## **Original Article**

# Modeling the Long-term Antibody Response and Duration of Immune Protection Induced by an Inactivated, Preservative-free Hepatitis A Vaccine (Healive) in Children<sup>\*</sup>



YU Yong Pei<sup>1,2,&</sup>, CHEN Jiang Ting<sup>3,&</sup>, JIANG Zhi Wei<sup>4</sup>, WANG Ling<sup>1</sup>, YU Cheng Kai<sup>1</sup>, YAN Xiao Yan<sup>2</sup>, YAO Chen<sup>2</sup>, and XIA Jie Lai<sup>1,#</sup>

College of Military Preventive Medicine, the Fourth Military Medical University, Xi'an 710032, Shaanxi, China;
 Peking University Clinical Research Institute, Peking University Health Science Center, Beijing 100191, China;
 Sinovac Biotech, Beijing 100089, China; 4. Beijing Keytech Statistical Consulting, Beijing 100020, China

## Abstract

**Objective** Long-term seroprotection *via* the hepatitis A vaccine is essential for the prevention of disease from the hepatitis A virus (HAV). Due to documented difficulties during decade-long follow-ups after receiving vaccines, statistical-modeling approaches have been applied to predict the duration of immune protection.

**Methods** Based on five-year follow-up data from a randomized positive-controlled trial among Chinese children (1–8 years old) following a 0, 6 months vaccination schedule, a power-law model accounting for the kinetics of B-cell turnover, as well as a modified power-law model considering a memory-B-cell subpopulation, were fitted to predict the long-term immune responses induced by HAV vaccination (Healive or Havrix). Anti-HAV levels of each individual and seroconversion rates up to 30 years after vaccination were predicted.

**Results** A total of 375 participants who completed the two-dose vaccination were included in the analysis. Both models predicted that, over a life-long period, participants vaccinated with Healive would have close but slightly higher antibody titers than those of participants vaccinated with Havrix. Additionally, consistent with previous studies, more than 90% of participants were predicted to maintain seroconversion for at least 30 years. Moreover, the modified power-law model predicted that the antibody titers would reach a plateau level after nearly 15 years post-vaccination.

**Conclusions** Based on the results of our modeling, Healive may adequately induce long-term immune responses following a 0, 6 months vaccination schedule in children *via* induction of memory B cells to provide stable and durable immune protection.

Key words: Hepatitis A; Inactivated hepatitis A vaccine; Modeling; Antibody persistence; Long-term follow-up

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<sup>&</sup>lt;sup>&</sup>These authors contributed equally to this work.

<sup>&</sup>lt;sup>#</sup>Correspondence should be addressed to XIA Jie Lai, Tel: 86-13571999716, E-mail: xiajielai@fmmu.edu.cn

Biographical notes of the first authors: YU Yong Pei, male, born in 1984, MS, majoring in biostatistics; CHEN Jiang Ting, female, born in 1962, MD, majoring in medical science and clinical study.

#### INTRODUCTION

epatitis A is a common acute infectious disease caused by the hepatitis A virus (HAV). The HAV is primarily transmitted by the fecal-oral route, usually through the ingestion of contaminated food or water or through contact with an infectious person<sup>[1]</sup>. Hepatitis A is usually self-limited and can cause patients to be asymptomatic to severe illness<sup>[2]</sup>. The infection of HAV usually causes more severe symptoms in older children and adults. HAV patients, particularly older than 40 years old, may experience several weeks of jaundice and may be at risk of acute liver failure and death<sup>[3,4]</sup>. The epidemic of hepatitis A is primarily determined by sanitary and socioeconomic conditions, and the consequent burden of medical care can be substantial, especially in developing countries<sup>[5]</sup>.

In recent decades, the epidemiology of hepatitis A has changed significantly. With the improvement of hygienic conditions in several resource-poor countries, the global incidence of hepatitis A continues to decrease and the average age at infection continues to increase<sup>[1,6]</sup>. Consequently, the incidence of severe illnesses related to hepatitis A—including fulminant hepatitis, hospitalizations, and deaths—is increasing, despite a reduction in the rate of HAV transmission<sup>[6]</sup>.

