

## **Long-term impact of 10-valent pneumococcal conjugate vaccination on invasive pneumococcal disease among children in Finland**

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## **Abstract**

### **Background**

The ten-valent pneumococcal conjugate vaccine (PCV10) was introduced into the Finnish National Vaccination Programme (NVP) in September 2010. The impact of PCV10 vaccination against invasive pneumococcal disease (IPD) in vaccine-eligible children has been high, possibly due to cross-protection against PCV10-related serotypes (serotypes in the same serogroups as the PCV10 types) in addition to protection against PCV10 serotypes. We evaluated the long-term impact of PCV10 vaccination against IPD in vaccine-eligible and older, unvaccinated children six years after PCV10 introduction with a special focus on cross-protection.

### **Methods**

We used data on IPD from the national, population-based surveillance. A target cohort of vaccine-eligible children (born June 2010 or later) was followed from 3 months of age until the end of 2016. To assess the indirect effect, another cohort of older, PCV10-ineligible children was followed from 2012 through 2016. IPD rates were compared with those of season- and age-matched reference cohorts before NVP introduction.

### **Results**

Among vaccine-eligible children, the incidence of all IPD decreased by 79% (95%CI 74 to 83%). There was a statistically significant reduction in the incidence of 6A IPD, but for 19A, the reduction was non-significant and the incidence of 19A increased towards the end of the study period in the older vaccine-eligible children. The increase in non-PCV10 related serotypes was non-significant.

In the unvaccinated older children, the incidence of all IPD decreased by 33% (95%CI 8 to 52%) compared to the reference cohort. There was no impact on serotype 6A or 19A IPD in the unvaccinated cohort.

## **Conclusion**

Overall, the impact of PCV10 vaccination on IPD was very high in vaccine-eligible children, with a major reduction in vaccine-type disease, and without notable replacement by other serotype groups. Our data suggest that PCV10 results in long-lasting direct cross-protection against 6A IPD. For 19A, no net reduction was observed six years after NVP introduction in the vaccine-eligible cohort. The indirect impact on IPD in unvaccinated children sustained.

## 1 **Introduction**

2 It has been estimated that *Streptococcus pneumoniae* (the pneumococcus) caused about 14.5 million  
3 episodes of serious pneumococcal disease, including pneumonia, meningitis and febrile  
4 bacteraemia, worldwide in 2000.<sup>1</sup> The introduction of pneumococcal conjugate vaccines (PCVs)  
5 since 2000 has afforded excellent direct protection for vaccinated children against invasive  
6 pneumococcal disease (IPD) caused by serotypes included in the vaccines. At the same time, the  
7 impact has extended to unvaccinated populations through indirect protection due to reduced  
8 vaccine-type carriage in vaccinated children and the subsequently reduced transmission. However,  
9 replacement in carriage and subsequent disease by serotypes not included in the vaccines has partly  
10 eroded the direct and indirect benefits of vaccination across all age groups and is a growing  
11 concern.<sup>2-8</sup>

12 In addition to protection against the vaccine serotypes, PCVs have been documented to provide  
13 cross-protection against some vaccine-related serotypes, i.e. serotypes that belong to the same  
14 serogroups as the vaccine serotypes. Previous studies have shown that the seven-valent vaccine  
15 (PCV7, Prevenar, Pfizer), which included serotypes 4, 6B, 9V, 14, 18C, 19F, and 23F, reduced also  
16 IPD caused by 6A IPD, but not by 19A or 6C.<sup>3,9,10</sup> Serotype 19A IPD was among the most common  
17 causes of replacement disease after PCV7 introduction.<sup>3,4,11,12</sup> In 2010, PCV13 (Prevenar 13, Pfizer)  
18 with six additional serotypes (1, 3, 5, 6A, 7F, and 19A) replaced PCV7. Recently, cross-protection  
19 against 6C from the 6A component of PCV13 has been reported<sup>13</sup>.

20 The 10-valent pneumococcal conjugate vaccine (PCV10, Synflorix, GSK Vaccines) including  
21 serotypes in the PCV7 + 1, 5 and 7F was introduced into the Finnish National Vaccination  
22 Programme (NVP) after a public tender in September 2010. In addition to providing protection  
23 against vaccine serotypes, PCV10 also had reduced incidence of 6A and 19A IPD among vaccine-  
24 eligible children in Finland three years after vaccine introduction into NVP.<sup>6</sup> Here we report

25 updated data about the overall and indirect long-term impact of PCV10 vaccination against IPD  
26 among vaccine-eligible and unvaccinated older children, with a special focus on cross-protection  
27 against the vaccine-related serotypes.

## 28 **Methods**

### 29 *The vaccination programme*

30 PCV10 was introduced into the NVP in September 2010. Children born on or after June 1, 2010,  
31 are eligible for vaccination with a 2+1 schedule (primary series at 3 and 5 months and a booster  
32 dose at 12 months of age). There was no catch-up vaccination at PCV10 introduction. In the birth  
33 cohort of 2012, the uptake of at least one dose of PCV10 was estimated at 94% based on data in the  
34 National Vaccination Register.<sup>14</sup>

35 In 2009-2011, the effectiveness of PCV10 against vaccine-type IPD was investigated in a nation-  
36 wide FinIP vaccine trial, in which over 30,000 children born 2008 through May 2010 were  
37 vaccinated with PCV10.<sup>15</sup> During the FinIP trial, the PCV10 vaccine coverage varied regionally  
38 from 0 to 60%. Prior to the FinIP trial and the introduction of PCV10 into the NVP, no  
39 pneumococcal conjugate or polysaccharide vaccines were routinely used for healthy children and  
40 adults, and the vaccine uptake was estimated to be less than 2%.

41 Influenza vaccine was introduced into the NVP in the beginning of the influenza season 2007/2008  
42 for children aged 6 to 36 months. The uptake of this age group has been estimated to vary between  
43 15 and 40%.<sup>14</sup>

### 44 *Data sources and case definition*

45 IPD cases were identified from the National Infectious Diseases Register (NIDR), a population-  
46 based electronic laboratory surveillance system maintained by the National Institute for Health and

47 Welfare (THL). It is mandatory for all clinical microbiology laboratories to notify all isolations of  
48 *Streptococcus pneumoniae* from blood or cerebrospinal fluid to NIDR and the process has been  
49 automated to send electronic reports to the database. Furthermore, the corresponding case isolates  
50 are sent to the national reference laboratory at THL for confirming the species and serotyping.  
51 Currently, more than 97% of the case isolates are received.<sup>16</sup> IPD case was defined as isolation of *S.*  
52 *pneumoniae* from blood or cerebrospinal fluid. The IPD surveillance in Finland and the THL  
53 laboratory methods have been described earlier.<sup>6,17</sup>

54 Vaccination status of each IPD case was verified from the National Vaccination Register and local  
55 electronic vaccination cards. Data on comorbidities were obtained from the national hospital  
56 discharge register (the Care Register for Health Care at THL). The study population was determined  
57 by using data from the Finnish Population Information System. All register-based information was  
58 linked by using the unique national personal identity code which is assigned to all permanent  
59 residents in Finland.

60 IPD cases were categorized according to the causative serotype into three mutually exclusive  
61 groups: PCV10 serotypes, PCV10-related serotypes (i.e. serotypes belonging to the same  
62 serogroups as vaccine types; in the data: 6A, 6C, 7C, 9N, 18B, 19A, 23A, 23B), or non-PCV10  
63 serotypes. The impact of PCV10 vaccination on each of the three serotype groups was evaluated. In  
64 addition, the impact was assessed separately for serotypes 6A and 19A.

#### 65 *Overall impact of PCV10 vaccination in vaccine-eligible children*

66 To estimate the overall impact of PCV10 vaccination on IPD, the relative reduction in the incidence  
67 of IPD for each serotype group was estimated by comparing a target cohort, comprising all vaccine-  
68 eligible children irrespective of vaccination status and born between June 2010 and September  
69 2016, to a season- and age-matched reference cohort in years 2002-2008 (Figure 1, panel A). The

70 children who were enrolled in the FinIP trial in years 2009-2010 were excluded from the analysis.

