

1 Trends in serotypes and sequence types among cases of invasive pneumococcal
2 disease in Scotland, 1999-2010

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76 **Abstract**

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78 **Introduction**

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80 The 7-valent pneumococcal conjugate vaccine (Prevenar®, Wyeth; PCV7) was
81 introduced to the UK paediatric immunisation schedule in 2006. This study
82 investigates trends in serotypes and multi locus sequence types (STs) among cases of
83 invasive pneumococcal disease (IPD) in Scotland prior to, and following, the
84 introduction of PCV7.

85

86 **Methods**

87

88 Scottish Invasive Pneumococcal Disease Enhanced Surveillance has records of all
89 cases of invasive pneumococcal disease in Scotland since 1999. Cases diagnosed from
90 blood or cerebrospinal fluid isolates until 2010 were analysed. Logistic and poisson
91 regression modelling was used to assess trends prior to and following the introduction
92 of PCV7.

93

94 **Results**

95

96 Prior to PCV7 use, on average 650 cases of IPD were reported each year; 12%
97 occurred in those aged <5 years and 35% affected those aged over 65 years. Serotypes
98 in PCV7 represented 47% of cases (68% in <5 year olds). The serotype and ST
99 distribution was relatively stable with only serotype 1 and associated ST 306 showing
100 an increasing trend. PCV7 introduction was associated with a 69% (95% CI: 50%,
101 80%) reduction in the incidence of IPD among those aged <5 years, a 57% (95% CI:
102 47%, 66%) reduction among those aged 5-64 years but no significant change among
103 those aged 65 years and over where increases in non-PCV7 serotypes were observed.
104 Serotypes which became more prevalent post-PCV7 are those which were associated
105 with STs related to the PCV7 serotypes.

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107

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109 Conclusions

110

111 Routine serotyping and sequence typing in Scotland allowed the assessment of the
112 relationship between the capsule and the clones in the post vaccination era. Changes
113 in the distribution of serotypes post PCV7 introduction appear to be driven by
114 associations between serotypes and STs prior to PCV7 introduction. This has
115 implications for the possible effects of the introduction of higher valency vaccines and
116 could aid in predicting replacement serotypes in IPD

117

118 **Introduction**

119

120 *Streptococcus pneumoniae* (*S. pneumoniae*) is responsible for a substantial burden of
121 disease, accounting for an estimated 1.6 million deaths annually worldwide [1]. In
122 developed countries, the observed incidence of invasive pneumococcal disease (IPD)
123 is between 8 and 75 cases per 100,000 individuals [2], with studies showing that the
124 burden of disease is predominantly attributable to only around 20 or 30 of the [94+](#)
125 circulating pneumococcal serotypes [3]. Furthermore, it has been observed that
126 approximately two thirds of adult pneumococcal disease and 80% of disease in
127 children is attributable to between only around 8 to 10 serotypes [4].

128

129 Many recent studies of serotypes involved in IPD concern the comparison of a pre-
130 conjugate vaccination period to a post-vaccination period to examine any changes in
131 serotype distribution likely to be related to the use of [the 7-valent pneumococcal](#)
132 [conjugate vaccine \(PCV7\)](#). In the USA, and other countries subsequently, great
133 reductions in IPD were documented which were not limited to the vaccine targeted
134 age group [5]. However, increases in IPD caused by non-PCV7 serotypes, in
135 particular serotype 19A, following PCV7 use have been documented [5-11].

136

137 The capsule is thought to be the main determinant for carriage prevalence and
138 invasiveness of the pneumococcus and hence the determinant of prevalence amongst
139 invasive disease isolates before and after vaccination [12, 13]. However, concerns
140 have been raised that the increase in serotype 19A IPD in particular is perhaps
141 attributable to a capsular switch event after being found associated with a sequence
142 type (ST), ST695, which was previously only linked with vaccine serotype 4 [14, 15].

143 Other studies have documented increases due to the expansion of multi-drug resistant
144 STs such as ST276 and ST320 [16, 17]. Thus, it is becoming increasingly important
145 to examine both the STs and serotypes involved in IPD in order to determine the
146 potential effectiveness of serotype-specific pneumococcal vaccinations.

147

148 [In September 2006, PCV7 was introduced to the routine childhood immunisation](#)
149 [schedule in the United Kingdom in a three dose program at age 2, 4, and 13 months,](#)
150 [with a catch-up for those aged up to 2 years.](#) This study examines the trends in
151 serotype and ST distributions prior to the introduction of PCV7 in Scotland, adding to
152 existing reports on the pre-vaccine period in Scotland which describe increases in
153 serotype 1 and ST 306 [18, 19]; the effect of PCV7 on the incidence of IPD; trends in
154 the serotype and ST distribution post-vaccination; and the association between
155 serotype and ST pre- and post-vaccination.

156

157 **Methods**

158

159 *Data*

160

161 Data were obtained from the Scottish Invasive Pneumococcal Disease Enhanced
162 Surveillance (SPIDER) database on all cases of IPD, identified by blood or
163 cerebrospinal fluid, in Scotland between 1999 and 2010. The serogroup responsible
164 for each case of disease was available for all years, whilst serotype and ST
165 information was only available from 2002.

166

167 Clinical isolates (grown from blood or cerebrospinal fluid) of *S. pneumoniae* were
168 sent to Scottish Haemophilus, Legionella, Meningococcus and Pneumococcus
169 Reference Laboratory (SHLMPRL) after identification at diagnostic microbiology
170 laboratories in Scotland. At SHLMPRL, these isolates were grown on Columbia
171 blood agar (Oxoid, United Kingdom) at 37°C under anaerobic conditions by use of an
172 anaerobic pack (Oxoid, United Kingdom) and after a single subculture were stored at
173 -80°C on Protect beads (M-Tech Diagnostics, United Kingdom). Isolates were
174 serotyped by a coagglutination method described elsewhere [20]. MLST was
175 performed as described previously [21-23].

176

177 Epidemiological years ranging from winter of one year to the end of autumn of the
178 following year were used to ensure winter seasons were grouped together since IPD
179 predominantly occurs during this season.

