


# Interaction between *Streptococcus pneumoniae* and *Staphylococcus aureus* in paediatric patients suffering from an underlying chronic disease

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Susanna Esposito,<sup>1</sup> Gian Luigi Marseglia,<sup>2</sup> Carla Colombo,<sup>3</sup>  
Lorenzo Iughetti,<sup>4</sup> Leonardo Terranova,<sup>1</sup> Valentina Ierardi,<sup>1</sup>  
Monia Gambino<sup>1</sup> and Nicola Principi<sup>1</sup> for the Italian  
Pneumococcal Study Groups on Asthma, Cystic  
Fibrosis and Diabetes

## Abstract

Little is known about the interaction between *Streptococcus pneumoniae* and *Staphylococcus aureus* in school-age children and adolescents suffering from an underlying chronic disease. To increase our knowledge in this regard, an oropharyngeal swab was obtained from school-age children and adolescents suffering from asthma ( $n = 423$ ), cystic fibrosis (CF) ( $n = 212$ ) and type I diabetes mellitus (DMI) ( $n = 296$ ). *S. pneumoniae* detection and serotyping were performed using a real-time polymerase chain reaction, and *S. aureus* detection was performed using the RIDAGENE MRSA system. Among asthmatic, CF and DMI patients, both pathogens were identified in 65/423 (15.4%), 21/212 (9.9%) and 62/296 (20.9%) children, respectively; *S. pneumoniae* alone was identified in 127/434 (30.0%), 21/212 (9.9%) and 86/296 (29.1%), respectively; *S. aureus* alone was identified in 58/434 (13.7%), 78/212 (36.8%) and 49/296 (16.6%), respectively. *S. pneumoniae* colonisation rates were higher in younger children and declined with age, whereas the frequency of *S. aureus* colonisation was quite similar in the different age groups. Among asthmatic and CF patients aged 6–9 years, *S. aureus* carriage was significantly higher in children who were positive for *S. pneumoniae* ( $P < 0.05$ ). No significant association emerged between *S. aureus* carriage and carriage of *S. pneumoniae* serotypes included in the pneumococcal conjugate vaccines (PCVs). This study shows for the first time that school-age children and adolescents with asthma, CF and DMI are frequently colonised by *S. pneumoniae* and *S. aureus* and that no negative relationship seems to exist between these pathogens. Moreover, the supposed protection offered by PCV administration against *S. aureus* colonisation was not demonstrated.

## Keywords

pneumococcal colonisation, pneumococcal conjugate vaccine, staphylococcal colonisation, *Staphylococcus aureus*, *Streptococcus pneumoniae*

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<sup>1</sup>Pediatric Highly Intensive Care Unit, Department of Pathophysiology and Transplantation, Università degli Studi di Milano, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, Milan, Italy

<sup>2</sup>Pediatric Clinic, University of Pavia, IRCCS Policlinico "S. Matteo" Foundation, Pavia, Italy

<sup>3</sup>Cystic Fibrosis Center, Lombardia Region, Department of Pathophysiology and Transplantation, Università degli Studi di Milano, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, Milan, Italy

<sup>4</sup>Pediatric Clinic, University of Modena and Reggio Emilia, Modena, Italy

## Corresponding author:

Susanna Esposito, Pediatric Highly Intensive Care Unit, Department of Pathophysiology and Transplantation, Università degli Studi di Milano, Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico, Via Commenda 9, 20122 Milano, Italy.  
Email: susanna.esposito@unimi.it

## Introduction

The upper respiratory tract is the reservoir of a diverse community of commensal and potential pathogens. Pharyngeal colonisation is a dynamic process that in a balanced state has a major beneficial role for the human host.<sup>1</sup> However, an imbalance of the respiratory microbial community can be the basis for developing invasive and mucosal diseases due to an overgrowth of pre-existing pathogens or the acquisition of new infectious agents. *Streptococcus pneumoniae* and *Staphylococcus aureus* are common commensals of the upper respiratory tract and among the major causes of bacterial infections in infants and children when imbalance occurs.<sup>2,3</sup> A better understanding of the incidence of carriage and the relationship between these two pathogens, including their potential for mutual interference, is needed to evaluate the epidemiology of the disease caused by each of them and the impact of related preventative measures, including the pneumococcal conjugate vaccines (PCVs).

