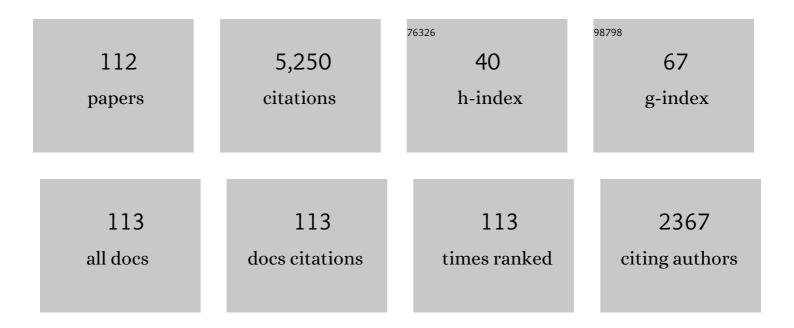
List of Publications by Year in descending order

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ANNE E SIMON

#	Article	IF	CITATIONS
1	Structural characterization of a new subclass of panicum mosaic virus-like 3′ cap-independent translation enhancer. Nucleic Acids Research, 2022, , .	14.5	2
2	Identification of Novel 5′ and 3′ Translation Enhancers in Umbravirus-Like Coat Protein-Deficient RNA Replicons. Journal of Virology, 2022, 96, e0173621.	3.4	5
3	Targeting of viral RNAs by Upf1-mediated RNA decay pathways. Current Opinion in Virology, 2021, 47, 1-8.	5.4	19
4	<i>Opium Poppy Mosaic Virus</i> Has an Xrn-Resistant, Translated Subgenomic RNA and a BTE 3′ CITE. Journal of Virology, 2021, 95, .	3.4	17
5	Structural Analysis and Whole Genome Mapping of a New Type of Plant Virus Subviral RNA: Umbravirus-Like Associated RNAs. Viruses, 2021, 13, 646.	3.3	24
6	Complete Nucleotide Sequence, Genome Organization, and Comparative Genomic Analyses of Citrus Yellow-Vein Associated Virus (CYVaV). Frontiers in Microbiology, 2021, 12, 683130.	3.5	18
7	Genome characterization of fig umbra-like virus. Virus Genes, 2021, 57, 566-570.	1.6	10
8	In Tribute to Michael Goodin. Viruses, 2021, 13, 78.	3.3	6
9	The Multifunctional Long-Distance Movement Protein of <i>Pea Enation Mosaic Virus 2</i> Protects Viral and Host Transcripts from Nonsense-Mediated Decay. MBio, 2020, 11, .	4.1	23
10	RNA2Drawer: geometrically strict drawing of nucleic acid structures with graphical structure editing and highlighting of complementary subsequences. RNA Biology, 2019, 16, 1667-1671.	3.1	51
11	RNA virus evasion of nonsense-mediated decay. PLoS Pathogens, 2018, 14, e1007459.	4.7	47
12	Unusual dicistronic expression from closely spaced initiation codons in an umbravirus subgenomic RNA. Nucleic Acids Research, 2018, 46, 11726-11742.	14.5	12
13	SELEX and SHAPE reveal that sequence motifs and an extended hairpin in the 5' portion of Turnip crinkle virus satellite RNA C mediate fitness in plants. Virology, 2018, 520, 137-152.	2.4	3
14	Trajectories of the ribosome as a Brownian nanomachine. journal of hand surgery Asian-Pacific volume, The, 2018, , 463-475.	0.4	2
15	A Sequence-Independent, Unstructured Internal Ribosome Entry Site Is Responsible for Internal Expression of the Coat Protein of Turnip Crinkle Virus. Journal of Virology, 2017, 91, .	3.4	19
16	Combined single molecule experimental and computational approaches for understanding the unfolding pathway of a viral translation enhancer that participates in a conformational switch. RNA Biology, 2017, 14, 1466-1472.	3.1	1
17	Differential use of 3'CITEs by the subgenomic RNA of Pea enation mosaic virus 2. Virology, 2017, 510, 194-204.	2.4	20
18	RNase III nucleases from diverse kingdoms serve as antiviral effectors. Nature, 2017, 547, 114-117.	27.8	57

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19	Concerted action of two 3′ cap-independent translation enhancers increases the competitive strength of translated viral genomes. Nucleic Acids Research, 2017, 45, 9558-9572.	14.5	23
20	Folding behavior of a T-shaped, ribosome-binding translation enhancer implicated in a wide-spread conformational switch. ELife, 2017, 6, .	6.0	15
21	Multiple Cis-acting elements modulate programmed -1 ribosomal frameshifting in Pea enation mosaic virus. Nucleic Acids Research, 2016, 44, 878-895.	14.5	31
22	An RNA Element That Facilitates Programmed Ribosomal Readthrough in Turnip Crinkle Virus Adopts Multiple Conformations. Journal of Virology, 2016, 90, 8575-8591.	3.4	33
