

*Streptococcus pneumoniae* to 22 selected antimicrobial agents.

<i>S. pneumoniae</i> isolates phenotype/genotype (no. isolates)				
<i>ermB</i> (n=157)	<i>mefE</i> (n=58)	<i>terM</i> (n=240)	Levofloxacin non-susceptible (n=25)	TMP/SMX-resistant (n=50)
MIC <sub>50</sub> MIC <sub>90</sub> Range				
0.25, 0.008-0.5	0.015, 0.125, 0.004-0.5	0.015, 0.25, 0.004-0.5	0.06, 0.25, 0.008-0.5	0.015, 0.125, 0.004-0.5
0.5, 2, <0.015>8	0.06, 1, <0.015>2	0.12, 2, <0.015>4	<0.015, 2, <0.015>2	0.5, 2, <0.015>8
0.5, 2, <0.06>16	<0.06, 2, <0.06>4	0.25, 2, <0.06>16	<0.06, 2, <0.06>16	0.5, 4, <0.06>16
0.2, 2, <0.12>16	<0.12, 2, <0.12>4	<0.12, 2, <0.12>16	0.25, 2, <0.12>16	0.5, 4, <0.12>16
>64, <0.12>64	0.5, 32, <0.12>64	1, >64, <0.12>64	8, 32, 0.12-32	16, 32, <0.12>64
0.2, 0.25, <0.03>0.4	4, >8, <0.03>8	1, >8, <0.25>8	1, 0.25, <0.25>8	1, >8, <0.25>8
1, 8, <0.06>8	0.25, 4, <0.06>8	0.25, 4, <0.06>8	0.25, 4, <0.06>8	2, 8, <0.06>16
16, <0.12>16	0.5, 8, <0.12>16	0.5, 16, <0.12>16	0.5, 16, <0.12>16	4, 16, <0.12>16
25, 2, <0.03>8	0.06, 2, <0.03>8	0.06, 2, <0.03>8	0.06, 2, <0.03>8	1, 2, <0.03>8
4, >4, 0.12>4	0.25, >4, 0.03>4	1, >4, <0.03>4	>4, >4, 0.25>4	>4, >4, <0.03>4
>4, >4, 1>4	>4, >4, 0.5>4	>4, >4, 0.5>4	>4, >4, 1>4	>4, >4, 1>4
12, 1, <0.004>2	0.06, 1, <0.004>2	0.06, 1, <0.004>2	0.06, 1, <0.004>1	0.25, 1, <0.004>4
16, <0.12>32	1, 8, <0.12>16	1, 16, <0.12>32	1, 8, <0.12>16	4, 16, <0.12>32
23, 1, <0.03>8	<0.03, 1, <0.03>8	0.12, 1, <0.03>8	0.12, 1, <0.03>1	0.5, 2, <0.03>8
1, 2, 0.12>4	1, 2, 0.12>4	1, 2, 0.12>4	4, 4, 4>4	1, 2, 0.06>4
1, 1, <0.25>2	1, 1, <0.25>2	1, 1, <0.25>2	2, 4, 2>4	1, 1, <0.25>2
<32, >32, >32	>32, >32, >32	>2, >32, <0.12>32	1, >32, <0.12>32	0.5, >16, <0.12>16
>16, >16, >16	2, >16, <0.12>16	8, >16, <0.12>16	1, >16, <0.12>16	0.5, >16, <0.12>16
<32, >32, >32	1, >32, <0.12>32	>2, >32, <0.12>32	1, >32, <0.12>32	1, >32, <0.12>32
>8, >8, >8	<0.06, 1, <0.06>8	<0.06, 1, <0.06>8	<0.06, >8, <0.06>8	<0.06, >8, <0.06>8
>6, >16, <1>16	8, >16, <1>16	>16, >16, >16	<1, >16, <1>16	16, >16, 1>16
1, 4, <0.25>8	<0.25, 4, <0.25>8	0.5, 4, <0.25>8	2, 4, <0.25>4	>8, >8, >8

Table 2: Susceptibility of *Moraxella catarrhalis* to 23 selected antimicrobial agents.