A number of studies have investigated the colonisation of both healthy and ill children by *S. pneumoniae* and *S. aureus*, but these studies have reported conflicting results.<sup>4-8</sup> However, most of these studies were carried out in healthy younger children.<sup>9</sup> Little is known about the interaction between these pathogens in school-age children and adolescents suffering from an underlying chronic disease although in some cases an increased risk of pneumococcal or staphylococcal infection in patients with asthma<sup>10</sup> or cystic fibrosis (CF)<sup>11</sup> is well known. Moreover, no data are available on the impact of vaccination with PCVs, given several years before, on combined *S. pneumoniae* and *S. aureus* colonisation of these subjects. To increase our knowledge in this regard, a study specifically designed to evaluate pneumococcal and staphylococcal pharyngeal colonisation rates in school-age children and adolescents suffering from asthma, CF and type 1 diabetes mellitus (DM1) was carried out.

## Materials and methods

### Swab collection

During the period 1 January 2014 to 30 June 2014, this study enrolled children and adolescents (age range, 6–17 years) suffering from documented asthma, CF and DM1 regularly followed in the

outpatient clinic of the University of Milan's Department of Pathophysiology and Transplantation and in a number of Department of Pediatrics sited in Pavia, Modena, Verona, Rome and Naples. The protocol was approved by the Ethics Committees of each participating centre. Moreover, written informed consent was obtained from the parent(s) or legal guardian(s) of each study participant and from participants aged older than 8 years. In asthmatic children, the characteristics of asthma were evaluated on the basis of the Global Initiative for Asthma criteria.<sup>12</sup> Only clinically stable patients with mild to moderate disease were considered suitable for enrolment. Subjects with CF had to be free from acute respiratory exacerbation. Finally, DM1 patients had to be in good metabolic equilibrium. All patients were clinically stable at the time of enrolment; patients with active respiratory infection, those with a chronic underlying disease other than those in the study, and those who had received antibiotic therapy during the previous 2 weeks were excluded. Data regarding the medical history of each subject, including vaccination, were collected by means of a questionnaire that was completed by parents/guardians for children <15 years and by the adolescents themselves for those ≥15 years. Children were considered fully vaccinated against *S. pneumoniae* if they have received three doses of the heptavalent pneumococcal conjugate vaccine (PCV7) in the first year of life or two doses in the second year or a single dose after the second until the age of 5 years, according to the recommendations of the Italian Ministry of Health ([http://www.salute.gov.it/portale/documentazione/p6\\_2\\_2\\_1.jsp?lingua=italiano&id=543](http://www.salute.gov.it/portale/documentazione/p6_2_2_1.jsp?lingua=italiano&id=543)).

In each centre, swabbing was carried out by a group of experienced paediatric nurses supervised by a paediatrician. The collection of pharyngeal secretions was performed by means of an oropharyngeal swab because recent studies have suggested that oropharyngeal samples are better for determining *S. pneumoniae* carrier status in school-age children and adolescents<sup>13</sup> and may give information regarding *S. aureus* colonisation rates not substantially different from those determined by traditional anterior nares samples.<sup>13</sup>

Oropharyngeal swabs were obtained using the ESwab kit and a polypropylene screw-cap tube with an internal conical shape filled with 1 mL liquid Amies medium (cat. Number 480CE, Brescia, Copan, Italy). Sampling was carried out using a

tongue spatula to press the tongue downward to floor of the mouth and swabbing both the tonsillar arches and the posterior pharynx, without touching the sides of the mouth. All swabs were immediately transported to a central laboratory and processed within 2 h for the identification of *S. pneumoniae* and *S. aureus*.

### *S. pneumoniae* and *S. aureus* identification

For *S. pneumoniae* identification, bacterial genomic DNA was directly extracted from the samples using a NucliSENS easyMAG automated extraction system (BioMérieux, Bagno a Ripoli, Florence, Italy), a 250 µL sample input and a generic protocol and was tested for the autolysin-A-encoding gene (*lytA*) and the *wzg* (*cpsA*) gene of *S. pneumoniae* by means of real-time PCR (RT-PCR) as previously described.<sup>14</sup> Each sample was tested in triplicate and was considered positive if at least two of the three tests revealed the presence of both genes. The level of detection of the test was 16 genome copies. All positive cases were serotyped using primers and probes designed on the basis of the GenBank database sequences (www.ncbi.nlm.nih.gov) of serotypes 1, 3, 4, 5, 6A, 6B, 7F, 9V, 14, 18C, 19A, 19F and 23F (i.e. those in the 13-valent pneumococcal conjugate vaccine [PCV13]) and synthesised by TIB Molbiol (Genoa, Italy) as previously described.<sup>14</sup> Analytical specificity was pre-evaluated by means of computer-aided analyses using Primer-BLAST (www.ncbi.nlm.nih.gov/tools/primer-blast) and BLAST (www.blast.ncbi.nlm.nih.gov/Blast.cgi) software to compare the sequences with all listed 'bacteria' and 'homo sapiens' sequences.

