Sexual Failure Decreases Sweet Taste Perception in Male *Drosophila* via Dopaminergic Signaling

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Abstract

Sweet taste perception, a critical aspect of the initiation of feeding behavior, is primarily regulated by an animal's internal metabolic state. However, non-metabolic factors, such as motivational and emotional states, can also influence peripheral sensory processing and hence feeding behavior. While mating experience is known to induce motivational and emotional changes, its broader impact on other innate behaviors such as feeding remains largely uncharacterized. In this study, we demonstrated that mating failure of male fruit flies suppressed sweet taste perception via dopamine signaling in specific neural circuitry. Upon repetitive failure in courtship, male flies exhibited a sustained yet reversible decline of sweet taste perception, as measured by the proboscis extension reflex (PER) towards sweet tastants as well as the neuronal activity of sweet-sensing Gr5a$^+$ neurons in the proboscis. Mechanistically, we identified a small group of dopaminergic neurons projecting to the subesophageal zone (SEZ) and innervating with Gr5a$^+$ neurons as the key modulator. Repetitive sexual failure decreased the activity of these dopaminergic neurons and in turn suppressed Gr5a$^+$ neurons via Dop1R1 and Dop2R receptors. Our findings revealed a critical role for dopaminergic signaling in integrating reproductive experience with appetitive sensory processing, providing new insights into the complex interactions between different innate behaviors and the role of brain's reward systems in regulating internal motivational and emotional states.

Introduction

As an innate behavior, feeding is essential for the survival and reproduction of all animal species, regulated by a dynamic interplay of internal and external factors such as nutritional status, circadian rhythms, food quality, and emotional states (Atasoy et al., 2012; Hergarden et al., 2012; Lim et al., 2012; Morton et al., 2006; Park et al., 2016; Shiu et al., 2022; Xu et al., 2008; Yapici et al., 2016; Zhan et al., 2016). Beyond nutritive value, palatable food also serves as a powerful reward signal that regulates feeding behavior by influencing various aspects of learning, memory, and motivation (Dayan & Balleine, 2002; Wise, 2004). The function of dopamine system, a key regulator in reward processing, is evolutionarily conserved across species. Studies in both fruit flies and rodents demonstrate that the consumption of sugar or other highly appealing foods triggers a robust dopamine response. The neural substrates of these responses, such as the mushroom bodies in fruit flies and the ventral tegmental area in rodents, underscore the universality of this reward circuit (Burke et al., 2012; Heymann et al., 2020; Liu et al., 2012; Takahashi et al., 2016; Yokel & Wise, 1975).

Mating behavior, as another critical driver of the survival and reproduction of animal species, is also influenced by external stimuli and internal states (Clowney et al., 2015; Kallman et al., 2015; Kohatsu et al., 2011). Failure in mating behavior leads to various behavioral adaptations and even abnormalities. For example, male fruit flies exhibit courtship conditioning, where unsuccessful mating attempts reduce

subsequent courtship toward inappropriate mates, demonstrating behavioral plasticity

aimed at enhancing future reproductive success (Keleman et al., 2012). Meanwhile,

the effect of mating failure extends beyond reproductive behavior. Research has

shown that mating failure increases alcohol consumption in fruit flies (Shohat-Ophir

et al., 2012), a behavior that mirrors reward-seeking tendencies such as substance

abuse seen in higher organisms under stress. Additionally, mating itself has a

rewarding nature. In male mice, after interaction with females, chemical signals are

detected by the accessory olfactory bulb and transmitted to neural circuits that

regulate mating behavior (Decoster et al., 2024). These signals are then projected to

the ventral tegmental area, where serotonin neurons play a crucial role in encoding

both reward acquisition and anticipation (Hashikawa et al., 2016).

Given the established impact of mating experience on behavioral plasticity, we aimed

to investigate whether it also affected reward perception and processing in fruit flies.

Specifically, we examined how courtship failure influenced sweet sensitivity and

feeding behavior, both of which were tightly linked to reward. In our study, we found

that sexual failure led to decreased sweet sensitivity and feeding behavior in male

fruit flies, highlighting a close link between two important innate behaviors.

