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Prediction Model with Validation for Polio-seronegativity in Malnourished Children from Poliomyelitis Transmission High-Risk Area of the Democratic Republic of the Congo (DRC)

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Background: Malnutrition is identified as a risk factor for insufficient polio seroconversion in the context of a vaccine-derived poliovirus (VDPV) outbreak-prone region. In the Democratic Republic of Congo (DRC), underweight decreased from 31% (in 2001) to 26% (in 2018). Since 2004, VDPV serotype 2 outbreaks (cVDPV2) have been documented and were geographically limited around the Haut-Lomami and Tanganyika Provinces.

Methods: To develop and validate a predictive model for poliomyelitis vaccine response in malnourished infants, a cross-sectional household study was carried out in the Haut-Lomami and Tanganyika provinces. Healthy children aged 6 to 59 months (n=968) were enrolled from eight health zones (HZ) out of 27, in March 2018. We performed a bivariate and multivariate logistics analysis. Final models were selected using a stepwise Wald method, and variables were selected based on the criterion $p < 0.05$. The association between nutritional variables, explaining polio seronegativity for the three serotypes, was assessed using the receiver operating characteristic curve (ROC curve).

Results: Factors significantly associated with seronegativity to the three polio serotypes were underweight, non-administration of vitamin A, and the age group of 12 to 59 months. The sensitivity was 10.5%, and its specificity was 96.4% while the positive predictive values (PPV) and negative (PNV) were 62.7% and 65.3%, respectively. We found a convergence of the curves of the initial sample and two split samples. Based on the comparison of the overlapping confidence intervals of the ROC curve, we concluded that our prediction model is valid.

Conclusion: This study proposed the first tool which variables are easy to collect by any health worker in charge of vaccination or in charge of nutrition. It will bring on top, the collaboration between the Immunization and the Nutritional programs in DRC integration policy, and its replicability in other low- and middle-income countries with endemic poliovirus.

Keywords: prediction model, validation, child health, child nutrition, underweight, serotypes, polio serotype, seronegativity, poliomyelitis-neutralizing antibodies, malnutrition, DRC

Introduction

Malnutrition, including micronutrient insufficiency, like copper, iodine, vitamin A, zinc, iron, and selenium,^{1,2} is a major² global health problem, not just in low- and middle-income countries (LMIC).^{3,4} Additionally, malnutrition can act as an aggravating factor for other illnesses.⁵ The food security report reveals that, in 2022, about 26.4 million people living in the DRC were in severe food insecurity. Root causes have been attributed to generalized poverty, conflicts and population displacement, insecurity, low agricultural production, material cost inflation, and the lack of basic infrastructure.^{6,7} In DRC, the national prevalence of underweight decreased from 31% to 26% in rural and urban areas from 2001 to 2018.⁸

Malnutrition is responsible, both directly and indirectly, for 54% of the 10.8 million deaths per year and contributes to half of infectious disease-related deaths among children under five LMICs. Previous studies have shown that several immune mechanisms may be defective in malnourished infants suffering from severe infections and constitute the vicious infection-malnutrition cycle.⁹

Poliomyelitis is an infectious disease of viral origin that can lead to paralysis. It is caused by one of three polio serotypes (types 1, type 2, and type 3) and mainly affects children less than five years old.¹⁰ Only one out of 200 polio infections experience paralytic symptoms or Acute Flaccid Paralysis (AFP) cases.¹¹ Since August 2020, WHO certified the African Region free from all three types of wild poliovirus (WPV) when Nigeria became the last African country to interrupt wild poliovirus transmission.¹² The DRC has been certified polio-free since 2012, however, since 2004, it has documented continued transmission¹³ circulating derived poliovirus type 2 (cVDPV2) and type 1 (cVDPV1) cases.^{14,15} Since May 2017, there has been an ongoing outbreak of VDPVs in the DRC. In 2021, 28 AFP cases were confirmed for cVDPV21, and in 2022, 377 cVDPV2 cases and 143 cVDPV1 cases were confirmed as well as 11 samples with cVDPV2 from environmental surveillance.¹⁶

