

J Andrew Berglund

List of Publications by Year in descending order

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Version: 2024-02-01

36
papers

1,989
citations

394421

19
h-index

361022

35
g-index

40
all docs

40
docs citations

40
times ranked

1997
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular characterization of myotonic dystrophy fibroblast cell lines for use in small molecule screening. <i>Science</i> , 2022, 25, 104198.	4.1	6
2	RNA structure probing to characterize RNA-protein interactions on low abundance pre-mRNA in living cells. <i>Rna</i> , 2021, 27, 343-358.	3.5	6
3	Zebrafish <i>mbnl</i> mutants model physical and molecular phenotypes of myotonic dystrophy. <i>DMM Disease Models and Mechanisms</i> , 2021, 14, .	2.4	7
4	CCG-CCG interruptions in high-penetrance SCA8 families increase RAN translation and protein toxicity. <i>EMBO Molecular Medicine</i> , 2021, 13, e14095.	6.9	12
5	Repeat length increases disease penetrance and severity in <i>C9orf72</i> ALS/FTD BAC transgenic mice. <i>Human Molecular Genetics</i> , 2021, 29, 3900-3918.	2.9	7
6	The potential of engineered eukaryotic RNA-binding proteins as molecular tools and therapeutics. <i>Wiley Interdisciplinary Reviews RNA</i> , 2020, 11, e1573.	6.4	13
7	Drug Screen Tugs at Common Thread for Repeat Disorders. <i>Trends in Pharmacological Sciences</i> , 2020, 41, 71-73.	8.7	3
8	Asian Zika Virus Isolate Significantly Changes the Transcriptional Profile and Alternative RNA Splicing Events in a Neuroblastoma Cell Line. <i>Viruses</i> , 2020, 12, 510.	3.3	25
9	Repeat-associated RNA structure and aberrant splicing. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2019, 1862, 194405.	1.9	23
10	Combination Treatment of Erythromycin and Furamidine Provides Additive and Synergistic Rescue of Mis-splicing in Myotonic Dystrophy Type 1 Models. <i>ACS Pharmacology and Translational Science</i> , 2019, 2, 247-263.	4.9	20
11	A CTG repeat-selective chemical screen identifies microtubule inhibitors as selective modulators of toxic CUG RNA levels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 20991-21000.	7.1	20
12	Mitigating RNA Toxicity in Myotonic Dystrophy using Small Molecules. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4017.	4.1	17
13	Transcriptome alterations in myotonic dystrophy skeletal muscle and heart. <i>Human Molecular Genetics</i> , 2019, 28, 1312-1321.	2.9	104
14	An engineered RNA binding protein with improved splicing regulation. <i>Nucleic Acids Research</i> , 2018, 46, 3152-3168.	14.5	15
15	Furamidine Rescues Myotonic Dystrophy Type I Associated Mis-Splicing through Multiple Mechanisms. <i>ACS Chemical Biology</i> , 2018, 13, 2708-2718.	3.4	26
16	Pseudouridine Modification Inhibits Muscleblind-like 1 (MBNL1) Binding to CCUG Repeats and Minimally Structured RNA through Reduced RNA Flexibility. <i>Journal of Biological Chemistry</i> , 2017, 292, 4350-4357.	3.4	43
17	Dose-Dependent Regulation of Alternative Splicing by MBNL Proteins Reveals Biomarkers for Myotonic Dystrophy. <i>PLoS Genetics</i> , 2016, 12, e1006316.	3.5	79
18	Conservation of context-dependent splicing activity in distant Muscleblind homologs. <i>Nucleic Acids Research</i> , 2016, 44, 8352-8362.	14.5	11

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19	Actinomycin D Specifically Reduces Expanded CUG Repeat RNA in Myotonic Dystrophy Models. <i>Cell Reports</i> , 2015, 13, 2386-2394.	6.4	74
20	Synthesis of N -substituted aryl amidines by strong base activation of amines. <i>Tetrahedron Letters</i> , 2015, 56, 4109-4111.	1.4	14
21	Biological Efficacy and Toxicity of Diamidines in Myotonic Dystrophy Type 1 Models. <i>Journal of Medicinal Chemistry</i> , 2015, 58, 5770-5780.	6.4	31
22	Modifications to toxic CUG RNAs induce structural stability, rescue mis-splicing in a myotonic dystrophy cell model and reduce toxicity in a myotonic dystrophy zebrafish model. <i>Nucleic Acids Research</i> , 2014, 42, 12768-12778.	14.5	27
23	Reducing Levels of Toxic RNA with Small Molecules. <i>ACS Chemical Biology</i> , 2013, 8, 2528-2537.	3.4	71
24	Combinatorial Mutagenesis of MBNL1 Zinc Fingers Elucidates Distinct Classes of Regulatory Events. <i>Molecular and Cellular Biology</i> , 2012, 32, 4155-4167.	2.3	22
25	Utilizing the GAAA Tetraloop/Receptor To Facilitate Crystal Packing and Determination of the Structure of a CUG RNA Helix. <i>Biochemistry</i> , 2012, 51, 8330-8337.	2.5	36
26	The four Zn fingers of MBNL1 provide a flexible platform for recognition of its RNA binding elements. <i>BMC Molecular Biology</i> , 2011, 12, 20.	3.0	35
27	Autoregulated Splicing of muscleblind-like 1 (MBNL1) Pre-mRNA. <i>Journal of Biological Chemistry</i> , 2011, 286, 34224-34233.	3.4	62
28	Role of RNA structure in regulating pre-mRNA splicing. <i>Trends in Biochemical Sciences</i> , 2010, 35, 169-178.	7.5	273
29	MBNL1 binds GC motifs embedded in pyrimidines to regulate alternative splicing. <i>Nucleic Acids Research</i> , 2010, 38, 2467-2484.	14.5	127
30	The protein factors MBNL1 and U2AF65 bind alternative RNA structures to regulate splicing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 9203-9208.	7.1	128
31	Pentamidine reverses the splicing defects associated with myotonic dystrophy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 18551-18556.	7.1	234
32	Transposition of two amino acids changes a promiscuous RNA binding protein into a sequence-specific RNA binding protein. <i>Rna</i> , 2008, 14, 78-88.	3.5	9
33	MBNL binds similar RNA structures in the CUG repeats of myotonic dystrophy and its pre-mRNA substrate cardiac troponin T. <i>Rna</i> , 2007, 13, 2238-2251.	3.5	153
34	The structural basis of myotonic dystrophy from the crystal structure of CUG repeats. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 16626-16631.	7.1	161
35	Expanding the Structural Repertoire of G-Quadruplexes. <i>Structure</i> , 2003, 11, 1315-1316.	3.3	0
36	The structure of an RNA dodecamer shows how tandem Uâ€“U base pairs increase the range of stable RNA structures and the diversity of recognition sites. <i>Structure</i> , 1996, 4, 917-930.	3.3	93