Furthermore, we identified a small group of dopaminergic neurons that projected to

the SEZ of the fly brain, modulating the activity of sweet-sensing gustatory neurons in

a manner dependent on sexual experience. These dopaminergic neurons were

functionally epistatic to sweet-sensing neurons and promoted behavioral responses to

appetitive cues, but their activity decreased following courtship failure. As a result,

the activity of Gr5a$^+$ neurons in the proboscis was reduced via signaling through

Dop1R1 and Dop2R receptors, leading to reduced sweet sensitivity. These findings

suggest that courtship failure inhibits the dopamine-mediated reward system in the

brain, directly influencing sweet perception and feeding behavior. This provides key

insights into how reproductive stressors, such as mating failure, can affect broader

behavioral output, expanding our understanding of the complex interplay between

instinctive drives.

Results

Courtship failure reduced sweet sensitivity in male flies.

To investigate whether mating experience influenced feeding behavior in *Drosophila melanogaster*, we employed a modified courtship conditioning paradigm (McBride et al., 1999; Shohat-Ophir et al., 2012) (Figure 1A) and examined three groups of males with different mating experiences *in prior*. In *Drosophila*, mated females typically refuse further copulation, leading males exposed to mated females to experience repeated sexual rejection. In contrast, males paired with multiple virgin females could engage in multiple rounds of successful copulations. In our experiments, virgin males were isolated from females for 4 days, after which they were divided into three groups on the 5th day: the Naïve group (continued isolation for 4.5 hours), the Failed group (each male exposed to 25 mated females for 4.5 hours), and the Satisfied group (each male exposed to 25 virgin females for 4.5 hours). Following the treatment, we measured sucrose intake using the MAFE (MAual FEeding) assay we developed previously (Qi et al., 2015). Despite similar consumption volumes across groups (Figure 1-figure supplement 1A), the proportion of males ever consuming any sucrose was significantly lower in the Failed group (Figure 1B), suggesting a reduced likelihood of meal initiation among these males.

Since taste perception is closely linked to the initiation of feeding behavior, we next examined whether sexual experience affected sweet sensitivity using the PER assay.

This well-established method quantified the tendency to initiate feeding behavior

towards different gustatory stimuli (Scott, 2005, 2018). While three groups of male

flies all exhibited dose-dependent responses to varying sucrose concentrations, males

in the Failed group demonstrated a rightward shift of the concentration-response

curve, indicating significantly reduced sweet sensitivity (Figure 1C, left). Consistently,

the Mean Acceptance Threshold (MAT), the sucrose concentration at which 50% of

the population displayed a PER response, was significantly elevated in the Failed

group (Figure 1C, right). Furthermore, we found that the diminished sweet sensitivity

extended to other sweet compounds such as fructose and glucose (Figure 1-figure

supplement 1B-C), indicating a general reduction in sweet sensitivity and an overall

elevation in the threshold to initiate feeding upon appetitive cues.

To determine the time course of this effect, we monitored PER responses in male flies

at 24, 48, and 72 hours post-treatment. Sweet sensitivity remained suppressed for up

to 48 hours in Failed males but returned to baseline levels by 72 hours (Figure 1D). To

explore the mechanisms underlying this lasting change, we examined whether it could

be attributed to the exposure to cis-vaccenyl acetate (cVA), a pheromone present on

the cuticle of mated females but absent from virgin females (Ejima et al., 2007).

Conditioning males with either mated females or decapitated virgin females produced

similar reductions in sweet sensitivity (Figure 1E), suggesting that cVA is not

essential for this behavioral effect. We then tested whether the reduced sweet

sensitivity was a result of failed copulation. Failed males were then divided into two

groups for additional treatments: one group remained untreated, while the other was

allowed to copulate with virgin females. We found that the reduction in sweet

sensitivity caused by sexual failure could be reversed through subsequent successful

copulation. Such males showed significantly increased sugar sensitivity compared to

those that did not have a chance to copulate at all (Figure 1F). Overall, our findings

reveal that sexual failure exerted a long-lasting yet reversible effect to reduce sweet

sensitivity in male flies.

Dopamine signaling mediated the suppression of sweet sensitivity

following sexual failure.