In 2015, the Global Polio Eradication Initiative (GPEI) advocated for the introduction of an Inactivated Polio Vaccine (IPV) worldwide in all countries that had not yet introduced it. On 24 April 30th, 2016, the DRC along with every other country participated in the worldwide switch from the trivalent OPV (tOPV) to the bivalent OPV (bOPV), wherein type 2 was taken out of the tOPV. Multiple reasons are attributed to this switch including that PV2 was declared eradicated in September 2015 and studies suggest that bOPV led to significantly higher immunogenicity for type 1 and type 3 than tOPV.^{17,18}

The 2017–2018 Multiple Indicator Cluster Surveys (MICS) revealed that only 35.0% of children 12–23 months were fully vaccinated in the DRC. However, there was provincial-level variation, and Haut Lomami and Tanganyika were estimated to have vaccine coverage estimates for children 12–23 months of 35.7% and 21.2%, respectively. The primary reasons for not being fully vaccinated included lack of knowledge among mothers and caregivers (29.0%), lack of time (18.4%), mistrust (16.1%), and challenges related to service delivery (28.7%). These include vaccinator unavailability, vaccine stock-outs, long waiting times, and financial barriers. This MICS revealed poor nutritional status. In the Haut-Lomami province, the *Z* scores were successively -1.2 for the weight/age ratio, -1.8 for the height/age ratio, -0.2 for the weight/height ratio. In Tanganyika province, the *Z*-score scores were successively -0.9 for weight-for-age, -1.6 for height-for-age, 0.1 for weight for size.¹⁹

Serological studies on polio vaccinations have been recommended by the WHO to assess and orient polio vaccination activities,²⁰ evaluate the risk of poliovirus outbreaks, and identify immunity issues in the population. They have been performed in DRC,²¹ Nigeria,²² India,²³ Pakistan,²⁴ West Africa,²⁵ and Madagascar²⁶ and they contribute in poliomyelitis eradication efforts in these countries. Additionally, low immunogenicity has been identified in severe wasting cases, but this does not necessarily explain the reduced effectiveness of the vaccine against vaccine-preventable diseases, especially in countries where malnutrition is prevalent.^{27,28}

To develop a predictive model for poliomyelitis vaccine response in malnourished children in VDPV outbreak-prone areas, the cross-sectional study was conducted in two DRC south-eastern provinces.

Methods

Study Population

This was a community-based, cross-sectional survey carried out in 2018 in four of the 16 Hz in Haut-Lomami Province: Butumba, Lwamba, Malemba-Nkulu, and Mukanga and four of the 11 Hz in Tanganyika Province: Ankoro, Kabalo, Kongolo, and Manono.²⁹

Study procedures have been previously described by Halbrook, 2020,26. Children were selected using a three-stage cluster sampling technique. During the first stage, eight health zones were selected based on the number of cVDPV2 registered and mass campaign responses organized. Within each health zone, five villages were selected by stratified random sampling using settlement feature layers derived from satellite imagery. During the second stage, we selected villages using a GIS point methodology with buffers. The clusters (health areas) were randomly selected using ArcGIS software based on two parameters: 1) not being in the same health area and 2) being separated by at least 500 meters. Households in which children aged 6 to 59 months reside formed the subgroups of these clusters. Overall, 327 households in Haut-Lomami and 641 in Tanganyika were investigated by simple random sampling. All households in the cluster were given the opportunity to be surveyed until the expected sample size was reached. Households that refused to participate were marked as “refused” in the tablet-based questionnaire.