*S. aureus* was identified using the RIDAGENE MRSA system (R-Biopharm AG, Darmstadt, Germany). The RIDAGENE MRSA system is a multiplex real-time PCR for the direct, qualitative detection and differentiation of methicillin-resistant *S. aureus* (MRSA) and methicillin-sensitive *S. aureus* (MSSA). An internal control is added to the samples during extraction to determine possible PCR inhibition or DNA extraction failure. After DNA isolation, the *mecA/mecC* gene, *SCCmec/orfX* junction (type I, II, III, IV, V, VI, VII, IX and XI) and the *orfX* gene are amplified by TaqMan according to the manufacturer's instructions using the Agilent Stratagene Mx3005P real-time PCR platform. The samples were evaluated as follows: negative, no

amplification signal but internal control DNA positive; MRSA, positive for *mecA/mecC*, the *SCCmec/orfX* junction, and *orfX*; MSSA, positive for both the *SCCmec/orfX* junction and *orfX* or only *orfX*. The RIDAGENE MRSA system has a limit of detection of  $\leq 5$  DNA copies per reaction.

### Statistical analysis

A contingency table analysis with the chi-squared or Fisher's exact test, as appropriate, was used to compare differences between groups. For an easier evaluation of carriage modifications in the studied population, this was arbitrarily subdivided into three age groups (6–9 years, 10–14 years and  $\geq 15$  years). Subgroup analyses were performed based on age and PCV7 vaccination status. All tests were two-sided, and a *P* value  $< 0.05$  was considered statistically significant. Data were analysed using SAS, version 9.2 (SAS Institute, Cary, NC, USA).

## Results

### Asthmatic patients

The study enrolled 423 participants with asthma (70.9% boys; median age, 10.6 years). Among them 176 (41.6%) were aged between 6 and 9 years, 207 (48.9%) were aged between 10 and 14 years and 40 (9.5%) were aged  $\geq 15$  years. The carriage of *S. pneumoniae* and *S. aureus* according to age and PCV7 vaccination coverage is shown in Table 1. A total of 173 (40.9%) patients did not carry either *S. pneumoniae* or *S. aureus*. Both pathogens were identified in 65 (15.4%) children, *S. pneumoniae* alone and *S. aureus* alone in 127 (30.0%) and 58 (13.7%), respectively. *S. pneumoniae* colonisation rates were higher in younger children and declined with age (52.8% in children aged 6–9 years, 43.5% in those aged 10–14 years and 22.5% in those aged  $\geq 15$  years). In contrast, the frequency of *S. aureus* colonisation was quite similar in all age groups (27.8%, 30.4% and 27.5% in children aged 6–9 years, 10–14 years, and  $\geq 15$  years, respectively). *S. aureus* carriage was significantly higher in children who were positive for *S. pneumoniae* (65/192, 33.9% vs. 58/231, 25.1%; *P* = 0.049). However, when carriage was evaluated according to the age of enrolled children, it was found that this difference remained statistically significant only for children aged 6–9 years (32/93, 34.4% vs. 17/83, 20.5%; *P* = 0.04).

**Table 1.** Oropharyngeal carriage of *Streptococcus pneumoniae* and *Staphylococcus aureus* in 423 asthmatic children, according to selected subgroups and *S. pneumoniae* serotypes.