180

181 Serotypes and STs were classified according to their joint occurrence prior to the
182 introduction of PCV7 [\(1999-2005\)](#) and their emergence post-PCV7 [\(2006-2010\)](#). STs
183 were classified as associated with PCV7 serotypes if they occurred at least once in
184 conjunction with one of the seven PCV7 serotypes (labelled PCV7-ST); otherwise
185 they were classified as not associated with PCV7 serotypes (NonPCV7-ST). STs
186 which only occurred following the introduction of PCV7 were classified separately as
187 PostPCV7-ST. The PCV7-ST group was subdivided into two groups: a group of 12
188 STs (9, 36, 113, 124, 138, 156, 162, 176, 205, 206, 246, 311) with a high frequency of
189 co-occurrence with the PCV7 serotypes (labelled HF PCV7-ST), and a larger group
190 with low frequency co-occurrence (LF PCV7-ST). Serotypes were categorised in four
191 different groups: PCV7 serotypes (4, 6B, 9V, 14, 18C, 19F, 23F); serotypes not in

192 | PCV7 but associated with STs linked through co-occurrence ~~prior~~ to the PCV7
193 | serotypes, denoted PCV7-ST serotypes; serotypes not in PCV7 and not associated
194 | with the STs linked to PCV7 serotypes, denoted NonPCV7-ST serotypes; and
195 | serotypes which only occurred post-PCV7 vaccination, denoted PostPCV7 serotypes.

196 |

197 | *Statistical analysis*

198 |

199 | Logistic regression models were used to examine changes and linear trends in the
200 | serogroup, serotype and ST distributions, with year treated as a continuous variable
201 | ranging from 1999 to 2005. Only serogroups, serotypes and STs responsible for at
202 | least 1% of cases of IPD were considered. Analyses were carried out for the
203 | serogroups for age groups 0-4, 5-64, and ≥ 65 years separately. Bonferroni adjusted
204 | confidence intervals were calculated and the Benjamini and Hochberg adjustment for
205 | multiple testing used in the assessment of the significance of the linear trend [24].

206 |

207 | Poisson regression models were used to assess changes in IPD incidence. The
208 | percentage change in the incidence of PCV7 serotypes and NonPCV7 serotypes from
209 | the pre-vaccine period to the post-vaccine period was assessed by predicting the post-
210 | vaccination incidence allowing for a trend in the pre-vaccination years and comparing
211 | the observed cases with the predicted as suggested elsewhere [25, 26]; 95%
212 | confidence intervals were used. Cases with missing age (27, 0.4%) were omitted. For
213 | 637 cases (10.1%), no information on the serotype or serogroup was available. The
214 | number of vaccine type (VT) or non-vaccine type (NVT) serotypes was imputed,
215 | separately by year and age group, using the observed proportions of VT serotypes.
216 | Imputation of serotype, from serogroup, was also carried out when serotype

Comment [KL1]: Should I add further comment that 1999 refers to year 1999/00 ("winter year") and 2005 is 2005/06?

217 information was not available prior to 2002. This was based upon the observed
218 proportions of serotypes within serogroups in the period 2002-2006, separately by age
219 group. All analysis was carried out using R versions 2.8-2.12 [27].

220

221

222 **Results**

223

224 *Trends in the serotype and sequence type distributions prior to the introduction on* 225 *PCV7*

226

227 Between 1999/00 and 2005/06, on average approximately 650 cases of IPD per year
228 were reported in Scotland, rising from 538 in 1999/00 to 743 in 2002/03. A
229 subsequent drop occurred, primarily amongst those aged ≥ 65 years, following the
230 introduction of [the 23-valent pneumococcal polysaccharide vaccine \(PPV23\) in the](#)
231 [UK for those aged at least 65 years in 2003, with a coverage of 71%. This was](#) followed
232 by a rise to 739 in 2005/06. IPD was most common amongst the elderly during this
233 period, with 44% of all cases of IPD identified in those aged ≥ 65 years. 12% of cases
234 affected those aged < 5 years.

235

236 *Serogroup analysis*

237

238 In total, 36 different serogroups were identified in IPD between 1999/00 and 2005/06.
239 Serogroup 14 was the most common, accounting for approximately 17% of all cases.
240 Serogroups 9 and 1 were also common, causing around 9% and 8% of cases,
241 respectively. Serogroup 1 replaced serogroup 14 as the most common serogroup in

Comment [KL2]: Stefan/Chris- any idea of coverage?

242 2005/06. The proportion of IPD cases associated with serogroup 1 increased steadily
243 over the pre-PCV7 study period.

244

245 There was significant evidence of an increasing trend for serogroup 1 ($p<0.001$),
246 (Table 1, Part A). Serogroup 14 was borderline significant in the analysis for all age
247 groups after adjustment for multiple testing, with a decreasing trend between 1999/00
248 and 2005/06.

249 *Serotype analysis*

250

251 Between 2003/04 and 2005/06, 42 different serotypes were identified as causing IPD.
252 PCV7 serotypes accounted for 47% of cases in this period and were responsible for
253 68% of cases in those <5 years, 40% in those aged 5-64 years and 48% in those ≥ 65
254 years. The most common serotypes, 14 (15%), 1 (13%), 4 (7%), 9V (7%), 8 (6%), 3
255 (6%), 23F (5%), 6B (4%), 7F (4%) and 19F (4%), together account for 71% of IPD.

256

257 A statistically significant increasing trend in the distribution of IPD isolates was found
258 in the unadjusted test for serotype 1 IPD ($p=0.029$). However, no other serotypes were
259 found to have significant increasing or decreasing trends.

260

261 *Sequence type analysis*

262

263 The most common STs in IPD between 2003/04 and 2005/06 were 9 (9%), 306 (9%),
264 162 (6%), 53 (5%), 180 (4%), 191 (4%), 124 (4%), 218 (3%), 199 (3%) and 227
265 (3%). ST9 is commonly associated with serotype 14, with approximately 60% of
266 serotype 14 IPD during this period identified as ST9 whilst ST306 is commonly

267 | associated with serotype 1. [There were](#) 158 STs ~~were~~ found in IPD in 2003/04, 140 in
268 | 2004/05 and only 115 in 2005/06, showing a reduction in the diversity of STs over
269 | time.

