CMAJ

Research

Persistence of bactericidal antibodies following early infant vaccination with a serogroup B meningococcal vaccine and immunogenicity of a preschool booster dose

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Abstract

Background: The multicomponent serogroup B meningococcal (4CMenB) vaccine was recently licensed for use in Europe. There are currently no data on the persistence of bactericidal antibodies induced by use of this vaccine in infants. Our objective was to evaluate serogroup B-specific bactericidal antibodies in children aged 40–44 months previously vaccinated at 2, 4, 6 and 12 months of age.

Methods: Participants given 4 doses of 4CMenB as infants received a fifth dose of the vaccine at 40–44 months of age. Age-matched participants who were MenB vaccine–naive received 4CMenB and formed the control group. We evaluated human complement serum bactericidal activity (hSBA) titres at baseline and 1 month after each dose of 4CMenB.

Results: Before a booster dose at enrolment, 41%–76% of 17 participants previously vaccinated with 4CMenB in infancy had hSBA titres

vaccine against serogroup B meningococcus has recently been licensed for L use in Europe¹ and is being considered for licensure in Canada. This vaccine, known as multicomponent serogroup B meningococcal (4CMenB) vaccine, consists of 3 recombinant proteins: factor H binding protein (fHbp), Neisseria adhesin A (NadA) and Neisseria heparin binding antigen (NHBA) combined with detoxified outer membrane vesicles from the strain responsible for an epidemic of serogroup B meningococcal disease in New Zealand (NZ98/254). Clinical trials of 4CMenB have shown it to be immunogenic against reference strains selected to specifically express one of the vaccine antigens.²⁻⁶ On the basis of these trials, the approved schedule for infants aged 2 to 5 months is 3 doses given at least 1 month apart, with a booster dose given at 12 to 23 months of age.7 The persistence of vaccineof 4 or greater against 4 reference strains. Before vaccination in the control group (n = 40) these proportions were similar for strains 44/76-SL (63%) and M10713 (68%) but low for strains NZ98/254 (0%) and 5/99 (3%). A booster dose in the 4CMenB-primed participants generated greater increases in hSBA titres than in controls.

Interpretation: As has been observed with other meningococcal vaccines, bactericidal antibodies waned after vaccination with 4CMenB administered according to an approved infant vaccination schedule of 2, 4, 6 and 12 months of age, but there was an anamnestic response to a booster dose at 40–44 months of age. If 4CMenB were introduced into routine vaccination schedules, assessment of the need for a booster dose would require data on the impact of these declining titres on vaccine effectiveness. ClinicalTrials.gov, no. NCT01027351

Competing interests: See end of article for competing interests.

This article has been peer reviewed.

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CMAJ 2013. DOI:10.1503 /cmaj.130257

induced antibodies throughout childhood following this booster dose is unknown, but it is particularly relevant because the incidence of invasive serogroup B meningococcal disease in children aged 1 to 4 years is second only to the incidence in children less than 1 year of age.⁸

In this study, we assessed the persistence of these bactericidal antibodies in children aged 40– 44 months who had previously received either 4CMenB or a vaccine containing the recombinant proteins alone (recombinant protein serogroup B meningococcal [rMenB] vaccine) at 2, 4, 6 and 12 months of age.³ We also assessed the immunogenicity and reactogenicity of a booster dose.

Methods

Participants

Participants in the original phase II study were recruited at 2 sites: 12 at the Gloucester Vaccine

Evaluation Unit and 135 at the Oxford Vaccine Group, University of Oxford. In the original study, infants were randomly assigned 2:2:1:1 to receive 4CMenB at 2, 4, 6 and 12 months; rMenB at 2, 4, 6 and 12 months; 4CMenB at 12 months; or rMenB at 12 months (Figure 1). All 125 participants who completed the original study at the Oxford site were invited to take part in this follow-up study, and we planned to recruit 50 MenB vaccine–naive, age-matched participants as controls.

Inclusion criteria were healthy children aged 40–44 months who had completed the original study, or, for controls, children of this age who had not previously received a MenB vaccine. Exclusion criteria were previous meningococcal disease (or household or intimate contact with anyone with meningococcal disease), allergy to vaccine components, severe acute or chronic disease, immune dysfunction, receipt of blood products, planned receipt of nonstudy vaccines within 30 days of the study vaccines, enrolment in another clinical trial, recent antibiotic use, being a family member of research staff or antipyretic use within 6 hours before enrolment.

Written informed consent was obtained from participant's parents or legal guardians. Ethics approval was obtained from Oxfordshire Research Ethics Committee B (reference no. 09/H0605/89). The study was conducted from January to December 2010.

Procedures

Participants who had previously received 4 doses of 4CMenB or rMenB in the original study had blood samples taken before and 30 days after a booster dose of the respective vaccines (Figure 1). Participants previously given a single dose of 4CMenB or rMenB in the original study had 2 doses of the respective vaccine administered about 60 days apart. Control participants received 4CMenB at 40 and 42 months of age, thereby receiving 2 doses of 4CMenB at least 2 months apart, the approved dosing schedule for this age group. All participants had blood samples taken at enrolment and 30 days after each vaccination.

