Impact of meningococcal ACWY conjugate vaccines on pharyngeal carriage in adolescents: evidence for herd protection from the UK MenACWY programme

Jeremy P. Carr, MBBS, Jenny M. MacLennan, MA, Emma Plested, Holly B. Bratcher, DPhil, Odile B. Harrison, PhD, Parvinder K. Aley, PhD, James E. Bray, PhD, Susana Camara, PhD, Charlene M.C. Rodrigues, DPhil, Kimberly Davis, MBBS, Angela Bartolf, MBBS, David Baxter, PhD, J. Claire Cameron, FFPH PhD, Richard Cunningham, PhD, Saul N. Faust, PhD, Katy Fidler, PhD, Rohit Gowda, MBBS, Paul T. Heath, PhD, Stephen Hughes, PhD, Sujata Khajuria, MBBS, David Orr, FRCPATH, Mala Raman, MBBS, Andrew Smith, PhD, David PJ. Turner, PhD, Elizabeth Whittaker, PhD, Christopher J. Williams, MBBS, Christos S. Zipitis, MBBS, Andrew J. Pollard, FMedSci, Jennifer Oliver, PhD, Begonia Morales-Aza, BSc, Aiswarya Lekshmi, MSc, Stephen A. Clark, PhD, Ray Borrow, PhD, Hannah Christensen, PhD, Caroline Trotter, PhD, Adam Finn, PhD, Martin C.J. Maiden, FMedSci, Matthew D. Snape, MD, for the UKMenCar4 and 'Be on the TEAM' Study Collaborators

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Impact of meningococcal ACWY conjugate vaccines on pharyngeal carriage in adolescents: evidence for herd protection from the UK MenACWY programme

Jeremy P Carr MBBS

Oxford Vaccine Group, Department of Paediatrics, University of Oxford, and the National Institute for Health Research (NIHR) Oxford Biomedical Research Centre, Oxford, UK. Monash University, Melbourne, Australia. Monash Children's Hospital, Melbourne, Australia.

Jenny M MacLennan MA

Department of Zoology, University of Oxford, UK.

Emma Plested

Oxford Vaccine Group, Department of Paediatrics, University of Oxford, and the National Institute for Health Research (NIHR) Oxford Biomedical Research Centre, Oxford, UK.

Holly B Bratcher DPhil

Department of Zoology, University of Oxford, UK.

Odile B Harrison PhD

Department of Zoology, University of Oxford, UK.

Parvinder K Aley PhD

Oxford Vaccine Group, Department of Paediatrics, University of Oxford, and the National Institute for Health Research (NIHR) Oxford Biomedical Research Centre, Oxford, UK.

James E Bray PhD

Department of Zoology, University of Oxford, UK.

Susana Camara PhD

Oxford Vaccine Group, Department of Paediatrics, University of Oxford, and the National Institute for Health Research (NIHR) Oxford Biomedical Research Centre, Oxford, UK.

Charlene M C Rodrigues DPhil

Department of Zoology, University of Oxford, UK.

Department of Paediatric Infectious Diseases, Great Ormond Street Hospital for Children NHS Foundation Trust, UK.

Kimberly Davis MBBS

Oxford Vaccine Group, Department of Paediatrics, University of Oxford, and the National Institute for Health Research (NIHR) Oxford Biomedical Research Centre, Oxford, UK.

Angela Bartolf MBBS

St George's Vaccine Institute, Institute of Infection & Immunity St George's University of London, UK.

David Baxter PhD

Stockport NHS Foundation Trust, UK.

J. Claire Cameron FFPH PhD

Public Health Scotland, UK.

Richard Cunningham PhD

University Hospitals Plymouth NHS Trust, Plymouth, UK.

Saul N Faust PhD

NIHR Southampton Clinical Research Facility and NIHR Southampton Biomedical Research Centre, University Hospital Southampton NHS Foundation Trust; and Faculty of Medicine and Institute for Life Sciences, University of Southampton, Southampton, UK.

Katy Fidler PhD

Brighton and Sussex Medical School, UK.

Royal Alexandra Children's Hospital, University Hospital Sussex NHS Foundation Trust, Brighton, UK.

Rohit Gowda MBBS

Maidstone and Tunbridge Wells NHS Trust, UK.

Paul T. Heath PhD

St George's Vaccine Institute, Institute of Infection & Immunity, St George's University of London, UK.

Stephen Hughes PhD

Royal Manchester Children's Hospital Manchester University NHS Foundation Trust, UK

Sujata Khajuria MBBS

Northamptonshire Healthcare NHS Foundation Trust, UK.

David Orr FRCPATH

Lancashire Teaching Hospitals NHS Foundation Trust, UK.

Mala Raman MBBS

University Hospitals Plymouth NHS Foundation Trust, UK.

Andrew Smith PhD

Glasgow Dental Hospital & School, College of Medical, Veterinary & Life Sciences, University of Glasgow, UK.

David PJ Turner PhD

School of Life Sciences, University of Nottingham & Nottingham University Hospitals NHS Trust, UK

Elizabeth Whittaker PhD

Imperial College London, UK. Imperial College Healthcare NHS Trust, London, UK.

Christopher J. Williams MBBS

Communicable Disease Surveillance Centre, Public Health Wales, Cardiff, UK.

