Title: Kinetics of *Plasmodium* midgut invasion in *Anopheles* mosquitoes

Short Title: Live imaging of *Plasmodium* invasion of mosquito midguts

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Abstract

Malaria-causing *Plasmodium* parasites traverse the mosquito midgut cells to establish infection at the basal side of the midgut. This dynamic process is a determinant of mosquito vector competence, yet the kinetics of the parasite migration is not well understood. Here we used transgenic mosquitoes of two *Anopheles* species and a *Plasmodium berghei* fluorescence reporter line to track parasite passage through the mosquito tissues at high spatial resolution. We provide new quantitative insight into malaria parasite invasion in African and Indian *Anopheles* species and demonstrate that species-specific kinetics of *Plasmodium* invasion is shaped by the mosquito complement-like system.

Author Summary

The traversal of the mosquito midgut cells is one of the critical stages in the life cycle of malaria parasites. Motile parasite forms, called ookinetes, traverse the midgut epithelium in a dynamic process which is not fully understood.

Here, we harnessed transgenic reporters to track invasion of *Plasmodium* parasites in African and Indian mosquito species. We found important differences in parasite dynamics between the two anopheline species and demonstrated an unexpected role of mosquito complement-like system in regulation of parasite invasion.

47 # Introduction

48 Malaria is a vector-borne human infectious disease caused by protozoan
49 parasites of *Plasmodium* species. It is widespread in tropical and subtropical
50 regions, including parts of the Americas, Asia, and Africa. Approximately 200
51 million annual cases of malaria result in half a million deaths [1]. Malaria-
52 causing *Plasmodium* parasites are transmitted by *Anopheline* mosquitoes.
53 Among more than 400 of known *Anopheles* species, only 40 are vectors of
54 human malaria [2].

55 *Plasmodium* development in the mosquito begins with the ingestion of red
56 blood cells infected with sexual-stage gametocytes. In the mosquito midgut,
57 gametocytes differentiate into gametes that egress from the red blood cells and
58 fuse to form the zygotes that develop into motile ookinetes within 16-18 h. The
59 ookinetes penetrate the midgut epithelium 18 – 26 h after the infectious blood
60 meal and transform into vegetative oocysts on the basal side of the midgut [3].
61 After 12–14 days, mature oocysts rupture and release thousands of sporozoites
62 into the mosquito hemocoel. Released sporozoites invade the salivary glands,
63 where they reside inside the salivary ducts to be injected into a new host when
64 the infected mosquito feeds again [4].

65 The passage of the malaria parasite through the mosquito vector is
66 characterized by a major population bottleneck. Previous studies revealed that
67 mosquitoes kill the majority of invading *Plasmodium* parasites (reviewed by
68 [5,6]), predominantly during the ookinete stage at the basal side of the
69 epithelium [7].

70 The immune response of mosquitoes to *Plasmodium* parasites is multifaceted
71 and involves multiple processes. In the midgut, reactive oxygen and nitrogen

72 species, hemoglobin degradation products, as well as digestive enzymes and
73 ´bacterial flora, all affect the rate of *Plasmodium* development (Reviewed in [8]).
74 As parasites traverse midgut epithelial cells, the invaded cells produce high
75 levels of nitric oxide synthase and peroxidases, creating a toxic environment
76 for the parasites. As a result, some parasites undergo nitration which marks
77 them for killing by the mosquito complement-like system [9]. Furthermore,
78 intracellular parasites can trigger apoptosis causing extrusion and clearance of
79 invaded cells from the cellular layer into the midgut lumen [10]. As *Plasmodium*
80 tries to evade reactive oxygen and nitrogen species inside the cells, these toxic
81 molecules may shape the path taken by the parasite through the cellular layer.
82 When the surviving parasites finally reach the basal lamina, they encounter
83 soluble immune factors that circulate in the hemolymph. Complement-like
84 proteins TEP1 and leucine-rich repeat proteins APL1 and LRIM1 form a
85 complex that mediates parasite killing [11,12]. Histological studies have shown
86 that parasites crossing the cellular layer can be found both inside and in
87 between midgut cells [3,13]. However, it is not yet known whether some
88 parasites cross the cellular layer exclusively between cells, thus avoiding
89 nitration and subsequent recognition by TEP1.
90 Despite accumulating evidence of molecular processes that govern the
91 passage of motile ookinetes through mosquito tissues, the complexity and
92 diversity of this dynamic process remains to be deciphered. Three modes of
93 motility were reported for the invading ookinetes, namely spiraling, gliding and
94 stationary rotation [14][15]. Spiraling and gliding movements result in active
95 displacement of the parasite in space. In contrast, stationary rotation movement
96 was observed for prolonged periods of time and resulted in no displacement of

97 the ookinete. Because of the lack of markers of the entire midgut cellular layer,
98 previous studies did not establish how distinct types of movements correlate
99 with ookinete location in the midgut.
100 It has been previously demonstrated that *Anopheles* species differ in their
101 vector competence [16]. In the laboratory, *Anopheles stephensi* (*As*) and
102 *Anopheles gambiae (Ag)* can be infected with the murine parasite *Plasmodium*
103 *berghei*, albeit at different rates [17]. We set out to image *in vivo* migration of
104 the RFP-expressing *P. berghei* (Pb) ookinetes through the epithelial cells in
105 these two genetically-modified mosquito species that express GFP in the
106 midgut cells. Using high-speed spinning disk microscopy and automated image
107 analyses, we quantified parasite invasion dynamics at high spatial and temporal
108 resolution. Our data revealed unexpected differences in invasion of closely-
109 related mosquito species, pointing to important species-specific mechanisms
110 that regulate mosquito – parasite interactions. Moreover, silencing of the major
111 component of the mosquito complement-like system affected the parasite
112 invasion dynamics, suggesting a new function of TEP1 at the early stages of
113 the midgut invasion process.

114 Results and discussion

115 **TEP1 inhibits midgut invasion of *P. berghei* ookinetes.**

116 To study the passage of *Pb* ookinetes through the mosquito midgut, we
117 combined multiscale imaging techniques with high-throughput data analysis
118 and mining (Fig 1). We used transgenic mosquitoes expressing GFP under
119 midgut-specific promoters [18,19] to label mosquito midgut cells, and
120 transgenic rodent *Pb* parasites expressing RFP under a constitutive promoter

[20] (S1a Fig). We first made sure that expression of the reporters did not interfere with *Plasmodium* infection. As expected, a significant difference was observed in infection intensity between *As* and *Ag*. Regardless of the infection levels, *As* developed significantly higher oocysts numbers than *Ag* (S1b Fig). We concluded that the transgenic mosquito and *Pb* lines can be used for *in vivo* imaging.

