



Recent Spatial and Temporal Trends of Malaria in Korea

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Abstract: This study was done to provide an analytical overview on the latest malaria infection clusters by evaluating temporal trends during 2010-2019 in Korea. Incheon was the most likely cluster (MLC) for all cases of malaria during the total period. MLCs for *P. falciparum*, *vivax*, *malariae*, *ovale*, and clinically diagnosed malaria without parasitological confirmation were Jeollanam-do, Incheon, Gangwon-do, Gyeongsangnam-do, and Jeollabuk-do, respectively. Malaria was decreasing in most significant clusters, but Gwangju showed an increase for all cases of malaria, *P. vivax* and clinically diagnosed cases. Malaria overall, *P. falciparum* and *P. vivax* seem to be under control thanks to aggressive health measures. This study might provide a sound scientific basis for future control measures against malaria in Korea.

Key words: Malaria, health insurance review, assessment service, cluster, trend, spatial analysis, temporal analysis, SaTScan

INTRODUCTION

Diseases transmitted by arthropod vectors are major risk factors responsible for the global burden of infectious diseases [1]. Mosquito-borne diseases (MBDs) are important in this respect as they include malaria and several important arboviruses [2,3]. Aside from the high burden, they tend to emerge in new areas, and also re-emerge in regions known to have been eradicated.[2] Several factors such as globalization and climate changes are inducing shifts in communicable disease trends, and this is especially evident in MBDs [1,4] as its incidence seems to show an increasing trend in parts of Korea.

Malaria is one of the oldest diseases, and its influence has probably been greater than that of any other infectious agents [5]. Caused by the *Plasmodium* species of protozoan parasites, malaria is a major public health burden worldwide [6]. *P. falciparum* is the most prevalent malaria parasite in Africa, and it is the greatest menace because of its high mortality rate [7]. *P. vivax* is less lethal than *P. falciparum*, but more prevalent in Korea [6]. It was eradicated from Korea late 1970s, but re-emerged 1993 and continued prevalent despite the ongoing national eradication program [8]. *P. malariae* is a benign ma-

laria with several distinct clinical features [9]. *P. ovale* make the infected RBC swollen and oval, the margin fimbriated, and stained pale, morphologically distinct from the others. In Korea, majority of the imported malaria cases were of *P. falciparum* (from Africa) and *P. vivax* (from Southeast Asia), whereas *P. malariae* and *P. ovale* cases were very rare [10].

Korea has a complete full health-coverage of its population. The clinical data from the Health Insurance Review and Assessment Service (HIRA) of Korea were used in this study, so that the whole Korean population is represented. Our study was performed to figure out spatial infection clusters and trends of the malaria, *Plasmodium* infections during recent 10 years in Korea.

MATERIALS AND METHODS

Ethics statement

This study was performed under the regulation of the IRB Committee of The Catholic University of Korea (No. MC-20ZASE0155). This research adhered to the tenets of the Declaration of Helsinki.

Data acquisition and definition

The nationwide malaria cases for 10 years (2010-2019) from the 16 administrative districts in Korea were obtained from the Healthcare Bigdata Hub provided by the HIRA. (<https://open-data.hira.or.kr/home.do>. [cited 2020 October]) The HIRA classifies the *Plasmodium* infections as *P. falciparum* (B50), *P. vivax*

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(B51), *P. malariae* (B52), *P. ovale*, simian plasmodia (B53) and clinically diagnosed malaria without parasitological confirmation (B54).

The population data of 16 administrative divisions for the same period were obtained from the Korean Statistical Information Service (<https://kosis.kr/index/index.do>. [cited 2021 October]). The geographical locations of the cases were set to the latitude and longitude of administrative center as the search point since the case data were compiled by administrative district unit. The population and number of cases were

summed up as accumulated counts and numbers for the total period to evaluate overall malaria status. Case numbers were adjusted for the respected district population (cases/1,000,000) to facilitate comparison between the districts (Table 1).

Malaria disease according to individual classification were further evaluated by dividing the timeline into tertiles (2010-2012, 2013-2015, 2016-2019), and total period. The third tertile had 4 years of data to reflect more recent information, and to facilitate the evaluation of temporal characteristics.

Table 1. Population and malaria cases for 2010-2019 according to administrative district

