Carriage dynamics of pneumococcal serotypes in naturally colonised infants in a rural African setting during the first year of life

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Supplementary Table 1. Clearance and acquisition rates of pneumococcal serotypes in the Gambian infants.

| Serotype | Clearance rate (per day) | | | Time from birth to first | Time from birth to reacquisition |
|------------|--------------------------|------------------------|-------------------------|------------------------------|----------------------------------|
| | All episodes (95% CI) | First episode (95% CI) | Second episode (95% CI) | acquisition in days (95% CI) | in days (95% CI) |
| 6B | 0.0116 (0.0072,0.0187) | 0.0111 (0.0066,0.0187) | 0.0153 (0.0049,0.0475) | 0.0119 (0.007,0.02) | 0.0067 (0.0022,0.0208) |
| 4 | 0.0159 (0.0066,0.0381) | 0.0159 (0.0066,0.0381) | _ | 0.0057 (0.0024,0.0136) | - |
| 13 | 0.0168 (0.0113,0.0251) | 0.0175 (0.0115,0.0265) | _ | 0.0062 (0.0041,0.0095) | _ |
| 23F | 0.0174 (0.0103,0.0294) | 0.0169 (0.0094,0.0305) | 0.0195 (0.0063,0.0604) | 0.0095 (0.0052,0.0171) | 0.0045 (0.0015,0.014) |
| 19A | 0.0178 (0.0146,0.0217) | 0.0157 (0.0125,0.0198) | 0.0279 (0.019,0.041) | 0.0075 (0.0059,0.0094) | 0.0055 (0.0036,0.0085) |
| 9L | 0.0185 (0.0111,0.0306) | 0.0184 (0.0105,0.0325) | 0.0186 (0.006,0.0578) | 0.0101 (0.0057,0.0178) | - |
| 20 | 0.0209 (0.0112,0.0388) | 0.0177 (0.0089,0.0354) | | 0.0086 (0.0043,0.0172) | _ |
| 39 | 0.0213 (0.0101,0.0446) | 0.026 (0.0117,0.0578) | - | 0.0077 (0.0034,0.017) | - |
| 35B | 0.0215 (0.0148,0.0311) | 0.0257 (0.0174,0.038) | 0.0091 (0.0029,0.0283) | 0.006 (0.004,0.0089) | 0.0079 (0.0026,0.0246) |
| 21 | 0.0231 (0.0159,0.0334) | 0.0239 (0.016,0.0356) | 0.019 (0.0071,0.0508) | 0.0085 (0.0057,0.0126) | - |
| 6A | 0.0232 (0.0185,0.0291) | 0.0222 (0.0172,0.0286) | 0.0282 (0.017,0.0468) | 0.009 (0.007,0.0116) | 0.005 (0.003,0.0085) |
| 11A | 0.0241 (0.0164,0.0354) | 0.0403 (0.0265,0.0612) | 0.0075 (0.0028,0.02) | 0.0059 (0.0039,0.009) | 0.0048 (0.0018,0.0129) |
| 10A | 0.0243 (0.0151,0.0391) | 0.0223 (0.0135,0.037) | _ | 0.0051 (0.0031,0.0084) | _ |
| 23B | 0.0247 (0.0175,0.0349) | 0.0288 (0.0197,0.042) | 0.014 (0.0058,0.0336) | 0.0088 (0.006,0.0128) | 0.0058 (0.0024,0.0138) |
| 22A | 0.026 (0.0148,0.0457) | 0.0286 (0.0154,0.0531) | - | 0.0087 (0.0047,0.0162) | _ |
| 14 | 0.026 (0.0176,0.0384) | 0.0284 (0.0181,0.0446) | 0.0204 (0.0092,0.0454) | 0.0083 (0.0053,0.0131) | 0.0081 (0.0034,0.0195) |
| 16F | 0.0271 (0.0182,0.0404) | 0.0336 (0.0221,0.051) | - | 0.0065 (0.0043,0.0099) | _ |
| 15B/C | 0.0276 (0.0204,0.0374) | 0.031 (0.0223,0.043) | 0.0168 (0.0076,0.0374) | 0.0055 (0.004,0.0076) | 0.004 (0.0018,0.0089) |
| 15A | 0.028 (0.0176,0.0444) | 0.0337 (0.021,0.0543) | _ | 0.0068 (0.0043,0.011) | _ |
| 19F | 0.0281 (0.0199,0.0395) | 0.0286 (0.02,0.0409) | 0.0238 (0.0077,0.0738) | 0.0059 (0.0042,0.0085) | 0.0043 (0.0014,0.0134) |
| 9A | 0.0286 (0.0136,0.0599) | 0.0286 (0.0136,0.0599) | | 0.0085 (0.004,0.0178) | _ |
| 34 | 0.0293 (0.0206,0.0417) | 0.0282 (0.0196,0.0406) | - | 0.0085 (0.0059,0.0123) | - |
| 48 | 0.0306 (0.0138,0.0681) | 0.0275 (0.0114,0.066) | — | 0.0143 (0.0059,0.0343) | - |
| 18A | 0.0312 (0.014,0.0694) | 0.0312 (0.014,0.0694) | — | 0.0193 (0.0087,0.0429) | - |
| 7 F | 0.0317 (0.0119,0.0846) | 0.0317 (0.0119,0.0846) | — | 0.0053 (0.002,0.0142) | _ |
| 17F | 0.0317 (0.0176,0.0573) | 0.0314 (0.0169,0.0584) | — | 0.0051 (0.0027,0.0095) | - |
| 9V | 0.0336 (0.0168,0.0672) | 0.0313 (0.0149,0.0656) | _ | 0.0059 (0.0028,0.0124) | _ |
| 40 | 0.0342 (0.0189,0.0617) | 0.0325 (0.0175,0.0603) | - | 0.0065 (0.0035,0.012) | - |
| 12F | 0.0357 (0.0179,0.0714) | 0.0357 (0.0179,0.0714) | - | 0.0085 (0.0043,0.0171) | - |
| 23A | 0.0371 (0.0216,0.064) | 0.0357 (0.0203,0.0629) | - | 0.0086 (0.0049,0.0152) | - |
| 58 | 0.03/8 (0.019/,0.0727) | 0.0408 (0.0204,0.0816) | - | 0.0069 (0.0034,0.0138) | - |
| | 0.0381(0.0191,0.0762) | 0.0357 (0.017,0.0749) | - | 0.0053 (0.0025,0.011) | - |
| 3 | 0.0401(0.0201,0.0802) | 0.05/1 (0.02/2,0.1199) | - | 0.0086(0.0041, 0.018) | - |
| 180 | 0.0417(0.0199,0.0874) | 0.039 (0.0175,0.0867) | - | 0.0126 (0.0057,0.0281) | - |
| 11D 255 | 0.0429 (0.0138,0.1329) | 0.0429 (0.0138,0.1329) | - | 0.0055 (0.0018,0.017) | - |
| 35F | 0.0476 (0.0179,0.1269) | 0.04/6 (0.01/9,0.1269) | - | 0.006 (0.0022,0.0159) | - |

