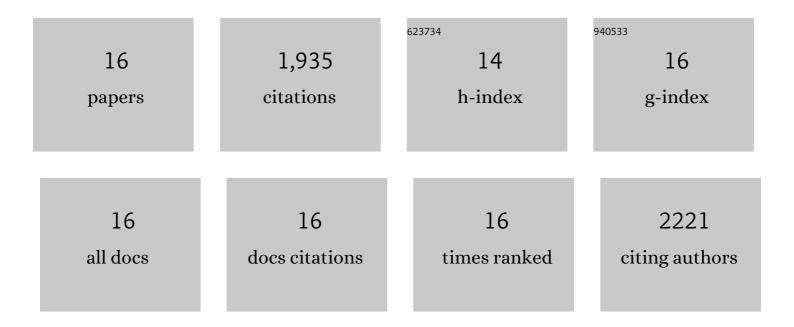
## Seon Young Choi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Evidence for several waves of global transmission in the seventh cholera pandemic. Nature, 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> . Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 15442-15447.	7.1	351
3	Shigella sonnei genome sequencing and phylogenetic analysis indicate recent global dissemination from Europe. Nature Genetics, 2012, 44, 1056-1059.	21.4	278
4	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
5	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
6	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
7	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
8	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
9	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
10	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
11	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
12	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
13	Phylogenetic Diversity of Vibrio cholerae Associated with Endemic Cholera in Mexico from 1991 to 2008. MBio, 2016, 7, e02160.	4.1	24
14	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
15	Hybrid Vibrio cholerae El Tor Lacking SXT Identified as the Cause of a Cholera Outbreak in the Philippines. MBio, 2015, 6, .	4.1	11
16	Genome Sequences of Clinical Vibrio cholerae Isolates from an Oyster-Borne Cholera Outbreak in Florida. Genome Announcements, 2013, 1, .	0.8	5