Both vaccines were manufactured and supplied by Novartis Vaccines and Diagnostics. Both contained 50 µg each of NadA allele 3, NHBA-GNA (genome-derived Neisseria antigen) 1030 fusion protein (containing NHBA peptide variant 2) and GNA2091-fHbp fusion protein (containing fHbp variant 1.1) as well as 1.5 mg aluminum hydroxide and 10 mM histidine. The 4CMenB vaccine also contained 25 µg of detoxified outer membrane vesicles from Neisseria meningitidis strain NZ98/254 (expressing the immunodominant antigen PorA [porin A] serosubtype P1.4). The vaccines were 0.5 mL in volume and were administered by intramuscular injection into the deltoid area of the nondominant arm. Participants also received a combination diphtheria-toxoid, tetanus-toxoid,

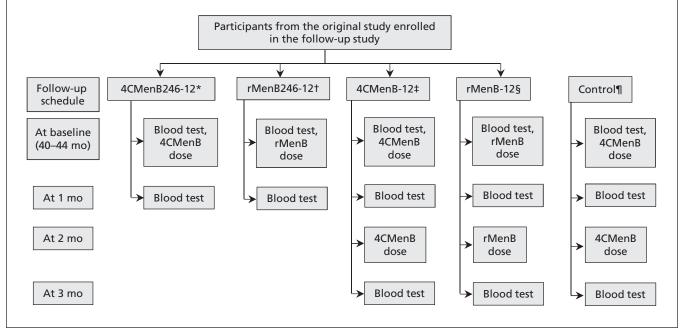


Figure 1: Study design. *Participants who received 4 doses of multicomponent serogroup B meningococcal (4CMenB) vaccine at 2, 4, 6 and 12 months. †Participants who received 4 doses of recombinant protein serogroup B meningococcal (rMenB) vaccine at 2, 4, 6 and 12 months. ‡Participants who received 1 dose of 4CMenB vaccine at 12 months. §Participants who received 1 dose of rMenB vaccine at 12 months. ¶Participants with no previous exposure to 4CMenB or rMenB vaccines.

acellular pertussis and inactivated polio vaccine (Repevax, Sanofi Pasteur MSD) and a combination measles, mumps and rubella vaccine (MMRvaxPro, Sanofi Pasteur MSD) at the final study visit. These vaccines were administered to keep participants vaccinated according to the routine schedule in the United Kingdom and did not form part of the study evaluation.

Immunogenicity evaluation

Sera were analyzed at the Vaccine Evaluation Unit of the Health Protection Agency in Manchester, UK, for serum bactericidal activity using a human complement source as previously described.9 Laboratory staff were unaware of participants' group allocation. The correlate of protection was a human complement serum bactericidal activity (hSBA) titre of 1:4 or greater. As previously described,³ reference hSBA strains were used to assess the immunogenicity of specific vaccine components: fHbp was assessed by strain 44/76-SL, NadA by strain 5/99 and PorA (the immunodominant antigen in the outer membrane vesicle) by NZ98/254 (Table 1). We used a novel strain (M10713) to assess the immunogenicity of NHBA because this expresses NHBA cross-reactive to that contained in the vaccine (peptide 10) but is mismatched for the vaccine components fHbp (subvariant 2.24) and PorA (16-3) and does not contain the gene for NadA. Therefore, in accordance with the principle underlying the use of the existing reference strains, any increase in hSBA titres against M10713 following vaccination may be attributed to the NHBA component of the vaccine. We assessed immunogenicity against additional strains (M00-242922, M01-240101, M01-240364) to evaluate the impact of antigenic variation and expression on susceptibility to vaccineinduced antibodies (Table 1). We specifically selected strain M01-240355 as a strain likely to be relatively resistant to vaccine-induced bactericidal antibodies because it was mismatched for all vaccine antigens.

Reactogenicity evaluation

Each day in the week after vaccination, parents recorded their child's axillary temperature, local reactions at the site of vaccination (pain, ery-thema, swelling and induration), solicited systemic reactions (fever, irritability, vomiting, diarrhea, sleepiness, anorexia, arthralgia, headache and rash) and the use of analgesic and antipyretic medication. Reactions were classified as severe if tenderness prevented daily activity, local ery-thema and induration were greater than 50 mm in diameter or the child's fever after vaccination was 40°C or higher.

Outcome measures

The primary outcome measure for immunogenicity was the percentage of children with an hSBA titre of 1:4 or greater among those who received 4 doses of either vaccine. The primary safety objective was to assess the safety and tolerability of 1 or 2 booster doses of these vaccines.

Secondary objectives were assessment of the persistence of bactericidal antibodies following vaccination with 1 dose of 4CMenB or rMenB at 12 months of age, increases in hSBA titres following 1 or 2 booster doses and adverse reactions following a 2-dose "catch-up" vaccination schedule of 4CMenB administered at enrolment.

Statistical analysis

We calculated the percentages of children in each study group with hSBA titres of 1:4 or greater, along with 2-sided 95% Clopper– Pearson confidence intervals (CIs).¹¹ Similarly, we calculated geometric mean titres of bactericidal antibodies and determined 95% CIs by exponentiating (base 10) the corresponding means and 95% CI of the log10 hSBA titre. Geometric mean ratios of hSBA geometric mean titres after vaccination to baseline and 95% CIs were also calculated. For these calculations, we allocated a value of 1 to hSBA titres below the lower limit of detection of 2.

The primary population for immunogenicity analysis was the intention-to-treat group, consisting of all participants who were randomly assigned, received a dose of either vaccine and provided at least 1 evaluable serum sample. Safety analysis was conducted on a population consisting of all participants who received a dose of either vaccine and provided safety data after baseline.

The sample size of the follow-up participants was determined by the number of participants who completed the original study at the Oxford site. All comparisons were descriptive. However, we calculated that if the percentage of participants with hSBA titres of 1:4 or greater was 40% in a follow-up group and 5% in the control group, a sample size of 30 participants in the follow-up group and 50 participants in the control group would be required for 95% power to show superiority.

