

Isabel ChillÃ³n

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

2,036
citations

840776

11
h-index

1199594

12
g-index

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all docs

14
docs citations

14
times ranked

1977
citing authors

#	ARTICLE	IF	CITATIONS
1	HOTAIR Forms an Intricate and Modular Secondary Structure. <i>Molecular Cell</i> , 2015, 58, 353-361.	9.7	299
2	Conserved Pseudoknots in lincRNA MEG3 Are Essential for Stimulation of the p53 Pathway. <i>Molecular Cell</i> , 2019, 75, 982-995.e9.	9.7	138
3	Splicing of the <i>Sinorhizobium meliloti</i> Rmlnt1 group II intron provides evidence of retroelement behavior. <i>Nucleic Acids Research</i> , 2011, 39, 1095-1104.	14.5	66
4	Inverted repeat elements in the human lincRNA-p21 adopt a conserved secondary structure that regulates RNA function. <i>Nucleic Acids Research</i> , 2016, 44, gkw599.	14.5	64
5	The molecular structure of long non-coding RNAs: emerging patterns and functional implications. <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 2020, 55, 662-690.	5.2	51
6	Native Purification and Analysis of Long RNAs. <i>Methods in Enzymology</i> , 2015, 558, 3-37.	1.0	49
7	Visualizing group II intron dynamics between the first and second steps of splicing. <i>Nature Communications</i> , 2020, 11, 2837.	12.8	31
8	Predicted group II intron lineages E and F comprise catalytically active ribozymes. <i>Rna</i> , 2013, 19, 1266-1278.	3.5	16
9	Visualizing the functional 3D shape and topography of long noncoding RNAs by single-particle atomic force microscopy and in-solution hydrodynamic techniques. <i>Nature Protocols</i> , 2020, 15, 2107-2139.	12.0	14
10	Exon sequence requirements for excision in vivo of the bacterial group II intron Rmlnt1. <i>BMC Molecular Biology</i> , 2011, 12, 24.	3.0	12
11	In vitro characterization of the splicing efficiency and fidelity of the Rmlnt1 group II intron as a means of controlling the dispersion of its host mobile element. <i>Rna</i> , 2014, 20, 2000-2010.	3.5	11
12	Abstract IA25: Regulatory control of lincRNA function through formation of complex RNA structural motifs. , 2016, , .		0