

# Michael A Dimattia

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10467599/publications.pdf>

Version: 2024-02-01

11

papers

1,294

citations

949033

11

h-index

1427216

11

g-index

11

all docs

11

docs citations

11

times ranked

2394

citing authors

#	ARTICLE	IF	CITATIONS
1	The Structure of HIV-1 Rev Filaments Suggests a Bilateral Model for Rev-RRE Assembly. <i>Structure</i> , 2016, 24, 1068-1080.	1.6	22
2	Identification and Mutagenesis of the Adeno-Associated Virus 5 Sialic Acid Binding Region. <i>Journal of Virology</i> , 2015, 89, 1660-1672.	1.5	41
3	A Cell-penetrating Antibody Fragment against HIV-1 Rev Has High Antiviral Activity. <i>Journal of Biological Chemistry</i> , 2014, 289, 20222-20233.	1.6	20
4	An activating NLRC4 inflammasome mutation causes autoinflammation with recurrent macrophage activation syndrome. <i>Nature Genetics</i> , 2014, 46, 1140-1146.	9.4	585
5	Antigenic Switching of Hepatitis B Virus by Alternative Dimerization of the Capsid Protein. <i>Structure</i> , 2013, 21, 133-142.	1.6	61
6	Structural Insights into Adeno-Associated Virus Serotype 5. <i>Journal of Virology</i> , 2013, 87, 11187-11199.	1.5	69
7	Capsid Antibodies to Different Adeno-Associated Virus Serotypes Bind Common Regions. <i>Journal of Virology</i> , 2013, 87, 9111-9124.	1.5	102
8	Structural Insight into the Unique Properties of Adeno-Associated Virus Serotype 9. <i>Journal of Virology</i> , 2012, 86, 6947-6958.	1.5	163
9	Implications of the HIV-1 Rev dimer structure at 3.2 Å resolution for multimeric binding to the Rev response element. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 5810-5814.	3.3	124
10	Human Bocavirus Capsid Structure: Insights into the Structural Repertoire of the <i>Parvoviridae</i> . <i>Journal of Virology</i> , 2010, 84, 5880-5889.	1.5	79
11	Generation and Characterization of a Chimeric Rabbit/Human Fab for Co-Crystallization of HIV-1 Rev. <i>Journal of Molecular Biology</i> , 2010, 397, 697-708.	2.0	28