

BACKGROUND

In the late 90s, respiratory fluoroquinolones were introduced in the treatment of (atypical) pneumonia. However, there was concern that short-term treatment success might be followed by emerging fluoroquinolone resistant (FQ-R) *Streptococcus pneumoniae* strains by accumulation of chromosomal mutations. In recent years, the role of efflux in low-level FQ-R has become more and more appreciated. Apart from a moderate increase in MIC, increased efflux is associated with rising mutational frequencies in the QRDRs [1]. Most clinical relevance is attributed to the reserpine-sensitive heterogenic ABC efflux pump PatAB [2].

Although most *S. pneumoniae* surveillance studies focus on bacteremia, recent work estimated that for every adult bacteremia case there are three non-invasive infections [3]. Here, we present data on FQ-R in 5,602 non-invasive pneumococci, collected during winter seasons between 1995 and 2014 across 15 Belgian clinical laboratories.

MAIN OBSERVATIONS

(i) Rising resistance to older fluoroquinolones since 2011

All isolates were assessed on FQ-R using broth microdilution (Table 1)

- High-level CIP resistance significantly increased to 9.0% in 2013 (P = 0.00025, X² trend analysis incl. Bonferroni's correction).
- Levofloxacin MIC₅₀ increased significantly from 0.5 to 1 µg/mL since 2012 (P < 10⁻⁶).
- Moxifloxacin is the compound with the highest intrinsic activity; Resistance arises only sporadically, and remained <1% throughout the entire study period.

Table 1: Yearly percentage of isolates displaying indicated MIC (µg/mL) against three fluoroquinolones. The MIC₅₀ values are coloured orange.

CIPROFLOXACIN		Sensitive (%)				Intermediate resistant (%)				Resistant (%)					% res.
Year	# strains	0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	32			
1995	143	-	0.7	2.8	18.9	34.3	35.7	7.7	-	-	-	-	0.0		
1997	162	-	-	-	6.8	17.9	61.7	11.7	2.5	0.6	-	-	3.1		
1999	227	-	0.4	0.4	6.2	30.8	47.1	13.2	1.8	-	-	-	1.8		
2001	334	-	-	0.9	12.9	38	38	6.6	3	0.6	-	-	3.6		
2003	391	-	0.5	3.1	11.3	25.1	46.3	9.5	2.6	1.8	-	-	4.4		
2004	424	0.2	1.2	1.9	14.2	37.3	36.3	6.6	2.1	-	0.2	-	2.3		
2005	447	0.2	1.1	2.5	12.8	35.6	40.5	6	0.9	0.2	0.2	-	1.3		
2006	430	-	0.2	1.4	7.4	28.6	53.7	8.1	0.5	-	-	-	0.5		
2007	413	-	0.2	1.5	7.7	30	56.7	1.7	1.5	0.2	0.5	-	2.2		
2008	448	-	0.2	0.4	4.7	16.1	73.4	4.7	-	-	0.4	-	0.4		
2009	413	-	-	1.9	6.5	44.1	44.1	1.9	1	0.2	0.2	-	1.4		
2010	370	-	0.8	2.7	10.8	26.2	55.1	1.9	2.2	-	-	0.3	2.5		
2011	368	-	0.3	0.5	4.6	14.9	46.2	29.6	2.2	1.1	0.5	-	3.8		
2012	351	-	-	0.3	1.1	14.2	46.4	29.9	7.1	0.6	0.3	-	8.0		
2013	369	-	-	-	3	12.5	38.8	36.9	7.3	1.1	0.3	0.3	9.0		
2014	312	-	-	-	0.6	9.9	49.7	33	6.4	-	0.3	-	6.7		