270

271 | ST306 was found to have a significant increasing trend in the distribution of IPD
272 | isolates between 2003/04 and 2005/06, comparing to the unadjusted significance level
273 | of 0.05 (Table 1, Part A). No other STs showed a significant increasing or decreasing
274 | trend.

Comment [KL3]: Remove unadjusted?

275

276 | ***The effect of PCV7 on the incidence of IPD***

277

278 | Following PCV7 use, the incidence of IPD caused by PCV7 serotypes declined by
279 | 97.4% in children aged <5 years (Table 2). Among those aged 5-64 years and ≥65
280 | years, a significant reduction of VT IPD of 86.3% and 80.4%, respectively, was
281 | observed. In those <5 years and those aged 5-64 years, there was no significant
282 | increase in NVT notifications in 2008/09 compared to the predicted incidence (Figure
283 | 1). In the population aged ≥65 years, a significant increase in NVT disease of 46.5%
284 | was observed. The reduction in VT incidence and increase in NVT incidence resulted
285 | in no change in all-type incidence in this age group.

286

287 | Almost all NVT serotypes exhibited an increase in disease incidence from the last two
288 | pre-vaccination years to 2008/09 (7F: 153.6%, 3: 26.2%, 8: 42.5%, 19A: 78.7%, 22F:
289 | 151.6%, 6A: 31.8%, 12F: 2.3%, 11A: 73.9%, 9N: 33.3%). The exception is serotype 1
290 | which showed a decrease despite the previously reported increasing trend pre-PCV7.
291 | However, only increases in 7F (128.5%; 95% CI (30%, 308.8%)) and 22F (126.7%;

292 95% CI (15%, 356.6%)) were found to be significant when allowing for pre-
293 vaccination trends. The significant decrease of serotype 1 after vaccination was
294 mainly driven by the age groups <5 years and 5-65 years.

295

296 *Trends in the serotype and sequence type distribution post vaccination*

297

298 Post-PCV7, seven serotypes not previously reported in Scotland were noted- 23B (12
299 times), 28 (6), 6C (5), 12 (1), 16A (1), 17A (1), and 35C (1), accounting for 27 of
300 2213 isolates typed. 164 STs which had not previously been reported were noted,
301 amounting to 222 reports of 2203 isolates sequenced. 10% of the isolates sequenced
302 were new STs whilst only 1% of the isolates typed gave rise to new serotypes.

303

304 Amongst the 14 serotypes each accounting for at least 1% of IPD cases post-PCV7
305 (Table 1, Part B), there were significant increasing trends in the distribution amongst
306 IPD isolates for 19A and 22F and decreasing trends for serotypes 1 and 20. Serotype 1
307 decreased at a rate of 29% per year and serotype 20 decreased at a rate of 36% per
308 year. Serotype 19A increased at a rate of 40% per year while 22F increased at a rate
309 of 34% per year.

310

311 There were 11 STs which accounted for more than 1% of all STs reported among IPD
312 cases post-PCV7. ST306 decreased significantly at a rate of 37% per year,
313 comparable with the decrease in its associated serotype 1. ST199 and ST433 both
314 exhibit significant increases post-PCV7 with 25% and 51% increases per year,
315 respectively. ST199 is principally associated with serotype 19A and, to a lesser

316 extent, with 15B whilst ST433 is almost universally associated with serotype 22F.
317 Serotype 20 is principally associated with ST235.

318

319 ***The association between serotypes and STs pre- and post-vaccination***

320

321 The association between serotypes and STs in the period prior to the introduction of
322 PCV7 is shown in Table 3. PCV7 serotypes are associated with a total of 166 STs,
323 however only 12 STs (9, 36, 113, 124, 138, 156, 162, 176, 205, 206, 246, 311)

324 account for the vast majority (74.3%) of the IPD cases. The PCV7 serotypes,
325 associated with these 12 STs (labelled PCV7-HF PCV7-ST), were responsible for 779
326 IPD cases. A further 269 cases of IPD were caused by PCV7 serotypes associated
327 with the remaining 154 STs (labelled PCV7-LF PCV7-ST). ~~In total, 25 serotypes~~
328 ~~(named PCV7-ST serotypes) not present in PCV7, were associated with the 166 STs~~
329 ~~linked to PCV7 serotypes.~~

330

331 ~~There are 708 entries in Table 3, of which only 25 are linked with HF PCV7 ST, 12~~
332 ~~are high frequency STs associated with PCV7. Regarding serotypes not present in~~
333 ~~PCV7, but associated with the 166 STs linked to PCV7, 25 different serotypes were~~
334 ~~responsible for 708 IPD cases, of which only 25 were linked with HF PCV7-ST. The~~
335 ~~and the~~ other 683 ~~were~~ associated with the remaining 154 low frequency STs (cross-
336 classification of PCV7-ST serotypes and LF PCV7-ST). The 25 PCV7-ST serotypes
337 ~~had~~ associations (353 cases) with 151 STs which ~~were~~ not directly associated with
338 PCV7 (cross-classification of PCV7 ST serotypes and NonPCV7-ST). Finally these
339 151 NonPCV7-STs ~~were~~ associated with 22 NonPCV7-ST serotypes (145 cases)
340 which ~~had~~ no direct link with any ST linked to PCV7.

Comment [KL4]: Check number

341

342 Trends in the distribution of groups of serotypes and STs are presented in Figure 2
343 and Figure 3, respectively. Both graphs show a relatively stable distribution in the
344 pre-PCV7 period. The serotype distribution has changed in favour of those serotypes
345 which were associated with STs shown to have had an association with serotypes in
346 the PCV7 vaccine– the PCV7-ST serotypes. Prior to 2006/07, these serotypes formed
347 ~40% of all serotypes but in 2009/10 they formed 80%. The NonPCV7-ST serotypes
348 formed 6% of serotypes prior to 2006/07 and rose to 8% in 2008/09 and 11% in
349 2009/10. The ratio of the percentage of NonPCV7-ST serotypes to the percentage of
350 PCV7-ST serotypes has remained relatively constant over the whole period. The ST
351 distribution has not changed as dramatically but the 12 high frequency STs associated
352 with the PCV7 serotypes are decreasing while the remaining low frequency STs
353 associated with PCV7 and the STs not associated with PCV7 have increased by about
354 10% each. New post PCV7 STs account for ~10% of STs in 2009-10.