Prevention of hepatitis A is preferable to symptomatic treatments, as no specific anti-HAV drug is currently available<sup>[7]</sup>. In the 1990s, several HAV vaccines became available and the protective efficacies of these vaccines have been demonstrated in different trials<sup>[8-10]</sup>. Among these studies, the cumulative rates of protective efficacy have been found to be 95%–100%. Long-term immunogenicity and immune persistence of inactivated hepatitis A vaccines have also been investigated in post-marketing studies<sup>[11-19]</sup>. Mosites et al.<sup>[15]</sup> performed a 22-year follow-up study and showed that more than 90% of subjects maintained protective levels of anti-HAV.

Although there is accumulated evidence of long-term immune persistence of hepatitis A vaccines, the precise duration of immune protection after vaccination remains unknown. This is because the estimate of duration requires decades of follow-ups to obtain individual immuneduration data. Alternatively, statistical-modeling approaches have been applied to provide a mathematical prediction of the duration of immune protection. Van Herck et al.<sup>[20]</sup> as well as Van Herck and Van Damme<sup>[21]</sup> used exponential models with fixed rates for serum-antibody kinetics estimate the long-term persistence of to inactivated HAV vaccines; these two studies yielded predictions of immune persistence of at least 10 years and 20-25 years, respectively. Another common framework for predictive modeling includes fractional-polynomial modeling and linear-mixed modeling with changepoints. In this framework, multiple phases with different antibody-decay rates are considered. A group of models are then fitted and selected based on fitting statistics. One or more models with the best fitting(s) would be used to perform the prediction. Using this framework, Hens et al.<sup>[22]</sup> predicted that at least 95% of subjects would remain anti-HAV seropositive for more than 25 years based on up to 17 years of follow-up data from two clinical trials. López et al.<sup>[23]</sup> reported a similar prediction that 88% of subjects would remain seroprotected for at least 30 years based on 15-year follow-up data. Models considering B-cell population kinetics have also been applied in predicting long-term antibody responses. Fraser et al.<sup>[24]</sup> proposed a power-law that considering B-cells model as two subpopulations, namely, active and memory B cells. Andraud et al.<sup>[25]</sup> applied the 'plasma-cell imprinted lifespan' model<sup>[26]</sup> on 10-year immunepersistence data and estimated that at least 95% of subjects would remain anti-HAV seropositive for more than 25 years.

China has been classified as a low-HAV endemic country<sup>[1]</sup>, where resources have remained invested into the prevention of infectious disease, including hepatitis A. Healive (Sinovac Biotech, Beijing, China), a new inactivated, preservative-free hepatitis A vaccine, was licensed in China in 2001. The protective efficacy and safety of Healive have been demonstrated in adults and children<sup>[27,28]</sup>. A five-year follow-up study showed that the seroprotection of Healive was consistent with that of a control vaccine<sup>[29]</sup>. However, information on the long-term persistence of immune protection of Healive remains limited.

The aim of the present study was to fit statistical models to the predicted long-term persistence of antibodies induced by HAV vaccines (Healive or Havrix), based on five-year follow-up data from a randomized positive-controlled trial<sup>[28,29]</sup>, and to predict the duration of immune protection in Chinese children after a two-dose vaccination schedule.

#### MATERIALS AND METHODS

#### **Trial Design and Population**

The present analysis was based on a doubleblind, randomized controlled trial (NCT00534885). The objective of the study was to evaluate the immunogenicity, safety, and lots-consistency of Healive. Details of the study design and outcomes of the trial have been described elsewhere<sup>[28,29]</sup>. Briefly, a total of 400 healthy children (1-8 years old; 203 males and 197 females, respectively) were randomly assigned into the following four treatment groups: three consecutive lots of Healive, and Havrix as a control vaccine. Participants were scheduled for vaccination according to a 0, 6 months schedule, among which 392 participants received the second dose. A total of 94, 95, and 94 subjects received lot 1, lot 2, and lot 3 of Healive, respectively, and 92 subjects received Havrix. Blood samples were collected at 1, 6, and 7 months after the first dose. Among the participants who complete the second vaccination, 375 volunteers participated in the follow-up phase, in which immunogenicity was monitored annually over the following five years.

A microparticle enzyme immunoassay (MEIA; AxSYM HAVAB 2.0 quantification kit, Abbott, Wiesbaden, Germany) was used to measure the serum anti-HAV antibody titers. Results are expressed in mIU/mL. The minimum detectable titer of anti-HAV was 5 mIU/mL, and titers above 20,000 mIU/mL were recorded as 20,000 mIU/mL. Anti-HAV  $\geq$  20 mIU/mL was categorized as seroconversion.