71 The follow-up period started at 3 months of age (the first scheduled vaccination dose) and lasted

72 until the end of December 2016 for the target cohort and December 2008 for the reference cohort.

73 The follow-up thus included children from 3 to 78 months of age.

74 To assess time trends in the age-specific risk of 19A IPD, age-specific cumulative hazards were

75 calculated for the target and reference cohorts.

### 76 *Indirect impact of PCV10 vaccination in unvaccinated children*

77 To estimate the indirect impact of PCV10 vaccination against IPD, the relative reduction in the

78 incidence of IPD for each serotype group was estimated by comparison of unvaccinated cohorts of

79 older children after and before PCV10 introduction (Figure 1, panel B). The unvaccinated target

80 cohort was chosen to comprise children born between January 2006 and May 2010. Children

81 vaccinated in the FinIP trial with PCV10 were excluded. The age- and season-matched reference

82 cohort comprised all children born between January 2000 and May 2004. The follow-up period

83 started in January 2012 (2006) and lasted until the end of December 2016 (2010) for the target

84 (reference) cohort. The follow-up included children from 19 to 131 months of age.

### 85 *Statistical methods*

86 Comparison of IPD incidence rates was performed by using Poisson regression. Vaccine impact

87 was defined as  $(1 - \text{incidence rate ratio}) * 100\%$ , comparing the target and reference cohorts. Absolute

88 rate reductions and the corresponding confidence intervals were calculated from the parameter

89 estimates with the delta method. No adjustments were made for comorbidities or influenza

90 vaccinations in the NVP because of the small number of cases, evenly distributed comorbidities in

91 the study cohorts and the low coverage of the influenza vaccinations. Statistical significance was

92 deemed at the 5% level. Statistical software R version 3.4.2<sup>18</sup> was used for all analyses.

93 *Ethical considerations*

94 As part of its statutory tasks, the National Institute for Health and Welfare (THL) is obliged to  
95 monitor the effectiveness and safety of the vaccines used in NVP. The study plan was approved by  
96 the THL institutional review board (May 23, 2013). Permissions to use the register data for research  
97 were obtained from the relevant register controllers at THL (THL/1090/6.02.00/2013).

98

99 **Results**

100 *Overall impact of PCV10 vaccination in vaccine-eligible children*

101 Table 1 presents the estimated overall impact of PCV10 vaccination by the serotype groups, based  
102 on comparisons of the entire vaccine-eligible cohort (years 2010-2016) with its reference cohort  
103 (years 2002-2008). Among children targeted for vaccination, the incidence of all IPD decreased by  
104 79% (95% CI 74 to 83%) from 42.9 to 9.2 cases per 100,000 person-years. The incidence of PCV10  
105 serotype IPD decreased by 94% (95% CI 91 to 96%) from 32.3 to 1.9 cases per 100,000 person-  
106 years. There was a borderline non-significant increase in non-PCV10 serotypes (IRR 1.53, 95% CI  
107 0.96 to 2.49). Figure 2 shows the evolution of the IPD incidence in children of vaccine-eligible age  
108 by epidemiological year (from July to June).

109 The incidence of IPD caused by PCV10-related serotypes decreased from 6.3 to 3.8 per 100,000  
110 person-years. This was mainly due to a decrease in the incidence of 6A with a relative reduction of  
111 95% (95% CI 75 to 100%). There was a statistically non-significant reduction of 26% (95% CI -13  
112 to 51%) in 19A.

113 Apart from the notably high incidence in the epidemiological years 2003/2004-2004/2005, the  
114 incidence of 19A fluctuated over the years with some increase towards the end of the follow-up



115 (Figure 3). The annual numbers of cases in 0-1 and 2-5 year-olds varied, respectively, between 0-17  
116 and 0-12. Cases of 19A IPD appeared to occur more often among older children aged 24-78 months  
117 in the target cohort compared to the reference cohort (Figure 4, Supplement Figure 1). The presence  
118 of underlying comorbidities in 19A cases was low and similar in both cohorts. The serotype-  
119 specific changes in incidences are presented in Supplement Table 1.

120 There were six breakthrough cases defined as vaccine-type IPD more than two weeks after  
121 administration of the booster dose at 12 months among the fully vaccinated children (received all 3  
122 doses). The causative serotypes were 19F (2 cases), 23F (2 cases), 6B and 14. No vaccine-type IPD  
123 cases occurred between two primary doses and a booster dose.

#### 124 *Indirect impact of PCV10 vaccination in unvaccinated children*

125 Table 2 presents the relative and absolute rate reductions in IPD incidence, based on the  
126 unvaccinated target cohort of older children (19-131 months of age) during 2012-2016 and its  
127 reference cohort in the years 2006-2010. The incidence of PCV10 serotype IPD decreased by 58%  
128 (95% CI 37 to 73%) from 5.9 cases in the reference cohort to 2.5 cases per 100,000 person-years in  
129 the target cohort, and the incidence of all IPD by 33% (95% CI 8 to 52%) from 7.3 to 4.8 per  
130 100,000 person-years (Figure 2). There was no indirect effect against the PCV10-related serotypes.  
131 The incidence of 19A IPD was 0.69 and 0.74 cases per 100,000 person-years, and that of 6A IPD  
132 0.00 and 0.25 cases per 100,000 person-years in the reference and target cohorts, respectively  
133 (Figure 3, Supplement Table 2). The incidence of non-PCV10 serotypes was five times higher in  
134 the target-cohort compared to the reference cohort, but the confidence intervals were wide and the  
135 incidence remained low.

#### 136 **Discussion**

137 In Finland, the introduction of PCV10 into the NVP has resulted in very high overall impact against  
138 IPD (79%) in the vaccine-eligible birth cohorts six years after PCV10 implementation. This is  
139 explained by the almost complete elimination of PCV10-type and 6A IPD and the low incidence of  
140 other PCV10-related and non-PCV10 serotypes.

141 In unvaccinated older children, the indirect PCV10 impact on IPD was less notable (33%) due to  
142 the smaller reduction in PCV10-serotype disease and small increases in PCV10-related and non-  
143 PCV10 disease.

144 According to a random-effect meta-analysis<sup>19</sup>, the pooled impact of PCV7 against all IPD was  
145 estimated to vary between 51 and 67% four to seven years after introduction among children <5  
146 years of age. The impact of the combined PCV7 and PCV13 programmes against all IPD among  
147 children <2 years of age have been reported to vary between 64% (Stockholm, Sweden) to 89%  
148 (Norway)<sup>7,20</sup> and the combined PCV7 and PCV10 programmes among children <5 years of age  
149 from 64% (Quebec) to 80% (the Netherlands)<sup>5,21</sup>. Thus, our estimate of 79% is consistent with  
150 other settings.

151 Our data suggest that PCV10 results in long-lasting direct cross-protection against 6A IPD. In  
152 previous studies, the incidence of serotype 6A carriage and invasive disease were largely reduced in  
153 vaccinated children after PCV7 implementation, suggesting direct cross-protection from 6B<sup>3,9,22</sup>,  
154 although not all studies have supported this.<sup>23</sup> After PCV10 introductions, a non-significant direct  
155 cross-protection against 6A was found in Brazil<sup>24</sup> and Sweden<sup>13</sup>, and significant protection in  
156 Finland<sup>6</sup> -- a finding that is further supported by the present study.

157 After PCV7 introduction in the USA, indirect cross-protection against 6A was observed among  
158 older children and adults<sup>9</sup> although the effect was smaller and appeared later than the indirect  
159 impact against the PCV7 serotypes.<sup>11</sup> In Sweden, a significant indirect cross-protection was

160 reported after PCV10 vaccination in all unvaccinated age-groups.<sup>13</sup> In our study, the number of  
161 cases of the older unvaccinated children was too small to assess indirect protection against 6A.  
162 However, 6A is not a replacing serotype among older children and adults in Finland in IPD<sup>25</sup> or in  
163 carriage in children<sup>26</sup> suggesting some degree of indirect impact.

