**Dietary L-3,4-dihydroxyphenylalanine (L-DOPA) augments cuticular melanization in *Anopheles* mosquitos while reducing their lifespan and malaria parasite burden**

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**ABSTRACT**

L-3,4-dihydroxyphenylalanine (L-DOPA), a naturally occurring tyrosine derivative, is prevalent in environments that include mosquito habitats, potentially serving as part of their diet. Given its role as a precursor for melanin synthesis we investigated the effect of dietary L-DOPA on mosquito physiology and immunity to *Plasmodium falciparum* and *Cryptococcus neoformans* infection. Dietary L-DOPA was incorporated into mosquito melanin via a non-canonical pathway and had profound transcriptional effects that were associated with enhanced immunity, increased pigmentation, and reduced lifespan. Increased melanization resulted in an enhanced capacity to absorb electromagnetic radiation that affected mosquito temperatures. Bacteria in the mosquito microbiome were sources of dopamine, which is a substrate for melanization. Our results illustrate how an environmentally abundant amino acid analogue can affect mosquito physiology and suggest its potential usefulness as an environmentally friendly vector control agent to reduce malaria transmission, warranting further research and field studies.

**INTRODUCTION**

Malaria is a major public health threat exacerbated by the impact of climate change [1]. Global warming has pushed mosquitoes to new latitudes and altitudes, prolonging the malaria transmission season and increasing the risk to previously unaffected communities [2,3]. In addition, human activities such as land development, intensive agricultural practices, and the widespread use of chemical pesticides significantly influence the proliferation and spread of mosquito populations [2]. As a result, the dynamics of disease transmission by these vectors are continually

evolving. A stark illustration of these public health consequences occurred last year when the United States of America (USA) experienced nine cases of locally acquired malaria –marking the first time in two decades that locally acquired transmission had occurred. Notably, eight of these cases were reported in Florida and Texas, with an additional case identified in Maryland. This alarming trend contrasts with the situation in 2003 when eight cases of locally acquired malaria were identified and all were restricted to Florida's Palm Beach County. The *Anopheles* mosquito vectors, responsible for transmitting malaria, are found throughout many regions of the US, posing a substantial risk if they encounter a malaria-infected person [4]. In Africa, climate change is fueling a perfect storm for malaria [3,5]. The combination of increased frequency of extreme weather events, deforestation, warmer temperatures at higher altitudes, rapid spread of insecticide-resistant *Anopheles* mosquitoes [6], and emergence of urban mosquito vectors like *Anopheles stephensi* [7] threaten progress in malaria control and pose greater risks to human health.

Most vector borne diseases have been mitigated through the implementation of efficacious vector control programs [8]. However, maintaining the long-term sustainability of such programs is difficult and often impractical. In the case of malaria, prior to the use of insecticides, vector control strategies  relied on understanding the ecology of local vectors such as changing the aquatic habitats where the vector developed (e.g., draining swamps, applying oil to open water bodies, cutting vegetation at edges of ponds/streams). These interventions were highly successful but labor intensive and they had huge negative environmental impact [9]. Consequently,  environmental management practices were replaced by insecticide-based vector control. Despite initial success – between 2000 to 2015, the employment of nets impregnated with residual insecticides led to a reduction in over 50% of malaria cases in Africa – these efforts were compromised by the emergence of insecticide-resistance in the mosquito vectors and drug-resistance in the malaria-carrying parasites which they harbor [10,11].

Considering the challenges outlined above, the World Health Organization (WHO) has urged the development of cost-effective, sustainable, and environmentally safe vector-control interventions. As adult mosquitoes of both sexes, including hematophagous species, feed from floral nectar as a common nutritional source [12], attractive toxic sugar baits (ATSBs) are alternative and potential new tools [13]. These mosquito control methods rely on appealing cues and sugar loaded with distinct agents such as transgenic entomopathogenic fungi [14], genetically modified bacteria (para-transgenesis) that confer mosquito refractoriness [15], or natural microbiota that result in increased resistance and/or decreased fitness [16].

In insects, melanin plays an essential role in three physiological processes: cuticular melanization, wound healing, and immunity [17]. Research on the complex nature of plant-insect interactions has shown that insects adapt to L-DOPA-rich diets [18-20]. For instance, the pea aphid *Acyrthosiphon pisum* uses plant-derived L-DOPA to promote melanin synthesis for wound healing and UV protection while sequestering high toxic compound levels [18]. Consequently, we aimed to investigate the potential of 3,4-dihydroxyphenylalanine (L-DOPA), a melanin precursor found naturally in flowers of the faba bean, *Vicia faba* L. (Fabaceae) [21,22]. We hypothesized that adult *A. gambiae* mosquitoes modulate melanin-related physiological processes in response to a sugar diet supplemented with L-DOPA, thereby reducing their susceptibility to foreign organisms that could pose a threat to humans. Consequently, we assessed the oral toxicity of L-DOPA concentrations in a sugar solution, defined the transcriptional profile of *A. gambiae* female mosquitoes in response to dietary L-DOPA and evaluated its functional impact on key vector life traits (**Fig. 1**). L-DOPA-fed *A. gambiae* female mosquitoes exhibited downregulation of essential immune-related genes, had reduced malaria parasite and fungal burden, enhanced cuticular melanization, shortened life span, and suppressed expression of critical genes involved in the synthesis of cuticular hydrocarbons. Our findings suggest a potential role for L-DOPA in malaria control strategies that target mosquito reproduction.

## RESULTS

**Effects of dietary L-DOPA on survival of *A. gambiae* female mosquitoes.**

*A. gambiae* females rely mainly on a blood meal to obtain phenylalanine as the initial substrate for the melanization process. In contrast, male mosquitoes depend exclusively on plant sources (**Fig. 2**). To investigate whether a sugar solution supplemented with L-DOPA had a beneficial or detrimental impact on melanin-related physiological processes of adult mosquitoes, we first assessed its effect as a supplement to sugar meals on longevity, as a measure of mosquito fitness. We provided 3–4-day-old female mosquitoes with a low sugar solution supplemented with L-DOPA at concentrations ranging from 0.2 to 10 mM (0.4 to 20% w/w) and then monitored their survival over a 14-d period. These L-DOPA concentrations were comparable to those reported in natural sources, which range from 0.2 to 7.7% (w/w) [21,23-28]. We observed that after 8 d of L-DOPA exposure, mosquitoes fed on L-DOPA solutions at most concentrations started to experience a drastic increase in mortality, except for those fed on 1 mM (2%) L-DOPA, whose survival exceeded that of the control (**Fig. 3A**). By 14 d, mosquitoes fed with L-DOPA solutions at concentrations lower than 1 mM exhibited a mortality of ~50%, while at higher doses mortality

increased in a dose-response relationship. Notably, analysis of the accumulative survival at different time points revealed that dietary L-DOPA exhibited a clear biphasic dose-response (**Fig. 3B**). Conversely, when providing dopamine in the sugar meal the hormesis effect disappeared, with control mosquitoes and those fed with up to 1 mM dopamine-sugar dying at an equal pace (**Fig. 3C, 3D**). Consequently, we used 1 mM L-DOPA in subsequent experiments.

**Dietary L-DOPA affects expression of mosquito immune- and cuticle-related genes.**

To identify genes whose expression was affected by a non-toxic concentration of L-DOPA, we collected female *A. gambiae* mosquitoes post 1 mM L-DOPA feeding of the sugar meal during 4 d and performed RNA sequencing (mRNA-seq) on whole mosquitoes (headless) and midguts. We observed 902 and 165 differentially expressed genes between L-DOPA-fed versus sucrose-fed controls (≥ 2-fold change, $P$ < 0.05) in the whole mosquitoes and midgut samples, respectively (**Fig. 4**). In whole mosquitoes, these included 808 downregulated and 94 upregulated genes, both overrepresented in the functional category of stress-related (R/S/M; oxidoreductive, stress-related, mitochondrial) (**Fig. 4A, 4B**). In midguts, we observed 50 downregulated and 115 upregulated genes. Downregulated genes were overrepresented in the functional categories of transport, DNA-related (R/T/T; replication, transcription, translation), and stress-related (**Fig. 4C, 4D**, **Supplementary Table 1, Supplementary Fig. 1**). Unexpectedly, key genes involved in the melanin biosynthesis pathway such as prophenoloxidase 5 (PPO5) (AGAP012616) and dopachrome-conversion enzyme (DCE) (AGAP005959) were significantly downregulated. Diverse immune-related gene families involved in pattern recognition receptors (PRRs) (e.g., *peptidoglycan recognition proteins* (PGRP), *scavenger receptor* (SCR), and *galectins* (GALE)), signal modulation and amplification (e.g., *clip domain serine proteases* (CLIP) and *serpins* (SRPN)), signal transduction pathways (e.g., TOLL), and other effector systems (e.g., *caspases* (CASP) also exhibited significant downregulation. Furthermore, two critical genes for dopamine metabolism in insects, dopamine beta-monooxygenase (DBH) (AGAP010485) and dopa decarboxylase (DDC) (AGAP009091), which participate in the synthesis of octopamine (main neurotransmitter) and cuticle, respectively, were also downregulated. By contrast, the immune-related dally protein (AGAP005799) involved in signal transduction was significantly upregulated (**Fig. 5A,** upper panel, **Supplementary Table 1**). In the midguts, the most significantly upregulated gene corresponded to phosphoinositide-3-kinase, regulatory subunit (PI3K) (AGAP005583) that participates in bacterial phagocytosis and apoptosis, and autophagy in *Drosophila*[29] (**Fig. 5A**, lower panel; **Supplementary Table 1**).

Regarding cuticle-related genes, transcriptional profiles of whole mosquitoes showed that L-DOPA-supplementation was associated with an extensive downregulation of gene families involved in cuticle regulation, metabolic detoxification, and synthesis of cuticular hydrocarbons. These included: cuticular proteins (e.g., AGAP000820 and AGAP00098); chitinase (CHT5-5, AGAP013120); chitin synthase 1 (CHS1A, AGAP001748); cytochrome P450s; propionyl-CoA-synthase (AGAP001473); and elongases (AGAP005511, AGAP005512, AGAP013219). Notably, upregulated genes in L-DOPA-fed mosquitoes corresponded to hexosaminidase D (AGAP004270) and glutathione S-transferase epsilon class 6 (GSTE6) (AGAP009191) (**Fig. 5B,** upper panel, **Supplementary Table 1**). The latter gene is associated with the development of insecticide resistance in the malaria vector *Anopheles funestus* [30]. Similarly, in the midgut, cytochrome P450s and chitin-related genes were downregulated while *Jheh2*, a gene coding for juvenile hormone epoxide hydrolase, was upregulated (**Fig. 5B,** upper panel, **Supplementary Table 1**).

Notably, almost a third (255 out of 902) of the genes differentially regulated in response to L-DOPA corresponded to hemocyte-associated genes. Most of these were oenocytoid-related (76.9%) versus granulocyte-related immune cell subtypes (19.2%), as defined by scRNA-seq hemocytes clustering performed by Kwon *et al* [31]. These genes are involved in multiple functional categories including immunity, lipid metabolism, proteolysis, transport, and chemosensory systems such as odorant binding proteins (OBPs) and G-coupled protein receptors (GPCRs) (**Supplementary Table 1**). Collectively, our transcriptional analysis suggests that dietary L-DOPA was associated with a global modulation of genes beyond those related to melanin-associated processes.

**Dietary L-DOPA promotes mosquito resistance to malaria parasite and fungal infection.**

To investigate whether a L-DOPA-supplemented sugar meal could modulate *P. falciparum* infection in *A. gambiae* mosquitoes, we fed *Anopheles* female mosquitoes on L-DOPA containing solutions (0.2 mM; 0.6 mM; and 1 mM) for 4 d prior to *P. falciparum* infection. The intensity of the parasite infection was assessed by enumeration of *P. falciparum* oocysts at 8 d post-infection (dpi). Mosquitoes fed with a sugar meal supplemented with L-DOPA demonstrated a dose-dependent reduction of the *P. falciparum* parasite burden with a significant reduction for the 1 mM L-DOPA in comparison to the sugar-fed control (median number of oocysts 2 vs 5, *p*= 0.0032). L-DOPA supplementation also reduced the infection prevalence (77.5% vs 87.3% at 1 mM in

comparison to the control) (**Fig. 6A**). Using dopamine instead of L-DOPA in the sugar meal was also associated with a significant reduction of *P. falciparum* infection burden in dose-dependent manner relative to the sugar-fed controls. At the highest dose, 1 mM dopamine, the median number of oocysts was 2 vs 7, *p*=0.0006 in comparison to the control, and the prevalence of infection was also reduced (81.1% vs 87.5%) but to a lesser degree than in the presence of an L-DOPA-supplemented sugar meal (**Fig. 6B**). No melanized parasites were visualized in these experiments at 2 or 8 dpi, either in absence or presence of L-DOPA- or dopamine-supplemented meals after the parasite infection (data not shown), indicating that reduced parasite infection resulted by a non-melanin-based immune response. Given that nectars provide mosquitoes with L-DOPA, we investigated whether L-DOPA directly exhibits *in vitro* antiplasmodial activity to both sexual and asexual stages of the human malaria parasite. At 1 mM L-DOPA, sexual stages of *P. falciparum* exhibited significant toxicity (32.4% survival vs sugar-fed control, *p* < 0.0001), while the viability of asexual stages of the parasite was nearly abolished at 1 mM L-DOPA and drastically reduced at 0.5 mM L-DOPA (26.3% survival vs control) (**Fig. 6C**).

*A. gambiae* triggers a melanin-based immune response against fungi [32,33]. To ascertain whether L-DOPA supplementation promoted fungal melanization, we challenged mosquitoes with an intrathoracic injection of *Cryptococcus neoformans* (H99-GFP strain), prior to L-DOPA exposure for 4 d. Fungal viability was assessed by determining the number of fungal colony-forming units (CFUs) 24 h following injection and by tracking GFP fluorescence. L-DOPA-fed mosquitoes showed reduced number of fungal CFUs in comparison to the control (**Fig. 6D**) and revealed efficiently melanized yeast cells (**Fig 6E**). Furthermore, we found that despite downregulation of dopa decarboxylase (DDC), a key gene on the melanization pathway, other genes involved in *Plasmodium* parasite killing and survival —specifically the antimicrobial peptide gambicin (GAM)[34] and the leucine-rich immunomodulator 15 (LRIM15)[35]— were significantly deregulated, affecting *Plasmodium* parasite development (**Fig. 6F**). Our analyses demonstrated that despite broad downregulation of canonical melanization-related genes that was validated by qRT-PCR, L-DOPA-supplementation promoted an antiplasmodial and anti-fungal response in *A. gambiae* female mosquitoes.

