

Characteristics of *Stenotrophomonas maltophilia* infection in children in Sichuan, China, from 2010 to 2017

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Abstract

Stenotrophomonas maltophilia (*S. maltophilia*) is an important nosocomial bacterial pathogen. However, the clinical features of children with *S. maltophilia* infection, the predisposing factors, and the antibiotic susceptibility of the bacteria have not been fully evaluated.

In this study, the data of children with *S. maltophilia* infection from the West China Second University Hospital of Sichuan University (Chengdu, China) between July 2010 and October 2017 were collected and analyzed. The clinical features of enrolled children, the predisposing factors, and the antibiotic susceptibility were reported.

In total, infection of *S. maltophilia* was identified in 128 patients. Most of these patients were under 1 year old (67.2%) and were mainly diagnosed as pneumonia (69%). A large proportion had underlying diseases (45.3%), received immunosuppressive therapy (53.1%), had undergone invasive operations (41.4%), had a history of carbapenem antibiotics use within 7 days before culture acquisition (54.7%), history of intensive care unit (ICU) hospitalization within previous 30 days (34.4%), and other risk factors. In particular, invasive operation (95% confidence interval [CI]: 1.125–14.324, $P=.032$), especially mechanical ventilation (95% CI: 1.277–20.469, $P=.021$), and ICU admission (95% CI: 1.743–22.956, $P=.005$) were independent risk factors for the children to develop severe *S. maltophilia* infection. As for antibiotic susceptibility, trimethoprim sulfamethoxazole (TMP-SMX), piperacillin tazobactam, ticarcillin clavulanate, and ceftazidime exhibited strong antibacterial activities against *S. maltophilia*, the susceptibility rates were 97.5%, 86.7%, 92.9%, and 81.5%, respectively.

We report the clinical features of children with *S. maltophilia* infection, the predisposing factors and the antibiotic susceptibility. TMP-SMX can continue to be the first choice for the treatment of *S. maltophilia* infection. Piperacillin tazobactam, ticarcillin clavulanate, and the third generation cephalosporins can be used as alternative drugs.

Abbreviations: BALF = bronchoalveolar lavage fluid, HPF = high power field, ICU = intensive care unit, *S. maltophilia* = *Stenotrophomonas maltophilia*, TMP-SMX = trimethoprim-sulfamethoxazole.

Keywords: children, clinical characteristic, drug susceptibility, *Stenotrophomonas maltophilia*

Editor: Babak Abdinia.

This study was partially supported by grant 2015SZ0152 from the Science & Technology Department of Sichuan Province; grant KL036 from the West China Second University Hospital of Sichuan University.

The authors have no conflicts of interest to disclose.

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How to cite this article: Wang L, Zhou W, Cao Y, Yang C, Liu H, Chen T, Chen L. Characteristics of *Stenotrophomonas maltophilia* infection in children in Sichuan, China, from 2010 to 2017. *Medicine* 2020;99:8(e19250).

Received: 6 August 2019 / Received in final form: 22 December 2019 / Accepted: 16 January 2020

<http://dx.doi.org/10.1097/MD.00000000000019250>

1. Introduction

Stenotrophomonas maltophilia (*S. maltophilia*) is a Gram-negative, nonfermentative organism.^[1] It is one of the opportunistic pathogens of nosocomial infections, causing such serious infections in immunocompromised patients as pneumonia, septicemia, as well as infections of the skin and soft tissue, surgical wounds, and the urinary tract.^[1,2] Pediatric mortality due to *S. maltophilia* bacteremia was reported to be 6% to 40%.^[3,4] Due to aminoglycoside acetyl-transferase and enzymes that inactivate erythromycin and genes encoding efflux pumps, *S. maltophilia* strains are intrinsically resistant to a variety of antibiotics.^[5] Therefore, selection of an appropriate antimicrobial regimen for the treatment of *S. maltophilia* infection is a challenge for clinicians.

Thus far, most clinical studies of *S. maltophilia* have focused on the adult population, and only a limited number of studies describing the infection in children have been reported.^[6–8] Overall, there is a dearth of data on the clinical characteristics of this infection in Chinese children. In this study, we aimed to close this gap and explore the clinical features of children with *S. maltophilia* infection, the predisposing factors, and the antibiotic susceptibility. We reasoned that the results should be helpful for

early recognition and initiation of appropriate treatment of *S. maltophilia* infection in the clinical setting.