The protocol was reviewed and approved by the ethics review committee of the Changzhou Center for Disease Control and Prevention. Written informed consent was obtained from parents or guardians of participants prior to their enrollment.

#### Statistical Modeling of Antibody Duration

Anti-HAV titers at 1, 12, 24, 36, 48, and 60 months after the second HAV vaccination were used to model the long-term duration of antibody responses induced by hepatitis A vaccines. Considering the fact that low-level antibodies may persist up to a life-long period after vaccination, models that accounted for a turnover of memory B-cell pools or long-lived plasma cells would be preferable when predicting the long-term immune duration.

In the present study, we used two nonlinear mixed-effects models—namely, the power-law

model and a modified power-law model proposed by Fraser et al.<sup>[24]</sup>—that took into account the rates of B-cell decay to describe antibody kinetics. For simplicity, we refer to these two models as model 1 (i.e., the power-law model) and model 2 (i.e., the modified power-law model). Assuming that the rate of B-cell decay follows a gamma distribution, model 1 is given by the following:

$$f(t) = k - a \log_{10} (c + t)$$
(1)

where f(t) is the log<sub>10</sub> antibody titer at time t, k is the peak antibody titer (log<sub>10</sub> mIU/mL), a is the decay rate, and c is an arbitrary small constant (often set to zero).

Fraser et al.<sup>[24]</sup> proposed a modified power-law model, in which B cells are considered to be comprised of two subpopulations, namely activated and memory B-cells. The model assumes that the amount of activated B cells decreases over time, while the amount of memory B cells is constant over time due to a rapid turnover of memory B cells<sup>[30]</sup>. By including the component that accounts for memory B cells, the model would allow for a long-term antibody plateau. Model 2 is given by the following:

$$f(t) = k + \log_{10} \left[ (1 - \pi) t^{-a} + \pi \right]$$
(2)

where  $\pi$  is the relative level of antibodies produced in the long-term plateau, ranging from 0 to 1. Parameter  $\pi$  is an indicator of long-term antibody persistence. Thus, a long-term seroprotective effect could be tested based on interval analysis of  $\pi$ .

In both models, k and a are assigned as random effects and follow a bivariate normal distribution. The parameter  $\pi$  in the modified power-law model is assigned as a fixed effect. In this way, the participant-specific time-antibody functions would be obtained to perform predictions of antibody dynamics for each participant. To analysis the influence of age at vaccination and gender, supportive analyses were conducted: in both models, age at vaccination and gender were included as covariates of the peak antibody titer (k) and the decay rate (a).

Models were fitted independently for each vaccine. Model parameters were estimated through dual Quasi-Newton algorithms using the SAS nlmixed procedure. Goodness of fit was evaluated by Akaike's Information Criterion (AIC), Schwarz's Bayesian Information Criterion (BIC), and the adjusted coefficient of determination ( $R_{adj}^2$ ), which is given by the following:

$$R_{adj}^{2} = 1 - \frac{\sum_{j=1}^{m} \left[ \sum_{i=1}^{n_{j}} \left( y_{ij} - \hat{y}_{ij} \right)^{2} / (n_{j} - p - 1) \right]}{\sum_{j=1}^{m} \left[ \sum_{i=1}^{n_{j}} \left( y_{ij} - \bar{y}_{j} \right)^{2} / (n_{j} - 1) \right]}$$
(3)

where  $y_{ij}$  and  $\hat{y}_{ij}$  are the observed and predicted log10 antibody titer of *i*th participant at *j*th time point, respectively;  $\bar{y}_j$  and  $n_j$  are the mean of observed antibody titer and corresponding number of participants at *j*th time point; m is the number of time points; and *p* is the number of parameters in the prediction model. The parameter  $R_{adj}^2$  ranges from 0 to 1, where a larger value of  $R_{adj}^2$  indicates a higher agreement of observed data and predicted values. Models with lower AIC and BIC, and larger  $R_{adj}^2$  would be preferable.