23	Editorial overview: Virus–vector interactions. Current Opinion in Virology, 2015, 15, iv-vi.	5.4	1
24	In Vivo, Large-Scale Preparation of Uniformly 15N- and Site-Specifically 13C-Labeled Homogeneous, Recombinant RNA for NMR Studies. Methods in Enzymology, 2015, 565, 495-535.	1.0	8
25	Complete Genome Sequence and Genome Analysis of <i>Eggplant mottled dwarf virus</i> â€ŀranian Isolate. Journal of Phytopathology, 2015, 163, 331-341.	1.0	7
26	3′UTRs of carmoviruses. Virus Research, 2015, 206, 27-36.	2.2	33
27	Rapid evolution of in vivo-selected sequences and structures replacing 20% of a subviral RNA. Virology, 2015, 483, 149-162.	2.4	7
28	Requirement for Host RNA-Silencing Components and the Virus-Silencing Suppressor when Second-Site Mutations Compensate for Structural Defects in the 3′ Untranslated Region. Journal of Virology, 2015, 89, 11603-11618.	3.4	6
29	Trajectories of the ribosome as a Brownian nanomachine. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 17492-17497.	7.1	218
30	The 3′ Untranslated Region of Pea Enation Mosaic Virus Contains Two T-Shaped, Ribosome-Binding, Cap-Independent Translation Enhancers. Journal of Virology, 2014, 88, 11696-11712.	3.4	43
31	Position of the kissing-loop interaction associated with PTE-type 3′CITEs can affect enhancement of cap-independent translation. Virology, 2014, 458-459, 43-52.	2.4	11
32	3′ Cap-Independent Translation Enhancers of Plant Viruses. Annual Review of Microbiology, 2013, 67, 21-42.	7.3	176
33	The Kissing-Loop T-Shaped Structure Translational Enhancer of Pea Enation Mosaic Virus Can Bind Simultaneously to Ribosomes and a 5′ Proximal Hairpin. Journal of Virology, 2013, 87, 11987-12002.	3.4	40
34	Preparation of biologically active Arabidopsis ribosomes and comparison with yeast ribosomes for binding to a tRNA-mimic that enhances translation of plant plus-strand RNA viruses. Frontiers in Plant Science, 2013, 4, 271.	3.6	6
35	A Local, Interactive Network of 3′ RNA Elements Supports Translation and Replication of Turnip Crinkle Virus. Journal of Virology, 2012, 86, 4065-4081.	3.4	23
36	A Ribosome-Binding, 3′ Translational Enhancer Has a T-Shaped Structure and Engages in a Long-Distance RNA-RNA Interaction. Journal of Virology, 2012, 86, 9828-9842.	3.4	60

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37	Building bridges between plant and animal viruses. Current Opinion in Virology, 2011, 1, 319-321.	5.4	2
38	Long-distance kissing loop interactions between a 3′ proximal Y-shaped structure and apical loops of 5′ hairpins enhance translation of Saguaro cactus virus. Virology, 2011, 417, 113-125.	2.4	45
39	Evolution of a helper virus-derived, ribosome binding translational enhancer in an untranslated satellite RNA of Turnip crinkle virus. Virology, 2011, 419, 10-16.	2.4	5
40	Ribosome Binding to a 5′ Translational Enhancer Is Altered in the Presence of the 3′ Untranslated Region in Cap-Independent Translation of Turnip Crinkle Virus. Journal of Virology, 2011, 85, 4638-4653.	3.4	62
41	Carmovirus. , 2011, , 1885-1894.		0
42	EMBO World Lecture Course â€~Virus–Host: Partners in Pathogenesis'. Future Virology, 2010, 5, 379-383.	1.8	0
43	The terminal loop of a 3′ proximal hairpin plays a critical role in replication and the structure of the 3′ region of Turnip crinkle virus. Virology, 2010, 402, 271-280.	2.4	21
44	Solution structure of the cap-independent translational enhancer and ribosome-binding element in the 3 [′] UTR of turnip crinkle virus. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 1385-1390.	7.1	89