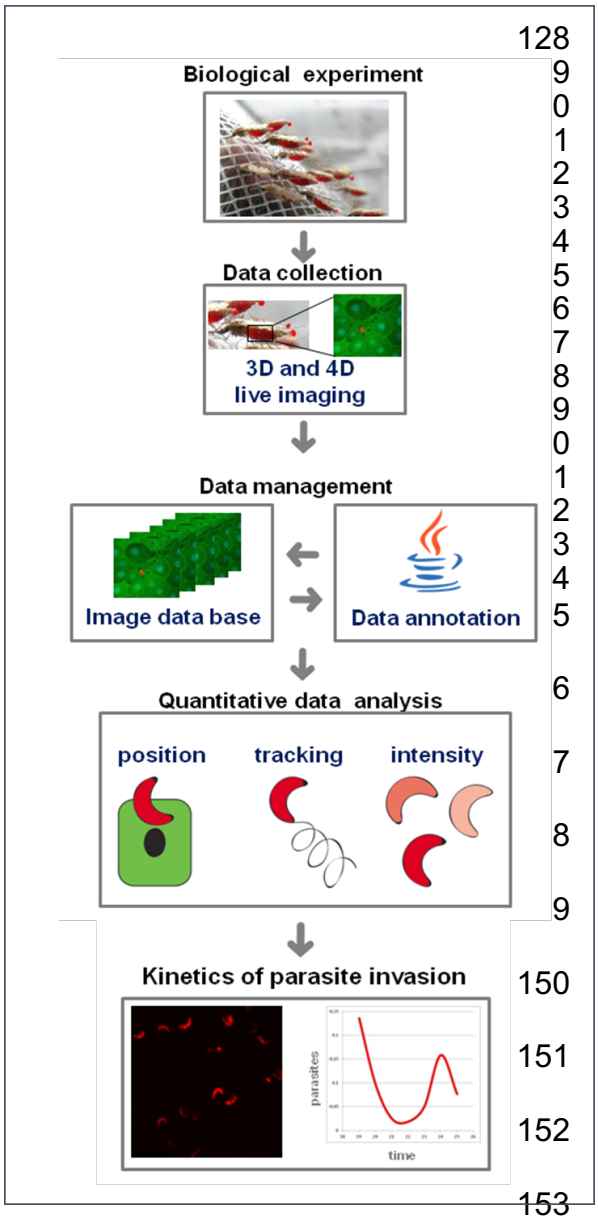

Figure 1. Workflow and experimental settings. *A. stephensi* (*As*) and *A. gambiae* (*Ag*) mosquitoes were blood fed on *P. berghei* infected mice, their midguts dissected and visualized using fast confocal microscopy. Images from all experiments collected at different time points after infection were uploaded into an image database and annotated. Quantitative data was extracted from the images in the database regarding the number, position and intensity of visualized parasites. The results of the data analysis reveal the kinetics of parasite invasion.

As the transgenic mosquito lines expressed GFP in the entire midgut cell, we measured the exact position of RFP-expressing parasites relative to the cellular

156 layer (Fig 2). To this end, we collected large series of z-stack images of live
157 parasites inside the dissected mosquito midguts at different time points after
158 infection and time-lapse images of selected parasites. These tools enabled us
159 to study the parasite invasion process at two time-scales: one was based on
160 statistical analysis of parasites in three dimensional (3D) snapshots of the state
161 of infection between 18 and 25 h post infection, the second tracked single
162 parasites 18 to 25 h post infection over a time of 20 min to 2 h.

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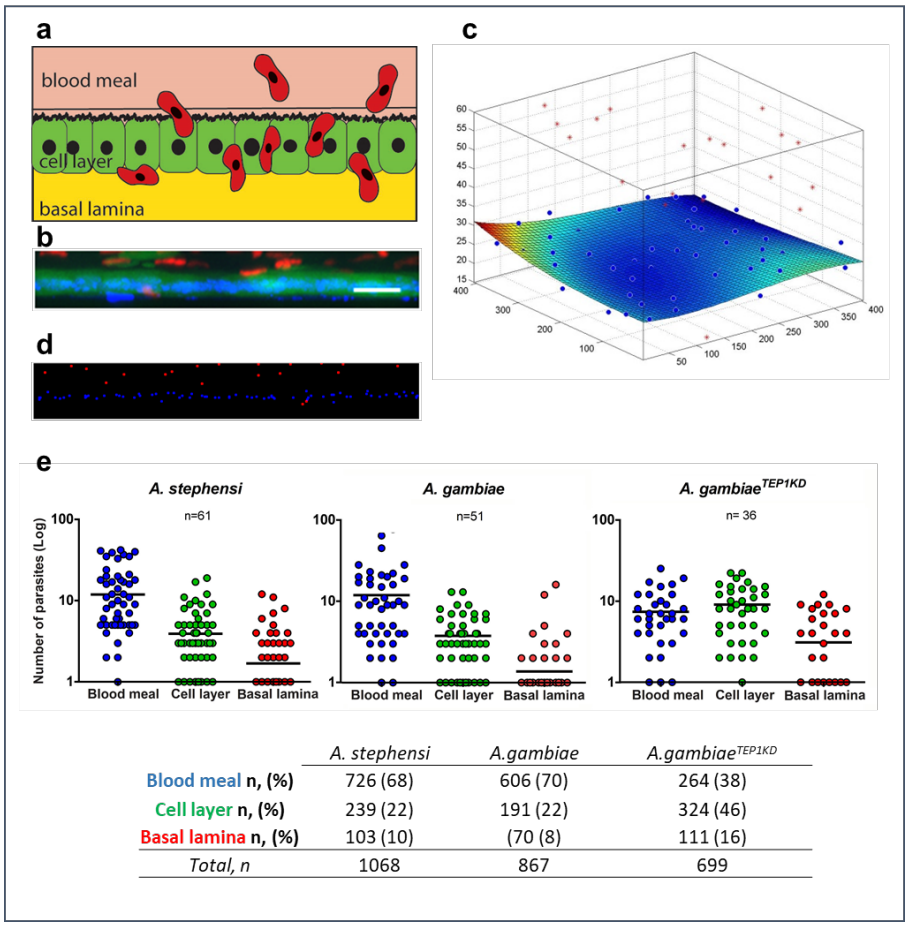