355

356 Discussion

357

358 Prior to the introduction of PCV7, the distribution of serotypes and STs among
359 Scottish IPD cases was fairly static, only the proportion of serotype 1 was found to
360 significantly increase, along with a corresponding increase in ST306, among IPD
361 cases. Introduction of routine vaccination with PCV7 drastically reduced the burden
362 of VT IPD in Scotland not only among children targeted for vaccination but also for
363 the rest of the population. Little evidence of serotype replacement was found except
364 for the elderly where the increase in NVT IPD outbalanced the decrease in VT IPD.
365 The major replacement serotypes were 19A and 22F along with the STs 199 and 433.

366 The routine collection of information for both the genetic background and the
367 expressed capsular serotype further allowed an analysis of the relationship in response
368 to vaccine implementation. Interestingly, the proportional increase of serotypes after
369 vaccination was greatly attributable mostly confined to those serotypes which are
370 associated with PCV7 STs.

371

372 One of the key strengths of this study is that the IPD data for Scotland can be
373 considered as a complete national data set as more than 90% of pneumococci
374 isolated from IPD patients in Scotland are sent to the SHLMPRL [28]. Our use of
375 logistic and poisson regression to model linear trends in the serotype and ST
376 distribution enables the identification of changes in the serotype epidemiology.

Comment [KL5]: This is a bit clumsy.
Suggest better phrasing?

377 Our findings on pre and post-vaccination trends of specific serotypes and STs mainly
378 correspond to existing literature. In particular the distribution of serotypes and STs in
379 Scotland prior to the introduction of PCV7 has similarly been reported by Jefferies et
380 al. [18]. Also serotype 1 bacteraemia was found to increase over time in the UK and
381 Ireland [29] as well as serotype 1 associated IPD in England and Wales [25].
382 Furthermore, the increase we observe amongst the proportion of serotype 19A has
383 been widely observed [14-17, 30-32].

384

385 Within four years following PCV7 use, VT serotypes were almost eliminated from
386 IPD cases in those aged <5 years, providing clear evidence of a strong vaccine effect
387 in the targeted age group, as has been documented in other countries [33-35]. In
388 addition, there appears to be evidence of herd protection in those aged 5-64 years, as
389 well as those aged ≥ 65 years. This corresponds with herd protection observed
390 elsewhere, with sustained benefits of PCV7 use in preventing VT serotypes recently

391 | [documented](#) [36]. However, among those aged ≥ 65 years, there is evidence of
392 | serotype replacement with an increase in NVT incidence, as was also shown in the
393 | United States and elsewhere [37, 38]. [It is possible that serotype replacement in those](#)
394 | [aged over 65 years could be attributable to the introduction of PPV23 in this age](#)
395 | [group, however, it does not appear that the timing of the observed decline in VT IPD](#)
396 | [corresponds with the introduction of PPV.](#) Among those aged < 5 years and 5-64
397 | years, the impact of serotype replacement is less clear and is masked by the effect of
398 | serotype 1 which was increasing prior to the introduction of PCV7 and then
399 | decreased. However, even accounting for this, serotype replacement in these age
400 | groups has been less pronounced in Scotland than reported in England and Wales [25]
401 | and elsewhere [39, 40]. It is not clear why the pattern in Scotland is different from
402 | that in England and Wales. A possible reason could be the replacement in the
403 | nasopharynx of Scottish residents by mainly opportunistic NVTs which
404 | predominantly cause invasive disease in those ≥ 65 years of age. Studying changes in
405 | nasopharyngeal carriage before and after introduction of PCV7 as done elsewhere [41,
406 | 42] could shed more light on this.

407

408 | Amongst the non-PCV7 serotypes and the STs not primarily associated with these
409 | serotypes, there is some evidence of a change in the distribution. In particular, as
410 | mentioned, serotype 1 decreased following intervention and was mirrored with a
411 | decrease in the incidence of IPD attributable to ST306. The NVT serotypes, 19A and
412 | 22F were both observed to increase in IPD, whilst serotype 20 showed a significant
413 | decreasing trend. Serotypes 19A and 22F are in the group of serotypes (PCV7-ST
414 | serotypes) linked to the low frequency STs associated with PCV7 (LF PCV7-ST) and
415 | this is the group of serotypes which were shown to increase. Serotype 20 is in the

416 group of serotypes not linked to PCV7 by STs (Non PCV7-ST serotypes) and as a
417 whole this group of serotypes is relatively static in comparison with PCV7-ST
418 serotypes. This implies that there is a possible role of the ST in determining the fitness
419 of a pneumococcus and that it may be possible to predict the serotypes which are
420 likely to increase the most as a result of the introduction of increased valency
421 vaccines. [In interpreting these results, however, it is important to note that the STs](#)
422 [linked to the common disease causing serotypes in the developing world do not](#)
423 [necessarily correspond with those in the developed world \(for example, outbreaks](#)
424 [attributable to serotype 1 in sub-Saharan Africa have been found to be associated with](#)
425 [ST 618 and 217, not 306 and 227 as in the developed world\) \[43\]. Therefore, the](#)
426 [results presented here may not be applicable worldwide.](#)

427

428 The 13-valent PCV contains the seven serotypes found in PCV7, as well as serotypes
429 1, 3, 5, 6A, 7F and 19A. This vaccine was introduced to the paediatric vaccination
430 schedule in the United Kingdom in 2010 and should aid in the prevention of further
431 IPD in Scotland, however as there will be serotypes linked to those in PCV13 through
432 STs associated with PCV13 serotypes, a change in the serotype distribution can
433 perhaps be anticipated due to an increase in those linked serotypes. It is therefore of
434 clear importance to continue to monitor the STs, as well as the serotypes, associated
435 with cases of IPD to aid in determining the potential long-term effectiveness of
436 serotype-specific vaccine interventions and to guide the development of future
437 [vaccinations.](#)

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585

586 Table 1: Results from A) the logistic regression models of serogroups and STs
 587 responsible for at least 1% of IPD between 1999/00 and 2005/06 and between
 588 2003/04 and 2005/06, respectively; B) the logistic regression models of serotypes and
 589 STs responsible for at least 1% of IPD between 2006/2007 and 2009/2010, [examining](#)
 590 [evidence of significant trends in the proportion of IPD attributable to each serogroup,](#)
 591 [serotype and ST.-](#)

