## Mark J Young

## List of Publications by Year in descending order

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214721 218592 4,156 44 26 47 citations g-index h-index papers 49 49 49 4733 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Gut bacteriophage dynamics during fecal microbial transplantation in subjects with metabolic syndrome. Gut Microbes, $2021, 13, 1-15$ .	4.3	24
2	Effect of Inactivation Methods on SARS-CoV-2 Virion Protein and Structure. Viruses, 2021, 13, 562.	1.5	33
3	Bacterial Viruses Subcommittee and Archaeal Viruses Subcommittee of the ICTV: update of taxonomy changes in 2021. Archives of Virology, 2021, 166, 3239-3244.	0.9	24
4	An Uncultivated Virus Infecting a Nanoarchaeal Parasite in the Hot Springs of Yellowstone National Park. Journal of Virology, 2020, 94, .	1.5	10
5	The intriguing world of archaeal viruses. PLoS Pathogens, 2020, 16, e1008574.	2.1	16
6	Discovery and Characterization of Thermoproteus Spherical Piliferous Virus 1: a Spherical Archaeal Virus Decorated with Unusual Filaments. Journal of Virology, 2020, 94, .	1.5	2
7	Survey of high-resolution archaeal virus structures. Current Opinion in Virology, 2019, 36, 74-83.	2.6	10
8	The Molecular Mechanism of Cellular Attachment for an Archaeal Virus. Structure, 2019, 27, 1634-1646.e3.	1.6	21
9	Minimum Information about an Uncultivated Virus Genome (MIUViG). Nature Biotechnology, 2019, 37, 29-37.	9.4	414
10	A virus or more in (nearly) every cell: ubiquitous networks of virus–host interactions in extreme environments. ISME Journal, 2018, 12, 1706-1714.	4.4	94
11	Structural studies of <i>Acidianus</i> tailed spindle virus reveal a structural paradigm used in the assembly of spindle-shaped viruses. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 2120-2125.	3.3	29
12	Discovering novel hydrolases from hot environments. Biotechnology Advances, 2018, 36, 2077-2100.	6.0	38
13	Single-cell genomics of co-sorted Nanoarchaeota suggests novel putative host associations and diversification of proteins involved in symbiosis. Microbiome, 2018, 6, 161.	4.9	44
14	Archaeal Viruses from High-Temperature Environments. Genes, 2018, 9, 128.	1.0	54
15	Isolation and Characterization of Metallosphaera Turreted Icosahedral Virus, a Founding Member of a New Family of Archaeal Viruses. Journal of Virology, 2017, 91, .	1.5	19
16	The transcript cleavage factor paralogue TFS4 is a potent RNA polymerase inhibitor. Nature Communications, 2017, 8, 1914.	5.8	18
17	The Human Gut Phage Community and Its Implications for Health and Disease. Viruses, 2017, 9, 141.	1.5	206
18	Coupling Peptide Antigens to Virus-Like Particles or to Protein Carriers Influences the Th1/Th2 Polarity of the Resulting Immune Response. Vaccines, 2016, 4, 15.	2.1	20