	A. stephensi	*A.gambiae*	*A.gambiae*^{TEP1KD}
Blood meal n, (%)	726 (68)	606 (70)	264 (38)
Cell layer n, (%)	239 (22)	191 (22)	324 (46)
Basal lamina n, (%)	103 (10)	(70 (8)	111 (16)
Total, n	1068	867	699

164

165

166 **Figure 2. Positions of the parasites relative to the midgut cells.** **a**. Schematic
167 representation of topology in the mosquito midgut. Motile ookinetes (red) traverse the
168 mosquito midgut cells and establish infection on the basal side under the basal lamina.
169 **b**. A representative projection of a cross section of *A. stephensi* midgut, scale bar - 50
170 µm. GFP-positive midgut cells are in green, RFP-positive *P. berghei* parasites are in
171 red, nuclei are labeled by DAPI in blue. **c**. Schematic 3D representation of the same
172 midgut as in (**b**), where the position of the cellular layer is calculated relative to the

173 nuclei. Positions of parasites are indicated as red dots, nuclei as blue dots. Deviation
174 of the cell layer from a flat surface is color-coded from blue to red (blue no deviation,
175 red - 10 µm). Note the blood meal location of the majority of parasites (above the cell
176 layer). **d**. Representation of nuclei (blue) and parasites (red) in the same midgut as (**b)**
177 after segmentation. **e**. Pooled positions of the parasites from all records at all time
178 points are shown for three layers relative to the midgut cells (blood meal, cellular layer
179 or basal lamina) for *A. stephensi*, *A. gambiae* and *A. gambiae* mosquitoes silenced for
180 *TEP1* (*A. gambiae*^{TEP1KD}). Each dot represents the number of parasites at a given
181 position in a single midgut. The numbers of midguts analyzed (n) are indicated above
182 the graph. Horizontal lines depict the mean number of parasites per position. The table
183 below summarizes parasite distribution inside the mosquito midguts at 18-25 hpi. The
184 percentage of ookinetes in the midguts of *A. stephensi*, *A. gambiae* and *Ag*^{TEP1KD} at
185 each location (blood meal, cellular layer and basal lamina) is given in parenthesis. n is
186 the number of parasites at each position, total n is the total number of analyzed
187 parasites.
188

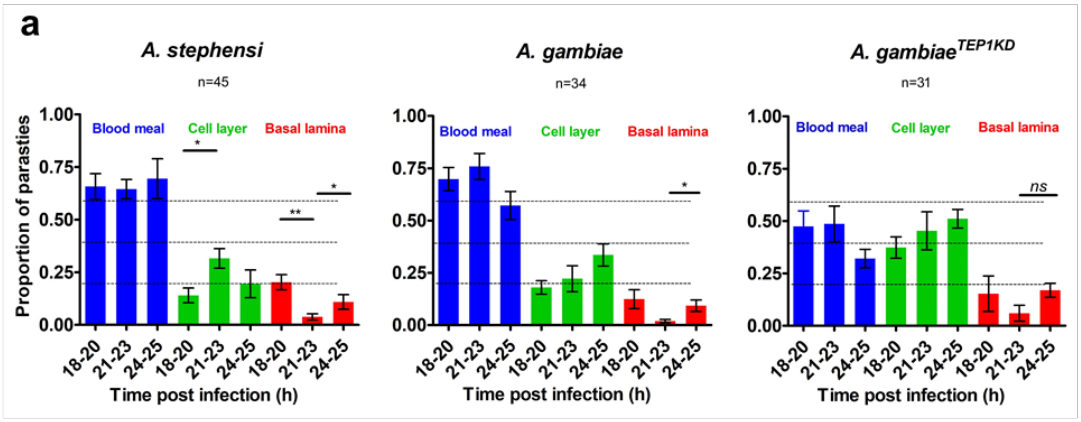
189 For each record, parasites and nuclei of the midgut cells were segmented and
190 their positions in 3D space were calculated relative to the cellular layer at each
191 examined time point after infection (Fig 2). The position of parasites relative to
192 the cellular layer was determined by fitting the midgut cell nuclei position by a
193 cubic spline surface. This surface was then considered as the central position
194 of the cellular layer (normalized z=0). An average thickness of 5 µm above and
195 below this surface defined the average cellular layer position.
196 We next examined whether the dynamics of parasite invasion was similar in
197 two *Anopheles* species. To this end, we measured the number of parasites at
198 each position (blood meal, cellular layer and basal lamina) in *As* and *Ag*.
199 Analyses of all time points did not detect significant differences in parasite
200 localization between the two species (Fig 2e). The majority of ookinetes were
201 detected in the blood meal (70%) and in the cellular level (20%). Only few
202 ookinetes crossed the midgut and reached the basal side (10%). Interestingly,
203 silencing of the major antiparasitic factor *TEP1* in *Ag* (*AgTEP1*^{KD}) significantly
204 changed spatial distribution of the parasites with only 40% of ookinetes
205 observed in the blood meal, 45% in the cellular layer and 15% at the basal side.

206 This difference in the dynamics of *Pb* invasion in *TEP1*-depleted mosquitoes
207 was suggestive of an additional role of TEP1 in inhibition of ookinete midgut
208 invasion. Previous studies reported *TEP1* expression in the larval gastric ceaca
209 and adult midguts [21,22]. In line with these reports, silencing of *TEP1* also
210 affected midgut microbiota by an as yet unknown mechanism [23]. Our findings
211 extend these observations to the early stages of parasite invasion and suggest
212 that in addition to parasite killing at the basal side, TEP1 directly or indirectly
213 inhibits *Plasmodium* midgut traversal.
214

