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## RESEARCH ARTICLE



# Epidemiology and antibiotic resistance profile of bacterial meningitis in Morocco from 2015 to 2018

YOUSSEF IKKEN<sup>1,2\*</sup> , RÉDA CHAROF<sup>2</sup>,  
AMINA BENAOUA<sup>3</sup>, FARIDA HILALI<sup>1</sup> ,  
SANAE AKKAOUT<sup>4</sup> , MOSTAFA ELOUENNASS<sup>5</sup> and  
YASSINE SEKHSOKH<sup>1</sup>

<sup>1</sup> Biosafety Level 3 and Research Laboratory, Mohammed V Military Teaching Hospital, Faculty of Medicine and Pharmacy, Mohammed V University in Rabat, Rabat 10 000, Morocco

<sup>2</sup> Laboratory of Medical Bacteriology, National Institute of Hygiene in Rabat, 27, Avenue Ibn Batouta, B.P. 769, Rabat 10 000, Morocco

<sup>3</sup> Laboratory of Microbiology, Cheick-Zaid University Hospital, University Internationale Abulcasis of Sciences and Health in Rabat, B.P. 6533, Avenue Allal El Fassi, Madinat Al Irfane, Rabat 10 000, Morocco

<sup>4</sup> Research Laboratory in Oral Biology and Biotechnology, Faculty of Dental Medicine, Mohammed V University in Rabat, Rabat 10 000, Morocco

<sup>5</sup> Laboratory of Bacteriology, Mohammed V Military Teaching Hospital, Faculty of Medicine and Pharmacy, Mohammed V University in Rabat, Rabat 10 000, Morocco

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## ABSTRACT

Over a 4-year study period from 2015 to 2018, altogether 183 isolates of bacterial meningitis were collected from 12 hospitals covering the entire Moroccan territory. *Neisseria meningitidis* represented 58.5%, *Streptococcus pneumoniae* 35.5%, and *Haemophilus influenzae type b* 6%. *H. influenzae type b* mainly affected 5-year-olds and unvaccinated adults. *N. meningitidis serogroup B* represented 90.7% followed by *serogroup W135* with 6.5%. Decreased susceptibility to penicillin G (DSPG) for all isolates accounted for 15.7%, with 11.6% being resistant to penicillin G (PG) and 4.1% decreased susceptibility. Cumulative results of all strains showed 2.7% decreased susceptibility to amoxicillin and 3.3% resistant, 2.2% of isolates were resistant to third-generation cephalosporin and 2.2% were decreased susceptible, 5.5% were resistant to chloramphenicol and 2.7% were resistant to rifampin. The frequency of DSPG observed in our study is more common in *S. pneumoniae* than in *N. meningitidis* ( $P < 0.05$ ). These isolates have been found to be highly susceptible to antibiotics used for treatment and prophylaxis chemotherapy and the observed resistance remains rare. The impact of introduction of conjugate vaccines against *H. influenzae type b* and *S. pneumoniae* (PCVs) is an advantage in reducing meningitis cases due to these two species.

## KEYWORDS

antimicrobial resistance, co-resistance, *Haemophilus influenzae*, epidemiology meningitis, *Neisseria meningitidis*, *Streptococcus pneumoniae*, vaccination

## INTRODUCTION

The most common causes of bacterial meningitis are *Neisseria meningitidis*, *Streptococcus pneumoniae*, and *Haemophilus influenzae type b*, and these are responsible for 75,000, 118,400, and 83,000 deaths worldwide, respectively and lead to severe neurological morbidity, despite advances in antimicrobial therapy [1]. In sub-Saharan Africa, the so-called “meningitis

\*Corresponding author. National Institute of Hygiene in Rabat, 27, Avenue Ibn Batouta, B.P. 769, Rabat 10 000, Morocco.  
Tel.: +212 05 37 77 19 02 65 30;  
fax: +212 05 37 77 20 67.  
E-mail: ikkenyoussef@gmail.com