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19	Novel viral genomes identified from six metagenomes reveal wide distribution of archaeal viruses and high viral diversity in terrestrial hot springs. Environmental Microbiology, 2016, 18, 863-874.	1.8	53
20	Healthy human gut phageome. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 10400-10405.	3.3	439
21	Structure-Based Mutagenesis of Sulfolobus Turreted Icosahedral Virus B204 Reveals Essential Residues in the Virion-Associated DNA-Packaging ATPase. Journal of Virology, 2016, 90, 2729-2739.	1.5	8
22	Acidianus Tailed Spindle Virus: a New Archaeal Large Tailed Spindle Virus Discovered by Culture-Independent Methods. Journal of Virology, 2016, 90, 3458-3468.	1.5	27
23	Viral assemblage composition in Yellowstone acidic hot springs assessed by network analysis. ISME Journal, 2015, 9, 2162-2177.	4.4	48
24	40 Years of archaeal virology: Expanding viral diversity. Virology, 2015, 479-480, 369-378.	1.1	41
25	Nanoarchaeota, Their Sulfolobales Host, and Nanoarchaeota Virus Distribution across Yellowstone National Park Hot Springs. Applied and Environmental Microbiology, 2015, 81, 7860-7868.	1.4	63
26	Large Tailed Spindle Viruses of Archaea: a New Way of Doing Viral Business. Journal of Virology, 2015, 89, 9146-9149.	1.5	19
27	CRISPR-Induced Distributed Immunity in Microbial Populations. PLoS ONE, 2014, 9, e101710.	1.1	67
28	A Survey of Protein Structures from Archaeal Viruses. Life, 2013, 3, 118-130.	1.1	6
29	Identification of Novel Positive-Strand RNA Viruses by Metagenomic Analysis of Archaea-Dominated Yellowstone Hot Springs. Journal of Virology, 2012, 86, 5562-5573.	1.5	107
30	Development of a genetic system for the archaeal virus Sulfolobus turreted icosahedral virus (STIV). Virology, 2011, 415, 6-11.	1.1	29
31	Two-component magnetic structure of iron oxide nanoparticles mineralized in <i>Listeria innocua</i> protein cages. Journal of Applied Physics, 2010, 107, .	1.1	13
32	Particle Assembly and Ultrastructural Features Associated with Replication of the Lytic Archaeal Virus <i>Sulfolobus</i> Turreted Icosahedral Virus. Journal of Virology, 2009, 83, 5964-5970.	1.5	96
33	From Metal Binding to Nanoparticle Formation: Monitoring Biomimetic Iron Oxide Synthesis within Protein Cages using Mass Spectrometry. Angewandte Chemie - International Edition, 2009, 48, 4772-4776.	7.2	26
34	Monitoring Biomimetic Platinum Nanocluster Formation Using Mass Spectrometry and Clusterâ€Dependent H <sub>2</sub> Production. Angewandte Chemie - International Edition, 2008, 47, 7845-7848.	7.2	40
35	Plant Viruses as Biotemplates for Materials and Their Use in Nanotechnology. Annual Review of Phytopathology, 2008, 46, 361-384.	3.5	233
36	High-Density Targeting of a Viral Multifunctional Nanoplatform to a Pathogenic, Biofilm-Forming Bacterium. Chemistry and Biology, 2007, 14, 387-398.	6.2	58

#	Article	IF	CITATIONS
37	Controlled Ligand Display on a Symmetrical Protein-Cage Architecture Through Mixed Assembly. Small, 2006, 2, 962-966.	5.2	61
38	Melanoma and Lymphocyte Cell-Specific Targeting Incorporated into a Heat Shock Protein Cage Architecture. Chemistry and Biology, 2006, 13, 161-170.	6.2	146
39	Paramagnetic viral nanoparticles as potential high-relaxivity magnetic resonance contrast agents. Magnetic Resonance in Medicine, 2005, 54, 807-812.	1.9	198
40	From The Cover: The structure of a thermophilic archaeal virus shows a double-stranded DNA viral capsid type that spans all domains of life. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 7716-7720.	3.3	219
41	Heterologous expression of the modified coat protein of Cowpea chlorotic mottle bromovirus results in the assembly of protein cages with altered architectures and function. Journal of General Virology, 2004, 85, 1049-1053.	1.3	96
42	Metal binding to cowpea chlorotic mottle virus using terbium(III) fluorescence. Journal of Biological Inorganic Chemistry, 2003, 8, 721-725.	1.1	52
43	Viruses of hyperthermophilic Archaea. Research in Microbiology, 2003, 154, 474-482.	1.0	33
44	Host–guest encapsulation of materials by assembled virus protein cages. Nature, 1998, 393, 152-155.	13.7	887