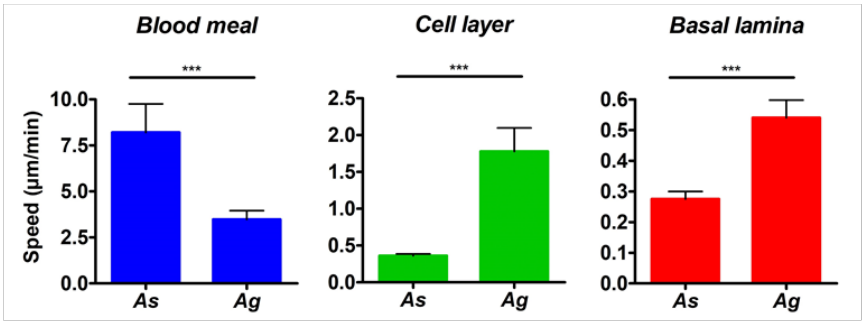
215 **Dynamics of the ookinete midgut invasion**

216 We next focused on *P. berghei* ookinete passage through the mosquito midgut
217 cells at different time points after infection and examined the proportion of
218 parasites at each position (blood meal, cellular layer and basal lamina). To this
219 end, we calculated the average proportion of parasites at each position at the
220 early (18 – 20 h post infection, hpi), intermediate (21 -23 hpi) and late (24 – 25
221 hpi) intervals after infection (Fig 3a, S4Fig).
222 We observed that in *As* mosquitoes the proportion of blood bolus-residing
223 parasites did not change over time. The proportion of parasites within the
224 cellular layer significantly increased between the early (14% at 18-20 hpi) and
225 intermediate (32% at 21-23 hpi) time intervals. However, this increase did not
226 cause accumulation of the ookinetes at the basal lamina. Instead, a significant
227 decrease in the proportion of basally located parasites was detected between
228 the early (20% at 18-20 hpi) and intermediate (4% at 21-23 hpi) time intervals.
229 Strikingly, this decrease was temporal, as the proportion of parasites in the
230 basal lamina significantly increased at the late time interval (10% at 24-25 hpi).

231 Similar decrease in the proportion of basally located ookinetes was detected in
232 *Ag*, where the proportion of parasites at the basal lamina declined from 12% at
233 18-20 hpi to 3% at 21-23 hpi, and then increased again to 14% at the late time
234 interval.



	Time post infection (h)				
	18-20	21-23	24-25	Midguts (n)	Parasites (n)
As	8	27	10	45	1068
Ag	12	11	11	34	796
Ag^{TEP1KD}	9	3	19	31	699



	Blood meal	Cell layer	Basal lamina
A. stepehensi			
n	25	84	288
Speed (µm/min)	8.2	0.36	0.28
A. gambiae			
n	54	10	23
Speed (µm/min)	3.47	1.78	0.54
p value	0.0001	<0.0001	<0.0001

235

236 **Figure 3. Kinetics of *P. berghei* invasion of *A. stephensi* and *A. gambiae* midguts.**
237 **a**. Positions of parasites in *A. stephensi* (*As*), *A. gambiae* (*Ag*) and in *A. gambiae*
238 mosquitoes silenced for *TEP1* (*A. gambiae*^{TEP1KD}) between 18 and 25 h post infection
239 (hpi). Plots show the proportion of parasites at each position (blood meal, cellular layer

and basal lamina) for three different time intervals (18-20, 21-23 and 24-25 hpi). Each bar represents the average proportion of parasites in midguts with at least 10 parasites. Parasite positions were calculated by the distance from the cellular layer: blood meal for ookinetes detected more than 5 µm above the cellular layer; basal lamina for parasites observed more than 5 µm below the cellular layer. Statistical analyses were performed by a non-parametric Mann-Whitney test. The table below shows the number of midguts analyzed at each time interval for each mosquito type. **b**. Speed of parasites as function of the parasite position in *As* and *Ag*. Speed (µm/min) was determined by tracking the parasites position over time from the time-lapse series. Four time-lapse experiments were used: guid 1615 and guid 1628 for *As* and guid 1622 and guid 2109 for *Ag*. The table below details the number of frames (n) used for speed calculations. Statistical significance of differences in the average speed at each given position between *As* and *Ag* were examined by the non-parametric Mann-Whitney test and $p \leq 0.0001$ are shown by three asterisks.

Since the mosquito immune system targets the ookinetes at the basal side of the midgut [24], we examined whether the observed decrease in the proportion of basally located ookinetes was rescued by *TEP1* knockdown. *TEP1* silencing eliminated the decrease in the basally located ookinetes observed in *As* and *Ag* mosquitoes and at the same time increased the proportion of parasites within the cellular layer (Fig 3a). These results suggest that the first wave of invading ookinetes is rapidly killed and lysed by the mosquito immune system. As the parasites that reach the basal lamina at later time points do accumulate, it is possible that asynchronous midgut invasion by *Pb* exhausts the components of the mosquito immune system and, thereby, benefits the establishment of infection by the second wave of the parasites. These results may also explain why not all parasites are recognized and killed by TEP1 at the basal lamina. We suggest that early crossing parasites may serve as pioneers that attract and locally deplete TEP1, allowing later-coming parasites to survive the immune attack.

To better understand *Pb* invasion dynamics, we measured ookinete motility in time-lapse experiments. The blood-filled midguts were dissected from infected mosquitoes and mounted *ex vivo* for imaging by spinning disk microscopy for

274 20 to 120 min. In line with the previous work [14], we observed four distinct
275 ookinete motility modes: (i) passive floating within the blood bolus (guid 2107,
276 guid 1615, S1 Table), (ii) gliding within the cellular layer (guid 1628, S1 Table)
277 (iii) spiraling in the blood meal and within the cellular layer (guid 1622, guid
278 1624, S2 Table) and (iv) stationary rotation without translocation within the
279 cellular layer (guid 2115, S1 Table). Some ookinetes were observed within a
280 midgut cell for more than one hour, suggesting that the parasites may remain
281 intracellular for relatively long periods of time without inducing cellular
282 apoptosis. By measuring the parasite speed in the blood meal, cellular layer,
283 and at the basal lamina, we found that the speed of ookinetes carried by the
284 bolus content was the highest as compared to other locations (Fig 3b).
285 Interestingly, the speed of the ookinetes in the blood bolus differed between *As*
286 (8.2 μm/min) and *Ag* (3.4 μm/min) midguts, suggesting some differences in the
287 blood bolus environment. The ookinete spiraling motility in the cellular layer was
288 much slower in both mosquito species, namely 0.36 μm/min in *As* and 1.78
289 μm/min in *Ag*. The slowest stationary rotation movement of parasites was
290 observed at the basal lamina (in *As*, average speed 0.28 μm/min, guid 2113,
291 S1 Table, in *Ag*, average speed 0.54 μm/min, guid 1622, S2 Table). We noted
292 that the speed of ookinetes within the cellular layer and at the basal lamina was
293 faster in *Ag* mosquitoes than in *As* mosquitoes. This observation indicates
294 important differences in the cellular organization of midguts of the closely
295 related mosquito species.