Christos S. Zipitis MBBS

Wrightington, Wigan and Leigh Teaching Hospitals NHS Foundation Trust, UK.

Andrew J Pollard FMedSci

Oxford Vaccine Group, Department of Paediatrics, University of Oxford, and the National Institute for Health Research (NIHR) Oxford Biomedical Research Centre, Oxford, UK.

Jennifer Oliver PhD

Bristol Children's Vaccine Centre, University of Bristol, UK.

Begonia Morales-Aza BSc

Bristol Children's Vaccine Centre, University of Bristol, UK.

Aiswarya Lekshmi MSc

UK Health Security Agency Meningococcal Reference Unit, Manchester Royal Infirmary Manchester, UK.

Stephen A. Clark PhD

UK Health Security Agency Meningococcal Reference Unit, Manchester Royal Infirmary Manchester, UK.

Ray Borrow PhD

UK Health Security Agency Meningococcal Reference Unit, Manchester Royal Infirmary Manchester, UK.

Hannah Christensen PhD

School of Population Health Sciences, Bristol Medical School, University of Bristol, UK.

Caroline Trotter PhD

Department of Veterinary Medicine, University of Cambridge, UK.

Adam Finn PhD

School of Population Health Sciences, Bristol Medical School, University of Bristol, UK.

Martin C J Maiden FMedSci

Corresponding Author

Department of Zoology, University of Oxford, 11a Mansfield Road, Oxford, OX1 3SZ, United Kingdom +44 1865 271284 Matthew D Snape MD

Oxford Vaccine Group, Department of Paediatrics, University of Oxford, and the National Institute for Health Research (NIHR) Oxford Biomedical Research Centre, Oxford, UK.

for the UKMenCar4 and 'Be on the TEAM' Study Collaborators

1 Abstract

Objectives Serogroup W and Y invasive meningococcal disease (IMD) increased globally from 2000 onwards. Responding to a rapid increase in serogroup W clonal complex 11 (W:cc11) IMD, the UK replaced an adolescent booster dose of meningococcal C conjugate vaccine with quadrivalent MenACWY conjugate vaccine in 2015. By 2018, vaccine coverage in the eligible school cohorts aged 14-19 years-old was 84%. We assessed the impact of the MenACWY vaccination programme on meningococcal carriage.

8 **Methods** An observational study of culture-defined oropharyngeal meningococcal carriage 9 prevalence before and after the start of the MenACWY vaccination programme in UK school 10 students, aged 15–19 years, using two cross-sectional studies: 2014–15 "UKMenCar4" and 11 2018 "Be on the TEAM" (ISRCTN75858406).

Results A total of 10625 participants pre-implementation and 13434 post-implementation were included. Carriage of genogroups C, W, and Y (combined) decreased from 2.03% to 0.71%(OR 0.34 [95% CI 0.27-0.44] p<0.001). Carriage of genogroup B meningococci did not change (1.26% vs 1.23% [95% CI 0.77-1.22] p=0.80) and genogroup C remained rare (n = 7/10625 vs 17/13488, p=0.135). The proportion of serogroup positive isolates, *i.e.*, those expressing capsule, decreased for genogroup W by 53.8% (95% CI -5.0%-79.8%, p=0.016) and for genogroup Y by 30.1% (95% CI 8.9%-46.3%, p=0.0025).

19 Conclusions The UK MenACWY vaccination programme reduced carriage acquisition of 20 genogroup and serogroup Y and W meningococci and sustained low levels of genogroup C 21 carriage. These data support the use of quadrivalent MenACWY conjugate vaccine for indirect 22 (herd) protection.

23 Introduction

During the first two decades of the 21st century, serogroup W invasive meningococcal disease 24 (IMD) increased globally. Previously confined to small local outbreaks, serogroup W expanded 25 from 2% to 40% of IMD in the UK between 2008 and 2015 (Figure 1).¹ Serogroup Y IMD also 26 increased, albeit with lower prevalence than serogroup W IMD and predominantly affecting 27 the elderly.² These changes led to some countries introducing the quadrivalent protein-28 conjugate (MenACWY) vaccine.^{1,2} The licenced MenACWY conjugate vaccines were 29 anticipated, but not robustly demonstrated, to induce herd protection. Large-scale carriage 30 31 studies accompanying the introduction of the monovalent Meningococcal C conjugate (MCC, 32 1999) and the serogroup A conjugate (MenAfriVac®, 2011) vaccines demonstrated strong herd protective effect by reducing carriage acquisition of the targeted serogroup.³⁻⁵ However, in 33 early studies MenACWY immunogenicity against serogroup C was reduced when compared 34 with the monovalent MCC.⁶ A recent systematic review concluded that there was a lack of 35 evidence supporting herd protective effects from multivalent MenACWY vaccines.⁷ 36

The absence of large-scale carriage studies represented an important evidence gap and this uncertainty affected the immunisation programmes introduced. For example, Chile introduced an infant-only programme providing direct protection to the age group at highest risk of IMD,⁸ whereas the United Kingdom targeted adolescents as the age-group with both a 'peak' in IMD incidence and the highest asymptomatic pharyngeal carriage rates. This strategy aimed to interrupt transmission and provide protection to unvaccinated age groups.⁹