belt” *N. meningitidis*, *S. pneumoniae*, and *H. influenzae type b* are the most common bacteria [2]. In Morocco, the cumulative incidence of meningitis is 2.9 per 100,000 population and 2/3 of all meningitis cases were considered meningococcal [3]. However, it is also important to diagnose and begin treatment as soon as possible to avoid any adverse consequences. The treatment of choice is the administration of amoxicillin and/or third generation cephalosporin as an empirical treatment regimen for all patients with bacterial meningitis [4]. In Morocco, the management of cases of syndromic bacterial meningitis is based on empirical antibiotic therapy which usually includes a third generation cephalosporin (C3G), which remains a reference treatment and this treatment must be started in front of any cerebrospinal fluid (CSF) purulent [5]. This empirical antibiotic therapy is given urgently even before the identification stage of the three isolates (*N. meningitidis*, *S. pneumoniae*, and *H. influenzae type b*) and not yet known their susceptibility to first-line antibiotics. A reflection on bacteriological profile and the current situation of antibiotic resistance of three bacterial pathogens responsible for syndromic bacterial meningitis in our national context is necessary. The objective of this study is therefore to study the phenotypic profile and to determine the antimicrobial susceptibility of the three bacteria (*N. meningitidis*, *S. pneumoniae*, and *H. influenzae type b*) responsible for meningitis in Morocco. The treatment and chemoprophylaxis as well as the target vaccine will be highlighted in this study.

## MATERIALS AND METHODS

### Statement of ethics

Isolates from this study were collected as part of the national meningitis surveillance system in Morocco. The laboratory of medical bacteriology at the National Institute of Hygiene is the national laboratory of meningitis. All patient identity information was treated confidentially.

### Data

The data was entered using Excel software and the statistical calculations were performed using SPSS (Statistical Package for Social Sciences version 13.0, SPSS Inc., Chicago, IL, USA). The quantitative variables were expressed in median and quartiles. The qualitative variables were expressed in terms of number and percentage. The comparison of the qualitative variables was made by the  $\chi^2$  test and the exact Fisher test. A  $P$ -value < 0.05 was considered as statistically significant.

### Design of the study

It is a case study of bacterial isolates analyzed at the level of laboratory of medical bacteriology at the National Institute of Hygiene in Morocco. These isolates were collected over a 4-year period from 2015 to 2018 from 12 hospitals covering the entire territory of the Kingdom. For this reason, 183 isolates were collected, including 107 *N. meningitidis*, 65 *S.*

*pneumoniae*, and 11 *H. influenzae type b* from the CSFs of culture-confirmed bacterial meningitis. Only one isolate per patient was included.

### Microbiological identification

Samples of CSF sent to laboratories were treated according to the usual methods including: chemistry and microscopic examination. The bacterial culture was performed on poly-aminated chocolate agar and Columbia agar added with 5% defibrinated horse blood (Oxoid). The agar was incubated in jars containing 5% CO<sub>2</sub> in an oven at 37 °C for a maximum of 18–72 h. Biochemical identification of these isolates was done by the API-NH and API Strep (Bio-merieux), oxidase test (Oxoid), optochine test (Oxoid), and cefinase test (Oxoid).

### Serogroup and agglutination tests

Soluble antigens were tested on the CSF supernatant by latex agglutination for *S. pneumoniae*, *H. influenzae type b*, *N. meningitidis* serogroup A, B, C, Y, W135, *Escherichia coli* K1, and Group B *Streptococcus* (Oxoid, Wellcogen, Remel Europe Ltd., UK). Confirmation of *S. pneumoniae* and *H. influenzae type b* was done by the same kit from positive cultures. Subsequently and from the cultures, *N. meningitidis* isolates were serogrouped by agglutination test on the slide with serogroup A, B, C, 29E, Y, and W135 antisera (Oxoid). Both procedures were performed in accordance with the manufacturer’s instructions.

### Antibiotic susceptibility

The antibiotic susceptibility test was performed by the agar diffusion method using a standard dilution suspension of 0.5 McFarland, antibiotic discs, and MIC on Mueller–Hinton agar supplemented with 5% defibrinated horse blood (Oxoid), culture was performed under aerobic condition with increased CO<sub>2</sub> pressure. Antibiotic susceptibility testing and interpretation of results were done according to EUCAST (2019). Quality control of all antibiotics tested was examined using *Staphylococcus aureus* (ATCC 25923) and *E. coli* (ATCC 25922) as control strains.