164 In contrast to 6A and our earlier observation based on a three-year follow-up post introduction<sup>6</sup>, we  
165 observed no significant effect of PCV10 against 19A IPD. There were, however, differences by age  
166 and calendar time in 19A disease between the cohorts, as the cases of the vaccine-eligible cohort  
167 tended to be older and occur more often towards the end of the follow-up than those of the reference  
168 cohort. One explanation could be the increasing infection pressure from 19A, also seen in adult age  
169 groups 2012-2013 onwards<sup>25</sup>; another speculation is that the 19F component in PCV10 induces  
170 initial cross-protection which wanes with age thus rendering vaccinated children susceptible again  
171 to 19A in lack of specific 19A immunity. The small number of cases and year-to-year variability  
172 make firm conclusions difficult.

173 After PCV7 introduction in the USA and several countries in Europe, 19A carriage and the  
174 incidence of 19A IPD increased rapidly among children targeted to vaccination and also in older  
175 age groups.<sup>3,4,7,11,22</sup> Our results are in contrast with the 3- to 4-fold increases of 19A IPD post-PCV7  
176 in the USA in children<sup>11</sup>, although the increases in adults have been similar.<sup>11,25</sup> In the US, the  
177 increase in 19A disease was mainly due to the increase in antimicrobial-resistant clonal complex  
178 (CC) 320 and a concurrent decrease in antimicrobial-susceptible CC199, suggesting that antibiotic  
179 selection pressure could have contributed to the 19A emergence<sup>27,28</sup>. In Finland, the outpatient  
180 antimicrobial use is lower than in USA<sup>29</sup>, and also lower than the average for all European Union  
181 countries<sup>30</sup>, although higher compared to the Nordic neighbors. The introduction of PCV10 into the  
182 vaccination program has resulted in further reduction of antimicrobial use in children.<sup>31</sup> The  
183 antimicrobial non-susceptibility of pneumococci has also decreased in all age-groups and especially

184 among young children during the PCV10 program.<sup>32,33</sup> For 19A, antimicrobial susceptible clones  
185 (especially ST994 and ST199) have become dominant both in children<sup>34</sup> and among elderly  
186 people<sup>35</sup> during the post-PCV10 period in Finland resulting in low proportion of antimicrobial  
187 resistance.

188 In our first evaluation three years after PCV10 implementation in Finland, the 19A IPD incidence  
189 had declined significantly by 62% in the vaccine-eligible cohort (children aged 3-42 months).<sup>6</sup>  
190 Statistically significant direct effectiveness against 19A IPD was reported with matched case-  
191 control studies in Brazil (82%, children aged 2.6-53.1 months) and Quebec (79%, children aged 2-  
192 59 months) after at least one PCV10 dose.<sup>24,36</sup> In the Netherlands, where PCV10 was introduced  
193 about 5 years after PCV7 with a 3+1 schedule, the incidence of 19A IPD was lower in the PCV10-  
194 eligible cohort than in the PCV7-eligible cohort. However, the cross-protective effect of 19A was  
195 inconclusive, as the incidence of both PCV10-related and non-related serotypes was low.<sup>5</sup> In  
196 Sweden, where PCV7 was first introduced in the years 2007 through 2009 depending on the county  
197 and then replaced by PCV10 or PCV13 in 2009-2010, no direct cross-protection against 19A was  
198 reported in the PCV10 areas.<sup>13</sup>

199 The strengths of the current study setting include high data completeness ( $\approx 80\%$  of isolates in years  
200 2002-2008 were available for serotyping, and  $\approx 100\%$  after that) and a well-established and  
201 consistent reporting in a population-based nationwide surveillance system. The personal identity  
202 code allowed linkage between different registers to find vaccination histories and health statuses for  
203 the IPD cases.

204 An apparent limitation of our study design, based on a before-after comparison, is its inclination to  
205 bias due to secular trends. In addition, the small numbers of cases caution against over-  
206 interpretation of analyses on individual serotypes. The age- and calendar-time related changes  
207 observed in 19A IPD may equally well be explained by chance alone: the annual numbers of 19A

208 cases in under 2-year-olds in 2002-2016 ranged between 0 and 17, with median of 5. Therefore,  
209 firm conclusions can only be made on the overall IPD level, where randomness and serotype-  
210 specific secular trends are balanced out with sufficient numbers and follow-up time. In addition, the  
211 vaccine-induced dynamics between serotypes in the post-vaccination period may limit comparisons  
212 with the reference cohort. Depending on the levels of competition between serotypes and vaccine  
213 coverage, replacement and herd protection in carriage and disease may occur differently.<sup>37</sup>  
214 Moreover, influenza vaccinations that started in fall 2007 for children 6 to 36 months of age, may  
215 have prevented some IPD cases. However, the vaccination coverage was 40% in 2007 to 2009 but  
216 less than 20% during the PCV10 program<sup>14</sup>, which might result in a conservative estimate of the  
217 PCV10 impact.

218 In conclusion, PCV10 vaccination has resulted in a very high impact against PCV10 serotypes, no  
219 notable replacement due to non-PCV10 serotypes, and long-lasting direct cross-protection against  
220 6A IPD among vaccine-eligible children. Most importantly, the net impact on IPD irrespective of  
221 serotypes six years after PCV implementation in Finland is among the largest reported  
222 globally.<sup>5,7,8,19</sup>

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348 **Conflict of Interest Statement for all authors.**

349 National Institute for Health and Welfare has received research funding from GlaxoSmithKline  
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351 conjugate vaccine. HR-K, AAP, MT, LS, and JJ are co-investigators in these studies. The other  
352 authors have no conflicts to disclose. Current study was entirely publicly funded.

353 **Authorship contributions**

354 HR-K coordinated the data collection, carried out the analyses, interpreted the data and drafted the  
355 manuscript; AAP, KA, JPN, HN, and JJ designed the study, interpreted the data and reviewed and  
356 revised the manuscript; MT, LS, and MJV supervised the microbiological data quality and reviewed  
357 and revised the manuscript; and all authors approved the final manuscript as submitted.

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360 **Tables and figures**

361 Table 1. Incidence rates of invasive pneumococcal disease (IPD) and the corresponding relative and  
362 absolute incidence rate reductions, based on the comparison of the PCV10-eligible target cohort  
363 (2010-2016) vs. a reference cohort (2002-2008).

364

365 Table 2. Incidence rates of invasive pneumococcal disease (IPD) and the corresponding relative and  
366 absolute incidence rate reductions, based on the comparison of an unvaccinated target cohort (2012-  
367 2016) vs. a reference cohort (2006-2010).

368

369 Figure 1. Target and reference cohorts for estimating the overall impact of PCV10 on IPD among  
370 vaccine-eligible children (panel A) and unvaccinated children (panel B).

371

372 Figure 2. Incidence rates of invasive pneumococcal disease (IPD) in age groups <2 years (panel A)  
373 and 2-5 years (panel B) by serotype group in epidemiological years 6/2002–6/2017.

374

375 Figure 3. Incidence rates of 19A and 6A IPD in age groups <2 years (panel A) and 2-5 years (panel  
376 B) in epidemiological years 6/2002–6/2017.

377

378 Figure 4. Incidence of all IPD (first panel) and 19A IPD (second panel) in the PCV10-eligible target  
379 and reference cohorts by age group.

380

381

382 Supplement table 1. Incidence rates of invasive pneumococcal disease (IPD) of individual PCV10  
383 serotypes and the corresponding relative and absolute incidence rate reductions, based on the  
384 comparison of the PCV10-eligible target cohort (2010-2016) vs. a reference cohort (2002-2008).

385

386 Supplement table 2. Incidence rates of invasive pneumococcal disease (IPD) of individual PCV10  
387 serotypes and the corresponding relative and absolute incidence rate reductions, based on the  
388 comparison of an unvaccinated target cohort (2012-2016) vs. a reference cohort (2006-2010).

389

390 Supplement Figure 1. Age-specific cumulative hazards of 19A IPD in the PCV10-eligible target  
391 cohort and the reference cohort.