296 **Ookinete invasion routes**

297 To characterize ookinete invasion routes, intra- or extracellular location of the
298 ookinetes at the cellular layer was examined in more detail. To this end, we
299 developed an algorithm that classified intracellular, extracellular and
300 intercellular parasites based on the score of their 3D distance to the four
301 nearest neighboring nuclei of the midgut cells. The score was calculated for
302 each parasite (Fig 4a,b). The parasites with the score between 0 - 0.45 were
303 defined as extracellular, 0.45-0.55 - as intercellular, and higher than 0.55 - as
304 intracellular. We noticed a proportion of parasites that was extracellular at all
305 time points in both species (Fig 4c).

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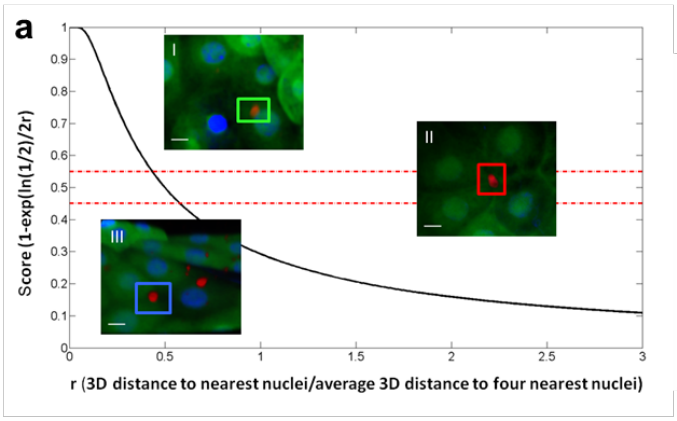
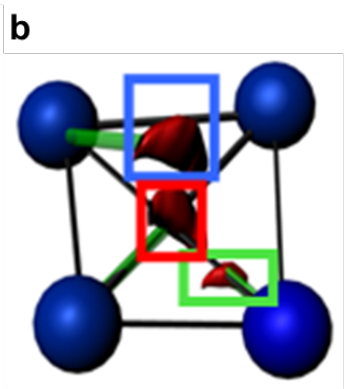
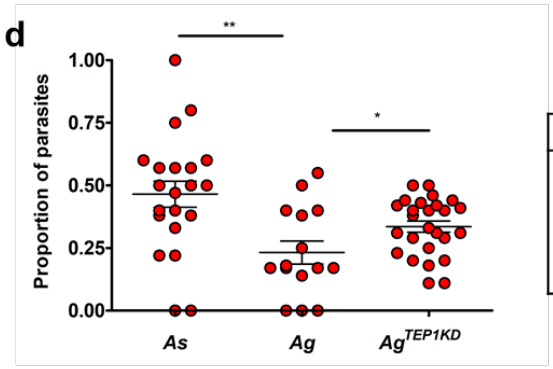




c

	Intercellular (%)	Intracellular (%)	Extracellular (%)	Parasites (n)
As (n=61)	39,0	46,7	14,2	246
Ag (n=51)	27,0	56,7	16,3	187
Ag^{TEP1KD} (n=36)	28,3	56,9	14,8	332



	As	*Ag*	*Ag^{TEP1KD}*
n	21	15	26
Intercellular	0.46	0.23	0.33
Intracellular	0.54	0.77	0.67
p value		0.0033	0.03

307 **Figure 4. Parasite distribution in the mosquito midgut**. Parasite positions within
308 the cellular layer calculated relative to the distance of each parasites to the nuclei of
309 surrounding midgut cells. **a**. Calculations of the distance of parasites from the nuclei
310 of the nearest neighboring midgut cell. The score (s) determines whether the parasite
311 is intercellular ($0.45 \leq s \leq 0.55$), extracellular ($s < 0.45$), or intracellular ($s>0.55$).
312 Example images from a z stack, scale bar = 20 µm: (I) s = 0.74, the parasite (green
313 arrow) is intracellular; (II) s = 0.45 (red arrow) the parasite is intercellular and (III) s =
314 0.36, the parasite is extracellular (blue arrow). **b**. Schematic representation of parasite
315 (red) and nuclei (blue) positions with distances (green lines) used to calculate
316 distances from the nuclei. **c**. Positions of parasites within the cell layer over time in *A.*
317 *stephensi* (*As*), *A. gambiae* (*Ag*) and *A. gambiae* mosquitoes silenced for *TEP1* (*A.*
318 *gambiae*^{TEP1KD}). The table indicates the percentage of parasites at each position for
319 each mosquito. The number *(n)* indicates the number of midguts analyzed for each
320 mosquito genotype. **d.** Comparison of the proportion of intercellular parasites between
321 *As*, *Ag* and *Ag*^{TEP1KD}. Each dot represents the proportion of parasites detected between
322 cells in a single midgut. Midguts (n) with at least six parasites within the cellular layer
323 were used for analyses. Statistically significant differences between *As* and *Ag* and
324 between *Ag* and *Ag*^{TEP1KD} revealed by a non-parametric t-test (Mann-Whitney) are
325 indicated by asterisks (p = 0.03 (*); p = 0.003(**)). The table details the mean
326 proportion values for parasites in each midgut and for each position for *n* mosquitoes.
327