Data Collection and Study Variables

Informed consent and a questionnaire were administered orally in the participants’ preferred language (French or Swahili) by trained interviewers. For selected households, we used the concept “healthy child” to mean a child who does not present the specific signs of serious illness (such as lethargy, unconsciousness, and convulsion),² disability, cough or difficulty breathing, dehydration or persistent diarrhea, fever, edema of both feet, palmar pallor at the assessment time. Nine hundred and sixty-eight children fit this operational definition.

Consenting parents or guardians of children were administered a questionnaire designed to collect basic demographic data, as well as data on the participants’ work practices and health and child immunization data. A tablet-based questionnaire was used to collect data on current health status, anthropometric measures (height, weight), and other behaviors, including knowledge of vaccinations and utilization of the routine immunization system to receive vaccinations, which could be associated with the diseases.² Height, weight, and mid-upper arm circumference (MUAC) were measured using a wooden infant-cum-stadiometer, SECA 874 digital weighing scales, and tricolor MUAC tapes, respectively. Standard methods were followed to take weight, height/length, and MUAC as recommended by the WHO. Additionally, a dried blood spot (DBS) sample was obtained through a finger prick. Questionnaires and collected DBS specimens were assigned linking barcode numbers to facilitate data analysis.

The following quantitative and qualitative variables were used: age, sex, marital status, tribe, level of education, religion, main occupation, nutritional status, and seroprevalence. Household density was estimated at six per household based on estimates from previous studies.⁴ We generated z-scores using ENA (Emergency Nutrition Assessment) software for SMART (Standardized Monitoring and Assessment of Relief and Transitions). Anthropometric index based on the WHO 2006 standards (stunting, wasting, and undernutrition) were evaluated using ENA software (July 2015 version Manufactured by Action Against Hunger Canada). Using WHO classification of nutritional status of infants and children. Underweight, wasting, and stunting were respectively defined as weight-for-age Z-score ≤ -2 , weight-for-height Z-score ≤ -2 , and height-for-age Z-score ≤ -2 standard deviations of the WHO Child Growth Standards median.

Laboratory Analysis

Testing for neutralizing antibodies against poliovirus types 1(PV1), 2(PV2), and 3(PV3) was conducted at the US Centers for Disease Control and Prevention (CDC), Atlanta, GA. A modified poliovirus microneutralization assay was used to measure the ability of antibodies in serum or eluted from DBS punches to block the infectivity of poliovirus in an in vitro cell culture system. Following collection and in-country processing, the DBS were shipped at ambient temperature and stored at -20°C . The testing process has been previously described elsewhere. In brief, equivalent to approximately 60 ul of sera were collected from each card and processed for the low-volume poliovirus neutralization assay. A series of dilutions of DBS elute were combined with a fixed amount of virus before inoculation of poliovirus-susceptible cells. After five days of incubation, a luminescent cell viability reagent was added to detect live cells. The presence of live cells indicates protection from the virus cytopathic effect, which is the neutralization of virus infectivity. Neutralization titers are reported in a log₂ format, with 2.5 log₂ as the lower limit of detection and 10.5 log₂ as the upper limit of detection. Neutralizing antibodies have been evaluated against PV1, PV2, and PV3, and titers ≥ 3.0 log₂ are considered evidence of seroprotection.^{3,6}

Statistical Analysis

Data from this study were analyzed using SPSS 23 and STATA 14.0 software. After calculating the prevalence of malnutrition, the different frequencies were compared using the Pearson Chi-square test, and a p-value <0.05 was considered statistically significant. Univariate analyses were expressed as frequency distributions and percentages as appropriate. In the bivariate analysis, the chi-square test was used to assess the link between nutritional status and seronegativity.

Logistic regression using the stepwise Wald method was used to select final variables to be included in the model. For model validation, we used the split-sample model validation method. We randomly selected 75% of nutritional factors explaining seronegativity to the different polio serotypes instead of selecting all the factors found in the initial model.