Cluster (code)	Cumulated Population	Cases					
		<i>P. falciparum</i>	<i>P. vivax</i>	<i>P. malariae</i>	<i>P. ovale</i>	Clinically diagnosed	All cases
Seoul (SE)	100,309,373	492	999	37	113	2,617	4,258
Busan (BU)	35,042,646	89	97	4	17	503	710
Daegu (DG)	24,866,860	56	67	1	10	352	486
Incheon (IN)	28,915,035	75	568	9	18	991	1,661
Gwangju (GW)	14,657,097	51	42	4	32	357	486
Daejeon (DJ)	15,108,648	57	53	2	6	217	335
Ulsan (UL)	11,546,517	23	30	0	6	89	148
Gyeonggi-do (GY)	124,839,896	414	2,126	80	141	3,561	6,322
Gangwon-do (GA)	15,426,610	84	114	37	15	425	675
Chungcheongbuk-do (CB)	15,798,992	45	46	9	9	323	432
Chungcheongnam-do (CN)	20,856,616	51	39	12	18	537	657
Jeollabuk-do (JB)	18,605,718	50	69	13	35	880	1,047
Jeollanam-do (JN)	19,016,443	271	43	5	14	608	941
Gyeongsangbuk-do (GB)	26,925,299	73	66	5	31	794	969
Gyeongsangnam-do (GN)	33,458,210	99	66	10	61	653	889
Jeju-do (JE)	6,193,531	4	11	4	4	195	218
Total	511,567,491	1,934	4,436	232	530	13,102	20,234
Adjusted cases*							
Seoul (SE)		4.9	9.96	0.37	1.13	26.09	42.45
Busan (BU)		2.54	2.77	0.11	0.49	14.35	20.26
Daegu (DG)		2.25	2.69	0.04	0.40	14.16	19.54
Incheon (IN)		2.59	19.64	0.31	0.62	34.27	57.44
Gwangju (GW)		3.48	2.87	0.27	2.18	24.36	33.16
Daejeon (DJ)		3.77	3.51	0.13	0.4	14.36	22.17
Ulsan (UL)		1.99	2.60	0.00	0.52	7.71	12.82
Gyeonggi-do (GY)		3.32	17.03	0.64	1.13	28.52	50.64
Gangwon-do (GA)		5.45	7.39	2.40	0.97	27.55	43.76
Chungcheongbuk-do (CB)		2.85	2.91	0.57	0.57	20.44	27.34
Chungcheongnam-do (CN)		2.45	1.87	0.58	0.86	25.75	31.50
Jeollabuk-do (JB)		2.69	3.71	0.70	1.88	47.3	56.27
Jeollanam-do (JN)		14.25	2.26	0.26	0.74	31.97	49.48
Gyeongsangbuk-do (GB)		2.71	2.45	0.19	1.15	29.49	35.99
Gyeongsangnam-do (GN)		2.96	1.97	0.30	1.82	19.52	26.57
Jeju-do (JE)		0.65	1.78	0.65	0.65	31.48	35.20
Total		59	85	8	16	397	565

*Adjusted cumulated infection rate (cases/1,000,000).

Spatiotemporal statistical analyses

Spatial scan analysis detects clusters with maximum likelihood ratio by creating a circular window on a map and scanning the study area by varying the window size. The window size determines a percentage of the population at risk within its boundaries [11]. Spatial scan statistic works best for detecting spatial clusters, and may be effective in the study of small numbered cases, such as novel or infrequent outbreaks [12, 13]. SaTScan (v10.0) [14] software was used to detect these clusters and evaluate their significance through simulation. The discrete Poisson model was used since the malaria case data were linked to their background population at risk.

Disease trend was determined through spatial variation in temporal trend analysis, by scanning for clusters with either increasing or decreasing rates. Data for *P. malariae* in Ulsan was missing and was adjusted for known relative risks according to the SaTScan software guide.

It is important to find an appropriate set value of cluster size because a large value could hide effect of small core clusters, while a small value could overlook the regional pattern of clusters [15]. Preliminary studies with 15 and 25% spatial window sizes showed that the 15% window size was most effective in finding clusters. The present study selected this window size for statistical analyses.

Statistical significance of the clusters was calculated using the Monte Carlo simulations with an inference of 9,999 [16] and expressed as *P*-value. A significance level of $\alpha < 0.05$ was used as a standard. QGIS software (v3.16) was used to visualize cluster patterns on a map. The clusters and trends are shown in order of log likelihood ratio (LLR).

RESULTS

General characteristics

Gyeonggi-do had the highest population (124,839,896) and highest number of cases (31.3%) during the study period. After adjusting the case numbers for population, Incheon (57.4) and Jeollabuk-do (56.3) showed the highest rate of infection.

Spatial clusters

When considering for all cases of malaria, Incheon was the most likely cluster (MLC) with a relative risk (RR) and LLR of 1.49 and 109.579, respectively, for the whole study period. The joint cluster of Jeollanam-do/Jeollabuk-do was next (RR = 1.37,

LLR = 82.628), followed by Jeollabuk-do (RR = 1.45, LLR = 60.557), and Jeollanam-do (RR = 1.26, LLR = 22.861) (Table 2). Gangwon-do was a significant cluster at the first tertile of 2010-2012, and Gwangju was also significant at the mid tertile of 2013-2015 (Fig. 1).

Among clusters detected for *P. falciparum*, Jeollanam-do was the MLC (RR = 4.22, LLR = 171.545) during the whole period. The next likely cluster was Gangwon-do (RR = 1.46, LLR = 5.146) (Table 2). Jeollanam-do was a significant at all tertiles, while Gangwon-do (RR = 1.46, LLR = 5.146) was only significant at the earlier tertile (Fig. 2). Incheon was the MLC (RR = 2.45, LLR = 159.540) for *P. vivax* during the whole period (Table 2), and also the only significant cluster at all tertiles (Fig. 3). The MLC for *P. malariae* was Gangwon-do (RR = 5.96, LLR = 33.013) during the whole period (Table 2). Gyeongsangnam-do was the MLC (RR = 1.86, LLR = 8.853) for *P. ovale* during the whole period. Gwangju was the second (RR = 2.18, LLR = 7.316), and Jeollabuk-do was the third likely cluster during the whole period (RR = 1.87, LLR = 5.398) (Table 2).

