

Julia Morales

List of Publications by Year in descending order

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3,077
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331670

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docs citations

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times ranked

4187
citing authors

#	ARTICLE	IF	CITATIONS
1	A Peak of H3T3 Phosphorylation Occurs in Synchrony with Mitosis in Sea Urchin Early Embryos. <i>Cells</i> , 2020, 9, 898.	4.1	4
2	mTOR Signaling at the Crossroad between Metazoan Regeneration and Human Diseases. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2718.	4.1	26
3	Translational Control of Canonical and Non-Canonical Translation Initiation Factors at the Sea Urchin Egg to Embryo Transition. <i>International Journal of Molecular Sciences</i> , 2019, 20, 626.	4.1	5
4	In vivo analysis of protein translation activity in sea urchin eggs and embryos. <i>Methods in Cell Biology</i> , 2019, 151, 335-352.	1.1	4
5	Translatome analysis at the egg-to-embryo transition in sea urchin. <i>Nucleic Acids Research</i> , 2018, 46, 4607-4621.	14.5	19
6	Toward Multiscale Modeling of Molecular and Biochemical Events Occurring at Fertilization Time in Sea Urchins. <i>Results and Problems in Cell Differentiation</i> , 2018, 65, 69-89.	0.7	2
7	Analysis of translation using polysome profiling. <i>Nucleic Acids Research</i> , 2017, 45, gkw907.	14.5	119
8	MAPK/ERK activity is required for the successful progression of mitosis in sea urchin embryos. <i>Developmental Biology</i> , 2017, 421, 194-203.	2.0	12
9	Translational Control in Echinoderms: The Calm Before the Storm. , 2016, , 413-434.		5
10	Model of the delayed translation of cyclin B maternal mRNA after sea urchin fertilization. <i>Molecular Reproduction and Development</i> , 2016, 83, 1070-1082.	2.0	4
11	Cyclin B Translation Depends on mTOR Activity after Fertilization in Sea Urchin Embryos. <i>PLoS ONE</i> , 2016, 11, e0150318.	2.5	18
12	Modelization of the regulation of protein synthesis following fertilization in sea urchin shows requirement of two processes: a destabilization of eIF4E:4E-BP complex and a great stimulation of the 4E-BP-degradation mechanism, both rapamycin-sensitive. <i>Frontiers in Genetics</i> , 2014, 5, 117.	2.3	7
13	Activation of a GPCR leads to eIF4G phosphorylation at the 5' cap and to IRES-dependent translation. <i>Journal of Molecular Endocrinology</i> , 2014, 52, 373-382.	2.5	9
14	Tracking a refined eIF4E-binding motif reveals Angel1 as a new partner of eIF4E. <i>Nucleic Acids Research</i> , 2013, 41, 7783-7792.	14.5	25
15	The Ectocarpus Genome and Brown Algal Genomics. <i>Advances in Botanical Research</i> , 2012, 64, 141-184.	1.1	18
16	mRNA-Selective Translation Induced by FSH in Primary Sertoli Cells. <i>Molecular Endocrinology</i> , 2012, 26, 669-680.	3.7	29
17	Dephosphorylation of eIF2 α is essential for protein synthesis increase and cell cycle progression after sea urchin fertilization. <i>Developmental Biology</i> , 2012, 365, 303-309.	2.0	15
18	The Ectocarpus genome and the independent evolution of multicellularity in brown algae. <i>Nature</i> , 2010, 465, 617-621.	27.8	774

