

Raffaella Santoro

List of Publications by Year in descending order

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47
papers

3,730
citations

186265

28
h-index

214800

47
g-index

52
all docs

52
docs citations

52
times ranked

4973
citing authors

#	ARTICLE	IF	CITATIONS
1	The nucleolar remodeling complex NoRC mediates heterochromatin formation and silencing of ribosomal gene transcription. <i>Nature Genetics</i> , 2002, 32, 393-396.	21.4	393
2	Intergenic Transcripts Regulate the Epigenetic State of rRNA Genes. <i>Molecular Cell</i> , 2006, 22, 351-361.	9.7	302
3	The epigenetic modifier EZH2 controls melanoma growth and metastasis through silencing of distinct tumour suppressors. <i>Nature Communications</i> , 2015, 6, 6051.	12.8	281
4	Molecular Mechanisms Mediating Methylation-Dependent Silencing of Ribosomal Gene Transcription. <i>Molecular Cell</i> , 2001, 8, 719-725.	9.7	231
5	The chromatin remodeling complex NoRC targets HDAC1 to the ribosomal gene promoter and represses RNA polymerase I transcription. <i>EMBO Journal</i> , 2002, 21, 4632-4640.	7.8	212
6	The NoRC complex mediates the heterochromatin formation and stability of silent rRNA genes and centromeric repeats. <i>EMBO Journal</i> , 2010, 29, 2135-2146.	7.8	170
7	Epigenetic Mechanism of rRNA Gene Silencing: Temporal Order of NoRC-Mediated Histone Modification, Chromatin Remodeling, and DNA Methylation. <i>Molecular and Cellular Biology</i> , 2005, 25, 2539-2546.	2.3	156
8	BAZ2A (TIP5) is involved in epigenetic alterations in prostate cancer and its overexpression predicts disease recurrence. <i>Nature Genetics</i> , 2015, 47, 22-30.	21.4	141
9	Inheritance of Silent rDNA Chromatin Is Mediated by PARP1 via Noncoding RNA. <i>Molecular Cell</i> , 2012, 45, 790-800.	9.7	136
10	Epigenetic disruption of ribosomal RNA genes and nucleolar architecture in DNA methyltransferase 1 (Dnmt1) deficient cells. <i>Nucleic Acids Research</i> , 2007, 35, 2191-2198.	14.5	128
11	Inflammasome-Activated Caspase 7 Cleaves PARP1 to Enhance the Expression of a Subset of NF- κ B Target Genes. <i>Molecular Cell</i> , 2012, 46, 200-211.	9.7	128
12	lncRNA Maturation to Initiate Heterochromatin Formation in the Nucleolus Is Required for Exit from Pluripotency in ESCs. <i>Cell Stem Cell</i> , 2014, 15, 720-734.	11.1	124
13	EZH2-Mediated Primary Cilium Deconstruction Drives Metastatic Melanoma Formation. <i>Cancer Cell</i> , 2018, 34, 69-84.e14.	16.8	123
14	Intergenic transcripts originating from a subclass of ribosomal DNA repeats silence ribosomal RNA genes in <i>trans</i> . <i>EMBO Reports</i> , 2010, 11, 52-58.	4.5	106
15	The chromatin remodeling complex NoRC controls replication timing of rRNA genes. <i>EMBO Journal</i> , 2005, 24, 120-127.	7.8	98
16	Genome Organization in and around the Nucleolus. <i>Cells</i> , 2019, 8, 579.	4.1	92
17	Antagonistic Cross-Regulation between Sox9 and Sox10 Controls an Anti-tumorigenic Program in Melanoma. <i>PLoS Genetics</i> , 2015, 11, e1004877.	3.5	85
18	Formation of nuclear heterochromatin. <i>Epigenetics</i> , 2012, 7, 811-814.	2.7	78