45	The Capsid Protein of Turnip Crinkle Virus Overcomes Two Separate Defense Barriers To Facilitate Systemic Movement of the Virus in Arabidopsis. Journal of Virology, 2010, 84, 7793-7802.	3.4	74
46	Structural Plasticity and Rapid Evolution in a Viral RNA Revealed by In Vivo Genetic Selection. Journal of Virology, 2009, 83, 927-939.	3.4	21
47	The 3′ end of Turnip crinkle virus contains a highly interactive structure including a translational enhancer that is disrupted by binding to the RNA-dependent RNA polymerase. Rna, 2009, 15, 1849-1864.	3.5	42
48	RNA conformational changes in the life cycles of RNA viruses, viroids, and virus-associated RNAs. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2009, 1789, 571-583.	1.9	47
49	Importance of coat protein and RNA silencing in satellite RNA/virus interactions. Virology, 2008, 379, 161-167.	2.4	20
50	Structural Domains within the 3′ Untranslated Region of Turnip Crinkle Virus. Journal of Virology, 2008, 82, 8706-8720.	3.4	69
51	The 3′ proximal translational enhancer of Turnip crinkle virus binds to 60S ribosomal subunits. Rna, 2008, 14, 2379-2393.	3.5	92
52	Callus Cultures ofArabidopsis. , 2006, Chapter 16, 16D.1.1-16D.1.9.		10
53	Evolution of virus-derived sequences for high-level replication of a subviral RNA. Virology, 2006, 351, 476-488.	2.4	24
54	A cis-replication element functions in both orientations to enhance replication of Turnip crinkle virus. Virology, 2006, 352, 39-51.	2.4	35

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55	Conformational changes involved in initiation of minus-strand synthesis of a virus-associated RNA. Rna, 2006, 12, 147-162.	3.5	35
56	A Pseudoknot in a Preactive Form of a Viral RNA Is Part of a Structural Switch Activating Minus-Strand Synthesis. Journal of Virology, 2006, 80, 9181-9191.	3.4	43
57	Importance of sequence and structural elements within a viral replication repressor. Virology, 2005, 333, 301-315.	2.4	28
58	Short Internal Sequences Involved in Replication and Virion Accumulation in a Subviral RNA of Turnip Crinkle Virus. Journal of Virology, 2005, 79, 512-524.	3.4	18
59	Repression and Derepression of Minus-Strand Synthesis in a Plus-Strand RNA Virus Replicon. Journal of Virology, 2004, 78, 7619-7633.	3.4	61
60	Biased Hypermutagenesis Associated with Mutations in an Untranslated Hairpin of an RNA Virus. Journal of Virology, 2004, 78, 7813-7817.	3.4	20
61	Analysis of a viral replication repressor: sequence requirements for a large symmetrical internal loop. Virology, 2004, 326, 90-102.	2.4	23
62	PLANT VIRUS SATELLITE AND DEFECTIVE INTERFERING RNAS: New Paradigms for a New Century. Annual Review of Phytopathology, 2004, 42, 415-437.	7.8	209
63	Enhanced viral pathogenesis associated with a virulent mutant virus or a virulent satellite RNA correlates with reduced virion accumulation and abundance of free coat protein. Virology, 2003, 312, 8-13.	2.4	47
64	A novel procedure for the localization of viral RNAs in protoplasts and whole plants. Plant Journal, 2003, 35, 665-673.	5.7	27
65	A Multifunctional Turnip Crinkle Virus Replication Enhancer Revealed by in vivo Functional SELEX. Journal of Molecular Biology, 2003, 326, 35-48.	4.2	31
66	Fitness of a Turnip Crinkle Virus Satellite RNA Correlates with a Sequence-Nonspecific Hairpin and Flanking Sequences That Enhance Replication and Repress the Accumulation of Virions. Journal of Virology, 2003, 77, 7880-7889.	3.4	17
