Phase III, randomized controlled trial in girls 9-15 years old to evaluate lot consistency of a novel nine-valent human papillomavirus L1 virus-like particle vaccine

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A 9-valent human papillomavirus (6/11/16/18/31/33/45/52/58) VLP (9vHPV) vaccine has recently been proven highly efficacious in preventing disease associated with vaccine HPV types in a pivotal Phase III study. The demonstration of lot-to-lot consistency to confirm the reliability of the manufacturing process is a regulatory requirement for vaccine licensure in the United States. A randomized trial was conducted to demonstrate that three lots of 9vHPV vaccine elicit equivalent antibody response for all 9 vaccine types. The study required thorough planning because it required success on 27 separate statistical comparisons. An innovative statistical approach was used taking into account between-lot variance for more conservative power calculations. The study demonstrated equivalence of three lots of 9vHPV vaccine for all 9 vaccine types.

Introduction

The 9-valent human papillomavirus (6/11/16/18/31/33/45/ 52/58) VLP (9vHPV) vaccine addresses the 4 HPV types covered by the licensed quadrivalent human papillomavirus (6/11/16/18) VLP (qHPV) vaccine (GARDASILTM/SILGARDTM) plus five additional oncogenic types for increased cervical cancer coverage. A pivotal study in young women 16 to 26 years of age demonstrated that the 9vHPV vaccine is highly efficacious in preventing disease associated with vaccine HPV types.¹ The 9vHPV vaccine was licensed in 2014 in the United States under the name GARDASILTM9.

The demonstration of lot-to-lot consistency to support the reliability of the manufacturing process is generally an obligatory step in vaccine development.^{2,3} Per regulatory guidance, consistency of manufacture needs to be demonstrated based on at least three manufacturing lots of the final manufacturing process (FMP).⁴⁻⁶ As part of this requirement, a lot consistency clinical study must be performed to show that at least three FMP lots are equivalent with respect to immunogenicity.⁷

A Phase III study was conducted to assess 9vHPV vaccine lot consistency in girls 9 to 15 years of age. This article summarizes the design and results of this study.

Study design

The V503-002 study (NCT #00943722) was designed to enroll 1800 girls 9 to 15 years of age, 600 boys 9 to 15 years of age, and 400 young women 16 to 26 years of age. The study was comprised of two immunogenicity substudies (Fig. 1): 1. An adult-adolescent immunobridging substudy to compare 9vHPV vaccine immunogenicity at Month 7 in girls versus young women, and boys versus young women; 2. A lot consistency substudy to demonstrate consistent immunogenicity at Month 7 in subjects randomized to three different vaccine lots of the final manufacturing process.

Methods

The lot consistency substudy was conducted in girls. All enrolled girls 9 to 15 years of age were equally randomized to three vaccine lots (termed lots 1, 2, and 3). Boys and young women did not participate in the lot consistency substudy and were all assigned to lot 1. As shown in Figure 1, girls who received lot 1 participated in both substudies. This report is focused on the lot consistency substudy. The results of the adult-adolescent immunobridging substudy as well as the safety findings have been reported elsewhere.⁸

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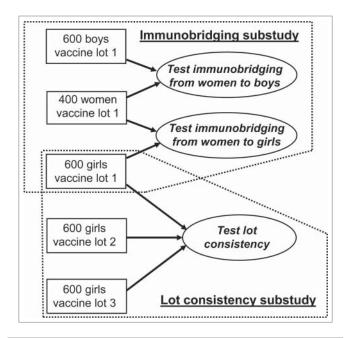


Figure 1. Subject distribution between the adult-adolescent substudy and the lot consistency substudy. The V503-002 study (NCT #00943722) was designed to enroll 1800 girls 9 to 15 years of age, 600 boys 9 to 15 years of age, and 400 young women 16 to 26 years of age. The study was comprised of two immunogenicity substudies: 1. An adult-adolescent immunobridging substudy to compare 9vHPV vaccine immunogenicity at Month 7 in girls versus young women, and boys versus young women; 2. A lot consistency substudy to demonstrate consistent immunogenicity at Month 7 in subjects randomized to three different vaccine lots of the final manufacturing process.