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19	Methylation-dependent SOX9 expression mediates invasion in human melanoma cells and is a negative prognostic factor in advanced melanoma. <i>Genome Biology</i> , 2015, 16, 42.	8.8	76
20	Oxidative stress in sperm affects the epigenetic reprogramming in early embryonic development. <i>Epigenetics and Chromatin</i> , 2018, 11, 60.	3.9	70
21	Nuclear envelope alterations generate an aging-like epigenetic pattern in mice deficient in Zmpste24 metalloprotease. <i>Aging Cell</i> , 2010, 9, 947-957.	6.7	50
22	The expanding role of PARPs in the establishment and maintenance of heterochromatin. <i>FEBS Journal</i> , 2013, 280, 3508-3518.	4.7	50
23	Helicase CHD4 is an epigenetic coregulator of PAX3-FOXO1 in alveolar rhabdomyosarcoma. <i>Journal of Clinical Investigation</i> , 2016, 126, 4237-4249.	8.2	46
24	ARTD2 activity is stimulated by RNA. <i>Nucleic Acids Research</i> , 2014, 42, 5072-5082.	14.5	42
25	The RNA helicase DHX9 establishes nucleolar heterochromatin, and this activity is required for embryonic stem cell differentiation. <i>EMBO Reports</i> , 2017, 18, 1248-1262.	4.5	42
26	Nucleolus and rRNA Gene Chromatin in Early Embryo Development. <i>Trends in Genetics</i> , 2019, 35, 868-879.	6.7	38
27	Challenges in the analysis of long noncoding RNA functionality. <i>FEBS Letters</i> , 2016, 590, 2342-2353.	2.8	37
28	Regulation and Roles of the Nucleolus in Embryonic Stem Cells: From Ribosome Biogenesis to Genome Organization. <i>Stem Cell Reports</i> , 2020, 15, 1206-1219.	4.8	37
29	Pramel7 mediates ground-state pluripotency through proteasomal epigenetic combined pathways. <i>Nature Cell Biology</i> , 2017, 19, 763-773.	10.3	33
30	Genome-wide maps of nucleolus interactions reveal distinct layers of repressive chromatin domains. <i>Nature Communications</i> , 2022, 13, 1483.	12.8	32
31	Epigenetic Engineering of Ribosomal RNA Genes Enhances Protein Production. <i>PLoS ONE</i> , 2009, 4, e6653.	2.5	24
32	BAZ2A-mediated repression via H3K14ac-marked enhancers promotes prostate cancer stem cells. <i>EMBO Reports</i> , 2021, 22, e53014.	4.5	19
33	Analysis of Chromatin Composition of Repetitive Sequences: The ChIP-Chop Assay. <i>Methods in Molecular Biology</i> , 2014, 1094, 319-328.	0.9	19
34	Many players, one goal: how chromatin states are inherited during cell division. <i>Biochemistry and Cell Biology</i> , 2005, 83, 332-343.	2.0	18
35	TIP5 primes prostate luminal cells for the oncogenic transformation mediated by PTEN loss. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 3637-3647.	7.1	17
36	Identification of cis- and trans-acting elements regulating calretinin expression in mesothelioma cells. <i>Oncotarget</i> , 2016, 7, 21272-21286.	1.8	17

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37	Epigenetic control of melanoma cell invasiveness by the stem cell factor SALL4. Nature Communications, 2021, 12, 5056.	12.8	15
38	<scp>BAZ</scp> 2A safeguards genome architecture of groundâ€state pluripotent stem cells. EMBO Journal, 2020, 39, e105606.	7.8	14
39	Specific Inhibitory Effect of H1e Histone Somatic Variant on in Vitro DNA-Methylation Process. Biochemical and Biophysical Research Communications, 1996, 220, 102-107.	2.1	12
40	Noncoding RNAs link PARP1 to heterochromatin. Cell Cycle, 2012, 11, 2217-2218.	2.6	8
41	Does hypomethylation of linker DNA play a role in chromatin condensation?. Gene, 1995, 157, 247-251.	2.2	5
42	Specific variants of H1 histone regulate CpG methylation in eukaryotic DNA. Gene, 1995, 157, 253-256.	2.2	4
43	UV-Laser Induced Protein/DNA Crosslinking Reveals Sequence Variations of DNA Elements Bound by c-Jun in Vivo. Biochemical and Biophysical Research Communications, 1999, 256, 68-74.	2.1	4
44	H1â€™H1 Cross-Linking Efficiency Depends on Genomic DNA Methylation. Biochemical and Biophysical Research Communications, 1996, 227, 768-774.	2.1	1
45	The NoRC complex mediates heterochromatin formation and stability of silent rRNA genes and centromeric repeats. EMBO Journal, 2010, 29, 2253-2253.	7.8	1
46	Epigenetics and Cancer. Learning Materials in Biosciences, 2021, , 151-177.	0.4	1
47	Looking for a job in a dynamic and collaborative working place? LncRNAs are recruiting!. Cell, 2021, 184, 6019-6021.	28.9	1