The antibiotics included in this study for the three isolates are: (1) *S. pneumoniae*: Discs: oxacillin (1 µg), trimethoprim-sulfamethoxazole or cotrimoxazole (1.25–23.75 µg), chloramphenicol (30 µg), norfloxacin (10 µg), levofloxacin (5 µg), rifampin (5 µg), tetracycline (30 µg), erythromycin (15 µg) (Oxoid). MIC: penicillin G, amoxicillin, cefotaxime, ceftriaxone, vancomycin (Liofilchem). (2) *N. meningitidis*: Discs: chloramphenicol (30 µg), rifampin (5 µg) (Oxoid). MIC: penicillin G, amoxicillin, cefotaxime, ceftriaxone, ciprofloxacin (Liofilchem). (3) *H. influenzae type b*: Discs: Penicillin G (1 unit), trimethoprim-sulfamethoxazole (1.25–23.75 µg), chloramphenicol (30 µg), rifampin (5 µg), tetracycline (30 µg), erythromycin (15 µg), ciprofloxacin (5 µg), levofloxacin (5 µg). MIC: amoxicillin, ampicillin, cefotaxime, ceftriaxone, ciprofloxacin (Liofilchem).



Table 1. Characteristics of the population of isolates studied

Characteristics	(Effective)*	<i>H. influenzae</i> type <i>b</i>	<i>N. meningitidis</i>			<i>S. pneumoniae</i>	<i>P</i> value
			<i>N. meningitidis</i> NG	<i>N. meningitidis</i> B	<i>N. meningitidis</i> W135		
Isolates*	100(183)	6.0(11)	1.6(3)	53.0(97)	3.8(7)	35.5(65)	–
Sex*	100(163)						
Women	39.3(64)	4.7(3)	0.0(0)	60.9(39)	1.6(1)	32.8(21)	0.450
Man	60.7(99)	6.1(6)	3.0(3)	49.5(49)	3.0(3)	38.4(38)	
Age * in year	100(102)						
<2	24.5(25)	4.0(1)	8.0(2)	56.0(14)	4.0(1)	28.0(7)	0.278
[2,5]	36.3(37)	8.1(3)	2.7(1)	56.8(21)	5.4(2)	27.0(10)	
[6, 15]	11.8(12)	8.3(1)	0.0(0)	75.0(9)	0.0(0)	16.7(2)	
[16, 25]	8.8(9)	11.1(1)	0.0(0)	55.6(5)	0.0(0)	33.3(3)	
[26, 49]	11.8(12)	0.0(0)	0.0(0)	41.7(5)	16.7(2)	41.7(5)	
≥50	6.9(7)	0.0(0)	0.0(0)	0.0(0)	14.3(1)	85.7(6)	

\* Percentage % (expressed in effective). *N. meningitidis*: *Neisseria meningitidis*; *S. pneumoniae*: *Streptococcus pneumoniae*; *H. influenzae* type *b*: *Haemophilus influenzae* type *b*.

## RESULTS

### Characteristics of the isolates (Table 1)

Of the 183 isolates collected, *N. meningitidis* represented 58.4% including 53% (97/107) of *N. meningitidis* serogroup B (*N. meningitidis* B), 3.8% (7/107) of *N. meningitidis* serogroup W135 (*N. meningitidis* W135), and 1.6% (3/107) of non-serogrouped *N. meningitidis* (*N. meningitidis* NG). *S. pneumoniae* and *H. influenzae* type *b* represented respectively 35.5% (65/183) and 6% (11/183). Regarding sex, 60.7% were male and 39.3% were female (Sex Ratio: 1.55). The median age of the sample studied was 3.9 [1.8; 17] years, ranging from 3 days to 65 years. In infant's <2 years of age, *N. meningitidis* represented 68% of meningitis cases followed by 28% of *S. pneumoniae*. In the children of [2; 5] years, *N. meningitidis* represented 64.9%, followed by *S. pneumoniae* with 27%. In older children [6; 15] years old, *N. meningitidis* represented 75% and *S. pneumoniae* was 16.7%. In adults [16; 25] years old, *N. meningitidis* represented 55.6% and *S. pneumoniae* was 33.3%. In older adults [26; 49] years old, the *N. meningitidis* represented 58.4% and *S. pneumoniae* was at 41.7%. Beyond the age of 50, *N. meningitidis* was 14.3% while *pneumoniae* was 85.7%. However, at age ≤5 years, we have 66.7% (4/7) of *H. influenzae* type *b* included in the study who reached this age.