Part A:									
Serogroup	Count	OR	95% CI	p-value	ST	Count	OR	95% CI	p-value
14	673	0.94	(0.883, 0.997)	0.003	9	213	1.06	(0.804, 1.402)	0.539
9	364	0.97	(0.892, 1.047)	0.230	306	174	1.40	(1.042, 1.869)	0.001
1	331	1.36	(1.238, 1.493)	<0.001	162	145	1.03	(0.741, 1.432)	0.797
6	301	0.97	(0.891, 1.057)	0.328	53	126	1.01	(0.700, 1.464)	0.924
19	290	0.98	(0.900, 1.074)	0.595	180	96	1.01	(0.678, 1.508)	0.939
4	273	1.04	(0.946, 1.134)	0.284	191	95	1.24	(0.823, 1.875)	0.134
8	247	0.95	(0.865, 1.044)	0.135	124	85	1.00	(0.658, 1.515)	0.987
23	242	0.94	(0.858, 1.035)	0.084	218	74	1.24	(0.784, 1.956)	0.183
3	220	1.00	(0.902, 1.100)	0.917	199	68	0.72	(0.444, 1.151)	0.045
7	162	1.04	(0.926, 1.167)	0.357	227	63	1.22	(0.750, 1.990)	0.244
18	158	0.98	(0.876, 1.104)	0.685	311	63	0.92	(0.561, 1.498)	0.615
12	131	1.04	(0.914, 1.183)	0.400	246	58	0.97	(0.593, 1.601)	0.883
22	107	0.98	(0.849, 1.125)	0.648	433	48	0.60	(0.314, 1.132)	0.022
20	83	0.95	(0.808, 1.113)	0.360	205	41	1.21	(0.654, 2.228)	0.386
33	71	0.94	(0.789, 1.110)	0.285	176	41	1.18	(0.620, 2.237)	0.468
11	65	0.97	(0.811, 1.160)	0.638	206	40	1.22	(0.652, 2.271)	0.372
15	51	1.01	(0.829, 1.239)	0.864	113	38	1.02	(0.530, 1.965)	0.930
					235	35	0.77	(0.390, 1.531)	0.284
					36	32	1.19	(0.572, 2.475)	0.499
					138	30	1.16	(0.581, 2.295)	0.552
					62	30	1.14	(0.531, 2.429)	0.636
					65	25	1.63	(0.691, 3.838)	0.106
Part B:									
Serotype	Count	OR	95% CI	p-value	ST	Count	OR	95% CI	p-value
1	210	0.71	(0.58, 0.86)	<0.001	306	139	0.73	(0.58, 0.92)	<0.001
8	162	0.85	(0.68, 1.05)	0.026	191	217	1.16	(0.96, 1.39)	0.026
7F	240	1.11	(0.93, 1.34)	0.091	53	123	0.90	(0.71, 1.14)	0.195
3	173	1.00	(0.81, 1.23)	0.997	180	135	1.08	(0.86, 1.35)	0.343
19A	165	1.40	(1.11, 1.75)	<0.001	199	128	1.25	(0.98, 1.58)	0.008
22F	130	1.34	(1.04, 1.72)	<0.001	433	88	1.51	(1.12, 2.04)	<0.001
12F	79	1.10	(0.81, 1.49)	0.372	218	59	1.00	(0.72, 1.40)	0.995
6A	74	0.86	(0.63, 1.17)	0.161	227	46	0.85	(0.58, 1.25)	0.238
9N	48	0.90	(0.62, 1.32)	0.434	62	45	1.15	(0.78, 1.69)	0.306
11A	54	1.08	(0.75, 1.56)	0.514	235	23	0.80	(0.47, 1.36)	0.232

20	35	0.64	(0.40, 1.02)	0.005		65	23	0.74	(0.43, 1.28)	0.119
33F	43	1.13	(0.75, 1.70)	0.397						
15B	31	1.16	(0.72, 1.89)	0.362						
23A	28	0.96	(0.58, 1.57)	0.800						

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594 Note: Count is the number of serogroups and STs among IPD cases in the pre-PCV7
595 period in Part A and serotypes and STs among IPD cases in the post-PCV7 period in
596 Part B; OR – Odds Ratio associated with a one year change; CI- 95% Bonferroni
597 adjusted confidence interval; *p*-value is the unadjusted *p*-value. The entries in bold
598 typeface are those with *p*-values below the Benjamini and Hochberg adjusted *p*-value.

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600 Table 2: Incidence rates of the most common non-vaccine type (NVT) IPD serotypes
 601 and vaccine type (VT) serotypes in Scotland from 2004/05 to 2009/10.

602

	Incidence 2004/05	Incidence 2005/06	Incidence 2009/10	Change 2009/10 predicted compared to observed	Change 2009/10 predicted compared to observed (serotype 1 excluded)
0-4 years					
All	38.23	27.35	14.18	-68.5% (-80.4, -50.0)	-69.8% (-81.7, -50.8)
NVT	12.87	7.12	13.49	-39.6% (-71.4, 28.6)	-13.4% (-62.4, 102.7)
VT	25.36	20.24	0.69	-97.4% (-99.6, -91.3)	-97.5% (-99.6, -91.4)
5-64 years					
All	8.11	9.52	6.59	-57.2% (-65.5, -46.9)	-42.1% (-54.1, -26.9)
NVT	4.73	6.22	5.92	-45.6% (-58.3, -29.1)	3.4% (-23.7, 40.3)
VT	3.38	3.30	0.67	-86.3% (-91.6, -78.4)	-87.0% (-92.0, -79.5)
65+ years					
All	31.09	32.08	26.48	-4.9% (-24.4, 19.5)	0.0% (-20.7, 26.1)
NVT	15.06	17.00	24.12	+46.5% (9.0, 97.4)	64.7% (21.4, 124.0)
VT	16.03	15.08	2.30	-80.4% (-88.6, -67.9)	-80.5% (-88.6, -68.0)

603

604 Notes: The percentage change is a comparison of the predicted incidence in 2009/10
 605 to the observed incidence in 2009/10 adjusting for the temporal trend pre-vaccination
 606 (see Methods). 95% confidence intervals for the percentage changes are derived from
 607 the Poisson regression model. When serotype 1 is excluded, the percentage changes
 608 for VT differ slightly because inflation in this case is assumed to distribute over all
 609 types.