**Dietary L-DOPA enhances mosquito cuticular melanization.**

The insect cuticle protects against physical trauma (rigid cuticle) and flexibility to support joint areas allowing mobility (soft cuticle) [36]. Melanin is primarily thought to be involved in the formation of rigid cuticles, though its participation in the biogenesis of flexible cuticles is uncertain. While

assessing daily mortality, we serendipitously noticed that the 1 mM L-DOPA-fed mosquitoes displayed a greater darkening of the cuticle pigmentation in comparison to sucrose-fed controls (**Fig. 7A,** upper panel). Thus, we explored the contribution of dietary L-DOPA to insect cuticular melanization by allowing *A. gambiae* female mosquitoes to feed with a sugar solution supplemented with L-DOPA (0.2 mM to 1 mM) for up to 14 d. At 8 and 14 d, we observed and measured a dose-dependent darkening of the mosquito cuticle; 1 mM L-DOPA-fed mosquitoes exhibited significantly darker pigmentation relative to sugar-fed control mosquitoes (**Fig. 7A,** lower panel)**.** To further assess the mosquito cuticular ultrastructure and chemical nature of its pigment, we used transmission electron microscopy (TEM) to examine the cuticular architecture of legs from *A. gambiae* female mosquitoes after 8 d of exposure to the L-DOPA diet. Micrographs from female L-DOPA-fed mosquitoes revealed a distinct microstructure of the leg endocuticle layer in comparison to the sucrose-fed control (**Fig. 7B,** top panel). The mean thickness of the total cuticle was significantly greater (mean = 2,424.11 ± 467.97 nm) than that of the control (mean = 1,962.63 ± 302.91 nm) **(Fig. 7B,** bottom panel). In response to the L-DOPA diet, female mosquitoes showed cuticle enrichment promoted by an increase of both procuticle layers (endocuticle and exocuticle). In addition, we determined the accumulation of melanin in whole mosquitos using solid-state NMR with $^{13}$C-isotopically-labeled L-DOPA. Adult female *A. gambiae* mosquitoes were fed with 1 mM ring-$^{13}$C-L-DOPA in sucrose for 8 d to augment the ssNMR signal intensity. To estimate the relative amounts of long-chain lipids, polysaccharides, and melanin in *A. gambiae* mosquitoes fed with or without labeled L-DOPA, we acquired quantitatively reliable $^{13}$C direct-polarization magic-angle spinning (DPMAS) NMR data. In comparison to the control, spectra from ring-$^{13}$C-L-DOPA-fed female mosquitoes displayed increased signal intensities at 110-160 ppm which are characteristic of the indole-based melanin pigment (**Fig. 7C,** green arrows in left panel), suggesting that mosquitoes used ring-$^{13}$C-L-DOPA to synthesize cuticular melanin. To determine whether mosquitoes broke down ring-$^{13}$C-L-DOPA for nutritional purposes, we also acquired $^{13}$C cross-polarization magic-angle spinning (CPMAS) spectra that favored the most rigid constituents. This latter study corroborated the nearly exclusive use of ring-$^{13}$C-L-DOPA for pigment production: signals corresponding to polysaccharides (~55-105 ppm) as well as those corresponding to aliphatic carbons (10-40 ppm) were largely unchanged in integrated intensity upon feeding with ring-$^{13}$C-L-DOPA (**Fig. 7C,** right panel). Interestingly, L-DOPA supplementation also increased the content of indole groups but reduced the cuticular thickness of legs in male *A. gambiae* mosquitoes (**Supplementary Fig. 2**). These mosquitoes also displayed a negligible increase in their chitin content (0.64 ± 0.07 nmoles of glucosamine per mg of whole mosquito dry weight) in comparison to the sucrose-fed controls (0.46 ± 0.06 nmoles of glucosamine per mg of

whole mosquito dry weight) (**Fig. 7D**). Furthermore, we tested whether the augmented indole-pigment content corresponded to melanin by performing electron paramagnetic resonance (EPR) measurements. EPR is a particularly effective methodology that detects a remarkable characteristic feature of this biopolymer, the presence of a population of stable organic free radicals [37]. Analysis of the acid-resistant material isolated from L-DOPA-fed mosquitoes revealed a spin concentration that decreased to 0.81-fold of its value (14.55 vs 17.75, scaled by mass) compared with sugar-fed controls (**Fig. 7E**)." Lastly, we demonstrated that melanin synthesis was not driven by DCE providing a L-DOPA diet to the *Aedes aegypti* yellow (DCE2, AAEL006830) mutant strain[38], which is defective for eumelanin synthesis. This mutant mosquito strain also exhibited a spin concentration that diminished to 0.3-fold of its value (2.36 vs 7.60) compared with its sugar-fed control. The lower spin concentrations of L-DOPA supplemented samples in comparison to controls is suggestive of an enhanced radical scavenging activity— a key melanin property[39]. Notably, the high spin concentration exhibited by the *A. gambiae* L-DOPA supplemented samples compared with the *Aedes* L-DOPA-fed mutant mosquito strain (14.55 vs 2.36, respectively) could imply the isolation of different types of melanins (a mixture of pheomelanin/eumelanin vs eumelanin) with distinct antioxidant properties. These results agree with our fungal eumelanin sample, used a control, which displayed a spin concentration of 4.26. Further studies will be required to gain a deeper understanding of this system.

In sum, oral L-DOPA promoted mosquito cuticle melanization via a non-canonical pathway independent of L-DOPA conversion into dopachrome, impacting the thickness of both endocuticle and exocuticle layers.

**Enhanced cuticle melanization fails to promote insecticide resistance yet raises mosquito body temperature and reduces lifespan.**

Previous studies in *Anopheles* mosquitoes have shown that changes in the cuticle structure are associated with insecticide-resistance [40,41]. Thus, we set out to determine whether enhanced cuticular melanization impacted female *A. gambiae* resistance to pyrethroids, the only WHO-approved insecticide used on long-lasting insecticide-treated nets (LLINs). Using WHO-approved papers impregnated with 0.05% deltamethrin for 1 h exposure, L-DOPA-fed *A. gambiae* female mosquitoes demonstrated susceptibility to pyrethroids (98.7± 3.3% mortality rate in 24 h). Despite their highly melanized and thicker legs, L-DOPA-fed *A. gambiae* female mosquitoes were susceptible to 0.05% deltamethrin.

Dark pigmentation is associated with enhanced ability to capture heat from electromagnetic radiation on animals, plants, and fungi [42]. In insects, melanization (melanism) is associated with negative or positive consequences in thermal adaptation, mating, cuticle strength, etc. that impact their fitness and ecology [43-45]. To probe mosquitoes' thermal response to visible electromagnetic radiation, we used a thermal camera to measure the maximum average body temperature of L-DOPA-fed individual mosquitoes after being exposed to a white light mimicking sunlight. Female *A. gambiae* mosquitoes fed with L-DOPA for 8 d maintained an apparent average body temperature that was $0.5 - 2.0$ °C higher that sugar-fed control mosquitoes, following 1 min of irradiation (**Fig. 8A, Supplementary Fig. 3**). Notably, 1 mM L-DOPA-fed mosquitoes were warmer than control. These mosquitoes exhibited the highest temperature, both pre- and post-irradiation (**Fig. 8B, Supplementary Fig. 3**). Because trade-offs between heat tolerance and life-history are observed in insects [46], we further investigated whether longevity of L-DOPA-fed mosquitoes differed from those of controls. We measured the longevity of female mosquitoes when maintained on the L-DOPA diet (**Fig. 8C**). The lifespan of female mosquitoes was significantly decreased when maintained on a sucrose meal supplemented with 1 mM L-DOPA (Wilcoxon test, Chi-square, 13.74, $p = 0.0002$). These mosquitoes exhibited no apparent changes in reproductive fitness: number of eggs laid (**Fig. 8D**) and hatching rate (**Fig. 8E**).

**The mosquito microbiome includes bacteria that produce dopamine.**

Unlike mammals, which can use L-DOPA as a melanin precursor [47], insects rely on dopamine [48]. This led us to investigate how dietary L-DOPA was metabolized in female mosquitoes and whether the mosquito microbiome included dopamine synthesizing bacteria. We observed the intriguing trend whereby as L-DOPA levels increased in the sugar meal, dopamine levels in mosquitoes decreased (**Fig. 9A**). Leveraging our experience with studying fungal melanization in *Cryptococcus* spp., which can be triggered by dopamine synthesized by bacteria [49], we explored whether gut microbiota isolated from field-caught mosquitoes have the potential to serve as an endogenous source of dopamine. *Cryptococcus* spp., known to be members of the mosquito gut mycobiome [50,51], depend exclusively on exogeneous precursors for melanin synthesis (e.g., L-DOPA, dopamine, epinephrine, and norepinephrine) [52]. We selected thirteen bacterial strains that had been previously isolated from the midgut of field-caught *Anopheles* [16] and *Aedes* [53] mosquitoes plus a strain isolated from the soil[54,55] (**Table 1**) and conducted a plate confrontational assay whereby bacteria that produce dopamine trigger melanization in *Cryptococcus neoformans* [49]. After eye inspection of the plates, we noticed that five bacterial isolates consistently induced

melanization (brown pigmentation) in both of our positive control strains (*C. gattii* WM179 and *C. neoformans* H99), at both room temperature and 30°C. As expected, no pigmentation was observed in the *C. neoformans* LAC1 mutant strain (negative control), in which melanin synthesis is compromised by the deletion of its laccase gene (**Fig. 9B**). The melanin-inducing bacteria were identified as: *Comamonas* sp., *Enterobacter hormaechei*, *Aeromonas hydrophila*, *Serratia marcescens*, and *Chromobacterium subtsugae* Δ*vioS* (Δ*vioS*). To verify the chemical nature of the soluble compound released by mosquito-derived bacteria that promoted cryptococcal melanization, we used a Fast Protein Liquid Chromatography (FPLC) protocol for recovering dopamine from bacterial supernatants grown in Luria-Bertani (LB) media for up to 96 h at 30°C. The chromatograms displayed a peak between 4.39 and 5.66 min matching the position of the dopamine spike peak. We observed that peaks varied depending on the bacterial species and tended to be larger as the incubation time increased (**Supplementary Fig. 4**). Therefore, we collected peaks at 96 h of culture incubation, removed cells by quick centrifugation, filter-sterilized the supernatants, acidified them to pH 4.0 with 1 M HCl to maintain dopamine stability, and kept them on ice, and away from light. Samples were further identified as dopamine and quantified by Electrospray Ionization (ESI) Mass Spectrometry (MS). Dopamine production was verified from four strains at concentrations ranging from 7 ng/ml to 67.5 ng/ml. These were the dopaminergic bacteria identified (concentration of dopamine detected): *Aeromonas hydrophila* (7.10 ± 0.4 ng/ml), *Comamonas sp.* (7.00 ± 0.1 ng/ml) and *Enterobacter hormaechei* (67.5 ± 2.4 ng/ml). No dopamine was quantified for *Serratia marcescens* under the tested conditions. The limit of quantification of the ESI MS was 2.5 ng/ml.

**Discussion**

In insects, melanin is synthesized both in the exoskeleton (cuticle) and inside the body, including by the immune system, which uses melanization as an antimicrobial mechanism [56-58]. The cuticle is crucial for insect survival, making it a primary target for pest and disease vector control. Its mechanical properties (i.e., rigidity and flexibility) are related to various chitins, proteins, and the type and abundance of sclerotization or protein crosslinking occurring within [59]. This study demonstrates that *Anopheles* mosquitoes can use dietary L-DOPA to enhance cuticle melanization, leading to increased body pigmentation, cuticular thickness in the legs, overall indole and chitin content, and a key EPR signal (**Fig 7**). Given the abundance of L-DOPA in environmental and the potential microbiome sources of dopamine, our results have important implications for mosquito biology and potential control strategies.

Cuticular melanogenesis is closely related to cuticular hardening or sclerotization. Both are oxidative processes occurring in the same tissues and use L-DOPA or dopamine as precursors; however, the participation of laccase 2 (AGAP008731) and dopamine derivatives such as *N*-acetyl-dopamine (NADA) and *N*-β-alanyldopamine (NBAD) in mediating covalent crosslinks between cuticular proteins and/or between cuticular proteins and chitin fibers is exclusive to the sclerotization process [17,48]. Our data demonstrate that mosquitoes can metabolize L-DOPA via a non-canonical pathway using 3,4-dihydroxylphenylacetaldehyde (DOPAL) synthase (**Fig. 10A**). As a detoxification mechanism, L-DOPA is turned directly into DOPAL and hydrogen peroxide which further oxidizes the DOPAL catechol ring to form a melanized and flexible cuticle that is crosslinked to cuticular proteins while avoiding the conversion of L-DOPA to dopamine [59,60]. DOPAL buildup promotes oligomerization and quinone adduct formation ("quinonization") inhibiting its own metabolism and the activity of other enzymes involved in the melanin biosynthesis and supporting the observed downregulation of critical genes encoding enzymes involved in the melanization pathway (e.g., DDC, PPO5, DCE, and DOPAL synthase) [61] (**Fig 10B**).

Dietary L-DOPA affects multiple mosquito physiological pathways in a dose-dependent manner (**Figs. 3, 6, 7, 8, 9**). The strong allelopathic activity of L-DOPA may explain our observations [25], Depending on the cellular redox potential, L-DOPA behaves as a prooxidant (at alkaline pH) promoting oxidative damage and covalent modification of proteins, or as an antioxidant (at acid and neutral pH) scavenging initiating radicals by electrostatic interactions and chelating metal ions to prevent their participation in radical formation [62]. Our analyses demonstrate that L-DOPA-supplemented sucrose meals induced a biphasic dose-response (**Fig. 3B**). This dose-response relationship, known as hormesis, is observed in response to moderate environmental or self-imposed challenges, which causes a system to improve its functionality and/or tolerance when faced with more severe challenges [63,64]. The occurrence of hormesis is widespread in nature and has been reported for nitric oxide, prostaglandins, and dopamine, among other molecules[65]. Hormetic-like responses describe the simultaneous stimulation of many independent cellular functions/endpoints and define the limits of biological plasticity [64].

Despite marked downregulation of immune-related genes (**Fig. 5A**), oral L-DOPA promoted antimicrobial activity in *A. gambiae* female mosquitoes (**Fig. 6**). Our *in vitro* analyses demonstrate that above 0.5 mM, L-DOPA impairs *P. falciparum* development of both sexual and asexual stages, a finding that correlates with a previous report demonstrating induction of apoptosis to *P. berghei* ookinetes under similar conditions [66]. Conversely, given the particularity of our L-DOPA

diet (an amino acid analog within a low sucrose concentration) and the complexity of the mosquito digestion system (sugar and protein meals have distinct compartmentalization) [67], the enhanced resistance to parasite burden exhibited by L-DOPA-fed mosquitoes might reflect multifactorial causes. At least two possible scenarios, non-mutually exclusive, could explain the results. First, L-DOPA-mediated acidification of midgut pH could block parasite development at the midgut lumen thus reducing midgut invasion [68]; mean pH measurements of 1 mM L-DOPA-sugar solution dropped from 8.3 to 6.7 in 72 h (**Supplementary Fig 5**). Second, the restrictive sugar diet (1.5% sucrose) may induce a metabolic switch that functionally primes the mosquito to resist pathogen invasion [69]. In line with the latter scenario, the upregulation of the Dally protein (AGAP005799), phosphoinositide-3-kinase (PI3K) (AGAP005583), and juvenile hormone (JH) (AGAP008685) could reflect an integrative response from the insulin/TOR signal transduction pathway, sensing not only nutritional information but also other environmental cues such as temperature and foreign organisms to fine-tune allocation of resources and promote fitness [70-73].