For clinically diagnosed malaria, Jeollabuk-do was the MLC (RR = 1.91, LLR = 142.846) for the whole period, followed by Incheon (RR = 1.37, LLR = 40.797), Jeju-do/Jeollanam-do (RR = 1.26, LLR = 18.776) and Gyeongsangbuk-do (RR = 1.16, LLR = 7.973) (Table 2).

Temporal trends

Temporal trends analysis for the whole study period showed that malaria infection rate was decreasing in most clusters (Fig. 4) Gwangju showed an increasing rate for all cases of malaria (+8.8%, LLR = 73.128), *P. vivax* (+6.03%, LLR = 6.866) and clinically diagnosed cases (+17.25%, LLR = 85.970). *P. falciparum* showed decreasing rates in Jeollanam-do/Jeollabuk-do/Jeju-do/Daejeon (-4.70%, LLR = 20.700) and Gyeongsangbuk-do/Daegu/Chungcheongbuk-do (-23.70%, LLR = 9.456) joint clusters. Other clusters detected by purely spatial analysis showed varying degrees of decreasing rates (Table 3).

DISCUSSION

The present study provides an analytical overview of the latest malaria status in Korea by determining disease clusters and evaluating temporal trends during 2010-2019.

Korea is a peninsula, but the northern border is a demilitarized zone (DMZ) and cut off from the rest of Asia by North Korea. The result is virtually an isolated island, and contact

Table 2. Clusters detected malaria

	District	Cases	Expected	Relative Risk	Log Likelihood Ratio	P-value
<i>All cases</i>						
2010-2012	Incheon	694	471	1.52	49.274	<0.001
	Jeollanam-do, Jeollabuk-do	733	636	1.17	7.674	0.00
	Gangwon-do	314	258	1.23	5.914	0.01
2013-2015	Incheon	497	322	1.60	43.556	<0.001
	Gyeongsangbuk-do	406	300	1.38	17.984	<0.001
	Gwangju	202	164	1.24	4.342	0.04
2016-2019	Jeollanam-do, Jeollabuk-do	823	438	2.02	147.836	<0.001
	Jeollabuk-do	438	216	2.11	91.604	<0.001
	Jeollanam-do	385	221	1.79	51.661	<0.001
Total period	Incheon	470	346	1.39	21.285	<0.001
	Incheon	1,661	1,144	1.49	109.579	<0.001
	Jeollanam-do, Jeollabuk-do	1,988	1,488	1.37	82.628	<0.001
	Jeollabuk-do	1,047	736	1.45	60.557	<0.001
	Jeollanam-do	941	752	1.26	22.861	<0.001
<i>P. falciparum</i>						
2010-2012	Jeollanam-do	95	35	2.91	37.006	<0.001
	Gangwon-do	47	28	1.71	5.52	0.01
2013-2015	Jeollanam-do	65	19	3.81	36.544	<0.001
2016-2019	Jeollanam-do	111	18	7.46	116.334	<0.001
Total period	Jeollanam-do	271	72	4.22	171.545	<0.001
	Gangwon-do	84	58	1.46	5.146	0.02
<i>P. vivax</i>						
2010-2012	Incheon	264	117	2.42	72.733	<0.001
2013-2015	Incheon	169	59	3.22	73.935	<0.001
2016-2019	Incheon	135	73	1.96	22.949	<0.001
Total period	Incheon	568	251	2.45	159.54	<0.001
<i>P. malariae</i>						
Total period	Gangwon-do	37	7	5.96	33.013	<0.001
<i>P. ovale</i>						
Total period	Gyeongsangnam-do	61	35	1.86	8.853	<0.001
	Gwangju	32	15	2.18	7.316	0.00
	Jeollabuk-do	35	19	1.87	5.398	0.01
<i>Clinically diagnosed malaria</i>						
Total period	Jeollabuk-do	880	477	1.91	142.846	<0.001
	Incheon	991	741	1.37	40.797	<0.001
	Jeju-do, Jeollanam-do	803	646	1.26	18.776	<0.001
	Gyeongsangbuk-do	794	690	1.16	7.973	0.00

*Decreasing order of Log Likelihood Ratio.

with the outside world is limited to air- and sea-routes. The western and southern parts have numerous islands that number around 5,000, and a fair number of them are inhabited. This means limited resources stretched over a large area with isolated populations, a potential health hazard for the related authorities.

In terms of climate, Korea is mostly located in the temperate zone, except for Jeju-do, but global warming has caused semi-tropical climate changes in the southern parts of the mainland.

These factors might have caused an influence in the increased rate of various infectious diseases.

