# Spatial and Temporal Patterns of Malaria in Phu Yen Province, Vietnam, from 2005 to 2016

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*Abstract.* Malaria in Vietnam has become focal to a few provinces, including Phu Yen. This study aimed to assess correlations between intervention (population proportion protected by insecticide-treated nets and indoor residual spraying) and climatic variables with malaria incidence in Phu Yen Province. The Vietnam National Institute of Malariology, Parasitology, and Entomology provided incidence data for *Plasmodium falciparum* and *Plasmodium vivax* for 104 communes of Phu Yen Province from January 2005 to December 2016. A multivariable, zero-inflated Poisson regression model was developed with a conditional autoregressive prior structure to identify the underlying spatial structure of the data and quantify associations with covariates. There were a total of 2,778 *P. falciparum* and 1,770 *P. vivax* cases during the study period. *Plasmodium falciparum* and *P. vivax* incidence increased by 5.4% (95% credible interval [Crl] 5.1%, 5.7%) and 3.2% (95% Crl 2.9%, 3.5%) for a 10-mm increase in precipitation without lag, respectively. *Plasmodium falciparum* and *P. vivax* incidence decreased by 7.7% (95% Crl 5.6%, 9.7%) and 10.5% (95% Crl 8.3%, 12.6%) for a 1°C increase in minimum temperature without lag, respectively. There was a > 95% probability of a higher than provincial average trend of *P. falciparum* and *P. vivax* in Song Cau and Song Hoa districts. There was a > 95% probability of a lower than provincial average trend in Tuy Dong Xuan and Hoa districts for both species. Targeted distribution of resources, including intensified interventions, in this part of the province will be required for local malaria elimination.

## INTRODUCTION

Vietnam has made tremendous progress in reducing mortality and morbidity associated with malaria in recent years.<sup>1–4</sup> A successful ramping-up of interventions including improvements in early and accurate diagnosis, free access to treatment with artemisinin-based combination therapies (ACTs), widespread and routine distribution of insecticide-treated mosquito nets (ITNs), and targeted and reactive indoor residual spraying (IRS) has seen a reduction in malaria cases and deaths by 97% and 99.8%, respectively, between 1991 and 2014.<sup>3,5,6</sup> As a result of the significant reduction in malaria incidence, the Vietnam National Institute of Malariology, Parasitology, and Entomology (NIMPE) is pursuing an agenda of progressive elimination with a goal to eliminate local transmission by 2030.<sup>1–4,7</sup>

Since 1991, malaria control in Vietnam has been based on free early diagnosis and treatment with ACT, vector control through the free distribution of ITNs/long-lasting insecticidal nets (LLINs), and IRS.<sup>8</sup> Quinine and chloroquine were the main treatments for *Plasmodium falciparum* and *Plasmodium vivax* until 1991. Between 1992 and 1994, artemisinin derivatives were introduced in all districts. In 1999–2000, a fixed combination of dihydroartemisinin, piperaquine, trimethoprim, and primaquine became the first-line treatment.<sup>9</sup>

Vector control in Vietnam underwent many changes. In 1992 and 1993, dichlorodiphenyltrichloroethane was used for IRS. Because of the rapid decline of the malaria incidence after the introduction of ITNs, IRS was largely abandoned after 1995.<sup>8</sup> Since 2009, ITNs have been progressively replaced by LLINs as funds have become available through the Global Fund to Fight AIDS, Tuberculosis, and Malaria. Supplemental single LLINs or long-lasting insecticide-treated hammock nets are also now provided to mobile and migrant populations and forest-goers.  $^{10}\,$ 

In recent years, malaria has become more geographically confined to provinces in Central and Central-Southern Vietnam, including Phu Yen Province.<sup>11</sup> In these areas, surges in cases have been attributed to a number of factors including the presence of exophagic and anthropophilic vectors (*Anopheles dirus*),<sup>12,13</sup> barriers to control activities due to remote mountainous and forested areas,<sup>2</sup> forest-related economic activities,<sup>14–16</sup> and poverty.<sup>14,17</sup> Furthermore, the spread of artemisinin-resistant *P. falciparum* in the Greater Mekong Subregion (GMS) poses a serious threat to malaria elimination in Vietnam.<sup>18–22</sup>

The aims of this study were to identify malaria clusters by species in Phu Yen Province at the commune level and assess correlations between intervention-related variables (proportion of the population protected by ITNs and IRS) and environmental variables, with malaria incidence at the commune level. The findings from this study can be used for focused interventions of malaria by the malaria program officials of Phu Yen and by malaria elimination countries.

