakpoint Regions. Cytogenetic and Genome Research, 2015, 145, 59-67.	1.1	11
17	Expansion of the HSFY gene family in pig lineages. BMC Genomics, 2015, 16, 442.	2.8	10
18	Evolution of the rapidly mutating human salivary agglutinin gene (<i>DMBT1</i>) and population subsistence strategy. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 5105-5110.	7.1	35

#	Article	IF	CITATION
19	Quantitative Genetics of CTCF Binding Reveal Local Sequence Effects and Different Modes of X-Chromosome Association. PLoS Genetics, 2014, 10, e1004798.	3.5	55
20	Praomys tullbergi (Muridae, Rodentia) genome architecture decoded by comparative chromosome painting with Mus and Rattus. Chromosome Research, 2012, 20, 673-683.	2.2	10
21	Defining the Sister Rat Mammary Tumor Cell Lines HH-16 cl.2/1 and HH-16.cl.4 as an In Vitro Cell Model for Erbb2. PLoS ONE, 2012, 7, e29923.	2.5	24
22	Different evolutionary trails in the related genomes Cricetus cricetus and Peromyscus eremicus (Rodentia, Cricetidae) uncovered by orthologous satellite DNA repositioning. Micron, 2008, 39, 1149-1155.	2.2	13