Results

Of the 125 participants who completed the original study at the Oxford site, 70 were enrolled in the follow-up study, of whom 68 were included in the intention-to-treat analysis. In addition, 43 MenB vaccine–naive participants were enrolled, of whom 40 were included in the intention-to-

Table 1: Strair	is used to	assess hur	man comp	plement ser	um bactericidal	activity of a sero	group B me	iningococcal vac	Table 1: Strains used to assess human complement serum bactericidal activity of a serogroup B meningococcal vaccine and description of vaccine target antigens	on of vaccine	et antigen	S
					fHbp			NadA			NHBA	
Strain	ST	С	PorA (VR2)	Novartis variant*	Expression†	GenBank accession no.‡	Variant	Expression†	GenBank accession no.‡	Peptide	Expression†	GenBank accession no.‡
44/76-SL	32	32	16	1.1§	‡	AY548370	×	×	×	m	+	AF226436
5/99	1349	∞	2	2.2	I	GQ302858	2§	++++	GQ302859	20	-/+	GQ302856
NZ98/254	42	41/44	4§	1.1	+	AY548375		×	×	2	+	GQ302855
M10713	136	41/44	16-3	2.2	+	NA	×	×	×	10§	+	NA
M00-242922	41	41/44	4§	1.4	+	FJ750977	×	×	×	2	+	FJ750982
M01-240101	1049	269	15	1.2	+	FJ750976	×	×	×	21	-/+	FJ750981
M01-240355	213	213	14	3.3	-/+	EU541888	5	-/+	FJ750979	18	-/+	GQ302857
M01-240364	11	11	2	3.3	-/+	FJ750978	2§	+	FJ619643	28	-/+	FJ750980
Note: CC = clonal complex, fHbp = factor H binding protein, NadA = <i>Neisseria</i> adhes VR2 = variable region 2, x = does not harbour the NadA gene. *The fHbp variant nomenclature displayed is the one used by Novartis obtained by. #Expression of antigen as determined by binding of murine antibodies raised again #GenBank accession number identifying unique sequence at www.ncbi.nlm.nih.gov §These vaccine targets are homologous (PorA, fHbp, NHBA) or cross-reactive (NadA)	complex, fH gion 2, x = dı : nomenclatu tigen as detu on number iu gets are hor	bp = factor oes not harb ure displayed smined by t dentifying u nologous (P	H binding p bour the Nar d is the one binding of r indue seque orA, fHbp, I	rrotein, NadA = dA gene. used by Novar nurine antiboc ence at www.r NHBA) or cross	 Neisseria adhesin rtis obtained by ad fies raised against i ncbi.nlm.nih.gov. reactive (NadA) tt 	Note: CC = clonal complex, fHbp = factor H binding protein, NadA = <i>Neisseria</i> adhesin A, NA = GenBank accession number not assigned, NHBA = <i>Neisseria</i> H VR2 = variable region 2, x = does not harbour the NadA gene. *The fHbp variant nomenclature displayed is the one used by Novartis obtained by adding a prefix defining the main variant (1, 2 or 3) to the subvariant. ¹⁰ #Expression of antigen as determined by binding of murine antibodies raised against each of the vaccine antigen(s): – low expression, +/– medium expressic #GenBank accession number identifying unique sequence at www.ncbi.nlm.nih.gov.	:ession numbe j the main var ntigen(s): – lov łCMenB.	rr not assigned, NHE iant (1, 2 or 3) to th v expression, +/- me	Note: CC = clonal complex, fHbp = factor H binding protein, NadA = Neisseria adhesin A, NA = GenBank accession number not assigned, NHBA = Neisseria heparin binding antigen, PorA = porin A, ST = sequence type, VR2 = variable region 2, x = does not harbour the NadA gene. *The fHbp variant nomenclature displayed is the one used by Novartis obtained by adding a prefix defining the main variant (1, 2 or 3) to the subvariant. ¹⁰ *Expression of antigen as determined by binding of murine antibodies raised against each of the varcine antigen(s): – low expression, +/– medium expression, ++ very high expression. #GenBank accession number identifying unique sequence at www.ncbi.nlm.nlh.gov.	binding antige th expression, +	1, PorA = porin A, S .+ very high expres:	T = sequence type, ion.

treat analysis (Table 2). The demographic characteristics of the participants were similar across all groups. The median age at enrolment was 41 (range 40–44) months, 50% were male (compared with 57% in the original study) and 92% of participants were white (compared with 94% in the original study).

Immunogenicity

Waning of antibodies was observed at baseline following vaccination in early infancy with either vaccine. At least 65% of participants who received 4CMenB at 2, 4, 6 and 12 months had hSBA titres of 1:4 or greater at baseline for strains 44/76-SL (evaluating fHbp), 5/99 (NadA) and M10713 (NHBA) compared with 41% for NZ98/254 (PorA) (Table 3; Appendix 1, available at www.cmaj.ca/lookup/suppl/doi:10.1503 /cmaj.130257/-/DC1). Among participants who received rMenB at 2, 4, 6 and 12 months of age, at least 43% showed hSBA titres of 1:4 or greater for these strains, except for NZ98/254 (PorA) (3%). As expected, the percentage of participants with hSBA titres of 1:4 or greater for strain M01-240355 was low, regardless of the vaccine received.

At baseline, 25% and 38% of children who received 4CMenB at 12 months had hSBA titres of 1:4 or greater for strains M10713 (NHBA) and 44/76-SL (fHbp), respectively, whereas for strains 5/99 (NadA) and NZ98/254 (PorA), these proportions were 0% (Table 3, Appendix 1).

Administration of a booster dose of 4CMenB at enrolment resulted in 86%-100% of 4CMenB-primed participants achieving hSBA titres of 1:4 or greater for all strains except M01-240355.