### Antimicrobial resistance of isolates to antibiotics (Tables 2 and 3)

For all isolates tested of *N. meningitidis*, *S. pneumoniae*, and *H. influenzae* type *b*; the decreased susceptibility to penicillin G (DSPG) represented 15.7% (27/183), with 11.6% (20/183) which were resistant to penicillin G (PG) and 4.1% (7/183) intermediate. Amoxicillin represents 2.7% (5/183) which were intermediate and 3.3% (6/183) resistant. 2.2% (2/183) of isolates were resistant to cefotaxime and ceftriaxone and 2.2% (2/183) were intermediate. 5.5% (10/183) were resistant to chloramphenicol and 2.7% (5/183) were resistant to rifampin. The frequency of DSPG observed in our study was

more common in *S. pneumoniae* than *N. meningitidis* ( $P < 0.05$ ). Of the 107 strains of *N. meningitidis*, 9.3% (10/107) were DSPG of which 2.8% (3/107) were resistant, 5.6% (6/107) were reduced susceptibility to amoxicillin with a strain that was resistant 0.9% (1/107), one strain was resistant to ceftriaxone and cefotaxime 0.9% (1/107), 3.7% (4/107) strains were resistant to chloramphenicol and 1.9% (2/107) resistant to rifampin. All strains are susceptible to ciprofloxacin and no strains produced beta-lactamase. The level of DSPG and amoxicillin resistance in *N. meningitidis* B and *N. meningitidis* W135 were respectively 7.5% (8/107) vs. 1.8% (2/107) and 3.7% (4/107) vs. 1.9% (2/107). While resistance to ceftriaxone, cefotaxime, rifampin, and chloramphenicol was only recorded in *N. meningitidis* B. In our sample, resistance to amoxicillin was more common in *N. meningitidis* B than *N. meningitidis* W135 ( $P < 0.05$ ). All strains of *N. meningitidis* with DSPG have MIC of 0.064 µg/mL at 0.25 µg/mL and the resistant have MIC of 0.38 µg/mL at 1 µg/mL. MIC amoxicillin in intermediate isolates range from 0.19 µg/mL to 0.75 µg/mL and the only resistant isolate was a MIC of 1 µg/mL. The MIC of ceftriaxone and cefotaxime for the resistant strain was 0.25 µg/mL.

Of the 65 strains of *S. pneumoniae*, 26.2% (17/65) were of DSPG and 7.7% (5/65) were resistant to amoxicillin, 4.6% (3/65) were of reduced susceptibility to ceftriaxone and cefotaxime with 3.1% (2/65) were intermediate and one strain 1.5% (1/65) was resistant, 9.2% (6/65) strains were resistant to chloramphenicol and 3.1% (2/65) were resistant to rifampin. Cotrimoxazole recorded 30% (18/65) of reduced susceptibility of which 20% (12/65) were resistant and 10% (8/65) were intermediate. Levofloxacin and erythromycin represented 2.7% (2/65) and 12.7% (8/65) of resistance, respectively. Tetracycline represented 15% (9/65) of resistance and 5% (3/65) of intermediate resistance. All strains are susceptible to vancomycin. Strains of *S. pneumoniae* that were resistant to PG have MIC between 0.125 µg/mL and 2 µg/mL. The MIC of resistant amoxicillin was 0.75–1 µg/mL. The intermediate MIC of cefotaxime and ceftriaxone was 0.75 µg/mL at 1 µg/mL, and that of the resistant strain was 2 µg/mL.

