

Eleonora De Klerk

List of Publications by Year in descending order

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Version: 2024-02-01

9
papers

723
citations

1163117

8
h-index

1474206

9
g-index

9
all docs

9
docs citations

9
times ranked

1633
citing authors

#	ARTICLE	IF	CITATIONS
1	Full-length mRNA sequencing uncovers a widespread coupling between transcription initiation and mRNA processing. <i>Genome Biology</i> , 2018, 19, 46.	8.8	106
2	An alanine expanded PABPN1 causes increased utilization of intronic polyadenylation sites. <i>Npj Aging and Mechanisms of Disease</i> , 2017, 3, 6.	4.5	15
3	Converging disease genes in ICF syndrome: <i>ZBTB24</i> controls expression of <i>CDCA7</i> in mammals. <i>Human Molecular Genetics</i> , 2016, 25, 4041-4051.	2.9	49
4	Alternative mRNA transcription, processing, and translation: insights from RNA sequencing. <i>Trends in Genetics</i> , 2015, 31, 128-139.	6.7	283
5	Assessing the translational landscape of myogenic differentiation by ribosome profiling. <i>Nucleic Acids Research</i> , 2015, 43, 4408-4428.	14.5	43
6	RNA sequencing: from tag-based profiling to resolving complete transcript structure. <i>Cellular and Molecular Life Sciences</i> , 2014, 71, 3537-3551.	5.4	33
7	Ribosome Profiling Uncovers the Role of uORFs in Translational Control of Gene Expression during Erythroblast Differentiation. <i>Blood</i> , 2014, 124, 2658-2658.	1.4	1
8	DeepSAGE Reveals Genetic Variants Associated with Alternative Polyadenylation and Expression of Coding and Non-coding Transcripts. <i>PLoS Genetics</i> , 2013, 9, e1003594.	3.5	45
9	Poly(A) binding protein nuclear 1 levels affect alternative polyadenylation. <i>Nucleic Acids Research</i> , 2012, 40, 9089-9101.	14.5	148