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614 Table 3: The association between ST and serotype among IPD cases in Scotland in the
 615 period prior to the introduction of PCV7 in September 2006.

616

		STs associated with PCV7 Serotypes													STs not associated with PCV7 Serotypes	
		HF PCV7-ST [12 STs]												LF PCV7-ST	Non PCV7-ST	
		9	36	113	124	138	156	162	176	205	206	246	311	[154 STs]	[151 STs]	
PCV7 Serotypes	PNE004	0	0	0	0	0	0	0	0	40	37	56	0	269 entries	0	
	PNE06B	0	0	0	0	25	1	0	37	0	0	0	0			
	PNE09V	0	0	0	1	0	13	102	0	0	1	0	0			
	PNE014	208	0	0	84	0	4	1	0	0	0	0	1			
	PNE18C	0	0	36	0	0	0	0	1	0	0	0	0			
	PNE19F	1	0	0	0	0	0	35	0	0	0	0	0			
	PNE23F	0	32	0	0	0	0	0	1	0	0	0	62			
PCV7-ST serotypes	Serotypes not in PCV7 but associated with the STs linked to PCV7	25 entries over all serotypes and all 12 sequence types												683 entries	353 entries	
NonPCV7-ST serotypes	Serotypes not associated with any ST linked to PCV7	0												0	145 entries	

Serotypes not in PCV7 but associated with the STs linked to PCV7:

PNE001 PNE003 PNE005 PNE008 PNE020 PNE034 PNE038 PNE06A PNE07A PNE07C PNE07F PNE09A PNE09N PNE11A PNE15A PNE15B PNE15C PNE16F PNE17F PNE18B PNE18F PNE19A PNE22F PNE33C PNE33F

Serotypes not associated with any ST linked to PCV7:

PNE002 PNE013 PNE021 PNE024 PNE027 PNE029 PNE031 PNE037 PNE041 PNE042 PNE10A PNE10F PNE12A PNE12B PNE12F PNE18A PNE22A PNE23A PNE24F PNE28F PNE35B PNE35F

STs in the HF PCV7-ST group had more than 10 reports of co-occurrence with PCV7 serotypes. There are 779 entries for these 12 STs. In the full matrix there are 1048 reports among 166 STs associated with PCV7.

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		STs associated with PCV7 Serotypes												STs not associated with PCV7 Serotypes	
		HF PCV7-ST [12 STs]											LF PCV7-ST	Non PCV7-ST	
		9	36	113	124	138	156	162	176	205	206	246	311	[154 STs]	[151 STs]
PCV7 Serotypes	PNE004	0	0	0	0	0	0	0	0	40	37	56	0	269 entries	0
	PNE06B	0	0	0	0	25	1	0	37	0	0	0	0		
	PNE09V	0	0	0	1	0	13	102	0	0	1	0	0		
	PNE014	208	0	0	84	0	4	1	0	0	0	0	1		
	PNE18C	0	0	36	0	0	0	0	1	0	0	0	0		
	PNE19F	1	0	0	0	0	0	35	0	0	0	0	0		
	PNE23F	0	32	0	0	0	0	0	1	0	0	0	62		
PCV7-ST serotypes	25 Serotypes not in PCV7 but associated with the STs linked to PCV7	25 entries over all 25 serotypes and all 12 sequence types											683 entries	353 entries	
NonPCV7-ST serotypes	22 different Serotypes not associated with any ST linked to PCV7	0											0	145 entries	

Serotypes not in PCV7 but associated with the STs linked to PCV7:

PNE001 PNE003 PNE005 PNE008 PNE020 PNE034 PNE038 PNE06A PNE07A PNE07C PNE07F PNE09A PNE09N PNE11A PNE15A PNE15B PNE15C PNE16F PNE17F PNE18B PNE18F PNE19A PNE22F PNE33C PNE33F

Serotypes not associated with any ST linked to PCV7:

PNE002 PNE013 PNE021 PNE024 PNE027 PNE029 PNE031 PNE037 PNE041 PNE042 PNE10A PNE10F PNE12A PNE12B PNE12F PNE18A PNE22A PNE23A PNE24F PNE28F PNE35B PNE35F

STs in the HF PCV7-ST group had more than 10 reports of co-occurrence with PCV7 serotypes. There are 779 entries for these 12 STs. In the full matrix there are 1048 reports among 166 STs associated with PCV7.

