

Naomi Attar

List of Publications by Year in descending order

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Version: 2024-02-01

58
papers

1,031
citations

1162367

8
h-index

433756

31
g-index

95
all docs

95
docs citations

95
times ranked

1957
citing authors

#	ARTICLE	IF	CITATIONS
1	The Retromer Coat Complex Coordinates Endosomal Sorting and Dynein-Mediated Transport, with Carrier Recognition by the trans-Golgi Network. <i>Developmental Cell</i> , 2009, 17, 110-122.	3.1	252
2	SNX4 coordinates endosomal sorting of TfnR with dynein-mediated transport into the endocytic recycling compartment. <i>Nature Cell Biology</i> , 2007, 9, 1370-1380.	4.6	233
3	A loss-of-function screen reveals SNX5 and SNX6 as potential components of the mammalian retromer. <i>Journal of Cell Science</i> , 2007, 120, 45-54.	1.2	210
4	Sorting nexin-1 defines an early phase of <i>Salmonella</i> -containing vacuole-remodeling during <i>Salmonella</i> infection. <i>Journal of Cell Science</i> , 2008, 121, 2027-2036.	1.2	92
5	The retromer complex. <i>Advances in Enzyme Regulation</i> , 2010, 50, 216-236.	2.9	76
6	ZIKA virus circulates in new regions. <i>Nature Reviews Microbiology</i> , 2016, 14, 62-62.	13.6	52
7	A heterodimeric SNX4:SNX7 SNX-BAR autophagy complex coordinates ATG9A trafficking for efficient autophagosome assembly. <i>Journal of Cell Science</i> , 2020, 133, .	1.2	19
8	The allure of the epigenome. <i>Genome Biology</i> , 2012, 13, 419.	13.9	8
9	Raymond Gosling: the man who crystallized genes. <i>Genome Biology</i> , 2013, 14, 402.	13.9	6
10	Gut bugs in the lung link sepsis to ARDS. <i>Nature Reviews Microbiology</i> , 2016, 14, 546-546.	13.6	6
11	Have exploding cells blown up MV dogma?. <i>Nature Reviews Microbiology</i> , 2016, 14, 334-335.	13.6	5
12	Transgenerational missing taxa. <i>Nature Reviews Microbiology</i> , 2016, 14, 132-133.	13.6	5
13	The RBPome: where the brains meet the brawn. <i>Genome Biology</i> , 2014, 15, 402.	13.9	4
14	FISHing in the oral microbiota. <i>Nature Reviews Microbiology</i> , 2016, 14, 133-133.	13.6	4
15	How CRISPR captures spacer invaders. <i>Nature Reviews Microbiology</i> , 2015, 13, 738-739.	13.6	3
16	CPR breathes new air into the tree of life. <i>Nature Reviews Microbiology</i> , 2016, 14, 332-332.	13.6	3
17	Animal viruses in pieces. <i>Nature Reviews Microbiology</i> , 2016, 14, 607-607.	13.6	3
18	Strain competition keeps a lid on gut pathogens. <i>Nature Reviews Microbiology</i> , 2016, 14, 546-546.	13.6	3