Discrimination between nutritional variables explaining polio seronegativity to the three polio serotypes was assessed using the ROC curve. We determined sensitivity, specificity, positive and negative predictive values. The performance measure of this logistic model is numerically equivalent to the area under the ROC curve.

Ethical Considerations

Ethical clearance was obtained from the ethics committee of UCLA Institutional Review Board (IRB#18-000303) and the Kinshasa School of Public Health (approval letter No: ESP/CE/164/2021), University of Kinshasa, DRC.

Our study complies with the Declaration of Helsinki and verbal informed consent was acceptable and approved by the ethics committees.

Results

Bivariate Analysis of Seronegativity to the Three Polio Serotypes and Nutritional Status of the Children Aged 6 to 59 from Poliomyelitis Transmission High-Risk Area of the DRC

An association between underweight ($p=0.003$) and non-administration of vitamin A ($p<0.001$) was observed significantly with seronegativity to the three polio serotypes. While no significant association was found between seronegativity to the three polio serotypes with chronic malnutrition ($p=0.97$) and acute malnutrition ($p=0.11$) (Table 1).

Table 1 Bivariate Analysis of Seronegativity to the Three Polio Serotypes and Nutritional Status of the Children Aged 6 to 59 from Poliomyelitis Transmission High-Risk Area of the DRC (n=968)

Nutritional Status	Seronegativity to the Three Polio Serotypes	
	n (%)	p-value
Underweight		
Yes	133 (55.6)	0.003*
No	483 (66.3)	
Chronic malnutrition		
Yes	193 (36.4)	0.97
No	153 (34.9)	
Acute malnutrition		
Yes	68 (45.6)	0.11
No	284 (34.7)	
Reception of Vit A		
Yes	139 (28.3)	<0.001*
No	213 (44.7)	

Notes: *Pearson's chi-square test at significance level $p<0.05$.

Logistic Regression Model of Nutritional Variables Explaining Seronegativity to the Three Polio Serotypes

After logistic regression, three variables stand out as the predictors of seronegativity to the three polio serotypes: not receiving Vitamin A, being underweight and the age group of 12 to 59 months (Table 2). Thus, the model for predicting seronegativity to the three polio serotypes can be written as below:

$$P = (Y = \text{Seronegativity to the three polio serotypes} / X = xi) = \frac{e^{-0,38+0,63.x1+0,41x2+0,51.x3}}{1 + e^{-0,38+0,63.x1+0,41x2+0,51.x3}}$$

The predictive value of seronegativity to the three polio serotypes was extracted from the matrix of the regression prediction model (Table 3). The value of the area under the ROC curve is 0.627 (Figure 1), which indicates a discrimination of its ability to predict the onset of seronegativity to the three polio serotypes.

After internal validation of this model, three variables stand out as the predictors of seronegativity to the three polio serotypes: not receiving Vitamin A, being underweight and the age group of 12 to 59 months (Table 4). Thus, the model for predicting seronegativity to the three polio serotypes can be written below:

$$P = (Y = \text{Seronegativity to the three polio serotypes} / X = xi) = \frac{e^{-0,49+0,62.x1+0,53x2+0,49.x3}}{1 + e^{-0,49+0,62.x1+0,53x2+0,49.x3}}$$

The value of the area under the curve is 0.627 (Figure 2), which indicates the ability of the split model discrimination to predict the onset of seronegativity to the three polio serotypes. Comparing the ROC curves of the initial sample and the split samples, the value of the area under the two curves is 0.627 (Figure 3).