In most insects, genes encoding enzymes involved in melanization and sclerotization are referred to as pleotropic [74-76], supporting the hypothesis that pleiotropy (i.e., expression of multiple traits by a single gene) drives the coevolution of correlated traits [77]. Such is the case for cuticle pigmentation, which was significantly darkened in 1 mM L-DOPA-fed mosquitoes (**Fig 7**). Its association with increased thickness in female mosquitoes immediately raises a concern for insecticide resistance given that several studies in disease vectors have reported this linkage [40,78-80]. Other mechanisms extensively shown to be involved in resistance to insecticides are alteration of target molecule, reduced cuticle permeability, and metabolic clearance via overexpression of detoxification protein families such as cytochrome P450 (CYPs), glutathione S transferases (GST), and esterases [81]. In agreement with *A. gambiae* susceptibility to deltamethrin upon L-DOPA feeding, these mosquitoes exhibit a drastic downregulation of GSTs, carboxylesterases, cuticular proteins, and CYPs including CYP4C27 (AGAP009246), which were previously associated with insecticide resistance in *A. gambiae* [82] (**Fig 5B**). It is worth mentioning that the slight increase of chitin content measured in whole L-DOPA fed mosquitoes corresponds to a mixture of chitin and chitosan, given that the samples were deacetylated before performing the colorimetric reaction. Regardless of the chemical processing, based on our previous work with melanized cryptococcal cell walls [39,83] and downregulation of chitin synthase 1 (CHS1A,B) (AGAP001748), we deduce that dietary L-DOPA induces an increase of chitosan within the mosquitoes legs which promotes melanin deposition and thus translates into augmented cuticular pigmentation. As predicted by the thermal melanism hypothesis [84], increased mosquito pigmentation was associated with augmented heat absorption for the 1 mM L-DOPA fed

mosquitoes (**Fig. 8A-B**) supporting an adaptative trait also described in insects from various orders [85-87]. *Anopheles* mosquitoes are poikilotherms, so ambient temperature strongly influences developmental processes, behaviors, and even survival [88]. These effects in thermoregulation could have an ecological impact; indeed, darker mosquitoes displayed a reduced longevity (**Fig. 8C**). This negative effect could be attributable to low tolerance to heat associated with starvation[46] given that our polyphenol-rich diet (e.g., dietary L-DOPA) with low sucrose concentration mimics calorie restriction [89].

Concomitant with dietary L-DOPA, our evidence that gut microbiota colonizing adult *Aedes* and *Anopheles* mosquitoes can synthesize dopamine (**Fig. 9B**) strongly suggests that the mosquito gut microbiome could also play a role in the effects reported here. Indeed, certain *Enterococcus* and *Bifidobacterium* species identified in in the gut of Parkinson's Disease patients can metabolize L-DOPA into dopamine or other metabolites like 3,4-dihydroxyphenyl pyruvic acid (DHPAA) [90,91]. The mosquito microbiome is a critical modulator of vector competence and vectorial capacity, also influenced by the mosquito diet [16,92,93]. The interactions between fungal and bacterial microbes shape fungal microbiome communities within the mosquito gut, which are dominated by Ascomycota and Basidiomycota. Among these are *Candida* and *Cryptococcus* spp.[50,51] which can use catechols (L-DOPA and dopamine) as substrates for melanin synthesis [52]. Therefore, as previously reported by other authors [94-96], midgut microbiota could also contribute to mosquito homeostasis by detoxifying toxic compounds that modulate symbiotic relationships between the gut microbiome and their host. In this line of thinking, the L-DOPA-modulated microbiome can also promote a certain pH at the midgut, thus favoring either prooxidant or antioxidant properties of L-DOPA [68]. Interestingly, recent research in *Aedes aegypti* demonstrated that higher levels of juvenile hormone (JH) in mated and virgin-JH treated in comparison to virgin mosquitoes promoted establishment of the midgut microbiota, midgut enlargement and allocation of energetic resources to enhance fecundity while inducing some immunosuppression. L-DOPA-fed female mosquitoes had no defects in reproduction (**Figs. 8D,E**). Recent evidence in the invertebrate model *Daphnia magna* showed that its gut microbiome is responsible for the synthesis of L-DOPA, which may serve as a signaling molecule modulating host-microbiome interactions relevant for key life traits [97]. The existence of a gut-to-brain signaling axis modulated by 5-hydroxytryptamine innervations in the midgut tissue of *Anopheles stephensi* mosquitoes was also recently suggested [98]. In addition, the contribution from hemocytes as key regulators of mosquito immunity cannot be ruled out. In *Drosophila*, hematopoietic progenitors can directly sense and respond to systemic (sugar/insulin) and nutritional (amino acids) signals via coordinated communication between organs [99].

Our findings could be incorporated into mosquito control strategies. Despite strenuous measures

to mitigate vector-borne diseases, these still account for more than 700,000 deaths annually, of

which over 50% are caused by malaria - 96% of them in Africa [100]. As proposed by other groups,

the use of genetically modified mosquitoes to control malaria offers great hope but also comes

with concerns and controversy [101,102]. To effectively address these global risks and safeguard the

biosphere on which all species depend, it is imperative to develop novel, safe, and biodegradable

insecticides. Solutions derived from natural plant products hold promise in mitigating the impact

of chemical pesticides on the environment. L-DOPA, the plant phenolic precursor of dopamine, is

a neurotransmitter involved in several complex behaviors and processes in invertebrates,

including locomotion, learning, courtship, development, and melanin synthesis [48,103,104]. Native

plants from Africa such as banana [24] (*Musa balbisiana* Colla) and Fava beans (*Vicia faba*) [21,23],

contains up to 5% of L-DOPA in their leaves, flowers, and fruit parts. This metabolite is also a

non-protein amino acid usually shown to be toxic to many insects [62,105] that could drive trade-offs

between enhancing tolerance and life-history traits [18,19,103]. By fostering the development and

implementation of eco-friendly alternatives such as L-DOPA supplemented sugar baits designed

to specifically lure defined mosquito species and using natural body odor chemoattracts [106], it may

be possible to develop a new tools for control of mosquito populations. Moreover, L-DOPA

supplementation of mosquito infected waters could enhance their immunity to reduce parasite

burden and shorten lifespan, thus interfering with lifecycle and reducing the opportunity for

parasite transmission.

In summary, feeding of a sugar meal supplemented with L-DOPA to adult *A. gambiae* mosquitoes

augmented cuticular melanization and reduced parasite burden and lifespan. In addition to direct

L-DOPA effects, increased melanization has the potential to affect insect thermoregulation.

Consequently, L-DOPA supplementation has the potential for development as a novel, effective,

and environmentally safe strategy to eliminate and/or reduce malaria transmission, which merits

additional investigation including field studies of its usefulness and feasibility.

**Methods**

**Ethical statement**

The Johns Hopkins University Animal Care and Use Committee has approved this protocol, with

permit number MO18H82. Commercial anonymous human blood was used for parasite cultures

and mosquito feeding, and informed consent was therefore not applicable. The Johns Hopkins School of Public Health Ethics Committee has approved this protocol. Mice for mosquito blood feeding were housed in environmentally controlled rooms at 72°F and 42% humidity, with a 14.5-h light/9.5-h dark cycle.

**Mosquito rearing**

Mosquitoes were obtained from colonies of *A. gambiae* (Keele strain) maintained in the insectary of the Johns Hopkins Malaria Research Institute (JHMRI) on a 10% sucrose solution with 12-h light/dark cycles at 27°C and 80% humidity either under normal, non-sterile conditions or placed into a sterile environment.

*Plasmodium falciparum* **culture**

*P. falciparum* NF54 asexual blood stage cultures were maintained *in vitro* according to the method described earlier by Trager and Jenson (PMID 351412). Briefly, parasites were cultured in O+ erythrocytes at a 4% hematocrit in RPMI 1640 (Corning) supplemented with 2.1mM L-glutamine, 25mM HEPES, 0.72 mM hypoxanthine (Sigma), 0.21% w/v sodium bicarbonate (Sigma), and 10% v/v heat inactivated human serum (Interstate blood bank). Cultures were maintained at 37°C in a glass candle jar. Gametocyte cultures were initiated at 0.5% asexual blood stage parasitemia and at 4% hematocrit and maintained by changing the media daily for 17 d without the addition of fresh blood to promote gametocytogenesis as described earlier [107]. Erythrocytes used for the study were obtained from healthy donors under a Johns Hopkins Institutional Review Board-approved protocol and were provided without identifiers to the laboratories.

**L-DOPA supplementation into sugar meal**

L-DOPA (3,4-Dihydroxy-L-phenylalanine) (Cat N° D9628) was obtained from Sigma-Aldrich. Sterile sucrose solutions at 3% were diluted with an equal volume of sterile L-DOPA solutions, and prepared from a 5 mM stock in distilled water to obtain sugar solutions supplemented with L-DOPA at concentrations ranging from 0.2 to 10 mM (0.4 to 20% w/w) L-DOPA. As control, 1.5% sucrose solutions were prepared.

**Survival assay**

Three- to four-day-old adult female *A. gambiae* were allowed to feed on sugar solutions prepared as described above. Survival of mosquitoes was monitored for 14 d, and dead mosquitoes were

removed daily. Solutions containing 1.5% sucrose (end concentration) with an equal ratio of L-

DOPA were freshly prepared and provided every 2-3 d.

**RNA extraction and transcriptional profiling**

Newly emerged adult female mosquitoes were allowed to feed with sugar solutions in the

presence and absence of 1 mM L-DOPA for 4 d. After this, mosquitoes were cold anesthetized to

remove their head using a tweezer over a glass plate on ice. Four groups of 10 headless

mosquitoes were transferred to a 1.5-ml tube. Midguts from four groups of 10 mosquitoes,

previously surface-sterilized, were dissected in a drop of sterile PBS and transferred to a 1.5-ml

tube containing 50 µl of sterile PBS. All samples were flash frozen in an ethanol/dry-ice bath and

stored at -80°C until RNA extraction.

Midguts and headless mosquitoes were homogenized in 1ml of Trizol with Lysing Matrix D Fast

Prep tubes in the FastPrep 24 (MP Bio), followed by RNA extraction using the PureLink RNA Mini

kit with on-column DNase treatment (ThermoFisher). RNA-Seq Libraries were prepared using the

Universal Plus mRNA-Seq Library prep kit (Tecan Genomics) incorporating unique dual indexes.

Libraries were assessed for quality by High Sensitivity D5000 ScreenTape on the 4200

TapeStation (Agilent Technologies). Quantification was performed with NuQuant reagent and by

Qubit High Sensitivity DNA assay, on Qubit 4 and Qubit Flex Fluorometers (Tecan

Genomics/ThermoFisher).

Libraries were diluted and an equimolar pool was prepared, according to the manufacturer's

protocol for appropriate sequencer. An Illumina iSeq Sequencer with iSeq100 i1 reagent V2 300

cycle kit was used for the final quality assessment of the library pool. For deep mRNA sequencing,

a 300 cycle (2 x 150 bp) Illumina NextSeq 500 (High) run was performed. Three-to-four biological

replicates were analyzed for each condition for a total of 15 sequencing libraries. RNA-seq data

was analyzed with Partek Flow NGS Software as follows: pre-alignment QA/QC; alignment to *A.*

*gambiae* Reference Index using STAR 2.7.8a; post-alignment QA/QC; quantification of gene

counts to annotation model (Partek E/M); filter and normalization of gene counts; and identification

and comparison of differentially expressed genes with DESeq2. Reference Genome: NCBI:

GCA_000005575.1_AgamP3. From the resulting gene lists, gene enrichment and pathway

enrichment were performed within Partek Flow (**Supplementary Fig. 1B**). Genes of interest were

validated by q-RTPCR. All sequence files and sample information have been deposited at NCBI

Sequence Read Archive, NCBI BioProject: PRJNA1161683.

***A. gambiae* infection with *P. falciparum*.**

Newly emerged adult female mosquitoes were allowed to feed on sugar solutions with up to 1 mM L-DOPA or dopamine for 4 d. On the fifth day, sugar meals were removed, and mosquitoes were starved for 5-6 h prior to offering a *Plasmodium falciparum* NF54-laced blood meal with gametocytemia 0.03%. Unfed mosquitoes were sorted, and blood-engorged mosquitoes were further incubated for 8 d at 27 °C for oocyst counting. Midguts from *Pf*-infected mosquitoes were dissected, stained with a 0.2% mercurochrome solution, and mounted on a slide with a coverslip gently pressed on top to facilitate oocyst visualization. Enumeration of oocysts was performed using a 20X objective under a Leica light microscope. For infections under aseptic conditions, the artificial glass feeder membrane-feeding process was conducted under conditions as sterile as possible to prevent antibiotic-treated mosquitoes from acquiring a bacterial infection [92].

***In vitro* susceptibility of *P. falciparum* NF54 to L-DOPA**

To determine *P. falciparum* toxicity to L-DOPA, solutions at concentration up to 2 mM were freshly prepared from a 5 mM L-DOPA stock. *P. falciparum* NF54 asexual and gametocyte stage cultures were dispensed into 96 well plates and triplicate wells were treated with vehicle or L-DOPA at concentrations of 0.25, 0.5, 1 and 2 mM for a total of 48 h. Blood smears were prepared after drug treatment and asexual parasitemia or gametocytemia was calculated microscopically from Giemsa-stained slides. The effect of L-DOPA exposure was determined by calculating % decrease in number of parasites as compared to vehicle control.

**Fungal challenge**

*Cryptococcus neoformans* (strain H99-GFP) was cultured for 48 h in rich media at 30 °C with moderate shaking. Yeast cells were collected, washed twice with PBS, and adjusted to a final concentration of $1 \times 10^7$ cells/ ml.  In parallel, newly emerged adult female mosquitoes were allowed to feed with sugar solutions in the presence or absence of 1 mM L-DOPA for 4 d. Using a nanoinjector, adult mosquitoes were injected intrathoracically with 69 µl of the fungal suspension and then incubated at 19°C. To assess the midgut fungal infection, mosquitoes were surface sterilized 24 h post injection in 70% ethanol for 3 min and rinsed twice in PBS for 2 min each time. Individual mosquitoes were transferred to 1.5-ml tubes containing 100 µl PBS and homogenized for 20 s using a sterile pestle. Two serial dilutions of each sample were plated in Bird Seed agar (HiMedia Laboratories, PA), a selective media for isolation and differentiation of *Cryptococcus* spp, and incubated at 30°C for 7 d, after which the number of fungal colony-forming units (CFUs)

were counted. Mosquito abdomens were also dissected to visualize the presence of live or

melanized *C. neoformans* yeasts, and images were captured using a Leica DM 2500 microscope.

**Cuticle pigmentation**

Newly emerged adult female mosquitoes were allowed to feed with L-DOPA solutions for up to 14

d as described above. For photography, mosquitoes were cold anesthetized, placed ventral side

up on a slide under a dissection microscope, and imaged with an Iphone X under constant

exposure and lighting conditions. Image analysis and pigment measurement were done as

described previously [108]. Using the Fiji software [109], RGB color images were converted to 8-bit

color, entire abdomens were selected using a freehand selection tool, and mean gray values were

analyzed with the Measure tool. Pure black corresponds to 0 gray value whereas pure white equals

255.

**Transmission Electron Microscopy**

Newly emerged adult mosquitoes were allowed to feed with sucrose solutions, with and without

supplementation of 1 mM L-DOPA for 7 d. Groups of 10 midlegs from each condition were

dissected on a glass petri dish. To reduce hydrophobicity, legs were rinsed once in 70% ethanol,

twice in PBS for 1 min each step, and then incubated overnight at 4°C in fixing solution containing

2% glutaraldehyde, 0.1 M sodium cacodylate (pH 7.4) and 2% paraformaldehyde in distilled

water. After buffer rinse, samples were postfixed in 1% osmium tetroxide, 0.8% potassium

ferrocyanide in 0.1 M sodium cacodylate for at least one hour (no more than two) on ice in the

dark. After osmium, samples were rinsed in 100 mM maleate buffer, followed by uranyl acetate

(2%) in 100 mM maleate  (0.22 μm filtered, 1 h dark), dehydrated in a graded series of ethanol

and embedded in a SPURR's/Eponate 12 (Ted Pella) resin. Samples were polymerized at

60C overnight. Thin sections, 60 to 90 nm, were cut with a diamond knife on the Reichert-Jung

Ultracut E ultramicrotome and picked up with 2x1 mm formvar copper slot grids. Grids were

stained with 2% uranyl acetate in 50% methanol followed by lead citrate and observed with

a ThermoFisher Talos L120C at 120 kV.   Images were captured with a ThermoFisher Ceta CCD

(16 megapixel CMOS, 16-bit).

