



Characteristics of the 2023-2024 *Mycoplasma pneumoniae* epidemic in adults, Southeast France

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ABSTRACT

Objectives: Analysis of current 2023-2024 *Mycoplasma pneumoniae* (Mp) infection characteristics in adults.

Methods: A retrospective case series analysis was performed on Mp polymerase chain reaction-positive adult patients admitted to the University Hospital of Marseille from April 2017 to June 2024. Clinical presentations, treatments, and outcomes were assessed. We compared the epidemiological and clinical characteristics of Mp infections between 2017 and 2022 with the current epidemic.

Results: Clinical and radiological characteristics and outcomes of patients with Mp infection did not differ significantly between the current epidemic (N = 108) and the 5 previous years (N = 94), except that patients in the current epidemic required less supplemental oxygen (odds ratio [95% confidence interval] = 0.48 [0.29-0.78]) and were less likely to present with fever on admission (odds ratio [95% confidence interval] = 0.22 [0.10-0.47]). In both periods, more than half of the patients hospitalized with Mp infection required supportive oxygen therapy. **Conclusions:** During the current 2023-2024 epidemic, more hospital admissions for Mp infection in adults were observed at the University Hospital of Marseille than in the previous 5 years. The clinical characteristics and outcomes of patients with Mp infection did not differ significantly. In our cohort, Mp infection was often severe, regardless of the study period.

Introduction

Mycoplasma pneumoniae (Mp) is a common cause of respiratory tract infections with community-acquired pneumonia (CAP) being the major disease burden, most commonly in children aged 5 to 15 years old [1]. Droplets containing the microorganism can be spread from person to person by coughing or sneezing. Infections occur sporadically throughout the year in many different climates around the world, with epidemic waves occurring every few years [1]. Previous data suggest an interval of 1-3 years between Mp epidemics in Europe [2]. The reported incidence of sporadic Mp in adults ranges from 4-8% of bacterial CAP rising to 20% and 70% during epidemics [3]. Several factors, including waning herd immunity or the introduction of new subtypes into the population, account for the periodic occurrence

of epidemics. The previous epidemic occurred in late 2019-early 2020 simultaneously in several countries, mainly in Europe and Asia [4].

Global prospective surveillance data show the re-emergence of Mp in Europe and Asia more than 3 years after the introduction of SARS-CoV-2 pandemic restrictions [2,4-7]. The first signs of the recent epidemic were detected in June 2023 and peaked in December 2023 [2]. This increased incidence of Mp was observed mainly in school-aged children and young adults [5,6,8,9]. Edouard *et al.* [10] observed a shift in the population affected by the epidemic, with adults becoming more affected from January 2024 (42% of infected patients vs 21% since the beginning of the outbreak), possibly due to massive transmission of the bacterium from infected children. To date, the burden of Mp infection in adults is poorly understood. In addition, it remains unclear whether

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or not the current epidemic is associated with a higher proportion of severe disease.

The aim of this study is to describe the clinical characteristics of *Mp* infections in adults during the current epidemic and to compare them with those of the previous 5 years.

Materials and methods

Following the STROBE guidelines and checklist, we retrospectively extracted the list of all respiratory specimens tested at the University Hospital of Marseille with one of the following specific quantitative polymerase chain reaction (qPCR) assays for *Mp*: (i) qPCR performed by point-of-care laboratories with a syndromic panel using the Biofire FilmArray Respiratory panel 2 plus assay (Biomérieux, Marcy-l'Étoile, France); (ii) qPCR performed routinely at the core laboratory with a syndromic approach using the FTD Respiratory pathogens 21 assay (Fast Track Diagnosis, Luxembourg) or with an in-house *Mp*-specific qPCR [11]. We did not include those diagnosed by serology because of the lack of immunoglobulin (Ig)M specificity and the long persistence of IgG [10].

The inclusion period started in April 2017 and ended on 31 May 2024. Two periods were defined within this time range: period 1 from 1 April 2017 to 31 March 2023, covering two previous *Mp* outbreaks as well as the beginning of the COVID pandemic, and period 2 from 1 April 2023 to 31 May 2024, covering the current *Mp* epidemic. Only patients over 15 years of age were included. Clinical, biological, and radiological data for each patient were prospectively extracted from our computerized medical record system (DPI Reflex®, aXigate), and pseudonymized.