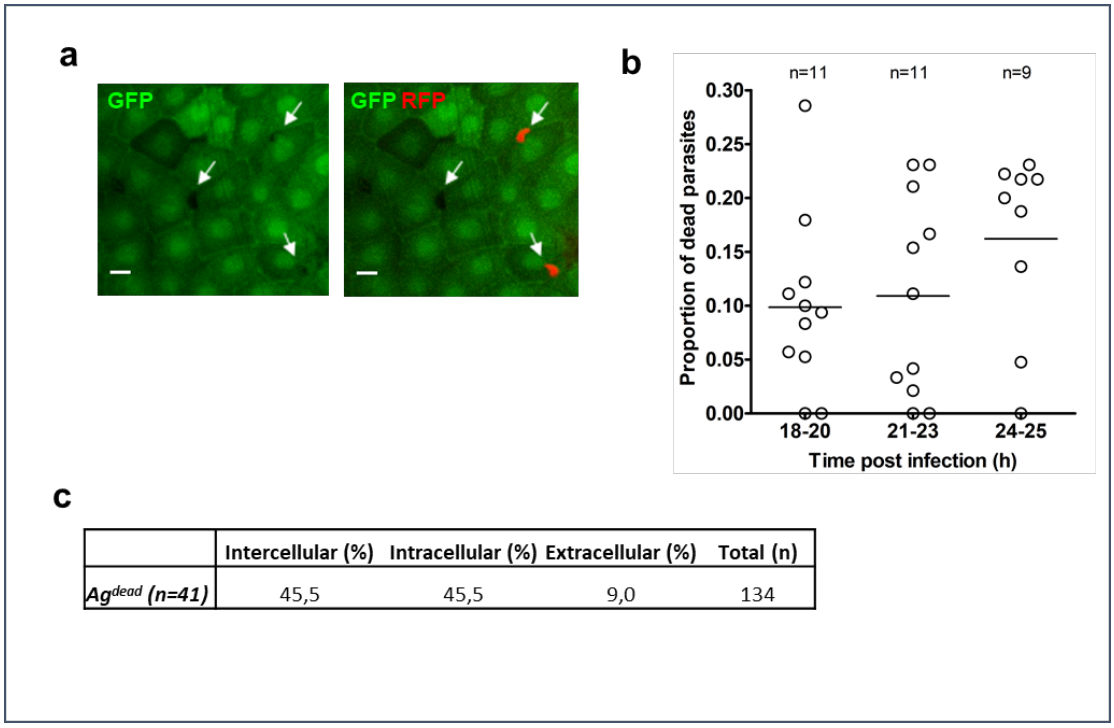
328 When comparing intercellular and intracellular parasite distribution, a higher
329 proportion of intercellular ookinetes was observed in *As* (40%) than in *Ag* (20%)
330 (Fig 4d, S6 Fig). These results point to intricate differences in parasite invasion
331 routes between the two related *Anopheline* species.

332

333 **Parasite viability within the midgut**

334 As the transgenic *P. berghei* line used in this study expressed the fluorescence
335 reporter under a constitutive promoter, we were surprised by high variability in
336 the reporter fluorescence levels observed between individual parasites in the
337 same midgut. We examined whether differences in fluorescence intensity
338 correlated with parasite localization and time post infection in two mosquito
339 species. To compare different experimental conditions, we normalized
340 fluorescence intensity of each parasite based on the highest and lowest
341 intensity of parasites in each image. We found only modest overall differences
342 in mean fluorescence intensities at different positions (basal lamina, cellular

343 layer, blood meal) over time and between the two species (*S7*-S9 Fig, Tables
344 S8-S9). Furthermore, we observed the parasites with very low levels of
345 fluorescence that appeared as a black hole on the background of the midgut
346 cells expressing GFP reporter in *Ag* mosquitoes (Fig 5a**)** that expressed GFP
347 uniformly in all midgut cells. In contrast, irregular patterns of GFP expression in
348 the midgut were reported for *As* [18] (S1 Fig). Hence, fluorescence-negative
349 parasites were only examined in *Ag* where parasites were clearly identified as
350 black shapes on fluorescent background.
351

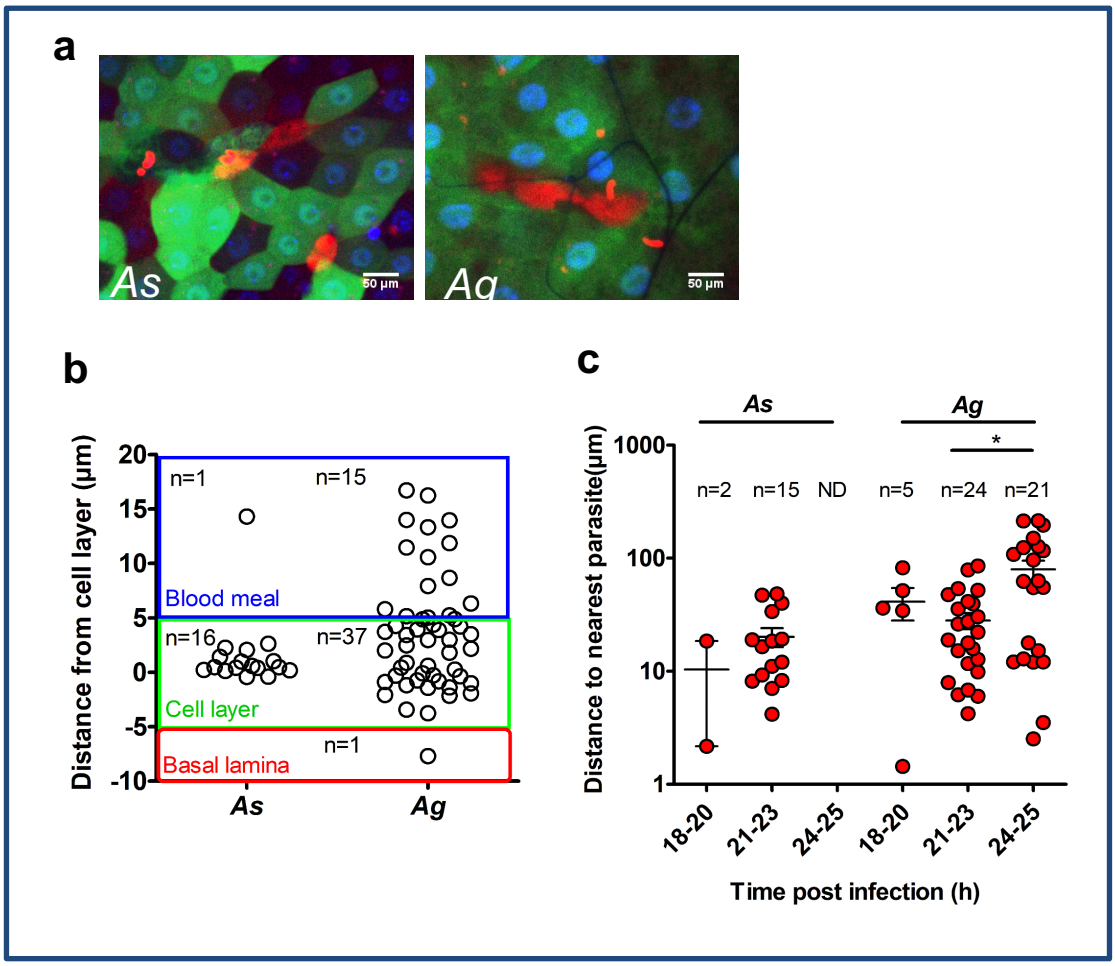


	Intercellular (%)	Intracellular (%)	Extracellular (%)	Total (n)
Ag^{dead} *(n=41)*	45,5	45,5	9,0	134