| NT | 0.0505 (0.0382,0.0669) | 0.0466 (0.0339,0.064) | 0.0714 (0.0396,0.129) | 0.0093 (0.0068,0.0128) | 0.0043 (0.0024,0.0077) |
|-----|------------------------|------------------------|-----------------------|------------------------|------------------------|
| 28F | 0.0571 (0.0272,0.1199) | 0.0571 (0.0272,0.1199) | _ | 0.009 (0.0043,0.019) | _ |
| 8 | 0.0595 (0.0248,0.143) | 0.0595 (0.0248,0.143) | _ | 0.0089 (0.0037,0.0215) | _ |
| 9N | 0.0625 (0.0298,0.1311) | 0.0612 (0.0275,0.1363) | _ | 0.0057 (0.0026,0.0127) | _ |
| 1 | 0.0635 (0.0318,0.127) | 0.0635 (0.0318,0.127) | _ | 0.0106 (0.0053,0.0212) | _ |
| 41 | 0.0714 (0.023,0.2215) | _ | - | - | - |
| 19B | 0.0714 (0.0268,0.1903) | 0.0714 (0.0268,0.1903) | _ | 0.0063 (0.0024,0.0167) | _ |
| 47 | 0.0714 (0.0268,0.1903) | 0.0714 (0.023,0.2215) | _ | 0.0102 (0.0033,0.0316) | _ |
| 47A | 0.0714 (0.0268,0.1903) | 0.0714 (0.0268,0.1903) | _ | 0.0136 (0.0051,0.0363) | _ |
| 5 | 0.0714 (0.0268,0.1903) | 0.0714 (0.0268,0.1903) | - | 0.0053 (0.002,0.0141) | - |
| 10F | 0.0714 (0.0297,0.1716) | 0.0714 (0.0297,0.1716) | _ | 0.0069 (0.0029,0.0167) | _ |
| 16A | 0.0714 (0.0297,0.1716) | 0.0714 (0.0297,0.1716) | _ | 0.0038 (0.0016,0.009) | _ |
| 36 | 0.0714 (0.0321,0.159) | 0.0714 (0.0321,0.159) | _ | 0.0126 (0.0057,0.0281) | _ |
| 19C | 0.0714 (0.0396,0.129) | 0.0714 (0.0396,0.129) | _ | 0.0047 (0.0026,0.0085) | _ |
| 47F | 0.0779 (0.0251,0.2416) | 0.0779 (0.0251,0.2416) | _ | 0.007 (0.0022,0.0216) | _ |

Cells marked by "–" designate serotypes where no or very few data were available for analysis.