Study variables included general demographic and medical background data such as age, sex, body mass index, immunosuppression status, and the components of the Charlson comorbidity index. Clinical data at enrollment included vital signs, oxygen flow at enrollment and peak oxygen flow, pulmonary symptoms (cough, sputum), and the presence of extrapulmonary symptoms. Patients were considered to have severe *Mp* if they required oxygen support. Oxygen saturation rate <95% (or 92% in patients with chronic respiratory failure) using pulse oximetry on room air. In addition, we recorded radiological findings, blood test results (erythrocyte, platelet, and leukocyte count, lymphocyte/neutrophil ratio, C-reactive protein level, transaminase, bilirubin, fibrinogen, lactate dehydrogenase, creatinine and urea levels, D-dimer). Finally, outcome data were recorded, including hospitalization, intensive care unit admission, length of hospital stay, and death.

Continuous variables are presented as: mean (standard deviation). Qualitative variables are presented as percentages (n/N). For the univariate analysis, Fisher's exact test was applied to categorical variables, and the Wilcoxon-Mann-Whitney test was used for continuous variables to compare patient groups. To identify the most significant factors associated with the period (1/2), we employed a logistic Bayesian model averaging [12]. Model selection was based on the highest posterior probability and the inclusion of variables in the final model was determined by their posterior inclusion probabilities. Variables with posterior inclusion probabilities greater than 0.5 were considered significant predictors. Initial variable selection was based on univariate analyses, including those with $P < 0.10$ as candidates for Bayesian model averaging. Variables with more than 20% missing data or low frequencies were excluded to avoid biases and instability. For the remaining candidate variables, missing values were imputed using multiple imputation techniques via the mice R package (REF). We assessed the missing at-random assumption through both visual analysis and logistic regression models, examining whether the probability of missingness was associated with observed data. The missing at-random assumption was supported by these analyses. Additionally, we performed sensitivity analyses to evaluate the robustness of our findings under different imputation scenarios, confirming the stability of our results. All statistical analyses were performed using R (R Core Team [2023] Foundation for Statistical Computing, Vienna, Austria. URL <https://www.R-project.org/>).

Results

Between April 2017 and June 2024, 118,427 respiratory specimens from 101,548 patients were tested for *Mp* by qPCR at the University Hospital of Marseille. The incidence of *Mp* infection over time is shown in Figure 1. Overall, 535 patients tested positive, of which 202 were 15 years or older, and had available data in the computerized medical record system (Figure S1). The characteristics of the 202 patients are shown in Table 1. The most common clinical presentation of diagnosed *Mp* infection was CAP. Among the 202 included patients, 81.5% presented with at least a cough, 44.4% had sputum expectoration and 31.7% were febrile on admission. Upper respiratory tract (acute coryza, pharyngitis) and gastrointestinal symptoms (nausea, abdominal pain) were also observed in 14.8% and 15.8% of cases, respectively. Mucocutaneous lesions have also been observed in 5.1% of cases (10 patients) consistent with *Mp*-induced rash and mucositis [13] (Figure 2). Other extrapulmonary symptoms were most commonly headache, malaise, and diffuse myalgia (21.4%). More than half of the patients (50.5%) re-