2. Materials and methods

2.1. Subjects and ethics statement

This study was conducted retrospectively at the West China Second University Hospital, Sichuan University (Chengdu, China). Data were collected between July 2010 and October 2017 from electronic medical records. The cultures that contained viable *S. maltophilia* were identified. Patients who satisfied the following criteria of *S. maltophilia* infection were included for further analysis (Fig. 1).

The Institutional Review Board/Ethics Committee affiliated with West China Second University Hospital, Sichuan University, approved this study, which was performed in accordance with the ethical standards of the Declaration of Helsinki.

2.2. Definitions

2.2.1. Infection of *S. maltophilia*. The criteria for diagnosis of *S. maltophilia* infection were as follows:

- (1) Site of isolation: isolates from nonrespiratory sites were included for analysis. The respiratory isolates were included only if they were from bronchoalveolar lavage fluid (BALF), tracheal intubation secretion or high quality sputum, that is, the number of leukocytes in the sputum smear was more than 25/high power field (HPF), and the epithelial cell number was less than 10/HPF, or the ratio of leukocyte to epithelial cell

was no less than 10.^[9] A BALF culture was considered positive when the number of *S. maltophilia* colonies was $\geq 10^4$ CFU/mL. As for the high quality specimen of sputum or tracheal secretion, it was also considered significant if *S. maltophilia* isolate was the sole bacteria or the dominant bacteria in the mixed flora.

- (2) Manifestations: patients should have clinical symptoms or signs of the corresponding site infection (such as shortness of breath, dyspnea, headache, etc), with associated laboratory results such as increased leukocytes and increased percentage of neutrophils or increased procalcitonin level, and elevated nucleated cells in the cerebrospinal fluid.^[10] Conversely, if patients did not have any infective symptoms or associated laboratory manifestations, their isolates were considered to be bacterial colonization or specimen contamination, and therefore, not included in the analysis. Appropriate empirical therapy was defined as microorganism susceptibility to one of several antimicrobial agents administered within 72 hours after the onset of bacterial infection.

2.2.2. Severe infection of *S. maltophilia*. The severity of illness was assessed by the Acute Physiology and Chronic Health Evaluation II score. The Charlson comorbidity index was used as an aggregate measure of comorbidities.^[7,11] Those who met any of the following criteria were classified as suffering severe *S. maltophilia* infection:

- (1) Patients died during this hospitalization; the attributable mortality (bacteremia-related death) was judged by 2 infectious diseases physicians, when the patient had no other

