## Stefania Millevoi

List of Publications by Year in descending order

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STEEANIA MULEVOL

#	Article	IF	CITATIONS
1	Analysis of mRNA Translation by Polysome Profiling. Methods in Molecular Biology, 2022, 2404, 69-81.	0.4	2
2	Translational Regulation by hnRNP H/F Is Essential for the Proliferation and Survival of Glioblastoma. Cancers, 2022, 14, 1283.	1.7	3
3	G-Quadruplexes in RNA Biology: Recent Advances and Future Directions. Trends in Biochemical Sciences, 2021, 46, 270-283.	3.7	95
4	Activation of the integrated stress response confers vulnerability to mitoribosome-targeting antibiotics in melanoma. Journal of Experimental Medicine, 2021, 218, .	4.2	31
5	The NMD Pathway Regulates GABARAPL1 mRNA during the EMT. Biomedicines, 2021, 9, 1302.	1.4	1
6	Translation reprogramming by eIF3 linked to glioblastoma resistance. NAR Cancer, 2020, 2, zcaa020.	1.6	9
7	hnRNP H/F drive RNA G-quadruplex-mediated translation linked to genomic instability and therapy resistance in glioblastoma. Nature Communications, 2020, 11, 2661.	5.8	62
8	Microtubule-Driven Stress Granule Dynamics Regulate Inhibitory Immune Checkpoint Expression in T Cells. Cell Reports, 2019, 26, 94-107.e7.	2.9	42
9	RNA G-quadruplexes: emerging mechanisms in disease. Nucleic Acids Research, 2017, 45, gkw1280.	6.5	153
10	A novel cytoprotective function for the <scp>DNA</scp> repair protein Ku in regulating p53 <scp>mRNA</scp> translation andÂfunction. EMBO Reports, 2016, 17, 508-518.	2.0	25
11	hnRNP A1-mediated translational regulation of the G quadruplex-containing RON receptor tyrosine kinase mRNA linked to tumor progression. Oncotarget, 2016, 7, 16793-16805.	0.8	30
12	Stabilization of the G-quadruplex at the VEGF IRES represses cap-independent translation. RNA Biology, 2015, 12, 320-329.	1.5	53
13	Int6/eIF3e Is Essential for Proliferation and Survival of Human Glioblastoma Cells. International Journal of Molecular Sciences, 2014, 15, 2172-2190.	1.8	34
14	Destabilization of nucleophosmin mRNA by the HuR/KSRP complex is required for muscle fibre formation. Nature Communications, 2014, 5, 4190.	5.8	56
15	VEGF-A mRNA processing, stability and translation: a paradigm for intricate regulation of gene expression at the post-transcriptional level. Nucleic Acids Research, 2013, 41, 7997-8010.	6.5	190
16	Decreased efficiency of <i>MSH6</i> mRNA polyadenylation linked to a 20-base-pair duplication in Lynch syndrome families. Cell Cycle, 2012, 11, 2578-2580.	1.3	8
17	Gâ€quadruplexes in RNA biology. Wiley Interdisciplinary Reviews RNA, 2012, 3, 495-507.	3.2	247
18	Essential role for the interaction between hnRNP H/F and a G quadruplex in maintaining p53 pre-mRNA 3′-end processing and function during DNA damage. Genes and Development, 2011, 25, 220-225.	2.7	155

STEFANIA MILLEVOI

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19	The c.5242C>A BRCA1 missense variant induces exon skipping by increasing splicing repressors binding. Breast Cancer Research and Treatment, 2010, 120, 391-399.	1.1	23
20	Molecular mechanisms of eukaryotic pre-mRNA 3′ end processing regulation. Nucleic Acids Research, 2010, 38, 2757-2774.	6.5	322
21	Exon-Based Clustering of Murine Breast Tumor Transcriptomes Reveals Alternative Exons Whose Expression Is Associated with Metastasis. Cancer Research, 2010, 70, 896-905.	0.4	59
22	Widespread Estrogen-Dependent Repression of microRNAs Involved in Breast Tumor Cell Growth. Cancer Research, 2009, 69, 8332-8340.	0.4	225
23	A physical and functional link between splicing factors promotes pre-mRNA 3′ end processing. Nucleic Acids Research, 2009, 37, 4672-4683.	6.5	68
24	An interaction between U2AF 65 and CF Im links the splicing and 3′ end processing machineries. EMBO Journal, 2006, 25, 4854-4864.	3.5	179
25	A novel function for the U2AF 65 splicing factor in promoting preâ€mRNA 3′â€end processing. EMBO Reports, 2002, 3, 869-874.	2.0	57
26	Atypical binding of the neuronal POU protein N-Oct3 to noncanonical DNA targets. FEBS Journal, 2001, 268, 781-791.	0.2	15
27	The titin cDNA sequence and partial genomic sequences: Insights into the molecular genetics, cell biology and physiology of the titin filament system. Reviews of Physiology, Biochemistry and Pharmacology, 1999, 138, 19-55.	0.9	5
28	Conformational stability studies of the pleckstrin DEP domain: definition of the domain boundaries. BBA - Proteins and Proteomics, 1998, 1385, 157-164.	2.1	32
29	A monomeric mutant of Clostridium symbiosum glutamate dehydrogenase: Comparison with a structured monomeric intermediate obtained during refolding. Protein Science, 1998, 7, 966-974.	3.1	6
30	SH3 in muscles: solution structure of the SH3 domain from nebulin. Journal of Molecular Biology, 1998, 276, 189-202.	2.0	40
31	Characterization of nebulette and nebulin and emerging concepts of their roles for vertebrate Z-discs. Journal of Molecular Biology, 1998, 282, 111-123.	2.0	139
32	A Survey of the Primary Structure and the Interspecies Conservation of I-Band Titin's Elastic Elements in Vertebrates. Journal of Structural Biology, 1998, 122, 206-215.	1.3	51
33	Refined Localisation of the Genes for Nebulin and Titin on Chromosome 2q Allows the Assignment of Nebulin as a Candidate Gene for Autosomal Recessive Nemaline Myopathy. European Journal of Human Genetics, 1997, 5, 229-234.	1.4	26
34	Refolding Pathway and Association Intermediates of Glutamate Dehydrogenase from the Hyperthermophile Pyrococcus furiosus. FEBS Journal, 1996, 239, 679-685.	0.2	12
35	NAD-dependent glutamate dehydrogenase from the thermophilic eubacterium Bacillus acidocaldarius. Comparative Biochemistry and Physiology Part B: Comparative Biochemistry, 1994, 109, 691-699.	0.2	3