Population

Subjects in the lot consistency substudy were enrolled from 66 sites in 17 countries (Austria, Belgium, Brazil, Chile, Colombia, Costa Rica, Finland, India, Peru, Poland, South Africa, South Korea, Spain, Sweden, Taiwan, Thailand and the United States). The study was conducted in accordance with principles of Good Clinical Practice and was approved by the appropriate institutional review boards and regulatory agencies. An external Data Monitoring Committee assessed safety findings throughout the study.

The lot consistency substudy was designed to enroll girls 9 to 15 years of age. Participants were required to be generally healthy and sexually naïve at enrollment and throughout the vaccination period (i.e., through month 7). Reasons for exclusions from the study included: pregnancy (determined by urine or serum β -human chorionic gonadotropin testing), known allergy to any vaccine component, thrombocytopenia, immunosuppression or prior immunosuppressive therapy, and previous receipt of an HPV vaccine.

Randomization and vaccination

Following informed consent and determination that all inclusion criteria and none of the exclusion criteria were met, eligible subjects received an allocation number and were randomized to a vaccination group. An Interactive Voice Response System (IVRS) was used to allocate study subjects and balance randomization between sites. The IVRS assigned the subject an allocation number from an allocation schedule. Girls were enrolled in two age strata (9 to 12 years of age and 13 to 15 years of age at enrollment) in approximately a 2:1 ratio to ensure that the immunogenicity and safety profile of the vaccine in younger subjects would be clearly defined. Girls enrolled in the lot consistency substudy were equally randomized in a double-blinded manner to three FMP vaccine lots. The randomized allocation schedule was based on balanced randomization blocks of size 6.

All subjects were administered a 3-dose regimen of 9vHPV vaccine (at day 1, month 2, and month 6). All subjects received the same formulation of 9vHPV vaccine administered (in the deltoid) as a 0.5-mL intramuscular injection. A 0.5-mL dose of 9vHPV vaccine contains 30 μ g HPV 6, 40 μ g HPV 11, 60 μ g HPV 16, 40 μ g HPV 18, 20 μ g HPV 31, 20 μ g HPV 33, 20 μ g HPV 45, 20 μ g HPV 52, 20 μ g HPV 58, and 500 μ g of amorphous aluminum hydroxyphosphate sulfate (AAHS). All participants were required to be afebrile (oral temperature <37.8 °C) within 24 hours before each injection. All participants underwent pregnancy testing that was based on urine or serum analyses for β -human chorionic gonadotropin before each vaccination. Participants who were found to be pregnant were not to be vaccinated. No subject was found pregnant in this study.

Assessment

All participants were assessed for antibody titers to all 9 vaccine HPV types at day 1 and month 7. Seropositivity at day 1 was not a reason for exclusion from the study. However, the results of this testing were part of the criteria to define the per protocol analysis population.

Serum samples obtained at day 1 and month 7 from all subjects were tested for anti-HPV 6, anti-HPV 11, anti-HPV 16, anti-HPV 18, anti-HPV 31, anti-HPV 33, anti-HPV 45, anti-HPV 52, and anti-HPV 58 by HPV 6, 11, 16,18, 31, 33, 45, 52, 58 competitive Luminex immunoassay (HPV-9 cLIA).⁹ A subject was defined to be anti-HPV 6, anti-HPV 11, anti-HPV 16, or anti-HPV 18 seropositive if the anti-HPV serum cLIA level was ≥ 30 , ≥ 16 , ≥ 20 , or ≥ 24 mMU/mL, respectively. A subject was defined to be seropositive to HPV Types 31, 33, 45, 52, or 58 if the anti-HPV serum cLIA level was ≥ 10 , ≥ 8 , ≥ 8 , ≥ 8 , or ≥ 8 mMU/mL, respectively.⁹