67	In Vivo and in Vitro Characterization of an RNA Replication Enhancer in a Satellite RNA Associated with Turnip crinkle virus. Virology, 2001, 288, 315-324.	2.4	32
68	Analysis of cis-Acting Sequences Involved in Plus-Strand Synthesis of a Turnip Crinkle Virus-Associated Satellite RNA Identifies a New Carmovirus Replication Element. Virology, 2000, 268, 345-354.	2.4	39
69	Requirement of a 5′-Proximal Linear Sequence on Minus Strands for Plus-Strand Synthesis of a Satellite RNA Associated with Turnip Crinkle Virus. Virology, 2000, 268, 355-363.	2.4	23
70	3′-End Stem-Loops of the Subviral RNAs Associated with Turnip Crinkle Virus Are Involved in Symptom Modulation and Coat Protein Binding. Journal of Virology, 2000, 74, 6528-6537.	3.4	14
71	CCA initiation boxes without unique promoter elements support in vitro transcription by three viral RNA-dependent RNA polymerases. Rna, 2000, 6, 698-707.	3.5	35
72	Polymerization of nontemplate bases before transcription initiation at the 3' ends of templates by an RNA-dependent RNA polymerase: An activity involved in 3' end repair of viral RNAs. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 12451-12456.	7.1	50

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73	Symptom Attenuation by a Satellite RNA in Vivo Is Dependent on Reduced Levels of Virus Coat Protein. Virology, 1999, 259, 234-245.	2.4	31
74	Minimal Sequence and Structural Requirements of a Subgenomic RNA Promoter for Turnip Crinkle Virus. Virology, 1999, 253, 327-336.	2.4	40
75	RNA elements required for RNA recombination function as replication enhancers in vitro and in vivo in a plus-strand RNA virus. EMBO Journal, 1999, 18, 5653-5665.	7.8	93
76	In VitroCharacterization of Late Steps of RNA Recombination in Turnip Crinkle Virus. Virology, 1998, 249, 379-392.	2.4	35
77	In VitroCharacterization of Late Steps of RNA Recombination in Turnip Crinkle Virus. Virology, 1998, 249, 393-405.	2.4	25
78	Dissecting RNA recombination invitro: role of RNA sequences and the viral replicase. EMBO Journal, 1998, 17, 2392-2403.	7.8	100
79	Analysis of sequences and predicted structures required for viral satellite RNA accumulation by in vivo genetic selection. Nucleic Acids Research, 1998, 26, 2426-2432.	14.5	58
80	Satellite RNA-Mediated Resistance to Turnip Crinkle Virus in Arabidopsis Involves a Reduction in Virus Movement. Plant Cell, 1997, 9, 2051.	6.6	11
81	A novel 3'-end repair mechanism in an RNA virus. Proceedings of the National Academy of Sciences of the United States of America, 1997, 94, 1113-1118.	7.1	100
82	Analysis of the Two Subgenomic RNA Promoters for Turnip Crinkle Virusin Vivoandin Vitro. Virology, 1997, 232, 174-186.	2.4	78
83	New Insights into the Mechanisms of RNA Recombination. Virology, 1997, 235, 1-9.	2.4	357
84	Analysisin Vivoof Turnip Crinkle Virus Satellite RNA C Variants with Mutations in the 3′-Terminal Minus-Strand Promoter. Virology, 1997, 238, 470-477.	2.4	47
85	The Coat Protein of Turnip Crinkle Virus Is Involved in Subviral RNA-Mediated Symptom Modulation and Accumulation. Virology, 1997, 238, 478-485.	2.4	29
86	Studies on RNA Recombination in vivo and in vitro. , 1997, , 33-39.		0
87	RNA recombination in turnip crinkle virus: its role in formation of chimeric RNAs, multimers, and in 3′-end repair. Seminars in Virology, 1996, 7, 373-379.	3.9	24
88	Changes in Locations of Crossover Sites over Time inde NovoGenerated RNA Recombinants. Virology, 1996, 223, 165-173.	2.4	10
89	In VivoRepair of 3′-End Deletions in a TCV Satellite RNA May Involve Two Abortive Synthesis and Priming Events. Virology, 1996, 226, 153-160.	2.4	27