LEVOFLOXACIN		Sensitive (%)				Resistant (%)				% res.		
Year	# strains	0.03	0.06	0.12	0.25	0.5	1	2	4		8	16
1995	143	-	1.4	2.8	19.6	40.6	30.1	5.6	-	-	-	0.0
1997	162	-	-	0.6	8	58.6	26.5	4.9	1.2	-	-	1.2
1999	227	-	0.4	-	2.6	37.9	44.1	13.2	1.8	-	-	1.8
2001	334	-	-	1.2	9	47.6	33.2	6.3	2.4	0.3	-	2.7
2003	391	-	0.5	3.6	13.6	31.7	41.4	5.9	1.8	1.5	-	3.3
2004	424	0.5	0.7	3.8	14.2	42.7	30.2	5.2	2.6	-	0.2	2.7
2005	447	0.9	2	4.5	22.6	48.1	15.9	5.4	0.4	-	0.2	0.6
2006	430	0.2	1.2	2.1	9.3	28.6	53.7	8.1	0.5	-	-	0.5
2007	413	0.2	0.5	2.2	13.8	58.1	23.5	0.7	0.2	0.7	-	0.9
2008	448	0.2	-	1.1	6.9	60.7	26.1	4.2	0.2	-	0.4	0.6
2009	413	-	1.2	5.3	30.8	46.2	15	0.7	0.2	0.5	-	0.7
2010	370	0.3	3.5	4.3	17	55.9	15.7	2.4	0.5	0.3	-	0.8
2011	368	0.3	0.5	3	10.1	37	41.3	6.8	0.5	0.5	-	1.0
2012	351	-	-	0.9	3.7	41.3	39	12	2.8	0.3	-	3.1
2013	369	-	-	1.4	2.7	35	49.3	10.3	0.8	0.3	0.3	1.4
2014	312	-	-	0.6	2.2	30.8	59.6	6.1	0.3	-	0.3	0.6

MOXIFLOXACIN		Sensitive (%)				Resistant (%)					% res.		
Year	# strains	0.008	0.015	0.03	0.06	0.12	0.25	0.5	1	2		4	8
1995	143	-	9.1	33.6	38.5	13.3	0.7	-	-	-	-	-	0.0
1997	162	-	0.6	12.3	38.9	44.4	3.1	0.6	-	-	-	-	0.0
1999	227	0.4	1.8	11	40.1	30.4	12.8	2.6	0.9	-	-	-	0.9
2001	334	0.6	6.3	9.3	43.7	32.3	5.4	1.5	0.6	0.3	-	-	0.9
2003	391	1	6.6	13.6	30.2	36.8	10.5	0.8	-	0.3	0.3	-	0.6
2004	424	0.5	4.5	17	39.4	30.2	8	0.2	-	0.2	-	-	0.2
2005	447	1.1	4	18.6	39.6	28.2	6.9	1.3	-	0.2	-	-	0.2
2006	430	1.8	4.7	17	41.4	30.9	-	0.2	-	-	-	-	0.0
2007	413	0.7	2.9	11.1	43.1	30	11.4	-	0.5	0.2	-	-	0.7
2008	448	0.2	0.9	7.4	38.6	46.4	6.9	-	-	0.2	-	0.2	0.4
2009	413	0.2	5.3	11.1	51.3	25.2	6.3	0.2	0.2	-	-	-	0.2
2010	370	-	5.4	11.6	49.5	26.8	5.7	0.8	-	-	0.3	-	0.3
2011	368	0.3	3.3	12.8	48.9	27.2	6.5	0.5	0.5	-	-	-	0.5
2012	351	-	2.3	5.4	48.1	36.5	6.6	0.9	0.3	-	-	-	0.3
2013	369	-	1.4	9.5	53.9	30.1	4.3	-	-	0.8	-	-	0.8
2014	312	-	0.3	8.3	52.6	34.3	4.2	-	-	0.3	-	-	0.3