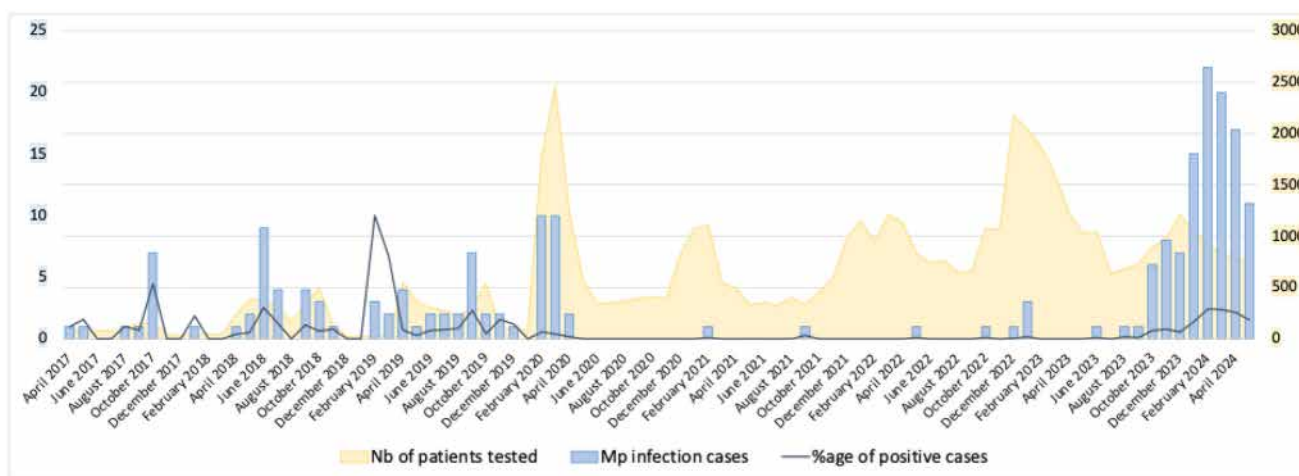


Figure 1. Incidence of *Mp* infection over time from April 2017 to June 2024.

Mp: *Mycoplasma pneumoniae*.

The number of samples tested over time and the percentage of positive cases are also reported.

Table 1
Characteristics of patients with *Mycoplasma pneumoniae* infection. Comparison of periods 1 (April 2017-March 2023) and 2 (April 2023-May 2024).

	All N = 202 ^a	Period 1 N = 94 (46.5%)	Period 2 N=108 (53.5%)	Missing data ^a	P-value ^b
Baseline characteristics					
Age (years)	43.4 (22.2)	44.9 (22.3)	42.2 (22.0)	-	0.416
Sex (female)	37.1% (75/202)	36.2% (34/94)	38.0% (41/108)	-	0.884
Body mass index (kg/m ²)	25.9 (7.3)	26.3 (7.4)	25.6 (7.3)	74 (37 ; 37)	0.989
Tobacco use	44.0% (84/191)	46.7% (42/90)	41.6% (42/101)	11 (4 ; 7)	0.559
Immunodepression	25.6% (51/199)	30.1% (28/93)	21.7% (23/106)	3 (1 ; 2)	0.195
Charlson comorbidity index	1.2 (1.9)	1.4 (2.1)	1.1 (1.7)	-	0.350
Clinical presentation					
Fever (>38°C)	31.7% (64/202)	42.6% (40/94)	22.2% (24/108)	-	0.002
Confusion (Glasgow)	5.9% (12/202)	4.2% (4/94)	7.4% (8/108)	-	0.388
Coma scale <15)					
Oxygen therapy	50.5% (102/202)	43.6% (41/94)	56.5% (61/108)	-	0.090
Heart rate	98.7 (20.3)	94.8 (18.8)	101.9 (21.0)	19 (13 ; 6)	0.012
Cough	81.5% (163/200)	83.7% (77/92)	79.6% (86/108)	2 (2 ; 0)	0.584
Sputum expectorations	44.4% (88/198)	37.4% (34/91)	50.5% (54/107)	4 (3 ; 1)	0.085
Extrapulmonary symptoms	43.1% (87/202)	35.1% (33/94)	50.0% (54/108)	5 (2 ; 3)	0.046
Mucocutaneous	5.1% (10/197)	4.3% (4/92)	5.7% (6/105)	5 (2 ; 3)	0.753
Upper respiratory tract	14.8% (29/196)	13.2% (12/91)	16.2% (17/105)	6 (3 ; 3)	0.687
Digestive	16.8% (33/196)	14.3% (13/91)	19.0% (20/105)	6 (3 ; 3)	0.446
Other	21.4% (42/196)	17.6% (16/91)	24.8% (26/105)	6 (3 ; 3)	0.295
Outcome					
Hospitalization	74.6% (151/202)	78.7% (74/94)	71.3% (77/108)	-	0.258
Intensive care unit admission	18.8% (38/202)	22.3% (21/94)	15.7% (17/108)	-	0.280
Length of hospital stay (days)	8.8 (14.5)	10.1 (19.1)	7.6 (8.5)		0.560
Death	4.5% (9/202)	7.4% (7/94)	1.9% (2/108)	-	0.085
Radiology					
Computed tomography-scan features					
Ground glass opacity	80.6% (83/103)	77.5% (31/40)	82.5% (52/63)	99 (54 ; 45)	0.612
Tree-in-bud	64.2% (61/95)	44.1% (15/34)	75.4% (46/61)	107 (60 ; 47)	0.004
Consolidation	64.6% (73/113)	59.1% (26/44)	68.1% (47/69)	89 (50 ; 39)	0.42
Pleural effusion	15.7% (20/127)	13.2% (7/53)	17.6% (13/74)	75 (41 ; 34)	0.624
Pulmonary embolism	10.3% (7/68)	21.7% (5/23)	4.4% (2/45)	134 (71 ; 63)	0.039
Microbiology					
Immunoglobulin M	13.8 (10.8)	10.8 (10.0)	17.6 (10.6)	122 (50 ; 72)	0.009
<i>Mycoplasma pneumoniae</i> index	0.6-3.3-9.3-27.0-27.0				
Co-infections	25.7% (52/202)	26.6% (25/94)	25.0% (27/108)	-	0.872
Biology					
Leucocytes (G/l)	10.0 (4.3)	10.0 (4.0)	10.0 (4.6)	10 (4 ; 6)	0.941
Lymphocytes (G/l)	0.9-7.0-9.5-13.0-22.6	1.5 (0.9)	1.5 (1.0)	43 (27 ; 16)	0.827
C-reactive protein (mg/l)	94.4 (81.1)	80.9 (80.0)	105.2 (80.8)	18 (12 ; 6)	0.006

