

Seon Young Choi

List of Publications by Year in descending order

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16
papers

1,935
citations

623734

14
h-index

940533

16
g-index

16
all docs

16
docs citations

16
times ranked

2221
citing authors

#	ARTICLE	IF	CITATIONS
1	Evidence for several waves of global transmission in the seventh cholera pandemic. <i>Nature</i> , 2011, 477, 462-465.	27.8	649
2	Comparative genomics reveals mechanism for short-term and long-term clonal transitions in pandemic <i>Vibrio cholerae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 15442-15447.	7.1	351
3	<i>Shigella sonnei</i> genome sequencing and phylogenetic analysis indicate recent global dissemination from Europe. <i>Nature Genetics</i> , 2012, 44, 1056-1059.	21.4	278
4	Genomic diversity of 2010 Haitian cholera outbreak strains. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, E2010-7.	7.1	173
5	Cholera Outbreaks Caused by an Altered <i>Vibrio cholerae</i> O1 El Tor Biotype Strain Producing Classical Cholera Toxin B in Vietnam in 2007 to 2008. <i>Journal of Clinical Microbiology</i> , 2009, 47, 1568-1571.	3.9	104
6	Multilocus sequence typing (MLST) analysis of <i>Vibrio cholerae</i> O1 El Tor isolates from Mozambique that harbour the classical CTX prophage. <i>Journal of Medical Microbiology</i> , 2006, 55, 165-170.	1.8	74
7	Deep-sea hydrothermal vent bacteria related to human pathogenic <i>Vibrio</i> species. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E2813-9.	7.1	63
8	Genomic and Phenotypic Characterization of <i>Vibrio cholerae</i> Non-O1 Isolates from a US Gulf Coast Cholera Outbreak. <i>PLoS ONE</i> , 2014, 9, e86264.	2.5	54
9	Multilocus variable-number tandem repeat analysis of <i>Vibrio cholerae</i> O1 El Tor strains harbouring classical toxin B. <i>Journal of Medical Microbiology</i> , 2010, 59, 763-769.	1.8	43
10	Comparative genomic analysis and characterization of incompatibility group FIB plasmid encoded virulence factors of <i>Salmonella enterica</i> isolated from food sources. <i>BMC Genomics</i> , 2017, 18, 570.	2.8	35
11	Classification of hybrid and altered <i>Vibrio cholerae</i> strains by CTX prophage and RS1 element structure. <i>Journal of Microbiology</i> , 2009, 47, 783-788.	2.8	27
12	Nontoxigenic <i>Vibrio cholerae</i> Non-O1/O139 Isolate from a Case of Human Gastroenteritis in the U.S. Gulf Coast. <i>Journal of Clinical Microbiology</i> , 2015, 53, 9-14.	3.9	25
13	Phylogenetic Diversity of <i>Vibrio cholerae</i> Associated with Endemic Cholera in Mexico from 1991 to 2008. <i>MBio</i> , 2016, 7, e02160.	4.1	24
14	Classical RS1 and environmental RS1 elements in <i>Vibrio cholerae</i> O1 El Tor strains harbouring a tandem repeat of CTX prophage: revisiting Mozambique in 2005. <i>Journal of Medical Microbiology</i> , 2010, 59, 302-308.	1.8	19
15	Hybrid <i>Vibrio cholerae</i> El Tor Lacking SXT Identified as the Cause of a Cholera Outbreak in the Philippines. <i>MBio</i> , 2015, 6, .	4.1	11
16	Genome Sequences of Clinical <i>Vibrio cholerae</i> Isolates from an Oyster-Borne Cholera Outbreak in Florida. <i>Genome Announcements</i> , 2013, 1, .	0.8	5