352
353
354 **Figure 5. Quantification of dead parasites in *A. gambiae*. a**. Detection of dead
355 parasites within the cellular layer. Due to uniform GFP expression with the midgut cells
356 of the *dmAct5C::GFP* line of *A. gambiae*, dead parasites that no longer express RFP
357 could be distinguished in the midgut by their negative signal and a characteristic
358 shape. Shown is a single z-section (scale bar - 20 µm) containing two live RFP-
359 expressing parasites and one dead parasite, indicated by arrows. **b.** The proportion of
360 dead parasites at different time points after *Ag* infection. Midguts (n) that contained at
361 least 10 parasites were used for analyses. Each dot represents a single midgut. **c**.
362 Distribution of dead parasites within the cellular layer. The table gives the number of
363 parasites at each position at all time points. The number (n) is the number of midguts
364 analyzed. Total (n) is the number of all analyzed parasites.
365

366 We considered the parasites that lost their fluorescence dying or dead [11,25].
367 On average, 10-15% of all recognized parasites had no fluorescence and were
368 classified as dead (Fig 5b). Differences in distribution were observed for live
369 and dead parasites within the cellular layer. More dead parasites were found to
370 be located extracellular or intercellular (compare Fig 5c and Fig 4d). This
371 observation points to more efficient parasite killing of extracellular parasites.
372 Interestingly, we hardly detected any dead parasites in Ag^{TEP1KD} mosquitoes,
373 suggesting that TEP1 may be involved in killing of parasites within the cellular
374 layer.
375
376 **Cell damage caused by parasite passage**
377 Midgut regeneration is a natural process of epithelia renovation after a blood
378 feeding, whether infective or not [26]. Blood meal generates a stressful
379 environment as it contains bacteria, reactive oxygen species and digestive
380 enzymes that may cause damage to the midgut cells. It has been previously
381 suggested that invaded midgut cells die after invasion and are expelled into the
382 midgut lumen [27] resulting in accumulation of hundreds of cells in highly
383 infected midguts. However, we only once observed GFP positive midgut cells
384 in the midgut lumen. This result indicates that either upon expel dead midgut
385 cells rapidly lose their GFP fluorescence, or that only few midgut cells are
386 expelled after invasion. To resolve these conjectures, we investigated the
387 integrity of the cell layer using high molecular weight Texas-Red conjugated
388 dextran which is trapped inside damaged cells [28]. In these experiments, the
389 fluorescent dextran was delivered into the midgut by blood feeding mosquitoes
390 on mice injected intravenously with fluorescent dextran several minutes before

391 mosquito feeding. We detected dextran filled cells (Fig 6a), calculated their
392 position (Fig 6b, S11 Fig) and measured the distance to the nearest parasite
393 (Fig 6c). The majority of dextran filled cells (70%) that contained a parasite in
394 *As* were predominantly detected in the cellular layer. In contrast in *Ag,* dextran
395 filled cells were observed both in the cellular layer and in the midgut lumen (Fig
396 6b). As many as 30% of dextran filled cells in *Ag*, were found in the midgut
397 lumen. Out of these, 50% contained a parasite (S10 Table). In contrast, we
398 found only one (5% of total) dextran filled cell the midgut lumen of *As*
399 mosquitoes.
400



401
402
403 **Figure 6. Quantification of damaged cells**. **a**. Detection of dextran-positive cells in
404 the midguts of *As* and *Ag* mosquitoes. Shown are single z-sections of GFP-expressing
405 dissected midguts. Mosquitoes were fed on mice injected with Texas-Red conjugated

406 dextran. Dextran-filled cells appeared red (scale bar - 50 μm). **b**. Positions of dextran-
407 filled cells in the midgut layers of *As* and *Ag*. Each dot represents a single dextran-
408 positive cell. The graph depicts positions of the dextran-positive cells within the midgut
409 layers. Each midgut layer is color coded: Blood meal (blue), cellular layer (green) and
410 basal lamina (red). The number of dextran filled cells (n) at each position is indicated.
411 **c**. Distances of dextran-filled cells to the nearest parasite at different time points after
412 infection of *Ag* and *As*. The number of dextran filled cells analyzed (n) is shown.
413 Statistical analysis was performed by a Mann-Whitney non-parametric t-test.
414
415 Interestingly in *Ag*, the distance between the dextran-positive cell and the
416 nearest parasite significantly increased at 24-25 hpi compared to the earlier
417 time intervals (Fig 6c). Furthermore, no dextran-positive cells were found in *As*
418 at the late time interval after infection (24-25 hpi). Taken together, these results
419 suggest that in *Ag* mosquitoes, damaged cells are readily extruded into the
420 midgut lumen with or without the parasites. It is important to note that while
421 some dextran filled cells contained a parasite, most midgut cells that we
422 observed to host a parasite were dextran-negative, indicating that ookinete
423 invasion damaged and killed only a small proportion of midgut cells.
424 Interestingly, in both *As* and *Ag*, we never observed more than one parasite in
425 a non-damaged midgut cell, indicating that parasites refrain from entering an
426 invaded cell. Here we also observed chains of several connected dextran-
427 positive cells, indicating that ookinetes can traverse several neighboring cells
428 before exiting on the basal side of the cellular layer. In conclusion, our results
429 led us to suggest that the route of ookinete invasion for the same parasite is
430 species-specific and shaped by midgut tissue morphology, physiology, damage
431 and immune responses. Future studies should examine how invasion
432 strategies of the human malaria *P. falciparum* parasites are affected by diverse
433 vector species.