43 The UK MenACWY programme was implemented in 2015 using MenACWY-TT
44 (Nimenrix®, Pfizer) and MenACWY-CRM₁₉₇ (Menveo®, GSK) vaccines (figure 1) to replace

the MCC school booster given at 13–14 years of age.¹⁰ In England, there was a phased roll-out 45 46 over three years, with a school based catch-up programme and a General Practitioner-based 47 programme up to 25 years of age. In Scotland, vaccination was delivered in schools within 48 one-year. In Wales, the catch-up programme was delivered in school and primary care over 49 two years. By mid-2018, MenACWY vaccine school-based coverage in England, Wales, and 50 Scotland, was 84%, ranging from 71-86% in each year cohort between 14 to 19 years of age.^{11,12} Vaccination coverage in the cohort aged 20 - 25 years was significantly lower; for 51 example, in England it was 35% to 39%. The MenACWY vaccine was not introduced for any 52 other age group. An infant-only '2 + 1' schedule of 4CMenB (Bexsero®, GSK) commenced 53 in September 2015,¹³ which was not anticipated to impact on meningococcal carriage.¹⁴ 54

The UK Meningococcal Carriage (UKMenCar1-4) surveys demonstrated the impact of herd protection from the MCC vaccine, and highlighted behaviours that increase the risk of meningococcal carriage and transmission.^{4,15,16} UKMenCar4 was conducted in 2014-2015 just prior to the introduction of the MenACWY vaccination programme. The subsequent 'Be on the TEAM (Teenagers Against Meningitis)' study (ISRCTN75858406) commenced in 2018 using compatible methods. We report the impact of the MenACWY campaign on meningococcal carriage in adolescents using these two studies.

62 Methods

63 Study Design and Sample Population

We compared two cross-sectional oropharyngeal carriage surveys of adolescents, taken before
(UKMenCar4: September 2014 – March 2015)¹⁵ and after ('Be on the TEAM', March –
November 2018)¹⁷ MenACWY vaccine introduction. Participants were aged 15–19 years

67 attending their penultimate year of school or college. Both studies were done by the same 68 research network with participants recruited in schools across multiple sites in England, Wales, 69 and Scotland. Characteristics of included schools reflected the diversity of educational settings 70 across the community.¹⁵ The 2018 cohort was eligible for MenACWY vaccination two to three years prior,¹⁸ while those sampled in 2014–15 were eligible for the adolescent MCC vaccine. 71 72 This study was approved by the NHS Research Ethics Committee (UKMenCar4 reference 14/SC/1163, Be on the TEAM reference 18/SC/0055). The study protocols have been 73 previously published^{17,19} and the 'Be on the TEAM' protocol is publicly available at 74 75 www.beontheteam.uk.

76 Sampling Collection & Laboratory Methods

Methods were compatible between the two surveys. After standard informed consent, 77 participants completed a meningococcal carriage risk factor questionnaire.^{15,17} Oropharyngeal 78 79 swab samples were taken using a standardised collection technique. Swab samples were either 80 directly plated (in some sites in 2014/15) or placed in STGG (skim milk, tryptone, glucose, glycerol) broth and frozen at -80°C within 4 hours of collection.¹⁵ After rethawing, swab 81 82 samples were incubated on GC-VCAT (vancomycin, colistin amphotericin B, trimethoprim; 83 ThermoFisher Scientific, Basingstoke, UK) at 37°C with 5% CO₂ for up to 48 hours. Putative 84 *Neisseria* isolates (oxidase-positive gram-negative colonies) were serogrouped by an in-house dot-blot ELISA method to identify serogroups B, C, W, and Y (UK Health Security Agency 85 Meningococcal Reference Unit).¹⁷ Following short-read whole genome sequencing (llumina 86 87 HiSeq or NovaSeq6000 platform, Wellcome Trust Oxford Genomics Centre), genomes were 88 assembled, characterised, and uploaded on the https://pubMLST.org/neisseria database, as described in the study protocols.^{15,17,19} 89

90

91 Statistical Methods

92 The survey sample sizes were pre-determined by a subset of the two cross-sectional studies: all Year 12 students (or S5 in Scotland) from the 2014–15 MenCar4 study (n=10625) and all 93 94 Year 12 (S5) students recruited in 2018 at the time of a planned interim analysis of the 'Be on 95 the TEAM' study (n=13341). In 2014–15, the carriage prevalence of genogroup W was 0.34%, 96 genogroup Y 1.60%, and genogroup C 0.07%. The sample size was estimated to provide a 97 detectable effect size of a 24% reduction in the carriage prevalence of genogroups C, W, and 98 Y combined at 80% power and alpha of 0.05. The primary analysis used logistic regression to 99 compare the difference in carriage prevalence between the pre- and post-implementation surveys, adjusting for study site. Several secondary analyses were done to assess the robustness 100 101 of our findings including: (i) clustering at the school level; (ii) restricting analysis to sites that 102 were included at both time points only; and (iii) multivariable logistic regression to adjust for 103 risk factors for any meningococcal carriage. Risk-factors included on the questionnaire were chosen based on significance from prior carriage studies.¹⁵ These included: age; gender; self-104 105 reported ethnicity; smoking and vaping status; household-smokers; current sore throat; current 106 or recent use of antibiotics; attendances at clubs, pubs or parties in the last week; number of 107 people kissed in the last week; relationship status and regular partner smoking status. Risk 108 factors with p values of < 0.20 in the univariable models were included in the multivariable 109 regression model and retained in the full model if the p < 0.05. Carriage prevalence was 110 reported as odds ratios with 95% confidence intervals (Clopper-Pearson). Carriage prevalence 111 reduction was derived from the odds ratio using the formula $((1 - OR) \times 100)$. Missing data 112 was not imputed and assumed to be missing at random. Statistical analyses were performed 113 using Stata version 17.