**Quantification of $^{13}$C L-DOPA Melanin by Solid-State NMR**

Newly emerged adult mosquitoes were allowed to feed with sucrose solutions, with and without supplementation with 1 mM of stable-isotope enriched $^{13}C$ L-DOPA (RING-$^{13}C_6$, 99%, CLM-1007-0, Cambridge Isotope Laboratories, MA, USA) for 7 d. Next, mosquitoes were sorted by sex, collected in 15-ml Falcon tubes, and lyophilized for 3 d. Solid-state NMR measurements were conducted on 8-10 mg samples using a Varian (Agilent) DirectDrive2 spectrometer operating at a $^{13}C$ frequency of 150 MHz and equipped with a 1.6-mm T3 HXY fast magic-angle spinning (fastMAS) probe spinning at 15.00 ± 0.02 kHz and a regulated temperature of 25 °C. $^{13}C$ NMR spectra of the lyophilized mosquito samples were obtained with 90° pulse lengths of 1.2 and 1.4 µs for $^1H$ and $^{13}C$, respectively; 104-kHz heteronuclear decoupling using the small phase incremental alternation pulse sequence (SPINAL[110]) was applied during signal acquisition. Whereas a 10% linearly ramped $^1H$ amplitude, 1-ms time for $^1H$-to-$^{13}C$ polarization transfer, and 3-s delay between acquisitions were used to acquire CPMAS spectra that identified the carbon-containing functional groups, a longer 50-s delay was used for direct polarization (DPMAS) experiments that generated spectra with quantitatively reliable signal intensities. The CPMAS and DPMAS spectral runs required 4 and 56 hours, respectively.

**Chitin measurements**

The chitin content of adult mosquitoes post L-DOPA feeding for 7 d was determined based on the method of Lehmann and White (1975), [111] adapted as described previously used to quantify the chitin content of the cryptococcal cell wall material [112]. Using 15-ml tubes, 60 and 80 cold-anesthetized female and male mosquitoes, respectively, were freeze-dried, and lyophilized for 2-3 d. Mosquitoes were transferred to 1.5-ml microcentrifuge tubes, weighted (typically 14 and 11 mg, respectively), and homogenized in 200 µl of distilled water using a battery-operated homogenizer (3 20-sec cycles) and a horned sonicator at half amplitude (2 30-sec cycles). To solubilize and alkali-extract the samples, homogenates were centrifuged at 21,000 $g$ for 5 min at room temperature and the pellet of each sample was resuspended in an equal volume (200 µl) of 3% (w/v) of sodium dodecyl sulfate (SDS). The samples were then incubated at 100 °C for 15 min and centrifuged again for 5 min after cooling. Next, after each pellet was washed twice with 200 µl distilled water, it was resuspended in 150 µl 2.1 M KOH. Samples were incubated at 80 °C for 90 min and cooled down to room temperature. After 400 µl 75% ethanol was added to each sample, the sample was mixed and incubated on ice for 15 min. Sixty µl of a Celite 545 (Sigma-Aldrich) suspension (1g of Celite 545 in 12.5 ml 75% ethanol, rested for 2 min) was then added to each sample and centrifuged. The resultant cell matter was washed thoroughly once with 40%

ethanol, twice with PBS (pH 7), twice with distilled water, and resuspended in distilled water to a final concentration of 10 mg ml $^{-1}$. To estimate the total amount of glucosamine stemming from chitin plus chitosan, one 100 µl aliquot (equivalent to 1 mg dry weight) was transferred into a 2 ml self-locking microcentrifuge tube and subjected to deacetylation with 100 µl 1 M HCl in a heating block at 110 °C for 2 h. As a blank, one 100 µl of milliQ water was used instead of the mosquito suspension. In a fume hood, sample deamination was achieved by adding 400 µl 2.5% NaNO$_2$, then mixed and incubated for 15 min at room temperature to allow nitrogen oxides to dissipate. After that, 200 µl of 12.5% ammonium sulfate was slowly added to each sample, vortexed, and let stand at room temperature for 5 min. Measurement of hexosamines (amino sugars) from the deacetylated samples was done by treatment with 3-methyl-2-benzothiazolone hydrazine hydrochloride (MBTH, Sigma-Aldrich) under mildly acidic conditions. A 200 µl aliquot from each sample was transferred to a 96-well plate and the absorbance at 650nm was recorded. Concentrations were determined by comparison to a standard curve prepared from a 1mM GlcNAc stock (Sigma-Aldrich) and ranging from 0 to 100 nM.

**Tarsal contact bioassay**

A susceptibility test against deltamethrin (0.05%) was done following the WHO protocol [113]. The choice of deltamethrin, a pyrethroid insecticide, was justified by being one of the only two insecticide classes approved for use on insecticide-treated bed nets (ITNs). Using an aspirator 22-25 *A. gambiae* female mosquitoes aged 2-5 d and previously allowed to feed on sucrose with or without 1 mM L-DOPA supplementation for 4 d were introduced into two WHO holding tubes (one test and one control) that contained untreated papers. Mosquitoes were allowed to settle inside the tube for 15 min in an upright position, then with a quick and gentle tap mosquitoes were transferred into the exposure tubes containing the insecticide-impregnated papers. Following one-hour exposure at room temperature, mosquitoes were transferred back into holding tubes, provided with a cotton pad moistened with 10 % sugar, and incubated at 27 °C. The number of dead mosquitoes at 24 h was recorded. Control tubes contained oil-treated paper.

**Electron Paramagnetic Resonance (EPR) Spectroscopy**

Acid-resistant material attributable to melanin was isolated from *A. gambiae* mosquitoes fed with and without 1 mM L-DOPA supplementation for 7 d following the protocol described previously [114]. *Aedes aegypti* (mutant strain Yellow) and fungal eumelanin from *C. neoformans* were used as negative and positive controls, respectively. EPR measurements were conducted at the

Caltech EPR Facility using a Bruker EMX X-band continuous-wave (CW) EPR spectrometer equipped with a Bruker SHQE resonator. Samples were measured at room temperature under non-saturating microwave power (4.4 μW) using a field modulation amplitude of 1 Gauss and conversion time of 5 ms.

**Infrared thermography for mosquito bodies**

L-DOPA fed mosquitoes for 7 d were cold anesthetized, clustered into groups of ten on top of a cooling block set at 0°C and arranged from left to right for increasing L-DOPA concentrations for up to 1 mM. Basal mosquito temperatures were recorded in dark conditions using an infrared (IR) camera (FLIR Systems, Wilsonville, OR) mounted 40 cm above the cooling block setup on a benchtop in a temperature-controlled room (22 ± 5°C, 50% relative humidity). Irradiations were conducted for 4 min using a white LED lamp (250 W, 1850 Lumens, PAR38HO-E26-19W-4000K-25°, Green Creative) mounted 40 cm above the cooling block and resulting in an average luminance of ~415 LUX. The optimal LED bulb positioning for homogeneous light exposure was calculated based on the light intensity distribution using graph paper and a light meter (**Supplementary Fig. 5**).

**Fecundity, egg hatching rate and life span.**

For the fecundity assay, newly emerged adult female *A. gambiae* were allowed to feed on a sugar solution supplemented with 1mM L-DOPA for 7 d, whereas the controls were fed 1.5% sucrose without L-DOPA. Mosquitoes were allowed to blood-feed on mice for 20 min and then anesthetized on ice; all non-engorged mosquitoes were discarded. At 3 d post blood meal, individual mosquitoes were transferred to a 50-ml Falcon tube containing a paper cup in the bottom halfway submerged into clean distilled water for oviposition. The number of eggs laid by each mosquito was recorded for up to 4 d under a light microscope. Female mosquitoes that died before oviposition were not included in the study. After each count, adult mosquitoes were removed from the vials, eggs were completely submerged into larval rearing water, and vials were incubated for 3-4 d at 27°C and 30°C, making sure that the paper cups remained moistened. The first and second instar larvae were counted under a light microscope.

The life span of *A. gambiae* females during continued exposure to a sugar meal supplemented with 1 mM L-DOPA was measured by placing newly emerged mosquitoes into paper cups with a sugar solution that was freshly prepared every 2-3 d. Dead mosquitoes were recorded and

removed daily until no live mosquitoes remained. Survival analysis represents pooled data from three biological replicates of 35 mosquitoes each.

**Dopamine measurements**

Dopamine quantification was conducted from whole female mosquitoes post L-DOPA feeding and from bacterial supernatants.

**Sample collection**: For mosquito samples, newly emerged adult mosquitoes were allowed to feed with sugar solutions supplemented with L-DOPA at three different concentrations (0.2 mM, 0.6 mM, and 1 mM) for 4 d. At the assayed time, five whole female mosquitoes were collected in 1.5-ml brown tubes and flash-frozen in an ethanol-dry ice bath. Mosquito supernatants containing both hemolymph and cuticle dopamine were obtained by adding 200 µl ice-cold 0.2 M perchloric acid to each tube, homogenizing for 20 s on ice, and centrifuging at 20,000 g for 10 min at 4 °C [115]. Supernatants (~100 µl) were transferred to a new brown 1.5-ml tube and kept at -80 °C until further processing was conducted. For bacterial samples, a single colony of each bacterial strain, previously shown to induce fungal melanization, was grown in LB broth at 30 ˚C for 96 h. Filter sterilized, cell-free culture supernatants were acidified to pH 4 with 1 M HCl, fractionated by FPLC using an AKTA Pure liquid chromatograph with a UV-900 monitor to measure absorbance at 214 nm (Amersham Biosciences) and a C18 reverse-phase column (Phenomenex Luna 5 µm, 100 Å, LC Column 150 x 4.6 mm) using an isocratic mobile phase of 25 mM potassium phosphate buffer (pH 3.1) at a flow rate of 1.0 ml/min [49]. Fractions of 200 µl were collected into a new brown 1.5-ml tube and kept at -80 °C until further processing was conducted; a 50 mM dopamine (Sigma-Aldrich) solution was used as standard.

**Sample preparation and derivatization:** Dopamine standards were prepared in 10 µg/mL sodium metabisulfite in water. Samples were derivatized as described previously, with the following adjustments: 20 µL of standards and samples were transferred to an extraction plate; 10 µL of 100 mM sodium tetraborate in water and 5 µL of a cold internal standard solution consisting of 500 ng/mL dopamine-d3 in acetonitrile (v/v) were added to each sample and mixed for 1 min on plate shaker at 1,600 rpm; 10 µL of 2% benzoyl chloride in acetonitrile (v/v) was then added to each sample and the plate was mixed for 2 min at 1,600 rpm. 10µL of 1% formic acid in water (v/v) was added to quench the reaction. 1µL of each sample was injected for analysis.

**Sample analysis:** Samples were analyzed using an Agilent 6540 QTOF with a Jet Stream Electrospray Ionization Source (ESI) and an Agilent 1290 UHPLC (Agilent Technologies, Santa

Clara, CA). Solvents were 10 mM ammonium formate in water (A) and acetonitrile (B). The samples were kept at 10°C during analysis. Chromatographic separation was achieved over 3.0 minutes using a Phenomenex Luna Omega 2.1 x 100mm, 1.6μm, C18 column at 40°C with a binary gradient starting with 10% B. The LC gradient was as follows: 0–1.2 min, 10% - 95% B; 1.2 - 2.0 min, 95% B; 2.01, 10% B. The flow rate was 400 μl min$^{-1}$. Mass spectral acquisition was performed using full scan MS from *m/z* 100 to 1000 with the following source conditions: Drying and Sheath Gas temperatures of 350°C and 400°C, respectively; both gas flows at 12L/min; Nebulizer- 45psig, VCap, Nozzle and Fragmenter voltages at 3000 V, 600 V and 100 V, respectively. Proton adducts of the triple benzoylated dopamine and dopamine-d$_3$ (*m/z* 466.1649 and 469.1837, respectively) were extracted within a +/- 20 ppm window. Analyte peak areas were determined from the extracted ion chromatograms and concentrations were calculated by linear regression analysis using Agilent Masshunter Quantitative Analysis Software (B.08.00). Limit of quantification was 1.0 ng/ml.

**Plate confrontation assays**

As a proxy for the ability to synthesize catecholamines from bacteria isolated from the midgut of wild-caught *An. aranbiensis* and *Ae. aegypti* adult mosquitoes as well as an environmental source relevant to mosquito aquatic stages, we monitored Cryptococcal melanization as previously reported [49].

**Statistical Analysis**

Data were graphed and analyzed for statistical significance using GraphPad Prism 10 (version 10.0.1) software (GraphPad Software, LLC). Particular tests used for experiments are indicated in the legend of each respective figure.

**Data availability**

All data generated or analyzed during this study are included in this published article (and its Supplementary files). The RNAseq data was submitted to the Sequence Read Archive (SRA) database (PRJNA1161683).

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## Author contributions

E.C.: conceptualization, methodology, validation, formal analysis, investigation, project administration, funding acquisition, writing—original draft, writing-review and editing, visualization. Y.D., C.C. R.J.B.C. R.G.S., Y.A-R., D.F.Q.S., E.J., I.H., J.A.P-M., M.DP., A.D., A.J., B.S., G.M., A.T.: methodology, validation, formal analysis, investigation, writing— writing-review and editing, visualization. N.A.B.: Resources, writing-review, and editing. R.E.S.: Resources, supervision, writing-review and editing, and funding acquisition. D.G.: Resources, supervision, validation, writing-review, and editing. A.C.: conceptualization, supervision, project administration, resources, supervision, writing-review and editing, and funding acquisition.

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## Tables and Figures

**Table 1.** Detection of melanin precursors released by diverse bacterial species.

**Fig 1. Experimental overview for investigating the impact of L-DOPA-supplemented sugar meal on *A. gambiae* mosquitoes.**

**Fig 2. Melanization pathways in insects.** In *Anopheles* female mosquitoes, melanin synthesis relies mostly on dopamine produced from the blood meal-derived phenylalanine; however, other precursors from plant sources such as tyrosine or L-DOPA are used by both female and male mosquitoes for the generation of this biopolymer. Most melanins synthesized in insects derive from L-DOPA or dopamine in a reaction initiated by phenoloxidases present in the hemolymph; however, there is one possible minor pathway from a degradation product of dopamine: 3,4-dihydroxyphenylaldehide (DOPAL). DOPAL is a highly unstable compound that reacts with primary amines, leading to protein crosslinking and inactivation. Current literature suggests that its toxicity outweighs its benefits; evidence for DOPAL melanin formation in any biological system remains to be established. Simultaneously but via an independent pathway, dopamine serves as substrate for a process known as sclerotization which promotes hardening of insect cuticles (exoskeleton) and prevents desiccation, dehydration, and invasion by foreign organisms. This process is catalyzed by cuticle-bound phenoloxidases [12,17]. Figure created with Biorender.com

**Fig 3. L-DOPA supplementation exerts a hormesis effect. A)** Daily proportion of surviving mosquitoes after L-DOPA feeding in 1.5% sugar meal at various supplementation concentrations. Each survival curve represents pooled data from three independent experiments (n= 75). Survival curves for L-DOPA-fed mosquitoes were compared statistically to the survival curves of control mosquitoes using Log-rank Mantel-Cox tests (NS, not significant, $*p = 0.0113$, $**p = 0.0040$, $****p < 0.0001$). **B)** Cumulative survival percentage at days 4, 8 and 14 shows how mosquitoes adapt to the toxic effects of L-DOPA at concentrations below and above 1 mM over time. **C)** Daily proportion of surviving mosquitoes after dopamine feeding in 1.5% sugar meal at various supplementation concentrations. Each survival curve represents pooled data from three independent experiments (n= 75). Survival curves for dopamine-fed mosquitoes were compared statistically to the survival curves of control mosquitoes using Log-rank Mantel-Cox tests (NS, not significant, $****p < 0.0001$). **D)** Cumulative survival percentage at days 4, 8 and 14 shows highly toxic effects of dopamine over time and striking death observations for sugar-fed control mosquitoes.