## MATERIALS AND METHODS

**Study sites and data sources.** Phu Yen is located in the South Central Coastal region of Vietnam. Phu Yen is administratively divided into nine districts and 104 communes (Figure 1). The total population of Phu Yen in 2016 was 875,387. Numbers of reported *P. falciparum* and *P. vivax* cases by commune and by month from January 2005 to December 2016 and ITN/IRS data were provided by the NIMPE. Commune-level population data were provided by Phu Yen provincial council. Commune population was imputed by month as follows: the difference in the district population in 2004 and 2005 was calculated and then divided by 12 to allow for a monthly population increase in 2005. A similar approach was used to calculate the monthly population of the rest of the study period (2006–2016). High-resolution (1 km<sup>2</sup> [30 arc-

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FIGURE 1. Map of Phu Yen Province in Vietnam with districts and communes. This figure appears in color at www.ajtmh.org.

seconds]) raster maps of interpolated long-term (1950–2000) average monthly precipitation and temperature were obtained from the WORLDCLIM website.<sup>23</sup> Precipitation and temperature maps were imported into a geographical information system (GIS) (GIS; ArcMap version 10.5, ESRI, Redlands, CA) and linked spatially to a digitized boundary map of the 104 communes of Phu Yen Province. The monthly mean of precipitation and temperature were extracted for each study

commune using Zonal Statistics functions in ArcMap (ESRI, Redlands, CA).

**Exploration of seasonal patterns and temporal trends.** The monthly malaria incidences by *Plasmodium* species were calculated for the full time series (January 2005–December 2016). The time series of malaria incidence was decomposed using seasonal-trend decomposition based on locally weighted regression to show the seasonal pattern, the temporal trend, and the residual variability. The time series data, the seasonal component, the trend component, and the remainder component are denoted by  $Y_t$ ,  $S_t$ ,  $T_t$ , and  $R_t$ , respectively, for month t = 1 to N, and

$$Y_t = S_t + T_t + R_t$$

The parameter setting "periodic" was used for the seasonal extraction, and all other parameters were by default. In the study, logarithmic transformations were used for the time series data.<sup>24,25</sup>

**Crude standardized morbidity ratios (SMRs).** Crude SMRs for each commune were calculated by

$$Y_i = \frac{O_i}{E_i}$$

where *Y* is the SMR in commune *i*, *O* is the observed number of malaria cases in the commune *i*, and *E* is the expected number of malaria cases in the commune *i*, across the study period. The expected number of malaria case was calculated by multiplying the provincial malaria incidence by the average population for each commune over the entire study period.

**Independent variable selection.** Initially, a preliminary Poisson regression of total malaria cases was undertaken to select the significant covariates; of these, the best fit covariates were selected with the lowest Akaike's information criterion (AIC). Climatic variables, namely, precipitation and minimum temperature (°C) without a lag, were selected for inclusion into the final model because these climatic variables had the best fit (Supplemental Table 1). Selected covariates for the final model were tested for collinearity, and no collinearity was found (Supplemental Table 2).

**Spatial autocorrelation analysis.** Spatial autocorrelation was explored at a global scale using Moran's I statistic, and at a local scale estimated using the Anselin Local Moran's I statistic (local indicators of spatial association [LISA]) and the Getis-Ord Gi\* statistics. The global Moran's I statistic was used to assess the presence and strength of spatial autocorrelation over the whole study area and to test the assumption of spatial independence in the implementation of the spatial pattern analysis. The LISA and the Getis-Ord Gi\* statistics were used to detect local clustering of malaria and to identify the locations of hotspots. These analyses were conducted using tools provided in ArcMap software.<sup>26</sup>