All participants were observed for at least 30 minutes after each vaccination for any immediate reaction, with particular attention to any evidence of a hypersensitivity reaction. All subjects received a vaccination report card (VRC) at the day 1, month 2, and month 6 study vaccination visits. On the VRC, the parent/guardian of the subject was asked to record adverse experiences. Safety findings for this study have been reported.⁸

Statistical methodology

The primary approach to the analyses of immunogenicity was per-protocol. Each vaccine component was analyzed separately. To be included in the primary immunogenicity analysis for the HPV 6 and HPV 11 components, subjects had to be seronegative to both HPV 6 and 11 at day 1 (because of extensive cross-reactivity due to the high aminoacid sequence identity [92%] between HPV 6 and HPV 11 L1 proteins. {2583}. To be included in the primary immunogenicity analysis for the other vaccine HPV types, subjects were only required to be seronegative at day 1 for the HPV type being analyzed. In addition, subjects had to receive all 3 doses of the correct clinical material within acceptable day ranges (dose 2: days 36 to 84 and dose 3: days 148 to 218, relative to day 1) and have at least 1 post-dose 3 serology result within acceptable day ranges (days 21 to 49 post-dose 3) and not violate the protocol.

The relevant parameters for assessing lot consistency are geometric mean titers (GMTs) and seroconversion rates. The primary immunogenicity objective of the lot consistency substudy was to demonstrate that the FMP results in lots of the 9vHPV vaccine induce consistent Month 7 GMTs for serum anti-HPV 6, anti-HPV 11, anti-HPV 16,

anti-HPV 18, anti-HPV 31, anti-HPV 33, anti-HPV 45, anti-HPV 52, and anti-HPV 58. The statistical criterion for equivalence requires that the two-sided 95% confidence interval (CI) of GMT ratio for each of the three pairs of lots (lot 1 vs. lot 2, lot 1 vs. lot 3, and lot 2 vs. lot 3) be contained entirely within the interval (0.5, 2.0) for each HPV type. The secondary immunogenicity objective of the lot consistency substudy was to demonstrate that the FMP results in lots of the 9vHPV vaccine that induce consistent Month 7 seroconversion percentages to HPV types 6, 11, 16, 18, 31, 33, 45, 52, and 58. These tests were conducted using the method of Miettinen and Nurminen [213]. The statistical criterion for equivalence requires that the two-sided 95% CI of the difference between seroconversion percentages for each of three pairs of lots (lot 1 vs. lot 2, lot 1 vs. lot 3, and lot 2 vs. lot 3) be contained entirely within the interval (-5%, 5%) for each HPV type.

Margin for lot consistency

The primary analysis of lot consistency required 27 pairs of tests of equivalence for the 3 manufacturing lots and 9 HPV types. Study success was defined as meeting all 27 pairs of equivalence tests. To determine the sample size, the study power calculation was based on total variance (i.e., between-lot variance as well as within-lot variance). Between-lot variability has been generally ignored in vaccine lot consistency studies. The critical importance of considering between-lot variability in study power determination has been recently recognized.^{10,11} Some between-lot variability is expected under any manufacturing practices and, although small, it can have substantial impact on the sample size

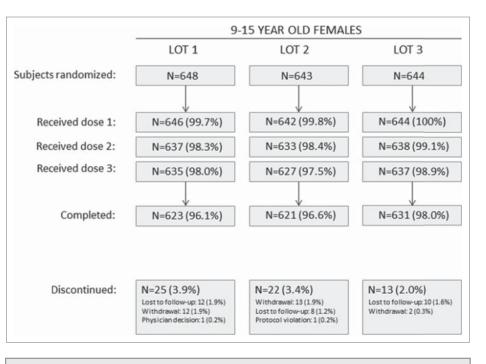


Figure 2. Accounting of subjects in the lot consistency substudy. A total of 1935 subjects were randomized from 65 sites in Africa (South Africa), Asia (India, Korea, Taiwan, Thailand), Europe (Austria, Belgium, Finland, Poland, Sweden, Spain), Latin America (Brazil, Chile, Colombia, Costa Rica, Peru) and North America (United States). Withdrawal = patient withdrew consent.