**Fig 4. Comparative transcriptome analysis of *A. gambiae* gene expression for L-DOPA-fed and sugar-fed female mosquitoes.** **A)** Volcano plot of differentially expressed genes (DEGs) in whole (headless) mosquitoes. Significantly upregulated and downregulated genes are highlighted in red and blue spots, respectively. **B)** DEGs by gene ontology category in whole L-DOPA-fed female mosquitoes compared with sugar-fed (control) females. Differential gene expression was considered significant when the expression fold change (L-DOPA-fed versus sugar-fed) was ≥1.3 on a -$\log_{10}$ scale (P < 0.05). Categories are R/S/M, oxidoreductive, stress-related, and mitochondrial **C)** Volcano plot of differentially expressed genes (DEGs) in mosquito midguts. Significantly upregulated and downregulated genes are highlighted in red and blue spots, respectively. **D)** DEGs by gene ontology category in midguts from L-DOPA-fed female mosquitoes compared with sugar-fed females. Differential gene expression was considered significant when the expression fold change (L-DOPA-fed versus sugar-fed) was ≥1.3 on a -$\log_{10}$ scale (P < 0.05). Categories are R/T/T, replication, transcription, and translation; R/S/M, oxidoreductive, stress-related, and mitochondrial.

**Fig 5. Differential transcript abundances of melanin-related physiological processes in *A. gambiae* female mosquitoes following feeding of a sugar-meal supplemented with 1 mM L-DOPA for 4 d.** $\log_2$ scale indicates transcripts that were found at significantly higher (red) or lower (blue) levels in whole (headless) mosquitoes (upper panel) and midguts (lower panel) from *A. gambiae* post L-DOPA-feeding versus sugar-fed (controls). **A)** Set of immune-related transcripts. **B)** Set of cuticle-related transcripts.

**Fig 6. L-DOPA exhibits antiparasitic and antifungal activity. A)** Intensity of *P. falciparum* oocyst load according to L-DOPA concentration in adult female mosquitoes. Each circle indicates the number of oocysts on an individual midgut; horizontal bars represent the median value of oocysts. n=3, NS, not significant, **$p$ = 0.0081, One-way analysis of variance (ANOVA) nonparametric Kluskal-Wallis test. Statistical significance for prevalence of infection was assessed with a Fisher's exact test, ns. **B)** Intensity of *P. falciparum* infection according to dopamine concentration in adult female mosquitoes. n=3, NS, not significant, **$p$ = 0.0019, One-way ANOVA nonparametric Kluskal-Wallis test. Statistical significance for prevalence of infection was assessed with a Fisher's exact test, ns. **C)** *In vitro* assessment of L-DOPA toxicity to sexual and asexual stages of *P. falciparu*m parasite. Use of L-DOPA concentrations ranging from 0.25 mM to 2 mM during in vitro parasite culture, appeared more toxic to asexual stages of the parasite than to sexual stages. n=3, NS, not significant, ****$p$< 0.0001, Two-way ANOVA with Dunnett's multiple comparisons tests against individual controls. **D)** Fungal load in L-DOPA-fed female

mosquitoes shows a reduction compared to control. Each circle corresponds to a whole mosquito

homogenate; horizontal red bars represent the median value of fungal CFUs. n=2, NS, not

significant, unpaired nonparametric Mann-Whitney test. **E)** Cryptococcal yeasts are successfully

melanized (red arrow) in the abdomens of *A. gambiae* female mosquitoes 24 h post-injection of

*C. neoformans* H99-GFP strain (1 x $10^7$ yeasts/ml). Live yeast cells (green arrows), Dead yeast

cells (red arrows). **F)** qRT-PCR analysis of mRNA abundance in whole mosquitoes from key

genes [dopa decarboxylase (DDC), gambicin (GAM), phenoloxidase 1 ( PPO1), and leucine-rich

immunomodulator 15 (LRIM15)] involved in the mosquito melanization pathway and *Plasmodium*

parasite development. Data represent the mean ± SE of three or more independent replicates,

analyzed using unpaired by one-way ANOVA and multiple comparison test to determine gene

expression relative to housekeeping S7 gene. NS, not significant, **$p$ = 0.0041, *$p$ = 0.0235.

**Fig 7. Oral L-DOPA boosts mosquito cuticular melanization via a non-canonical pathway.**

**A)** Dietary L-DOPA enhances cuticle pigmentation of *A. gambiae* female mosquitoes in a dose-

dependent manner. *Upper panel*, Representative images of *A. gambiae* female mosquitoes fed

with a sugar meal supplemented with different concentrations of L-DOPA for up to 14 d. Scale

bar, 1 mm; *Lower panel*, Quantification of cuticle darkening at 8 and 14 d. Each dot corresponds

to an individual mosquito. Representative data from 2-3 biological replicates. Statistical

significance was determined by an unpaired two-tailed *t*-test with two or more data sets (Left

panel, *$p$ = 0.0342; Right panel, ns = not significant, *$p$ = 0.0189, ****$p$ < 0.0001) **B)** *Top panel,*

TEM micrographs exhibiting distinct electron-dense bands within the endocuticle of L-DOPA-fed

mosquitoes that are not seen in the control. Scale bar, 500 nm. *Lower panel,* Measurements of

cuticular thickness from legs of female mosquitoes fed with L-DOPA supplemented sugar meals

and control show increased values for exo and endocuticle.****$p$<0.0001; Two-way analysis of

variance (ANOVA) with Tukey's multiple comparisons test. **C)** L-DOPA supplementation

augments content of indole groups. *Left panel*, DPMAS $^{13}$C ssNMR spectra of $^{13}$C-L-DOPA fed

mosquitoes display pigment peaks (green arrows) with greater relative intensities than their

respective controls. Spectra were normalized to the tallest peak (lipid peak at ~30 ppm); Right

panel, CPMAS $^{13}$C ssNMR spectra of $^{13}$C-L-DOPA fed mosquitoes corroborate the assertion that

$^{13}$C-L-DOPA has been nearly-exclusively used to make melanin instead of being broken down

and used as a nutrient source. Spectra were normalized to the tallest peak (polysaccharide peak

at ~72 ppm). Representative data from 2 biological replicates. **D)** Quantification of chitin content

in whole *A. gambiae* female mosquitoes demonstrates no significant differences between sugar-

fed (control) and L-DOPA-fed mosquitoes. Data correspond to 3 biological replicates. Floating

bars (min to max) Line at mean. NS, not significant; unpaired, nonparametric Mann-Whitney test.

**E)** EPR spectroscopy analysis of acid-resistant material isolated from *An. gambiae* and *Ae.*

*aegypti* yellow mutant strain L-DOPA-fed (red spectra) and sugar-fed (blue spectra).

Representative data from 2 biological replicates. Fungal eumelanin was used as positive control

(black spectrum).

**Fig. 8. Dietary L-DOPA darkening of cuticle increases heat-exposed mosquito**

**temperatures following light exposure and decreases lifespan. A)** *A. gambiae* female

mosquitoes fed with a sugar meal supplemented with different concentrations of L-DOPA (0.2

mM to 1 mM) for 8 d displayed a dose-dependent ability to absorb heat. **B)** Pre-exposure to a

direct light source, 1mM L-DOPA- fed mosquitoes (blue circle) exhibited a warmer body

temperature than sugar-fed control (red square). Post-irradiation for 4 min, L-DOPA-fed

mosquitoes (blue circle) were distinctly hotter than sugar-fed controls (red square), starting to cool

down but remaining warmer than control. **C)** Lifespan of mosquitoes maintained on 1.5% sugar

with 1 mM L-DOPA was diminished in comparison to sugar-only fed mosquitoes. Representative

data from 3 biological replicates. **D)** Number of eggs laid by female mosquitoes after feeding with

a sugar meal supplemented with 1 mM L-DOPA during 4 d before a blood meal on mice. Mann-

Whitney test, NS, not significant, $p$ = 0.1987. Red lines denote mean. **E)** Hatching rate at 27 °C

indicating average percentage of eggs giving rise to 1st and 2nd instar larvae. Mann-Whitney test,

NS, not significant. $p$ = 0.9390. Red lines denote mean.

**Fig. 9 Dietary L-DOPA metabolism and probing for dopaminergic bacteria amongst gut**

**microbiota from field-caught mosquitoes.   A)** Quantification of dopamine by HPLC from whole

*Anopheles* mosquitoes post 4 d of dietary L-DOPA suggests tight regulation of L-DOPA

conversion into dopamine.  Data correspond to 2 independent biological replicates from 5-pooled

mosquitoes per condition. A 2-way analysis of variance (ANOVA) with Tukey's multiple

comparisons test showed no significant differences among and between groups. **B)** Plate

confrontational assays showed induced fungal melanization due to melanin precursors released

by bacterial isolates. A) *Comamonas* sp.; B) *Enterococcus hormaechei*; C) *Aeromonas*

*hydrophila*; D) *Serratia marcescens* E) *Chromobacterium subtsugae* Δ*vioS* F) *Enterococcus* sp.

*Cryptococcus* strains (top: *C. gattii* WM179; middle: *C. neoformans* ΔLAC1; bottom: *C.*

*neoformans* H99) were streaked perpendicular to bacterial isolates in SM plates. The plates were

incubated at 30˚C for 5-7 days. Images are representative of 3 biological replicates.

**Fig. 10 Alternative pathway for L-DOPA incorporation to the melanin synthesis in mosquitoes. A)** L-DOPA serves as a substrate for cuticular melanization via DOPAL synthase as a detoxification mechanism given a cellular redox environment that promotes L-DOPA antioxidant properties (neutral or acid pH) [62], therefore preventing its conversion to dopamine via a dopa decarboxylase. L-DOPA conversion into DOPAL leads into DOPAL buildup that promotes oligomerization and quinone adduct formation ("quinonization") inhibiting its own metabolism and the activity of other enzymes involved in the melanin biosynthesis such as dopa decarboxylase (DDC), phenoloxidase 5 (PPO5), and dopachrome conversion enzyme (DCE)[61]. **B)** Model for dietary L-DOPA-mediated cuticular melanization proposes that *Anopheles gambiae* female mosquitoes use L-DOPA supplement in the sugar meal as a substrate for synthesizing cuticular melanin. L-DOPA is internalized into the midgut epithelium and exported to the hemolymph by sharing the same transporters with amino acids such as $Na^+$-driven amino acid transporter, alpha2 transporter, and/or SLC protein. Within regions of flexible cuticle, in the presence of DOPAL synthase, L-DOPA is turned into DOPAL and $H_2O_2$ promoting oxidation of DOPAL catechol ring that results in the formation of a melanized and protective flexible cuticle via crosslinking to cuticular proteins[59,60]. The contribution of gut microbiota as well as from hemocytes as key regulators of mosquito vectorial capacity and immunity, post a L-DOPA diet is under investigation. Figure created with Biorender.com

**Supplementary Table 1. Differentially Expressed Genes between 1mM L-DOPA-fed and sucrose-fed *A. gambiae* female mosquitoes. 10.6084/m9.figshare.27122907**

**Supplementary Fig 1. Partek Flow analysis. A)** Principal component analysis (PCA) of 4 L-DOPA-fed mosquito groups and 3 sugar-fed controls. Blue spheres represent control replicates; red spheres represent L-DOPA-fed replicates. **B)** Top 20 KEGG Pathway enrichment analysis of transcriptional data.

**Supplementary Fig 2. L-DOPA supplementation also enhances cuticular melanization in *Anopheles gambiae* male mosquitoes. A)** ^{13}C DPMAS NMR spectra of ^{13}C-L-DOPA fed mosquitoes display pigment peaks (green arrows) with greater relative intensities than their controls. Spectra were normalized to the tallest peak (lipid peak at ~30 ppm). **B)** *Top panel,* TEM micrographs displaying distinct electron-dense bands within the endocuticle of L-DOPA-fed mosquitoes not seen in those of control. Scale bar, 500 nm. *Lower panel,* Measurements of cuticular thickness from legs of male mosquitoes fed with a 1 mM L-DOPA supplemented sugar

meal show reduced exocuticle in comparison to controls (sugar-fed). NS, not significant; ****$p <$

0.0001; Two-way analysis of variance (ANOVA) with Tukey's multiple comparisons test.

**Supplementary Fig 3. Mapping white LED intensity distribution on the cooling block and**

**mosquitoes' temperature variance. A)** Light intensity (LUX) distribution on the sample platform

shows that luminous flux decreases around the edge of the cooling block. **B)** Apparent

temperature of mosquitoes fed with up to 1 mM L-DOPA consistently demonstrated that 1 mM L-

DOPA-fed mosquitoes exhibited a warmer temperature than sugar-fed controls.

**Supplementary Fig 3. Mapping white LED intensity distribution on the cooling block and**

**mosquitoes' temperature variance. A)** Apparent temperature of group of 20 mosquitoes fed

with up to 1 mM L-DOPA demonstrated that 1 mM L-DOPA-fed mosquitoes exhibited a warmer

temperature than sugar-fed controls (Trial 1). **B)** Apparent temperature of group of 20 mosquitoes

fed with 1 mM L-DOPA consistently demonstrated that 1 mM L-DOPA-fed mosquitoes exhibited

a warmer temperature that sugar-fed controls (Trial 2). **C)** Apparent temperature of individual

mosquitoes (n = 6) fed with 1 mM L-DOPA consistently demonstrated that 1 mM L-DOPA-fed

mosquitoes exhibited a warmer temperature that sugar-fed controls (Trial 3). **D)** Light intensity

(LUX) distribution on sample platform showing that luminous flux decreases around the edge of

the cooling block.

**Supplementary Fig 4. Gut microbiota can secrete dopamine.** Chromatographs of cell-free

bacterial supernatants after growth for 24, 48, 72 and 96 h Luria Bertani media at 30°C.

Overlapped dotted chromatograph corresponds to 1 mM spike dopamine. Red arrows show

points at which samples were collected. The elution time was measured in minutes and the

absorbance is measured at 214 nm in arbitrary units. For Fast Protein Liquid Chromatography

(FPLC), we used a C18 Reverse-Phase column to observe the dopamine peaks. 25 mM

Potassium Phosphate buffer (pH 3.1) was used as an isocratic mobile phase at a flow rate of 1

ml/min.

**Supplementary Fig 5. L-DOPA-sucrose solutions pH monitored over time.** Measurement of

pH from sucrose solution while incubated at 27°C inside the mosquito chamber displays

progressive acidification. Supplementation with L-DOPA accentuates this trend. Data

corresponding to three independent measurements for all conditions. Bars correspond to mean

with 95% confidence interval. Statistical analyses were performed using a Two-way analysis of

variance (ANOVA) with Sidak's multiple comparisons test.

# References

Klepac, P. *et al.* Climate change, malaria and neglected tropical diseases: a scoping review. *Trans R Soc Trop Med Hyg*, doi:10.1093/trstmh/trae026 (2024).

Caminade, C. *et al.* Impact of climate change on global malaria distribution. *Proc Natl Acad Sci U S A* **111**, 3286-3291, doi:10.1073/pnas.1302089111 (2014).

Kulkarni, M. A. *et al.* 10 Years of Environmental Change on the Slopes of Mount Kilimanjaro and Its Associated Shift in Malaria Vector Distributions. *Front Public Health* **4**, 281, doi:10.3389/fpubh.2016.00281 (2016).

Bagcchi, S. Locally acquired malaria cases in the USA. *Lancet Infect Dis* **23**, e401, doi:10.1016/S1473-3099(23)00581-9 (2023).