Supplementary Figure 1. Proportion of colonisation episodes associated with each serotype.



Supplementary Figure 2. Clearance and acquisition rates of pneumococcal serotypes in Gambian infants. Graph showing (a) clearance rate, (b) initial acquisition rate of serotypes, (b) acquisition rate of the second serotype, (c) rates of acquisition of a serotype for the first and second episode and (d) clearance rates of the first and second episode of a serotype.



Supplementary Figure 3. The Kaplan-Meier survival curves and exponential fit to the carriage duration data of each serotype. The plots show the overall duration from acquisition to clearance of each serotypes regardless of acquisition time. The black curve represents the Kaplan-Meier estimates while the yellow line is the fitted survival curve. The rate parameter in the exponential model is shown at the top of each plot represents the mean clearance rate whose inverse equates to mean the carriage duration.



Supplementary Figure 4. The Kaplan-Meier survival curves and exponential fit to the time to first acquisition data of each serotype. The plots show duration from birth of an infant to first acquisition of each serotype. The black curve represents the Kaplan-Meier estimates while the yellow line is the fitted survival curve. The rate parameter in the exponential model is shown at the top of each plot represents the mean acquisition rate whose inverse equates to mean the time to first acquisition of each serotype.



Supplementary Figure 5. The Kaplan-Meier survival curves and exponential fit to the carriage duration data after first acquisition for each serotype. The plots show duration from first acquisition until clearance of each serotype. The black curve represents the Kaplan-Meier estimates while the yellow line is the fitted survival curve. The rate parameter in the exponential model is shown at the top of each plot represents the mean clearance rate after first episode whose inverse equates to mean the time to acquisition of each serotype.



Supplementary Figure 6. The Kaplan-Meier survival curves and exponential fit to the time to second acquisition or reacquisition data for each serotype. The plots show duration from birth of an infant until second acquisition of each serotype. The black curve represents the Kaplan-Meier estimates while the yellow line is the fitted survival curve. The rate parameter in the exponential model is shown at the top of each plot represents the mean reacquisition rate after whose inverse equates to mean the time to reacquisition of each serotype.



Supplementary Figure 7. The Kaplan-Meier survival curves and exponential fit to the carriage duration data after second acquisition or reacquisition for each serotype. The plots show duration from first acquisition until clearance of each serotype. The black curve represents the Kaplan-Meier estimates while the yellow line is the fitted survival curve. The rate parameter in the exponential model is shown at the top of each plot represents the mean clearance rate after first episode whose inverse equates to mean the time to acquisition of each serotype.



Supplementary Figure 8. The Kaplan-Meier survival curves comparing the carriage duration of the first episode and second episode of serotypes acquired multiple times. The *P*-values were calculated using the log-rank test.



Supplementary Figure 9. The Kaplan-Meier survival curves comparing the carriage duration for each serotype in different communities. The infants in group 1 were from PCV-unexposed (unvaccinated) communities and received PCV7 after 6 months. The infants in group 2 were also from PCV-unexposed (unvaccinated) communities where the infants received PCV7 at 2, 3 and 4 months. The infants from group 3 originated from PCV-exposed (vaccinated) communities and they received PCV7 at 2, 3 and 4 months. The P-values were calculated using the log-rank test. Only serotypes found in both communities in >3 colonisation episodes are shown in the figure.



Supplementary Figure 10. The Kaplan-Meier survival curves comparing the time to acquisition for each serotype in different communities. The infants in community 1 were from PCV-unexposed (unvaccinated) villages and received PCV7 after 6 months. The infants in community 2 were also from PCV-unexposed (unvaccinated) villages where the infants received PCV7 at 2, 3 and 4 months. The infants from community 3 originated from PCV-exposed (vaccinated) villages and they received PCV7 at 2, 3 and 4 months. The *P*-values were calculated using the log-rank test. Only serotypes found in both communities in >3 colonisation episodes are shown in the figure.