114 **Results**

115 In total 24062 oropharyngeal swab samples were included: 10624 from the pre-implementation 116 survey (2014–15) and 13438 from the post-implementation survey (2018). The cohorts were 117 demographically similar, with minor reductions in carriage risk factors consistent with previously reported trends (Table 1).¹⁵ In 2014-2015, 5.80% of participants were carriers of 118 any meningococcus compared with 4.49% in 2018 (Table 2). There was a significant decrease 119 120 in combined genogroup C, W, and Y meningococcal carriage from 2.03% to 0.71% (OR 0.34 121 [95% CI 0.27-0.44] p<0.001) (Table 2). Genogroup W carriage decreased from 0.34% to 122 0.09% (OR 0.27 [95% CI 0.14-0.51] p<0.001) and genogroup Y carriage decreased from 1.60% to 0.50% (OR 0.31 [95% CI 0.23-0.41] p<0.001). Genogroup C remained rare, with 123 124 no evidence of a difference in carriage prevalence between the time points, 0.07% (n=7/10625) 125 to 0.13% (n=17/13488) (OR 1.96 [95% CI 0.81-4.73] p=0.135). There was no evidence of a 126 change in the carriage prevalence of genogroup B (1.26% vs 1.24%, OR 0.97 [95% CI 0.77-127 1.22 p=0.81), other genogroups (OR 1.01 [95% CI 0.87–1.39] p=0.45), or capsule null (*cnl*) 128 meningococci (OR 0.90 [95% CI 0.72–1.12] *p*=0.36).

For the secondary analysis including meningococcal carriage risk factors, the final multivariable logistic regression model included smoking status, party/club/pub attendances, gender, recent antibiotic use, self-reported ethnicity, vaping, regular partner, partner smoking status. Whilst key carriage risk factors such as smoking decreased between 2014–15 and 2018, inclusion of carriage risk factors in the model did not substantially change the outcome measurements (supplemental figure 1). Outcome was also unaffected by adjustment of clustering in schools, or inclusion of only sites in both pre- and post-implementation surveys

(supplemental figure 1). Carriage showed between-site variation, as seen in previous studies
(supplemental table 1).^{15,16}

138 The odds ratio for serogroup W carriage in 2018 was 0.16 (95% CI 0.06–0.40, p<0.001) and 139 0.22 (95% CI 0.15-0.32, p < 0.001) for serogroup Y (Table 2). Capsule expression, defined 140 by serogroup positive isolates, decreased as a proportion of all genogroup W isolates from 141 72.2% to 33.3% (relative difference in proportions 53.8% [95% CI -5.0%-79.8%] p=0.016) 142 and for genogroup Y, from 69.4% to 48.5% (relative difference in proportions 30.1% [95% CI 143 8.9%-46.3%] p=0.0025) (figure 2). In 2014–15, one of seven genogroup C isolates expressed 144 capsule compared with none of the 17 genogroup C meningococci in 2018. There was no 145 significant change in the proportion of genogroup B that were expressing capsules (relative 146 difference in proportions 16.6% [95% CI -4.8%-33.5%] p=0.120).

The change in the carriage of the major ccs reflected the serogroup-specific vaccine effects (Figure 3): there were decreases in cc11 and cc22, associated with genogroups W, and cc-23, cc-167 and cc-174, associated with genogroup Y. However the proportion of clonal complexes associated with genogroups W, Y, and C remained similar (table 3). The carriage of W:cc-11 in 2014–15 was 0.22%, despite its high disease burden, in contrast to Y:cc23 (1.34%), which has a lower IMD incidence but was carried more commonly.

153 Discussion

Three years after the introduction of the UK MenACWY vaccination programme, crosssectional oropharyngeal carriage surveys showed: (i) a sustained low carriage of genogroup C meningococci; (ii) a 73% reduction in carriage of genogroup W meningococci; and (iii) a 69% reduction of genogroup Y meningococci. There was no change in genogroup B meningococcal

