

# Age of the Association between *Helicobacter pylori* and

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Multilocus Sequence Typing as a Replacement for Serotyping in <i>Salmonella enterica</i> . <i>PLoS Pathogens</i> , 2012, 8, e1002776.	2.1	574
2	<i>In Vivo</i> Sequence Variation in HopZ, a Phase-Variable Outer Membrane Protein of <i>Helicobacter pylori</i> . <i>Infection and Immunity</i> , 2012, 80, 4364-4373.	1.0	41
3	Genomic Variation in Seven Khoe-San Groups Reveals Adaptation and Complex African History. <i>Science</i> , 2012, 338, 374-379.	6.0	364
4	Survival in hostile territory: the microbiota of the stomach. <i>FEMS Microbiology Reviews</i> , 2013, 37, 736-761.	3.9	126
5	Mitochondrial diversity in human head louse populations across the Americas. <i>American Journal of Physical Anthropology</i> , 2013, 152, 118-129.	2.1	22
6	Diagnosis and Epidemiology of <i>Helicobacter pylori</i> Infection. <i>Helicobacter</i> , 2013, 18, 5-11.	1.6	114
7	Genomic evolution and transmission of <i>Helicobacter pylori</i> in two South African families. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 13880-13885.	3.3	115
8	Modulation of autophagy by <i>Helicobacter pylori</i> and its role in gastric carcinogenesis. <i>Trends in Microbiology</i> , 2013, 21, 602-612.	3.5	86
9	Phylogeographic evidence of cognate recognition site patterns and transformation efficiency differences in <i>H. pylori</i> : theory of strain dominance. <i>BMC Microbiology</i> , 2013, 13, 211.	1.3	11
10	Genome Sequences of Three hpAfrica2 Strains of <i>Helicobacter pylori</i> . <i>Genome Announcements</i> , 2013, 1, .	0.8	11
11	Recent Acquisition of <i>Helicobacter pylori</i> by Baka Pygmies. <i>PLoS Genetics</i> , 2013, 9, e1003775.	1.5	35
12	Bridging disciplines to better elucidate the evolution of early <i>Homo sapiens</i> in southern Africa. <i>South African Journal of Science</i> , 2013, 109, 8.	0.3	15
13	Chromosome Painting In Silico in a Bacterial Species Reveals Fine Population Structure. <i>Molecular Biology and Evolution</i> , 2013, 30, 1454-1464.	3.5	87
14	Gastric and Enterohepatic Non- <i>Helicobacter pylori</i> Helicobacters. <i>Helicobacter</i> , 2013, 18, 66-72.	1.6	28
15	Echoes of a Distant Past: The <i>cag</i> Pathogenicity Island of <i>Helicobacter pylori</i> . <i>Cold Spring Harbor Perspectives in Medicine</i> , 2013, 3, a010355-a010355.	2.9	12
16	Comparative Genomic Analysis of East Asian and Non-Asian <i>Helicobacter pylori</i> Strains Identifies Rapidly Evolving Genes. <i>PLoS ONE</i> , 2013, 8, e55120.	1.1	27
17	Population Genetic Structure and Isolation by Distance of <i>Helicobacter pylori</i> in Senegal and Madagascar. <i>PLoS ONE</i> , 2014, 9, e87355.	1.1	17
18	<i>Streptococcus mitis</i> Induces Conversion of <i>Helicobacter pylori</i> to Coccoid Cells during Co-Culture In Vitro. <i>PLoS ONE</i> , 2014, 9, e112214.	1.1	27

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20	<i>Helicobacter pylori</i> DNA methyltransferases and the epigenetic field effect in cancerization. <i>Frontiers in Microbiology</i> , 2014, 5, 115.	1.5	24
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38	Dormant phages of <i>Helicobacter pylori</i> reveal distinct populations in Europe. <i>Scientific Reports</i> , 2015, 5, 14333.	1.6	37

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