<i>S. pneumoniae</i>	<i>S. aureus</i>		P value
	Negative	Positive <sup>a</sup>	
	n (row %)	n (row %)	
<i>All subjects</i>			
Negative	173 (74.9)	58 (25.1)	0.049
Positive	127 (66.1)	65 (33.9)	
<i>Age subgroups</i>			
6–9 years (n = 176)			
Negative	66 (79.5)	17 (20.5)	0.04
Positive	61 (65.6)	32 (34.4)	
10–14 years (n = 207)			
Negative	85 (72.6)	32 (27.4)	0.27
Positive	59 (65.6)	31 (34.4)	
15–17 years (n = 40)			
Negative	22 (71.0)	9 (29.0)	0.99
Positive	7 (77.8)	2 (22.2)	
<i>Vaccination with PCV7</i>			
Unvaccinated (n = 228)			
Negative	93 (72.1)	36 (27.9)	0.47
Positive	67 (67.7)	32 (32.3)	
Vaccinated (n = 195)			
Negative	80 (78.4)	22 (21.6)	0.03
Positive	60 (64.5)	33 (35.5)	
<i>Serotypes of PCV7<sup>b</sup></i>			
Positive for any PCV7 serotype	119 (66.5)	60 (33.5)	0.77
Negative for PCV7 serotypes, but positive for <i>S. pneumoniae</i>	8 (61.5)	5 (38.5)	
<i>Serotypes of PCV13<sup>b</sup></i>			
Positive for any PCV13 serotype	119 (65.4)	63 (34.6)	0.50
Negative for PCV13 serotypes, but positive for <i>S. pneumoniae</i>	8 (80.0)	2 (20.0)	
<i>Specific serotypes<sup>c</sup></i>			
Negative for serotype 3	283 (70.7)	117 (29.3)	0.75
Positive for serotype 3	17 (73.9)	6 (26.1)	
Negative for serotype 4	255 (71.2)	103 (28.8)	0.74
Positive for serotype 4	45 (69.2)	20 (30.8)	
Negative for serotype 5	277 (71.9)	108 (28.1)	0.14
Positive for serotype 5	23 (60.5)	15 (39.5)	
Negative for serotype 6A	296 (71.8)	116 (28.2)	0.02
Positive for serotype 6A	4 (36.4)	7 (63.6)	
Negative for serotype 9V	274 (71.5)	109 (28.5)	0.39
Positive for serotype 9V	26 (65.0)	14 (35.0)	
Negative for serotype 19A	296 (71.7)	117 (28.3)	0.07
Positive for serotype 19A	4 (40.0)	6 (60.0)	
Negative for serotype 19F	189 (73.8)	67 (26.2)	0.10
Positive for serotype 19F	111 (66.5)	56 (33.5)	

<sup>a</sup>Among *S. aureus* cases, 95.4% were methicillin-sensitive and 4.6% methicillin-resistant. Seven patients were positive for staphylococcal cassette chromosome mec.

<sup>b</sup>Among 192 subjects positive for *S. pneumoniae*.

<sup>c</sup>Only serotypes of *S. pneumoniae* for which at least 10 subjects were positive are presented in the table. PCV7, 7-valent pneumococcal conjugate vaccine; PCV13, 13-valent pneumococcal conjugate vaccine.

Among vaccinated subjects, *S. aureus* was identified more frequently in children who were positive for *S. pneumoniae* than in those who were negative (33/93, 35.5% vs. 22/102, 21.6%;  $P = 0.03$ ). In contrast, among unvaccinated subjects, the percentage of those colonised by *S. aureus* was not materially different in subjects positive and negative for *S. pneumoniae* (32/99, 32.3% vs. 36/129, 27.9%,  $P = 0.47$ ). Among 192 subjects positive for *S. pneumoniae*, no significant association emerged between *S. aureus* carriage and carriage of *S. pneumoniae* serotypes included in PCV7 ( $P = 0.77$ ) nor PCV13 ( $P = 0.50$ ).

### CF patients

A total of 212 CF patients were enrolled (48.1% boys; median age, 11.7 years). Among them, 63 (29.7%) were aged 6–9 years, 102 (48.1%) were aged 10–14 years and 47 (22.2%) were aged  $\geq 15$  years. The carriage of *S. pneumoniae* and *S. aureus* according to age and PCV7 vaccination coverage is shown in Table 2. A great number of patients were not colonised by the studied pathogens (92, 43.4%). Both pathogens were identified in 21 (9.9%) children, *S. pneumoniae* alone and *S. aureus* alone in 21 (9.9%) and 78 (36.8%), respectively. *S. pneumoniae* colonisation rates were higher in younger children and declined with age (28.6% in children aged 6–9 years, 17.6% in those aged 10–14 years and 12.8% in those aged  $\geq 15$  years). The frequency of *S. aureus* colonisation was slightly higher in children aged 10–14 years (51/102, 50.0%) in comparison with patients aged 6–9 years and  $\geq 15$  years (29/63, 46.0% and 19/47, 40.4%, respectively). *S. aureus* carriage was quite similar in children positive and negative for *S. pneumoniae* (21/42, 50.0% vs. 78/170, 45.9%;  $P = 0.63$ ). However, in younger children, colonisation by *S. aureus* was significantly higher in patients positive for *S. pneumoniae* than in those who were negative (12/18, 66.6% vs. 17/45, 37.8%;  $P = 0.04$ ).

