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1 **Activation of complement-like antiparasitic responses in** *Anopheles* **mosquitoes**

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15 **Abstract**

 $\frac{16}{17}$ During their development in mosquitoes, malaria parasites undergo massive losses that are 18 in part due to a potent antiparasitic response mounted by the vector. The most efficient and
19 best characterized response relies on a complement-like system particularly effective against best characterized response relies on a complement-like system particularly effective against 20 parasites as they cross the mosquito midgut epithelium. While our vision of the molecular 21 and cellular events that lead to parasite elimination is still partial, our understanding of the 22 steps triggering complement activation at the surface of invading parasites has considerably
23 progressed, not only through the identification of novel contributing genes, but also with the progressed, not only through the identification of novel contributing genes, but also with the 24 recent in-depth characterization of the different mosquito blood cell types, and the ability to
25 track them in live mosquitoes. Here, we propose a simple model based on the time of 25 track them in live mosquitoes. Here, we propose a simple model based on the time of 26 invasion to explain how parasites may escape complement-like responses during midgut 27 infection.

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31 31 **Glossary**

 $\frac{32}{33}$ **Hemocytes**: Immune cells in insects. They circulate in the body cavity, bathed in the hemolymph. Some of them are sessile, i.e. stay attached to tissues. In mosquitoes three main types can be distinguished: prohemocytes, oenocytoids and granulocytes. The last type is composed of several subtypes presenting specific transcriptomic signatures, including larger cells called megacytes. They fulfill several essential functions such as phagocytosis and melanisation.

- 39 40 **Hemolymph:** extracellular fluid in insects, analogous to vertebrate blood. Insect have an 41 open circulatory system with hemolymph bathing all tissues and kept in movement by a
42 heart. Beside hemocytes, hemolymph carries important soluble immune factors such as 42 heart. Beside hemocytes, hemolymph carries important soluble immune factors such as
43 antiparasitic TEP1. LRIM1 and APL1C. and antimicrobial peptides. antiparasitic TEP1, LRIM1 and APL1C, and antimicrobial peptides.
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Melanisation: deposition of a dark pigment called melanin. Physiological process playing important roles in insects, including wound healing, clot formation, encapsulation of invading pathogens. It results from the activation of a proteolytic cascade involving prophenoloxidases (produced by oenocytoids) that synthesize melanin.

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Detection of parasite invasion in the mosquito gut

 Plasmodium parasites causing human malaria are transmitted by mosquitoes of the genus *Anopheles*. *P. falciparum* is responsible for the most severe symptoms of the disease, and 54 for most deaths. While feeding on a parasite carrier, mosquitoes ingest infected red blood
55 cells of which only gametocytes, the sexual forms of the parasite, are able to further develop cells, of which only gametocytes, the sexual forms of the parasite, are able to further develop in the vector. Gametocytes become activated in the blood bolus, and release gametes that proceed to mate, forming a zygote that becomes motile and is then called ookinete. Within 18 to 32h, the ookinete traverses the midgut epithelium (Figure 1) and undergoes meiosis. It then resides between the midgut epithelium and the basal lamina, forming a cyst where it goes through multiple rounds of replication for 8-14 days before cytokinesis occurs, giving rise to sporozoites. These egress in batches upon rupture of the oocyst membrane, resulting in the release of thousands of sporozoites in the mosquito open circulatory system. Part of these invade the salivary glands and can then be transmitted to a naive host upon a subsequent blood meal. Still, mosquitoes are not mere needles that transmit parasites, they mount potent immune responses that limit their development, especially during the first days after infection, when most parasite losses occur [1]. After a blood meal, midgut cells secrete a peritrophic matrix, an acellular semi-permeable layer of chitin polymers, to protect cells from direct contact with the blood bolus and proliferating bacteria during digestion. This protection is reinforced by the Duox/IMPer module (Dual oxidase / IMmune heme Peroxidase Hpx15) that catalyzes crosslinking of mucins in the ectoperitrophic space located between the peritrophic matrix and the midgut epithelium, further protecting midgut cells from contact with immune elicitors [2]. Hence, after forming inside the mosquito midgut, ookinetes first 73 traverse the peritrophic matrix before crossing the epithelium (Figure 1). This breaching of mosquito intestinal barriers allows both parasites and bacterial elicitors to come in contact mosquito intestinal barriers allows both parasites and bacterial elicitors to come in contact with the midgut cells, triggering a series of cellular and humoral reactions that ultimately lead to complement activation at the surface of ookinetes.