In MenB vaccine-naive children, administration of 2 doses of 4CMenB 2 months apart resulted in 89%-100% achieving hSBA titres of 1:4 or greater for all strains except M01-240355 (69%). More than half of these children had hSBA titres greater than 1:4 to strains 44/76-SL (fHbp), M10713 (NHBA), M01-240101 and M01-240364 at baseline, with hSBA geometric mean titres similar to those of participants who received 4CMenB at 2, 4, 6 and 12 months. The geometric rise in hSBA titres for all these strains following a 4CMenB booster dose was higher in the group that received 4CMenB at 2, 4, 6 and 12 months than in controls, as determined by nonoverlapping CIs (Table 4). The hSBA geometric mean titres for all time points in this and the original study are displayed in Appendix 2 (available at www.cmaj.ca/lookup/suppl/doi:10 .1503/cmaj.130257/-/DC1).

Reactogenicity

Most recipients of either vaccine had local pain and erythema at the injection site, but few reactions were severe (Appendices 3 and 4, available at www.cmaj.ca/lookup/suppl/doi:10.1503/cmaj .130257/-/DC1). Few participants reported fever.

One serious adverse event was experienced by a participant in the control group in whom otitis media developed 12 days after vaccination with 4CMenB, followed by cervical lymphadenitis. He required hospital admission for intravenous antibiotics. This was considered unrelated to the vaccination.

	Origin	al study			Foll	ow-up study		
				At baseline		At 1 mo	At 2 mo	At 3 mo
Group		Completed (Oxford)*	Enrolled	Included in ITT analysis for persistence†	Received first dose	Included in ITT analysis for immunogenicity†	Received second dose	Included in ITT analysis for immunogenicity†
4CMenB246-12‡	50 (46)	45 (41)	19	17	19	19	NA	NA
rMenB246-12§	48 (44)	44 (40)	29	29	29	28	NA	NA
4CMenB-12¶	24 (22)	23 (21)	8	8	8	7	8	8
rMenB-12**	25 (23)	24 (23)	14	14	14	13	14	13
Control++	NA	NA	43	40	42	38	41	36

4CMenB = multicomponent serogroup B meningococcal vaccine, ITT = intention-to-treat, NA = not applicable, rMenB = recombinant protein serogroup B meningococcal vaccine.

*Because the follow-up study was a single-centre study, only participants recruited from Oxford (in parentheses) in the original study were invited to participate in the follow-up study.

tExclusions from the ITT analysis are due to lack of evaluable serum sample, except for 1 participant in the control group who withdrew consent.

Participants who received 4 doses of 4CMenB vaccine at 2, 4, 6 and 12 months.

§Participants who received 4 doses of rMenB vaccine at 2, 4, 6 and 12 months.

¶Participants who received 1 dose of 4CMenB at 12 months.
**Participants who received 1 dose of rMenB vaccine at 12 months.

††Participants with no previous exposure to 4CMenB or rMenB vaccines.

Strain		At baseline	e	At 1 mo		At 3 mo	
(antigen assessed)	Group	% (95% CI)	n/N	% (95% CI)	n/N	% (95% CI)	n/N
44/76-SL	4CMenB246-12*	65 (38–86)	11/17	100 (82–100)	19/19	NA	NA
(fHbp)	rMenB246-12†	45 (26–64)	13/29	100 (88–100)	28/28	NA	NA
	4CMenB-12‡	38 (9–76)	3/8	100 (59–100)	7/7	100 (63–100)	8/8
	rMenB-12§	57 (29–82)	8/14	100 (75–100)	13/13	100 (75–100)	13/13
	Control¶	63 (46–77)	25/40	89 (75–97)	34/38	100 (90–100)	36/36
5/99	4CMenB246-12*	76 (50–93)	13/17	100 (81–100)	18/18	NA	NA
(NadA)	rMenB246-12†	43 (24–63)	12/28	100 (88–100)	28/28	NA	NA
	4CMenB-12‡	0 (0–37)	0/8	100 (59–100)	7/7	100 (63–100)	8/8
	rMenB-12§	57 (29–82)	8/14	100 (75–100)	13/13	100 (75–100)	13/13
	Control¶	3 (0–13)	1/40	76 (60–89)	29/38	100 (90–100)	36/36
NZ98/254	4CMenB246-12*	41 (18–67)	7/17	89 (67–99)	17/19	NA	NA
(PorA)	rMenB246-12†	3 (0–18)	1/29	14 (4–33)	4/28	NA	NA
	4CMenB-12‡	0 (0–37)	0/8	100 (59–100)	7/7	100 (63–100)	8/8
	rMenB-12§	7 (0–34)	1/14	15 (2–45)	2/13	15 (2–45)	2/13
	Control¶	0 (0–9)	1/40	66 (49–80)	25/38	94 (81–99)	34/36
M10713 (NHBA) M00-	4CMenB246-12*	67 (38–88)	10/15	94 (73–100)	17/18	NA	NA
	rMenB246-12†	68 (48–84)	19/28	96 (82–100)	27/28	NA	NA
	4CMenB-12‡	25 (3–65)	2/8	86 (42–100)	6/7	100 (63–100)	8/8
	rMenB-12§	54 (25–81)	7/13	100 (75–100)	13/13	83 (52–98)	10/12
	Control¶	68 (51–81)	27/40	76 (60–89)	29/38	89 (74–97)	32/36
M00- 242922	4CMenB246-12*	53 (28–77)	9/17	100 (81–100)	18/18	NA	NA
	rMenB246-12†	18 (6–37)	5/28	75 (55–89)	21/28	NA	NA
	4CMenB-12‡	0 (0–37)	0/8	100 (59–100)	7/7	100 (63–100)	8/8
	rMenB-12§	8 (0–36)	1/13	54 (25–81)	7/13	42 (15–72)	5/12
	Control¶	18 (8–34)	7/39	49 (32–66)	18/37	94 (81–99)	33/35
M01-	4CMenB246-12*	71 (44–90)	12/17	100 (81–100)	18/18	NA	NA
240101	rMenB246-12†	71 (51–87)	20/28	100 (88–100)	28/28	NA	NA
	4CMenB-12‡	63 (24–91)	5/8	100 (59–100)	7/7	100 (63–100)	8/8
	rMenB-12§	77 (46–95)	10/13	100 (75–100)	13/13	100 (74–100)	12/12
	Control¶	74 (58–87)	29/39	97 (86–100)	36/37	100 (90–100)	35/35
M01-	4CMenB246-12*	6 (0–29)	1/17	50 (26–74)	9/18	NA	NA
240355	rMenB246-12†	25 (11–45)	7/28	29 (13–49)	8/28	NA	NA
	4CMenB-12‡	25 (3–65)	2/8	71 (29–96)	5/7	63 (24–91)	5/8
	rMenB-12§	15 (2–45)	2/13	31 (9–61)	4/13	8 (0–38)	1/12
	Control¶	18 (8–34)	7/39	30 (16–47)	11/37	69 (51–83)	24/35
M01-	4CMenB246-12*	88 (64–99)	15/17	94 (73–100)	17/18	NA	NA
240364	rMenB246-12†	61 (41–78)	17/28	82 (63–94)	23/28	NA	NA
	4CMenB-12‡	50 (16–84)	4/8	100 (59–100)	7/7	100 (63–100)	8/8
	rMenB-12§	77 (46–95)	10/13	100 (75–100)	13/13	100 (74–100)	12/12
	Control¶	82 (66–92)	31/38	86 (71–95)	32/37	100 (90–100)	34/34