TAKE-HOME MESSAGES

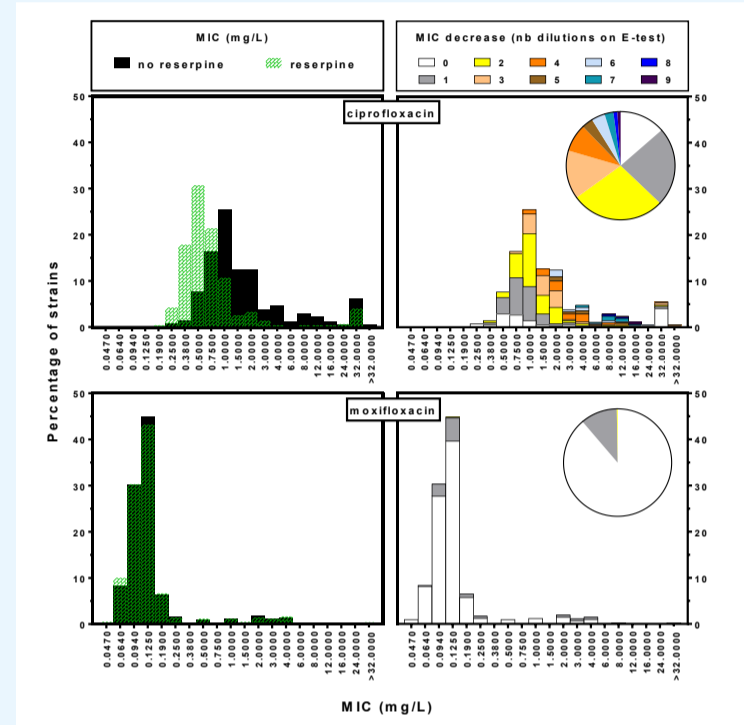
- From 2011 onwards, we observe a trend towards increased resistance to ciprofloxacin and (although only visible at the MIC₅₀ level) levofloxacin. Although a similar increase in ciprofloxacin resistance was reported in Canada, this is not confirmed in other European, American or Asian surveillance studies.
- The preference of first-step mutations in ParC is reflected by the 4:1 ratio of single QRDR mutations in the Topoisomerase IV subunits compared to the DNA Gyrase.
- While target topoisomerase mutations and efflux pump (over)expression clearly contribute to FQ-R, we add novel isolates to the existing collection of strains deprived of known molecular mechanisms of fluoroquinolone resistance. It would be of great value to unravel their resistance mechanisms through profound, comparative molecular characterization at the genomic, transcriptomic and proteomic level.

(ii) Efflux is pivotal in low-level ciprofloxacin resistance

- For 229 out of 422 isolates (54.2%), at least a twofold decrease in MIC₅₀ was achieved upon addition of 20mg/L of the efflux pump inhibitor reserpine.
- Moxifloxacin is much less sensitive to reserpine.
- Overexpression of the ABC pump PatAB was found in 48% of reserpine-sensitive strains, but could only be linked to disruptive terminator mutations in a fraction of these (data not shown).

Fig 1

MIC distribution of ciprofloxacin and moxifloxacin, based on E-test of 422 non-invasive *S. pneumoniae* isolates. Left-hand panels: MIC distributions determined in the absence (control; black) or presence (green) of 20 mg/L reserpine. Right-hand panels: reduction of MIC (in blocks of 0.5 log₂ dilutions from 0 to 3 log₂ dilutions) after addition of 20 mg/L reserpine and plotted as a function of the MIC distribution of the isolates in the absence of reserpine.



(iii) Topoisomerase mutations explain a lot in high-level resistance, but not all

- Classical topoisomerase mutations in *gyrA* (n=25), *parC* (n=46) and *parE* (n=3) were found in varying combinations, arguing against clonal expansion of FQ-R.
- Marginal impact of recombination with co-habiting commensal streptococci on FQ-R (10.4%).
- A rare combination of DNA Gyrase mutations (*GyrA_S81L/ GyrB_P454S*) suffices for high-level moxifloxacin resistance, which contrasts the current model.
- We encountered a number of isolates displaying high-level ciprofloxacin resistance, but are deprived of any known molecular resistance mechanism.

Table 2

Overview of the various FQ-R genotypes encountered in 422 clinical pneumococcal strains. Signature residues of the viridans group of streptococci are shaded in grey.

MIC _{CIP+R} (µg/ml)	MIC _{MOX+R} (µg/ml)	No. isolates	S81	E85	S114	P454	S52	N91	ParC D78	S79	D83	ParE D435	
< 1 (n=311)	0.064-0.19	289											
		10			G								
		6			G			D					
		2						D					
		2						D					
		1				G							
		1			F								
		1											
≤ 2 (n = 80)	0.125-0.25*	47											
		12								F			
		1							N				
		1	F										
		1						D			F		
		1										N/Y	
		1									F		K
		1				G			D		F		
		5				G							
		5				G			D				
		4				G			D				
2-4 (n = 12)	0.19-1	3											
		4									Y/F		
		1	F									N	
		1						D			F		
		1										G	
		1				G							
		1				G			D				
> 4 (n = 29)	1-32	1											
		3	F										
		1	F									N	
		12	F							F/Y			
		2	F								G/Y		
		2			G			D		F			
		1	Y		G			D		Y			
		1	G										
		2									N		
		2									F/Y		
1		K							Y				
1	L		G		S								

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