Geometric mean titers (GMTs) of predicted antibody titers and corresponding 95% confidence intervals, as well as the predicted proportion of participants maintaining antibody titers above seroconversion thresholds (Anti-HAV  $\geq$  20 mIU/mL) and corresponding 95% confidence intervals (95% *CI*), were calculated.

### RESULTS

#### **Population Characteristics**

Among the 400 participants who were initially enrolled, 375 participants completed the two-dose vaccination and participated in the follow-up phase. The average age of the Healive and Havrix groups were 3.8 and 3.7 years old, respectively. The male/female ratio was 142/141 for the Healive group and 48/44 for the Havrix group. At each timepoint of the follow-up phase, the Healive group showed higher antibody titers and the differences between the two groups were statistically significant. Since the antibody titers of most subjects in both the Healive and Havrix groups are above 20 mIU/mL, the seroconversion rates of both groups were around 100%. The differences in seroconversion rate were not statistically significant<sup>[28,29]</sup>.

## Modeling the Persistence of Antibody Responses

To estimate the duration of antibody protection, two models (model 1 and model 2) considering longterm antibody responses were fitted based upon five years of antibody titers. Model parameters and fitting statistics are presented in Table 1. AIC and BIC of model 2 were slightly lower than those of model 1. The adjusted coefficients of determination of both models were very close with one another. Figure 1 shows the observational plots and curves for each model. Model 2 showed better fitting, as the curve overlaid the observed data points more closely than that of model 1.

For each vaccine, the peak antibody titers (*k*), as well as decay rates (*a*), were close between the two models. In both models, the peak antibody titers of Healive were higher than those of Havrix, while the antibody-decay rates were higher for Healive, compared with those of Havrix. The half-lives and durations of immune protection were also derived from the models. Model 1 estimated half-lives of 91.58 d and 124.21 d for Healive and Havrix, respectively. Model 2 yielded similar outcomes, as presented in Table 2. Note that in both models, the 95% confidence intervals of Healive and Havrix did not overlap, which indicated that the decay rates of the two vaccines were significantly different from one another.

Figure 2 presents the predictions of long-term antibody responses. Both models predicted life-long durations of immune protection (Table 2 and Figure 2), as the predicted geometric means of both Healive and Havrix remained larger than 20 mIU/mL at 50 years. The curves of Healive were above those of Havrix (Figure 2). Model 1 predicted that the antibody titers of both Healive and Havrix would decline rapidly over the first 10 years after two-dose

	Model parameters			Goodness of fit		
Group -	k	а	π	R <sup>2</sup>	AIC	BIC
Power-law mode	1					
Healive	3.5115	0.6211	-	0.8786	204.88	226.75
Havrix	3.1530	0.4879	-	0.9118	-66.59	-51.46
Modified power-	law model					
Healive	3.5270	0.7097	0.0190	0.8851	156.82	182.33
Havrix	3.1532	0.4890	0.0005	0.9106	-64.60	-46.95

Table 1. Model parameters and fitting statistics

vaccinations, and would then decline slowly in the following decades. Model 2 predicted similar antibody-decline curves but predicted that the antibody titers would reach a plateau level at nearly 15 years after the two-dose vaccination. The estimated long-term antibody plateaus were 63.866 mIU/mL (95% *CI*: 43.4359–84.2961 mIU/mL) for Healive and 0.7681 mIU/mL (95% *CI*: 20.3874–21.9237 mIU/mL) for Havrix.

In both models, age at vaccination and gender were not statistically significant for the peak antibody titers, as well as decay rates, for both Healive and Havrix, as presented in Supplementary Table S1 available in www.besjournal.com.

#### Seroprotection Rates Predicted by Models

Predictions of antibody titers of each individual participant were also estimated, as well as proportions of seroconversion. As shown in Table 3, the geometric means of predicted antibody titers by both models were close to the observed data over five years in both Healive and Havrix groups. Up to five years after the two-dose vaccination, the seroconversion rate of Healive and Havrix were

Power-law mode Modified power-law model Geomentric mean titre (mlU/mL) 3,000 2,000 1,000 20 mlU/mL 20 mlU/mL 0 24 36 48 601 12 24 36 1 12 48 60 Time post vaccination (months) Healive (predicted) ---- Havrix (predicted) Healive (measured) o Havrix (measured)

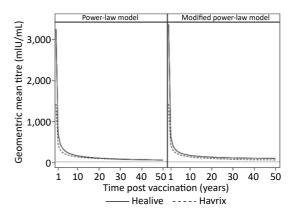
**Figure 1.** Observed and model-based prediction of geometric mean titers over time.