For continuous variables: Mean (SD); For qualitative variables: % (n/N).

^a Missing data: Total (N missing for period 1. N missing for period 2).

^b Fisher's exact test for frequencies and Wilcoxon-Mann-Whitney test for continuous variables.

quired supportive oxygen therapy, including 6.4% on high-flow therapy, 4.0% on mechanical ventilation, and 3.5% on extracorporeal membrane oxygenation. Most of the patients (74.6%) were admitted to the hospital with a median length of stay of 5 days. The in-hospital mortality rate was 4.45% (9/202) (period 1, N = 7 and period 2, N = 1).

Clinical characteristics and outcomes of patients with *Mp* infection did not differ between the periods before (N = 94) and after (N = 108) the COVID-19 pandemic, except that patients in the 2023-2024 outbreak required less supplemental oxygen (odds ratio [OR] [95% confidence interval: CI] = 0.48 [0.29-0.78]), were less likely to present with fever on admission (OR [95% CI] = 0.22 [0.10-0.47]), and their heart rate was significantly higher (OR [95% CI] = 1.93 [1.31-2.83]) (Figure S2). Although, during period 1, the proportion of patients with *Mp* infection having undergone a chest computed tomographic scan was significantly higher (OR [95% CI] = 2.84 [1.35-5.97]), the radiological characteristics were broadly similar between the two periods

(Table 1). Co-infections were diagnosed in 25.7% of patients, the most frequent pathogen being rhinovirus, accounting for a quarter of all the co-pathogen detections (Table S1). The rate of co-infection did not differ significantly between both periods.