and power.¹⁰ Therefore, the study was designed to account for between-lot variability. For example, even with a small between-lot variance of less than 1% of the total variance, an equivalence

Table 1 Subject characteristics

	Lo	t 1	Lo	t 2	Lo	t 3
	n	(%)	n	(%)	n	(%)
Subjects in population	648		643		644	
Age (Years)						
9 to 12 Years of Age	440	(67.9)	432	(67.2)	432	(67.1)
13 to 15 Years of Age	208	(32.1)	211	(32.8)	212	(32.9)
Mean	11.7		11.6		11.6	
SD	1.8		1.8		1.9	
Median	12.0		11.0		11.0	
Range	9 to 15		9 to 15	5	9 to 15	5
Race						
American Indian Or Alaska Native	1	(0.2)	1	(0.2)	0	(0.0)
Asian	150	(23.1)	141	(21.9)	139	(21.6)
Black Or African American	50	(7.7)	59	(9.2)	52	(8.1)
Multi-Racial	81	(12.5)	91	(14.2)	86	(13.4)
White	366	(56.5)	351	(54.6)	367	(57.0)
Ethnicity						
Hispanic Or Latino	176	(27.2)	191	(29.7)	193	(30.0)
Not Hispanic Or Latino	472	(72.8)	452	(70.3)	451	(70.0)
Region						
Africa	32	(4.9)	34	(5.3)	29	(4.5)
Asia-Pacific	148	(22.8)	137	(21.3)	138	(21.4)
Europe	206	(31.8)	182	(28.3)	185	(28.7)
Latin America	125	(19.3)	147	(22.9)	136	(21.1)
North America	137	(21.1)	143	(22.2)	156	(24.2)