Table 2. Analysis of common antibiotics in NM, SP, and Hib isolates

ATB		* <i>H. influenzae</i> type b (11)	* <i>N. meningitidis</i> (107)	* <i>S. pneumoniae</i> (65) <sup>a,b,c</sup>	*Cumulative	P value
Discs						
Oxacillin	S	NT	94.4(101)	76.9(50)	87.8(151)	0.001
	I	NT	–	–	–	
	R	NT	5.6(6)	23.1(15)	12.2(21)	
Penicillin G	S	NT	94.4(101)	76.9(50)	87.8(151)	0.001
	I	NT	–	–	–	
	R	NT	5.6(6)	23.1(15)	12.2(21)	
<sup>a</sup> Trimethoprim-sulfamethoxazol	S	90.9(10)	NT	70.0(42)	73.2(52)	0.323
	I	0(0)	NT	10.0(6)	8.5(6)	
	R	9.1(1)	NT	20.0(12)	18.3(13)	
Chloramphenicol	S	100(11)	96.3(103)	90.8(59)	94.5(173)	0.219
	I	–	–	–	–	
	R	0(0)	3.7(4)	9.2(6)	5.5(10)	
<sup>b</sup> Levofloxacin	S	100(11)	NT	96.9(62)	97.3(73)	0.552
	I	–	NT	–	–	
	R	0(0)	NT	3.1(2)	2.7(2)	
Rifampin	S	90.9(10)	98.1(105)	96.9(63)	97.3(178)	0.367
	I	–	–	0(0)	0(0)	
	R	9.1(1)	1.9(2)	3.1(2)	2.7(5)	
<sup>a</sup> Tetracycline	S	100(11)	NT	80.0(48)	83.1(59)	0.266
	I	0(0)	NT	5.0(3)	4.2(3)	
	R	0(0)	NT	15.0(9)	12.7(9)	
<sup>c</sup> Erythromycin	S	90.9(10)	NT	87.3(55)	87.8(65)	0.736
	I	0(0)	NT	0(0)	0(0)	
	R	9.1(1)	NT	12.7(8)	12.2(9)	
MIC						
Penicillin G	S	NT	90.7(97)	73.8(48)	84.3(145)	0.000
	I	NT	6.5(7)	–	4.1(7)	
	R	NT	2.8(3)	26.2(17)	11.6(20)	
Cefotaxime	S	100(11)	99.1(106)	95.4(62)	97.8(179)	0.412
	I	–	–	3.1(2)	1.1(2)	
	R	0(0)	0.9(1)	1.5(1)	1.1(2)	
Ceftriaxone	S	100(11)	99.1(106)	95.4(62)	97.8(179)	0.412
	I	–	–	3.1(2)	1.1(2)	
	R	0(0)	0.9(1)	1.5(1)	1.1(2)	
Amoxicillin	S	100(11)	94.4(101)	92.3(60)	94(172)	0.047
	I	–	4.7(5)	–	2.7(5)	
	R	0(0)	0.9(1)	7.7(5)	3.3(6)	
Ciprofloxacin	S	100(11)	100(107)	NT	100(118)	–
	I	–	–	NT	–	
	R	0(0)	0(0)	NT	0(0)	
Vancomycin	S	–	–	100(65)	100(65)	–
	I	–	–	–	–	
	R	–	–	0	0(0)	

\*Percentage% (expressed in effective); a, b, c: the number tested [a(60), b(64), c(63)]; NT: Not Tested.

All *H. influenzae* type b strains in our study were susceptible to the majority of antibiotics except one strain that was resistant to erythromycin, rifampin, and cotrimoxazole. No strains produced beta-lactamase.

#### Co-resistance to PG and other antibiotics (Table 4)

Of the strains of *S. pneumoniae* DSPG: 8.3% were resistant to cotrimoxazole, 3.1% were resistant to levofloxacin, 1.54% were resistant to rifampin, 10% were resistant to tetracycline,

4.6% were resistant to cefotaxime and ceftriaxone, and 7.7% were resistant to amoxicillin. This association of co-resistance between PG, C3G, and amoxicillin in *S. pneumoniae* was statistically significant ( $P < 0.05$ ).

In strains of *N. meningitidis* DSPG: 1.9% chloramphenicol, 0.9% rifampin, 0.9% ceftriaxone, 0.9% cefotaxime, and 3.7% amoxicillin were resistant, respectively. This association of co-resistance between PG, Beta-lactam antibiotics, rifampin, and chloramphenicol in *N. meningitidis* was statistically significant ( $P < 0.05$ ).



Table 3. Resistance Analysis in *Neisseria meningitidis* serogroup

ATB(107)		* <i>N. meningitidis</i> NG 2.8(3)	* <i>N. meningitidis</i> B 90.7(97)	* <i>N. meningitidis</i> W135 6.5(7)	*Cumulative	P value
Discs						
Oxacillin	S	2.8(3)	86(92)	5.6(6)	94.4(101)	0.546
	I	–	–	–	–	
	R	0(0)	4.7(5)	0.9(1)	5.6(6)	
Penicillin G	S	2.8(3)	86(92)	5.6(6)	94.4(101)	0.546
	I	–	–	–	–	
	R	0(0)	4.7(5)	0.9(1)	5.6(6)	
Chloramphenicol	S	2.8(3)	86.9(93)	6.5(7)	96.3(103)	0.807
	I	–	–	–	–	
	R	0(0)	3.7(4)	0(0)	3.7(4)	
Rifampin	S	2.8(3)	88.8(95)	6.5(7)	98.1(105)	0.900
	I	–	–	–	–	
	R	0(0)	1.9(2)	0(0)	1.9(2)	
MIC						
Penicillin G	S	2.8(3)	83.2(89)	4.7(5)	90.7(97)	0.314
	I	0(0)	5.6(6)	0.9(1)	6.5(7)	
	R	0(0)	1.9(2)	0.9(1)	2.8(3)	
Cefotaxime	S	2.8(3)	89.7(96)	6.5(7)	99.1(106)	0.949
	I	–	–	–	–	
	R	0(0)	0.9(1)	0(0)	0.9(1)	
Ceftriaxone	S	2.8(3)	89.7(96)	6.5(7)	99.1(106)	0.949
	I	–	–	–	–	
	R	0(0)	0.9(1)	0(0)	0.9(1)	
Amoxicillin	S	2.8(3)	86.9(93)	4.7(5)	94.4(101)	0.045
	I	0(0)	2.8(3)	1.9(2)	4.7(5)	
	R	0(0)	0.9(1)	0(0)	0.9(1)	
Ciprofloxacin	S	2.8(3)	90.7(97)	6.5(7)	100(107)	–
	I	–	–	–	–	
	R	0(0)	0(0)	0(0)	0(0)	