Carlson, C. J., Bannon, E., Mendenhall, E., Newfield, T. & Bansal, S. Rapid range shifts in African Anopheles mosquitoes over the last century. *Biol Lett* **19**, 20220365, doi:10.1098/rsbl.2022.0365 (2023).

Suh, P. F. *et al.* Impact of insecticide resistance on malaria vector competence: a literature review. *Malar J* **22**, 19, doi:10.1186/s12936-023-04444-2 (2023).

Takken, W. & Lindsay, S. Increased Threat of Urban Malaria from Anopheles stephensi Mosquitoes, Africa. *Emerg Infect Dis* **25**, 1431-1433, doi:10.3201/eid2507.190301 (2019).

Organization, W. H. Global Vector Control Response 2017–2030. . (Geneva, Switzerland: World Health Organization. , 2017).

Wilson, A. L. *et al.* The importance of vector control for the control and elimination of vector-borne diseases. *PLoS Negl Trop Dis* **14**, e0007831, doi:10.1371/journal.pntd.0007831 (2020).

Bhatt, S. *et al.* The effect of malaria control on Plasmodium falciparum in Africa between 2000 and 2015. *Nature* **526**, 207-211, doi:10.1038/nature15535 (2015).

Protopopoff, N. *et al.* Effectiveness of a long-lasting piperonyl butoxide-treated insecticidal net and indoor residual spray interventions, separately and together, against malaria transmitted by pyrethroid-resistant mosquitoes: a cluster, randomised controlled, two-by-two factorial design trial. *Lancet* **391**, 1577-1588, doi:10.1016/S0140-6736(18)30427-6 (2018).

Barredo, E. & DeGennaro, M. Not Just from Blood: Mosquito Nutrient Acquisition from Nectar Sources. *Trends Parasitol* **36**, 473-484, doi:10.1016/j.pt.2020.02.003 (2020).

Fiorenzano, J. M., Koehler, P. G. & Xue, R. D. Attractive Toxic Sugar Bait (ATSB) For Control of Mosquitoes and Its Impact on Non-Target Organisms: A Review. *Int J Environ Res Public Health* **14**, doi:10.3390/ijerph14040398 (2017).

Lovett, B. *et al.* Transgenic Metarhizium rapidly kills mosquitoes in a malaria-endemic region of Burkina Faso. *Science* **364**, 894-897, doi:10.1126/science.aaw8737 (2019).

Bilgo, E. *et al.* Field assessment of potential sugar feeding stations for disseminating bacteria in a paratransgenic approach to control malaria. *Malar J* **17**, 367, doi:10.1186/s12936-018-2516-x (2018).

Cirimotich, C. M. *et al.* Natural microbe-mediated refractoriness to Plasmodium infection in Anopheles gambiae. *Science* **332**, 855-858, doi:10.1126/science.1201618 (2011).

Sugumaran, M. & Barek, H. Critical Analysis of the Melanogenic Pathway in Insects and Higher Animals. *Int J Mol Sci* **17**, doi:10.3390/ijms17101753 (2016).

Zhang, Y. *et al.* Pea aphid Acyrthosiphon pisum sequesters plant-derived secondary metabolite L-DOPA for wound healing and UVA resistance. *Sci Rep* **6**, 23618, doi:10.1038/srep23618 (2016).

Lanno, S. M. *et al.* Genomics Analysis of L-DOPA Exposure in Drosophila sechellia. *G3 (Bethesda)* **9**, 3973-3980, doi:10.1534/g3.119.400552 (2019).

Wu, K. *et al.* Plant phenolics are detoxified by prophenoloxidase in the insect gut. *Sci Rep* **5**, 16823, doi:10.1038/srep16823 (2015).

Topal, N., Bulduk, I., Mut, Z., Bozoglu, H. & Tosun, Y. K. Flowers, Pollen and Honey for Use in the Treatment of Parkinson's Disease. *Rev Chim* **71**, 308-319, doi:https://doi.org/10.37358/RC.20.9.8341 (2020).

Barda, M. *et al.* Flower morphology and nectar quality traits in faba bean affect attraction to bees. *Journal of Applied Entomology* **148**, 57-73, doi:https://doi.org/10.1111/jen.13208 (2023).

Gish, M., Mescher, M. C. & De Moraes, C. M. Targeted predation of extrafloral nectaries by insects despite localized chemical defences. *Proc Biol Sci* **282**, 20151835, doi:10.1098/rspb.2015.1835 (2015).

Khayhan, K. *et al.* Banana blossom agar (BABA), a new medium to isolate members of the Cryptococcus neoformans/Cryptococcus gattii species complex useful for resource limited countries. *Mycoses* **61**, 959-962, doi:10.1111/myc.12833 (2018).

Soares, A. R. *et al.* The role of L-DOPA in plants. *Plant Signal Behav* **9**, e28275 (2014).

Yang, X., Zhang, X. & Zhou, R. Determination of l-Dopa Content and Other Significant Nitrogenous Compounds in the Seeds of Seven Mucuna and Stizolobium Species in China. *Pharmaceutical Biology* **39**, 312-316, doi:10.1076/phbi.39.4.312.5905 (2001).

Goyoaga, C. *et al.* Content and distribution of vicine, convicine and L-DOPA during germination and seedling growth of two Vicia faba L. varieties. *Eur Food Res Technol* **227**, 1537-1542, doi:10.1007/s00217-008-0876-0 (2008).

Etemadi, F., Hashemi, M., Randhir, R., ZandVakili, O. & Ebadi, A. Accumulation of l-DOPA in various organs of faba bean and influence of drought, nitrogen stress, and processing methods on l-DOPA yield. `*The Crop Journal* **6**, 426-434, doi:https://doi.org/10.1016/j.cj.2017.12.001 (2018).

Zhang, W. *et al.* Regulators and signalling in insect antimicrobial innate immunity: Functional molecules and cellular pathways. *Cell Signal* **83**, 110003, doi:10.1016/j.cellsig.2021.110003 (2021).

Kouamo, M. F. M. *et al.* Genome-Wide Transcriptional Analysis and Functional Validation Linked a Cluster of Epsilon Glutathione S-Transferases with Insecticide Resistance in the Major Malaria Vector Anopheles funestus across Africa. *Genes (Basel)* **12**, doi:10.3390/genes12040561 (2021).

Kwon, H., Mohammed, M., Franzen, O., Ankarklev, J. & Smith, R. C. Single-cell analysis of mosquito hemocytes identifies signatures of immune cell subtypes and cell differentiation. *Elife* **10**, doi:10.7554/eLife.66192 (2021).

Yassine, H., Kamareddine, L. & Osta, M. A. The mosquito melanization response is implicated in defense against the entomopathogenic fungus Beauveria bassiana. *PLoS Pathog* **8**, e1003029, doi:10.1371/journal.ppat.1003029 (2012).

El Moussawi, L., Nakhleh, J., Kamareddine, L. & Osta, M. A. The mosquito melanization response requires hierarchical activation of non-catalytic clip domain serine protease homologs. *PLoS Pathog* **15**, e1008194, doi:10.1371/journal.ppat.1008194 (2019).

Vizioli, J. *et al.* Gambicin: a novel immune responsive antimicrobial peptide from the malaria vector Anopheles gambiae. *Proc Natl Acad Sci U S A* **98**, 12630-12635, doi:10.1073/pnas.221466798 (2001).

Smith, R. C. *et al.* Molecular Profiling of Phagocytic Immune Cells in Anopheles gambiae Reveals Integral Roles for Hemocytes in Mosquito Innate Immunity. *Mol Cell Proteomics* **15**, 3373-3387, doi:10.1074/mcp.M116.060723 (2016).

Wigglesworth, V. B. The insect cuticle. *Biol Rev Camb Philos Soc* **23**, 408-451, doi:10.1111/j.1469-185x.1948.tb00566.x (1948).

Sealy, R. C., Hyde, J. S., Felix, C. C., Menon, I. A. & Prota, G. Eumelanins and pheomelanins: characterization by electron spin resonance spectroscopy. *Science* **217**, 545-547, doi:10.1126/science.6283638 (1982).

Li, M. *et al.* Germline Cas9 expression yields highly efficient genome engineering in a major worldwide disease vector, Aedes aegypti. *Proc Natl Acad Sci U S A* **114**, E10540-E10549, doi:10.1073/pnas.1711538114 (2017).

Camacho, E. *et al.* N-acetylglucosamine affects Cryptococcus neoformans cell-wall composition and melanin architecture. *Microbiology* **163**, 1540-1556, doi:10.1099/mic.0.000552 (2017).

Wood, O., Hanrahan, S., Coetzee, M., Koekemoer, L. & Brooke, B. Cuticle thickening associated with pyrethroid resistance in the major malaria vector Anopheles funestus. *Parasit Vectors* **3**, 67, doi:10.1186/1756-3305-3-67 (2010).

Yahouedo, G. A. *et al.* Contributions of cuticle permeability and enzyme detoxification to pyrethroid resistance in the major malaria vector Anopheles gambiae. *Sci Rep* **7**, 11091, doi:10.1038/s41598-017-11357-z (2017).

Cordero, R. J. B. *et al.* Impact of Yeast Pigmentation on Heat Capture and Latitudinal Distribution. *Curr Biol* **28**, 2657-2664 e2653, doi:10.1016/j.cub.2018.06.034 (2018).

True, J. R. Insect melanism: the molecules matter. *Trends in Ecology and Evolution* **18**, 640-647 (2003).

Zeuss, D., Brandl, R., Brandle, M., Rahbek, C. & Brunzel, S. Global warming favours light-coloured insects in Europe. *Nat Commun* **5**, 3874, doi:10.1038/ncomms4874 (2014).

Haque, M. T., Khan, M. K. & Herberstein, M. E. Current evidence of climate-driven colour change in insects and its impact on sexual signals. *Ecol Evol* **14**, e11623, doi:10.1002/ece3.11623 (2024).

Gonzalez-Tokman, D. *et al.* Insect responses to heat: physiological mechanisms, evolution and ecological implications in a warming world. *Biol Rev Camb Philos Soc* **95**, 802-821, doi:10.1111/brv.12588 (2020).

d'Ischia, M. *et al.* Melanins and melanogenesis: from pigment cells to human health and technological applications. *Pigment Cell Melanoma Res* **28**, 520-544, doi:10.1111/pcmr.12393 (2015).

Barek, H., Sugumaran, M., Ito, S. & Wakamatsu, K. Insect cuticular melanins are distinctly different from those of mammalian epidermal melanins. *Pigment Cell Melanoma Res* **31**, 384-392, doi:10.1111/pcmr.12672 (2018).

Frases, S., Chaskes, S., Dadachova, E. & Casadevall, A. Induction by Klebsiella aerogenes of a melanin-like pigment in Cryptococcus neoformans. *Appl Environ Microbiol* **72**, 1542-1550, doi:10.1128/AEM.72.2.1542-1550.2006 (2006).

Hegde, S. *et al.* Interkingdom interactions shape the fungal microbiome of mosquitoes. *Anim Microbiome* **6**, 11, doi:10.1186/s42523-024-00298-4 (2024).

Tawidian, P., Coon, K. L., Jumpponen, A., Cohnstaedt, L. W. & Michel, K. Host-Environment Interplay Shapes Fungal Diversity in Mosquitoes. *mSphere* **6**, e0064621, doi:10.1128/mSphere.00646-21 (2021).

Chatterjee, S., Prados-Rosales, R., Itin, B., Casadevall, A. & Stark, R. E. Solid-state NMR Reveals the Carbon-based Molecular Architecture of Cryptococcus neoformans Fungal Eumelanins in the Cell Wall. *J Biol Chem* **290**, 13779-13790, doi:10.1074/jbc.M114.618389 (2015).

Ramirez, J. L. *et al.* Reciprocal tripartite interactions between the Aedes aegypti midgut microbiota, innate immune system and dengue virus influences vector competence. *PLoS Negl Trop Dis* **6**, e1561, doi:10.1371/journal.pntd.0001561 (2012).

Wells, J. S. *et al.* SQ 26,180, a novel monobactam. I Taxonomy, fermentation and biological properties. *J Antibiot (Tokyo)* **35**, 184-188, doi:10.7164/antibiotics.35.184 (1982).

Chernin, L. S. *et al.* Chitinolytic activity in Chromobacterium violaceum: substrate analysis and regulation by quorum sensing. *J Bacteriol* **180**, 4435-4441, doi:10.1128/JB.180.17.4435-4441.1998 (1998).

Hillyer, J. F. Insect immunology and hematopoiesis. *Dev Comp Immunol* **58**, 102-118, doi:10.1016/j.dci.2015.12.006 (2016).

Clayton, A. M., Dong, Y. & Dimopoulos, G. The Anopheles innate immune system in the defense against malaria infection. *J Innate Immun* **6**, 169-181, doi:10.1159/000353602 (2014).

Saraiva, R. G., Kang, S., Simoes, M. L., Anglero-Rodriguez, Y. I. & Dimopoulos, G. Mosquito gut antiparasitic and antiviral immunity. *Dev Comp Immunol* **64**, 53-64, doi:10.1016/j.dci.2016.01.015 (2016).

Liao, C., Upadhyay, A., Liang, J., Han, Q. & Li, J. 3,4-Dihydroxyphenylacetaldehyde synthase and cuticle formation in insects. *Dev Comp Immunol* **83**, 44-50, doi:10.1016/j.dci.2017.11.007 (2018).

Vavricka, C. *et al.* From L-dopa to dihydroxyphenylacetaldehyde: a toxic biochemical pathway plays a vital physiological function in insects. *PLoS One* **6**, e16124, doi:10.1371/journal.pone.0016124 (2011).

Goldstein, D. S. The Catecholaldehyde Hypothesis for the Pathogenesis of Catecholaminergic Neurodegeneration: What We Know and What We Do Not Know. *Int J Mol Sci* **22**, doi:10.3390/ijms22115999 (2021).

Renault, D., Dorrah, M. A., Mohamed, A. A., Abdelfattah, E. A. & Bassal, T. T. Assessment of oxidative stress and activities of antioxidant enzymes depicts the negative systemic effect of iron-containing fertilizers and plant phenolic compounds in the desert locust. *Environ Sci Pollut Res Int* **23**, 21989-22000, doi:10.1007/s11356-016-7391-9 (2016).

Calabrese, E. J. & Baldwin, L. A. Agonist consentration gradients as a generalizable regulatory implementation Strategy. *Crit Rev Toxicol* **31**, 471-473, doi:10.1080/20014091111758 (2001).

Calabrese, E. J. & Mattson, M. P. How does hormesis impact biology, toxicology, and medicine? *NPJ Aging Mech Dis* **3**, 13, doi:10.1038/s41514-017-0013-z (2017).

Schirrmacher, V. Less Can Be More: The Hormesis Theory of Stress Adaptation in the Global Biosphere and Its Implications. *Biomedicines* **9**, doi:10.3390/biomedicines9030293 (2021).

Ali, M., Al-Olayan, E. M., Lewis, S., Matthews, H. & Hurd, H. Naturally occurring triggers that induce apoptosis-like programmed cell death in Plasmodium berghei ookinetes. *PLoS One* **5**, doi:10.1371/journal.pone.0012634 (2010).

Foster, W. A. Mosquito sugar feeding and reproductive energetics. *Annu Rev Entomol* **40**, 443-474, doi:10.1146/annurev.en.40.010195.002303 (1995).

Wang, M. *et al.* Glucose-mediated proliferation of a gut commensal bacterium promotes Plasmodium infection by increasing mosquito midgut pH. *Cell Rep* **35**, 108992, doi:10.1016/j.celrep.2021.108992 (2021).

Magistrado, D., El-Dougdoug, N. K. & Short, S. M. Sugar restriction and blood ingestion shape divergent immune defense trajectories in the mosquito Aedes aegypti. *Sci Rep* **13**, 12368, doi:10.1038/s41598-023-39067-9 (2023).

