Eleonora De Klerk

List of Publications by Year in descending order

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