90	Symptom Attenuation by a Normally Virulent Satellite RNA of Turnip Crinkle Virus Is Associated with the Coat Protein Open Reading Frame. Plant Cell, 1995, 7, 1625.	6.6	9

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91	???. Journal of Molecular Biology, 1995, 245, 608-622.	4.2	90
92	Requirement of a 3′-Terminal Stem-loop inin VitroTranscription by an RNA-dependent RNA Polymerase. Journal of Molecular Biology, 1995, 254, 6-14.	4.2	126
93	Synthesis of novel products in vitro by an RNA-dependent RNA polymerase. Journal of Virology, 1995, 69, 4020-4028.	3.4	25
94	RNA-RNA Recombination and Evolution in Virus-Infected Plants. Annual Review of Phytopathology, 1994, 32, 337-362.	7.8	153
95	Recombination between Plus and Minus Strands of Turnip Crinkle Virus. Virology, 1994, 201, 419-423.	2.4	17
96	RNA-dependent RNA polymerase from plants infected with turnip crinkle virus can transcribe (+)- and (-)-strands of virus-associated RNAs Proceedings of the National Academy of Sciences of the United States of America, 1994, 91, 8792-8796.	7.1	83
97	Sequences and structures required for recombination between virus-associated RNAs. Science, 1993, 260, 801-805.	12.6	128
98	Susceptibility and Resistance of <i>Arabidopsis thaliana</i> to Turnip Crinkle Virus. Molecular Plant-Microbe Interactions, 1992, 5, 496.	2.6	66
99	Effects of Defective Interfering Viruses on Virus Replication and Pathogenesis In Vitro and In Vivo. Advances in Virus Research, 1991, 40, 181-211.	2.1	225
100	Recombination between satellite and genomic RNAs of turnip crinkle virus. Virology, 1991, 184, 791-794.	2.4	49
101	Formation of multimers of linear satellite RNAs. Virology, 1991, 183, 586-594.	2.4	40
102	Mutations in a satellite RNA of turnip crinkle virus result in addition of poly(U) in vivo. Virology, 1991, 183, 595-601.	2.4	13
103	In vivo accumulation of a turnip crinkle virus defective interfering RNA is affected by alterations in size and sequence. Journal of Virology, 1991, 65, 4582-4590.	3.4	42
104	Recombination between satellite RNAs of turnip crinkle virus EMBO Journal, 1990, 9, 1709-1715.	7.8	122
105	Symptom Intensification on Cruciferous Hosts by the Virulent Satellite RNA of Turnip Crinkle Virus. Phytopathology, 1990, 80, 238.	2.2	50
106	Turnip crinkle virus defective interfering RNAs intensify viral symptoms and are generated de novo Proceedings of the National Academy of Sciences of the United States of America, 1989, 86, 9173-9177.	7.1	125
107	Satellite RNAs of plant viruses. Plant Molecular Biology Reporter, 1988, 6, 240-252.	1.8	24
108	Synthesis in vitro of infectious RNA copies of the virulent satellite of turnip crinkle virus. Virology, 1987, 156, 146-152.	2.4	41

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109	The virulent satellite RNA of turnip crinkle virus has a major domain homologous to the 3′ end of the helper virus genome. EMBO Journal, 1986, 5, 3423-3428.	7.8	114
110	Nucleotide sequence of a cDNA clone of Brassica napus 12S storage protein shows homology with legumin from Pisum sativum. Plant Molecular Biology, 1985, 5, 191-201.	3.9	102
111	High-frequency mutation at the adenine phosphoribosyltransferase locus in Chinese hamster ovary cells due to deletion of the gene Proceedings of the National Academy of Sciences of the United States of America, 1983, 80, 810-814.	7.1	50
112	Two new umbravirus-like associated RNAs (ulaRNAs) discovered in maize and johnsongrass from Ecuador. Archives of Virology, 0, , .	2.1	4