Maite Huarte

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/8978079/maite-huarte-publications-by-year.pdf

Version: 2024-04-05

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

52	14,690	30	57
papers	citations	h-index	g-index
57	17,489 ext. citations	20.4	7.27
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
52	Subcellular Distribution of p53 by the p53-Responsive lncRNA Determines Chemotherapeutic Response in Neuroblastoma. <i>Cancer Research</i> , 2021 , 81, 1457-1471	10.1	10
51	Gene regulation by long non-coding RNAs and its biological functions. <i>Nature Reviews Molecular Cell Biology</i> , 2021 , 22, 96-118	48.7	597
50	The DNA damage inducible lncRNA SCAT7 regulates genomic integrity and topoisomerase 1 turnover in lung adenocarcinoma. <i>NAR Cancer</i> , 2021 , 3, zcab002	5.2	2
49	A lncRNA-SWI/SNF complex crosstalk controls transcriptional activation at specific promoter regions. <i>Nature Communications</i> , 2020 , 11, 936	17.4	28
48	Analysis of copy number alterations reveals the lncRNA ALAL-1 as a regulator of lung cancer immune evasion. <i>Journal of Cell Biology</i> , 2020 , 219,	7.3	16
47	SNHG15 is a bifunctional MYC-regulated noncoding locus encoding a lncRNA that promotes cell proliferation, invasion and drug resistance in colorectal cancer by interacting with AIF. <i>Journal of Experimental and Clinical Cancer Research</i> , 2019 , 38, 172	12.8	45
46	hCLE/RTRAF-HSPC117-DDX1-FAM98B: A New Cap-Binding Complex That Activates mRNA Translation. <i>Frontiers in Physiology</i> , 2019 , 10, 92	4.6	4
45	LncRNA-OIS1 regulates DPP4 activation to modulate senescence induced by RAS. <i>Nucleic Acids Research</i> , 2018 , 46, 4213-4227	20.1	28
44	The aberrant splicing of BAF45d links splicing regulation and transcription in glioblastoma. <i>Neuro-Oncology</i> , 2018 , 20, 930-941	1	13
43	A lncRNA GUARDINg genome integrity. <i>Nature Cell Biology</i> , 2018 , 20, 371-372	23.4	1
42	Sense-Antisense lncRNA Pair Encoded by Locus 6p22.3 Determines Neuroblastoma Susceptibility via the USP36-CHD7-SOX9 Regulatory Axis. <i>Cancer Cell</i> , 2018 , 33, 417-434.e7	24.3	86
41	PR-LncRNA signature regulates glioma cell activity through expression of SOX factors. <i>Scientific Reports</i> , 2018 , 8, 12746	4.9	11
40	A "Counter-Enhancer" in Tumor Suppression. <i>Cell</i> , 2018 , 173, 1318-1319	56.2	3
39	Noncoding RNAs as effective markers in cancer-care management. <i>Nature Medicine</i> , 2017 , 23, 1122-112	2 3 50.5	1
38	The human lncRNA LINC-PINT inhibits tumor cell invasion through a highly conserved sequence element. <i>Genome Biology</i> , 2017 , 18, 202	18.3	122
37	The multidimensional mechanisms of long noncoding RNA function. <i>Genome Biology</i> , 2017 , 18, 206	18.3	540
36	Long Noncoding RNA PURPL Suppresses Basal p53 Levels and Promotes Tumorigenicity in Colorectal Cancer. <i>Cell Reports</i> , 2017 , 20, 2408-2423	10.6	77

(2011-2016)