Discussion

This study suggests that being underweight, non-administration of vitamin A, and age group of 12 to 59 months are the powerful negative predictors of seroconversion to the three polio serotypes. These results are supported by Carol, Chandir and the Stoffel team in underweight children.^{30–32}

Table 2 Logistic Regression Model of Nutritional Variables Explaining Seronegativity to the Three Polio Serotypes

Factors	A	ES	Wald	p	EXP (B)	CI for Exp(B) 95%	
						Inferior	Superior
Have not received Vit A (No vs Yes) (x1)	0.633	0.139	20.726	0.000	1.882	1.434	2.472
Underweight (Yes vs No) (x2)	0.410	0.156	6.964	0.008	1.507	1.111	2.045
Age range (12–59 months vs 6–11 months) (x3)	0.509	0.144	12.394	0.000	1.663	1.253	2.208
Constant	–0.384	0.167	5.269	0.022	0.681		

Abbreviations: A, regression coefficient; ES, standard error of the regression coefficient; Wald, Wald test; p, adjusted p-value; Exp(B), Adjusted odds ratio; CI, Confidence interval of Exp(B).

Table 3 Logistic Regression Prediction Model Confusion Matrix

	Seronegativity to the Three Expected Polio serotypes	Seropositivity to the Three Expected Polio serotypes	Total	
Seronegativity to the three polio serotypes observed	37	315	352	Se=10.5%
Seropositivity to the three polio serotypes observed	22	594	616	Sp=96.4%
Total	59	909	968	
	PPV=62.7%	NPV=65.3%		

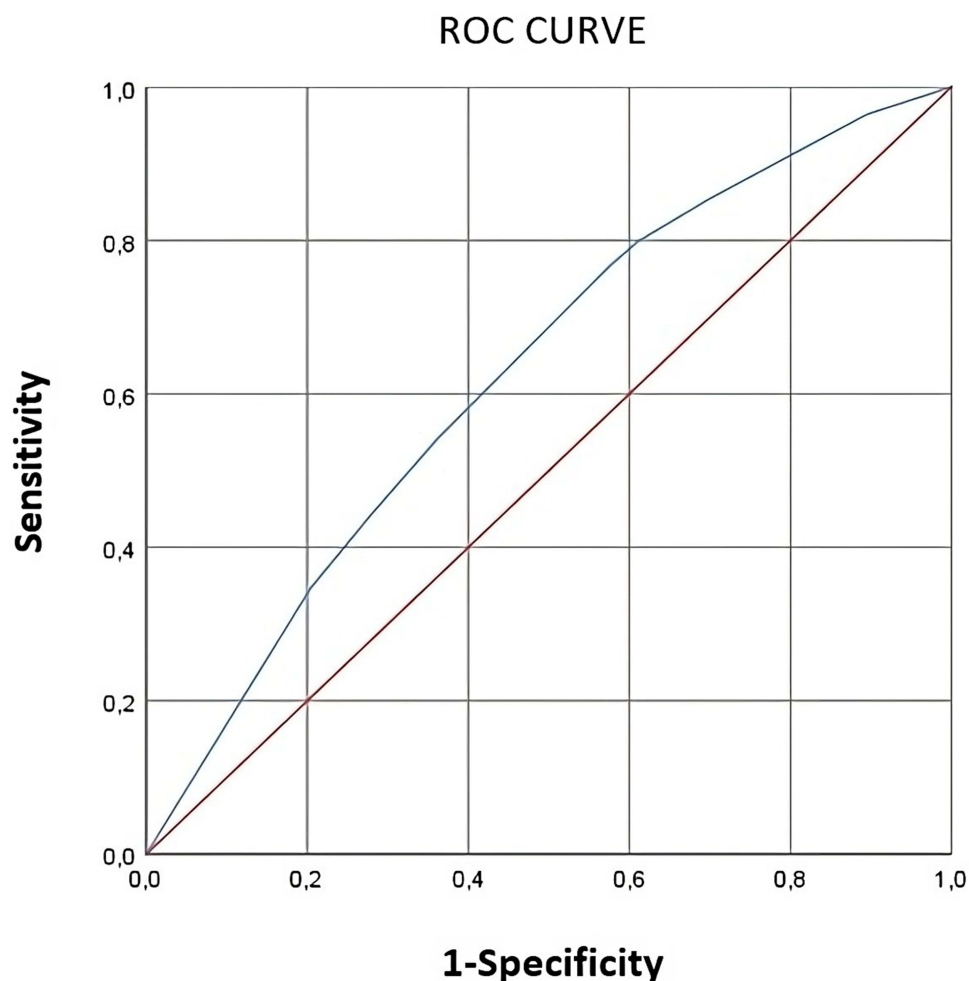


Figure 1 ROC curves of nutritional factors explaining seronegativity to the three polio serotypes in the initial model.