158 carriage. Reductions of carriage were observed at both genogroup level, *i.e.*, those 159 meningococci with cps regions encoding these capsules, and to a greater extent the serogroup level, *i.e.*, those meningococci expressing the capsule. These results were consistent with 160 161 studies of monovalent meningococcal conjugate polysaccharide vaccines, notably in the UKMenCar1-3 and MenAfriCar Studies. ^{3,4} Am impact of MenACWY on carriage was aso 162 163 seen in observational studies conducted in Polish soldiers, and an individually randomised controlled trial in UK university students.^{5,20,21} In this RCT, MenACWY-_{CRM} vaccinated 164 participants had relative genogroup B C, W and Y carriage reduction of 27.1% (95% CI 6.9 – 165 42.9%) at any timepoint between 2 - 12 months after vaccination compared with controls. In 166 this present population-level study, the magnitude of the reduction in carriage was greater, and 167 was commensurate with the impact of both the monovalent MCC and serogroup A vaccines.^{3,4} 168 A 2020 systematic review reported an absence of evidence supporting carriage effects of 169 MenACWY conjugate vaccines.⁷ However, this review included only the primary outcome at 170 1-month after vaccination for the UK student RCT, which did not differ between control and 171 vaccine groups.²² The other negative studies reviewed were in university populations in the 172 173 USA and the UK. In the USA MenACWY studies, meningococcal Y and W carriage rates were 174 so low that any inference about the impact of vaccination on carriage could not be considered definitive with respect to these serogroups.²³⁻²⁵ By contrast, in the UK student populations 175 analysed, carriage prevalence of genogroup W increased during the surveillance period after 176 MenACWY vaccination. This was likely due to an inadequate time interval between 177 178 vaccination and the commencement of the university year, where the maximum frequency of transmission occurs.^{5,26} An observational carriage study in Norwegian teenagers did not 179 demonstrate any impact of individual MenACWY vaccination status on serogroup-specific 180 carriage, although vaccination coverage in the cohort was less than 30%.²⁷ 181

The reduction in IMD incidence in the UK in both immunised and unimmunised cohorts 182 following the introduction of the MenACWY programme²⁸ was consistent with the 183 maintenance of direct and herd protection against serogroup C IMD, and the introduction of 184 185 direct and herd protection against serogroup W and serogroup Y IMD. The number of lives under five-years of age saved by herd protection against serogroup W in England and Wales 186 was estimated to be between 114 to 899 over four years.²⁸ Infant 4CMenB immunisation may 187 have provided some direct protection against serogroup W:cc11 IMD.²⁸ Serogroup W IMD, 188 which had been rising rapidly until 2015, plateaued and fell (figure 1). Whilst there is natural 189 190 fluctuations in meningococcal carriage, in our study the interval between pre and post-191 intervention was short and the carriage impact was shown to be specific to the dominant vaccine capsular targets genogroups Y and W and was demonstrated across multiple clonal 192 complexes associated with these genogroups. In contrast, genogroup B carriage was 193 194 unchanged.

195 The strengths of the present study included large scale, consistent methodology, and the use of age and school year-level matched cohorts. The culture-defined endpoint of meningococcal 196 197 carriage permitted high-resolution phenotypic and genotypic isolate characterisation, 198 strengthening the evidence that the trends in UK IMD are due to a capsule-specific vaccine 199 impact, rather than secular changes in meningococcal epidemiology. The existence of a 200 national immunisation program precluded a cluster-randomised approach and the study remains observational. The timing of sampling of each survey differed for pragmatic reasons, 201 202 with more sampling during winter months in the pre-implementation study. Meningococcal 203 carriage, unlike IMD, has not been shown to exhibit seasonal variation and even if present, it 204 would not disproportionally affect specific carriage of genogroup W and Y. The absence of

205 sampling from Northern Ireland was unlikely to have biased the results, given the similarity 206 between the populations in these countries. Whilst vaccine roll-out was more rapid in Scotland 207 than in England or Wales, Scotland was only in the pre-implementation survey, hence these 208 differences in roll-out did not impact on the post-implementation carriage results. Finally, while small changes in risk factors for meningococcal carriage reach statistical significance, 209 this is a factor of the study sample size, and the magnitude of differences is clinically small. 210 211 Inclusion of risks fact in the secondary analysis did not change the vaccine impact. It is unlikely that these differences between groups would effect carriage of genogroups W and Y 212 213 and not B.

214 We have demonstrated that the UK meningococcal ACWY conjugate vaccine programme 215 maintained low carriage of genogroup C meningococci, and reduced carriage of serogroup and genogroup W and Y meningococci. This herd protection effect reduced IMD in all age groups, 216 217 affirming the UK Joint Committee on Vaccines and Immunisation strategy of deploying 218 meningococcal conjugate vaccines in age groups with high meningococcal transmission.¹³ As 219 with monovalent meningococcal serogroup C and A conjugate vaccines, public health 220 interventions that leverage herd protection from MenACWY vaccines will result in greater 221 reductions in IMD across all age-cohorts, more rapid impact, and favourable cost-effectiveness. 222 By contrast, there remains no evidence of indirect protection from outer membrane protein 223 vaccines designed for group B meningococci, thus carriage studies are essential to complement 224 IMD surveillance and inform meningococcal vaccine policy.