With recent economic prosperity large numbers of Koreans are enjoying travelling, both domestic and international. And especially among the young generation, their interest has shifted from urban travel to seeking out nature as it is. Trekking, hiking, wildlife experiences, and related outdoor life have become the trend. This may also have an effect on the spread of malaria.

We included the data on clinically diagnosed malaria with-

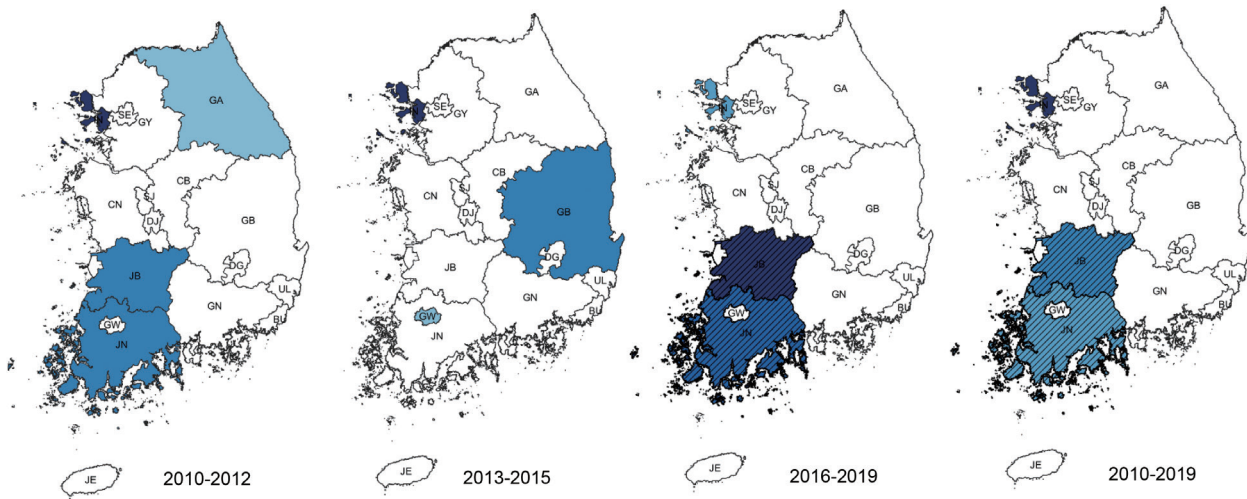


Fig. 1. Clusters detected for all cases of malaria. Area codes represent administrative districts of the Korean government. Light to dark blue gradient indicates increasing order of log likelihood ratio. Dashed lines represent a joint cluster.

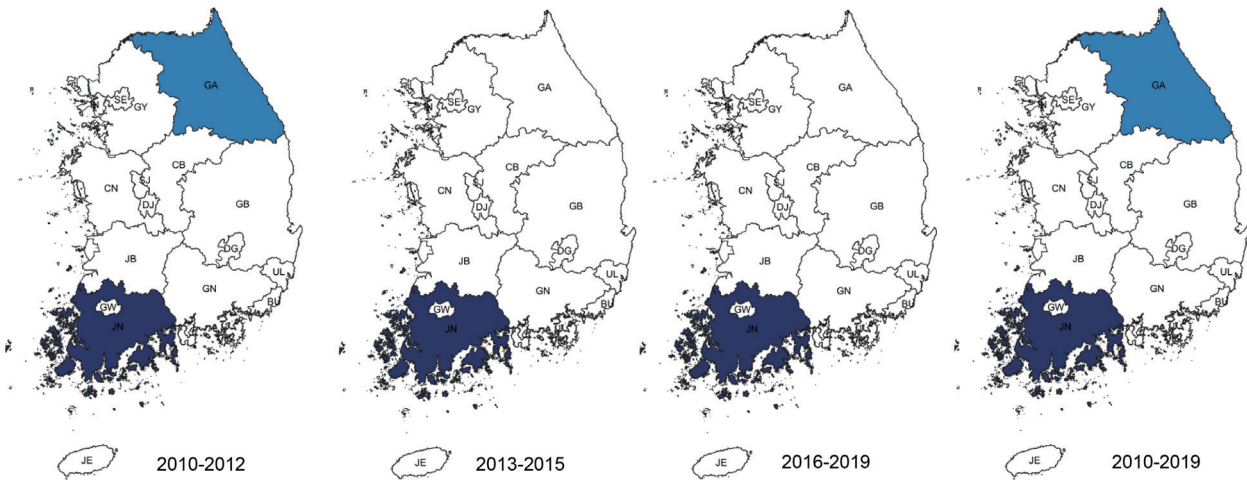


Fig. 2. Clusters detected for *P. falciparum*. Area codes represent administrative districts. Light to dark blue gradient indicates increasing order of log likelihood ratio.