392

393

Table 1. Incidence rates of invasive pneumococcal disease (IPD) and the corresponding relative and absolute incidence rate reductions, based on the comparison of the PCV10-eligible target cohort (2010-2016) vs. a reference cohort (2002-2008).

Table 2. Incidence rates of invasive pneumococcal disease (IPD) and the corresponding relative and absolute incidence rate reductions, based on the comparison of an unvaccinated target cohort (2012-2016) vs. a reference cohort (2006-2010).

Table 1.

	Incidence per 100,000 person-years (N)		Target cohort vs. reference cohort	
	Reference cohort 2002-2008* Follow-up yrs 1211504	Target cohort 2010-2016** Follow-up yrs 1243145	Relative rate reduction, % (95% CI)	Absolute rate reduction per 100,000 person-years (95% CI)
Any culture confirmed IPD	42.9 (520)	9.2 (114)	78.6 (73.9, 82.6)	33.8 (29.7, 37.8)
PCV10 serotypes	32.3 (391)	1.9 (23)	94.3 (91.5, 96.3)	30.4 (27.1, 33.7)
PCV10-related serotypes	6.3 (76)	3.8 (47)	39.7 (13.7, 58.4)	2.5 (0.72, 4.3)
Non-PCV10 serotypes	2.3 (28)	3.5 (44)	-53.1 (-148.6, 4.1)	-1.2 (-2.6, 0.12)
Not known	2.1 (25)	0 (0)	NA	NA

\*Age 3-78 months, born Jun 2002 – Sep 2008

\*\*Age 3-78 months, born Jun 2010 – Sep 2016

NA = not applicable

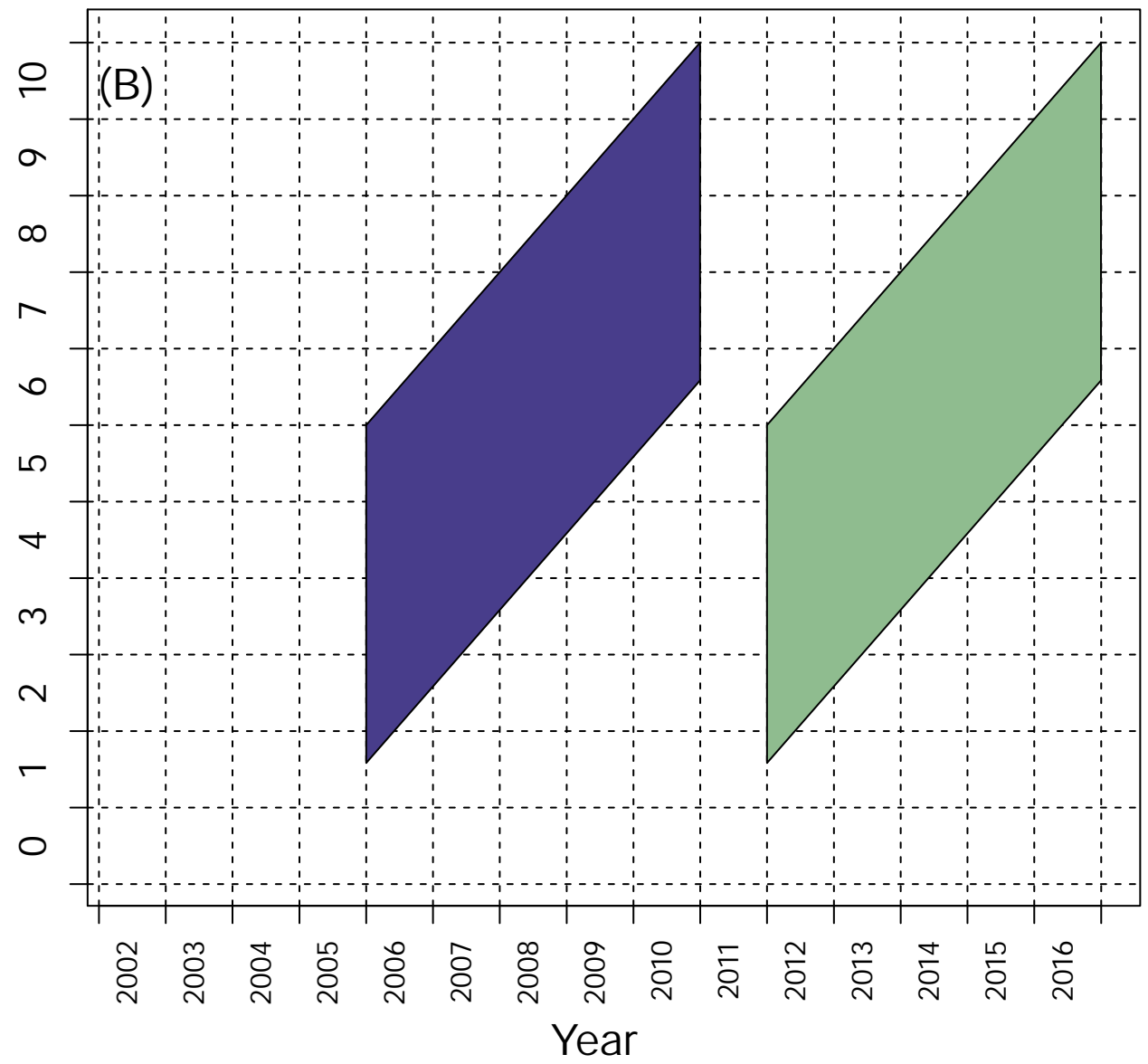
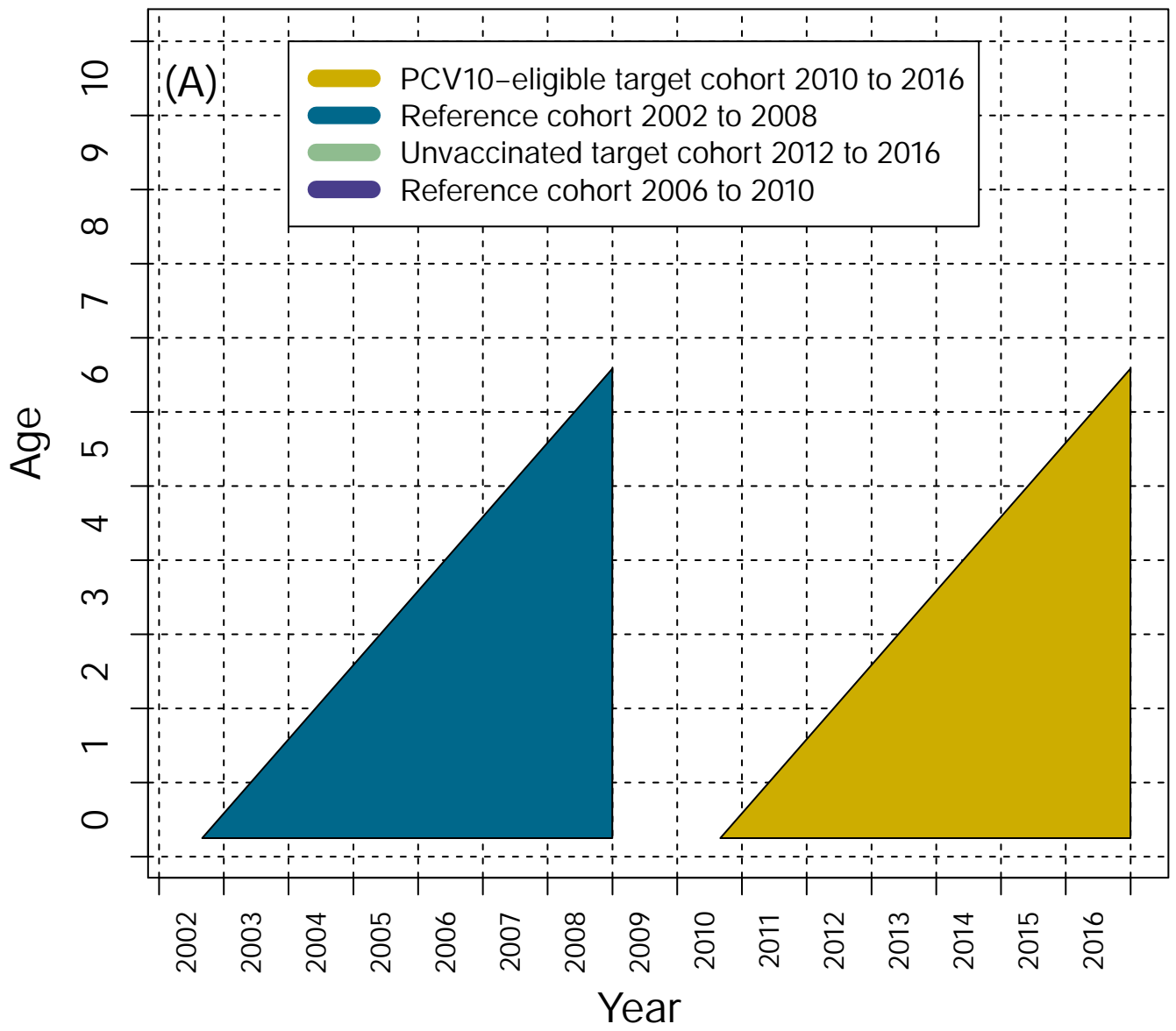
Table 2.