99.13% and 97.47%, respectively. Both models yielded similar predictions to the observed seroconversion rates (Table 3).

Table 4 presents the long-term prediction of the geometric mean of antibody titers and the proportion of seroconversion based on the estimated participant-specific functions. Model 1 predicted that at 30 years, more than 90% of participants would have seroconversion (anti-HAV  $\geq$  20 mIU/mL). In model 2, which showed better fitting, the predicted seroconversion rate of Healive remained above 95% for at least up to 35 years (Table 4 and Figure 3).

#### DISCUSSION

Clinical evidence on the duration of HAV vaccine protection has been rare due to difficulties in obtaining decade-long follow-ups after patients have received vaccines. Therefore, modeling methods have been the primary approach for investigating long-term immune persistence<sup>[20-24,31]</sup>. In the present study, we used the power-law model and a modified power-law model to investigate the long-term



**Figure 2.** Model-based prediction of geometric mean titers for up to 50 years.

<b>6</b>	Half-life of antibody decay (d)		Duration of immune protection (years)	
Group —	Estimate	95% CI	Estimate	95% Cl
Power-law model				
Healive	91.58	87.06-96.11	301.92	208.66-395.18
Havrix	124.21	107.03-141.40	521.24	128.20-914.28
Aodified power-law mo	del			
Healive	81.89	76.82-86.96	-	-
Havrix	123.95	105.31-142.59	553.41	-471.41-1578.24

Table 2. Model-based estimation of half-lives of antibody decays and durations of immune protection

antibody responses to Healive, an inactivated preservative-free HAV vaccine, among Chinese children. Both models were well-fitted with five-year follow-up data, as indicated by fitting statistics (AIC, BIC, and  $R_{adi}^2$ ).

Based on statistical modeling on the five-year follow-up data after the original trial, we projected a duration of antibody protection of at least 30 years for more than 90% of vaccinated individuals. Moreover, both models predicted that Healive would induce stronger immunogenicity compared with that of Havrix, as the peak of antibody response induced by Healive was larger than that of Havrix. The antibody-decay rate of Healive was also larger than that of Havrix. Over a life-long period, participants vaccinated with Healive were predicted to have close but slightly higher antibody titers than those of individuals vaccine with Havrix. The modified power-law model indicated that both Healive and Havrix may induce memory B-cells and would lead to a plateau of antibody titers, which may maintain a life-long period of immune protection. The predicted plateau-antibody titer of Healive was 63.866 mIU/mL (P < 0.001), which was larger than the threshold of seroconversion (20 mIU/mL) and indicated that Healive may trigger a protective anamnestic response.