Among vaccinated subjects, *S. aureus* was identified with similar frequency in children who were positive and negative for *S. pneumoniae* (3/10, 30.0% vs. 10/25, 40.0%;  $P = 0.71$ ). Similarly, among unvaccinated subjects, the percentage of those colonised by *S. aureus* was not materially different in subjects positive and negative for

*S. pneumoniae* (18/32, 56.3% vs. 68/145, 46.9%;  $P = 0.34$ ).

Among 42 subjects positive for *S. pneumoniae*, no significant association emerged between *S. aureus* carriage and carriage of the PCV7 and PCV13 *S. pneumoniae* serotypes, although the number of subjects positive for serotypes other than PCV7 was small ( $n = 3$ ).

### Diabetic patients

A total of 296 DM1 patients were enrolled (51.3% boys; median age, 12.7 years). Among them, 61 (20.6%) were aged 6–9 years, 154 (52.0%) were aged 10–14 years and 81 (27.4%) were aged  $\geq 15$  years. The carriage of *S. pneumoniae* and *S. aureus* according to age and PCV7 vaccination coverage is shown in Table 3. Approximately one-third of the patients were not colonised by the studied bacteria (99, 33.4%). Both pathogens were identified in 62 (20.9%) children, and *S. pneumoniae* alone and *S. aureus* alone were identified in 86 (29.1%) and 49 (16.6%), respectively. *S. pneumoniae* colonisation rates were higher in younger children and declined with age (62.3% in children aged 6–9 years, 53.9% in those aged 10–14 years and 33.3% in those aged  $\geq 15$  years). The frequency of *S. aureus* colonisation was quite similar in the three age groups even if slightly higher in children aged 6–9 years (25/61, 41.0%) in comparison with patients aged 10–14 years and  $\geq 15$  years (56/154, 36.4% and 30/81, 37.0%, respectively). *S. aureus* carriage was not significantly different between children positive and negative for *S. pneumoniae* (62/148, 41.9% vs. 49/148, 33.1%;  $P = 0.12$ ), even when different age groups were considered.

Among vaccinated subjects, *S. aureus* was identified more frequently in children who were positive for *S. pneumoniae* than those who were negative (26/56, 46.4% vs. 14/53, 26.4%;  $P = 0.03$ ). In contrast, among unvaccinated subjects, the percentage of those colonised by *S. aureus* was similar in subjects positive and negative for *S. pneumoniae* (35/88, 39.8% vs. 32/86, 37.2%;  $P = 0.73$ ).

Among 148 subjects positive for *S. pneumoniae*, no significant association emerged between *S. aureus* carriage and carriage of the PCV7 and PCV13 *S. pneumoniae* serotypes, although the number of subjects positive for serotypes other than PCV7 was small ( $n = 4$ ).

**Table 2.** Oropharyngeal carriage of *Streptococcus pneumoniae* and *Staphylococcus aureus* in 212 children with cystic fibrosis, also according to selected subgroups and *Streptococcus pneumoniae* serotypes.

<i>S. pneumoniae</i>	<i>S. aureus</i>		P value
	Negative	Positive <sup>a</sup>	
	n (row %)	n (row %)	
<i>All subjects</i>			
Negative	92 (54.1)	78 (45.9)	0.63
Positive	21 (50.0)	21 (50.0)	
<i>Age subgroups</i>			
6–9 years (n = 63)			
Negative	28 (62.2)	17 (37.8)	0.04
Positive	6 (33.3)	12 (66.7)	
10–14 years (n = 102)			
Negative	40 (47.6)	44 (52.4)	0.30
Positive	11 (61.1)	7 (38.9)	
15–17 years (n = 47)			
Negative	24 (58.5)	17 (41.5)	0.99
Positive	4 (66.7)	2 (33.3)	
<i>Vaccination with PCV7</i>			
Unvaccinated (n = 177)			
Negative	77 (53.1)	68 (46.9)	0.34
Positive	14 (43.7)	18 (56.3)	
Vaccinated (n = 35)			
Negative	15 (60.0)	10 (40.0)	0.71
Positive	7 (70.0)	3 (30.0)	
<i>Serotypes of PCV7<sup>b</sup></i>			
Positive for any PCV7 serotype	20 (51.3)	19 (48.7)	0.99
Negative for PCV7 serotypes, but positive for <i>S. pneumoniae</i>	1 (33.3)	2 (66.7)	
<i>Serotypes of PCV13<sup>b</sup></i>			
Positive for any PCV13 serotype	20 (50.0)	20 (50.0)	0.99
Negative for PCV13 serotypes, but positive for <i>S. pneumoniae</i>	1 (50.0)	1 (50.0)	
<i>Specific serotypes<sup>c</sup></i>			
Negative for serotype 5	104 (51.5)	98 (48.5)	0.02
Positive for serotype 5	9 (90.0)	1 (10.0)	
Negative for serotype 19F	95 (53.7)	82 (46.3)	0.81
Positive for serotype 19F	18 (51.4)	17 (48.6)	

