

Sequence Analysis of the p1 Adhesin Gene of Mycoplasma Collected in Beijing in 2008 to 2009

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Sequence Variation within the P1 Gene of <i>Mycoplasma pneumoniae</i> . <i>Journal of Clinical Microbiology</i> , 2011, 49, 3723-3724.	1.8	4
2	Complete Genome Sequence of <i>Mycoplasma pneumoniae</i> Type 2a Strain 309, Isolated in Japan. <i>Journal of Bacteriology</i> , 2012, 194, 1253-1254.	1.0	24
3	Antibiotic Sensitivity of 40 <i>Mycoplasma pneumoniae</i> Isolates and Molecular Analysis of Macrolide-Resistant Isolates from Beijing, China. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 1108-1109.	1.4	55
4	Surveillance of Macrolide-Resistant <i>Mycoplasma pneumoniae</i> in Beijing, China, from 2008 to 2012. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 1521-1523.	1.4	116
5	Multiple-Locus Variable-Number Tandem-Repeat Analysis of 201 <i>Mycoplasma pneumoniae</i> Isolates from Beijing, China, from 2008 to 2011. <i>Journal of Clinical Microbiology</i> , 2013, 51, 636-639.	1.8	35
6	Detection of <i>Mycoplasma pneumoniae</i> by colorimetric loop-mediated isothermal amplification. <i>Acta Microbiologica Et Immunologica Hungarica</i> , 2013, 60, 1-9.	0.4	6
7	Multiple-Locus Variable-Number Tandem-Repeat Analysis of <i>Mycoplasma pneumoniae</i> Clinical Specimens and Proposal for Amendment of MLVA Nomenclature. <i>PLoS ONE</i> , 2013, 8, e64607.	1.1	51
8	Molecular Characterizations of PCR-Positive <i>Mycoplasma pneumoniae</i> Specimens Collected from Australia and China. <i>Journal of Clinical Microbiology</i> , 2014, 52, 1478-1482.	1.8	36
9	Novel Strategy for Typing <i>Mycoplasma pneumoniae</i> Isolates by Use of Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry Coupled with ClinProTools. <i>Journal of Clinical Microbiology</i> , 2014, 52, 3038-3043.	1.8	28
10	Extensive Variation and Rapid Shift of the MG192 Sequence in <i>Mycoplasma genitalium</i> Strains from Patients with Chronic Infection. <i>Infection and Immunity</i> , 2014, 82, 1326-1334.	1.0	21
11	Detection of <i>Mycoplasma pneumoniae</i> P1 subtype variations by denaturing gradient gel electrophoresis. <i>Diagnostic Microbiology and Infectious Disease</i> , 2014, 78, 24-28.	0.8	21
13	Molecular Typing of <i>Mycoplasma pneumoniae</i> Isolated from Pediatric Patients in Tokyo, Japan. <i>Japanese Journal of Infectious Diseases</i> , 2015, 68, 76-78.	0.5	8
14	Specificity and Strain-Typing Capabilities of Nanorod Array-Surface Enhanced Raman Spectroscopy for <i>Mycoplasma pneumoniae</i> Detection. <i>PLoS ONE</i> , 2015, 10, e0131831.	1.1	19
15	Comparative genome analysis of <i>Mycoplasma pneumoniae</i> . <i>BMC Genomics</i> , 2015, 16, 610.	1.2	59
16	<i>Mycoplasma pneumoniae</i> and <i>Chlamydia</i> spp. Infection in Community-Acquired Pneumonia, Germany, 2011–2012. <i>Emerging Infectious Diseases</i> , 2015, 21, 426-434.	2.0	99
17	<i>Mycoplasma pneumoniae</i> Monoclonal P1 Type 2c Outbreak, Russia, 2013. <i>Emerging Infectious Diseases</i> , 2016, 22, 348-350.	2.0	14
18	The Evolution of Advanced Molecular Diagnostics for the Detection and Characterization of <i>Mycoplasma pneumoniae</i> . <i>Frontiers in Microbiology</i> , 2016, 7, 232.	1.5	40
19	Epidemiology of <i>Mycoplasma pneumoniae</i> Infections in Japan and Therapeutic Strategies for Macrolide-Resistant <i>M. pneumoniae</i> . <i>Frontiers in Microbiology</i> , 2016, 7, 693.	1.5	90