**Table 3.** Observed and predicted geometric mean titers and seroconversion rates forup to five-years post vaccination

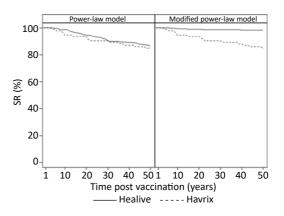
	GMT, mlU	/mL (95% <i>Cl</i> )	SR, % (95% <i>Cl</i> )		
Models	Healive	Havrix	Healive	Havrix	
1 month					
Observed	3427.18 (3036.53–3868.10)	1441.88 (1191.08–1745.49)	100.00 (98.69–100.00)	100.00 (96.07–100.00)	
Power-law model	3247.03 (2949.59–3574.46)	1422.40 (1206.28–1677.24)	100.00 (98.70–100.00)	100.00 (96.07–100.00)	
Modified power-law model	3300.66 (2984.44–3650.39)	1422.53 (1206.36–1677.44)	100.00 (98.70–100.00)	100.00 (96.07–100.00)	
1 year					
Observed	571.95 (519.59–629.59)	385.98 (323.68–460.28)	100.00 (98.57–100.00)	100.00 (95.85–100.00)	
Power-law model	693.80 (633.04–760.38)	423.20 (359.94–497.58)	100.00 (98.70–100.00)	100.00 (96.07–100.00)	
Modified power-law model	650.78 (592.92–714.29)	422.87 (359.64–497.21)	100.00 (98.70–100.00)	100.00 (96.07–100.00)	
2 years					
Observed	468.66 (423.21–518.99)	300.12 (250.21–360.00)	100.00 (98.51–100.00)	100.00 (95.55–100.00)	
Power-law model	451.10 (410.20–496.07)	301.78 (254.31–358.09)	100.00 (98.70–100.00)	100.00 (96.07–100.00)	
Modified power-law model	435.10 (395.28–478.95)	301.64 (254.19–357.95)	100.00 (98.70–100.00)	100.00 (96.07–100.00)	
3 years					
Observed	341.16 (298.91–389.38)	234.28 (187.20–293.21)	99.59 (97.73–99.99)	100.00 (95.65–100.00)	
Power-law model	350.67 (317.98–386.73)	247.61 (207.20–295.91)	100.00 (98.70–100.00)	100.00 (96.07–100.00)	
Modified power-law model	350.32 (317.70–386.28)	247.61 (207.20–295.91)	100.00 (98.70–100.00)	100.00 (96.07–100.00)	
4 years					
Observed	322.26 (284.35–365.23)	222.26 (179.06–275.90)	100.00 (98.40–100.00)	100.00 (95.32–100.00)	
Power-law model	293.30 (265.33–324.21)	215.19 (179.05–258.62)	100.00 (98.70–100.00)	98.91 (94.09–99.97)	
Modified power-law model	303.34 (274.80–334.84)	215.29 (179.14–258.73)	100.00 (98.70–100.00)	98.91 (94.09–99.97)	
5 years					
Observed	257.14 (226.89–291.42)	168.12 (135.64–208.38)	99.13 (96.89–99.89)	97.47 (91.15–99.69)	
Power-law model	255.34 (230.53–282.82)	193.00 (159.83–233.05)	99.65 (98.05–99.99)	98.91 (94.09–99.97)	
Modified power-law model	272.91 (247.06–301.48)	193.17 (159.99–233.24)	100.00 (98.70–100.00)	98.91 (94.09–99.97)	

*Note.* Predictions are based on the estimated participant-specific functions. GMT: geometric mean titer; SR: seroconversion rate.

follow-up studies of immune Long-term persistence have revealed that most vaccines remain seropositive for nearly 20 years. The predictions in the present work are in line with the outcomes of these previous studies. Van Herck et al.<sup>[20]</sup> evaluated antibody persistence using data from two of the longest documented follow-up studies and showed that 100% of vaccinated individuals remained seropositive for up to 17 years following a 0, 6 months vaccination schedule. A 22-year follow-up study carried out by Mosites et al.<sup>[15]</sup> found that 15 out of 16 vaccinated individuals maintained protective levels of anti-HAV following a 0, 6 months vaccination schedule. López et al.<sup>[23,32]</sup> investigated long-term persistence of anti-HAV in children less than 15 years old and showed that 100% of vaccinated individuals maintained seroconversion for at 15 years after vaccination.

The advantage of the power-law model is that the kinetics of B-cell turnover is accounted for and may lead to a reasonable interpretation for the longterm duration of antibody persistence<sup>[33,34]</sup>. The modified power-law model further considers a memory-B-cell subpopulation and an antibody plateau<sup>[24]</sup>, which are important in modeling the long-term antibody memory. The suitability of the modified power-law model is supported by the longterm persistence of HAV-specific memory B cells, which has been verified in a cohort study<sup>[11]</sup>. Additionally, inter-subject variabilities are accounted for in both models.

In this study, antibody titers above 20 mIU/mL were considered as seroconversion for conservative concern, although the detection limit was 5 mIU/mL. The cut-off value of 20 mIU/mL was referred from the minimal detectable concentrations in previous studies<sup>[19,35]</sup>. The minimal detectable concentrations were generally derived from animal studies. Lower cut-off value for seroconversion, such as 10 mIU/mL and 15 mIU/mL, are used in some studies in which