<sup>a</sup>Among *S. aureus* cases, 91.9% were methicillin-sensitive and 8.1% methicillin-resistant. Six patients were positive for staphylococcal cassette chromosome mec.

<sup>b</sup>Among 42 subjects positive for *S. pneumoniae*.

<sup>c</sup>Only serotypes of *S. pneumoniae* for which at least 10 subjects were positive are presented in the Table. PCV7, 7-valent pneumococcal conjugate vaccine; PCV13, 13-valent pneumococcal conjugate vaccine.

## Discussion

Whether bacteria can colonize or not is determined by many ecological factors including the availability of resources (i.e. nutrients, space, attachment space), host immune responses and the presence of toxins or harmful substances.<sup>15</sup> Moreover, the interference between bacterial resident populations producing harmful substances (i.e. bacteriocins)<sup>15,16</sup> or inducing an immune

response<sup>17,18</sup> can play a role at this regard. In the case of *S. pneumoniae* and *S. aureus*, epidemiological studies showed that co-colonisation is rarer than expected and suggested that *S. pneumoniae* colonisation is reduced in presence of *S. pneumoniae*.<sup>19,20</sup> Knowledge of carriage characteristics of these pathogens, which is important in healthy children, is even more relevant in subjects with chronic underlying disease because applicable

**Table 3.** Oropharyngeal carriage of *Streptococcus pneumoniae* and *Staphylococcus aureus* in 296 children with type I diabetes mellitus, also according to selected subgroups and *S. pneumoniae* serotypes.

<i>S. pneumoniae</i>	<i>S. aureus</i>		P value
	Negative	Positive <sup>a</sup>	
	n (row %)	n (row %)	
All subjects			
Negative	99 (66.9)	49 (33.1)	0.12
Positive	86 (58.1)	62 (41.9)	
Age subgroups			
6–9 years (n = 61)			
Negative	15 (65.2)	8 (34.8)	0.44
Positive	21 (55.3)	17 (44.7)	
10–14 years (n = 154)			
Negative	51 (71.8)	20 (28.2)	0.051
Positive	47 (56.6)	36 (43.4)	
15–17 years (n = 81)			
Negative	33 (61.1)	21 (38.9)	0.63
Positive	18 (66.7)	9 (33.3)	
Vaccination with PCV7 <sup>b</sup>			
Unvaccinated (n = 174)			
Negative	54 (62.8)	32 (37.2)	0.73
Positive	53 (60.2)	35 (39.8)	
Vaccinated (n = 109)			
Negative	39 (73.6)	14 (26.4)	0.03
Positive	30 (53.6)	26 (46.4)	
Serotypes of PCV7 <sup>c</sup>			
Positive for any PCV7 serotype	83 (57.6)	61 (42.4)	0.64
Negative for PCV7 serotypes, but positive for <i>S. pneumoniae</i>	3 (75.0)	1 (25.0)	
Serotypes of PCV13 <sup>c</sup>			
Positive for any PCV13 serotype	83 (57.2)	62 (42.8)	0.26
Negative for PCV13 serotypes, but positive for <i>S. pneumoniae</i>	3 (100.0)	0 (0.0)	
Specific serotypes <sup>d</sup>			
Negative for serotype 4	166 (64.3)	92 (35.7)	0.09
Positive for serotype 4	19 (50.0)	19 (50.0)	
Negative for serotype 5	168 (62.7)	100 (37.3)	0.84
Positive for serotype 5	17 (60.7)	11 (39.3)	
Negative for serotype 9V	153 (61.7)	95 (38.3)	0.51
Positive for serotype 9V	32 (66.7)	16 (33.3)	
Negative for serotype 19A	176 (62.2)	107 (37.8)	0.77
Positive for serotype 19A	9 (69.2)	4 (30.8)	
Negative for serotype 19F	104 (66.7)	52 (33.3)	0.12
Positive for serotype 19F	81 (57.9)	59 (42.1)	

<sup>a</sup>Among *S. aureus* cases, 97.3% were methicillin-sensitive and 2.7% methicillin-resistant. Five patients were positive for staphylococcal cassette chromosome mec.