434 # Conclusions

435 By combining live imaging techniques with quantitative bioimage analysis
436 workflow, we uncovered differences in ookinete invasion strategies in two
437 related mosquito species. We showed that in both species, the "pioneer"
438 parasites that first reach the basal side of the midgut were rapidly eliminated by
439 the mosquito immune system, and that colonization of the mosquito midgut was
440 initiated at later stages of the infection. High throughput image data analyses
441 of two *Anopheles* species revealed important differences in parasite invasion
442 routes. We showed that the average ookinete speed in the cellular layer is lower
443 in *As* compared to *Ag* mosquitoes. Moreover, *As* midguts contained more
444 intercellular parasites and displayed higher numbers of damaged parasite-
445 harboring cells. These results indicate that faster ookinete speeds and
446 preference for intracellular route may impede parasite survival during invasion
447 in *Ag*, the mosquito species which is more resistant to *P. berghei* infection.
448 The reported here combination of live imaging and automated image analysis
449 is highly adaptable and can be extended to functional analyses of gene
450 knockdowns, mutations, and drug treatments. Moreover, the image data base
451 and image analysis tools generated by this study offer a powerful tool for
452 studying *Plasmodium* motility in *Anopheles* mosquitoes.

453 # Materials and methods

454 **Mosquito rearing**

455 Transgenic *Anopheles stephensi* mosquitoes expressing GFP under the
456 midgut-specific G12 promoter (*pG12::EGFP transgenic line* [18]) and
457 *Anopheles gambiae* expressing GFP under the *Drosophila Acti5c* promoter

458 (*dmActin5c::dsx-eGFP*) line [19]) were reared in the lab as previously described
459 [29]. Briefly, mosquitoes were maintained in standard conditions (28°C, 75–
460 80% humidity, 12-hr/12-hr light/dark cycle). Larvae were raised in deionized
461 water and fed finely ground TetraMin fish food. Adults were fed on 10% sucrose
462 *ad libitum* and females were blood fed on anaesthetized mice. To obtain *Ag*
463 mosquitoes that do not express *TEP1*, the dominant *TEP1* knockdown Ag^{TEP1KD}
464 transgenic line [30] was crossed to *dmActin5c::dsx*-e*GFP* mosquitoes. The F1
465 progeny had reduced TEP1 levels while expressing GFP in the midgut [30].

466 ***P. berghei* infections**

467 For infections, mosquitoes were blood fed on *P. berghei* infected mice as
468 previously described [31]. *P. berghei* pyrimethamine resistant strain (RMgm
469 296) constitutively expressed RFP [20]. For the visualization of damaged
470 mosquito cells, mice were injected in the tail vein with 0.1 ml of 5% dextran
471 (3,000 kDa Texas Red conjugated, Invitrogen) diluted in PBS 10 min prior to
472 blood feeding. Mosquitoes were blood fed for 20 min on anesthetized mice and
473 dissected between 18-24 h after blood feeding, as indicated in each
474 experiment.

475 **Confocal microscopy**

476 Immediately prior to visualization, infected mosquitoes were dissected on ice in
477 PBS buffer supplemented with 0.02% DAPI (Thermo Fisher, 4′,6-diamidino-2-
478 phenylindole, 5 mg/mL), and with 0.2% tricaine (Sigma), 0.02% tetramisole
479 (Sigma) to prevent midgut contraction during image acquisition. Blood-filled
480 midguts were placed on 35 mm plastic dishes with glass bottom (Nunc,
481 ThermoFisher). Dishes were mounted on inverted DMI6000 Leica Microscope,
482 equipped with a Nipkow Disk confocal module (Andor Revolution), 20X

483 objective. For time-lapse experiments, samples were visualized for up to two
484 hours at 1 min intervals. The number of 1 µm-stacks, annotated for each image,
485 ranged between 24 and 95 depending on tissue thickness. We noticed that *As*
486 midguts were rigid and sturdy, allowing for longer live imaging. *Ag* midguts were
487 more fragile and tended to move and tear during image acquisition. We were
488 able to collect live imaging data from 16 *As* (S1 Table) and 5 *Ag* midguts (S2
489 Table). We were not able to follow parasites in mosquitoes lacking the immune
490 protein TEP1 due to high midgut fragility of Ag^{TEP1KD} midguts.

491 **Image analyses**

492 All images were uploaded to a database where they were annotated according
493 to mosquito species and other experimental conditions. Images were subjected
494 to bulk analysis as well as manual verification. The annotated image database
495 is accessible to JAVA programming using the Strand Avadis IManage data
496 management software. All data (images and extracted data as text files) are
497 available on cid.curie.fr, Project "Malaria parasite invasion in the mosquito
498 tissues" at https://cid.curie.fr/iManage/standard/login.html. The META data is
499 managed using OpenImadis https://strandls.github.io/openimadis/. Companion
500 scripts are available here:
501 https://github.com/PerrineGilloteaux/MalariaParasiteinMosquito.
502 The api documentation is available under API tab
503 https://cid.curie.fr/iManage/api/client/. The api client jar is available at
504 https://cid.curie.fr/iManage/standard/downloads.html.
505 Companion scripts include segmentation of parasites, nuclei and quantification
506 of intensities corrected by background was performed using a set of ImageJ

507 Plugins in Java. Analysis of the position of parasites relative to cell layer and
508 statistics are performed with MATLAB.
509 The data set was collected from 110 experiments including a total of 2,557
510 parasites (*As* - 45 midguts, 1,068 parasites; *Ag* - 34 midguts, 796 parasites;
511 and Ag^{Tep1KD} – 31 midguts, 693 parasites). There was no bias in the number of
512 parasites per midgut across different time points and mosquito species (S2
513 Fig). Furthermore, we found that infection levels (low, intermediate, or high) had
514 some effect on the result of parasite distribution, specifically in at low infection
515 levels (S3 Fig). Consequently, we used for our analysis only images that
516 contained at least 10 parasites per image.

517 **Ethics statement**

518 The animal work described in this study received agreement #E67-482-2 from
519 the veterinary services of the region Bas-Rhin, France (Direction
520 départementale des services vétérinaires).

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528 Author contributions

529 Conceived and designed the experiments: GV, EAL. Performed the
530 experiments: GV, JŠ, JuS. Analyzed the data: GV, PPG, JeS, EAL. Contributed
531 reagents/materials/analysis tools: PPG, JeS. Wrote the paper: GV, EAL.

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