

## **Researchers discover new set of channels connecting malaria parasite and blood cells**

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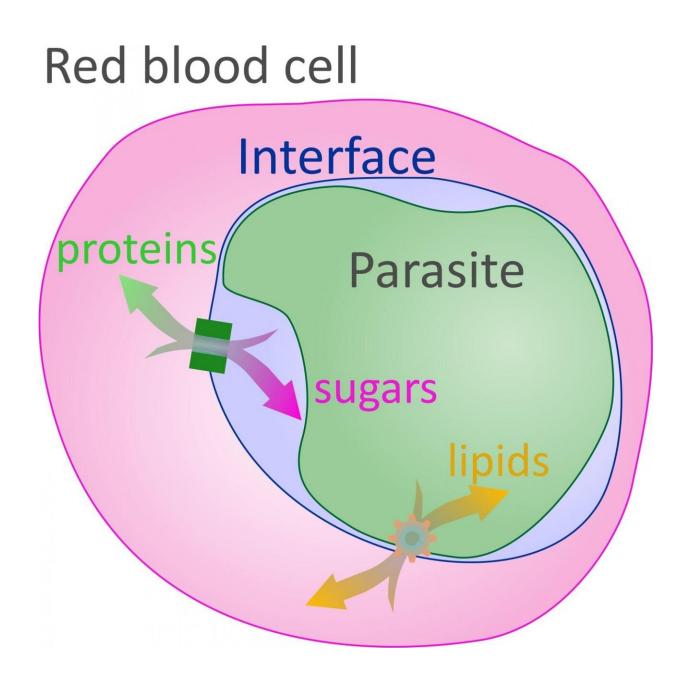




Diagram of the two membrane channels between the malaria parasite vacuole and the red blood cell. Credit: National Institutes of Health

Researchers at the National Institutes of Health and other institutions have discovered another set of pore-like holes, or channels, traversing the membrane-bound sac that encloses the deadliest malaria parasite as it infects red blood cells. The channels enable the transport of lipids—fatlike molecules—between the blood cell and parasite, Plasmodium falciparum. The parasite draws lipids from the cell to sustain its growth and may also secrete other types of lipids to hijack cell functions to meet its needs.

The finding follows an earlier discovery of another set of channels through the membrane enabling the two-way flow of proteins and nonfatty nutrients between the parasite and <u>red blood cells</u>. Together, the discoveries raise the possibility of treatments that block the flow of nutrients to starve the parasite.

The research team was led by Joshua Zimmerberg, M.D., Ph.D., a senior investigator in the Section on Integrative Biophysics at NIH's Eunice Kennedy Shriver National Institute of Child Health and Human Development (NICHD). The study appears in *Nature Communications*.

In 2018, there were 228 million cases of malaria worldwide, leading to more than 400,000 deaths, 67% of which were among children under 5, according to the World Health Organization. In the current study, researchers determined that the channels through the sac, or vacuole, that encloses the parasite are made of Niemann-Pick C1-related protein (PfNCR1). The PfNCR1 channels are restricted to locations where the vacuole membrane touches the parasite's membrane. The channels the team discovered in the previous study are formed by exported protein 2



(EXP2). Areas of the vacuole membrane containing EXP2 are located far from the parasite's membrane, at an average distance of 20 to 40 nanometers. The researchers believe that the parasite may use this variation in distance to separate the two transport systems.

**More information:** Matthias Garten et al, Contacting domains segregate a lipid transporter from a solute transporter in the malarial host–parasite interface, *Nature Communications* (2020). <u>DOI:</u> 10.1038/s41467-020-17506-9

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