78 Nitric oxide synthase (NOS), whose expression is rapidly induced in cells that are traversed
79 by ookinetes [3], produces nitric oxide (NO) that can diffuse in nearby tissues and possibly 79 by ookinetes [3], produces nitric oxide (NO) that can diffuse in nearby tissues and possibly
80 act as a signaling molecule. Locally, NO and derivatives (such as nitrites) can also react with act as a signaling molecule. Locally, NO and derivatives (such as nitrites) can also react with H_2O_2 produced by the NADPH oxidase NOX5 and heme peroxidase Hpx2 whose expression 82 is upregulated in infected midguts, leading to protein nitration at the site of parasite invasion is upregulated in infected midguts, leading to protein nitration at the site of parasite invasion [4] (Figure 2). This modification has been evidenced in some midgut cells that were traversed by parasites [5] and was proposed to affect the parasite surface, making it visible to the complement-like system [4]. However, whether the parasite surface itself is a target of nitration remains to be demonstrated experimentally. In addition, parasites traverse the midgut epithelium within minutes [6], often invading several cells in a row before exiting and resting below the basal lamina, which is not compatible with the time required for their 89 nitration. Indeed, the upregulation of NOS, NOX5 and Hpx2 activity in invaded cells likely
80 takes several hours, which is consistent with the observation that nitro-tyrosines are usually 90 takes several hours, which is consistent with the observation that nitro-tyrosines are usually
91 detected only in the first invaded cell of a series at 24h post infection [5] and suggests protein 91 detected only in the first invaded cell of a series at 24h post infection [5] and suggests protein
92 nitration occurs long after the parasite has traversed the midgut cells (Figure 1). Alternatively. nitration occurs long after the parasite has traversed the midgut cells (Figure 1). Alternatively, nitration of midgut cells could be detected as a danger signal by patrolling **hemocytes** that would activate the complement system, or even represent a mark of extensive cellular damage with no specific role in the antiparasitic response. In the latter case, the fact that NOX5 and Hpx2 are essential for an efficient antiparasitic response [4] would indicate that 97 they may contribute to it in a different manner. For instance, parasite infection or bacteria
98 ingestion also trigger secretion of prostaglandins (PGs) by midgut cells, and silencing of 98 ingestion also trigger secretion of prostaglandins (PGs) by midgut cells, and silencing of 99 Hox7 or Hox8 decreases PG release, suggesting these enzymes are involved in PG Hpx7 or Hpx8 decreases PG release, suggesting these enzymes are involved in PG synthesis [7] (Figures 1, 2). It would be interesting to investigate the contribution of NOX5, Hpx2 and other induced Hpxs in PG synthesis. Among PGs, prostaglandin E2 seems to have the strongest effect on parasite infection: (1) it contributes to recruit **hemocytes** to the infected midgut, (2) it signals to fat body cells to produce a **hemocyte** differentiating factor (HDF) that is composed of a lipid, lipoxin A4, bound to Evokin, a proteic lipid carrier [6–8].

Hemocytes, key actors of the TEP1-mediated response

 Hemocytes are immune cells of insects that circulate in the **hemolymph**, the equivalent of vertebrate blood. Single cell transcriptome profiling has identified two main **hemocyte** lineages in *Anopheles gambiae*, the oenocytoids expressing prophenoloxidases that are involved in the **melanisation** cascade, and several granulocyte subtypes including large megacytes and expressing distinct sets of effector genes [9,10]. Both lineages are likely derived from prohemocyte precursors [10].

113
114 114 Upon infection, midgut cells increase their secretion of PGE2, a strong and rapid attractant to
115 hemocytes that also enhances their mobility. Impressive *in vivo* images of granulocytes hemocytes that also enhances their mobility. Impressive *in vivo* images of granulocytes patrolling around the site of infection have been obtained [6]. Importantly, some granulocytes attracted to the parasite- or the bacteria-infected midgut, undergo apoptosis and release **hemocyte**-derived microvesicules (HdMv) that are strongly associated with parasite-infected midgut cells [6,11] (Figure 1). Uptake of synthetic beads by **hemocytes** reduces the number 120 of midgut-associated HdMvs, but also the labeling of parasites by the key antiparasitic 121 effector ThioEster-containing Protein 1 (TEP1) (see below) [11], the later phenotype being 121 effector ThioEster-containing Protein 1 (TEP1) (see below) [11], the later phenotype being
122 also observed upon hemocyte depletion by clodronate liposomes [12]. As **hemocytes** do not also observed upon hemocyte depletion by clodronate liposomes [12]. As **hemocytes** do not produce large quantities of TEP1 [13], it is tempting to speculate HdMvs may instead deposit factors that are essential for TEP1 activation on the microbial surface. It would be interesting to determine the exact content of HdMvs by mass spectrometry.