Discussion

In this study, we described the epidemiological and clinical characteristics of *Mp* infections in adults during the current 2023-2024 epidemic (period 2) and compared them with those of the previous 5 years (period 1), in a French University Hospital. We observed that more cases of *Mp* infection in adults occurred during period 2 (N = 108) than during period 1 (N = 94). The re-emergence of *Mp* could be explained, at least in part, by the waning herd and individual immunity. Knowing that outbreaks of *Mp* infection occur cyclically every 3-5 years, the current epidemic could be the usual periodic recurrence but marked by an

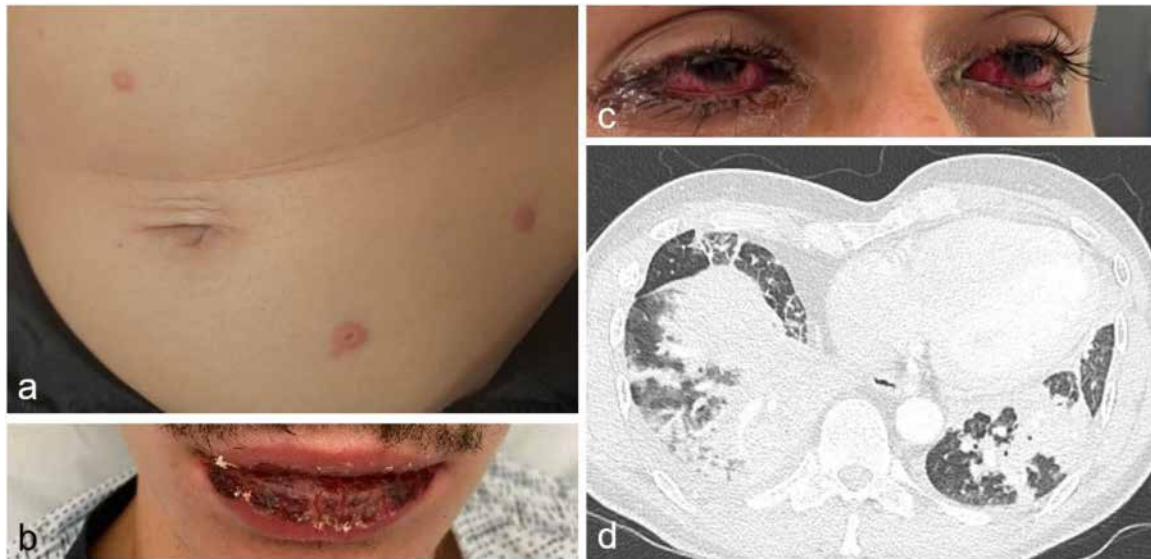


Figure 2. Extrapulmonary and pulmonary presentations of *Mycoplasma pneumoniae* infection. (a) Target-like lesions including a central bulla on the abdomen consistent with erythema multiforme in a 36-year-old female; (b) Mucositis in a 27-year-old male patient; (c) Bilateral conjunctivitis; (d) Bilateral *M. pneumoniae* pleuropneumonia with reactive mediastinal lymph nodes, in a 57-year-old male (axial computed tomographic-scan).

exacerbation due to a period of low exposure to respiratory pathogens because of restrictive measures due to the COVID-19 pandemic. Another explanation could be the emergence of a new *Mp* strain. A recent work suggested that while *Mp* genotypes may not determine specific clinical outcomes, they vary over time and geographic location [14].

While the clinical spectrum of *Mp* infections is diverse, including atypical presentations and potential complications, warranting heightened clinical awareness, we did not observe substantial differences between the current epidemic and the previous ones in our center. In most European countries, current empiric first-line treatment regimens for non-severe CAP mainly consist of beta-lactam antibiotics such as amoxicillin or cephalosporins, which are ineffective against *Mp* [15]. It is therefore imperative that clinicians have a high index of suspicion for atypical micro-organisms causing CAP such as *Mp*, particularly those with atypical clinical presentations or insufficient response to empiric antibiotic treatment for CAP. Increased access to *Mp* PCR testing could facilitate targeted treatment and prevent complications.

We also observed a high rate of co-infection (almost 25%), higher than the 18% reported in the Netherlands [6]. A high rate of co-infection with *Mp* and other respiratory pathogens has been previously described in 65% of children and 34% of adults presenting with acute respiratory infection in the United States [16]. Knowing that asymptomatic carriage could be frequent [17], clinical signs could not be specifically attributed to *Mp* in the case of co-infection. This may explain, at least in part, why more than half of patients hospitalized with *Mp* infection in our study required supportive oxygen therapy, which is higher than in other reports [18].

