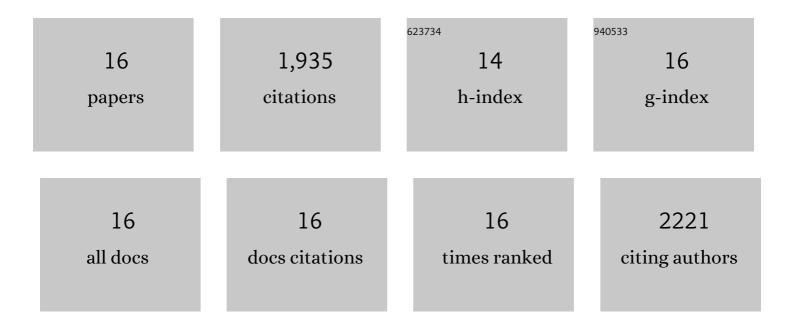
Seon Young Choi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/12177230/publications.pdf Version: 2024-02-01



SEON YOUNG CHOL

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