

# Alejandro Villar-Briones

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4428971/publications.pdf>

Version: 2024-02-01

8  
papers

3,348  
citations

1307594

7  
h-index

1720034

7  
g-index

9  
all docs

9  
docs citations

9  
times ranked

6387  
citing authors

#	ARTICLE	IF	CITATIONS
1	Addendum: Aird, S.D. et al. Coralsnake Venomics: Analyses of Venom Gland Transcriptomes and Proteomes of Six Brazilian Taxa. <i>Toxins</i> 2017, 9(6), 187. <i>Toxins</i> , 2018, 10, 172.	3.4	0
2	Organic and Peptidyl Constituents of Snake Venoms: The Picture Is Vastly More Complex Than We Imagined. <i>Toxins</i> , 2018, 10, 392.	3.4	20
3	Coralsnake Venomics: Analyses of Venom Gland Transcriptomes and Proteomes of Six Brazilian Taxa. <i>Toxins</i> , 2017, 9, 187.	3.4	66
4	Polyamines as Snake Toxins and Their Probable Pharmacological Functions in Envenomation. <i>Toxins</i> , 2016, 8, 279.	3.4	8
5	Developmental Progression in the Coral <i>Acropora digitifera</i> Is Controlled by Differential Expression of Distinct Regulatory Gene Networks. <i>Genome Biology and Evolution</i> , 2016, 8, 851-870.	2.5	27
6	Snake venoms are integrated systems, but abundant venom proteins evolve more rapidly. <i>BMC Genomics</i> , 2015, 16, 647.	2.8	66
7	Quantitative high-throughput profiling of snake venom gland transcriptomes and proteomes ( <i>Ovophis okinavensis</i> and <i>Protobothrops flavoviridis</i> ). <i>BMC Genomics</i> , 2013, 14, 790.	2.8	130
8	MZmine 2: Modular framework for processing, visualizing, and analyzing mass spectrometry-based molecular profile data. <i>BMC Bioinformatics</i> , 2010, 11, 395.	2.6	3,031