CI = confidence interval, fHbp = factor H binding protein, NA = not applicable, NadA = Neisseria adhesin A, NHBA = Neisseria heparin binding antigen, PorA = porin A. *Participants who received 4 doses of 4CMenB vaccine at 2, 4, 6 and 12 months.

†Participants who received 4 doses of rMenB vaccine at 2, 4, 6 and 12 months. ‡Participants who received 1 dose of 4CMenB at 12 months.

§Participants who received 1 dose of rMenB vaccine at 12 months. ¶Participants with no previous exposure to 4CMenB or rMenB vaccines.

Table 4: Geometric mean titres and geometric mean ratios of human complement serum bactericidal activity, intention-to-treat population

Strain (antigon		Geon	netric mean titres (9	95% CI)	Geometric mea	n ratios (95% CI)
Strain (antigen assessed)	Group	At baseline	At 1 mo	At 3 mo	1 mo:baseline	3 mo:baseline
44/76-SL (fHbp)	4CMenB 246-12*	5.3 (3.3–8.8)	89 (68–116)	NA	17 (11–27)	NA
	rMenB246-12†	3.2 (2.4–4.5)	99 (62–158)	NA	30 (18–50)	NA
	4CMenB-12‡	3.5 (0.9–14)	76 (27–210)	145 (74–283)	21 (4.2–110)	42 (12–147)
	rMenB-12§	3.6 (2.1–6.2)	94 (46–192)	127 (81–200)	27 (13–57)	37 (21–64)
	Control¶	4.2 (3.2–5.6)	12 (8.0–19)	88 (66–117)	3.2 (2.1–5.0)	20 (15–28)
5/99 (NadA)	4CMenB246-12*	28 (9.4–83)	1708 (774–3771)	NA	70 (29–167)	NA
	rMenB246-12†	5.1 (2.3–11)	778 (461–1311)	NA	148 (90–245)	NA
	4CMenB-12‡	1.0 (1.0–1.0)	509 (178–1456)	2413 (988–5892)	509 (178–1456)	2413 (988–5892)
	rMenB-12§	9.6 (4.2–22)	2379 (995–5685)	5240 (3216–8538)	299 (129–694)	501 (143–1748)
	Control¶	1.1 (0.7–1.8)	22 (13–36)	1019 (760–1367)	19 (10–36)	910 (594–1394)
NZ98/254 (PorA)	4CMenB246-12*	2.8 (1.4–5.6)	47 (20–107)	NA	20 (7.7–51)	NA
	rMenB246-12†	1.1 (1.0–1.2)	1.6 (1.0–2.6)	NA	1.5 (1.0–2.4)	NA
	4CMenB-12‡	1.0 (1.0–1.0)	148 (66–330)	65 (22–189)	148 (66–330)	65 (22–189)
	rMenB-12§	1.2 (0.9–1.7)	1.7 (0.8–3.8)	1.9 (0.9–3.7)	1.4 (0.6–3.1)	1.5 (0.7–3.1)
	Control¶	1.0 (1.0–1.0)	7.7 (4.6–13)	47 (31–72)	7.6 (4.5–13)	47 (31–74)
M10713 (NHBA)	4CMenB246-12*	5.3 (2.3–12)	39 (22–69)	NA	10 (4.3–24)	NA
	rMenB246-12†	9.2 (4.9–17)	38 (24–59)	NA	4.7 (2.3–9.4)	NA
	4CMenB-12‡	3.0 (0.8–12)	30 (9.5–96)	36 (21–61)	10 (2.0–49)	12 (3.0–48)
	rMenB-12§	3.3 (1.6–6.6)	35 (19–67)	21 (7.4–60)	11 (4.2–30)	5.5 (1.5–20)
	Control¶	8.8 (5.2–15)	11 (6.7–19)	33 (22–51)	1.3 (0.9–1.8)	3.9 (2.1–7.2)
M00-242922	4CMenB246-12*	3.5 (1.7–7.0)	64 (27–155)	NA	22 (8.3–56)	NA
	rMenB246-12†	1.6 (1.2–2.2)	7.0 (4.4–11)	NA	4.6 (2.9–7.5)	NA
	4CMenB-12‡	1.0 (1.0–1.0)	206 (33–1308)	53 (21–131)	206 (33–1308)	53 (21–131)
	rMenB-12§	1.2 (0.9–1.7)	4.6 (1.6–13)	3.0 (1.6–5.7)	3.8 (1.5–9.6)	2.4 (1.2–5.0)
	Control¶	1.4 (1.2–1.8)	3.6 (2.6–5.2)	31 (22–44)	2.7 (2.0–3.8)	23 (16–34)
M01-240101	4CMenB246-12*	7.6 (4.3–13)	159 (80–314)	NA	20 (6.8–59)	NA
	rMenB246-12†	7.0 (4.0–12)	76 (40–143)	NA	11 (5.7–23)	NA
	4CMenB-12‡	4.0 (1.7–9.2)	59 (14–248)	92 (40–213)	16 (2.4–105)	23 (5.4–100)
	rMenB-12§	12 (3.4–41)	82 (35–190)	43 (21–89)	6.9 (2.8–17)	4.2 (1.1–16)
	Control¶	6.6 (4.4–9.9)	87 (46–165)	207 (120–358)	12 (6.5–23)	33 (17–64)
M01-240355	4CMenB246-12*	1.2 (0.8–1.7)	5.0 (2.6–9.3)	NA	4.2 (2.2–8.2)	NA
	rMenB246-12†	1.8 (1.2–2.8)	1.9 (1.3–2.8)	NA	1.0 (0.7–1.6)	NA
	4CMenB-12‡	1.7 (0.9–3.0)	5.9 (3.1–11)	5.5 (1.8–17)	3.3 (1.3–8.5)	3.3 (1.4–7.7)
	rMenB-12§	1.4 (0.8–2.4)	1.7 (1.0–2.9)	1.3 (0.8–2.0)	1.2 (0.8–1.7)	0.9 (0.6–1.3)
	Control¶	1.5 (1.2–2.0)	2.5 (1.5–4.1)	6.2 (3.6–11)	1.7 (1.2–2.5)	4.1 (2.7–6.3)
M01-240364	4CMenB246-12*	17 (8.3–36)	134 (56–322)	NA	8.3 (3.0–23)	NA
	rMenB246-12†	6.5 (3.6–12)	41 (17–96)	NA	6.2 (2.6–15)	NA
	4CMenB-12‡	5.0 (1.6–16)	52 (14–193)	107 (26–443)	8.2 (1.6–43)	21 (5.0–91)
	rMenB-12§	12 (4.9–31)	275 (144–528)	616 (191–1980)	22 (11–44)	56 (18–172)
	Control¶	9.4 (6.5–14)	11 (7.6–16)	104 (68–159)	1.2 (0.8–1.7)	11 (6.4–18)