This study has several limitations. First, this was a single-center retrospective study, so our results could not be generalized. Second, the study population included adult patients assessed in hospitals, not outpatients. Insights into *Mp* infection in children were beyond the scope of the analysis. However, community health management remains critical to pandemic mitigation from a public health perspective. Third, an assessment of drug resistance was not conducted. Reported macrolide resistance is low in Europe, including France [19]. However, caution should be maintained as macrolide-resistance-associated genotypes have emerged as the predominant type in many Asian countries. Given the significance of genotype information in epidemiological investigations, ongoing surveillance is essential.

In conclusion, the clinical spectrum of *Mp* infections is diverse, including atypical presentations and potential complications including hypoxemic pneumonia, warranting heightened clinical awareness. However, in our center, epidemiological and clinical characteristics did not vary substantially between the current epidemic and the 5 previous years. The resurgence of *Mp* post-COVID-19 adds a layer of complexity to respiratory infections. Strategies addressing accurate diagnosis, prudent antibiotic use, and surveillance are critical to managing *Mp* infections worldwide.

Declarations of competing interest

The authors have no competing interests to declare.

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Ethical statement

This study was approved by the local ethics committee (Assistance Publique Hôpitaux de Marseille) and registered under the number PADS24-75. Written informed consent was given for the publication of clinical images when needed.

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Author contributions

All authors have made substantial contributions to one or all of the following: (1) the conception and design of the study (NC), the acquisition of data (AW, SE, BM, PL, JP, and AB), the analysis and interpretation of data (AW, SC, and NC), (2) drafting the article or revising it critically for important intellectual content (AW, SC, SH, FF, JCL, and NC), (3) final approval of the version to be submitted (AW, SE, BM, PL, JP, AB, SC, SH, FF, JCL, and NC).

Data availability

Data are available from the corresponding author upon reasonable request.

Supplementary materials

Supplementary material associated with this article can be found, in the online version, at [doi:10.1016/j.ijregi.2024.100548](https://doi.org/10.1016/j.ijregi.2024.100548).