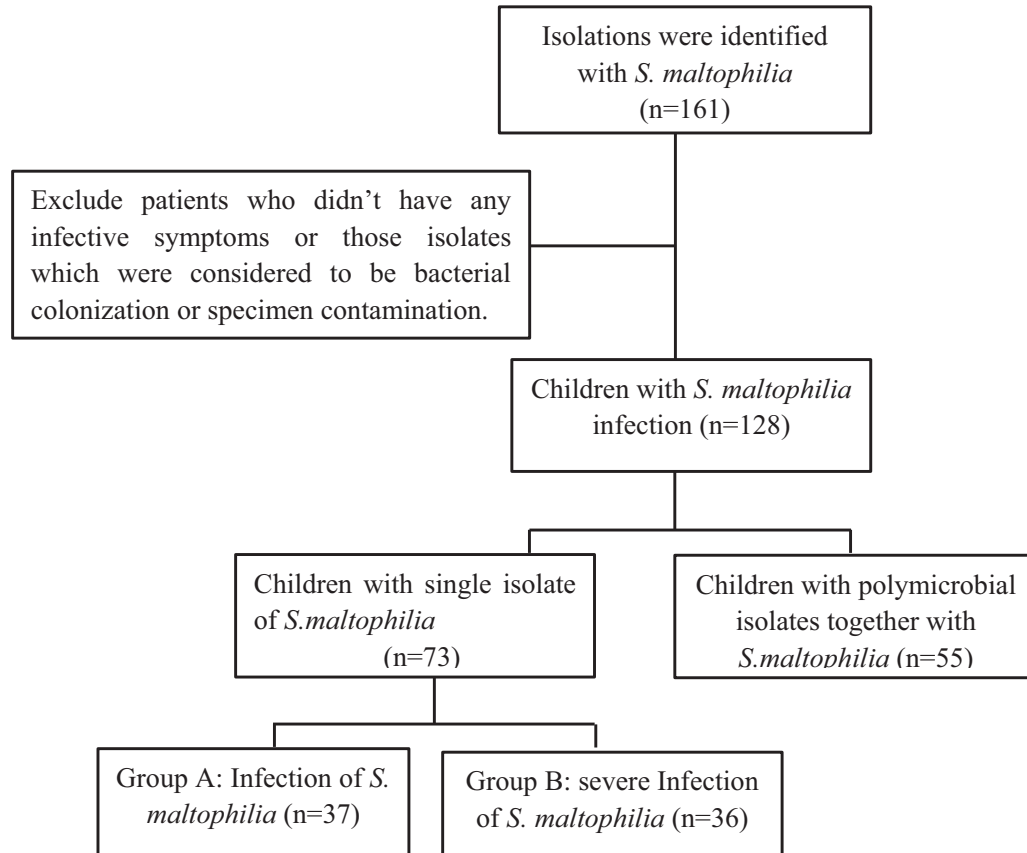


Figure 1. Study inclusion and exclusion criteria applied for patients.

Table 1**Site of isolation of *Stenotrophomonas maltophilia* (n = 161).**

Site of isolation	Number(%) of isolates			Total
	Infecting	Colonizing or contaminating	Undetermined	
BALF	9 (5.6%)	1 (0.6%)		10 (6.2%)
Tracheal intubation secretion	18 (11.2%)	1 (0.6%)	1 (0.6%)	20 (12.4%)
CSF	1 (0.6%)	0		1 (0.6%)
Blood	9 (5.6%)	0		9 (5.6%)
Sputum	90 (56%)	7 (4.3%)	22 (13.6%)	119 (73.9%)
Peritoneal Drainage fluid	1 (0.6%)	1 (0.6%)		2 (1.2%)
Total	128 (79.5%)	10 (6.2%)	23 (14.3%)	161 (100%)

BALF = bronchoalveolar lavage fluid, CSF = cerebrospinal fluid.

identifiable reason for death.^[6] If the death occurred within 7 days of isolation of *S. maltophilia*, it was considered related to the infection regardless of the presence of comorbid conditions that could potentially account for death.^[12]

(2) Patients were diagnosed as severe pneumonia, respiratory failure, heart failure or multiple organ dysfunction, and the association with *S. maltophilia* was evaluated by 2 infectious diseases physicians.

2.2.3. Underlying diseases. Renal diseases included nephrotic syndrome, renal insufficiency, and so on. Respiratory diseases included bronchopulmonary dysplasia, tracheobronchomalacia, and tracheobronchial stenosis. Heart diseases mainly included congenital heart diseases. Autoimmune diseases included systemic lupus erythematosus, and so on. Neurologic diseases included epilepsy, neonatal hypoxic and ischemic encephalopathy, and so on. Gastrointestinal diseases mainly included gastroesophageal reflux.

2.3. Microbiology

An automatic identification system, the Vitek2 system (bioMe'ieux, Marcy l'Etoile, France) was used to identify isolates of *S. maltophilia*. The antimicrobial susceptibilities were measured with ATB PSE 5, 25 STRIPS (bioMe'ieux, Marcy l'Etoile), and were interpreted according to the latest Clinical Laboratory Standard Institute M100 guideline.

2.4. Data collection and statistical analysis

The age, sex, manifestations, auxiliary examination, diagnosis, treatment, and other clinical information of the children were collected. Data were analyzed using the SPSS 19.0 software package (IBM, Armonk, NY). Continuous variables were compared using Student *t* test or the nonparametric Mann-Whitney *U* test, and categorical variables were compared using the Chi-squared (χ^2) or Fisher exact test. A logistic regression analysis was performed to study the associations between variables and disease severity. Two-sided *P* values of <.05 were considered statistically significant.

3. Results

3.1. Isolation rates and clinical characteristics of patients

A total of 161 children had positive isolates of *S. maltophilia* between July 2010 and October 2017. Among them, infection of *S. maltophilia* was identified in 128 patients (128/161, 79.5%), whereas 10 (10/161, 6.2%) isolates were considered as coloniza-

tion or contamination and 23 (23/161, 14.3%) were of undetermined origin. The respiratory tract was identified as the main infection site. The proportion of infecting and colonizing or contaminating isolates varied according to the site of isolation (Table 1).