\* Percentage % (expressed in effective); S: Sensitive, I: Intermediate, R: Resistant.

## DISCUSSION

This national study conducted in Morocco focuses on both the epidemiological profile of the three bacteria responsible for bacterial meningitis, which were *N. meningitidis*, *S. pneumoniae*, and *H. influenzae type b*, and their resistance profile. For this reason, we collected 183 isolates from different regions and cities across the country from culture-confirmed meningitis cases. Isolates are divided into 58.5% (107/183) *N. meningitidis*, 35.5% (65/183) *S. pneumoniae*, and 6% (11/183) *H. influenzae type b*. The age group most affected was that  $\leq 5$  years old, which represented 60.8%. This study showed that in Moroccan children and adults, meningococcal disease causes more meningitis followed by pneumococcal disease. *H. influenzae type b* come last with very low rates and mainly reaches children  $\leq 5$  years old and unvaccinated adults. The man is more affected than the woman. The technical difficulties associated with the high rates of negative culture and the demanding transport conditions of these three bacteria are technical problems limiting the collections of these isolates. As a result, the rate of positive cultivation between regions of the country varies between 7 and 12%.

In the Maghreb countries of northern Africa, despite the poverty of data, pneumococcus was the one that causes more

meningitis than meningococcal [6, 7]. In Europe and the African meningitis belt, meningococcal disease and pneumococcus were the two most dominant bacterial species in children and adults [8, 9].

The most dominant *N. meningitidis* serogroup in our study was *N. meningitidis* B followed by *N. meningitidis* W135. Serogroup commonly detected in the African meningitis belt, serogroup C, W, and X are those that cause epidemics [8]; In Europe *N. meningitidis* B was responsible for most cases of meningococcal meningitis [9, 10]; while in South Africa are *N. meningitidis* W135, in Saudi Arabia were *N. meningitidis* A and *N. meningitidis* W135 [11]. The bacteriological and epidemiological profile of meningitis in Morocco was different from neighboring countries, the African meningitis belt and even countries with a tourist destination (Turkey) or spiritual (Saudi Arabia) visited by Moroccan citizens. But this profile was very similar to the profile found in Europe.

As presented by our results, there was a dramatic regression in cases of meningitis due to *H. influenzae type b*. This decrease in cases of *H. influenzae type b* meningitis in children  $\leq 5$  years was the result of the introduction in 2007 into the national immunization program of the *H. influenzae type b* conjugate vaccine for children from the age of 2 months [12].



Table 4. Co-resistance of common antibiotics relative to sensitivity and resistance of PG in *Neisseria meningitidis* and *Streptococcus pneumoniae*