Serotype group	Incidence per 100,000 person-years (N)		Unvaccinated target cohort vs. reference cohort	
	Reference cohort 2006-2010* Follow-up yrs 1309618	Target cohort 2012-2016** Follow-up yrs 1221933	Relative rate reduction, % (95% CI)	Absolute rate reduction per 100,000 person-years (95% CI)
Any culture confirmed IPD	7.3 (95)	4.8 (59)	33.4 (8.2, 52.1)	2.4 (0.52, 4.3)
PCV10-serotypes	5.9 (77)	2.5 (30)	58.2 (37.1, 73.0)	3.4 (1.8, 5)
PCV10-related serotypes	0.69 (9)	1.2 (15)	-78.6 (-325.3, 20.5)	-0.54 (-1.3, 0.23)
Non-PCV10 serotypes	0.23 (3)	1.2 (14)	-400.2 (-2069.8, -63.3)	-0.92 (-1.8, -0.26)
Not known	0.46 (6)	0 (0)	NA	NA

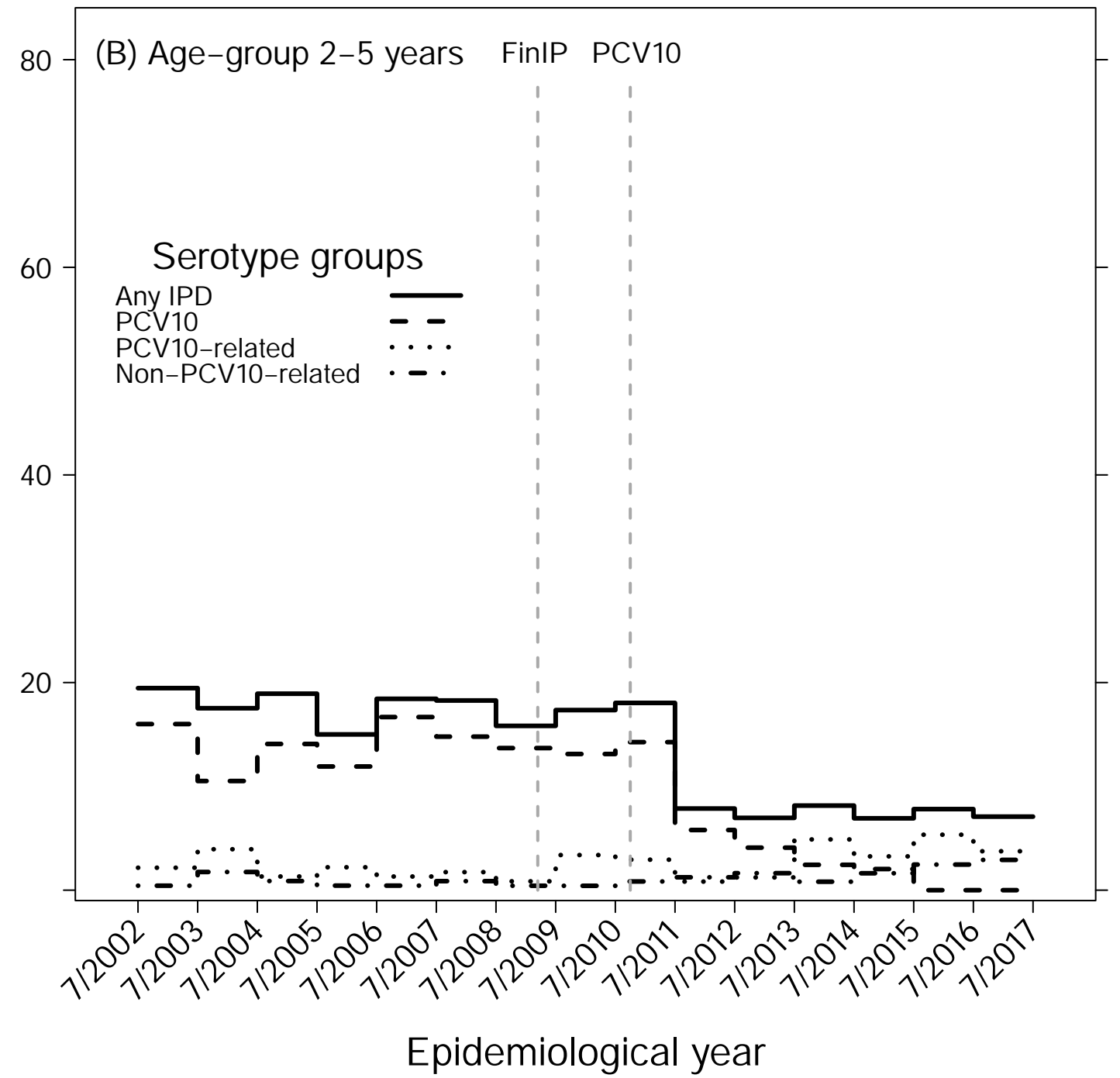
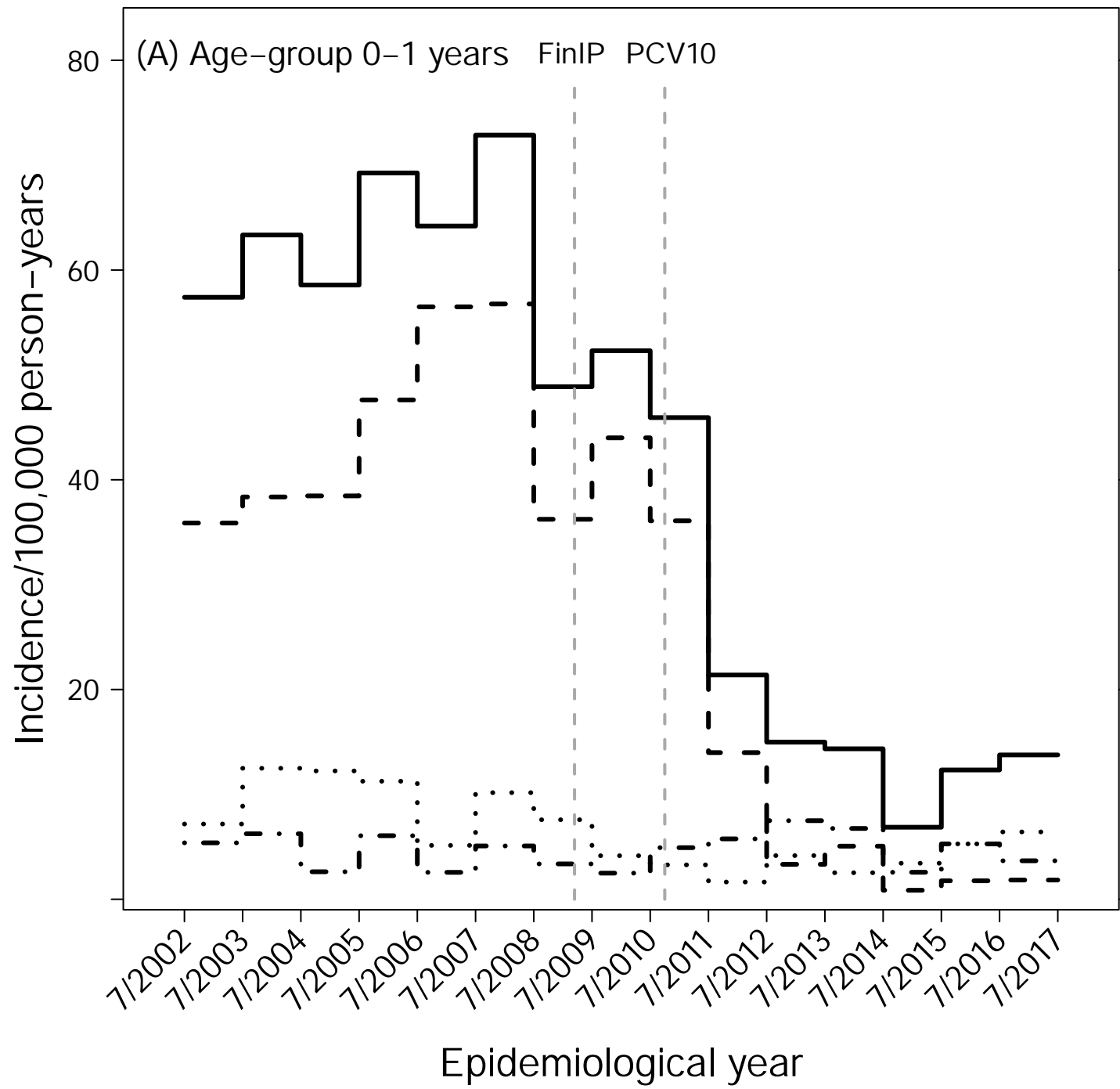
\*Age 19-131 months, born Jan 2000 – May 2004

