

Discovery of key molecules involved in severe malaria

– A new target for malaria vaccine –

❖ Points

- Malaria, one of the world's three major infectious diseases alongside tuberculosis and AIDS, has been reported to affect approximately 300 million people every year, accounting for about 500,000 deaths, and there is still no effective vaccine.
- We found that proteins called RIFIN expressed on erythrocytes infected with *Plasmodium falciparum* (*P. falciparum*) help the parasite to suppress the host immune response, causing severe malaria.
- These findings are expected to contribute to the development of effective vaccines and therapeutic drugs against malaria.

❖ Summary

Fumiji Saito (Research fellow), Kouyuki Hirayasu (Assistant Professor), Hisashi Arase (Professor) (Osaka University, Japan) and others have revealed a novel molecular mechanism that *P. falciparum* suppresses host's immune response and causes severe malaria. This result is expected to greatly contribute to the development of therapeutic drug and vaccine against malaria.

❖ Background

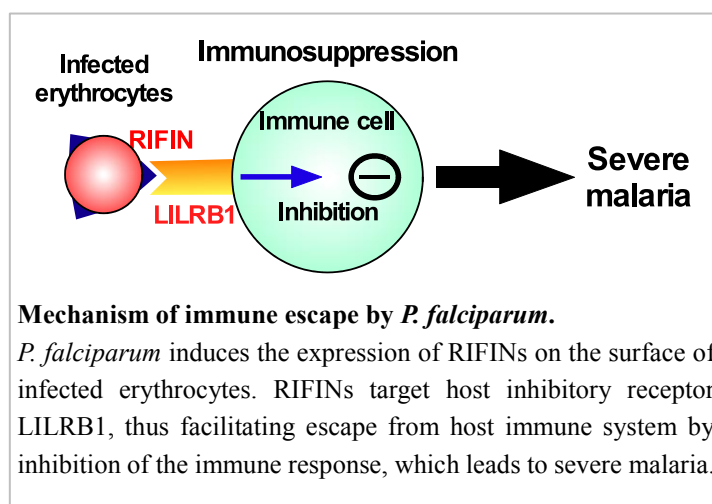
Malaria^{*1} is one of three major infectious diseases^{*2} affecting approximately 300 million people every year, accounting for about 500,000 deaths, but effective vaccine development has not been successful. Among malaria parasites infecting humans, *P. falciparum*^{*3} causes especially severe disease. In addition, acquired immunity to malaria is inefficient, even after repeated exposures to *P. falciparum*, but the immune regulatory mechanisms used by *P. falciparum* remain largely unclear. Therefore, malaria parasites appear to have a mechanism to escape our immune system.

❖ Findings

Malaria parasites infect mainly erythrocytes in the host and proliferate within infected erythrocytes. The research group found that proteins called RIFIN^{*4} expressed on *P. falciparum*-infected erythrocytes bind to a host inhibitory receptor LILRB1^{*5}. Furthermore, RIFIN suppresses the immune response to malaria, resulting in severe complications of malaria (upper figure).

❖ Significance of the findings

This research disclosed for the first time in the world that *P. falciparum* has a new mechanism to suppress the host immune response by using an inhibitory receptor, contributing to the pathogenesis of severe malaria. The results of this research are expected to greatly contribute to the development of therapeutic drug and vaccine against malaria.



❖ Technical terms

***1 : Malaria**

Infectious diseases caused by malaria parasites

***2 : Three major infectious diseases**

Malaria, tuberculosis, and HIV/AIDS

3 : *Plasmodium falciparum

Malaria parasites infecting humans, and causing the most severe complications of malaria

***4 : RIFIN**

RIFIN proteins are encoded by the *rif* (repetitive interspersed family) genes of *P. falciparum*. There are about 150 *rif* genes per parasite genome. However, their functions have been still unclear.

***5 : LILRB1**

One of the immune inhibitory receptors that suppress the activation of immune cells and prevent autoimmune responses by recognizing self-molecules like major histocompatibility complex (MHC) class I. Human cytomegalovirus is also known to have a viral MHC class I-like molecule (UL18) that suppresses the immune response via LILRB1 for immune escape.

❖ Article information

Title: Immune evasion of *Plasmodium falciparum* by RIFIN via inhibitory receptors

Authors: Fumiji Saito*, Kouyuki Hirayasu*, Takeshi Satoh, Christian W. Wang, John Lusingu, Takao Arimori, Kyoko Shida, Nirianne Marie Q. Palacpac, Sawako Itagaki, Shiroh Iwanaga, Eizo Takashima, Takafumi Tsuboi, Masako Kohyama, Tadahiro Suenaga, Marco Colonna, Junichi Takagi, Thomas Lavstsen, Toshihiro Horii, Hisashi Arase

*These authors equally contributed to this work.

Journal: Nature (Nov, 29, 2017, London)

❖ Fund

This research was supported by Japanese Initiative for Progress of Research on Infectious Disease for global Epidemic funded by the Japan Agency for Medical Research and Development (AMED).