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19	A Variant Mimicking Hyperphosphorylated 4E-BP Inhibits Protein Synthesis in a Sea Urchin Cell-Free, Cap-Dependent Translation System. <i>PLoS ONE</i> , 2009, 4, e5070.	2.5	31
20	Inhibition of translation and modification of translation factors during apoptosis induced by the DNA-damaging agent MMS in sea urchin embryos. <i>Experimental Cell Research</i> , 2008, 314, 961-968.	2.6	17
21	Chromium(III) Triggers the DNA-Damaged Checkpoint of the Cell Cycle and Induces a Functional Increase of 4E-BP. <i>Chemical Research in Toxicology</i> , 2008, 21, 542-549.	3.3	12
22	After fertilization of sea urchin eggs, eIF4G is post-translationally modified and associated with the cap-binding protein eIF4E. <i>Journal of Cell Science</i> , 2007, 120, 425-434.	2.0	19
23	The Genome of the Sea Urchin <i>Strongylocentrotus purpuratus</i> . <i>Science</i> , 2006, 314, 941-952.	12.6	1,018
24	Translational control genes in the sea urchin genome. <i>Developmental Biology</i> , 2006, 300, 293-307.	2.0	33
25	The sea urchin kinome: A first look. <i>Developmental Biology</i> , 2006, 300, 180-193.	2.0	84
26	The genomic repertoire for cell cycle control and DNA metabolism in <i>S. purpuratus</i> . <i>Developmental Biology</i> , 2006, 300, 238-251.	2.0	48
27	eEF1B: At the dawn of the 21st century. <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 2006, 1759, 13-31.	2.4	98
28	A glyphosate-based pesticide impinges on transcription. <i>Toxicology and Applied Pharmacology</i> , 2005, 203, 1-8.	2.8	55
29	Embryonic-stage-dependent changes in the level of eIF4E-binding proteins during early development of sea urchin embryos. <i>Journal of Cell Science</i> , 2005, 118, 1385-1394.	2.0	30
30	Translational control during mitosis. <i>Biochimie</i> , 2005, 87, 805-811.	2.6	43
31	Formulated Glyphosate Activates the DNA-Response Checkpoint of the Cell Cycle Leading to the Prevention of G2/M Transition. <i>Toxicological Sciences</i> , 2004, 82, 436-442.	3.1	42
32	Signal transduction pathways that contribute to CDK1/cyclin B activation during the first mitotic division in sea urchin embryos. <i>Experimental Cell Research</i> , 2004, 296, 347-357.	2.6	24
33	Characterization of carbonic anhydrases from <i>Riftia pachyptila</i> , a symbiotic invertebrate from deep-sea hydrothermal vents. <i>Proteins: Structure, Function and Bioinformatics</i> , 2003, 51, 327-339.	2.6	24
34	M-phase regulation of the recruitment of mRNAs onto polysomes using the CDK1/cyclin B inhibitor aminopurvalanol. <i>Biochemical and Biophysical Research Communications</i> , 2003, 306, 880-886.	2.1	6
35	EIF4E/4E-BP dissociation and 4E-BP degradation in the first mitotic division of the sea urchin embryo. <i>Developmental Biology</i> , 2003, 255, 428-439.	2.0	50
36	eIF4E Association with 4E-BP Decreases Rapidly Following Fertilization in Sea Urchin. <i>Developmental Biology</i> , 2001, 232, 275-283.	2.0	48

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37	Protein translation during early cell divisions of sea urchin embryos regulated at the level of polypeptide chain elongation and highly sensitive to natural polyamines. <i>Zygote</i> , 2001, 9, 229-236.	1.1	15
38	Assembly of the $\hat{\pm}$ -Globin mRNA Stability Complex Reflects Binary Interaction between the Pyrimidine-Rich 3' UTR Untranslated Region Determinant and Poly(C) Binding Protein $\hat{\pm}$ CP. <i>Molecular and Cellular Biology</i> , 1999, 19, 4572-4581.	2.3	123
39	Sequence Divergence in the 3' UTR Untranslated Regions of Human $\hat{\pm}$ - and $\hat{\pm}$ -Globin mRNAs Mediates a Difference in Their Stabilities and Contributes to Efficient $\hat{\pm}$ -to- $\hat{\pm}$ Gene Developmental Switching. <i>Molecular and Cellular Biology</i> , 1998, 18, 2173-2183.	2.3	39
40	Destabilization of Human $\hat{\pm}$ -Globin mRNA by Translation Anti-termination Is Controlled during Erythroid Differentiation and Is Paralleled by Phased Shortening of the Poly(A) Tail. <i>Journal of Biological Chemistry</i> , 1997, 272, 6607-6613.	3.4	59
41	Phosphorylation of elongation factor-1 (EF-1) by cdc2 kinase. , 1995, 1, 265-270.		21
42	Expression of elongation factor $\hat{\pm}$ (EF- $\hat{\pm}$) and $\hat{\pm}$ (EF- $\hat{\pm}$) are uncoupled in early <i>Xenopus</i> embryos. <i>Genesis</i> , 1993, 14, 440-448.	2.1	5
43	Targets of MPF during meiotic cell division. <i>Biology of the Cell</i> , 1992, 76, 218-218.	2.0	0
44	Phosphorylation of <i>Xenopus</i> elongation factor- $\hat{\pm}$ by cdc2 protein kinase: Identification of the phosphorylation site. <i>Experimental Cell Research</i> , 1992, 202, 549-551.	2.6	21