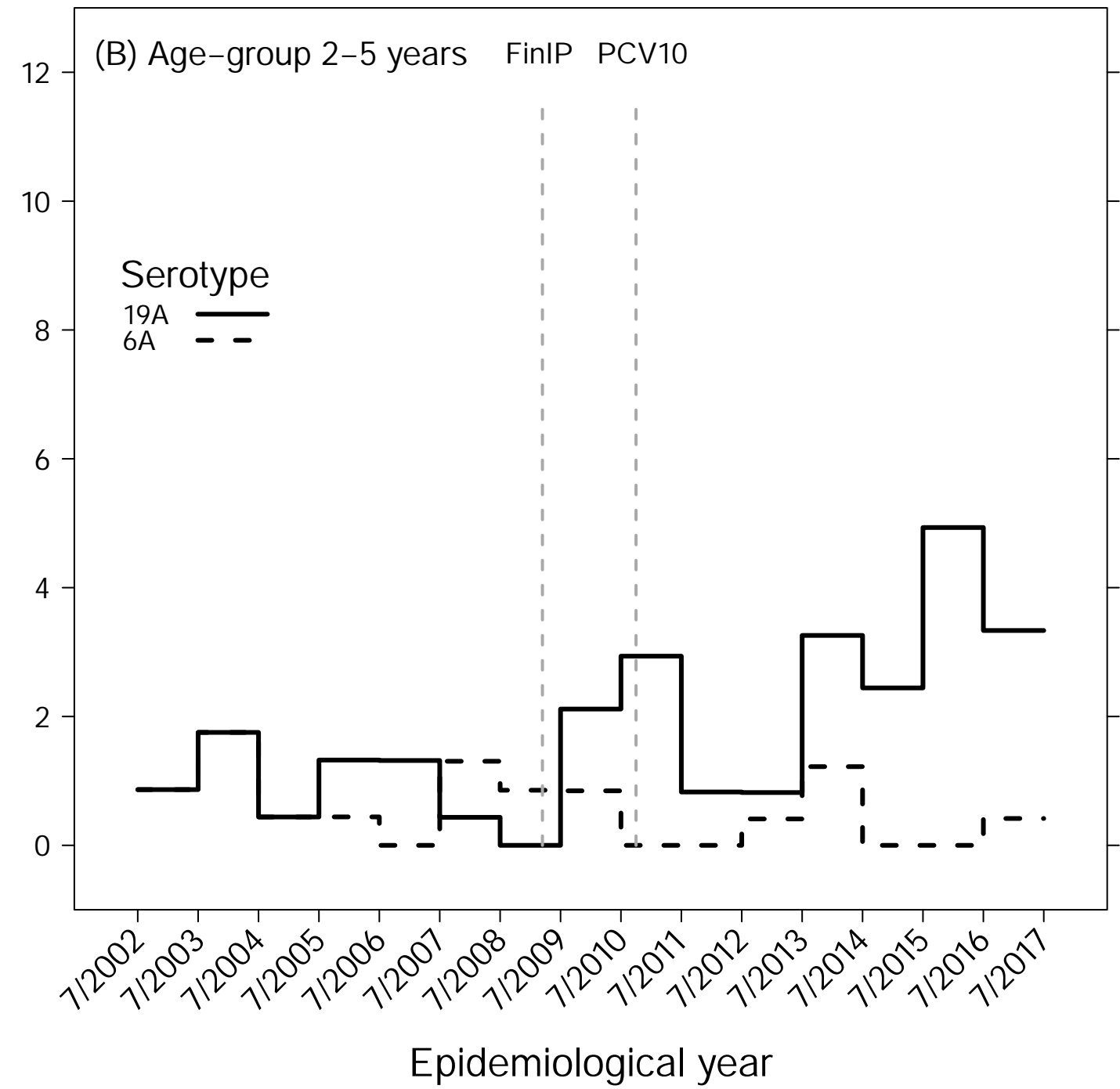
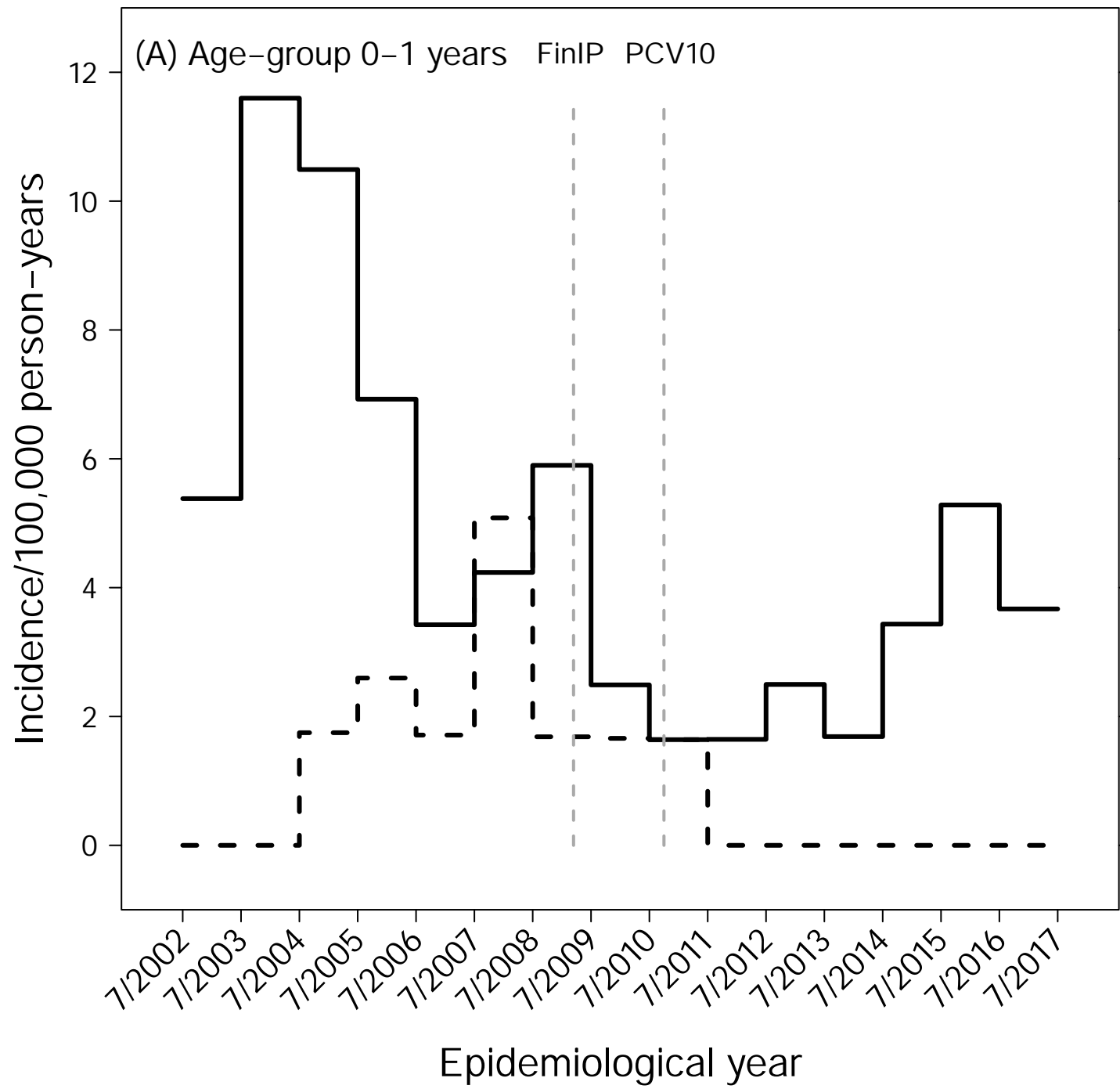
\*\*Age 19-131 months, born Jan 2006 – May 2010

