Sarah M Mcdonald

List of Publications by Year in descending order

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687363 677142 22 984 13 22 citations h-index g-index papers 23 23 23 1320 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	<i>In Vitro</i> Double-Stranded RNA Synthesis by Rotavirus Polymerase Mutants with Lesions at Core Shell Contact Sites. Journal of Virology, 2019, 93, .	3.4	8
2	Cryo-EM Reveals Architectural Diversity in Active Rotavirus Particles. Computational and Structural Biotechnology Journal, 2019, 17, 1178-1183.	4.1	5
3	Cryoâ€EMâ€Onâ€aâ€Chip: Customâ€Designed Substrates for the 3D Analysis of Macromolecules. Small, 2019, 1 1900918.	¹⁵ 10.0	5
4	Group A Rotavirus VP1 Polymerase and VP2 Core Shell Proteins: Intergenotypic Sequence Variation and <i>In Vitro</i> Functional Compatibility. Journal of Virology, 2019, 93, .	3.4	17
5	A Temperature-Sensitive Lesion in the N-Terminal Domain of the Rotavirus Polymerase Affects Its Intracellular Localization and Enzymatic Activity. Journal of Virology, 2017, 91, .	3.4	12
6	Genetic determinants restricting the reassortment of heterologous NSP2 genes into the simian rotavirus SA11 genome. Scientific Reports, 2017, 7, 9301.	3.3	9
7	Rotavirus genome replication: Some assembly required. PLoS Pathogens, 2017, 13, e1006242.	4.7	30
8	Determinants of VH1-46 Cross-Reactivity to Pemphigus Vulgaris Autoantigen Desmoglein 3 and Rotavirus Antigen VP6. Journal of Immunology, 2016, 197, 1065-1073.	0.8	21
9	Distinguishing the genotype 1 genes and proteins of human Wa-like rotaviruses vs. porcine rotaviruses. Infection, Genetics and Evolution, 2016, 43, 6-14.	2.3	8
10	Reassortment in segmented RNA viruses: mechanisms and outcomes. Nature Reviews Microbiology, 2016, 14, 448-460.	28.6	259
11	Electron microscopic analysis of rotavirus assembly-replication intermediates. Virology, 2015, 477, 32-41.	2.4	21
12	A Non-Symmetric Reconstruction Technique for Transcriptionally-Active Viral Assemblies. Journal of Analytical & Molecular Techniques, 2015, 2, .	0.0	3
13	Structural dynamics of viral nanomachines. Technology, 2014, 02, 44-48.	1.4	5
14	Analysis of Human Rotaviruses from a Single Location Over an 18-Year Time Span Suggests that Protein Coadaption Influences Gene Constellations. Journal of Virology, 2014, 88, 9842-9863.	3.4	23
15	PCR-based approach to distinguish group A human rotavirus genotype $1\mathrm{vs.}$ genotype $2\mathrm{genes.}$ Journal of Virological Methods, $2013,194,197\text{-}205.$	2.1	6
16	RNA synthetic mechanisms employed by diverse families of RNA viruses. Wiley Interdisciplinary Reviews RNA, 2013, 4, 351-367.	6.4	77
17	Rotavirus core shell subdomains involved in polymerase encapsidation into virus-like particles. Journal of General Virology, 2013, 94, 1818-1826.	2.9	17
18	Diversity and Relationships of Cocirculating Modern Human Rotaviruses Revealed Using Large-Scale Comparative Genomics. Journal of Virology, 2012, 86, 9148-9162.	3.4	45

#	Article	IF	CITATION
19	Structural insights into the coupling of virion assembly and rotavirus replication. Nature Reviews Microbiology, 2012, 10, 165-177.	28.6	182
20	Intra-genotypic diversity of archival G4P[8] human rotaviruses from Washington, DC. Infection, Genetics and Evolution, 2011, 11, 1586-1594.	2.3	26
21	Rotavirus VP2 Core Shell Regions Critical for Viral Polymerase Activation. Journal of Virology, 2011, 85, 3095-3105.	3.4	57
22	Mechanism for Coordinated RNA Packaging and Genome Replication by Rotavirus Polymerase VP1. Structure, 2008, 16, 1678-1688.	3.3	148