Spatiotemporal model. Of the 14,976 observations, there were 13,350 (89.1%) zero counts for P. falciparum and 13,864 (92.6%) zero counts for P. vivax. Zero counts can arise from two processes: "excess zeros" (also called structural zeros), for which the process of their occurrence is different from the "random zeros" that arise as part of the Poisson process that generates the malaria counts. One possible explanation for the excess zeros is that they arise in communes that were unable to support malaria transmission during the study period for a variety of epidemiological reasons such as vector habitat unsuitability or isolation from areas with ongoing transmission. Zero-inflated Poisson (ZIP) regression was a better model with lower AIC and Bavesian Information Criterion (BIC) than the standard Poisson regression, and the Vuong test showed the two models were statically different (Supplemental Table 3). Bayesian statistical software Win-BUGS version 1.4 (Medical Research Council, Cambridge, United Kingdom, and Imperial College London, London, United Kingdom) was used to develop ZIP regression models

for *P. falciparum* and *P. vivax* separately. They contained a mixing probability  $\omega$  that the observation is an excess zero count. The model included climatic variables (minimum temperature and precipitation); proportion of the population covered by ITNs and IRS, as explanatory variables; and spatially structured and unstructured random effects.

For the count of malaria cases Y, in the *i*th commune (i = 1...104) and the *j*th month (January 2005–December 2016), the model was structured as follows:

$$\begin{split} & P\left(Y_{ij} = y_{ij}\right) = \begin{cases} \omega + 1 \ (1 - \omega)e^{-\mu}, & y_{ij} = 0\\ (1 - \omega)e^{-\mu} \ \mu_{ij}^{y_{ij}} \ / y_{ij}, & y_{ij} > 0; \end{cases} \\ & Y_{ij} \sim \text{Poisson} \ (\mu_{ij}), \\ & \log \ (\mu_{ij}) = \ \log(E_{ij}) + \theta_{ij}, \\ & \theta_{ij} = \alpha + \beta_1 \times \text{Pop protected}_{ij} + \beta_2 \\ & \times \ \text{Precipitation}_{ij} + \beta_3 \times \text{Tmin}_{ij} + \beta_4 \times \text{trend}_j \\ & + u_i + s_i + w_{ij}, \end{cases} \end{split}$$

where  $E_{ij}$  is the expected number of cases (acting as an offset to control for population size) in commune *i* and month *j*, and  $\theta_{ij}$ is the mean log relative risk (RR);  $\alpha$  is the intercept; and  $\beta_1$ ,  $\beta_2$ ,  $\beta_3$ , and  $\beta_4$  are the coefficients of proportion of population covered by ITNs and IRS, the overall temporal trend of malaria precipitation and minimum temperature; unstructured, spatially structured, and spatiotemporal random effect were denoted by  $u_i$ ,  $s_i$ , and  $w_{ii}$  which assumed to a variance  $\sigma_s^2$  and mean of zero.

A conditional autoregressive prior structure was used to model the spatially structured random effect. Spatial relationships between the communes were determined using a queen contiguity. For two communes sharing a border, an adjacency weight of 1 was assigned, whereas if they did not, the weight was 0. An unbounded uniform (i.e., flat) prior distribution was specified for the intercept, whereas a non-informative normal prior distribution (i.e., with a wide variance,  $\sigma^2 = 1,000$ ) was used for the coefficients. The priors for the precision of unstructured and spatially structured random effects  $(1/\sigma_u^2)$  and  $1/\sigma_s^2$  were specified using non-informative gamma distributions, with shape and scale parameters equal to 0.01.

An initial 10,000 burn-in iterations were discarded. Convergence was examined by running the subsequent blocks of 20,000 iterations, by visual inspection of posterior density and history plots, and occurred at approximately 100,000 iterations for each model. The posterior distributions of each model's parameters were stored after the convergence (100,000 iterations). The summary of the analysis was performed with the posterior mean and 95% credible intervals (Crls). In all analyses, an  $\alpha$ -level of 0.05 was adopted to indicate statistical significance (as indicated by 95% Crl for RR that excluded 1). ArcMap 10.5.1 software (ESRI, Redlands, CA) was used to generate the maps of spatial distribution of posterior means of the unstructured and structured random effects obtained from the three models.