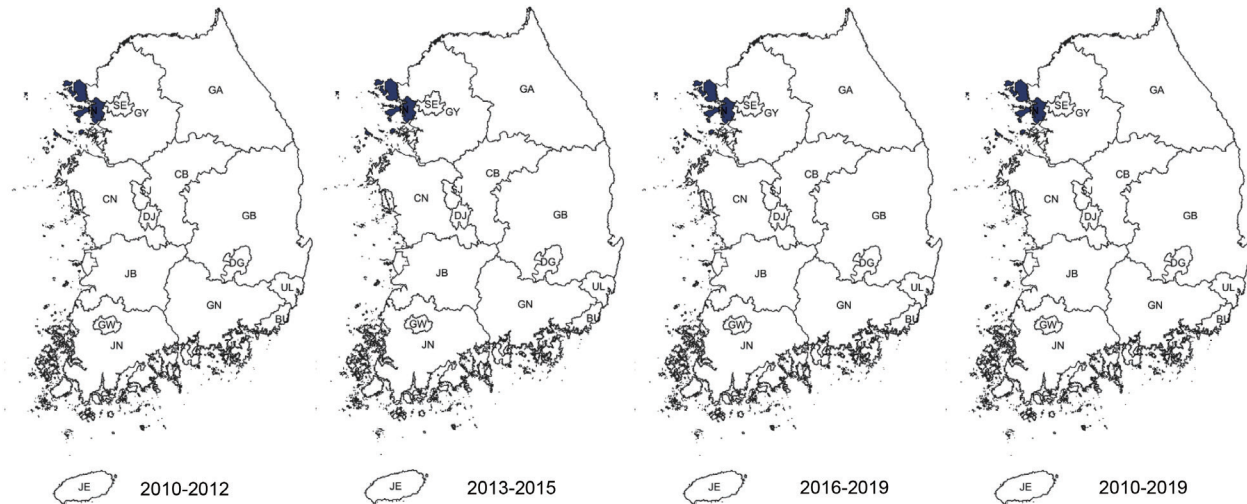


Fig. 3. Clusters detected for *P. vivax*. Area codes represent administrative districts. Incheon was the most likely cluster.

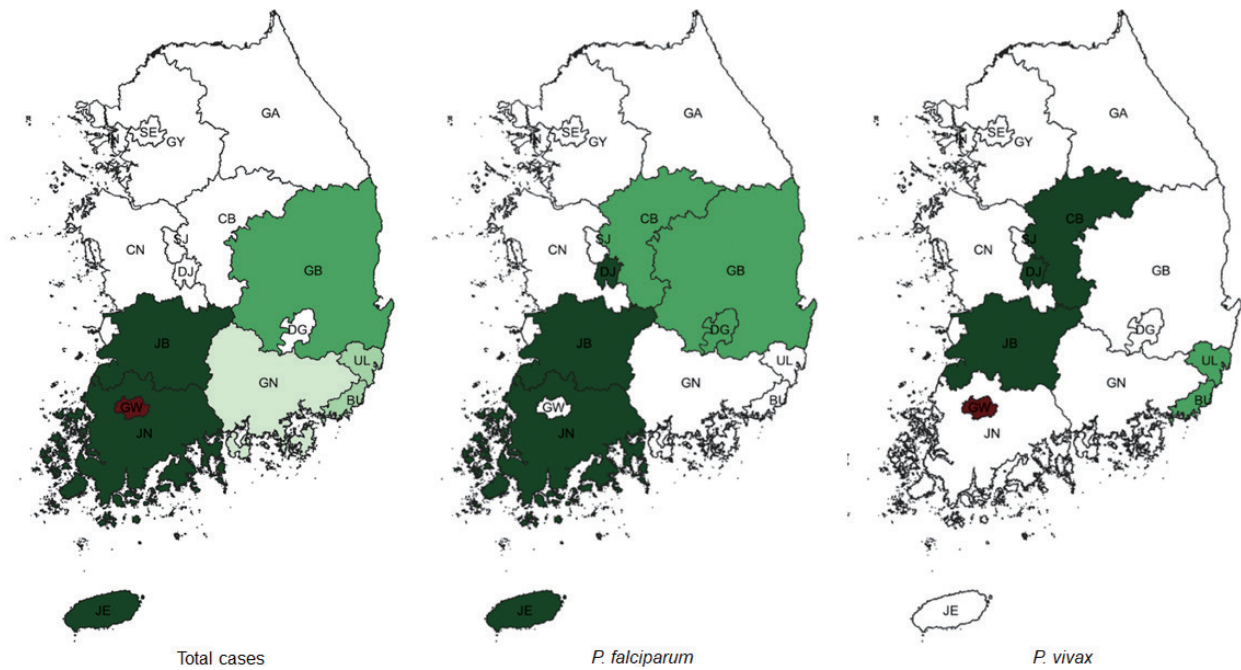


Fig. 4. Spatiotemporal trends of malaria. Area codes represent administrative districts. Green represents clusters that showed decreased rate over the period of 2010-2019. Light to dark green gradient indicates increasing order of log likelihood ratio. Red represents clusters that showed increase over the same period.