As for the effect of Vitamin A on immunogenicity, several studies have shown that vitamin A during immunization session improves immune responses,^{33–38} and does not alter the immunogenicity of vaccines.³⁹

We were not aware of any study on the model for predicting seronegativity in malnourished children at the time of writing this paper. Based on the comparison of the overlapping confidence intervals of the ROC curve, we concluded that our prediction model is valid.

Table 4 Logistic Regression of Nutritional Variables Explaining Seronegativity to the Three Polio Serotypes by the Split Sample Validation of the Initial Sample

Factors	A	ES	Wald	p	EXP (B)	CI for Exp(B) 95%	
						Inferior	Superior
Have received Vit A (No vs Yes) (x1)	0.619	0.159	15.233	0.000	1.858	1.361	2.536
Underweight (Yes vs No) (x2)	0.534	0.176	9.218	0.002	1.705	1.208	2.407
Age range (12–59 months vs 6–11 months) (x3)	0.492	0.165	8.831	0.003	1.635	1.182	2.261
Constant	–0.492	0.193	6.477	0.011	0.612		

Abbreviations: A, regression coefficient; ES, standard error of the regression coefficient; Wald, Wald test; p, adjusted p-value; Exp(B), Adjusted odds ratio, CI, Confidence interval of Exp(B).