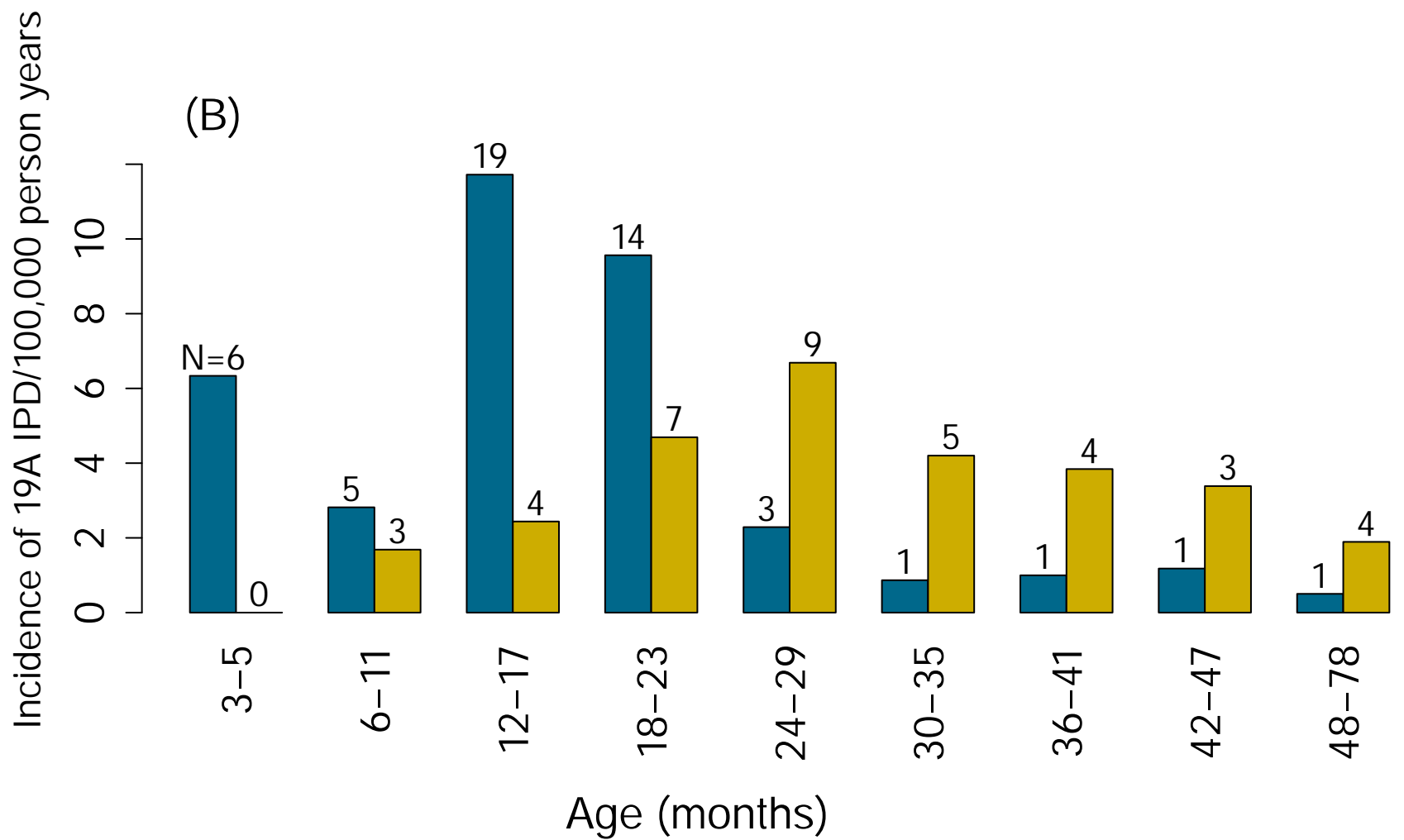
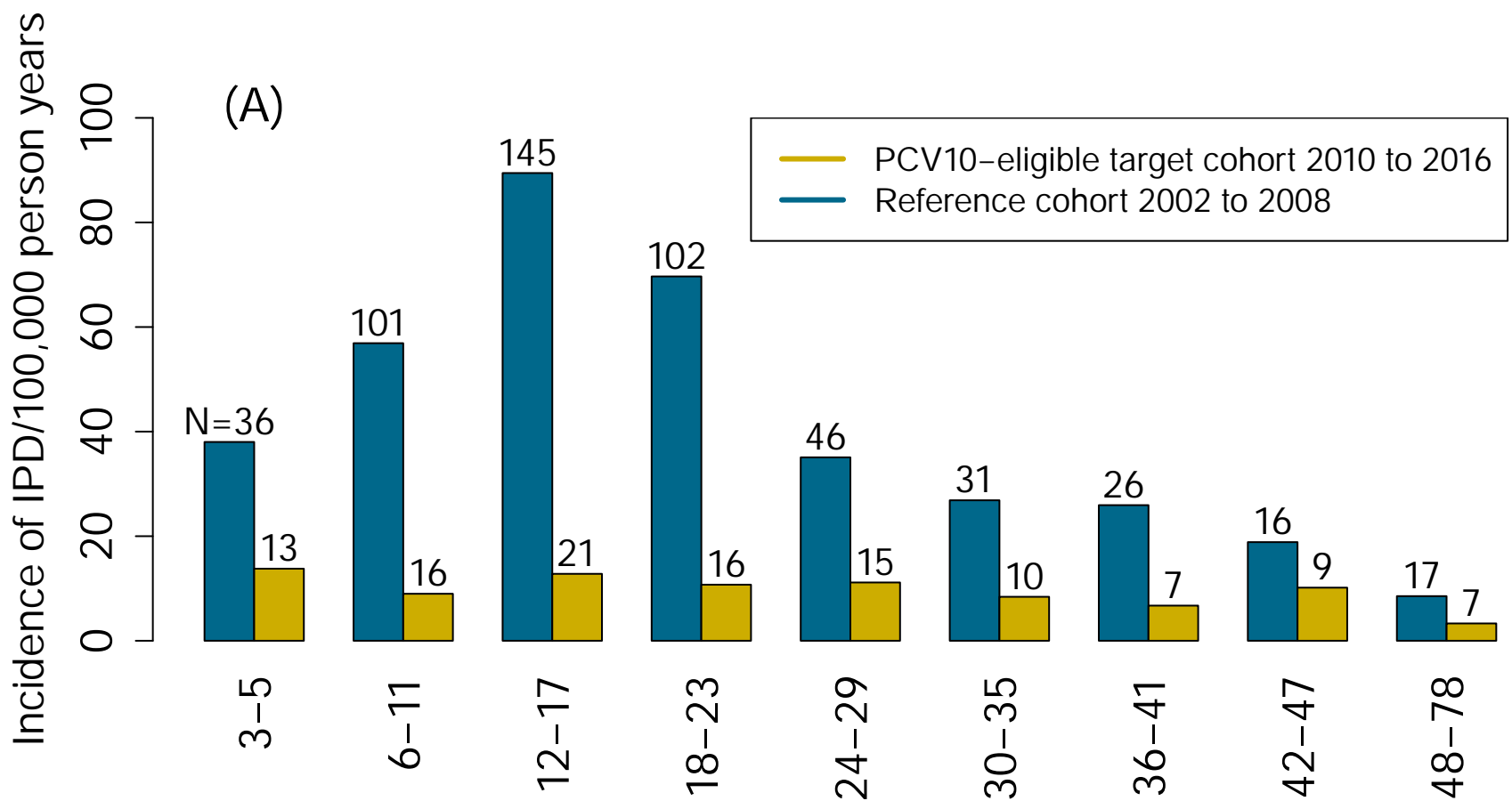
NA = not applicable