During the study period, patients with *S. maltophilia* infection were mainly newborns and infants younger than 1-year-old, and 57% (73/128) of them were male. The mean length of hospitalization was 20 days (range, 1–134 days). The clinical manifestations of children infected with *S. maltophilia* were diverse. Besides fever and other systemic symptoms, respiratory symptoms such as dyspnea and cough were the most common manifestations. Overall, 69% of the children were primarily diagnosed as pneumonia, and 45.3% had underlying diseases (mainly due to premature birth, heart diseases, and hematological malignancy). In addition, 53.1% of the children received immunosuppressant therapy, among them 45.3% were given glucocorticoids. The ratio of patients who underwent invasive operations was 41.4%. The most common procedures included mechanical ventilation (37.5%), and central venous catheterization (9.4%). More than one-third of children had a history of intensive care unit (ICU) admission within the previous 30 days, and 54.7% of the patients received carbapenem within 7 days before culture acquisition (Table 2).

The antibiotic susceptibility of *S. maltophilia* isolates was summarized in Table 3. The highest antibiotic sensitivity were shown towards trimethoprim sulfamethoxazole and ticarcillin clavulanate. Piperacillin tazobactam and ceftazidime also had substantial antibacterial activities against the bacterium, the susceptibility rates being 86.7% and 81.5%, respectively. All isolates were resistant to imipenem and meropenem, whereas 75% of the isolates were resistant to ampicillin sulbactam.

3.2. Risk factors associated with severe *S. maltophilia* infection

Co-isolated bacterium, along with *S. maltophilia*, were found in 25 specimens (19.5%); of these, *Klebsiella pneumoniae* was the most common one (5.5%), followed by *Acinetobacter baumannii* (4.7%) and *Pseudomonas aeruginosa* (3.9%). Once the patients infected with multiple pathogens (such as other bacterium, fungus, mycoplasma, etc) were excluded, there were 73 children left with monomicrobial isolation. Based on the severity of the disease, these 73 patients were divided into severe and nonsevere groups, and the characteristics of the 2 groups were compared. There were no significant differences in age, sex, length of hospitalization, underlying comorbidities, and immunosuppressive therapy between the 2 groups. However, there were

Table 2
Demographic and clinical characteristics of patients with *Stenotrophomonas maltophilia* infection (n = 128).

Clinical backgrounds	No. (%)
Age, mo	6 (0–165)*
0–1	18 (14.1)
~3	25 (19.5)
~12	43 (33.6)
~36	14 (10.9)
36–165	27 (21.1)
Sex	
Male	73 (57)
Female	55 (43)
Length of hospitalization, d	20 (1–134)†
Manifestation	
Fever	72 (56.3)
Cough	77 (60.2)
Dyspnea	65 (50.8)
Wheeze	20 (15.6)
Septic shock	4 (3.1)
Convulsion	8 (6.3)
Conscious disturbance	7 (5.5)
Main diagnosis	
Pneumonia	88 (69)
Hematological malignancy	12 (9.4)
Neurologic diseases‡	11 (8.6)
Autoimmune diseases	2 (1.6)
Sepsis	3 (2.3)
Others§	12 (9.3)
Underlying diseases	58 (45.3)
Hematological malignancy	14 (11)
Renal diseases	2 (1.6)
Heart diseases	29 (22.7)
Respiratory diseases	15 (11.7)
Autoimmune diseases	1 (0.8)
Immune deficiency	3 (2.4)
Gastrointestinal diseases	3 (2.4)
Neurologic diseases	5 (3.9)
Premature	28 (21.9)
Malnutrition	13 (10.2)
Immunosuppressive therapy	68 (53.1)
Bone marrow transplantation within previous 30 d	2 (1.6)
Chemotherapy within previous 30 d	8 (6.3)
Use of glucocorticoid	58 (45.3)
Surgery within previous 30 d	9 (7)
Invasive operation	53 (41.4)
Operation number = 1	38 (29.7)
Operation number ≥2	15 (11.7)
Mechanical ventilation	48 (37.5)
Central venous catheter	12 (9.4)
Blood - purifying therapy	4 (3.1)
Gastric tube	6 (4.7)
Drainage tubing	3 (2.3)
Urethral catheter	1 (0.8)
ICU admission within previous 30 d	44 (34.4)
Neutropenia	1 (0.8)
Previous use of carbapenems	70 (54.7)
Mortality during hospitalization	5 (3.9)

* Range of age.