Weger, A. A. & Rittschof, C. C. The diverse roles of insulin signaling in insect behavior. *Front Insect Sci* **4**, 1360320, doi:10.3389/finsc.2024.1360320 (2024).

Ferreira, A. & Milan, M. Dally Proteoglycan Mediates the Autonomous and Nonautonomous Effects on Tissue Growth Caused by Activation of the PI3K and TOR Pathways. *PLoS Biol* **13**, e1002239, doi:10.1371/journal.pbio.1002239 (2015).

Perez-Hedo, M., Rivera-Perez, C. & Noriega, F. G. The insulin/TOR signal transduction pathway is involved in the nutritional regulation of juvenile hormone synthesis in Aedes aegypti. *Insect Biochem Mol Biol* **43**, 495-500, doi:10.1016/j.ibmb.2013.03.008 (2013).

Regan, J. C., Froy, H., Walling, C. A., Moatt, J. P. & Nussey, D. H. Dietary restriction and insulin-like signalling pathways as adaptive plasticity: A synthesis and re-evaluation. *Functional Ecology* **34**, 107-128, doi:https://doi.org/10.1111/1365-2435.13418 (2019).

Huang, C. Y., Chou, S. Y., Bartholomay, L. C., Christensen, B. M. & Chen, C. C. The use of gene silencing to study the role of dopa decarboxylase in mosquito melanization reactions. *Insect Mol Biol* **14**, 237-244, doi:10.1111/j.1365-2583.2004.00552.x (2005).

Takahashi, A. Pigmentation and behavior: potential association through pleiotropic genes in Drosophila. *Genes Genet Syst* **88**, 165-174, doi:10.1266/ggs.88.165 (2013).

Li, J., Zhao, X. & Christensen, B. M. Dopachrome conversion activity in Aedes aegypti: significance during melanotic encapsulation of parasites and cuticular tanning. *Insect Biochem Mol Biol* **24**, 1043-1049, doi:10.1016/0965-1748(94)90142-2 (1994).

McKay, J. K., Richards, J. H. & Mitchell-Olds, T. Genetics of drought adaptation in Arabidopsis thaliana: I. Pleiotropy contributes to genetic correlations among ecological traits. *Mol Ecol* **12**, 1137-1151, doi:10.1046/j.1365-294x.2003.01833.x (2003).

Balabanidou, V. *et al.* Cytochrome P450 associated with insecticide resistance catalyzes cuticular hydrocarbon production in Anopheles gambiae. *Proc Natl Acad Sci U S A* **113**, 9268-9273, doi:10.1073/pnas.1608295113 (2016).

Balabanidou, V. *et al.* Mosquitoes cloak their legs to resist insecticides. *Proc Biol Sci* **286**, 20191091, doi:10.1098/rspb.2019.1091 (2019).

Pedrini, N. *et al.* Control of pyrethroid-resistant Chagas disease vectors with entomopathogenic fungi. *PLoS Negl Trop Dis* **3**, e434, doi:10.1371/journal.pntd.0000434 (2009).

Liu, N. Insecticide resistance in mosquitoes: impact, mechanisms, and research directions. *Annu Rev Entomol* **60**, 537-559, doi:10.1146/annurev-ento-010814-020828 (2015).

Bonizzoni, M. *et al.* RNA-seq analyses of changes in the Anopheles gambiae transcriptome associated with resistance to pyrethroids in Kenya: identification of candidate-resistance genes and candidate-resistance SNPs. *Parasit Vectors* **8**, 474, doi:10.1186/s13071-015-1083-z (2015).

Chrissian, C. *et al.* Melanin deposition in two Cryptococcus species depends on cell-wall composition and flexibility. *J Biol Chem* **295**, 1815-1828, doi:10.1074/jbc.RA119.011949 (2020).

Stuart-Fox, D., Newton, E. & Clusella-Trullas, S. Thermal consequences of colour and near-infrared reflectance. *Philos Trans R Soc Lond B Biol Sci* **372**, doi:10.1098/rstb.2016.0345 (2017).

Freoa, L. *et al.* Drosophilids with darker cuticle have higher body temperature under light. *Sci Rep* **13**, 3513, doi:10.1038/s41598-023-30652-6 (2023).

Sibilia, C. D. *et al.* Thermal Physiology and Developmental Plasticity of Pigmentation in the Harlequin Bug (Hemiptera: Pentatomidae). *J Insect Sci* **18**, doi:10.1093/jisesa/iey066 (2018).

Lopez, V. M. *et al.* Color lightness of velvet ants (Hymenoptera: Mutillidae) follows an environmental gradient. *J Therm Biol* **100**, 103030, doi:10.1016/j.jtherbio.2021.103030 (2021).

Agyekum, T. P. *et al.* A Systematic Review of the Effects of Temperature on Anopheles Mosquito Development and Survival: Implications for Malaria Control in a Future Warmer Climate. *Int J Environ Res Public Health* **18**, doi:10.3390/ijerph18147255 (2021).

Nunes, R. D. *et al.* Polyphenol-Rich Diets Exacerbate AMPK-Mediated Autophagy, Decreasing Proliferation of Mosquito Midgut Microbiota, and Extending Vector Lifespan. *PLoS Negl Trop Dis* **10**, e0005034, doi:10.1371/journal.pntd.0005034 (2016).

Cirstea, M. S. *et al.* A novel pathway of levodopa metabolism by commensal Bifidobacteria. *Sci Rep* **13**, 19155, doi:10.1038/s41598-023-45953-z (2023).

van Kessel, S. P. *et al.* Gut bacterial tyrosine decarboxylases restrict levels of levodopa in the treatment of Parkinson's disease. *Nat Commun* **10**, 310, doi:10.1038/s41467-019-08294-y (2019).

Dong, Y., Manfredini, F. & Dimopoulos, G. Implication of the mosquito midgut microbiota in the defense against malaria parasites. *PLoS Pathog* **5**, e1000423, doi:10.1371/journal.ppat.1000423 (2009).

Saraiva, R. G. & Dimopoulos, G. Bacterial natural products in the fight against mosquito-transmitted tropical diseases. *Nat Prod Rep* **37**, 338-354, doi:10.1039/c9np00042a (2020).

Roshchina, V. V. New Trends and Perspectives in the Evolution of Neurotransmitters in Microbial, Plant, and Animal Cells. *Adv Exp Med Biol* **874**, 25-77, doi:10.1007/978-3-319-20215-0_2 (2016).

Zhao, M., Lin, X. & Guo, X. The Role of Insect Symbiotic Bacteria in Metabolizing Phytochemicals and Agrochemicals. *Insects* **13**, doi:10.3390/insects13070583 (2022).

Shamjana, U., Vasu, D. A., Hembrom, P. S., Nayak, K. & Grace, T. The role of insect gut microbiota in host fitness, detoxification and nutrient supplementation. *Antonie Van Leeuwenhoek* **117**, 71, doi:10.1007/s10482-024-01970-0 (2024).

El-Shehawy, R., Luecke-Johansson, S., Ribbenstedt, A. & Gorokhova, E. Microbiota-Dependent and -Independent Production of l-Dopa in the Gut of Daphnia magna. *mSystems* **6**, e0089221, doi:10.1128/mSystems.00892-21 (2021).

Briggs, A. M. *et al.* Anopheles stephensi Feeding, Flight Behavior, and Infection With Malaria Parasites are Altered by Ingestion of Serotonin. *Front Physiol* **13**, 911097, doi:10.3389/fphys.2022.911097 (2022).

Shim, J., Mukherjee, T. & Banerjee, U. Direct sensing of systemic and nutritional signals by haematopoietic progenitors in Drosophila. *Nat Cell Biol* **14**, 394-400, doi:10.1038/ncb2453 (2012).

Organization, W. H. World Malaria Report 2021. 322 (World Health Organization Geneva, 2021).

Hartley, S. *et al.* Ugandan stakeholder hopes and concerns about gene drive mosquitoes for malaria control: new directions for gene drive risk governance. *Malar J* **20**, 149, doi:10.1186/s12936-021-03682-6 (2021).

Annas, G. J. *et al.* A Code of Ethics for Gene Drive Research. *CRISPR J* **4**, 19-24, doi:10.1089/crispr.2020.0096 (2021).

Dubovskiy, I. M. *et al.* More than a colour change: insect melanism, disease resistance and fecundity. *Proc Biol Sci* **280**, 20130584, doi:10.1098/rspb.2013.0584 (2013).

Mustard, J. A., Beggs, K. T. & Mercer, A. R. Molecular biology of the invertebrate dopamine receptors. *Arch Insect Biochem Physiol* **59**, 103-117, doi:10.1002/arch.20065 (2005).

Rehr, S. S., Janzen, D. H. & Feeny, P. P. L-dopa in legume seeds: a chemical barrier to insect attack. *Science* **181**, 81-82, doi:10.1126/science.181.4094.81 (1973).

Giraldo, D. *et al.* Human scent guides mosquito thermotaxis and host selection under naturalistic conditions. *Curr Biol* **33**, 2367-2382 e2367, doi:10.1016/j.cub.2023.04.050 (2023).

Tripathi, A. K., Mlambo, G., Kanatani, S., Sinnis, P. & Dimopoulos, G. Plasmodium falciparum Gametocyte Culture and Mosquito Infection Through Artificial Membrane Feeding. *J Vis Exp*, doi:10.3791/61426 (2020).

Smith, D. F. Q. *et al.* Glyphosate inhibits melanization and increases susceptibility to infection in insects. *PLoS Biol* **19**, e3001182, doi:10.1371/journal.pbio.3001182 (2021).

Schindelin, J. *et al.* Fiji: an open-source platform for biological-image analysis. *Nat Methods* **9**, 676-682, doi:10.1038/nmeth.2019 (2012).

Fung, B. M., Khitrin, A. K. & Ermolaev, K. An improved broadband decoupling sequence for liquid crystals and solids. *J Magn Reson* **142**, 97-101, doi:10.1006/jmre.1999.1896 (2000).

Lehmann, P. F. & White, L. O. Chitin assay used to demonstrate renal localization and cortisone-enhanced growth of Aspergillus fumigatus mycelium in mice. *Infect Immun* **12**, 987-992, doi:10.1128/iai.12.5.987-992.1975 (1975).

Camacho, E. *et al.* N-acetylglucosamine affects Cryptococcus neoformans cell-wall composition and melanin architecture. *Microbiology (Reading)* **163**, 1540-1556, doi:10.1099/mic.0.000552 (2017).

Organization, W. H. *Test procedures for insecticide resistance monitoring in malaria vector mosquitoes*. 2nd ed. edn,  (2018).

Jacobs, E. *et al.* Cuticular profiling of insecticide resistant Aedes aegypti. *Sci Rep* **13**, 10154, doi:10.1038/s41598-023-36926-3 (2023).

Matsumoto, H., Tanaka, K., Noguchi, H. & Hayakawa, Y. Cause of mortality in insects under severe stress. *Eur J Biochem* **270**, 3469-3476, doi:10.1046/j.1432-1033.2003.03745.x (2003).

**Fig 1. Experimental overview for investigating the impact of L-DOPA-supplemented sugar meal on *A. gambiae* mosquitoes.**

**Fig 2. Melanization pathways in insects.** In *Anopheles* female mosquitoes, melanin synthesis mostly depends on dopamine produced from the blood meal-derived phenylalanine; however, other precursors from plant sources such as tyrosine or L-DOPA are used by both female and male mosquitoes for the generation of this biopolymer. Most melanins synthesized in insects derive from L-DOPA or dopamine in a reaction initiated by phenoloxidases present in the hemolymph; however, there is one possible minor pathway from a degradation product of dopamine: 3,4-dihydroxyphenylaldehide (DOPAL). DOPAL is a highly unstable compound that reacts with primary amines, leading to protein crosslinking and inactivation. Current literature suggests that its toxicity outweighs its benefits, thus evidence of DOPAL melanin formation in any biological system remains to be determined. Simultaneously but via an independent pathway, dopamine serves as substrate for a process known as sclerotization which promotes hardening of insect cuticles (exoskeleton) and prevents desiccation, dehydration, and invasion from foreign organisms. This process is catalyzed by cuticle-bound phenoloxidases [12,17]. Figure created with Biorender.com

**Fig 3. L-DOPA supplementation exerts a hormesis effect. A)** Daily proportion of surviving mosquitoes after L-DOPA feeding in 1.5% sugar meal at various concentrations. Each survival curve represents pooled data from three independent experiments (n= 75). Survival curves for L-DOPA-fed mosquitoes were compared statistically to the survival curves of control mosquitoes using Log-rank Mantel-Cox tests (NS, not significant, *$p$ = 0.0113, **$p$ = 0.0040, ****$p$ < 0.0001). **B)** Cumulative survival percentage at days 4, 8 and 14 shows how mosquitoes adapt to toxic effects of L-DOPA at concentrations below and above 1 mM over time. **C)** Daily proportion of surviving mosquitoes after dopamine feeding in 1.5% sugar meal at various concentrations. Each survival curve represents pooled data from three independent experiments (n= 75). Survival curves for dopamine-fed mosquitoes were compared statistically to the survival curves of control mosquitoes using Log-rank Mantel-Cox tests (NS, not significant, ****$p$ < 0.0001). **D)** Cumulative survival percentage at days 4, 8 and 14 shows highly toxic effects of dopamine over time and striking death on sugar-fed control mosquitoes.

**Fig 4. Comparative transcriptome analysis of *A. gambiae* gene expression between L-DOPA-fed and sugar-fed female mosquitoes.** **A)** Volcano plot of differentially expressed genes (DEGs) in whole (headless) mosquitoes. Significantly upregulated and downregulated genes are highlighted in red and blue spots, respectively. **B)** DEGs by gene ontology category using VectorBase in whole L-DOPA-fed female mosquitoes compared with sugar-fed (control) females. Differential gene expression was considered significant when the expression fold change (L-DOPA-fed versus sugar-fed) was ≥1.3 on a  $-\log_{10}$ scale (P < 0.05). R/S/M, oxidoreductive, stress-related, and mitochondrial **C)** Volcano plot of differentially expressed genes (DEGs) in mosquito midguts. Significantly upregulated and downregulated genes are highlighted in red and blue spots, respectively. **D)** DEGs by gene ontology category in midguts from L-DOPA-fed female mosquitoes compared with sugar-fed females. Differential gene expression was considered significant when the expression fold change (L-DOPA-fed versus sugar-fed) was ≥1.3 on a $-\log_{10}$ scale (P < 0.05). R/T/T, replication, transcription, and translation; R/S/M, oxidoreductive, stress-related, and mitochondrial.