## Supplementary Appendix

Supplement to: Rinta-Kokko H, Palmu AA, Auranen K, et al. Long-term impact of 10-valent pneumococcal conjugate vaccination on invasive pneumococcal disease among children in Finland

**Supplement Table 1.** Incidence rates of invasive pneumococcal disease (IPD) of individual PCV10 serotypes and the corresponding relative and absolute incidence rate reductions, based on the comparison of the PCV10-eligible target cohort (2010-2016) vs. a reference cohort (2002-2008).

Serotype	Incidence per 100,000 person-years (N)		Target cohort vs. reference cohort	
	Reference cohort 2002-2008* Follow-up yrs 1211504	Target cohort 2010-2016** Follow-up yrs 1243145	Relative rate reduction, % (95% CI)	Absolute rate reduction per 100,000 person-years (95% CI)
1	0.08 (1)	0.08 (1)	2.6 (-2364.5, 96.2)	0 (-0.2, 0.2)
4	0.91 (11)	0 (0)	100 (80.8, 100)	0.9 (0.4, 1.4)
5	0 (0)	0 (0)	NA	0 (0, 0)
6B	10.9 (132)	0.24 (3)	97.8 (94.2, 99.5)	10.7 (8.8, 12.5)
7F	0.66 (8)	0.08 (1)	87.8 (33.6, 99.3)	0.6 (0.1, 1.1)
9V	1.24 (15)	0 (0)	100 (86.3, 100)	1.2 (0.6, 1.9)
14	8.58 (104)	0.64 (8)	92.5 (85.6, 96.6)	7.9 (6.2, 9.7)
18C	2.31 (28)	0.08 (1)	96.5 (83.7, 99.8)	2.2 (1.4, 3.1)
19F	3.71 (45)	0.4 (5)	89.2 (75.2, 96.3)	3.3 (2.2, 4.5)
23F	3.88 (47)	0.32 (4)	91.7 (79.6, 97.5)	3.6 (2.4, 4.7)
6A	1.57 (19)	0.08 (1)	94.9 (75.3, 99.7)	1.5 (0.8, 2.2)
19A	4.21 (51)	3.14 (39)	25.5 (-12.8, 51.2)	1.1 (-0.5, 2.6)

\*Age 3-78 months, born Jun 2002 – Sep 2008

\*\*Age 3-78 months, born Jun 2010 – Sep 2016

NA = not applicable

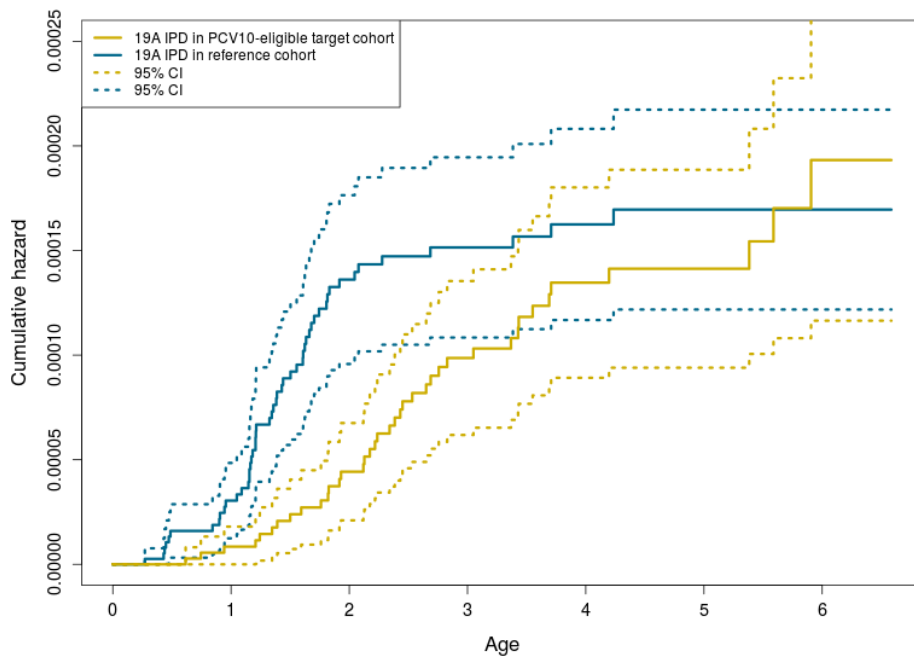
**Supplement Table 2.** Incidence rates of invasive pneumococcal disease (IPD) of individual PCV10 serotypes and the corresponding relative and absolute incidence rate reductions, based on the comparison of an unvaccinated target cohort (2012-2016) vs. a reference cohort (2006-2010).

Serotype	Incidence per 100,000 person-years (N)		Unvaccinated target cohort vs. reference cohort	
	Reference cohort 2006-2010* Follow-up yrs 1309618	Target cohort 2012-2016** Follow-up yrs 1221933	Relative rate reduction, % (95% CI)	Absolute rate reduction per 100,000 person-years (95% CI)
1	0 (0)	0 (0)	NA	0 (0, 0)
4	0.15 (2)	0 (0)	100 (-66.5, 100)	0.2 (-0.06, 0.4)
5	0 (0)	0 (0)	NA	0 (0, 0)
6B	0.76 (10)	0.41 (5)	46.4 (-50.8, 83.3)	0.4 (-0.2, 1.0)
7F	0.38 (5)	0.25 (3)	35.7 (-162.1, 86.8)	0.1 (-0.3, 0.6)
9V	0 (0)	0.16 (2)	-Inf (-Inf, 40)	-0.2 (-0.4, 0.06)
14	1.68 (22)	0.33 (4)	80.5 (49.1, 94.3)	1.4 (0.6, 2.1)
18C	1.3 (17)	0.49 (6)	62.2 (9, 86.4)	0.8 (0.08, 1.5)
19F	0.46 (6)	0.16 (2)	64.3 (-55.1, 94.8)	0.3 (-0.1, 0.7)
23F	1.15 (15)	0.65 (8)	42.8 (-31.6, 77)	0.5 (-0.3, 1.2)
6A	0 (0)	0.25 (3)	-Inf (-Inf, -7.3)	-0.3 (-0.5, 0.03)
19A	0.69 (9)	0.74 (9)	-7.2 (-174.5, 58.2)	-0.05 (-0.7, 0.6)

\*Age 19-131 months, born Jan 2000 – May 2004

\*\*Age 19-131 months, born Jan 2006 – May 2010

NA = not applicable



**Supplement Figure 1.** Age-specific cumulative hazards of 19A IPD in the PCV10-eligible target cohort and the reference cohort.