Jocelyn CÃ'té

List of Publications by Year in descending order

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52 papers

3,812 citations

30 h-index 51 g-index

55 all docs 55 docs citations

55 times ranked 5437 citing authors

#	Article	IF	CITATIONS
1	A novel CARM1–HuR axis involved in muscle differentiation and plasticity misregulated in spinal muscular atrophy. Human Molecular Genetics, 2022, 31, 1453-1470.	2.9	2
2	Differential regulation of autophagy by STAU1 in alveolar rhabdomyosarcoma and nonâ€transformed skeletal muscle cells. Cellular Oncology (Dordrecht), 2021, 44, 851-870.	4.4	7
3	Using affinity purification coupled with stable isotope labeling by amino acids in cell culture quantitative mass spectrometry to identify novel interactors/substrates of protein arginine methyltransferases. Methods, 2020, 175, 44-52.	3.8	3
4	PRMT7 methylates eukaryotic translation initiation factor $2\hat{l}_{\pm}$ and regulates its role in stress granule formation. Molecular Biology of the Cell, 2019, 30, 778-793.	2.1	31
5	A complex of C9ORF72 and p62 uses arginine methylation to eliminate stress granules by autophagy. Nature Communications, 2018, 9, 2794.	12.8	126
6	Novel Roles for Staufen1 in Embryonal and Alveolar Rhabdomyosarcoma via c-myc-dependent and -independent events. Scientific Reports, 2017, 7, 42342.	3.3	14
7	Muscle-specific expression of the RNA-binding protein Staufen1 induces progressive skeletal muscle atrophy via regulation of phosphatase tensin homolog. Human Molecular Genetics, 2017, 26, 1821-1838.	2.9	21
8	Misregulation of calcium-handling proteins promotes hyperactivation of calcineurin–NFAT signaling in skeletal muscle of DM1 mice. Human Molecular Genetics, 2017, 26, 2192-2206.	2.9	27
9	RNA binding protein RALY promotes Protein Arginine Methyltransferase 1 alternatively spliced isoform v2 relative expression and metastatic potential in breast cancer cells. International Journal of Biochemistry and Cell Biology, 2017, 91, 124-135.	2.8	27
10	Tudor Domain Containing Protein 3 Promotes Tumorigenesis and Invasive Capacity of Breast Cancer Cells. Scientific Reports, 2017, 7, 5153.	3.3	18
11	Atg5 Disassociates the V1V0-ATPase to Promote Exosome Production and Tumor Metastasis Independent of Canonical Macroautophagy. Developmental Cell, 2017, 43, 716-730.e7.	7.0	205
12	Lysine acetyltransferase NuA4 and acetyl-CoA regulate glucose-deprived stress granule formation in Saccharomyces cerevisiae. PLoS Genetics, 2017, 13, e1006626.	3.5	20
13	Staufen1 Regulates Multiple Alternative Splicing Events either Positively or Negatively in DM1 Indicating Its Role as a Disease Modifier. PLoS Genetics, 2016, 12, e1005827.	3.5	37
14	Staufen1 impairs stress granule formation in skeletal muscle cells from myotonic dystrophy type 1 patients. Molecular Biology of the Cell, 2016, 27, 1728-1739.	2.1	30
15	Staufen1s role as a splicing factor and a disease modifier in Myotonic Dystrophy Type I. Rare Diseases (Austin, Tex), 2016, 4, e1225644.	1.8	7
16	A novel role for CARM1 in promoting nonsense-mediated mRNA decay: potential implications for spinal muscular atrophy. Nucleic Acids Research, 2016, 44, 2661-2676.	14.5	29
17	Cross-talk between PRMT1-mediated methylation and ubiquitylation on RBM15 controls RNA splicing. ELife, 2015, 4, .	6.0	125
18	Identification of the PRMT1v1 and PRMT1v2 specific interactomes by quantitative mass spectrometry in breast cancer cells. Proteomics, 2015, 15, 2187-2197.	2.2	19