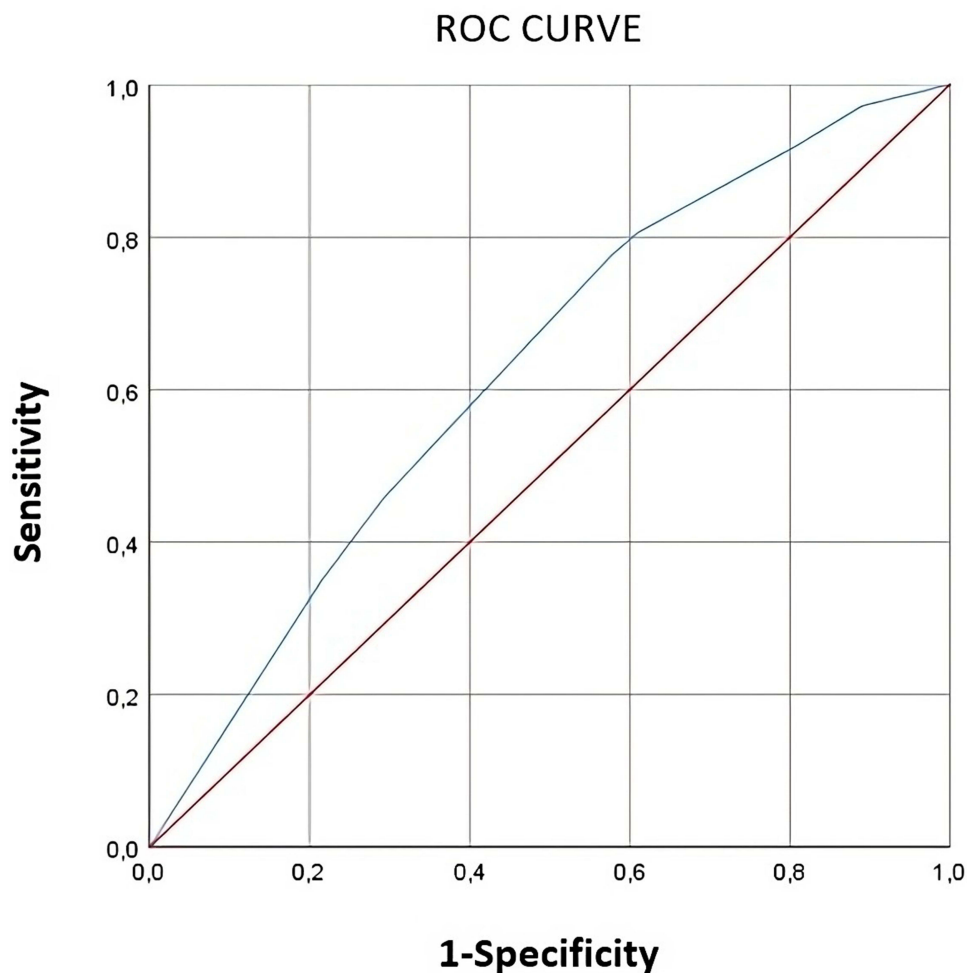


Figure 2 ROC curves of nutritional factors explaining seronegativity to the three polio serotypes in the split model.

The strength of our study is that, to our knowledge, it is the first study of the prediction model on seronegativity in malnourished children to the three polio serotypes. Steffel et al, in their study, on iron deficiency, predict seronegativity during administration of OPV and other antigens although their sample size was small compared to ours.³⁰

The other advantage is that the identification of seronegativity in a poliovirus risk environment can be achieved by taking weight for age alone during preschool visits. This model will bring together health workers who deal with immunization and nutrition in health facilities and in the community, on the ease of collecting these parameters (weight and age) during a preschool visit of routine or during vaccination campaigns by health workers. Whether in a hospital setting or in the community as supported by Olivier Mukuku and his collaborators in the prediction of the risk of severe acute malnutrition.⁴⁰ This model will bring on top the collaboration between the Immunization and the Nutritional programs in DRC integration policy and its replicability in other low- and middle-income countries with endemic polio virus.

Our survey is limited by its design, which reduced our ability to make causal inferences. Additionally, the laboratory method used did not allow us to make the difference between the presence of neutralizing antibodies due to vaccination or natural infection. Therefore, seroprevalence rates cannot be interpreted as only reflecting vaccination coverage in such cold chain challenging conditions.^{41,42} Since only 13.1% of participants had a vaccination card, did not allow us to carry out in-depth analyses. We selected children and their parents or guardians who were in the village at the moment of sample collection may have biased the study population. In addition, we had a low representation of children 35 months