Table 1 Statistical	analysis of equivalenc	e of geometric mean	titers at month 7

				9vHPV	Vacci	ne		Estimated			
		Con	nparis	on Group A	Con	nparis	on Group B	Fold Difference	p-Value for	Equivalence met (YES/NO))
	Comparison Group A vs.			GMT [‡]			GMT [‡]	Group A / Group B		Equivalence	•
Assay (cLIA)	Comparison Group B	Ν	n	(mMU/mL)	Ν	n	(mMU/mL)	(95% CI)	$\mathbf{Left}^{\mathbb{S}}$	Right	
Anti-HPV 6	Lot 1 vs. Lot 2	646	517	1,603.6	642	536	1,645.8	0.97 (0.88, 1.08)	<0.001	<0.001	YES
	Lot 1 vs. Lot 3	646	517	1,603.6	644	544	1,550.0	1.03 (0.93, 1.16)	< 0.001	<0.001	YES
	Lot 2 vs. Lot 3	642	536	1,645.8	644	544	1,550.0	1.06 (0.95, 1.19)	< 0.001	<0.001	YES
Anti-HPV 11	Lot 1 vs. Lot 2	646	517	1,221.9	642	536	1,223.2	1.00 (0.90, 1.11)	< 0.001	<0.001	YES
	Lot 1 vs. Lot 3	646	517	1,221.9	644	544	1,143.6	1.07 (0.95, 1.20)	< 0.001	<0.001	YES
	Lot 2 vs. Lot 3	642	536	1,223.2	644	544	1,143.6	1.07 (0.95, 1.20)	< 0.001	<0.001	YES
Anti-HPV 16	Lot 1 vs. Lot 2	646	529	6,465.1	642	542	6,764.7	0.96 (0.86, 1.06)	< 0.001	<0.001	YES
	Lot 1 vs. Lot 3	646	529	6,465.1	644	556	6,456.5	1.00 (0.90, 1.12)	< 0.001	<0.001	YES
	Lot 2 vs. Lot 3	642	542	6,764.7	644	556	6,456.5	1.05 (0.94, 1.17)	< 0.001	<0.001	YES
Anti-HPV 18	Lot 1 vs. Lot 2	646	531	1,976.8	642	547	1,969.5	1.00 (0.89, 1.14)	< 0.001	<0.001	YES
	Lot 1 vs. Lot 3	646	531	1,976.8	644	563	1,778.3	1.11 (0.98, 1.26)	< 0.001	<0.001	YES
	Lot 2 vs. Lot 3	642	547	1,969.5	644	563	1,778.3	1.11 (0.97, 1.26)	< 0.001	<0.001	YES
Anti-HPV 31	Lot 1 vs. Lot 2	646	522	1,742.3	642	542	1,736.2	1.00 (0.89, 1.13)	< 0.001	<0.001	YES
	Lot 1 vs. Lot 3	646	522	1,742.3	644	553	1,701.7	1.02 (0.91, 1.16)	< 0.001	<0.001	YES
	Lot 2 vs. Lot 3	642	542	1,736.2	644	553	1,701.7	1.02 (0.90, 1.15)	< 0.001	<0.001	YES
Anti-HPV 33	Lot 1 vs. Lot 2	646	534	913.6	642	543	873.0	1.05 (0.94, 1.16)	< 0.001	<0.001	YES
	Lot 1 vs. Lot 3	646	534	913.6	644	560	868.6	1.05 (0.94, 1.17)	< 0.001	<0.001	YES
	Lot 2 vs. Lot 3	642	543	873.0	644	560	868.6	1.01 (0.90, 1.12)	< 0.001	<0.001	YES
Anti-HPV 45	Lot 1 vs. Lot 2	646	534	643.3	642	548	770.3	0.84 (0.73, 0.95)	< 0.001	<0.001	YES
	Lot 1 vs. Lot 3	646	534	643.3	644	565	620.8	1.04 (0.91, 1.18)	< 0.001	<0.001	YES
	Lot 2 vs. Lot 3	642	548	770.3	644	565	620.8	1.24 (1.08, 1.42)	< 0.001	<0.001	YES
Anti-HPV 52	Lot 1 vs. Lot 2	646	533	862.7	642	547	934.1	0.92 (0.83, 1.03)	< 0.001	<0.001	YES
	Lot 1 vs. Lot 3	646	533	862.7	644	562	905.4	0.95 (0.85, 1.07)	< 0.001	< 0.001	YES
	Lot 2 vs. Lot 3	642	547	934.1	644	562	905.4	1.03 (0.92, 1.16)	< 0.001	<0.001	YES
Anti-HPV 58	Lot 1 vs. Lot 2	646	531	1,197.7	642	539	1,255.5	0.95 (0.86, 1.06)	< 0.001	< 0.001	YES
	Lot 1 vs. Lot 3	646	531	1,197.7	644	560	1,118.3	1.07 (0.96, 1.20)	< 0.001	< 0.001	YES
	Lot 2 vs. Lot 3	642	539	1,255.5	644	560	1,118.3	1.12 (1.00, 1.26)	< 0.001	<0.001	YES

[†]The per-protocol immunogenicity population includes all subjects who were not general protocol violators, received all 3 vaccinations within acceptable day ranges, were seronegative at Day 1 for the relevant HPV type(s), and had a Month 7 serum sample collected within an acceptable day range. [‡]Based on an ANCOVA model with a response of the natural log of individual titers and fixed effects for lots and age strata.

[§]p-value for the comparison of the GMT ratio to the lower bound (0.5).

¹¹p-value for the comparison of the GMT ratio to the upper bound (2.0).

A p-value (Left) ≤ 0.025 and a p-value (Right) ≤ 0.025 together support a conclusion of equivalence. If equivalence can be established in all 3 pairwise comparison for a given HPV type and endpoint, the 3 lots will be considered consistent for that HPV type. Equivalence must be established for all 9 HPV types for the 3 HPV vaccine lots to be considered consistent.

N = Number of subjects randomized to the respective vaccination group who received at least 1 injection.

n = Number of subjects with evaluable serology data and are eligible for the indicated analysis population.