 $\frac{126}{127}$

At later time points, the PGE2-induced and fat-body derived HDF triggers major changes in circulating **hemocytes**, notably their proliferation and an increase in the proportion of 129 circulating oenocytoids and granulocytes [8] (Figure 1). These changes likely occur too late
130 to have an effect on parasite development during a primary infection, but they are longto have an effect on parasite development during a primary infection, but they are long- lasting and provide extended protection to mosquitoes during subsequent infections or upon **hemolymph** transfer from previously infected mosquitoes, a phenomenon termed immune priming [8].

Parasite labeling and elimination mediated by complement-like protein TEP1

 The mosquito complement system is composed of soluble proteins that are mainly produced by the fat body [13] and circulate in the **hemolymph**. Its central component is TEP1, a 140 structural and functional homologue of vertebrate complement factor C3 [14]. Like C3, TEP1
141 opsonizes the surface of invading microorganisms (bacteria, parasites, fungi), marking them opsonizes the surface of invading microorganisms (bacteria, parasites, fungi), marking them for lysis and/or melanotic encapsulation [1]. TEP1 circulates constitutively in the **hemolymph** as a full-length and an activated, cleaved form. The protease(s) responsible for TEP1 cleavage are not known, however stability of the circulating cleaved form requires a complex 145 comprising two leucine-rich repeat proteins (LRRs), LRIM1 and APL1C. Deposition of TEP1
146 on foreign surfaces is mediated by a covalent thioester bond and must be tightly regulated, on foreign surfaces is mediated by a covalent thioester bond and must be tightly regulated, as experimental depletion of LRIM1 or APL1 triggers uncontrolled deposition on self-tissues 148 [15]. According to current models [1], the accumulation of TEP1 on the ookinete surface is 149 initiated by precocious, rapid binding of small amounts of circulating cleaved TEP1. In a initiated by precocious, rapid binding of small amounts of circulating cleaved TEP1. In a second and slower step, TEP1 deposition is amplified by mobilizing and processing the pool of circulating full length TEP1. This depends on the recruitment at the target surface of 152 SPCLIP1 (CLIPA30), a serine protease homolog that is not catalytically active. Therefore,
153 additional protease(s) are likely necessary to cleave further TEP1 molecules and expose 153 additional protease(s) are likely necessary to cleave further TEP1 molecules and expose
154 their thioester site, that can then react with nearby surfaces. The protein complex necessary their thioester site, that can then react with nearby surfaces. The protein complex necessary for this cascade of events, comprising some TEP1cut associated to (or freed from) LRRs; SPCLIP1; and unknown proteases(s), is called TEP1 convertase complex. An attractive hypothesis is that HdMvs deliver factors that potentiate, or take part in, this convertase. 158 Notably, genetic polymorphism in the *TEP1* locus contributes to determine parasite killing
159 efficiency, with the *TEP1^{*}R* allele being more efficient to kill *P. berghei* parasites [16]. efficiency, with the *TEP1*R* allele being more efficient to kill *P. berghei* parasites [16].

 Oocysts are relatively protected from TEP1 attack. Indeed, once formed, they undergo limited losses despite their long development time: over 10 days, only few oocysts are labeled with TEP1 and their elimination is TEP1-independent [17,18]. It is possible that oocyst proteins such as Cap380 [19], but also the circumsporozoite protein CSP [20], act as a shield, or as complement inhibitors, on the oocyst surface. In addition, the mosquito basal lamina under which oocysts reside could also provide some protection from soluble immune factors. Indeed, in the case of *P. berghei* but not *P. falciparum*, a subsequent blood meal 4 days after infection stretches the midgut basal lamina and makes it more permeable to **hemolymph** factors, leading to increased TEP1 labeling and elimination of oocysts [21].
169 These results suggest that P. falciparum oocysts are further protected from TEP1 activation. 169 These results suggest that *P. falciparum* oocysts are further protected from TEP1 activation.
170 Finally, despite their release in the **hemolymph** with no apparent protection, sporozoites are Finally, despite their release in the **hemolymph** with no apparent protection, sporozoites are 171 not labeled with TEP1, possibly because they are likewise shielded by CSP, a protein that 172 also protects them from melanisation [22]. also protects them from melanisation [22].