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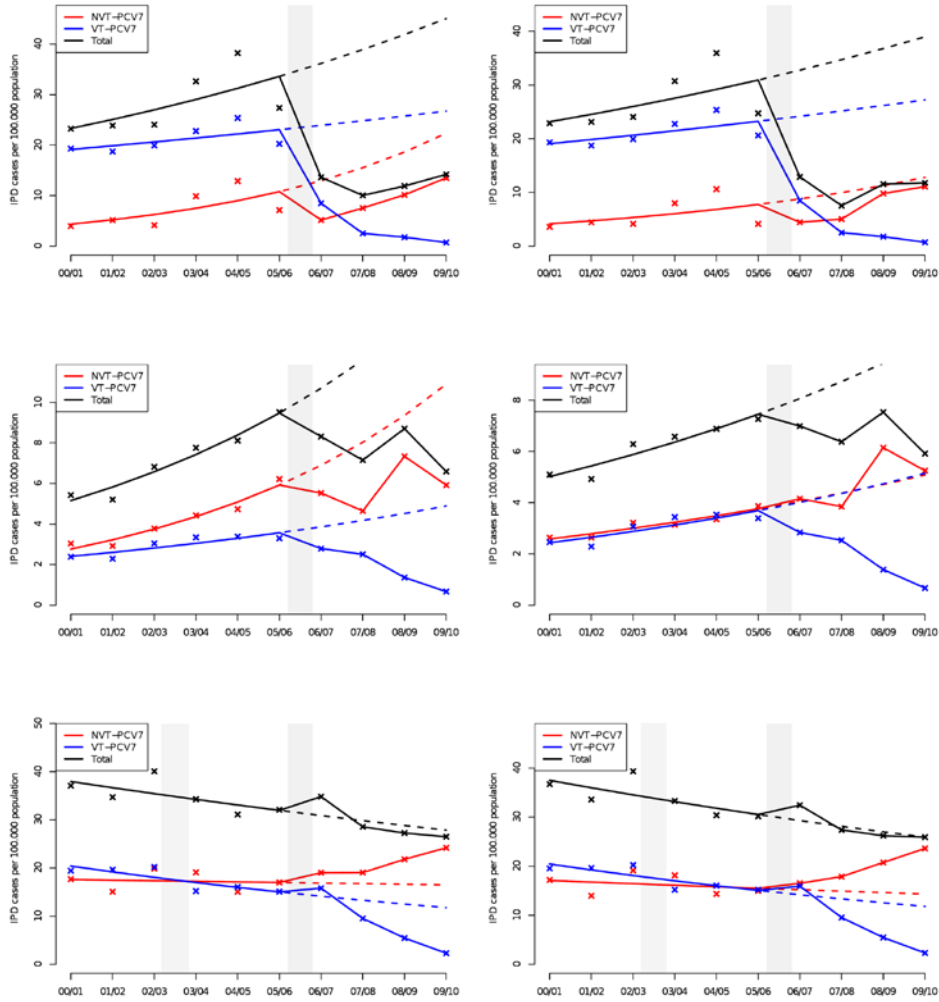
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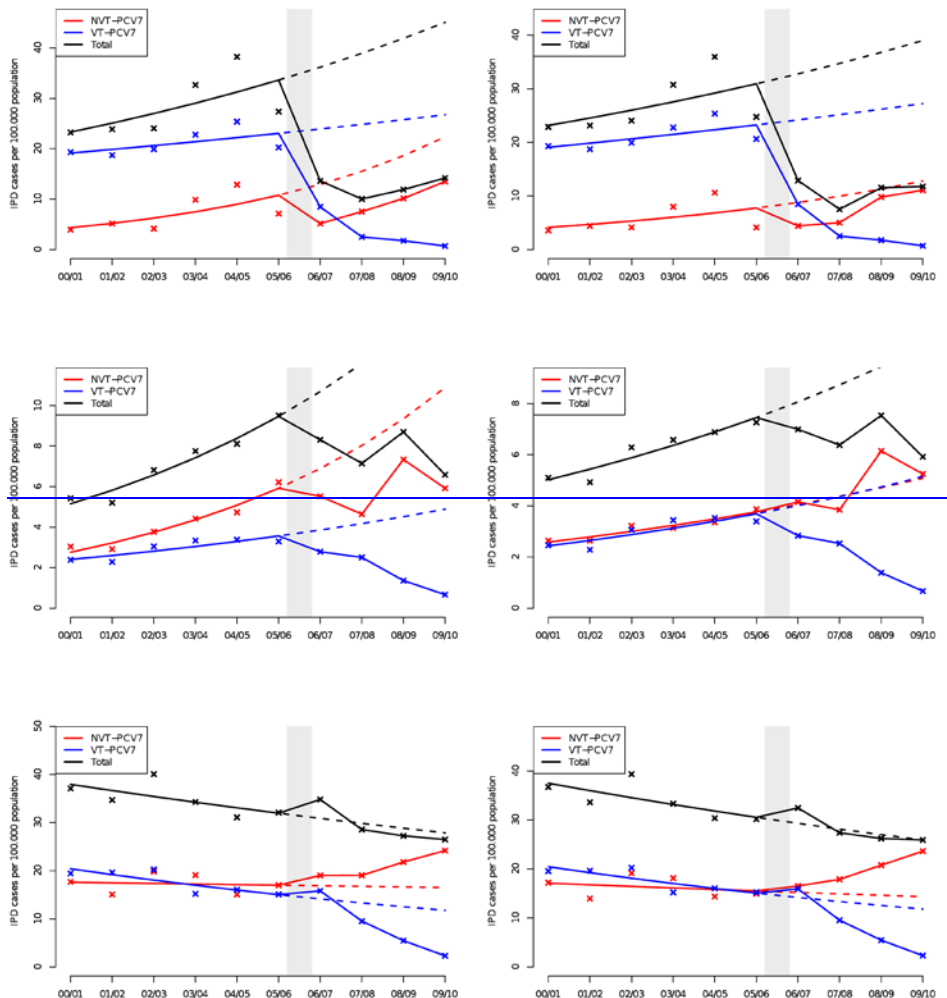
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628 Figure 1: Incidence rates of vaccine type (VT) and non-vaccine type (NVT) IPD in
 629 Scotland, by age group.



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632 Note: All serotypes are included in the left hand column of graphs while serotype 1 is
 633 excluded from the right hand column of graphs. The top row are for those aged 0-4
 634 years, the middle row for those aged 5-64 years and the bottom row for those aged
 635 65+ years. The grey vertical bar denotes the introduction of PCV7. [For those aged](#)
 636 [65+ years, the first grey vertical bar denotes the introduction of PPV23.](#) The data are
 637 plotted as crosses, the dashed lines show the predicted post vaccination incidence
 638 based on pre vaccination trends and the predicted values from the poisson regression
 639 model are the points joined by the lines.

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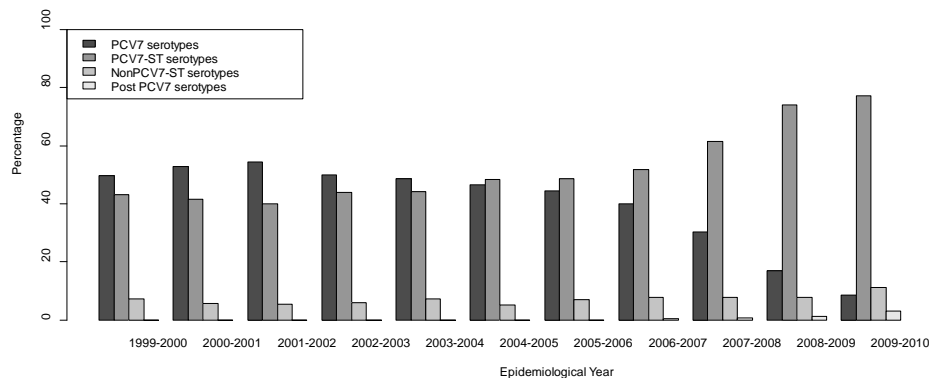
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663 Figure 2: Trends in the serotype distribution from 1999/2000 to 2009/2010.