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20	<i>Mycoplasma pneumoniae</i> : Current Knowledge on Macrolide Resistance and Treatment. <i>Frontiers in Microbiology</i> , 2016, 7, 974.	1.5	180
21	<i>Mycoplasma pneumoniae</i> from the Respiratory Tract and Beyond. <i>Clinical Microbiology Reviews</i> , 2017, 30, 747-809.	5.7	411
22	Pediatric clinical features of <i>Mycoplasma pneumoniae</i> infection are associated with bacterial P1 genotype. <i>Experimental and Therapeutic Medicine</i> , 2017, 14, 1892-1898.	0.8	14
23	Complete Genome Sequences of the <i>p1</i> Gene Type 2b and 2c Strains <i>Mycoplasma pneumoniae</i> KCH-402 and KCH-405. <i>Genome Announcements</i> , 2017, 5, .	0.8	1
24	Inter- and intra-strain variability of tandem repeats in <i>Mycoplasma pneumoniae</i> based on next-generation sequencing data. <i>Future Microbiology</i> , 2017, 12, 119-129.	1.0	7
25	Comprehensive bioinformatics analysis of <i>Mycoplasma pneumoniae</i> genomes to investigate underlying population structure and type-specific determinants. <i>PLoS ONE</i> , 2017, 12, e0174701.	1.1	27
26	Multiple-Locus Variable-Number Tandem-Repeat Analysis of <i>Mycoplasma pneumoniae</i> Isolates between 2004 and 2014 in Yamagata, Japan: Change in Molecular Characteristics during an 11-year Period. <i>Japanese Journal of Infectious Diseases</i> , 2017, 70, 642-646.	0.5	18
27	Genotyping and macrolide resistance of <i>Mycoplasma pneumoniae</i> identified in children with community-acquired pneumonia in Medellín, Colombia. <i>International Journal of Infectious Diseases</i> , 2018, 66, 113-120.	1.5	11
28	Clonal Expansion of Macrolide-Resistant Sequence Type 3 <i>Mycoplasma pneumoniae</i> , South Korea. <i>Emerging Infectious Diseases</i> , 2018, 24, 1465-1471.	2.0	26
29	Antimicrobial susceptibility and molecular characteristics of <i>Mycoplasma pneumoniae</i> isolates across different regions of China. <i>Antimicrobial Resistance and Infection Control</i> , 2019, 8, 143.	1.5	36
30	Molecular Typing of <i>Mycoplasma pneumoniae</i> Strains in Sweden from 1996 to 2017 and the Emergence of a New P1 Cytadhesin Gene, Variant 2e. <i>Journal of Clinical Microbiology</i> , 2019, 57, .	1.8	25
31	Comparative genomics of <i>Mycoplasma pneumoniae</i> isolated from children with pneumonia: South Korea, 2010–2016. <i>BMC Genomics</i> , 2019, 20, 910.	1.2	7
32	Periodic Genotype Shifts in Clinically Prevalent <i>Mycoplasma pneumoniae</i> Strains in Japan. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 385.	1.8	20
33	Molecular Characterization of <i>Mycoplasma pneumoniae</i> Isolates in the United States from 2012 to 2018. <i>Journal of Clinical Microbiology</i> , 2020, 58, .	1.8	32
34	<i>Mycoplasma pneumoniae</i> Infections: Pathogenesis and Vaccine Development. <i>Pathogens</i> , 2021, 10, 119.	1.2	42
35	Polyclonal spread of multiple genotypes of <i>Mycoplasma pneumoniae</i> in semi-closed settings in Yamagata, Japan. <i>Journal of Medical Microbiology</i> , 2019, 68, 785-790.	0.7	8
36	Identification of <i>Mycoplasma pneumoniae</i> type 2b variant strains in Japan. <i>Journal of Medical Microbiology</i> , 2012, 61, 1633-1635.	0.7	10
37	Culture-Independent Detection and Genotyping of <i>Mycoplasma pneumoniae</i> in Clinical Specimens from Beijing, China. <i>PLoS ONE</i> , 2015, 10, e0141702.	1.1	36

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38	Prevalence, genotyping and macrolide resistance of <i>Mycoplasma pneumoniae</i> among isolates of patients with respiratory tract infections, Central Slovenia, 2006 to 2014. <i>Eurosurveillance</i> , 2015, 20, .	3.9	39
39	Global Genome Diversity and Recombination in <i>Mycoplasma pneumoniae</i> . <i>Emerging Infectious Diseases</i> , 2022, 28, 111-117.	2.0	4
43	Molecular Tools for Typing <i>Mycoplasma pneumoniae</i> and <i>Mycoplasma genitalium</i> . <i>Frontiers in Microbiology</i> , 2022, 13, .	1.5	7
45	<i>Mycoplasma pneumoniae</i> multilocus variable-number tandem-repeat analysis genotypes are associated with inflammatory biomarker levels in children with lower respiratory tract infections. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 0, , .	1.3	3
46	The Association between <i>Mycoplasma pneumoniae</i> Genotype and Cutaneous Disease. <i>Microorganisms</i> , 2023, 11, 205.	1.6	1