

# Luis MarÃ-a Vaschetto

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6840831/publications.pdf>

Version: 2024-02-01

17  
papers

145  
citations

1307594

7  
h-index

1199594

12  
g-index

20  
all docs

20  
docs citations

20  
times ranked

239  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of leaf rust resistance genes in selected Argentinean bread wheat cultivars by gene postulation and molecular markers. <i>Electronic Journal of Biotechnology</i> , 2011, 14, .	2.2	31
2	miRNA activation is an endogenous gene expression pathway. <i>RNA Biology</i> , 2018, 15, 1-3.	3.1	29
3	Genetic structure of Argentinean hexaploid wheat germplasm. <i>Genetics and Molecular Biology</i> , 2013, 36, 391-399.	1.3	15
4	The disjunct pattern of the Neotropical harvestman <i>Discocyrtus dilatatus</i> (Gonyleptidae) explained by climate-driven range shifts in the Quaternary: Paleodistributional and molecular evidence. <i>PLoS ONE</i> , 2017, 12, e0187983.	2.5	11
5	Miniature Inverted-repeat Transposable Elements (MITEs) and their effects on the regulation of major genes in cereal grass genomes. <i>Molecular Breeding</i> , 2016, 36, 1.	2.1	8
6	The emerging importance of noncoding RNAs in the insecticide tolerance, with special emphasis on <i>Plutella xylostella</i> (Lepidoptera: Plutellidae). <i>Wiley Interdisciplinary Reviews RNA</i> , 2019, 10, e1539.	6.4	8
7	High genetic diversity in the harvestman <i>Geraecormobius sylvarum</i> (Arachnida, Opiliones). <i>Tj ETQq1 1 0.784314 rgBT /Overloc</i> sequences. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2015, 53, 211-218.	1.4	7
8	Modulating signaling networks by CRISPR/Cas9-mediated transposable element insertion. <i>Current Genetics</i> , 2018, 64, 405-412.	1.7	7
9	Palaeoclimatic distribution models predict Pleistocene refuges for the Neotropical harvestman <i>Geraecormobius sylvarum</i> (Arachnida: Opiliones: Gonyleptidae). <i>Journal of Natural History</i> , 2017, 51, 17-32.	0.5	6
10	Exploring an Emerging Issue: Crop Epigenetics. <i>Plant Molecular Biology Reporter</i> , 2015, 33, 751-755.	1.8	5
11	The Role of Sequence Duplication in Transcriptional Regulation and Genome Evolution. <i>Current Genomics</i> , 2019, 20, 405-408.	1.6	5
12	RNA Activation: A Diamond in the Rough for Genome Engineers. <i>Journal of Cellular Biochemistry</i> , 2018, 119, 247-249.	2.6	4
13	The Emergence of Non-coding RNAs as Versatile and Efficient Therapeutic Tools. <i>Current Gene Therapy</i> , 2019, 19, 289-289.	2.0	2
14	Small Activating RNAs as Promising Agents for Biotechnological Use. <i>Current Pharmaceutical Biotechnology</i> , 2018, 19, 602-603.	1.6	1
15	The Critical Role of Epigenetic Regulation in Developmental Programming of Higher Organisms. <i>Current Genomics</i> , 2019, 20, 403-404.	1.6	1
16	Understanding the role of protein interaction motifs in transcriptional regulators: implications for crop improvement. <i>Briefings in Functional Genomics</i> , 2016, 16, elw022.	2.7	0
17	Cereal Circular RNAs (circRNAs): An Overview of the Computational Resources for Identification and Analysis. <i>Methods in Molecular Biology</i> , 2020, 2072, 157-163.	0.9	0