<sup>b</sup>Thirteen subjects were excluded due to missing information on PCV7 status.

<sup>c</sup>Among 148 subjects positive for *S. pneumoniae*.

<sup>d</sup>Only serotypes of *S. pneumoniae* for which at least 10 subjects were positive are presented in the Table. PCV7, 7-valent pneumococcal conjugate vaccine; PCV13, 13-valent pneumococcal conjugate vaccine.

information can lead to more appropriate prophylactic and therapeutic measures. Unfortunately, *S. pneumoniae* and *S. aureus* carriage and reciprocal interactions in children with severe chronic

underlying disease have received poor attention in recent years. Only HIV-infected children have been studied with conflicting results. In a first cross-sectional study, it was indicated that no

association between pneumococcal and *S. aureus* colonisation could be demonstrated.<sup>21</sup> More recently, a longitudinal study found that a lower prevalence of pneumococcal colonisation could lead to an increase in *S. aureus* presence in nasopharyngeal niche.<sup>22</sup> This study adds some new information regarding older children with diseases, such as asthma, CF and DM1, for whom several studies have evidenced an increased risk of development of both pneumococcal and staphylococcal infections and for whom, at least in some cases, colonisation with these pathogens was expected to have a role in favouring the development of disease itself or its worsening.

Asthmatic patients, independent of asthma severity, suffer from pneumococcal diseases more frequently than healthy subjects.<sup>10</sup> Pneumococcal colonisation in the neonatal period is associated with frequent wheezing episodes in later paediatric life.<sup>22</sup> Finally, *S. aureus* colonisation is more common in asthmatic patients<sup>23</sup> and is associated with wheeze and asthma,<sup>24</sup> possibly through the bacterial enterotoxin sensitisation.<sup>25</sup>

In CF patients, *S. aureus* is a common coloniser of the respiratory tract and one of the most common pathogens responsible of the infective exacerbations and progressive decline of lung function.<sup>11</sup> Moreover, it has been recently demonstrated that children with this disease are frequently carriers of *S. pneumoniae*,<sup>2</sup> and CF samples can have peculiar characteristics that can lead to increased virulence and significant lung damage.<sup>2</sup> Colonising pneumococci form well-organised biofilm communities in the nasopharyngeal environment, with bacteria that are resistant to commonly prescribed anti-pneumococcal antibiotics. Recent studies have shown that changes in the nasopharyngeal environment caused by concomitant virus infection, modifications in the microflora, inflammation or other host assaults trigger active release of pneumococci from biofilms.<sup>26,27</sup> These dispersed bacteria have distinct phenotypic properties and transcriptional profiles different from both biofilm and broth-grown, planktonic bacteria, resulting in a significantly increased virulence *in vivo*.<sup>27</sup>

The higher risk of pneumococcal infections, including IPD, in DM1 patients compared with healthy subjects has been well known for many years and was recently confirmed by the observation that, despite the introduction of vaccine prophylaxis, it remained three times higher in the UK

than in the general population.<sup>28–30</sup> Finally, the incidence of staphylococcal infections in DM1 patients is not marginal and some of them can be particularly severe and difficult to treat.<sup>31</sup>

Regarding pneumococcal colonisation, this study shows that in all the study groups it decreases with age. This is not surprising because a decline of pneumococcal colonisation from infancy to adolescence has been repeatedly reported in healthy children and mainly ascribed to the maturation of the immune system.<sup>8,32</sup> However, the incidence of colonisation in the studied children was significantly higher than that reported in other studies involving younger children.<sup>32</sup> Host immune characteristics and the greater sensitivity of the molecular methods used in this study in comparison with cultural methods used in most of the studies carried out in healthy children could explain this finding. However, this finding deserves attention because it supports the use of prophylactic measures against pneumococcal infections for school-age children and adolescents with asthma, CF and DM1.<sup>33</sup>

