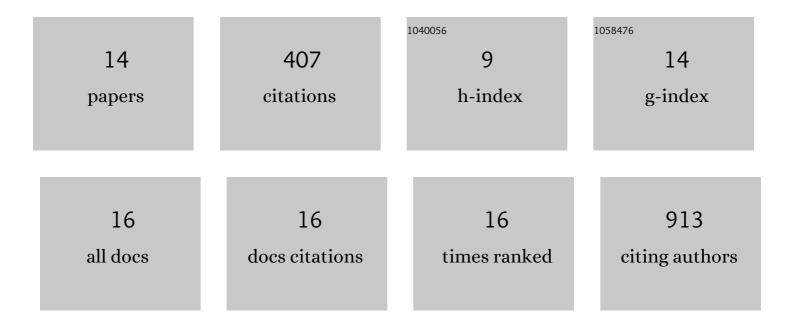
Clara Lopes Novo

List of Publications by Year in descending order

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CLARA LORES NOVO

#	Article	IF	CITATIONS
1	Satellite repeat transcripts modulate heterochromatin condensates and safeguard chromosome stability in mouse embryonic stem cells. Nature Communications, 2022, 13, .	12.8	16
2	A Tale of Two States: Pluripotency Regulation of Telomeres. Frontiers in Cell and Developmental Biology, 2021, 9, 703466.	3.7	4
3	Long-Range Enhancer Interactions Are Prevalent in Mouse Embryonic Stem Cells and Are Reorganized upon Pluripotent State Transition. Cell Reports, 2018, 22, 2615-2627.	6.4	99
4	Molecular profiling of aged neural progenitors identifies <i>Dbx2</i> as a candidate regulator of ageâ€associated neurogenic decline. Aging Cell, 2018, 17, e12745.	6.7	46
5	The pluripotency factor <i>Nanog</i> regulates pericentromeric heterochromatin organization in mouse embryonic stem cells. Genes and Development, 2016, 30, 1101-1115.	5.9	50
6	Crosstalk between pluripotency factors and higher-order chromatin organization. Nucleus, 2016, 7, 447-452.	2.2	7
7	Chromatin organization in pluripotent cells: emerging approaches to study and disrupt function. Briefings in Functional Genomics, 2016, 15, 305-314.	2.7	4
8	Recombinogenic Telomeres in Diploid Sorex granarius (Soricidae, Eulipotyphla) Fibroblast Cells. Molecular and Cellular Biology, 2014, 34, 2786-2799.	2.3	10
9	Telomeres and the nucleus. Seminars in Cancer Biology, 2013, 23, 116-124.	9.6	10
10	A new role for histone deacetylase 5 in the maintenance of long telomeres. FASEB Journal, 2013, 27, 3632-3642.	0.5	12
11	HOT1 is a mammalian direct telomere repeat-binding protein contributing to telomerase recruitment. EMBO Journal, 2013, 32, 1681-1701.	7.8	74
12	The heterochromatic chromosome caps in great apes impact telomere metabolism. Nucleic Acids Research, 2013, 41, 4792-4801.	14.5	11
13	The role of recombination in telomere length maintenance. Biochemical Society Transactions, 2009, 37, 589-595.	3.4	21
14	Telomere length maintenance – an ALTernative mechanism. Cytogenetic and Genome Research, 2008, 122, 281-291.	1.1	31