The long-lasting and reversible effect of sexual failure hinted certain changes in flies'

internal motivational state. Dopamine was reported to regulate PER behavior in a

metabolic state-dependent manner (Inagaki et al., 2012; Marella et al., 2012). We thus

hypothesized that dopamine might also play a key role in the taste plasticity triggered

by sexual failure. To test this, we first used a pharmacological approach to reduce

dopamine signaling in male flies by feeding them 3-iodotyrosine (3IY), a tyrosine

hydroxylase-specific inhibitor that blocked dopamine synthesis (Inagaki et al., 2012)

and then measured their PER responses. Flies treated with 3IY showed comparable

sweet responses across all three experimental groups, with an overall reduction in

sweet sensitivity compared to the control group, effectively abolishing the changes

induced by sexual failure (Figure 2A). To further confirm the necessity of dopamine

in modulating sexual experience-dependent sweet sensitivity, we employed a genetic

approach by using $Th^{-/-}$ mutant flies, in which tyrosine hydroxylase, the rate-limiting

enzyme in dopamine synthesis, was eliminated (Deng et al., 2019; Friggi-Grelin et al.,

2003; Marella et al., 2012). As expected, $Th^{-/-}$ mutant flies in the Failed group

exhibited sweet sensitivity indistinguishable from those in the Naïve and Satisfied

groups (Figure 2 -figure supplement 1A).

Next, we investigated whether elevating dopamine levels could restore diminished

sweet sensitivity in males from the Failed group. We first tested the effect of

exogenous dopamine on sweet sensitivity by injecting dopamine into the base of the

flies' proboscis. Consistent with previous findings (Inagaki et al., 2012; Marella et al.,

2012), increasing dopamine levels enhanced sweet sensitivity (Figure 2 -figure

supplement 1B). We then measured PER behavior in Naïve, Failed, and Satisfied flies

injected with either dopamine or ddH$_2$O as control. While ddH$_2$O-treated males from

the Failed group displayed reduced sweet sensitivity compared to Naïve and Satisfied

males, dopamine-treated Failed flies did not show such reduction (Figure 2B), further

suggesting that dopamine signaling is necessary for sexual failure to suppress sweet

sensitivity.

Besides dopamine, serotonin, another important biogenic amine, has been reported to

modulate motivational states and influence feeding behavior (Owald & Waddell, 2015;

Ries et al., 2017; Srinivasan et al., 2008). Therefore, we examined whether serotonin

also played a role in the sexual experience-dependent changes in sweet sensitivity.

Feeding male flies with DL-p-chlorophenylalanine (pCPA) or methysergide, both

blocking serotonin signaling (Engelman et al., 1967; McCorvy et al., 2018) , did not

affect the reduction in sweet sensitivity upon sexual failure (Figure 2 -figure

supplement 1C-D).

To further validate the involvement of dopamine signaling in sexual failure-induced

behavioral changes, we used *TH-Gal4* to drive *UAS-dTRPA1* or *UAS-Shibire^{ts}* (*Shi^{ts}*)

to transiently activate or inhibit dopaminergic neurons, respectively (Deng et al.,

2019). Activation of dopaminergic neurons through the temperature-sensitive cation

channel dTRPA1 (Pulver et al., 2009) increased sugar sensitivity in all male flies and

eliminated differences among groups with different sexual experiences. Meanwhile,

control genotypes expressing only *TH-Gal4* or *UAS-dTRPA1* still showed

significantly reduced sugar sensitivity in the Failed group (Figure 2C). Additionally,

males from the Failed group showed decreased sweet sensitivity across all genotypes

at the permissive temperature (20 °C) (Figure 2 -figure supplement 1E). In contrast,

blocking dopaminergic neurons via the temperature-sensitive dominant-negative

Dynamin *Shibire^{ts}* (*Shi^{ts}*) (Kitamoto, 2001) decreased sugar sensitivity across all three

groups and also completely suppressed the differences among groups (Figure 2D). In

comparison, in the two control genotypes and under permissive temperature (20 °C),

sexual failure could still suppress sweet sensitivity (Figure 2D, Figure 2 -figure

supplement 1F). These findings demonstrate that reduced sweet sensitivity in sexually

failed males is mediated by dopaminergic neurons.

Dopamine signaling directly modulated the activity of Gr5a$^+$ gustatory

neurons upon sexual failure.

We next asked how dopamine signaling modulated taste sensitivity. Gustatory

receptor neurons (GRNs) respond to specific taste stimuli, with Gr5a$^+$ neurons, which

express sweet taste receptors, playing a key role in sweet perception (Dahanukar et al.,

2007; Thorne et al., 2004; Wang et al., 2004). We hypothesized that behavioral

changes following sexual failure might be linked to altered activities of sweet-sensing

GRNs. To test this, we expressed calcium indicator GCaMP6m (Chen et al., 2013) in

Gr5a$^+$ neurons and conducted *in vivo* calcium imaging. Gr5a$^+$ neurons project to SEZ

in the fly brain, where we recorded calcium signals in response to sucrose stimulation.