## A — IMMUNE-RELATED

### WHOLE MOSQUITOES

| Gene ID | Log2 Fold Change | Gene name | P-value |
|---|---|---|---|
| AGAP005799 | 4.63 | Dally protein | 4.29E-14 |
| AGAP009091 | -2.37 | 3,4-dihydroxylphenylacetaldehyde (DOPAL) synthase | 4.10E-02 |
| AGAP007627 | -2.37 | Nitric oxide synthase protein (NOS) | 4.05E-02 |
| AGAP000536 | -2.44 | Peptidoglycan recognition protein (short) (PGRP-S1) | 3.47E-02 |
| AGAP012326 | -2.77 | TOLL-like receptor 7 (TOLL7) | 1.98E-02 |
| AGAP004806 | -2.79 | Galectin 6 (GAL6) | 1.47E-02 |
| AGAP012932 | -2.81 | Autophagy-related protein 9 | 3.03E-02 |
| AGAP012616 | -2.88 | prophenoloxidase 5 (PPO5 ) | 2.41E-02 |
| AGAP010833 | -2.88 | CLIP-domain serine protease (CLIPB14) | 3.97E-02 |
| AGAP010485 | -2.92 | Dopamine beta-monooxygenase (DBH) | 1.08E-02 |
| AGAP002451 | -4.32 | Class B Scavenger Receptor (SCRB1) | 3.54E-08 |
| AGAP002815 | -4.35 | CLIP-domain serine protease (CLIPA15) | 1.83E-08 |
| AGAP005959 | -4.40 | Dopachrome-conversion enzyme (DCE) | 6.55E-09 |
| AGAP007692 | -4.43 | Serine Protease Inhibitor (SRPN14) | 2.96E-10 |
| AGAP011952 | -4.47 | Short caspase 3 (CASPS3) | 1.17E-09 |
| AGAP007410 | -4.50 | C-type lectine mannose binding  (CTLMA5) | 4.68E-10 |
| AGAP011693 | -4.54 | Long caspase (CASPL1) | 6.92E-11 |

### MIDGUTS

| Gene ID | Log2 Fold Change | Gene name | P-value |
|---|---|---|---|
| AGAP005583 | 2.31 | Phosphoinositide-3-kinase, regulatory subunit (PPI3K) | 1.97E-02 |
| AGAP010817 | 2.19 | CUB domain-containing protein, predicted peroxidase | 1.12E-02 |
| AGAP004148 | 1.86 | CLIP domain-containing serine protease (CLIPB5) | 2.64E-02 |
| AGAP011360 | 0.42 | Kurtz arrestin-like protein (ARRK) | 3.17E-02 |
| AGAP011792 | -2.15 | CLIP domain-containing serine protease (CLIPA7) | 3.15E-02 |

## B — CUTICLE-RELATED

### WHOLE MOSQUITOES

| Gene ID | Log2 Fold Change | Gene name | P-value |
|---|---|---|---|
| AGAP004270 | 3.25 | Hexosaminidase D | 4.67E-03 |
| AGAP009191 | 3.21 | Glutathione S-transferase epsilon class 6 (GSTE6) | 6.01E-03 |
| AGAP013219 | -2.86 | Elongase, elongation of very long chain fatty acids protein 7 (ELO7) | 3.46E-02 |
| AGAP005371 | -2.88 | Carboxylesterase beta esterase (COEBE2C) | 4.33E-02 |
| AGAP005512 | -2.90 | Elongase, Elongation of very long chain fatty acids protein (ELO4) | 3.48E-02 |
| AGAP011936 | -3.04 | Serpentine (chitin deacetylase) | 1.36E-02 |
| AGAP001473 | -3.05 | Propionyl-CoA synthetase | 4.18E-04 |
| AGAP009195 | -3.12 | Glutathione S-transferase epsilon class 1 (GSTE1) | 6.89E-03 |
| AGAP004784 | -3.15 | Fatty acyl-CoA reductase 2 | 2.16E-03 |
| AGAP001748 | -3.33 | Chitin synthase 1 (CHS1A,B) | 2.80E-04 |
| AGAP000820 | -4.32 | Cuticular protein RR-1 (CPR125) | 3.97E-08 |
| AGAP005511 | -4.36 | Elongation of very long chain fatty acids protein 2 (ELO2) | 1.47E-08 |
| AGAP009241 | -4.41 | Cytochrome P450 (CYP4C36) | 4.65E-09 |
| AGAP007980 | -4.44 | Cuticular protein CPCFC family (CPCFC1) | 2.25E-09 |
| AGAP000986 | -4.47 | Cuticular protein (CPAP3-D) | 1.14E-09 |
| AGAP013120 | -4.57 | Chitinase (CHT5-5) | 2.13E-11 |
| AGAP006049 | -4.64 | Cytochrome P450 (CYP4J10) | 3.48E-12 |

### MIDGUTS

| Gene ID | Log2 Fold Change | Gene name | P-value |
|---|---|---|---|
| AGAP008685 | 2.29 | Juvenile hormone epoxide hydrolase 2 (Jheh2) | 6.75E-05 |
| AGAP005660 | -1.78 | Cytochrome P450 (CYP305A4) | 4.46E-02 |
| AGAP009246 | -2.70 | Cytochrome P450 (CYP4C27) | 3.21E-02 |
| AGAP010362 | -2.77 | Chitin binding domain-containing protein | 2.09E-02 |

**Fig 5. Differential transcripts abundances of melanin-related physiological processes in *A. gambiae* female mosquitoes following feeding of a sugar-meal supplemented with 1 mM L-DOPA for 4 d.** $Log_2$ scale indicates transcripts that were found at significantly higher (red) or lower (blue) levels in  whole (headless) mosquitoes (upper panel) and midguts (lower panel) from *A. gambiae* post L-DOPA-feeding versus sugar-fed (controls). **A)** Set of immune-related transcripts. **B)**  Set of cuticle-related transcripts.

**Fig 6. L-DOPA exhibits antiparasitic and antifungal activity. A)** Intensity of *P. falciparum* oocyst load according to L-DOPA concentration in adult female mosquitoes. Each circle indicates the number of oocysts on an individual midgut; horizontal bars represent the median value of oocysts. n=3, NS, not significant, **$p$ = 0.0081, One-way analysis of variance (ANOVA) nonparametric Kluskal-Wallis test. Statistical significance for Prevalence of infection was assessed with a Fisher's exact test, ns. **B)** Intensity of *P. falciparum* infection according to dopamine concentration in adult female mosquitoes. n=3, NS, not significant, **$p$ = 0.0019, One-way ANOVA nonparametric Kluskal-Wallis test. Statistical significance for Prevalence of infection was assessed with a Fisher's exact test, ns. **C)** *In vitro* assessment of L-DOPA toxicity to sexual and asexual stages of *P. falciparum* parasite. Use of L-DOPA concentrations ranging from 0.25 mM to 2 mM during in vitro parasite culture, appeared more toxic to asexual stages of the parasite than to sexual stages. n=3, NS, not significant, ****$p<$ 0.0001, Two-way ANOVA with Dunnett's multiple comparisons tests against individual controls.   **D)** Fungal load in L-DOPA-fed female mosquitoes shows a reduction comparison to control. Each circle corresponds to a whole mosquito homogenate; horizontal red bars represent the median value of fungal CFUs. n=2, NS, not significant, Unpaired nonparametric Mann-Whitney test. **E)** Cryptococcal yeasts are successfully melanized (red arrow) in the abdomens of *A. gambiae* female mosquitoes 24 h post injection of *C. neoformans* H99-GFP strain (1 x 10^7 yeasts/ml). Live yeast cells (green arrows), Dead yeast cells (red arrows). **F)** qRT-PCR analysis of mRNA abundance in whole mosquitoes from key genes [dopa decarboxylase (DDC), gambicin (GAM), phenoloxidase 1 ( PPO1), and leucine-rich immunomodulator 15 (LRIM15)] involved in the mosquito melanization pathway and *Plasmodium* parasite development. Data represent the mean ± SE of three or more independent replicates, analyzed using unpaired by one-way ANOVA and multiple comparison test to determine gene expression relative to housekeeping S7 gene. NS, not significant, **$p$ = 0.0041, *$p$ = 0.0235.

**Fig 7. Oral L-DOPA boosts mosquito cuticular melanization via a non-canonical pathway. A)** Dietary L-DOPA enhances cuticle pigmentation of *A. gambiae* female mosquitoes in a dose-dependent manner. *Upper panel*, Representative images of *A. gambiae* female mosquitoes fed with a sugar meal supplemented with different concentrations of L-DOPA for up to 14 d. Scale bar, 1 mm; *Lower panel*, Quantification of cuticle darkening at 8 and 14 d. Each dot corresponds to an individual mosquito. Data representative from 2-3 biological replicates. Statistical significance was determined by an Unpaired two-tailed t test with two or more data sets (Left panel, *$p$ = 0.0342; Right panel, ns = not significant, *$p$ = 0.0189, ****$p$ < 0.0001) **B)** *Top panel,* TEM micrographs exhibiting distinct electron-dense bands within the endocuticle of L-DOPA-fed mosquitoes not seen in the control. Scale bar, 500 nm. *Lower panel,* Measurements of cuticular thickness from legs of female mosquito fed with L-DOPA supplemented sugar meals and control show increased exo and endocuticle.****$p$<0.0001; Two-way analysis of variance (ANOVA) with Tukey's multiple comparisons test. **C)** L-DOPA supplementation augments content of indole groups. *Left panel*, DPMAS $^{13}$C ssNMR spectra of $^{13}$C-L DOPA fed mosquitoes display pigment peaks (green arrows) with greater relative intensities in comparison to their respective controls. Spectra were normalized to tallest peak (lipid peak at ~30 ppm); Right panel, CPMAS $^{13}$C ssNMR spectra of $^{13}$C-L-DOPA fed mosquitoes corroborate the assertion that $^{13}$C-L-DOPA has been nearly-exclusively used to make melanin instead of being broken down and used as a nutrient source. Spectra were normalized to tallest peak (polysaccharide peak at ~75 ppm). Data are representative from 2 biological replicates. **D)** Quantification of chitin content in whole *A. gambiae* female mosquitoes exhibits demonstrate no significant differences between sugar-fed (control) and L-DOPA-fed mosquitoes. Data correspond to 3 biological replicates. Floating bars (min to max) Line at mean. NS, not significant; Unpaired, nonparametric Mann-Whitney test. **E)** EPR spectroscopy analysis of acid-resistant material isolated from *An. gambiae* and *Ae. aegypti* yellow mutant strain L-DOPA-fed (red spectra) and sugar-fed (blue spectra). Fungal eumelanin was used as positive control (black spectrum).

**Fig. 8. Dietary L-DOPA darkening of cuticle increases heat-exposed mosquito temperatures following light exposure and decreases lifespan. A)** *A. gambiae* female mosquitoes fed with a sugar meal supplemented with 1 mM L-DOPA for 8 d recorded ability to absorb augmented ability to capture heat eat. **B)** Exposure to a direct light source during 1 min (shadowed area) demonstrated that 1mM L-DOPA- fed mosquitoes (red squares) exhibited a warmer body temperature than sugar-fed control (blue circles). Post-irradiation when light is turned of (unshadowed area), L-DOPA-fed mosquitoes (red squares) were distinctly hotter than sugar-fed controls (blue circles), starting to cool down but kept warmer than control The thermograph was obtained using the Teledyne FLIR E95 camera, and the temperature of each mosquito was extracted using the FLIR software, which provided the maximum, minimum, and average temperatures within a box selection for each mosquito. The maximum temperature for each mosquito was used to generate the graph, which represents the average (+/- Standard Deviation) of the six maximum temperatures corresponding to the two experimental groups (control sugar-fed and 1 mM L-DOPA-fed). **C)** Lifespan of mosquitoes maintained on 1.5% sugar with 1 mM L-DOPA was diminished in comparison to sugar-only fed mosquitoes. Data representative from 3 biological replicates. **D)** Number of eggs laid by female mosquitoes after feeding with a sugar meal supplemented with 1 mM L-DOPA during 4 d before a blood meal on mice. Mann-Whitney test, NS, not significant, $p = 0.1987$. Red lines denote mean. **E)** Hatching rate at 27 °C indicating average percentage of eggs giving rise to 1st and 2nd instar larvae. Mann-Whitney test, NS, not significant. $p = 0.9390$. Red lines denote mean.

**Fig. 9 Dietary L-DOPA metabolism and probing for dopaminergic bacteria amongst gut microbiota from field-caught mosquitoes.**
**A)** Quantification of dopamine by HPLC from whole *Anopheles* mosquitoes post 4 d of dietary L-DOPA suggest a tight regulation of L-DOPA conversion into dopamine. Data corresponds to 2 independent biological replicates from 5-pooled mosquitoes per condition. A 2-way analysis of variance (ANOVA) with Tukey's multiple comparisons test showed no significant differences among and between groups. **B)** Plate confrontational assays showed induced fungal melanization due to melanin-precursors released by bacterial isolates. A) *Comamonas* sp.; B) *Enterococcus hormaechei*; C) *Aeromonas hydrophila*; D) *Serratia marcescens* E) *Chromobacterium subtsugae ΔvioS* F) *Enterococcus* sp. *Cryptococcus* strains (top: *C. gattii* WM179; middle: *C. neoformans* ΔLAC1; bottom: *C. neoformans* H99) were streaked perpendicular to bacterial isolates in SM plates. The plates were incubated at 30°C for 5-7 days. Images are representative of 3 biological replicates.

**Table 1.** Detection of melanin precursors released by diverse bacterial species

| Phylum | Class | Family | Strain | Source | Host | Location | C. neoformans induced melanization | | Refs |
|---|---|---|---|---|---|---|---|---|---|
| | | | | | | | RT | 30°C | |
| Firmicutes | *Bacilli* | *Bacillaceae* | *Bacillus* sp. | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Negative | Negative | 15 |
| | | | *Staphylococcus* sp. A6 | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Negative | Negative | 15 |
| Proteobacteria | Betaproteobacteria | *Commonadaceae* | *Comamonas* sp. | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Positive | Positive | 15 |
| | | *Neisseriaceae* | *Chromobacerium subtsugae* ΔvioS | Soil/Water | None | New Jersey, USA | Positive | Positive | 46,47 |
| | Gammaproteobacteria | *Moraxellaceae* | *Acitenobacter* sp. PA6 | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Negative | Negative | 15 |
| | | *Aeromonadaceae* | *Aeromonas hydrophila* S1 | Midgut | Wild-caught *Ae aegypti* | Panama, Central America | Positive | Positive | 45 |
| | | *Enterobacteriaceae* | *Enterobacter* sp. | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Negative | Negative | 15 |
| | | | *Enterobacter hormaechei* PA | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Negative | Negative | 15 |
| | | | *Serratia marcescens* N1.14 | Midgut | Wild-caught *Ae aegypti* | Panama, Central America | Positive | Positive | 45 |
| | | | *Pantoea* sp. | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Negative | Negative | 15 |
| | | | *Proteus* sp. PAI3 | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Negative | Negative | 15 |
| | | *Pseudomonadaceae* | *Pseudomonas* sp. | Midgut | Wild-caught *An. arabiensis* | Zambia, Africa | Negative | Negative | 15 |

RT, room temperature

**Fig. 10 Alternative pathway for L-DOPA incorporation to the melanin synthesis in mosquitoes. A)** L-DOPA serves as a substrate for cuticular melanization via DOPAL synthase as a detoxification mechanism given a cellular redox environment that promotes L-DOPA antioxidant properties (neutral or acid pH) [60], therefore preventing its conversion to dopamine via a dopa decarboxylase. L-DOPA conversion into DOPAL leads into DOPAL buildup that promotes oligomerization and quinone adduct formation ("quinonization") inhibiting its own metabolism and the activity of other enzymes involved in the melanin biosynthesis such as dopa decarboxylase (DDC), phenoloxidase 5 (PPO5), and dopachrome conversion enzyme (DCE)[59] **B)** Model for dietary L-DOPA-mediated cuticular melanization proposes that *Anopheles gambiae* female mosquitoes use L-DOPA supplement in the sugar meal as a substrate for synthesizing cuticular melanin. L-DOPA is internalized into the midgut epithelium and exported to the hemolymph by sharing the same transporters with amino acids such as Na$^+$-driven amino acid transporter, alpha2 transporter, and/or SLC protein. Within regions of flexible cuticle, in the presence of DOPAL synthase, L-DOPA is turned into DOPAL and H$_2$O$_2$ promoting oxidation of DOPAL catechol ring that results in the formation of a melanized and protective flexible cuticle via crosslinking to cuticular proteins [57,58]. The contribution of gut microbiota as well as from hemocytes as key regulators of mosquito vectorial capacity and immunity, post a L-DOPA diet is under investigation. Figure created with Biorender.com