225 Collaborators

Keith A Jolley, Karen Ford & Hannah Roberts (Oxford), Karen Palmer (Preston), Debbie
Suggitt (Stockport), Nicola Pemberton (Wigan), Samantha Ray (Cardiff) Mandy Wootton
(Cardiff), Shamez N. Ladhani (UKHSA), Daniel Owens & Katrina Cathie (Southampton),
Simon Royal (The University of Nottingham Health Service), Neil Oldfield (School of Life
Sciences, University of Nottingham), Roisin Ure, (Meningococcal Reference Lab, Glasgow),
Jennifer Richards (Public Health Wales Microbiology), Rebecca Ramsay (Brighton),
Samantha Thomson Hill & Kaltun Duale (Bristol)

233

234 Transparency Declaration

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289

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317

318 Contribution

319 'UKMenCar4' conceptual design by JMM, CT, AJP, RB, SNL, and MCJM. 'Be on the TEAM'

320 study conceptual design and management committee: MDS, AF, MCJM, CT, HC, RB, SG,

321 JMM, PH, SF, KD, SC, PA, EP, JPC. Study sites - investigators, local management, 322 recruitment, microbiology processing and data handling MDS, JC, SC, EP, AL, RB, AF, JO, 323 SNF, PH, DPJT, EW, MR, SH, AS, JPC, MW, SR, CC, ASm, DO, DB, CW, RG, CSZ, PTH, 324 AB, AV, MDS, RS, RC, KP, DS, NP, JMM, HBB, KF and OBH. Bioinformatics and genomic analysis by HBB, JMM, CMCR, OH, KAJ, JEB. Data reviewed by JC HC, CT and MCJM 325 326 with primary statistical analysis by JPC, JMM, HC, CT, MCJM, MDS. Preparation of original 327 draft manuscript, figures and literature search by JPC, with conceptual design, original input and initial revisions by MCJM, MDS, JMM, CT, HC. All listed authors have reviewed, revised 328 329 and approved the manuscript.

330

331 Access to Data

332 UKMenCar4: All genomes and metadata for the UKMenCar4 study are available open access
333 through PubMLST, with short-read sequence data available from the European Nucleotide
334 Archive (reference PRJEB14319) with European Nucleotide Archive accession run identifiers
335 accessible on Pub MLST. Corresponding data from the Be on the TEAM study will be added
336 after publication of the final results.

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419

Table 1 Participant demographics & risk factors for meningococcal carriage

	Pre-Implementation Survey	Post Implementation Survey	Difference between groups	
	2014-15	2018	% (95% CI) unless stated	
Sample Size	10624	13438		
Sampling Period	Sept 2014 – March 2015	2018 Mar-May; Sept-Nov		
Vaccination Schedule	MCC boost	MCC boost (aged 13-14yo)		
	(aged 13-14yo)	MenACWY (from Sept 2015)		
MenACWY Vaccination Coverage (England; age-year specific cohort)	n/a	80.95%		
Age in years % of cohort	15 16 17 18 19 3.4 52.6 40.3 3.26 0.45	15 16 17 18 19 0.01 65.3 31.9 2.6 0.01		
mean	16.45 years	16.38 yrs	- 0.07 years (-0.06 to - 0.09 yrs) $p < 0.001$	
Gender M:F:non-binary	41% : 59% : n/a	38% : 61% : 0.5 %	Binary gender M:F -2.5% (-1.4% to - 3.9%) p < 0.001	
Self-identified Ethnicity			<i>p</i> < 0.001	
White	80.3%	79.2%		
Asian / Asian British	10.0%	9.6%		
Black/African/Caribbean/Black British	4.7%	4.7%		
Mixed/multiple ethnic	3.2%	4.2%		
Other ethnic group	1.6%	1.9%		
Not reported	0.1%	0.4%		
Current sore throat	23.5%	19.5%	- 4.0% (-3.0% to -5.0%) <i>p</i> < 0.001	
Antibiotics	13.5 %	10.1 %	-3.4% (-2.6% to $-4.2%$) $p < 0.001$	
currently or in the last month				
Cigarette smoking (any amount) in last	9.2%	6.8%	-2.4% (-1.7% to -3.1%) <i>p</i> < 0.001	
week				
Vape / E-cigarette use (any amount)	3.5%	6.2%	2.7% (2.2% to 3.2%) <i>p</i> < 0.001	
Household smokers (inside or outside)	27.1%	20.0%	-7.1% (-6.0% to -8.2%) <i>p</i> < 0.001	
Any attendance at club/pub/parties in the last week	33.2%	29.4%	-3.8% (-2.6% to - 5.0%) <i>p</i> < 0.001	
Intimate kissing in last week	33.5%	28.5%	-5.0% (-3.8% to - 6.2%) $p < 0.001$	
Regular Partner / Girlfriend / Boyfriend	26.6%	24.9%	-1.7% (-0.6% to $-2.8%$) $p < 0.001$	

Survey completion rate: 2014-15 99.7%; 2018 99.6%

	Ν	Aeningococcal (Carriage				
	2014/1	5	20	18			
	(n=1062	24)	(n= 1	3434)			
	n	%	n	%	Odds Ratio*	[95% CIs]	р
any N. meningitidis	616	5.80%	603	4.49%	0.76 [0.68	- 0.86]	< 0.00
genogroup C, W, Y**	216	2.03%	96	0.71%	0.35 [0.27	- 0.44]	<0.00
serogroup C, W, Y	147	1.38%	37	0.28%	0.20 [0.14	- 0.29]	<0.00
genogroup C	7	0.07%	17	0.13%	1.96 [0.81	- 4.73]	0.13
serogroup C	1	0.01%	0	0.00% .			
genogroup W	36	0.34%	12	0.09%	0.27 [0.14	- 0.51]	<0.00
serogroup W	26	0.24%	5	0.04%	0.16 [0.06	- 0.40]	<0.00
genogroup Y	170	1.60%	68	0.50%	0.31 [0.23	- 0.41]	< 0.00
serogroup Y	118	1.11%	33	0.25%	0.22 [0.15	- 0.32]	<0.00
genogroup B	134	1.26%	165	1.23%	0.97 [0.77	- 1.22]	0.80
serogroup B	73	0.69%	75	0.56%	0.81 [0.59	- 1.12]	0.20
other genogroups***	122	1.15%	168	1.25%	1.01 [0.87	- 1.39]	0.44
capsule null (cnl)	152	1.43%	174	1.29%	0.90 [0.72	- 1.12]	0.35