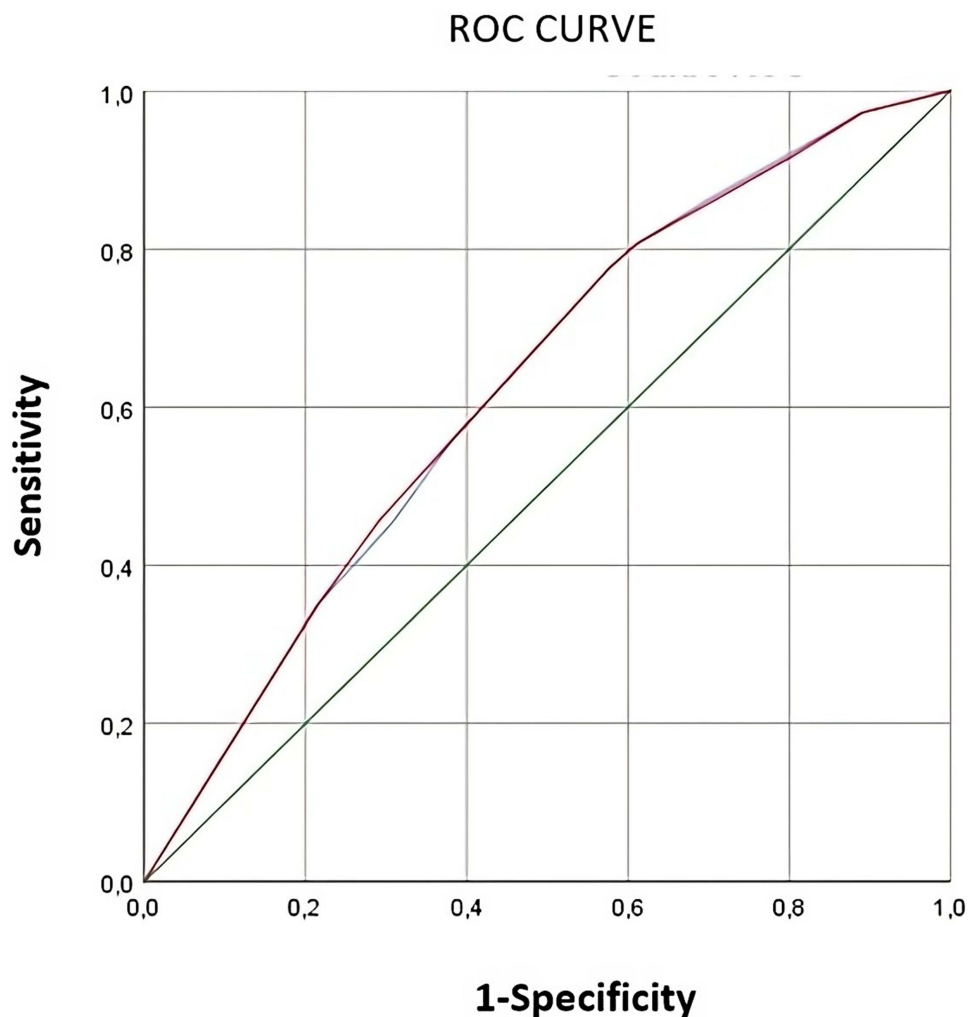


Figure 3 Comparison between ROC curve of the initial sample and the split sample.

or older, which could have lowered our estimates of immunogenicity in the study population⁴³. Data was not collected on the hemoglobin level or serum retinol level, birth site, or history of disease in the first months of life.

The lack of a model does not enable statistical comparisons to be made to assess the advantages or deficiencies of our model. However, compared to the standard proposed by J A Swets, with the area under the ROC curve between 0.5 and 0.7, the model could be considered not more informative.⁴⁴

The Clinical implications of our findings are that a large proportion of individuals with polio seronegativity also meet diagnostic criteria for at least one malnutrition disorders.^{25,42} Some existing researches suggest that malnutrition is the most common reason for clinical referral.^{45–47} It has also been shown that having malnutrition as well as polio seronegativity may cause additional burden to the individual and lead to worse functional outcomes.^{48,49}

Conclusion

This study of the predictive factors of seronegativity in malnourished children has made it possible to propose the first tool whose variables are easy to collect by any health worker in charge of vaccination or in charge of nutrition.

Not receiving Vitamin A, being underweight and the age group of 12 to 59 months stand out as the predictors of seronegativity to the three polio serotypes in this model.

Depending on the clinical implication, our results indicate that the assessment and diagnosis of malnutrition disorders should be a priority for clinicians caring for cases of acute flaccid paralysis, whether polio-related or not, because the

diagnosis of a disorder concomitant with malnutrition can lead to a treatment plan integrating several specific intervention programs.

So, this model will bring on top, the collaboration between the Nutritional and the Immunization programs in DRC integration policy and its replicability in other low- and middle-income countries with endemic polio virus. This model lends itself to being performed in a hospital setting or in the community.

Being the first model, it may have limitations. Its improvement requires applying external validation on a large sample to generalize the practice in the same settings. We also recommend randomized control trials to verify this model and overcome the current limitations.

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Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.³

All authors have read and agreed to the published version of the manuscript.

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Disclosure

The authors declare that they have no conflicts of interest.

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