CI = Confidence interval; cLIA = Competitive Luminex immunoassay; GMT = Geometric mean titer; mMU = Milli Merck units.

PCR = Polymerase chain reaction, HPV = Human papillomavirus, VLP = Virus-like particles.

margin of 1.5-fold would require a sample size up to 6 times greater than that utilizing an equivalence margin of 2-fold. Thus, to account for between-lot variability and conduct the study with an acceptable sample size and power, a 2-fold equivalence margin was found to be appropriate for this lot consistency substudy. Lot consistency is concluded if the two-sided 95% confidence interval of the ratio of geometric means of titers (GMTs) for each of the three pairs of lots is contained within the interval (0.5, 2.0).

Under the assumptions of approximately 20% exclusion rate for the per-protocol population in this age group and 2.0-fold equivalence margin for GMT ratio, the lot consistency substudy with 600 subjects per group would provide above 90% power to show consistency for the 3 lots for all 9 HPV types at an overall one-sided, 2.5% alpha-level as long as the between-lot variance is not above 1.2% of the total variance (between-lot variance and within-lot variance).

Results

A total of 1935 subjects were randomized from 66 sites in Africa (South Africa), Asia (India, Korea, Taiwan, Thailand), Europe (Austria, Belgium, Finland, Poland, Sweden, Spain), Latin America (Brazil, Chile, Colombia, Costa Rica, Peru) and North America (United States). A summary of the number of subjects who were randomized, vaccinated, who completed or discontinued during the study can be seen in Figure 2. No