References

- [1] Waites KB, Talkington DF. Mycoplasma pneumoniae and its role as a human pathogen. *Clin Microbiol Rev* 2004;17:697–728. doi:10.1128/CMR.17.4.697-728.2004.
- [2] Meyer Sauteur PM, Beeton ML. European Society of Clinical Microbiology and Infectious Diseases (ESCMID) Study Group for Mycoplasma and Chlamydia Infections (ESGMAC), and the ESGMAC Mycoplasma pneumoniae Surveillance (MAPS) study group. Mycoplasma pneumoniae: delayed re-emergence after COVID-19 pandemic restrictions. *Lancet Microbe* 2024;5:e100–1. doi:10.1016/S2666-5247(23)00344-0.
- [3] Bajantri B, Venkatram S, Diaz-Fuentes G. Mycoplasma pneumoniae: a potentially severe infection. *J Clin Med Res* 2018;10:535–44. doi:10.14740/jocmr3421w.
- [4] Meyer Sauteur PM, Beeton ML, Uldum SA, Bossuyt N, Vermeulen M, Loens K, et al. Mycoplasma pneumoniae detections before and during the COVID-19 pandemic: results of a global survey, 2017 to 2021. *Euro Surveill* 2022;27:2100746. doi:10.2807/1560-7917.ES.2022.27.19.2100746.
- [5] Bolluyt DC, Euser SM, Souverein D, van Rossum AM, Kalpoe J, van Westreenen M, et al. Increased incidence of Mycoplasma pneumoniae infections and hospital admissions in the Netherlands, November to December 2023. *Euro Surveill* 2024;29:2300724. doi:10.2807/1560-7917.ES.2024.29.4.2300724.
- [6] Nordholm AC, Søborg B, Jokelainen P, Lauenborg Møller K, Flink Sørensen L, Grove Krause T, et al. Mycoplasma pneumoniae epidemic in Denmark, October to December, 2023. *Euro Surveill* 2024;29:2300707. doi:10.2807/1560-7917.ES.2024.29.2.2300707.
- [7] Gong C, Huang F, Suo L, Guan X, Kang L, Xie H, et al. Increase of respiratory illnesses among children in Beijing, China, during the autumn and winter of 2023. *Euro Surveill* 2024;29:2300704. doi:10.2807/1560-7917.ES.2024.29.2.2300704.
- [8] Zayet S, Poloni S, Plantin J, Hamani A, Meckert Y, Lavoignet C-E, et al. Outbreak of Mycoplasma pneumoniae pneumonia in hospitalized patients: who is concerned? Nord Franche-Comté Hospital, France, 2023–2024. *Epidemiol Infect* 2024;152:e46. doi:10.1017/S0950268824000281.
- [9] Larcher R, Boudet A, Roger C, Villa F, Loubet P. Mycoplasma pneumoniae is back! Is it the next pandemic? *Anaesth Crit Care Pain Med* 2024;43:101338. doi:10.1016/j.accpm.2023.101338.
- [10] Edouard S, Boughammoura H, Colson P, La Scola B, Fournier P-E, Fenollar F. Large-scale outbreak of Mycoplasma pneumoniae infection, Marseille, France, 2023–2024. *Emerg Infect Dis* 2024;30:1481–4. doi:10.3201/eid3007.240315.
- [11] Morel AS, Dubourg G, Prudent E, Edouard S, Gouriet F, Casalta JP, et al. Complementarity between targeted real-time specific PCR and conventional broad-range 16S rDNA PCR in the syndrome-driven diagnosis of infectious diseases. *Eur J Clin Microbiol Infect Dis* 2015;34:561–70. doi:10.1007/s10096-014-2263-z.
- [12] Hoeting JA, Madigan D, Raftery AE, Volinsky CT. Bayesian model averaging: a tutorial (with comments by M. Clyde, David Draper and E. I. George, and a rejoinder by the authors). *Statist Sci* 1999;14:382–417. doi:10.1214/ss/1009212519.
- [13] Ramien ML. Reactive infectious mucocutaneous eruption: Mycoplasma pneumoniae-induced rash and mucositis and other parainfectious eruptions. *Clin Exp Dermatol* 2021;46:420–9. doi:10.1111/ced.14404.
- [14] Meyer Sauteur PM, Pánisová E, Seiler M, Theiler M, Berger C, Dumke R. Mycoplasma pneumoniae genotypes and clinical outcome in children. *J Clin Microbiol* 2021;59:e0074821. doi:10.1128/JCM.00748-21.
- [15] Waites KB, Xiao L, Liu Y, Balish MF, Atkinson TP. Mycoplasma pneumoniae from the respiratory tract and Beyond. *Clin Microbiol Rev* 2017;30:747–809. doi:10.1128/CMR.00114-16.
- [16] Diaz MH, Cross KE, Benitez AJ, Hicks LA, Kutty P, Bramley AM, et al. Identification of bacterial and viral Codetections with Mycoplasma pneumoniae using the Taq-Man array card in patients hospitalized with community-acquired pneumonia. *Open Forum Infect Dis* 2016;3:ofw071. doi:10.1093/ofid/ofw071.
- [17] Koenen MH, de Groot RCA, de Steenhuisen Pters WAA, Chu MLJN, Arp K, Hasrat R, et al. Mycoplasma pneumoniae carriage in children with recurrent respiratory tract infections is associated with a less diverse and altered microbiota. *EBiomedicine* 2023;98:104868. doi:10.1016/j.ebiom.2023.104868.
- [18] Kutty PK, Jain S, Diaz MH, Self WH, Williams D, Zhu Y, et al. Clinical and epidemiologic features of Mycoplasma pneumoniae infection among adults hospitalized with community-acquired pneumonia. *Int J Med Sci* 2024;21:3003–9. doi:10.7150/ijms.99233.
- [19] Kim K, Jung S, Kim M, Park S, Yang H-J, Lee E. Global trends in the proportion of macrolide-resistant Mycoplasma pneumoniae infections: A systematic review and meta-analysis. *JAMA Netw Open* 2022;5:e2220949. doi:10.1001/jamanetworkopen.2022.20949.