35	Expanding the p53 regulatory network: LncRNAs take up the challenge. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2016 , 1859, 200-8	6	65
34	Distinct Sets of lncRNAs are Differentially Modulated after Exposure to High and Low Doses of X Rays. <i>Radiation Research</i> , 2016 , 186, 549-558	3.1	4
33	p53 partners with RNA in the DNA damage response. <i>Nature Genetics</i> , 2016 , 48, 1298-1299	36.3	11
32	RNA. A lncRNA links genomic variation with celiac disease. <i>Science</i> , 2016 , 352, 43-4	33.3	9
31	A Long Noncoding RNA Regulates Sister Chromatid Cohesion. <i>Molecular Cell</i> , 2016 , 63, 397-407	17.6	62
30	The emerging role of lncRNAs in cancer. <i>Nature Medicine</i> , 2015 , 21, 1253-61	50.5	1711
29	RNA pulldown protocol for in vitro detection and identification of RNA-associated proteins. <i>Methods in Molecular Biology</i> , 2015 , 1206, 87-95	1.4	32
28	Long noncoding RNAs: from identification to functions and mechanisms. <i>Advances in Genomics and Genetics</i> , 2015 , 257		4
27	FAL1ing inside an amplicon. Cancer Cell, 2014 , 26, 303-304	24.3	6
26	Long non-coding RNAs and chromatin modifiers: their place in the epigenetic code. <i>Epigenetics</i> , 2014 , 9, 21-6	5.7	140
25	Genome-wide analysis of the human p53 transcriptional network unveils a lncRNA tumour suppressor signature. <i>Nature Communications</i> , 2014 , 5, 5812	17.4	137
24	Long non-coding RNAs: challenges for diagnosis and therapies. <i>Nucleic Acid Therapeutics</i> , 2013 , 23, 15-2	20 4.8	137
23	LncRNAs have a say in protein translation. Cell Research, 2013, 23, 449-51	24.7	25
22	LincRNA-p21 Suppresses Target mRNA Translation. <i>Molecular Cell</i> , 2013 , 50, 303	17.6	7
21	Pint lincRNA connects the p53 pathway with epigenetic silencing by the Polycomb repressive complex 2. <i>Genome Biology</i> , 2013 , 14, R104	18.3	182
20	Long Non-Coding RNAs and Their Roles in Cancer 2013 , 245-266		
19	LincRNA-p21 suppresses target mRNA translation. <i>Molecular Cell</i> , 2012 , 47, 648-55	17.6	728
18	To repress or not to repress: this is the guardian'd question. <i>Trends in Cell Biology</i> , 2011 , 21, 344-53	18.3	46

17	Histone H4K20/H3K9 demethylase PHF8 regulates zebrafish brain and craniofacial development. <i>Nature</i> , 2010 , 466, 503-7	50.4	232
16	The Emerging Non-Coding RNA World. <i>Modecular Medicine and Medicinal</i> , 2010 , 17-49		1
15	The DMM complex prevents spreading of DNA methylation from transposons to nearby genes in Neurospora crassa. <i>Genes and Development</i> , 2010 , 24, 443-54	12.6	42
14	Large non-coding RNAs: missing links in cancer?. Human Molecular Genetics, 2010, 19, R152-61	5.6	416
13	A large intergenic noncoding RNA induced by p53 mediates global gene repression in the p53 response. <i>Cell</i> , 2010 , 142, 409-19	56.2	1648
12	Chromatin signature reveals over a thousand highly conserved large non-coding RNAs in mammals. <i>Nature</i> , 2009 , 458, 223-7	50.4	3230
11	Journal club. A biologist looks at new functions for non-coding RNAs. <i>Nature</i> , 2009 , 459, 487	50.4	1
10	Many human large intergenic noncoding RNAs associate with chromatin-modifying complexes and affect gene expression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 11667-72	11.5	2331
9	Lid2 is required for coordinating H3K4 and H3K9 methylation of heterochromatin and euchromatin. <i>Cell</i> , 2008 , 135, 272-83	56.2	108
8	The fission yeast Jmj2 reverses histone H3 Lysine 4 trimethylation. <i>Journal of Biological Chemistry</i> , 2007 , 282, 21662-70	5.4	24
7	The X-linked mental retardation gene SMCX/JARID1C defines a family of histone H3 lysine 4 demethylases. <i>Cell</i> , 2007 , 128, 1077-88	56.2	524
6	S. pombe LSD1 homologs regulate heterochromatin propagation and euchromatic gene transcription. <i>Molecular Cell</i> , 2007 , 26, 89-101	17.6	83
5	Essential dosage-dependent functions of the transcription factor yin yang 1 in late embryonic development and cell cycle progression. <i>Molecular and Cellular Biology</i> , 2006 , 26, 3565-81	4.8	149
4	Reversal of histone lysine trimethylation by the JMJD2 family of histone demethylases. <i>Cell</i> , 2006 , 125, 467-81	56.2	794
3	hCLE/CGI-99, a human protein that interacts with the influenza virus polymerase, is a mRNA transcription modulator. <i>Journal of Molecular Biology</i> , 2006 , 362, 887-900	6.5	38
2	Threonine 157 of influenza virus PA polymerase subunit modulates RNA replication in infectious viruses. <i>Journal of Virology</i> , 2003 , 77, 6007-13	6.6	53
1	PA subunit from influenza virus polymerase complex interacts with a cellular protein with homology to a family of transcriptional activators. <i>Journal of Virology</i> , 2001 , 75, 8597-604	6.6	95