#### RESULTS

**Descriptive analysis.** There were 2,778 *P. falciparum* and 1,770 *P. vivax* cases during the study period. The proportion of

*P. falciparum* cases continued to decrease from 79.0% (211) in 2005 to 50.0% (42) in 2016, whereas the proportion of *P. vivax* increased from 21.0% (55) to 50.0% (42) during the same period. The annual parasite incidence for the study period was 0.28 and 0.18 cases per 1,000 person-years at risk for *P. falciparum* and *P. vivax*, respectively (Table 1). Both species of malaria displayed a strong seasonal pattern, with incidence increases starting in September and peaking in November (Figure 2, Supplemental Figure 1). Both were heterogeneously distributed across the province, with high SMRs in Dong Xuan and Song Hon districts (Supplemental Figure 2).

**Malaria clusters.** The Global Moran's I showed significant spatial autocorrelation for both *P. falciparum* (*z* score = 9.30; P < 0.0001) and *P. vivax* (*z* score = 5.94; P < 0.0001) (Supplemental Figures 3 and 4). Hotspot analysis using the Getis-Ord Gi\* statistic showed that 13 hotspots for *P. falciparum* and 11 hotspots *P. vivax* were located in the communes of Dong Xuan and Son Hoa districts, whereas 37 *P. falciparum* and 35 *P. vivax* coldspots were located in Dong Hoa, Phu Hoa, and Tuy Hoa districts. Nevertheless, cluster analysis using LISA showed only 18 *P. falciparum* and 5 *P. vivax* high–high clusters in Dong Xuan, Phu Hoa, and Son Hoa districts (Figure 3).

**Spatiotemporal model.** *Plasmodium falciparum* incidence decreased by 6.6% (95% Crl 0.6%, 13.3%) and *P. vivax* incidence increased by 89% (95% Crl 72.5%, 107.1%) every month during the study period. A 10-mm increase in precipitation was associated with an increase in *P. falciparum* and *P. vivax* by 5.4% (95% Crl 5.1%, 5.7%) and 3.2% (95% Crl 2.9%, 3.5%), respectively. A minimum temperature increase of 1°C was associated with a decrease in *P. falciparum* and *P. vivax* risk of 7.7% (95% Crl 5.6%, 9.7%) and 10.5% (95% Crl 8.3%, 12.6%), respectively. The model showed that every 10% increase in population protected by IRS and ITNs was associated with a decrease were not statistically significant (Table 2).

The spatially auto-correlated random effect ( $v_i$ ) smooths the spatial pattern of residual variation in malaria incidence after taking into account the fixed effects (Figure 4). Both types of malaria showed areas of lower than average residual malaria risk in Song Cau, Tuy An, Phu Hoa, Dong Hoa, and Tuy Hoa districts. For both types of malaria, areas of higher than

average residual malaria risk were found in Dong Xuan, Son Hoa, and Song Hinh districts.

There was > 95% probability of a higher than provincial average trend of *P. falciparum* in 10/104 communes, which were mostly located in the Son Hoa districts. Similarly, 11/104 communes had > 95% probability of a higher than national average increasing trend of *P. vivax*, also mostly located in Song Cau and Song Hoa districts. For both *P. falciparum* and *P. vivax*, 15/104 districts had > 95% probability of a trend less than the provincial average, mostly located in Tuy Dong Xuan and Hoa districts (Supplemental Figure 5).

## DISCUSSION

Using a surveillance dataset of 12 years (2005–2016), the present study has demonstrated substantial changes occurring with respect to annual trends and the geographical distribution of malaria in Phu Yen Province. In this study, we found that malaria hotspots for both species were found in Dong Xuan and Son Hoa districts. *Plasmodium falciparum* trend decreased, whereas *P. vivax* showed an increasing trend. Both species of malaria displayed a strong seasonal pattern. Prevention measures including LLINs and IRS were not significant predictors of malaria incidence. Minimum temperature was associated with reduction in malaria incidence, whereas precipitation was associated with increase in malaria incidence.