PG	<i>N. meningitidis</i> (107)				P value	<i>S. pneumoniae</i> (65) <sup>a, b, c</sup>				P value
	Resistant**		Sensible*			Resistant**		Sensible*		
	(R + I) 9.4(10)	R 2.8(3) I 6.5(7)	90.7 (97)			(R + I) 26.2(17)	R 26.2(17) I 0(0)	48 (73.85)		
Discs**										
SXT <sup>a</sup>	–	–	–	–	---	8.3(5)	5(3)	22(13)	15(9)	0.881
CHL	1.9(2)	1.9(2) 0(0)	1.9(2)	1.9(2) 0(0)	0.006	0(0)	0(0) 0(0)	9.2(6)	9.2(6) 0(0)	0.149
LEV <sup>b</sup>	–	–	–	–	---	3.1(2)	3.1(2) 0(0)	0(0)	0(0) 0(0)	0.067
RIF	0.9(1)	0.9(1) 0(0)	0.9(1)	0.9(1) 0(0)	0.020	1.5(1)	1.5(1) 0(0)	1.5(1)	1.5(1) 0(0)	0.458
TET <sup>a</sup>	–	–	–	–	---	10(6)	8.3(5) 1.7(1)	10(6)	6.7(4) 3.3(2)	0.037
ERY <sup>c</sup>	–	–	–	–	---	4.8(3)	4.8(3) 0(0)	7.9(5)	7.9(5) 0(0)	0.326
MIC**										
CTX	0.9(1)	0.9(1) 0(0)	0(0)	0(0) 0(0)	0.000	4.6(3)	1.5(1) 3.1(2)	0(0)	0(0) 0(0)	0.012
CRO	0.9(1)	0.9(1) 0(0)	0(0)	0(0) 0(0)	0.000	4.6(3)	1.5(1) 3.1(2)	0(0)	0(0) 0(0)	0.012
AMX	3.7(4)	0.9(1) 2.8(3)	1.9(2)	0(0) 1.9(2)	0.000	7.7(5)	7.7(5) 0(0)	0(0)	0(0) 0(0)	0.001
CIP	0(0)	0(0) 0(0)	0(0)	0(0) 0(0)	---	–	–	–	–	---

\*Expressed as percentage %(expressed in effective); \*\* R+I(%): R(%)/I(%); PG: Penicillin G, SXT: Trimethoprim-sulfamethoxazole; CHL: Chloramphenicol; LEV: Levofloxacin; RIF: Rifampin; TET: Tetracycline; ERY: Erythromycin; CTX: Cefotaxime; CRO: Ceftriaxone; AMX: Amoxicillin; CIP: Ciprofloxacin; MIC: Minimum inhibitory concentration.

<sup>a</sup>60 isolates tested for *S. pneumoniae*.

<sup>b</sup>64 isolates tested for *S. pneumoniae*.

<sup>c</sup>63 isolates tested for *S. pneumoniae*.

In Morocco, the pneumococcal PCV-13 vaccine was introduced into the national immunization program in October 2010 and subsequently replaced by PCV-10. Despite the lack of serotyping of the *S. pneumoniae* included in this study, there was a decrease and stability in the rate of pneumococcal meningitis disease during the periods studied after the introduction of the anti-pneumococcal vaccine in the immunization schedule. In a local study in Casablanca that covered the period before and after the implantation of PCV-10 and PCV-13, the incidence rate of invasive pneumococcal infections associated with vaccine serotypes decreased after the vaccine was introduced in children less than 2 years of age [13]. Impact of replacing PCV-13 vaccine with PCV-10 was not influencing and both contribute to the reduction of invasive infections in children fewer than 5 years of age, when they were implanted in national immunization programs of countries [12, 14, 15].

Currently, two vaccines are developed and available for active immunization against the *N. meningitidis* B serogroup [16, 17], but in Morocco the meningococcal B vaccine wasn't used yet and if this serogroup is identified, prevention will be limited to chemoprophylaxis. However, vaccination with

tetravalent vaccine ACYW135 became mandatory for all Hajjis and visitors to Mecca.

United Kingdom (UK) was the first country to introduce *N. meningitidis* B vaccine in 2015 for the ages of 2, 4, and 12 months, the impact of the introduction of *N. meningitidis* B vaccine in the UK infant vaccination program reduced the incidence of meningitis due to NMB [18].

Amoxicillin-resistant strains of *N. meningitidis* in this study were 5.6%. Whereas in previous studies at the local researches, no strains were found to be resistant to amoxicillin, while the *N. meningitidis* DSPG level is 9.3% in this study, which has changed over time in relation to these researches [19, 20]. Thus, the resistance reported here remained less severe than those reported elsewhere [18, 20]. In our study, we recorded resistance to cefotaxime, ceftriaxone, and rifampin, this information remained to our knowledge the first resistance declared for *N. meningitidis* in Morocco and the North African Maghreb countries [21, 22]. Resistance to chloramphenicol was similar to the rate in Tunisia [22]. And moreover resistance of rifampin was declared in a country of Latin America (Brazil) [23], while in the African meningitis belt the reported results are below ours [24]. The phenotypic profile of *N. meningitidis* DSPG

or PG-resistant is associated with mutations in the *penA* gene responsible for this reduced susceptibility to PG [25]. Therefore, one study has shown that increased resistance to C3G was observed in strains of decreased susceptibility to penicillin hosting a new modified *penA* gene allele “*penA327*”. These strains containing “*penA327*” allele belonged to the CC11 clonal complex and belonged to serogroup B and C [26]. In our study, *N. meningitidis* DSPG isolates were associated with co-resistance with chloramphenicol and rifampin ( $P < 0.05$ ) and resistance to third-generation cephalosporin – C3G – (cefotaxime and ceftriaxone) and amoxicillin ( $P < 0.001$ ).