Table 3. Temporal trends of malaria

Malaria	Districts	Cases	Trend inside cluster	Trend outside cluster	Log Likelihood Ratio	P-value
All cases	Gwangju	486	8.80	-10.47	73.128	<0.001
	Jeju-do, Jeollanam-do, Jeollabuk-do	2,206	-2.57	-10.93	63.799	<0.001
	Gyeongsangbuk-do	969	-19.70	-9.56	45.922	<0.001
	Ulsan, Busan	858	-15.23	-9.83	11.593	<0.001
	Gyeongsangnam-do	889	-6.58	-10.19	5.260	0.03
<i>P. falciparum</i>	Jeollanam-do, Jeollabuk-do, Jeju-do, Daejeon	382	-4.70	-16.47	20.700	<0.001
	Gyeongsangbuk-do, Daegu, Chungcheongbuk-do	193	-23.70	-13.15	9.456	0.00
<i>P. vivax</i>	Jeollabuk-do, Daejeon, Chungcheongbuk-do	168	-22.86	-12.80	7.965	0.00
	Ulsan, Busan	127	-23.66	-12.89	6.925	0.01
	Gwangju	42	6.03	-13.33	6.866	0.01
<i>P. malariae</i>	Jeollanam-do, Jeollabuk-do, Jeju-do, Daejeon	24	22.14	-6.27	5.741	0.02
<i>P. ovale</i>	Jeollabuk-do	35	0.80	-19.81	6.714	0.01
Clinically diagnosed	Gwangju	357	17.25	-8.78	85.97	<0.001
	Jeju-do, Jeollanam-do, Jeollabuk-do	1,683	-0.50	-9.25	50.545	<0.001
	Gyeongsangbuk-do	794	-18.45	-7.49	42.941	<0.001
	Daejeon	217	-21.79	-7.93	19.403	<0.001
	Ulsan, Busan	592	-13.11	-7.94	7.191	0.00
	Chungcheongbuk-do	323	-1.81	-8.31	6.027	0.01

*Trend inside/outside cluster denotes % increase/decrease over the period of 2010-2019.

**Decreasing order of Log Likelihood Ratio.

out parasitological confirmation. This was a point of debate among the authors. The purpose of this study was to determine the current status of malaria in Korea, so we decided to

include it.

When all cases of malaria during the whole study period were evaluated, significant clusters were observed in the west-

ern part of Korea, mainly Incheon and the Jeolla area (Jeollabuk-do and Jeollanam-do). Gangwon-do, and Gyeongsangbuk-do were clusters during the first tertile (2010-2012). Located in the western part of Korea, Incheon is the main access point for international travel and commerce, and its proximity with China has resulted in a large number of Chinese immigration. The outermost regions are also adjacent to North Korea. The Jeolla area is mainly an agricultural zone located in the west. Additionally, together with Incheon, fishery is also important in this area. Both are labor intensive, and with the Korean population declining and growing old, it has resulted in a large influx of foreigners, mainly from Southeast Asia, but also from Africa and South America.

Gangwon-do has a relatively small population spread over a large area. The DMZ is located in its northern part, which means an even smaller civilian population. Ironically, despite its name, this place is one of the most militarized area in the world, and generally off limits to civilians. The result is one of the most preserved natural habitats in the world abounding with wildlife. Because of its location between the two Koreas and the absence of humans, animals move freely into both countries. Apart from the preservation of nature that results in potential animal reservoirs of zoonosis, a large number of soldiers are stationed in this area. This has interesting consequences. Technically still at war, Korea has military conscription, so male citizens between the ages of 18 and 28 have to perform compulsory military service. A large number are also stationed in Incheon. After military service, these young men go back home, all throughout Korea. They might play a role in disease reservoir and spread of disease.

P. falciparum clusters were observed in Jeollanam-do and Gangwon-do. As mentioned above, Jeollanam-do is located in the southwestern part of Korea, and has an almost semi-tropical climate at times. Agriculture and fisheries require massive influx of foreign labor force, which might all be significant. Indigenous falciparum malaria has not been reported [7]. Gangwon-do was a cluster in the first tertile. This could have been caused by the large number of soldiers stationed in the DMZ. In 1979 the WHO officially certified that Korea was a malaria-free country [18], but vivax malaria re-emerged in 1993 in a soldier. Massive efforts were enforced to contain the disease, and this might have inadvertently caused the decrease in *P. falciparum*.

Vivax malaria has been an endemic infectious disease in the Korea for a long time [7]. Although it has re-emerged, renewed

eradication programs have been put in force. Our study shows that Incheon is a significant cluster. Most cases at the beginning of the vivax malaria outbreak occurred among soldiers stationed near the DMZ in the northern part of Gyeonggi and Gangwon-do, and among veterans who had been discharged. This suggested that North Korea might be a major reservoir of vivax malaria, and Incheon is also adjacent to North Korea. Another interesting point to consider is that the re-emerged *P. vivax* in Korea has genealogical origin in Southern China. As mentioned above, Incheon is major point of contact with China [19].

P. malariae causes the most benign form of malaria infection, and has several distinct clinical features [20]. It has been detected in the Greater Mekong Subregion (GMS) of Southeast Asia [21]. Foreign workers from this region can be seen frequently in Korea, and this population might have been reflected in the cluster found in our study. This region also happens to be a favorite destination for Korean travelers. A cluster was observed in Gangwon-do.