CI = confidence interval, fHbp = factor H binding protein, hSBA = human complement serum bactericidal activity, NA = not applicable, NadA = Neisseria adhesin A, NHBA = *Neisseria* heparin binding antigen, PorA = porin A. *Participants who received 4 doses of 4CMenB vaccine at 2, 4, 6 and 12 months.

Participants who received 4 doses of 4CMenB vaccine at 2, 4, 6 and 12 months. ‡Participants who received 1 dose of 4CMenB at 12 months. §Participants who received 1 dose of rMenB vaccine at 12 months. ¶Participants with no previous exposure to 4CMenB or rMenB vaccines.

Interpretation

The 4CMenB vaccine was licensed for use in Europe in January 2013, with approved dosage schedules including the 2, 4, 6, 12 month and 40, 42 month schedules evaluated in this study. In contrast to the single dose of 4CMenB given at 12 months of age to one of our study groups, the licensed indication for children 1–2 years of age is 2 doses of 4CMenB at least 2 months apart, with a booster dose given 1–2 years later.

Among children who received 4CMenB at 2, 4, 6, and 12 months of age in the original study, by 40 months of age the proportions with hSBA titres of 1:4 or greater were 65% for strain 44/76-SL (fHbp), 76% for strain 5/99 (NadA) and 41% for strain NZ98/254. These proportions represent a decline from 13 months of age (100%, 93%) and 96%, respectively).3 However, a good response to a booster dose of vaccine was observed. Experience with other meningococcal vaccines has shown that waning of bactericidal antibody titres was associated with a decline in vaccine effectiveness following infant vaccination with serogroup C meningococcal conjugate vaccines,12 adolescent vaccination with an investigational outer membrane vesicle vaccine in Norway¹³ and infant vaccination with the New Zealand outer membrane vesicle vaccine.^{10,14} Predicting the potential impact of a decline in bactericidal antibodies on vaccine effectiveness following vaccination with 4CMenB is less straightforward. The serogroup C meningococcal conjugate and outer membrane vesicle vaccines generate an immune response primarily directed against a single antigen (the capsular polysaccharide and PorA, respectively), whereas 4CMenB is a multicomponent vaccine that aims to induce antibodies against each component. The immunogenicity of 4CMenB antigens can therefore be assessed only by use of multiple meningococcal strains, and there is considerable interstrain variation in the rate of decline of hSBA titres. It is unclear whether this reflects true differences in the persistence of efficacy for each vaccine component, or if this simply reflects different susceptibilities of the strains to killing in the hSBA assay. However, it does raise the possibility that, after vaccination, susceptibility to meningococcal infection could develop at different rates for different strains, depending on whether they are expressing proteins recognized by persistent or waning antibodies. It is also notable that the bactericidal antibody concentrations against strains 44/76-SL (fHbp) and M10713 (NHBA) were similar in children previously vaccinated with 4CMenB and those who were MenB vaccine-naive. Although this may suggest the acquisition of natural immunity against these strains (evident in more than twothirds of the naive cohort), these children had a relatively poor increase in bactericidal antibodies following a single dose of 4CMenB when compared with the vaccine-primed children. This suggests previous vaccination with 4CMenB has resulted in effective priming of the immune systems of vaccine recipients, but whether this means they have any greater protection against serogroup B meningococcal disease than MenB vaccine–naive children with the same hSBA titres remains to be seen.