Strains of *S. pneumoniae* DSPG in our study recorded a rate of 26.2%. This rate was lower than that reported in a local study and other countries, which were conducted prior to the introduction of PCV vaccine [6, 21, 27]. Isolates resistant to amoxicillin were 7.7%, while C3G was 4.6% reduced susceptibility in our study and one strain 1.5% (1/65) was resistant. These rates appeared a little more than for amoxicillin and less than for C3G those reported in a other Moroccan study [27] and less than those reported elsewhere for both antibiotics [6, 21]. With the exception of cotrimoxazole (SXT), the observed resistance to chloramphenicol 9.2% and erythromycin 12.7% were more or less equivocal to those of the Moroccan study [27]. The resistance to levofloxacin presented here was already described in a study in Morocco that reported a rate of resistance of 1.2% [28]. However, the resistance to rifampin was 3.1%; this resistance remained to our knowledge the only rate so far reported in the Kingdom of Morocco and in the north of Africa. The results of our study noted a co-resistance between PG and beta-lactam antibiotics in *S. pneumoniae*, this association was statistically significant ( $P < 0.05$ ).

The proportions of resistance reported in this study for *S. pneumoniae* appear generally to be less moderate and stable than the proportions exposed here later in Morocco or elsewhere. This finding would be the result of the introduction of the pneumococcal vaccine PCVs into the national immunization program in 2010, as it was an additional benefit for reducing antibiotic resistance of *S. pneumoniae* [29].

Except one strain that was resistant to erythromycin, rifampin, and cotrimoxazole, the high susceptibility to antibiotics and the rarity of *H. influenzae type b* strains observed here was the result of the introduction of anti-*H. influenzae type b* vaccine into the national vaccination program [12].

For the treatment of syndromic bacterial meningitis, the national meningitis program recommended the use of cephalosporin as much as empirical treatment for the management of cases of confirmed bacterial meningitis and for probable bacterial meningitis [5]. In this study, the results of antibiotic susceptibility obtained for all isolates to cefotaxime, ceftriaxone, and vancomycin are very high; despite the C3G resistance observed in this study, which remained very rare. In addition, our study recommended the continued use of this empirical treatment for children, adolescents, and adults in the national meningitis program with surveillance of the development of resistance.

Chemoprophylaxis in Morocco was based on taking Rifampin 200 mg twice in two days for the eradication of *N. meningitidis* oropharyngeal carried for contacts cases [5]. Oropharyngeal portage of *N. meningitidis* varies by age and geographic area, it must be monitored laboratory-based to map the distribution of serogroup of *N. meningitidis* and determine the current potential need for vaccination [30, 31].

The resistance to rifampin observed in this study for the three isolates may have evolved as a result of the use of this prophylaxis for contact cases of *N. meningitidis* [32] and/or could be due to the use of rifampin in the treatment of tuberculosis patients in Morocco which remained a public health problem in the Kingdom [33]. Phenotypic studies and genotypic studies are useful for monitoring serogroup distribution and also assist national vaccination programs in the choice of vaccine to introduce [34].

All this would result in a review of the prophylactic vision in the contacts either by an adequate vaccine intervention or moderation in chemoprophylaxis with monitoring of developments of meningitis disease in these patients.

## CONCLUSION

In conclusion, syndromic bacterial meningitis of *N. meningitidis*, *S. pneumoniae*, and *H. influenzae type b* isolates in Morocco appeared to have a high susceptibility to antibiotics used for empirical treatment and chemoprophylaxis despite the appearance of some resistance to amoxicillin, rifampin, and cephalosporin which remained very low and very rare. The anti-*H. influenzae type b* and PCVs vaccination introduced into the national immunization program reflected good efficacy. For the *N. meningitidis* vaccine, this needs to be revised and adapted to the current situation. In the end, monitoring the evolution of antibiotic resistance in these three isolates and monitoring the effectiveness of long-term vaccines are necessary for disease control and for the adequacy of treatment and prophylaxis over time.

*Conflicts of interest:* The authors declare no conflict of interest.

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