P. ovale occurs mainly in sub-Saharan Africa and islands of the western Pacific [22], and often presents as mixed infections with other *Plasmodium* species. The Gyeongsang area has a large population, and thus has major international airports that connect to the West Pacific countries, another favorite destination for Korean travelers. As a result, in contrast to the other *Plasmodium* species, a *P. ovale* cluster was observed in Gyeongsangnam-do. Gwangju Metropolitan City is located roughly at the geographic center of the Jeolla area, and as its name implies, is host to major medical resources, so that complicated cases from this area might have been referred to Gwangju, thus resulting in a cluster.

In the case of clinically diagnosed malaria without parasitological confirmation, the symptoms and signs might have been obvious enough to have been diagnosed without confirmation. Clusters were observed throughout the country, except for most major metropolitan cities (Seoul, Daejeon, Daegu, Ulsan, Busan), Gyeonggi-do and the Chungcheong area. The major cities and Gyeonggi-do are far more industrialized and urbanized, with better access to medical resources. This may have helped in lowering the malaria risk.

On the other hand, while the Chungcheong area is situated in the middle of Korea, it tends to be isolated in some places, and its medical resource is far from satisfactory. Thus, its low rate of malaria is a paradox, but this might have been caused because its patients would simply just visit a medical facility in

a nearby city. Being in the middle of everywhere, travelling is facilitated by extensive road and railway infrastructure. And being an inland area, global warming might have had a lesser impact.

Significant clusters with either increasing or decreasing rates of malaria were mainly located in the southern part of Korea. Evaluation for all of cases of malaria showed decreasing trends in the southern parts, except for Gwangju. The city might function as the main medical support system for the surrounding area, so complicated cases would all end up here, resulting in an increasing trend (8.8%) compared to areas outside of the cluster (-10.47%).

Clinically diagnosed cases showed a similar trend, with Gwangju showing an increase of 17.25% in comparison to a decrease of -8.78% in other areas. General trends showed that these cases decreased initially, but increased again. This may have resulted in the significant increase observed in Gwangju. Despite being a major cause of malaria, *P. falciparum* showed a decreasing rate. Aggressive malaria containment measures seem to have been effective. *P. vivax* also showed a decreasing trend, but was increasing in Gwangju (6.03% vs. -13.33%). Its role as a health resources hub again seems to be responsible. Interestingly, the general trend of *P. vivax* showed that although cases were decreasing in number, the adjusted numbers were high and concentrated around Incheon, Gyeonggi-do and Seoul. This might potentially reflect the fact that aggressive anti-malaria health measures were effective in controlling vivax malaria elsewhere.

Although the cases are fewer in number, *P. malariae* showed an increasing trend in the Daejeon/Jeolla area/Jeju-do joint cluster (22.14% vs. -6.27%), while *P. ovale* was increasing in Jeollabuk-do (0.80% vs. -19.81%). A climatic factor, such as higher average temperature or rainfall might have been possible during this period, with facilitated survival of the pathogens. Also, an increased influx of potential hosts from afflicted areas, both Korean and foreign, might have been responsible for this changing trend. These variants might occupy an ever more significant niche, and in the future, even surpass vivax and falciparum malaria.

In conclusion, after taking account of indigenous malaria with various health measures, global warming seems to be the factor most responsible for malaria in Korea as the southern regions are most affected.

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CONFLICT OF INTEREST

The authors declare no conflict of interest related to this study.

REFERENCES

1. Franklins LHV, Jones KE, Redding DW, Abubakar I. The effect of global change on mosquito-borne disease. *Lancet Infect Dis* 2019; 19: 302-312. [https://doi.org/10.1016/S1473-3099\(19\)30161-6](https://doi.org/10.1016/S1473-3099(19)30161-6)
2. Huang YJS, Higgs S, Vanlandingham DL. Emergence and re-emergence of mosquito-borne arboviruses. *Curr Opin Virol* 2019; 34: 104-109 <https://doi.org/10.1016/j.coviro.2019.01.001>
3. Kilpatrick AM, Randolph SE. Drivers, dynamics, and control of emerging vector-borne zoonotic diseases. *Lancet* 2012; 380: 1946-1955. [https://doi.org/10.1016/S0140-6736\(12\)61151-9](https://doi.org/10.1016/S0140-6736(12)61151-9)
4. Bahk YY, Park SH, Kim-Jeon MD, Oh SS, Jung H, Jun H, Kim KA, Park JM, Ahn SK, Lee J, Choi EJ, Moon BS, Gong YW, Kwon MJ, Kim TS. Monitoring culicine mosquitoes (diptera: culicidae) as a vector of flavivirus in incheon metropolitan city and Hwaseong-Si, Gyeonggi-Do, Korea, during 2019. *Korean J Parasitol* 2020; 58: 551-558. <https://doi.org/10.3347/kjp.2020.58.5.551>
5. Carter R, Mendis KN. Evolutionary and historical aspects of the burden of malaria. *Clin Microbiol Rev* 2002; 15: 564-594. <https://doi.org/10.1128/cmr.15.4.564-594.2002>
6. Lee SK, Hu F, Firdaus ER, Park JH, Han JH, Lee SE, Shin HI, Cho SH, Park WS, Lu F, Han ET. Surveillance on the vivax malaria in endemic areas in the Republic of Korea based on molecular and serological analyses. *Korean J Parasitol* 2020; 58: 609-617. <https://doi.org/10.3347/kjp.2020.58.6.609>
7. Bahk YY, Lee HW, Na BK, Kim J, Jin K, Hong YS, Kim TS. Epidemiological characteristics of re-emerging vivax malaria in the Republic of Korea (1993-2017). *Korean J Parasitol* 2018; 56: 531-543. <https://doi.org/10.3347/kjp.2018.56.6.531>
8. Chai JY. Re-emerging *Plasmodium vivax* malaria in the Republic of Korea. *Korean J Parasitol* 1999; 37: 129-143. <https://doi.org/10.3347/kjp.1999.37.3.129>
9. Collins WE, Jeffery GM. *Plasmodium malariae*: parasite and disease. *Clin Microbiol Rev* 2007; 20: 579-592. <https://doi.org/10.1128/CMR.00027-07>
10. Shin HI, Ku B, Kim YJ, Kim TY, Cho SH, Lee SE. Diagnosis and mo-