Malaria cases showed a strong seasonal pattern, with cases increasing from September and peaking in November each year. This pattern was associated with the rainy season in Phu Yen, where two seasons (dry, from January to August, and rainy, from September to December) were distinguished. Other studies reported a similar association with rainfall in Bhutan, India, and other parts of the world.<sup>27-35</sup> This finding is consistent with that of the published literature from Vietnam.<sup>17,36</sup> During the study period, P. falciparum showed a decreasing trend, whereas the opposite was true for P. vivax, which is similar to the national trend.<sup>37</sup> As countries embark on malaria elimination, P. falciparum incidence declines more rapidly than the incidence of P. vivax because of the greater effectiveness of vector control interventions on the former.<sup>38</sup> Treating all stages of the parasite (radical cure) is a critical strategy for the successful control and ultimate elimination of P. vivax.39

		Plasmodium falciparum			Plasmodium vivax		
Year	Population	Cases	Proportion of total cases	API	Cases Propo	Proportion of total cases	API
2005	784,003	211	0.79	0.27	55	0.21	0.07
2006	791,922	352	0.86	0.44	56	0.14	0.07
2007	799,921	168	0.81	0.21	40	0.19	0.05
2008	808,001	156	0.92	0.19	13	0.08	0.02
2009	816,163	274	0.85	0.34	48	0.15	0.06
2010	824,407	222	0.80	0.27	57	0.20	0.07
2011	832,734	244	0.65	0.29	133	0.35	0.16
2012	841,146	331	0.54	0.39	281	0.46	0.33
2013	849,642	327	0.49	0.38	339	0.51	0.40
2014	858,138	348	0.45	0.41	419	0.55	0.49
2015	866,720	103	0.26	0.12	287	0.74	0.33
2016	875,387	42	0.50	0.05	42	0.50	0.05
Overall	9,948,183	2,778	0.61	0.28	1,770	0.39	0.18

TABLE 1 Malaria incidence during the study period (2005–2016)

API = annual parasite incidence.



FIGURE 2. Decomposed monthly (A) Plasmodium falciparum and (B) Plasmodium vivax incidence in Phu Yen Province, Vietnam, 2005–2016.

Malaria risk continues to decline in Vietnam, and transmission is becoming increasingly heterogeneous, with most cases now concentrated within a relatively small number of communes, including Phu Yen.<sup>11</sup> Such patterns are consistent with those of other countries with low malaria burden.<sup>27,35,40–43</sup> Therefore, spatially targeting interventions and associated resources are likely to achieve better results than a uniform approach to the distribution and delivery of malaria reduction interventions.44-46 Geographical information system-based spatial decision support systems (SDSSs) are one of the tools currently being used in countries in the Asia-Pacific region to support enhanced surveillance in priority areas, primarily as a means for locating malaria transmission, identifying and targeting appropriate foci-specific interventions, and ensuring these interventions are implemented at optimal levels of coverage.47-49 Combining the capacity to identify priority malaria clusters and associated appropriate foci-specific interventions with enhanced surveillance and intervention management tools such as a SDSS provides significant potential opportunity to target and strengthen elimination efforts where needed.45,50

This study found evidence of significant spatial variability in malaria incidence within Phu Yen Province. This likely arose because of two main processes: first, the effects of the covariates in the model (preventive coverage and climate) and, second, the residual effects of additional, unmeasured influences on malaria incidence that were captured by the random effects—these were both spatially structured and unstructured. Given that the areas of high residual risk were in the western part of the province, proximity to other high-transmission areas might partly explain this residual variation; however, further investigation is required.

One of the strengths of this study was the capacity to implement the spatial analysis at a relatively fine resolution, being the commune level. Traditionally, spatial patterns of malaria risk have been displayed at larger geographical units, such as at district, province, national, regional, and global scales.<sup>51,52</sup> However, such resolution may mask more localized underlying patterns of disease through averaging.<sup>53</sup> Therefore, use of finer geographic units such as communes may be necessary to observe important local variation in spatial patterns of malaria risk and to better guide disease control efforts and resource allocation, particularly when transmission declines to levels favorable to the pursuit of elimination.<sup>44</sup>

This study used Bayesian statistical methods to quantify seasonal and commune variations of *P. falciparum* and *P. vivax* and the effects of climatic factors. The finding from this analysis indicated that precipitation and temperature were important drivers of spatiotemporal patterns of malaria incidence in Phu Yen. Although there was a significant reduction in malaria incidence, this success has not been evenly distributed throughout Phu Yen, and spatial heterogeneities remain (Figure 4). Targeted distribution of resources should be implemented using evidence-based approaches, supported by spatiotemporal analytical methods, to assist more effective malaria control in Phu Yen Province where these resources are most needed.