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19	SMRT-seq reveals an epigenetic switch. Nature Reviews Microbiology, 2016, 14, 546-546.	13.6	3
20	Dual-core centrosomes power cell division. Nature Reviews Microbiology, 2015, 13, 252-252.	13.6	2
21	Phages' box of tricks for CRISPR. Nature Reviews Microbiology, 2015, 13, 660-661.	13.6	2
22	Archaeal virus escapology. Nature Reviews Microbiology, 2016, 14, 665-665.	13.6	2
23	De-chaperoning antivirals. Nature Reviews Microbiology, 2016, 14, 2-3.	13.6	2
24	Keeping a watchful eye on Ebola. Nature Reviews Microbiology, 2015, 13, 457-457.	13.6	1
25	Keeping a watchful eye on Ebola. Nature Reviews Genetics, 2015, 16, 437-437.	7.7	1
26	An interkingdom partnership. Nature Reviews Microbiology, 2015, 13, 400-400.	13.6	1
27	Tracking down HIV's hiding place. Nature Reviews Microbiology, 2015, 13, 187-187.	13.6	1
28	Phages' box of tricks for CRISPR. Nature Reviews Genetics, 2015, 16, 626-626.	7.7	1
29	The history of Lassa virus. Nature Reviews Microbiology, 2015, 13, 600-601.	13.6	1
30	Spore formation in the human gut microbiota. Nature Reviews Microbiology, 2016, 14, 403-403.	13.6	1
31	A new chaperone for regulatory sRNAs. Nature Reviews Microbiology, 2016, 14, 665-665.	13.6	1
32	Trypanosomes chew the fat. Nature Reviews Microbiology, 2016, 14, 403-403.	13.6	1
33	It's never too late for CRISPR RNases. Nature Reviews Microbiology, 2016, 14, 192-192.	13.6	1
34	Pushing the envelope on protein repair. Nature Reviews Microbiology, 2016, 14, 65-65.	13.6	1
35	How hemp got high. Genome Biology, 2011, 12, 409.	13.9	0
36	BioMed Central - open data guidance for authors. Nature Precedings, 2011, , .	0.1	0

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37	The Research Works Act: a comment. <i>Genome Biology</i> , 2012, 13, 416.	13.9	0
38	Cutting out the carBs feeds a bistable switch. <i>Nature Reviews Microbiology</i> , 2015, 13, 250-251.	13.6	0
39	Copying the bunyavirus genome. <i>Nature Reviews Microbiology</i> , 2015, 13, 399-399.	13.6	0
40	PINning down Theileria. <i>Nature Reviews Microbiology</i> , 2015, 13, 127-127.	13.6	0
41	The antisense antigen switch. <i>Nature Reviews Microbiology</i> , 2015, 13, 189-189.	13.6	0
42	Oscillations relieve the siege. <i>Nature Reviews Microbiology</i> , 2015, 13, 526-526.	13.6	0
43	Sifting out virulent bacteria. <i>Nature Reviews Microbiology</i> , 2015, 13, 330-330.	13.6	0
44	A CRISPR sense of self. <i>Nature Reviews Microbiology</i> , 2015, 13, 329-329.	13.6	0
45	Pick 'n' mix toxins. <i>Nature Reviews Microbiology</i> , 2015, 13, 601-601.	13.6	0
46	Mycobacteria that are long in the tooth. <i>Nature Reviews Microbiology</i> , 2016, 14, 475-475.	13.6	0
47	Why you're stuck with the skin you're in. <i>Nature Reviews Microbiology</i> , 2016, 14, 403-403.	13.6	0
48	ECT joins the rotary club. <i>Nature Reviews Microbiology</i> , 2016, 14, 265-265.	13.6	0
49	Autophagy genes link OMVs to IBD. <i>Nature Reviews Microbiology</i> , 2016, 14, 403-403.	13.6	0
50	MshEN: possible for c-di-GMP binding. <i>Nature Reviews Microbiology</i> , 2016, 14, 605-605.	13.6	0
51	3â€² UTRs: a paradigm for archaeal gene regulation?. <i>Nature Reviews Microbiology</i> , 2016, 14, 605-605.	13.6	0
52	A gateway protein for norovirus. <i>Nature Reviews Microbiology</i> , 2016, 14, 605-605.	13.6	0
53	A new class of Hfq-like sRNA chaperones?. <i>Nature Reviews Microbiology</i> , 2016, 14, 546-546.	13.6	0
54	How pathogens divide. <i>Nature Reviews Microbiology</i> , 2016, 14, 605-605.	13.6	0

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55	Microbial mobilomes differ between societies. Nature Reviews Microbiology, 2016, 14, 546-546.	13.6	0
56	A tale of two specificities. Nature Reviews Microbiology, 2016, 14, 405-405.	13.6	0
57	An antidote to persistence. Nature Reviews Microbiology, 2016, 14, 473-473.	13.6	0
58	One giant leap for Cas-kind. Nature Reviews Microbiology, 2016, 14, 194-194.	13.6	0