- lecular analysis on imported *Plasmodium ovale curtisi* and *P. ovale wallikeri* malaria cases from West and South Africa during 2013-2016. *Korean J Parasitol* 2020; 58: 61-65. <https://doi.org/10.3347/kjp.2020.58.1.61>
11. Kulldorff M, Nagarwalla N. Spatial disease clusters: detection and inference. *Stat Med* 1995; 14: 799-810. <https://doi.org/https://doi.org/10.1002/sim.4780140809>
 12. Mathes RW, Lall R, Levin-Rector A, Sell J, Paladini M, Konty KJ, Olson D, Weiss D. Evaluating and implementing temporal, spatial, and spatio-temporal methods for outbreak detection in a local syndromic surveillance system. *PLoS One* 2017; 12: e0184419. <https://doi.org/10.1371/journal.pone.0184419>
 13. Stelling J, Yih WK, Galas M, Kulldorff M, Pichel M, Terragno R, Tuduri E, Espetxe S, Binsztein N, O'Brien TF, Platt R. Automated use of WHONET and SaTScan to detect outbreaks of *Shigella* spp. using antimicrobial resistance phenotypes. *Epidemiol Infect* 2010; 138: 873-883. <https://doi.org/10.1017/S0950268809990884>
 14. Kulldorff M. A spatial scan statistic. *Commun Stat Theory Methods* 1997; 26: 1481-1496. <https://doi.org/10.1080/03610929708831995>
 15. Chen J, Roth RE, Naito AT, Lengerich EJ, MacEachren AM. Geovisual analytics to enhance spatial scan statistic interpretation: an analysis of U.S. cervical cancer mortality. *Int J Health Geogr* 2008; 7: 57. <https://doi.org/10.1186/1476-072X-7-57>
 16. Azage M, Kumie A, Worku A, Bagtzoglou AC. Childhood diarrhea exhibits spatiotemporal variation in Northwest Ethiopia: a SaTScan spatial statistical analysis. *PLoS One* 2015; 10: e0144690. <https://doi.org/10.1371/journal.pone.0144690>
 17. Park SH, Jegal S, Ahn SK, Jung H, Lee J, Na BK, Hong SJ, Bahk YY, Kim TS. Diagnostic performance of three rapid diagnostic test kits for malaria parasite *Plasmodium falciparum*. *Korean J Parasitol* 2020; 58: 147-152. <https://doi.org/10.3347/kjp.2020.58.2.147>
 18. Organization WH. Synopsis of the world malaria situation, 1979. *Wkly Epidem Rec* 1981; 56: 145-149. <https://apps.who.int/iris/handle/10665/223485>
 19. Iwagami M, Hwang SY, Fukumoto M, Hayakawa T, Tanabe K, Kim SH, Kho WG, Kano S. Geographical origin of *Plasmodium vivax* in the Republic of Korea: haplotype network analysis based on the parasite's mitochondrial genome. *Malar J* 2010; 9: 184. <https://doi.org/10.1186/1475-2875-9-184>
 20. Collins WE, Jeffery GM. *Plasmodium malariae*: parasite and disease. *Clin Microbiol Rev* 2007; 20: 579-592. <https://doi.org/10.1128/CMR.00027-07>
 21. Li P, Zhao Z, Xing H, Li W, Zhu X, Cao Y, Yang Z, Sattabongkot J, Yan G, Fan Q, Cui L. *Plasmodium malariae* and *Plasmodium ovale* infections in the China-Myanmar border area. *Malaria J* 2016; 15: 557. <https://doi.org/10.1186/s12936-016-1605-y>
 22. Kim G, Hong HL, Kim SY, Lee HR, Kim DG, Park S, Shin HS, Chin BS, Kim Y. Mixed infection with *Plasmodium falciparum* and *Plasmodium ovale* in a returned traveller: the first case in Korea. *J Korean Med Sci* 2019; 34: 23-23. <https://doi.org/10.3346/jkms.2019.34.e23>