A related issue is the proportion of serogroup B meningococcal strains likely to be prevented by vaccination with 4CMenB. Because the use of the hSBA assay for a very large panel of strains is not practical for technical reasons, attempts to predict coverage of meningococcal vaccines in development using surrogate assays more suitable for high throughput and standardization have been used. One example of this is the Meningococcal Antigen Typing System developed by Novartis.15 The system evaluates a representative panel of meningococcal strains to correlate the strength of binding (i.e., relative potency) of vaccine antigen-specific immunoglobulin G to the presence or absence of strainspecific bactericidal activity on the hSBA assay (performed using pooled serum obtained after vaccination at age 13 mo). Regional reference laboratories can then calculate the percentage of local invasive strains where the relative potency is above this positive bactericidal threshold for 1 or more of fHbp, NadA or NHBA (or where the PorA variant is homologous to the outer membrane vesicle component of 4CMenB). Using this technique, the Meningococcal Antigen Typing System predicts vaccination coverage of 78% of invasive serogroup B meningococcal strains in Europe,16 76% in Australia17 and 66% in Canada.¹⁸ Provisional data from the African meningitis belt predict 100% coverage for 9 serogroup X meningococcal strains.¹⁹ The demonstration of waning bactericidal antibodies in this study suggests that pooled sera obtained 2-3 years after vaccination would kill a smaller proportion of strains in the assay than that obtained at 13 months (1 mo after vaccination).

The above issues address the generation of direct protection by vaccination with 4CMenB. However, the impact of this vaccine will be determined at least as much by the presence or absence of herd immunity, which has been especially important for other meningococcal vaccines to ensure sustained protection in the face of waning immunity. Studies are currently underway to determine whether 4CMenB is likely to influence nasopharyngeal carriage and hence induce herd immunity by reducing circulation of the organism (NCT01214850). If not, the maintenance of adequate levels of bactericidal antibodies to provide direct protection may be even more critical than for conjugate vaccines. If vaccination with 4CMenB does influence nasopharyngeal carriage, then it will be important to monitor the population of "carried" strains to assess for the potential emergence of strains either lacking the vaccine antigens or expressing these at low levels, as this could herald the emergence of meningococci less susceptible to prevention by 4CMenB.

Limitations

There were a number of limitations in this study, including the relatively small numbers of participants and the fact that 45% of participants from the original study who were eligible to take part in the follow-up study did not do so. There is a potential for bias, in that children who tolerated the vaccine better in the original study may have been more likely to participate in the follow-up study. Furthermore, the control group underrecruited. Follow-up studies are currently being conducted from the larger phase IIb and III studies (NCT00944034, NCT00847145) and will provide additional data on antibody persistence and tolerability. There is as yet no information on persistence after booster doses in midchildhood; these data will become available soon because children from this study will be revisited when they reach 5 years of age (NCT01027351).

Conclusion

Consistent with other vaccines against meningococcal disease, a waning of hSBA titres was observed after infant vaccination with 4CMenB. A booster dose during preschool years was well tolerated. If 4CMenB were to be introduced into a routine vaccination schedule, measures such as adequate disease surveillance would be important to determine whether waning of antibodies might influence the effectiveness of a vaccination campaign against this bacterium.

References

- Summary of opinion: Bexsero. London (UK): European Medicines Agency; 2012. Available: www.ema.europa.eu/docs/en_GB /document_library/Summary_of_opinion_-_Initial_authorisation /human/002333/WC500134836.pdf (accessed 2013 May 8).
- Gossger N, Snape MD, Yu LM, et al. Immunogenicity and tolerability of recombinant serogroup B meningococcal vaccine administered with or without routine infant vaccinations according to different immunization schedules: a randomized controlled trial. JAMA 2012;307:573-82.
- Findlow J, Borrow R, Snape MD, et al. Multicenter, open-label, randomized phase II controlled trial of an investigational recombinant meningococcal serogroup B vaccine with and without outer membrane vesicles, administered in infancy. *Clin Infect Dis* 2010;51:1127-37.