Sexual failure significantly reduced the responsiveness of Gr5a$^+$ neurons to sucrose,

consistent with our behavioral observations (Figure 3A-D).

Given that dopamine signaling has been shown to modulate sweet sensitivity (Inagaki

et al., 2012; Marella et al., 2012), it was plausible that dopamine signaling directly

modulated the activity of Gr5a$^+$ neurons upon sexual failure. We thus examined

projection patterns of dopaminergic neurons relative to those of Gr5a$^+$ neurons. Using

TH-Gal4 and *Gr5a-LexA*, we expressed *UAS-mCD8GFP* and *LexAop-rCD2RFP* in

the same fly brains, respectively. The brain imaging data revealed that dopaminergic

projections were closely aligned with Gr5a$^+$ GRNs in the SEZ (Figure 3-figure

supplement 1A), suggesting possible synaptic connections of Gr5a$^+$ neurons and

dopaminergic neurons.

We confirmed this idea using enhanced trans-synaptic reconstruction (nSyb-GRASP)

(Feinberg et al., 2008; Gordon & Scott, 2009; Macpherson et al., 2015), which could

visualize direct and active synaptic innervations between two groups of neurons. The

nSyb-GRASP system relies on the expression of two split-GFP fragments on

membranes of two neuronal populations, which fluoresces only upon the reassembly

of these two fragments at active synapses. By employing the nSyb-GRASP system in

Gr5a$^+$ neurons and dopaminergic neurons, we observed GFP signals in the SEZ

(Figure 3-figure supplement 1B). Furthermore, in males that experienced mating

failure, synaptic connections between dopaminergic neurons and Gr5a$^+$ GRNs were

significantly weakened (Figure 3E-F). These results suggest that dopaminergic

neurons directly innervate with Gr5a$^+$ neurons and modulate their activity upon sexual

failure.

This idea was further assessed by other complementary approaches. We expressed the

photosensitive protein CsChrimson (Klapoetke et al., 2014) in dopaminergic neurons

and GCaMP6m in Gr5a$^+$ neurons. Light activation of dopaminergic neurons

significantly enhanced the calcium responses of Gr5a$^+$ neurons to sugar stimuli

(Figure 3-figure supplement 1C-D). Furthermore, by using a CaLexA reporter system

(Masuyama et al., 2012), we assessed the impact of sexual experience on neural

activity in male fruit flies. The results showed that sexual failure led to a significant

reduction in CaLexA-mediated GFP signals in dopaminergic neurons in the SEZ but

not in the antennal lobes (AL) (Figure 3G-I). In summary, these findings indicate that

mating experience modulates sweet taste perception in flies by inhibiting the activity

of specific dopamine neurons in the SEZ and reducing synaptic transmission between

dopaminergic neurons and sweet-sensing Gr5a$^+$ GRNs.

Two dopamine receptors, Dop1R1 and Dop2R, mediated the effect of

sexual failure.

We next explored the molecular mechanisms involved in the reduction of sweet

sensitivity upon sexual failure. Four dopamine receptors (Dop1R1, Dop1R2, Dop2R,

and DopEcR) have been identified in *Drosophila* (Deng et al., 2019; Feng et al., 1996;

Hearn et al., 2002; Srivastava et al., 2005; Sugamori et al., 1995). Through screening

of these receptors, we found that mutations in *Dop1R1* or *Dop2R* abolished the effect

of sexual failure on sweet sensitivity, while mutations in *Dop1R2* or *DopEcR* did not

prevent the decline in sweet sensitivity following sexual failure (Figure 4A-F). To

further confirm the roles of Dop1R1 and Dop2R, we used RNA interference (RNAi)

to knock down these receptors in flies' nervous system. In line with the above results,

knockdown of Dop1R1 or Dop2R blocked the influence of sexual experience on

sweet sensitivity, while knockdown of Dop1R2 or DopEcR had no significant effect

(Figure 4-figure supplement 1A-D). These results suggest that reductions in sweet

sensitivity induced by sexual failure are mediated by dopamine signaling acting on

Dop1R1 and Dop2R receptors.