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19	Arginine methyltransferases as novel therapeutic targets for breast cancer. Mutagenesis, 2015, 30, 177-189.	2.6	41
20	IGF-1R Reduction Triggers Neuroprotective Signaling Pathways in Spinal Muscular Atrophy Mice. Journal of Neuroscience, 2015, 35, 12063-12079.	3.6	38
21	Protein arginine methyltransferase 7 promotes breast cancer cell invasion through the induction of MMP9 expression. Oncotarget, 2015, 6, 3013-3032.	1.8	65
22	Converging pathways involving microRNA-206 and the RNA-binding protein KSRP control post-transcriptionally utrophin A expression in skeletal muscle. Nucleic Acids Research, 2014, 42, 3982-3997.	14.5	23
23	Role of PRMTs in cancer: Could minor isoforms be leaving a mark?. World Journal of Biological Chemistry, 2014, 5, 115-29.	4.3	62
24	Nuclear Fragile X Mental Retardation Protein Is localized to Cajal Bodies. PLoS Genetics, 2013, 9, e1003890.	3.5	38
25	Activation of p38 signaling increases utrophin A expression in skeletal muscle via the RNA-binding protein KSRP and inhibition of AU-rich element-mediated mRNA decay: implications for novel DMD therapeutics. Human Molecular Genetics, 2013, 22, 3093-3111.	2.9	36
26	A novel function for the survival motoneuron protein as a translational regulator. Human Molecular Genetics, 2013, 22, 668-684.	2.9	106
27	Abstract 3789: The alternatively spliced PRMT1 isoform PRMT1 ν 2 promotes breast cancer cell survival and invasiveness , 2013, , .		0
28	Alternatively spliced protein arginine methyltransferase 1 isoform PRMT1v2 promotes the survival and invasiveness of breast cancer cells. Cell Cycle, 2012, 11, 4597-4612.	2.6	68
29	The RNA-binding protein Staufen1 is increased in DM1 skeletal muscle and promotes alternative pre-mRNA splicing. Journal of Cell Biology, 2012, 196, 699-712.	5.2	104
30	HuD interacts with survival motor neuron protein and can rescue spinal muscular atrophy-like neuronal defects. Human Molecular Genetics, 2011, 20, 553-579.	2.9	121
31	<i>S. pombe</i> replication protein Cdc18 (Cdc6) interacts with Swi6 (HP1) heterochromatin protein. Cell Cycle, 2011, 10, 323-336.	2.6	29
32	The RNA binding protein KSRP negatively regulates utrophin A expression in skeletal muscle. FASEB Journal, 2011, 25, 663.10.	0.5	0
33	In Vivo NMDA Receptor Activation Accelerates Motor Unit Maturation, Protects Spinal Motor Neurons, and Enhances SMN2 Gene Expression in Severe Spinal Muscular Atrophy Mice. Journal of Neuroscience, 2010, 30, 11288-11299.	3.6	43
34	Protein Arginine Methylation Facilitates Cotranscriptional Recruitment of Pre-mRNA Splicing Factors. Molecular and Cellular Biology, 2010, 30, 5245-5256.	2.3	43
35	hnRNP A1 regulates UV-induced NF-κB signalling through destabilization of cIAP1 mRNA. Cell Death and Differentiation, 2009, 16, 244-252.	11.2	44
36	eEF1A Is a Novel Component of the Mammalian Nuclear Protein Export Machinery. Molecular Biology of the Cell, 2008, 19, 5296-5308.	2.1	72

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37	TDRD3, a novel Tudor domain-containing protein, localizes to cytoplasmic stress granules. Human Molecular Genetics, 2008, 17, 3055-3074.	2.9	100
38	Alternative Splicing Yields Protein Arginine Methyltransferase 1 Isoforms with Distinct Activity, Substrate Specificity, and Subcellular Localization. Journal of Biological Chemistry, 2007, 282, 33009-33021.	3.4	156
39	KH-type splicing regulatory protein interacts with survival motor neuron protein and is misregulated in spinal muscular atrophy. Human Molecular Genetics, 2007, 17, 506-524.	2.9	89
40	The Arginine Methyltransferase CARM1 Regulates the Coupling of Transcription and mRNA Processing. Molecular Cell, 2007, 25, 71-83.	9.7	323
41	Tudor Domains Bind Symmetrical Dimethylated Arginines. Journal of Biological Chemistry, 2005, 280, 28476-28483.	3.4	218
42	A Proteomic Analysis of Arginine-methylated Protein Complexes. Molecular and Cellular Proteomics, 2003, 2, 1319-1330.	3.8	323
43	Sam68 RNA Binding Protein Is an In Vivo Substrate for Protein ArginineN-Methyltransferase 1. Molecular Biology of the Cell, 2003, 14, 274-287.	2.1	237
44	The Product of the Survival of Motor Neuron(SMN) Gene is a Human Telomerase-associated Protein. Molecular Biology of the Cell, 2002, 13, 3192-3202.	2.1	60
45	Symmetrical dimethylarginine methylation is required for the localization of SMN in Cajal bodies and pre-mRNA splicing. Journal of Cell Biology, 2002, 159, 957-969.	5.2	175
46	A protein-domain microarray identifies novel protein–protein interactions. Biochemical Journal, 2002, 367, 697-702.	3.7	158
47	Identification of Sam68 Arginine Glycine-rich Sequences Capable of Conferring Nonspecific RNA Binding to the GSG Domain. Journal of Biological Chemistry, 2001, 276, 30803-30811.	3.4	21
48	Polypyrimidine Track-binding Protein Binding Downstream of Caspase-2 Alternative Exon 9 Represses Its Inclusion. Journal of Biological Chemistry, 2001, 276, 8535-8543.	3.4	62
49	Caspase-2 pre-mRNA alternative splicing: Identification of an intronic element containing a decoy 3' acceptor site. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 938-943.	7.1	35
50	Aberrant Splicing of tau Pre-mRNA Caused by Intronic Mutations Associated with the Inherited Dementia Frontotemporal Dementia with Parkinsonism Linked to Chromosome 17. Molecular and Cellular Biology, 2000, 20, 4036-4048.	2.3	121
51	An element in the 5' common exon of the NCAM alternative splicing unit interacts with SR proteins and modulates 5' splice site selection. Nucleic Acids Research, 1999, 27, 2529-2537.	14.5	25
52	The U1 Small Nuclear Ribonucleoprotein/5′ Splice Site Interaction Affects U2AF65 Binding to the Downstream 3′ Splice Site. Journal of Biological Chemistry, 1995, 270, 4031-4036.	3.4	25