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665 Note: In the period 1999-2002 when only serogroup was available, the serotypes of
 666 PCV7 serogroups were imputed based upon the distribution of serotypes within
 667 serogroups in the period 2003-2006. A sensitivity analysis showed that this had
 668 minimal impact on the distributions presented.

669 PCV7 – serotypes in the PCV7 vaccine.

670 PCV7 serotypes – serotypes in the PCV7 vaccine.

671 PCV7-ST serotypes – serotypes not in the PCV7 vaccine but which are associated
 672 with STs associated with the PCV7 serotypes.

673 NonPCV7-ST serotypes – the remaining serotypes present among IPD cases prior to
 674 the introduction of PCV7. These serotypes are not associated with any ST connected
 675 with the 7 PCV7 serotypes.

676 Post PCV7 serotypes – serotypes which have emerged post PCV7 (post September
 677 2006).

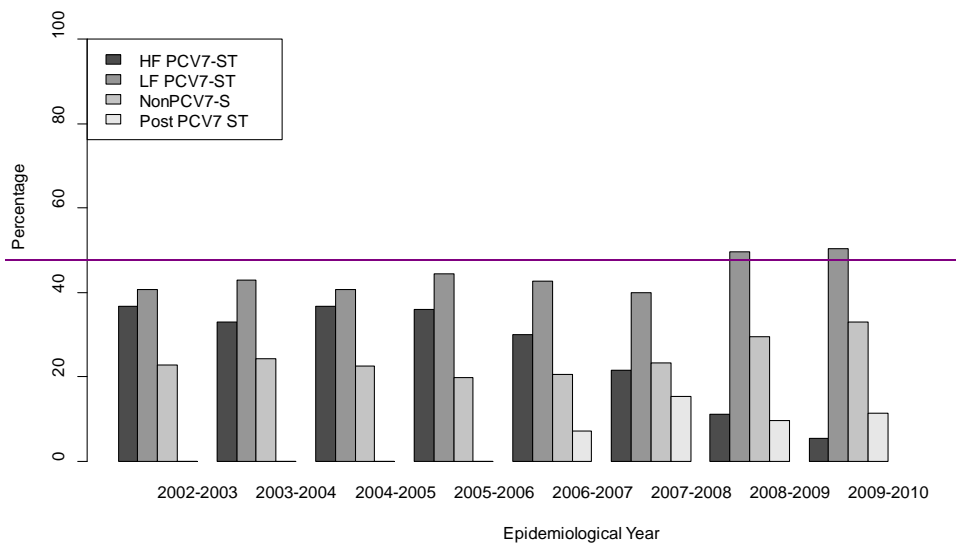
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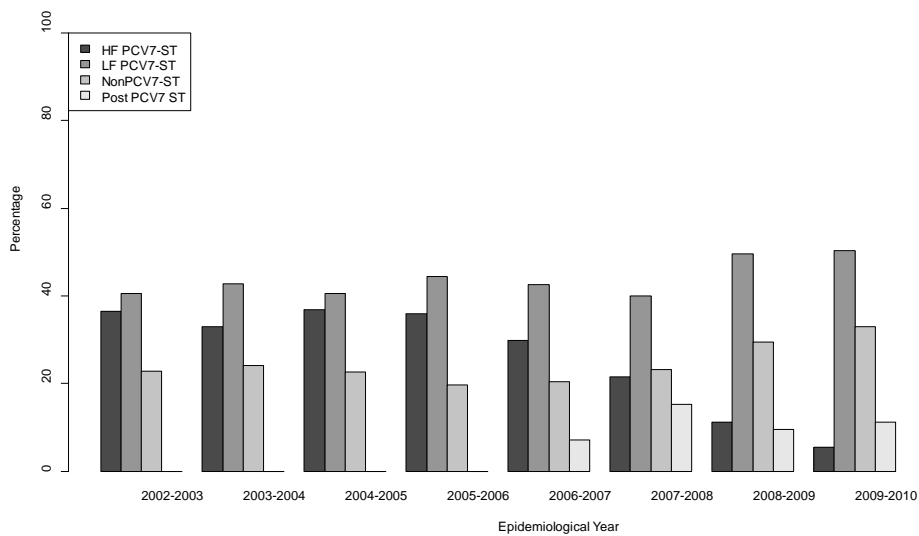
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682 Figure 3: Trends in the ST distribution from 2002/2003 to 2009/2010.



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685 Note: [HF PCV7-ST](#) ~~Main-PCV7~~ – The 12 STs with a strong association with the
 686 PCV7 serotypes in the PCV7 vaccine. [LF PCV7-ST](#) ~~Rest-PCV7~~ – The remaining STs
 687 associated with the PCV7 vaccine serotypes. [NonPCV7-ST](#) - STs not associated with
 688 any serotype in the PCV7 vaccine.

689 Post PCV7 – STs which have emerged post PCV7 (post September 2006).

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699 CONFLICTS OF INTEREST:

700 KEL was funded through an EPSRC CASE studentship with Wyeth Pharmaceuticals.

701 SF and MD have no conflicts to declare. DG has received funding to support a PhD

702 studentship from Wyeth Pharmaceuticals. SCC currently receives unrestricted

703 research funding from Pfizer Vaccines (previously Wyeth Vaccines). JMJ and SCC

704 have received consulting fees from GlaxoSmithKline and have received financial

705 assistance from vaccine manufacturers to attend conferences. All grants and honoraria

706 are paid into accounts within the respective NHS Trusts or Universities, or to

707 independent charities. JMJ, TJM, SCC, AS and GFSE previously received funding

708 from Wyeth Pharmaceuticals for a collaborative project with the Institute of

709 Biological Sciences, University of Glasgow and the Scottish Meningococcal and

710 Pneumococcal Reference Laboratory (2005-2007). BD, JM and EM have no conflicts

711 to declare. CR has received research funding from and has acted as a consultant for

712 Wyeth Pharmaceuticals.