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175 **Parasite success in crossing the midgut epithelium: a question of time?**

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177 177 *Plasmodium* and *Anopheles* are engaged in a co-evolution arm race to circumvent each 178 other's defense mechanisms. We propose a simple model, derived from the "time bomb
179 theory" [3], where invasion speed rather than inhibition of mosquito immunity during midgut theory" [3], where invasion speed rather than inhibition of mosquito immunity during midgut 180 traversal, may be key to *Plasmodium* success. This model predicts that the mosquito 181 immune system is activated by an infectious blood meal, however with a certain delay, 182 offering ookinetes a short window of time to invade and transform into protected oocysts. 183 Published data can be explained under this light. (i) P47, an ookinete surface protein, exists 184 as different alleles in P. falciparum that determine escape of the mosquito complement 184 as different alleles in *P. falciparum* that determine escape of the mosquito complement 185 system depending on the presence of a matching Pfs47 receptor, P47rec, on mosquito 186 midgut cells, this receptor being itself variable among *Anopheles* species [23–25]. Pfs47 is 187 thought to suppress activation of the JNK defense pathway, resulting in increased nitration 188 and parasite recognition by TEP1. However, artificially boosting JNK signaling by silencing a 189 negative regulator has no effect on *P. falciparum* [26]. An attractive alternative explanation is 190 that the Pfs47 / P47rec physical interaction may simply promote entry and traversal of the 191 parasite. speeding up the process. In agreement with this, P47rec aggregates at the site of 191 parasite, speeding up the process. In agreement with this, P47rec aggregates at the site of parasite entry [25], and Pfs47 belongs to the family of 6-cystein domain proteins comprising parasite entry [25], and Pfs47 belongs to the family of 6-cystein domain proteins comprising 193 notably the surface protein P36 expressed by sporozoites and mediating interactions with 194 CD81 and SR-BI, two liver cell receptors used by sporozoites to penetrate hepatocyte [27]. 195 We propose that if Pfs47 does not match its receptor, or if one of the two is absent or 196 expressed at low levels, traversal would be slower using alternative receptors, rendering 196 expressed at low levels, traversal would be slower using alternative receptors, rendering
197 ookinetes more vulnerable to the TEP1-dependent response, especially to the very efficient 197 ookinetes more vulnerable to the TEP1-dependent response, especially to the very efficient
198 TEP1*R allele. Of note, the increased JNK activation and nitration observed upon infection 198 *TEP1*R* allele. Of note, the increased JNK activation and nitration observed upon infection
199 with *Pfs47-KO* parasites [26] may be a consequence rather than a cause of the increased 199 with *Pfs47-KO* parasites [26] may be a consequence rather than a cause of the increased
200 parasite killing, or a consequence of prolonged midgut invasion. (ii) An alternative receptor 200 parasite killing, or a consequence of prolonged midgut invasion. (ii) An alternative receptor 201 could be the PIMMS43 ookinete surface protein whose depletion also sensitizes parasites to 201 could be the PIMMS43 ookinete surface protein whose depletion also sensitizes parasites to
202 elimination by the mosquito complement system [28]. (iii) The different temperatures at which 202 elimination by the mosquito complement system [28]. (iii) The different temperatures at which 203 infections with *P. falciparum* and *P. berghei* are taking place (26 vs. 21°C, respectively), and possibly less efficient Pbs47 and P47Rec combinations (not co-evolved), could also lead to 205 slower midgut traversal by *P. berghei* and explain why *P. falciparum* shows little sensitivity to 206 the complement pathway in contrast to *P. berghei*. Consistently with this, decreasing 207 temperature from 27 to 19ºC for *P. falciparum* infection elevates ookinetes susceptibility to 208 **melanisation** in *CTL4*-knockout mosquitoes [29]. In addition, temperature differences likely also affect midgut microbiota and mosquito immune reactions. (iv) Transgenic expression in 210 mosquito midgut cells of the Magainin peptide, decreasing mitochondrial activity and 211 therefore ATP production on which ookinete movement crucially depends, decreases oocyst 212 formation [30]. The two latter observations are consistent with increased vulnerability of 213 parasites to mosquito defense mechanisms when their speed is decreased, either by a drop 214 in temperature or by synthetic depletion of energy. In addition, delays in ookinete to oocyst 215 transformation are detrimental for oocyst development after both Magainin expression and
216 PIMMS43 knockout, suggesting that slow ookinete traversal could also result in long-lasting PIMMS43 knockout, suggesting that slow ookinete traversal could also result in long-lasting parasite growth defects.