		9vHPV Vaccine						Estimated		
		Seroconversion Rate Comparison Group A	Rate Compar	ison Group A	Seroconver	sion Rate Con	Seroconversion Rate Comparison Group B	Percentage Point		
			·	Estimated			Estimated	Difference	p-Value for	
	Comparison Group A vs.			Response			Response	Group A / Group B	Equivalence	
Assay (cLIA)	Comparison Group B	N	Ľ	(%)	N	Ľ	(%)	(95% CI)	Left [§] Right	Equivalence met (YES/NO)
cLIA 6	Lot 1 vs. Lot 2	517	516	99.8	536	535	99.8	-0.0 (-0.9, 0.9)	<0.001 <0.001	YES
	Lot 1 vs. Lot 3	517	516	99.8	544	540	99.3	0.5 (-0.4, 1.7)		YES
	Lot 2 vs. Lot 3	536	535	99.8	544	540	99.3	0.6 (-0.4, 1.7)	<0.001 <0.001	YES
cLIA 11	Lot 1 vs. Lot 2	517	517	100	536	536	100	0.0 (-0.7, 0.7)	<0.001 <0.001	YES
	Lot 1 vs. Lot 3	517	517	100	544	542	9.66	0.4 (-0.4, 1.3)	<0.001 <0.001	YES
	Lot 2 vs. Lot 3	536	536	100	544	542	9.66	0.4 (-0.4, 1.3)	<0.001 <0.001	YES
cLIA 16	Lot 1 vs. Lot 2	529	529	100	542	542	100	0.0 (-0.7, 0.7)		YES
	Lot 1 vs. Lot 3	529	529	100	556	554	9.66	0.4 (-0.4, 1.3)	<0.001 <0.001	YES
	Lot 2 vs. Lot 3	542	542	100	556	554	9.66	0.4 (-0.4, 1.3)	<0.001 <0.001	YES
cLIA 18	Lot 1 vs. Lot 2	531	530	99.8	547	547	100	-0.2 (-1.1, 0.5)		YES
	Lot 1 vs. Lot 3	531	530	99.8	563	561	99.7	0.2 (-0.7, 1.1)	<0.001 <0.001	YES
	Lot 2 vs. Lot 3	547	547	100	563	561	9.66	0.4 (-0.4, 1.3)	<0.001 <0.001	YES
cLIA 31	Lot 1 vs. Lot 2	522	522	100	542	542	100	0.0 (-0.7, 0.7)	<0.001 <0.001	YES
	Lot 1 vs. Lot 3	522	522	100	553	552	99.8	0.2 (-0.6, 1.0)	<0.001 <0.001	YES
	Lot 2 vs. Lot 3	542	542	100	553	552	99.8	0.2 (-0.5, 1.0)	·	YES
cLIA 33	Lot 1 vs. Lot 2	534	534	100	543	543	100	0.0 (-0.7, 0.7)	<0.001 <0.001	YES
	Lot 1 vs. Lot 3	534	534	100	560	558	9.66	0.4 (-0.4, 1.3)		YES
	Lot 2 vs. Lot 3	543	543	100	560	558	9.66	0.4 (-0.4, 1.3)		YES
cLIA 45	Lot 1 vs. Lot 2	534	533	99.8	548	548	100	-0.2 (-1.1, 0.5)		YES
	Lot 1 vs. Lot 3	534	533	99.8	565	563	99.7	0.2 (-0.7, 1.1)	•	YES
	Lot 2 vs. Lot 3	548	548	100	565	563	9.66	0.4 (-0.4, 1.3)	'	YES
cLIA 52	Lot 1 vs. Lot 2	533	533	100	547	547	100	0.0 (-0.7, 0.7)	•	YES
	Lot 1 vs. Lot 3	533	533	100	562	560	99.7	0.3 (-0.4, 1.3)	<0.001 <0.001	YES
	Lot 2 vs. Lot 3	547	547	100	562	560	9.66	0.4 (-0.4, 1.3)	<0.001 <0.001	YES
cLIA 58	Lot 1 vs. Lot 2	531	531	100	539	539	100	0.0 (-0.7, 0.7)	•	YES
	Lot 1 vs. Lot 3	531	531	100	560	558	9.66	0.4 (-0.4, 1.3)	<0.001 <0.001	YES
	Lot 2 vs. Lot 3	539	539	100	560	558	9.66	0.4 (-0.4, 1.3)	<0.001 <0.001	YES
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The per-prot	ocol immunogenicity popu	lation includes all	subjects who	were not gene	al protocol vic	olators, receiv	ed all 3 vaccination	s within acceptable da	ıy ranges, were s	The per-protocol immunogenicity population includes all subjects who were not general protocol violators, received all 3 vaccinations within acceptable day ranges, were seronegative at Day 1 for the
relevant mrv	relevant mrv type(s), and nad a Montri / serum sample conected with	serum sample collo	ected within a	n an acceptable day range.	ay range.					
p-value for tr	"p-value for the comparison of the difference of seroconversion rates to the lower bound (-5%).	ence of seroconver	sion rates to th	he lower bound	d (-5%).					

Table 2 Statistical analysis of equivalence of proportions of subjects who seroconverted by month 7

 $^{\parallel}$ p-value for the comparison of the difference of seroconversion rates to the upper bound (5%).

Seroconversion was defined as changing serostatus from seronegative to seropositive. Cutoff values for HPV seropositivity are \geq 30, 16, 20, 24, 10, 8, 8, 8, and 8 mMU/mL for HPV types 6, 11, 16, 18, 31, 33, 45, 52, and 58, respectively.

A p-value (Left) \leq 0.025 and a p-value (Right) \leq 0.025 together support a conclusion of equivalence. If equivalence can be established in all 3 pairwise comparison for a given HPV type and endpoint, the 3 lots will be considered consistent for that HPV type. Equivalence must be established for all 9 HPV types for the 3 HPV vaccine lots to be considered consistent.

The estimated proportions, proportion difference, associated confidence intervals, and p-values are based on the method of Miettinen and Nurminen, stratified by age strata using CMH weights. N = Number of subjects with evaluable serology data and are eligible for the indicated analysis population.

n = Number of subjects who had seroconversion.