† Range of length of hospitalization.

‡ Neurologic diseases include epilepsy, neonatal respiratory distress syndrome, hypoxic-ischemic encephalopathy, intracranial hemorrhage, viral encephalitis, Japanese encephalitis, and so on.

§ Other diagnoses include hemophagocytic syndrome, nephrotic syndrome, intestinal tuberculosis, necrotic enterocolitis, genetic metabolic diseases, and so on.

|| Previous use of carbapenems: administration of carbapenems within 7 d before culture acquisition.

Table 3
Antibiotic susceptibility of *Stenotrophomonas maltophilia* isolates.

Antibiotics	n	Susceptible (%)	Intermediately susceptible (%)	Resistant (%)
Amikacin	52	52.4	4.8	42.9
Ampicillin sulbactam	50	15	10	75
Cefepime	52	63.6	18.2	18.2
Ceftazidime	119	81.5	9.3	10.2
Ciprofloxacin	55	43.6	16.4	40
bacillosporin	49	42.1	–	57.9
Gentamicin	49	47.4	–	52.3
Imipenem	108	0	–	100
Meropenem	104	0	–	100
Piperacillin	47	64.7	–	35.3
Piperacillin tazobactam	60	86.7	–	13.3
Ticarillin	47	70.6	–	29.4
Ticarillin clavulanate	108	92.9	1	6.1
Tobramycin	49	52.6	–	47.4
Trimethoprim sulfamethoxazole	128	97.5	–	0.8
Levofloxacin	53	44.2	15.4	40.4

Not all antibiotics were tested for each isolate. n = the sample number for each antibiotic. The percentage of isolates = the number of isolates with different antibiotic susceptibilities/n.

significantly higher rates of invasive operations (24.3% vs 52.8%, $P = .012$), mainly mechanical ventilation (10.8% vs 44.4%, $P < .001$), ICU admission (13.5% vs 44.4%, $P = .004$), and previous use of carbapenems (37.8% vs 69.4%, $P = .007$) in the severe group compared with the nonsevere group (Table 4).

Through logistic regression analysis, we determined factors that were significantly associated with severe *S. maltophilia* infection. The results of the multivariate analysis indicated that invasive operations (95% CI: 1.125–14.324, $P = .032$), especially mechanical ventilation (95% CI: 1.277–20.469, $P = .021$), and ICU admission (95% CI: 1.743–22.956, $P = .005$) were independent risk factors for the development of severe *S. maltophilia* infection (Table 4).

4. Discussion

S. maltophilia is emerging as an opportunistic pathogen among hospitalized pediatric patients. To understand the characteristics of *S. maltophilia* infection in children, we conducted this retrospective analysis and analyzed the antibiotic susceptibility as well as risk factors for severe diseases.

S. maltophilia infection mainly causes respiratory symptoms such as dyspnea, cough, and so on. Patients with *S. maltophilia* infection tended to have prolonged hospitalization, underlying diseases, immunosuppressive therapy, invasive operation, history of ICU admission, and previous use of carbapenems, which were in agreement with the results of studies in adults.^[13,14] The most common underlying illness identified in our study were heart diseases, and the most common invasive operation was mechanical ventilation. *S. maltophilia* can adhere to medical materials (eg, tracheal intubation) with its biofilm and increases the chances of lower respiratory tract infection, which may be the reason why patients with *S. maltophilia* infection tend to have a history of medical invasive operation.^[22]

In our study, the majority of children with *S. maltophilia* infection suffered from pneumonia (n = 117). And 59% (69/117) of these children developed severe pneumonia. The reasonable treatment of *S. maltophilia* infection is of great important.

Table 4**Factors associated with severe disease in patients with single isolate of *Stenotrophomonas maltophilia* (n = 73).**