A limitation of the study included the use of routine case reports to measure incidence. Known issues exist surrounding completeness and representativeness of such data. It has been reported that routine reporting of malaria cases through the health information system in Vietnam underestimates the true number of cases.<sup>54</sup> Whether these factors affect the validity of our analysis depends on whether or not underreporting systematically differs between communes and time-periods. Second, populations of districts were projected and may have led to over or under estimation. Third, we used long-term interpolated climatic variables because there were no data that coincided with the study time period. This might have impacted the spatiotemporal distribution of malaria. Finally, unmeasured risk modifiers, such as socio-economic development, living standards, treatment, localized



FIGURE 3. Malaria cluster maps by communes of Phu Yen Province, Vietnam, 2005–2016. (A) Getis-Ord Gi\* statistics and (B) Anselin's Local Moran's I for *Plasmodium falciparum*. (C) Getis-Ord Gi\* statistics and (D) Anselin's Local Moran's I for *Plasmodium vivax*. This figure appears in color at www.ajtmh.org.

behavioral patterns, population mobility, imported cases, and bed net use, are unaccounted for in this study.  $^{55-57}$ 

## CONCLUSION

Minimum temperature was associated with decreased risk, whereas precipitation was associated with increased risk of both *P. falciparum* and *P. vivax*. A high residual risk area of malaria transmission (after accounting for intervention and climate variables) was identified in the northwestern part of Phu Yen Province. Hence, targeted distribution of resources, including intensified interventions, in this part of the province will be required for local malaria elimination. Similar approaches can be used for identifying spatial heterogeneity of malaria transmission for resource allocation by malaria elimination countries.

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Regression coefficients and 95% CrI from Bayesian spatial and nonspatial models of *P. falciparum* and *P. vivax* cases reported by month and communes in Phu Yen Province, Vietnam, 2005–2016

Variable	P. falciparum RR (95% Crl)	<i>P. vivax</i> RR (95% Crl)
Intercept*	-1.14 (-1.38, -0.94)	-1.23 (-1.48, -1.02)
Population protected (10% increase)†	0.999 (0.998, 1.00)	1.00 (0.998, 1.001)
Precipitation (10 mm increase)	1.054 (1.051, 1.057)	1.032 (1.029, 1.035)
Temperature minimum (°Celsius)	0.923 (0.903, 0.944)	0.895 (0.874, 0.917)
Mean monthly trend	0.934 (0.867, 1.006)	1.89 (1.725, 2.071)
Proportion of zero	0.218 (0.171, 0.264)	0.277 (0.226, 0.327)
Heterogeneity		
Unstructured	2.537 (0.689, 8.063)	2.038 (0.581, 6.879)
Structured (spatial)	0.153 (0.091, 0.258)	0.155 (0.084, 0.283)
Structured (trend)	2.903 (1.681, 4.675)	2.45 (1.364, 4.084)

Crl = credible interval; *P. falciparum* = *Plasmodium falciparum*; *P. vivax* = *Plasmodium vivax*; RR = relative risk.

\* Coefficients.

† Proportion of population protected by preventive measures.



FIGURE 4. Spatial distribution of the posterior means of structured and unstructured random effects from model in Phu Yen Province, Vietnam, 2005–2016. (A) Structured and (B) unstructured random effects of *Plasmodium falciparum*; (C) structured and (D) unstructured random effects of *Plasmodium vivax*. This figure appears in color at www.ajtmh.org.

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Note: Supplemental figures and tables appear at www.ajtmh.org.

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Disclaimer: The dataset and materials used for this study can be made available only upon approval by the National Institute of Malariology, Parasitology, and Entomology (NIMPE), Vietnam.

Disclosure: N. J. M. is a U.S. military service member or federal/ contract employee of the U.S. government. This work was prepared as part of his official duties. Title 17 U.S.C. 105 provides that "copyright protection under this title is not available for any work of the United States Government." Title 17 U.S.C. 101 defines a U.S. government work as work prepared by a military service member or employee of the U.S. government as part of that person's official duties. The study protocol was approved, and permission to use these data was granted by the Ministry of Health of Vietnam on November 20, 2017 (5234/QD-BYT).

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