CI = Confidence interval; cLIA = Competitive Luminex immunoassay; GMT = Geometric mean titer; mMU = Milli Merck units. HPV = Human papillomavirus; VLP = Virus-like particles.

subject discontinued study vaccination due to an adverse event. A summary of baseline subject characteristics is provided in **Table 1**. All vaccination groups were diverse with respect to geographic region, race, and ethnicity.

At enrollment, serum antibody titers for HPV 6, 11, 16, 18, 31, 33, 45, 52, or 58 that were greater than the predefined seropositive threshold values for one or more HPV type, indicative of previous exposure to the respective vaccine HPV types, were detected in 8.5% (163 of 1,920) of the study participants.

Table 1 displays the results of the statistical analyses of lot equivalence of month 7 HPV cLIA GMTs for each vaccine HPV type in the per-protocol immunogenicity (PPI) population. The lower bound of two sided 95% CI of GMT ratio for each of three pairs of lots (lot 1 vs. lot 2, lot 1 vs. lot 3, and lot 2 vs. lot 3) for all 9 vaccine types (representing a total of 27 comparisons) was contained entirely within the interval (0.5, 2.0) for each HPV type, with p-values <0.001, indicating cLIA GMT responses in groups who received the three vaccine lots were equivalent., The criterion for equivalent antibody responses for this secondary endpoint was met as well (**Table 2**). Overall, 99.6% of participants seroconverted by month 7 to all 9 HPV types.

Discussion

This study was designed to demonstrate manufacturing lot consistency of the 9vHPV vaccine. Three groups of girls 9 to 15 years of age were randomized to three lots of the FMP process. At one month after the completion of a 3-dose vaccination regimen, statistical equivalence of the GMTs between the 3 lots was established for all 9 vaccine types. Seroconversion rates were over 99% in all three groups and found to be statistically equivalent. These results support the conclusion that the 9vHPV vaccine from three lots of the FMP induces consistent antibody responses to all 9 vaccine HPV types.

A few subjects did not seroconvert at month 7. Because there is no known immune correlate of protection for HPV vaccines (e.g., minimum level of antibody that predicts protection against infection or disease), the absence of detectable antibodies is not equivalent to absence of protection. The minimum antibody titer needed for protection is not known but animal studies suggest that very low antibody titers (e.g., up to 100-fold lower than the threshold of detection of a standard pseudovirion-based neutralizing assay [PBNA]) may be protective.{2659} This result is also relevant to cLIA (the immunoassay used in this study) since PBNA and cLIA are highly correlated and similarly sensitive. {2736}

Lot consistency demonstration is generally required to support vaccine registration.¹⁻³ This entails a randomized clinical trial to establish that three different manufacturing lots provide equivalent immunogenicity. The specific requirements of each lot consistency study should be determined based on a thorough investigation considering the particulars of the study, as sample size and risk of failure can be dramatically increased if

requirements are too stringent. Study parameters to consider include number of vaccine genotypes, desired power to demonstrate lot consistency, assumed within-lot variance for each genotype, assumed between-lot variance for each genotype, and lot consistency criteria.

It is known that an increase in sample size reduces the withinlot variance of the mean, but does not affect the between-lot variance.¹⁰ Even under the best manufacturing practices, small variations between lots are expected. A small between-lot variance can greatly impact sample size and compromise study success as was the case in a previous lot consistency study of a meningococcal vaccine.^{10,11} In the present study, these issues were compounded by the need to demonstrate equivalence of three vaccine lots, each containing antigens from nine HPV types, which represents 27 separate comparisons. Based on a comprehensive investigation the study design described herein was found to be appropriate for demonstration of lot consistency. These considerations may be applicable to other vaccine development programs and may aid in designing efficient and successful vaccine lot consistency studies.

Disclosure of Potential Conflicts of Interest

Edson Moreira reports grants and personal fees from Merck & Co., Inc. Alain Luxembourg, Xiao Sun, Roger Maansson, Susan Christiano Erin Moeller and Joshua Chen are employees of Merck & Co., Inc. and may hold stock and/or stock options. Rudiwilai Samakoses and Kyung-Hyo Kim report no conflicts of interest.

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