Table 2 Meningococcal carriage prevalence and odds ratio of carriage after the introduction of the MenACWY adolescent vaccination programme in August 2015

*adjusted odds ratio with study site as a co-variable

** genogroup C, W, Y, includes 3 genogroup ambiguous W/Y isolates of which two were serogroup W, other single genogroup classifications do not included these isolates ***other includes genogroups E, H, K, L, X or Z or incomplete capsular group after manual capsular assignment

		2014/15			2018
Genogroup	Clonal Complex	n	% of genogroup	17	% of genogroup
C		n	genogroup	n	genogroup
C		2	50.00/	0	52.00/
	ST-269	3	50.0%	9	52.9%
	ST-41/44	0		4	23.5%
	ST-11	0		1	5.9%
	ST-1157	0		1	5.9%
	ST-162	1	16.7%	0	
	ST-231	1	16.7%	0	
	ST-35	0		1	5.9%
	not assigned	1	16.7%	1	5.9%
	Total	6		17	
W				0	
	ST-11	24	66.7%	7	58.3%
	ST-22	12	33.3%	3	25.0%
	ST-865	0		1	8.3%
	not assigned	0		1	8.3%
	Total	36		12	
Y					
	ST-23	143	84.1%	63	94.0%
	ST-167	14	8.2%	0	
	ST-174	4	2.4%	0	
	ST-1157	0		3	4.5%
	ST-103	1	0.6%	0	
	ST-41/44	0	0.0%	1	1.5%
	not defined	1	0.6%	0	
	not assigned	7	4.1%	1	1.5%
	Total	170		68	

Table 3 Carriage of clonal complexes listed by of genogroups C, Y, and W before and after implementation of the MenACWY immunisation programme

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Gender M:F:non-binary	41% : 59% : n/a	38% : 61% : 0.5 %	Binary gender M:F -2.5% (-1.4% to - 3.9%) p < 0.001	
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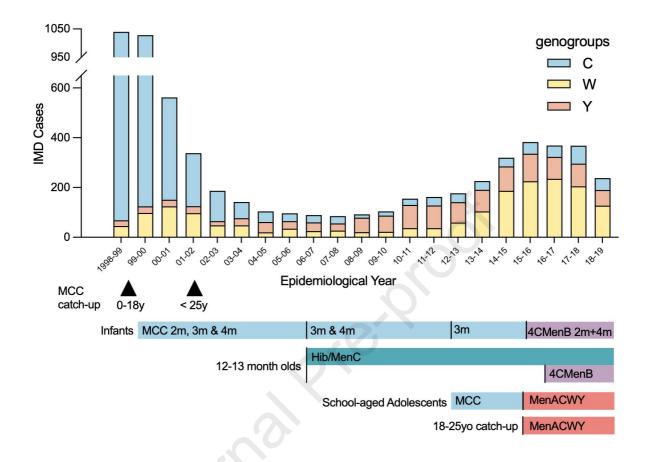
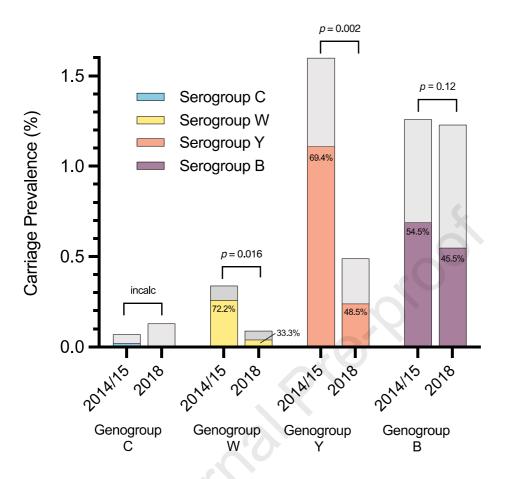
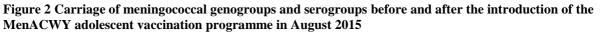


Figure 1 Meningococcal Vaccine Schedule & Invasive Meningococcal Disease due to genogroups C, W or Y in England & Wales 1998-2018 by epidemiological year

IMD - Invasive Meningococcal Disease (culture and/or PCR-confirmed, data collected by Public Health England Meningococcal Reference Unit); m - months; MCC - Meningococcal C conjugate vaccine.; MenACWY - quadrivalent meningococcal serogroup ACWY polysaccharide conjugate vaccine; Hib-MenC - Haemophilus influenzae type B & meningococcal C conjugate vaccine; 4CMenB - sub-capsular protein surrogate meningococcal B vaccine (Bexsero, GSK) introduced Sep 2015. MCC single-dose catch-up given in 1999 aged 0-18 years old and an expanded in 2002 to < 25 years old who were unvaccinated. Infant MCC ceased in July 2016. Adolescent vaccines given in schools at 13-14 years old. MenACWY commenced in schools in 2015, with a staggered catch-up over 1 year (Scotland) and 3 years (England, Wales) to vaccinate all those aged 14-18. A community / General Practitioner catch-up programme was implemented for school leavers up to 25 years old, but uptake was low.