Previous PCV7 vaccination did not influence either absolute pneumococcal carriage or carriage of pneumococcal serotypes included in the conjugate vaccine or in PCV13. Other studies have evidenced that PCV administration has a relevant impact on colonisation, significantly reducing carriage of serotypes included in the administered vaccine.<sup>34</sup> However, these studies were carried out only a few months after vaccine use and the waning of vaccine-induced immunity with time could explain the finding, as already suggested by the data recently collected in healthy older children and adolescents.<sup>35</sup> On the other hand, this could be a problem because the waning of immunity against carriage could favour the development of diseases due to the initial elimination of the same serotypes and suggests the use of further booster vaccine doses to maintain protection.<sup>33</sup>

Regarding *S. aureus* colonisation, it cannot be forgotten that, from a theoretical point of view, the presence of *S. pneumoniae* in the pharynx should reduce *S. aureus* carriage. *S. pneumoniae* produces hydrogen peroxide that inhibits *S. aureus*<sup>36</sup> and can further reduce staphylococcal carriage through immune mediated mechanisms based on cross-reactive antibodies against common conserved dehydrogenases.<sup>37</sup> Moreover, the expression of phosphorylcholine and neuraminidase production by *S. pneumoniae* may contribute



to competitive effects between these bacterial species.<sup>38</sup> Finally, pneumococci containing pilus-islands have been negatively associated with *S. aureus* colonisation.<sup>39</sup>

However, in this study in which the incidence of *S. aureus* colonisation was in line with what was previously reported in healthy older children,<sup>13</sup> pneumococcal colonisation was not associated with a reduced *S. aureus* colonisation. In asthmatic children and in those with CF aged 6–9 years, pharyngeal positivity for *S. pneumoniae* was associated with an increase in *S. aureus* colonisation. In children with DM1, the incidence of *S. aureus* carriage was similar in patients with or without pneumococcal colonisation. Moreover, no impact of PCV7 administration was evidenced in any of the groups. These findings are in contrast with those reported in healthy subjects who displayed a negative relationship between *S. pneumoniae* and *S. aureus* and a positive effect of PCV7 administration in conditioning *S. aureus* carriage.<sup>35</sup> The lack of a negative interaction between the two pathogens is difficult to explain. Bacterial colonisation with a single infectious agent depends on several factors. Host defences, environmental factors, and carriage of other bacteria and/or viruses may play a role. The importance of direct bacterial effectors, viral-induced bacterial adhesion, viral-derived disruption of the respiratory epithelium, production of viral products, and interference with the host immune system in conditioning type and degree of colonisation has been demonstrated.<sup>38</sup> All of these factors were not evaluated, and it is possible that their role could have influenced *S. pneumoniae* and *S. aureus* colonisation of the studied children. The interference of other unknown factors in conditioning carriage is also supported by the data regarding PCV7 administration and *S. aureus* colonisation, which contrast with the data from in healthy younger children. Previous studies have found that younger children who were colonised by primarily PCV7 serotypes were less frequent carriers of *S. aureus* compared with subjects colonised with other serotypes,<sup>9</sup> suggesting that the serotypes included in PCV7 could protect against *S. aureus* colonisation. In this study, children who had received PCV7 several years before were colonised by the same serotypes included in the vaccine, and despite this vaccination, they were frequently colonised by *S. aureus*. The waning of immune protection against pneumococcal colonisation some years after the

last vaccine dose has been already demonstrated in healthy children<sup>35</sup> and can be the cause of the persistence of carriage of PCV7 serotypes in the children enrolled in this study despite previous vaccination. Finally, with very few exceptions, *S. aureus* was not associated with any of the serotypes included in both PCV7 and PCV13, confirming that in this study *S. pneumoniae* and *S. aureus* colonisation were completely independent.

In conclusion, this study shows for the first time that school-age children and adolescents with asthma, CF and DM1 are frequently colonised by *S. pneumoniae* and *S. aureus* and that no negative relationship seems to exist between these pathogens. It is not known whether this colonisation depends on the immune characteristics and the infectious history of these patients or from other previously unidentified factors. Moreover, the supposed protection offered by PCV7 administration against *S. aureus* colonisation was not demonstrated and no substantial association between single pneumococcal serotypes and *S. aureus* was evidenced. Further studies are needed to clarify the relationships between bacterial pathogens and the role of PCV in conditioning staphylococcal colonisation.

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