Clinical backgrounds	Non-severe group (n = 37)	Severe group (n = 36)	P-value	Multivariate logistic regression analysis	
				Odds ratio (95% CI)	P-value
Age, mo	8 (0.03–165)	6.2 (0.03–160)	.310		
Male	21 (56.8%)	20 (55.6%)	.916		
Length of hospitalization, d	16 (1–83)	22.5 (3–134)	.106		
>28 d	8 (21.6%)	12 (33.3%)	.262		
Septic shock	1 (2.7%)	1 (2.8%)	–		
Underlying diseases	19 (51.4%)	12 (33.3%)	.119		
Immunosuppressive therapy	16 (43.2%)	21 (58.3%)	.197		
Invasive operations	9 (24.3%)	19 (52.8%)	.012*	4.015 (1.125–14.324)	.032*
Operation number = 1	6 (16.2%)	14 (38.9%)	–		
Operation number ≥2	3 (8.1%)	5 (13.9%)	–		
Mechanical ventilation	4 (10.8%)	16 (44.4%)	<.001*	5.113 (1.277–20.469)	.021*
ICU admission within previous 30 d	5 (13.5%)	16 (44.4%)	.004*	6.326 (1.743–22.956)	.005*
Neutropenia	0	1 (2.8%)	–		
Previous use of carbapenems	14 (37.8%)	25 (69.4%)	.007*	0.480 (0.147–1.565)	.224

* Two-sided P values of <.05 were considered statistically significant.
CI = confidence interval, ICU = intensive care unit.

Treatment of *S. maltophilia* infection is difficult, in part because the bacteria is resistant to a variety of antimicrobial agents. Trimethoprim-sulfamethoxazole (TMP-SMX) has been generally effective, based on in vitro susceptibility assays and reports of clinical outcomes.^[15–17] However, an increasing number of studies have reported the resistance of *S. maltophilia* to TMP-SMX,^[7,18] which presents a major challenge to clinical physicians. Therefore, we considered it important to determine the resistance rate of *S. maltophilia* to commonly used antibiotics, especially TMP-SMX in children. Fortunately, we found that *S. maltophilia* was highly susceptible to TMP-SMX, which was consistent with most studies published thus far.^[8,19,20] This suggests that TMP-SMX could still be the first choice for the treatment of *S. maltophilia* infection. Furthermore, a large proportion of isolates were also susceptible to ticarcillin clavulanate. A variety of studies have reported high resistance rates of *S. maltophilia* to cephalosporin antibiotics.^[7,20,21] However, ceftazidime showed substantial antibacterial activity against the bacteria in our study. The reason for the difference is currently unclear. The irrational use of cephalosporins maybe 1 reason. In addition, the sensitivity rate to piperacillin tazobactam was also found to be high. We, therefore, suggest that the third generation cephalosporins, as well as piperacillin tazobactam and ticarcillin clavulanate can be used as alternative drugs to patients who cannot tolerate TMP-SMX. Of note, *S. maltophilia* exhibits high-level intrinsic resistance to carbapenem antibiotics because of the production of the versatile L1 type β -lactamase (also called “carbapenemase”), which is capable of hydrolyzing carbapenem antibiotics.^[19]

To our knowledge, few studies have analyzed the risk factors for children to develop severe *S. maltophilia* infection. In the present study, we have analyzed the clinical characteristics of children with severe infection, and identified the associated risk factors. To exclude the effect of polymicrobial infection on the results of the study, we only analyzed the children with monomicrobial *S. maltophilia* infection. Invasive operations (especially mechanical ventilation), use of carbapenems within 7 days before culture acquisition, and ICU admission within the previous 30 days were found to be associated with severe infection. Multivariate logistic regression analysis identified

invasive operations (mainly consisting mechanical ventilation) and ICU admission as independent risk factors for the development of severe infection. This may be due to the fact that the clinical application of ventilator weakens the cough reflex and the mucosal cilia clearance function, and in parallel, promotes proliferation of the bronchial glands and increased secretion, which in turn increases the chances of respiratory infection.^[22] Thus, for children with long term ICU hospitalization and invasive operations such as mechanical ventilation, the clinicians should keep a particularly wary eye on *S. maltophilia* infection. Rational use of broad-spectrum antibiotics such as the carbapenems and practicing meticulous hand hygiene are also advised.

There were some potential limitations of our study. First, it was a single center study with a moderate sample size, which may not be generalizable. Second, because of the retrospective design, selection and observational bias may have affected the results. A more elaborate, match controlled study may be conducted in the future to further confirm our results.

In conclusion, TMP-SMX can continue to be the first choice for the treatment of *S. maltophilia* infection, while piperacillin tazobactam, ticarcillin clavulanate, and the third generation cephalosporins can be used as alternative drugs. To prevent the severe diseases, clinicians should use antibiotics more rationally, ensure good management of sterilization practices, and isolate children with high-risk factors.

Author contributions

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Software: Lili Wang.

Supervision: Hanmin Liu, Lina Chen.

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