Coloured shading – serogroup positive isolates. Grey shading serogroup negative isolates. Percentages in each column denote the proportion of genogroups that were serogroup positive with p-values shown for the difference in proportions of serogroup positive isolates at each time point. incalc – incalculable

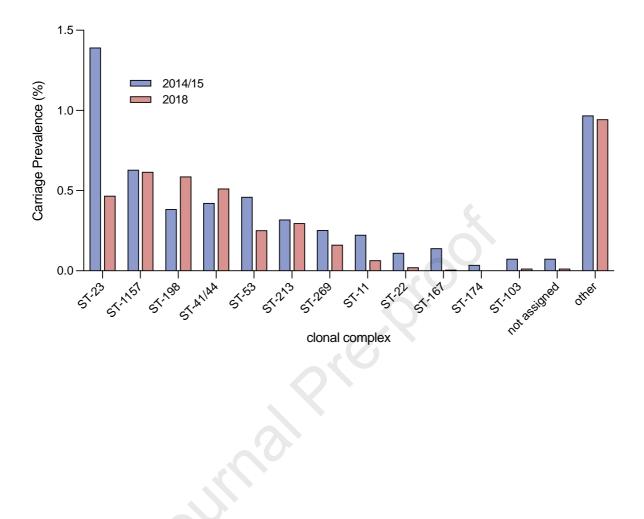


Figure 3 Carriage prevalence by clonal complex in adolescents aged 15 – 19yrs before and after the introduction of the adolescent MenACWY vaccination programme in August 2015

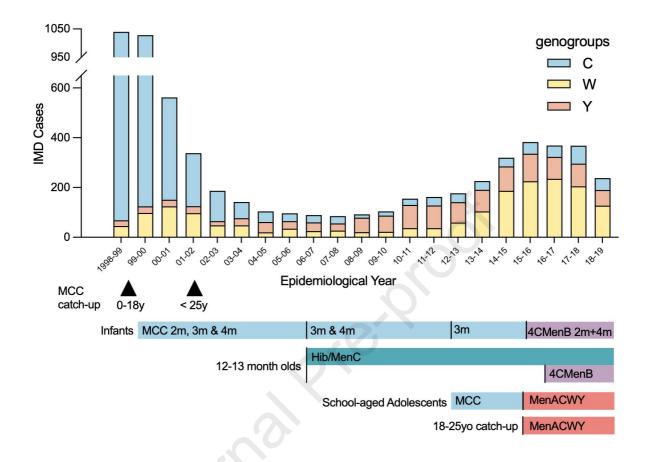
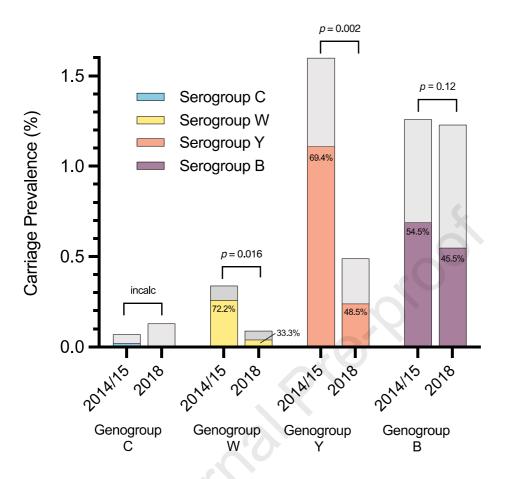
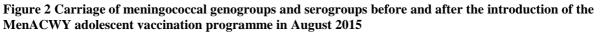


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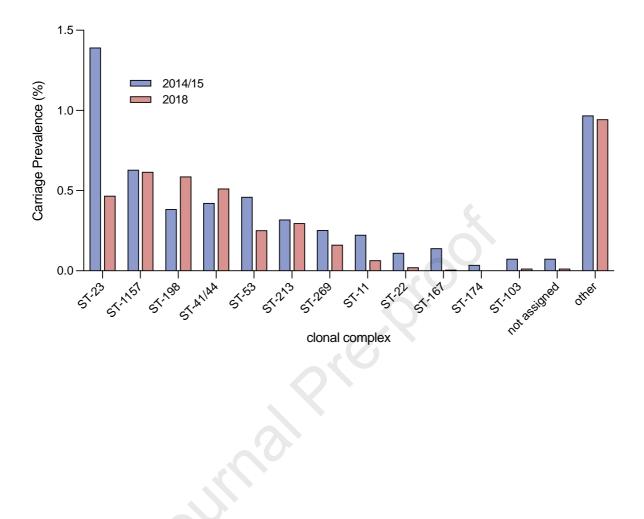


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