Antimicrobial Agent	<i>M. catarrhalis</i> isolates, Phenotype/Genotype (no. isolates)		
	Beta-lactamase-negative BRO-1 (n=34)	Beta-lactamase positive BRO-2 (n=363)	Beta-lactamase positive BRO-2 (n=22)
	MIC <sub>50</sub> MIC <sub>90</sub> Range [mg/L]		
Faropenem	0.03, 0.25, 0.03-0.25	0.12, 1, 0.03-1	0.03, 0.25, 0.03-0.5
Penicillin	<0.03, 0.25, <0.03>0.5	4, >4, <0.03>4	1, >4, <0.25>4
Amoxicillin	<0.06, 0.25, <0.06>0.5	2, 8, <0.06>8	0.25, 8, <0.06>8
Amoxicillin-clavulanate	<0.25, 0.25, <0.25>0.5	0.25, 0.25, <0.25>4	0.25, 0.25, <0.25>0.5
Cefaclor	0.35, 2, 0.25-2	0.5, 1, <0.25>2	<0.25, <0.25>0.25-0.5
Lorsacarb	0.25, 2, <0.25>2	1, 2, <0.25>16	1, 1, 0.5-2
Cefuroxime	0.5, 1, <0.06>2	1, 2, <0.06>4	0.5, 1, 0.12-2
Cefprozil	0.5, 4, 0.12-4	2, 4, 0.12-4	2, 4, 0.5-4
Cefpodoxime	0.12, 0.5, <0.03>0.5	0.5, 1, <0.03>4	0.25, 0.5, 0.06-1
Cefazime	<0.03, 0.03, <0.03>0.5	0.12, 0.25, <0.03>2	0.06, 0.5, <0.03>0.5
Cefbutin	0.12, 1, <0.03>4	1, 4, <0.03>4	0.25, 4, 0.06-4
Cefotaxime	0.08, 0.5, <0.008>1	0.25, 0.5, <0.008>4	0.06, 0.5, 0.03-1
Ceftriaxone	0.25, 0.25, <0.25>2	0.25, 0.25, <0.25>8	0.25, 0.25, <0.25>2
Cefepime	0.12, 0.5, <0.06>2	0.5, 1, <0.06>4	0.25, 1, 0.12-4
Ciprofloxacin	<0.015, 0.03, <0.0015>0.25	<0.015, 0.03, <0.0015>0.5	<0.015, 0.03, <0.0015>0.06
Levofloxacin	<0.5, 0.5, <0.5>1	<0.5, 0.5, <0.5>1	<0.5, 0.5, <0.5>1
Erythromycin	<0.25, <0.25>0.25, <0.25>0.25	<0.25, <0.25>0.25, <0.25>0.25	<0.25, <0.25>0.25-2
Azithromycin	<0.12, <0.12>0.12, <0.12>0.12	<0.12, <0.12>0.12, <0.12>0.12	<0.12, <0.12>0.12-16
Clarithromycin	<0.25, <0.25>0.25, <0.25>0.25	<0.25, <0.25>0.25, <0.25>0.25	<0.25, <0.25>0.25-2
Tellithromycin	<0.06, 0.12, <0.06>0.5	<0.06, 0.12, <0.06>0.5	<0.06, 0.12, <0.06>0.5
Tetracycline	<2, <2>2, <2>2	<2, <2>2, <2>2	<2, <2>2, <2>2
Rifampin	11, 51, 51, 1	11, 51, 51, 1	11, 51, 51, 1
TMP/SMX	<0.25, 0.25, <0.25>8	<0.25, 0.25, <0.25>8	<0.25, 0.25, <0.25>8

compounds collected, 385 (92%) were beta-BRO-1 (n=363) and BRO-2 (n=22) isolates

of beta-lactamase production occurred with the incidence of beta-lactamase-83% to 98% in the 13 participating

was detected at similar levels in both strains, regardless of specimen source or

ing isolates were collected from 10 of cated in Belgium, France, Germany, Switzerland, and the UK

- The comparative *in vitro* activities of the 23 compounds tested are listed in Table 2
- With the exception of penicillin, ampicillin, and amoxicillin, all other beta-lactam antimicrobials tested had excellent activity against *M. catarrhalis*, regardless of beta-lactamase production
- Faropenem exhibited potent *in vitro* activities with MIC<sub>50</sub> values of 1 mg/L for BRO-1 and 0.25 mg/L for BRO-2 and beta-lactamase-negative isolates
- All quinolones tested demonstrated excellent *in vitro* activities (ie, maximum MIC of 1 mg/L)

## CONCLUSIONS

- Faropenem had potent *in vitro* activity against 530 genetically characterized *S. pneumoniae* isolates, including those which exhibited multiple-antibiotic resistance (ie, MIC<sub>50</sub> = 0.25 mg/L)
- Antimicrobial susceptibility patterns for *M. catarrhalis* have changed little in recent years in Europe
- Faropenem demonstrated potent *in vitro* activity against both BRO-1 and BRO-2 beta-lactamase-producing *M. catarrhalis* isolates (ie, MIC<sub>50</sub> values ≤ 1 mg/L)
- Faropenem, a promising novel oral agent, warrants further clinical investigation for the treatment of RTIs

## REFERENCES

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- Mortensen JE, Egleton JH. Comparative *in vitro* activity of faropenem against aerobic bacteria isolated from pediatric patients. *Diagn Microbiol Infect Dis* 1995; 22:301-6.