- Snape MD, Dawson T, Oster P, et al. Immunogenicity of two investigational serogroup B meningococcal vaccines in the first year of life: a randomized comparative trial. *Pediatr Infect Dis J* 2010;29:e71-9.
- Santolaya ME, O'Ryan ML, Valenzuela MT, et al. Immunogenicity and tolerability of a multicomponent meningococcal serogroup B (4CMenB) vaccine in healthy adolescents in Chile: a phase 2b/3 randomised, observer-blind, placebo-controlled study. *Lancet* 2012;379:617-24.
- Vesikari T, Esposito S, Prymula R, et al. Immunogenicity and safety of an investigational multicomponent, recombinant, meningococcal serogroup B vaccine (4CMenB) administered concomitantly with routine infant and child vaccinations: results of two randomised trials. *Lancet* 2013;381:825-35.
- Summary of product characteristics: Bexsero. London (UK): European Medicines Agency; 2013. Available: www.ema.europa .eu/docs/en_GB/document_library/EPAR_-_Product_Information /human/002333/WC500137881.pdf (accessed 2013 May 8).
- Ramsay M. Current epidemiology of meningococcal disease in the UK and Europe. London (UK): Meningitis Research Foundation; 2011.
- Findlow J, Holland A, Andrews N, et al. Comparison of phenotypically indistinguishable but geographically distinct Neisseria meningitidis Group B isolates in a serum bactericidal antibody assay. *Clin Vaccine Immunol* 2007;14:1451-7.
- Jackson C, Lennon D, Wong S, et al. Antibody persistence following MeNZB vaccination of adults and children and response to a fourth dose in toddlers. *Arch Dis Child* 2011;96:744-51.
- Clopper CJ, Pearson ES. The use of confidence or fiducial limits illustrated in the case of the binomial. *Biometrika* 1934;26:404-13.
- Trotter CL, Andrews NJ, Kaczmarski EB, et al. Effectiveness of meningococcal serogroup C conjugate vaccine 4 years after introduction. *Lancet* 2004;364:365-7.
- Holst J, Feiring B, Fuglesang JE, et al. Serum bactericidal activity correlates with the vaccine efficacy of outer membrane vesicle vaccines against Neisseria meningitidis serogroup B disease. *Vaccine* 2003;21:734-7.
- Galloway Y, Stehr-Green P, McNicholas A, et al. Use of an observational cohort study to estimate the effectiveness of the New Zealand group B meningococcal vaccine in children aged under 5 years. *Int J Epidemiol* 2009;38:413-8.
- Donnelly J, Medini D, Boccadifuoco G, et al. Qualitative and quantitative assessment of meningococcal antigens to evaluate the potential strain coverage of protein-based vaccines. *Proc Natl Acad Sci U S A* 2010;107:19490-5.
- Donnelly J, Medini D, Giuliani MM, et al. Estimating the potential strain coverage in Europe of a multicomponent vaccine targeting serogroup B meningococci. Proceedings of the 11th meeting of the European Meningococcal Disease Society (EMGM); 2011 May 18–20; Ljubljana (Slovenia).
- Smith H, Nissen M, Sloots T, et al. Estimating the potential strain coverage in Australia of a multicomponent vaccine targeting serogroup B meningococci. Melbourne (Australia): World Society for Pediatric Infectious Diseases; 2011.
- Bettinger JA, Scheifele DW, Halperin SA, et al. Diversity of Canadian meningococcal serogroup B isolates and estimated coverage by an investigational meningococcal serogroup B vaccine (4CMenB). *Vaccine* 2013; Apr. 12 [Epub ahead of print].
- Hong E, Giuliani MM, Deghmane AE, et al. Could the multicomponent meningococcal serogroup B vaccine (4CMenB) control Neisseria meningitidis capsular group X outbreaks in Africa? Vaccine 2013;31:1113-6.

Competing interests: Matthew Snape and Andrew Pollard have conducted clinical trials on behalf of the University of Oxford, sponsored by Novartis Vaccines and Diagnostics, Pfizer, GlaxoSmithKline and Sanofi-Pasteur MSD. Andrew Pollard and Matthew Snape do not receive personal payments from vaccine manufacturers; grants for support of educational activities are paid to an educational/administrative fund held by the Department of Paediatrics, University of Oxford. Matthew Snape and Tessa John have received assistance to attend scientific meetings from Novartis, Pfizer and Glaxo-SmithKline. Matthew Snape has spent a period of secondment at Novartis. Andrew Pollard chairs the European Medicines Agency's Scientific Advisory Group on Vaccines and is a member of the UK Department of Health's Joint Committee on Vaccination and Immunisation Meningococcal Sub-committee. Huajun Wang, Peter Dull and Daniela Toneatto are employees of Novartis Vaccines and Diagnostics. Praveen Saroey, Hannah Robinson, Sarah Kelly, Nicoletta Gossger and Ly-Mee Yu have no competing interests to declare.

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Contributors: Peter Dull and Daniela Toneatto prepared the study protocol in collaboration with Matthew Snape, Andrew Pollard and Tessa John. The study organization, conduct and oversight was performed by Tessa John, Hannah Robinson, Sarah Kelly, Nicoletta Gossger, Matthew Snape and Andrew Pollard (chief investigator). Data analysis was performed by Huajun Wang and independently verified by Ly-Mee Yu. Interpretations were performed by Matthew Snape, Praveen Saroey, Peter Dull and Andrew Pollard. Matthew Snape and Praveen Saroey grepared the figures. All authors approved the final version of the manuscript submitted for publication.

Funding: This study was funded and sponsored by Novartis Vaccines and Diagnostics, which provided the study vaccine,

performed laboratory analysis of the sera and developed this protocol in collaboration with Andrew Pollard, Matthew Snape and Tessa John. Data were analyzed by Novartis Vaccines and Diagnostics and independently verified by Ly-Mee Yu. Editorial control over content was held by the investigators at the University of Oxford. Matthew Snape and Tessa John are supported by the NIHR (National Institute for Health Research) Oxford Biomedical Research Centre. The research was supported by the NIHR Oxford Biomedical Research Centre based at Oxford University Hospitals NHS (National Health Service) Trust and University of Oxford. The views expressed are those of the authors and not necessarily those of the National Health Service, the NIHR or the Department of Health. Andrew Pollard is a Jenner Institute investigator and a James Martin senior fellow.

Acknowledgments: The authors thank the study participants and their parents, without whom this study would not have been possible. The authors also thank the research nurses Jen Houlden and Aimee Field for their efforts in organizing and conducting study visits, Emma Plested for her outstanding administrative support and the staff of the Child Health Computer Departments in Oxfordshire, Buckinghamshire, Berkshire and Northamptonshire.