**Figure 3.** Predicted seroconversion rate based on the estimated participant-specific functions.

Maara	GMT, mlU	GMT, mIU/mL (95% <i>CI</i> )		SR, % (95% <i>Cl</i> )	
Years -	Healive	Havrix	Healive	Havrix	
ower-law	model				
10	166.02 (148.82–185.20)	137.62 (112.12–168.92)	98.59 (96.42–99.61)	94.57 (87.77–98.21	
15	129.06 (115.14–144.66)	112.92 (91.03–140.08)	96.47 (93.60–98.29)	93.48 (86.34–97.57	
20	107.94 (95.96–121.42)	98.14 (78.49–122.70)	94.70 (91.41–97.00)	93.48 (86.34–97.57	
25	93.97 (83.30–106.01)	88.01 (69.96–110.74)	92.93 (89.30–95.63)	90.22 (82.24–95.43	
30	83.91 (74.20–94.89)	80.52 (63.67–101.84)	90.46 (86.42–93.62)	90.22 (82.24–95.43	
35	76.25 (67.29–86.41)	74.69 (58.79–94.89)	89.75 (85.62–93.03)	89.13 (80.92–94.66	
Aodified p	ower-law model				
10	203.48 (183.91–225.14)	138.04 (112.51–169.37)	99.29 (97.47–99.91)	94.57 (87.77–98.21	
15	175.60 (158.62–194.40)	113.49 (91.55–140.68)	98.94 (96.93–99.78)	93.48 (86.34–97.57	
20	159.84 (144.35–176.99)	98.81 (79.11–123.41)	98.94 (96.93–99.78)	93.48 (86.34–97.57	
25	149.45 (134.94–165.51)	88.77 (70.65–111.54)	98.59 (96.42–99.61)	90.22 (82.24–95.43	
30	141.96 (128.17–157.23)	81.35 (64.42–102.72)	98.59 (96.42–99.61)	90.22 (82.24–95.43	
35	136.25 (123.01–150.92)	75.57 (59.59–95.83)	98.59 (96.42–99.61)	89.13 (80.92–94.66	

**Table 4.** Predicted geometric mean titers and seroconversion rates based on the estimated participant-specific functions

*Note.* GMT: geometric mean titer; SR: seroconversion rate.

more sensitive assays were applied<sup>[18,36,37]</sup>.

The primary limitation of the present study is in its absence of longer follow-up data. A longer followup period would increase the precision of prediction for long-term immune persistence. Additionally, models considering immunosenescence<sup>[25]</sup> could be applied if long-term data are available.

In conclusion, our modeling study suggests that Healive may adequately induce long-term immune responses following a 0, 6 months vaccination schedule in children. Additionally, our model suggests that memory B cells may be induced to provide stable and durable immune protection. Finally, based on statistical modeling on the five-year follow-up data, we could predict that more than 90% of participants would maintain seroconversion for at least 30 years.

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#### AUTHOR CONTRIBUTIONS

YPY performed the data analysis and drafted the manuscript. JTC conceived of research questions and revised the manuscript. ZWJ, LW, and CKY involved in data analysis and commented the manuscript. XYY and CY critically commented and revised the manuscript. JLX revised the manuscript. All authors read and approved the final version of the manuscript.

#### **CONFLICTS OF INTERESTS**

JTC is an employee of Sinovac Biotech. The other authors declare no conflicts of interest.

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	Power-la	aw model	Modified power-law model		
Parameter	Healive	Havrix	Healive	Havrix	
k	3.4625 (3.2697, 3.2697)	3.2136 (2.9105, 2.9105)	3.4614 (3.2612, 3.2612)	3.2138 (2.9105, 2.9105)	
Gender	0.0227 (-0.0751, -0.0751)	0.0010 (-0.1591, -0.1591)	0.0234 (-0.0784, -0.0784)	0.0010 (-0.1592, -0.1592)	
Age	0.0040 (-0.0265, -0.0265)	-0.0168 (-0.0654, -0.0654)	0.0081 (-0.0237, -0.0237)	-0.0168 (-0.0654, -0.0654)	
а	0.6112 (0.5031, 0.5031)	0.6500 (0.4739, 0.4739)	0.6783 (0.5294, 0.5294)	0.6524 (0.4680, 0.4680)	
Gender	-0.0006 (-0.0556, -0.0556)	-0.0543 (-0.1475, -0.1475)	0.0026 (-0.0726, -0.0726)	-0.0546 (-0.1487, -0.1487)	
Age	0.0029 (-0.0144, -0.0144)	-0.0222 (-0.0506, -0.0506)	0.0075 (-0.0162, -0.0162)	-0.0224 (-0.0510, -0.0510)	

Supplementary Table S1. Model parameters of age at vaccination and gender