Conclusion

223 Here we sought to propose an alternative, simple model to explain how parasites may
224 escape mosquito immune responses upon midgut invasion, which does not necessarily 224 escape mosquito immune responses upon midgut invasion, which does not necessarily
225 exclude other models positing that parasites actively dampen the antiparasitic response. In exclude other models positing that parasites actively dampen the antiparasitic response. In 226 our view, as in an earlier model called the time bomb theory, the speed of parasite invasion
227 is crucial for parasites to escape immune attack by mosquito soluble factors. However, rather is crucial for parasites to escape immune attack by mosquito soluble factors. However, rather than escaping nitration during midgut cell traversal, speed would allow parasites to timely reach the basal side and transform into oocysts, before hemocytes are activated and 230 stimulate parasite labeling with TEP1 which promotes their subsequent elimination.
231 Temperature during the invasion process may act as a first important determinant of parasite 231 Temperature during the invasion process may act as a first important determinant of parasite
232 successful escape, with faster and more successful invasion at higher temperature. 232 successful escape, with faster and more successful invasion at higher temperature.
233 Receptors on the parasite surface, such as Pfs47 and PIMMS43, if efficiently interacting with Receptors on the parasite surface, such as Pfs47 and PIMMS43, if efficiently interacting with their mosquito partner (a process dependent on genetic diversity in both organisms) would also speed up the invasion process and promote maximal ookinete survival. Live imaging of midgut invasion with P*. falciparum* at different temperatures to visualize and count ookinetes that have crossed the epithelium as a function of time, ideally *in vivo* and concomitant with 238 the tracking of fluorescence-expressing hemocytes, would shed more light on these complex
239 parasite-vector interactions. For this, genetic modifications that decrease cuticle parasite-vector interactions. For this, genetic modifications that decrease cuticle pigmentation, combined with the expression of fluorescent reporters in specific cell lineages or tissues, will be key to optimize visualization of mosquito/parasite interactions at the cellular level [31]. Should nitration of invaded midgut cells have no specific role in the antiparasitic response, other functions of NOX5 and Hpx2, e.g. in prostaglandin signaling and hemocyte 244 activation, should be investigated. Finally, delaying midgut invasion by ookinetes may be
245 considered as a promising transmission blocking strategy for instance by interfering with considered as a promising transmission blocking strategy, for instance by interfering with P47 [32,33] and PIMMS43 interactions with their respective receptors, ideally both at the 247 same time, through human vaccination against these receptors, or in genetically modified mosquitoes producing single chain antibodies targeting them in the midgut lumen.

$\frac{250}{251}$ **Conflict of interest statement**

We declare no conflict of interest.

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Figures & Legends

Figure 1

 Figure 1. How *Plasmodium* **invasion speed could determine evasion from the TEP1- dependent antiparasitic response in** *A. gambiae***.** Within the first 18h post infection, ookinetes breach through the peritrophic matrix and come in contact with the midgut 273 epithelium. Efficient entry into midgut cells is promoted by interactions of parasites receptors (e.g. P47, PIMMS43) with midgut cells apical surface proteins (e.g. AgP47Rec). By 24-32 hpi, bacteria and parasite elicitors contact midgut cells and trigger a massive immune response. The release of prostaglandins (PG) by invaded midgut cells recruits hemocytes to 277 the site of infection and induces the secretion of a hemocyte differentiation factor (HDF) by fat body cells. At the site of infection, granulocytes release hemocyte derived microvesicules (HdMvs), that could promote TEP1 binding (green dots on parasite surface) to late ookinetes that have not yet reached the less sensitive oocyst stage. Some cells, damaged by parasite traversal, undergo massive nitration (indicated in red) and will be extruded towards the midgut lumen. At later time points, HDF is responsible for an increase in the number of circulating hemocytes, notably granulocytes and oenocytoids, which will be protective upon a subsequent infection.

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Figure 2

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301 **Figure 2.** Processes induced in midgut cells damaged by parasite invasion. Ookinete 302 invasion of a midgut cell induces the expression of several genes encoding: (1) the nitric
303 oxide synthase (NOS) producing NO that can diffuse to nearby cells; (2) the NADPH oxidase 303 oxide synthase (NOS) producing NO that can diffuse to nearby cells; (2) the NADPH oxidase
304 NOX5 and heme peroxidase Hpx2, producing H₂O₂. Reaction of NO with H₂O₂ can lead to 304 NOX5 and heme peroxidase Hpx2, producing H₂O₂. Reaction of NO with H₂O₂ can lead to 305 massive nitration. (3) Heme peroxidases Hpx7, Hpx8 are induced as well, contributing the 305 massive nitration. (3) Heme peroxidases Hpx7, Hpx8 are induced as well, contributing the
306 synthesis of prostaglandins (PG). Whether Hpx2 is also involved in PG production remains to synthesis of prostaglandins (PG). Whether Hpx2 is also involved in PG production remains to 307 be tested.

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