Next, we activated the neurons expressing Dop1R1 or Dop2R by expressing the

bacterial sodium channel NaChBac (Nitabach et al., 2006). We found that consecutive

activating of $Dop1R1^+$ or $Dop2R^+$ neurons could prevent the decrease in sweet

sensitivity upon mating failure (Figure 5A-B). Meanwhile, activating $Dop1R2^+$ or

$DopEcR^+$ neurons did not reverse this effect (Figure 5-figure supplement 1A-C). We

also tried to silence $Dop1R1^+$ or $Dop2R^+$ neurons using the hyperpolarizing potassium

channel (Kir2.1) (Baines et al., 2001). When these neurons were silenced, male flies

that experienced mating failure no longer exhibited reduced sweet sensitivity

compared to Naive or Satisfied males (Figure 5C-D). These findings indicate that

sexual failure suppresses sweet sensitivity by regulating the activity of Dop1R1 and

Dop2R neurons.

Gr5a$^+$ neurons received dopaminergic modulation through Dop1R1 and

Dop2R.

The above discoveries raised a possibility that sweet-sensing Gr5a$^+$ neurons might

receive direct dopaminergic modulation via Dop1R1 and Dop2R receptors, and that

such mechanism might be critical for the modulation of sweet sensitivity upon sexual

failure. To investigate this possibility, we first examined the expression of Dop1R1

and Dop2R in Gr5a$^+$ neurons. Immunostaining confirmed that both receptors were

expressed in the labellum (Figure 6-figure supplement 1A). Moreover, co-labeling

experiments revealed a significant overlap of Dop1R1/Dop2R expression in Gr5a$^+$

neurons (Figure 6A-B).

Next, we assessed whether the expression of Dop1R1 and Dop2R in Gr5a$^+$ neurons

was functionally important for the reduction of sweet sensitivity upon sexual failure.

By using RNAi to reduce Dop1R1 or Dop2R expression in Gr5a$^+$ neurons, we found

that males experiencing sexual failure no longer exhibited a reduction in sweet

sensitivity compared to Naïve or successfully mated males (Figures 6C-D). In contrast,

reducing the expression of another dopamine receptor, DopEcR, which was also

partially expressed in Gr5a$^+$ neurons (Figure 6-figure supplement 1B), did not prevent

the reduction of sweet sensitivity upon sexual failure (Figure 6-figure supplement 1C).

These data suggest that Dop1R1 and Dop2R in Gr5a$^+$ neurons are indeed essential for

the observed behavioral changes.

We then investigated whether Dop1R1 and Dop2R were required for the modulation

of neuronal activity of Gr5a$^+$ neurons. Following sexual failure, the calcium responses

of Gr5a$^+$ neurons to sugar stimuli were suppressed (Figure 3C). But this suppression

was reversed when Dop1R1 or Dop2R expression was knocked down in Gr5a$^+$

neurons (Figures 6E-F), consistent with the above behavioral observations. In

summary, our results indicate that Dop1R1 and Dop2R in sweet-sensing Gr5a$^+$

neurons mediate the sexual experience-dependent regulation of sweet sensation by

modulating the activity of these neurons.

Discussion

Various innate behaviors are crucial for the survival and reproduction of animals, such as feeding, mating, aggression, sleep, and running away from predators (Amir et al., 2015; Decoster et al., 2024; Hall, 1994; Hashikawa et al., 2016; Hergarden et al., 2012; Nair et al., 2023; Weber & Dan, 2016; Yapici et al., 2016). However, when animals are exposed to competing and even conflicting needs of these innate behaviors, they have to adjust their behavioral output to balance these needs in a timely and precise manner. In the present study, we explored the potential link between two innate behaviors, mating and feeding. We found that failed experiences in sexual interactions suppressed sweet taste perception and hence feeding behavior in male fruit flies (Figure 7). The evolutionary significance of this behavioral plasticity remains unclear. It may represent an adaptive response where reproductive failure shifts time allocation away from feeding to conserve resources or seek alternative mating opportunities. Alternatively, it may indicate a maladaptive state of disrupted reward processing. Previous research has demonstrated that mating failure diminishes animals' motivational state, leading to widespread behavioral changes (Beckwith et al., 2017; Jung et al., 2020; Ryvkin et al., 2024; Shen et al., 2023; Shohat-Ophir et al., 2012). Our study extended this knowledge by showing that mating failure could also affect reward-related feeding behavior. This suggests that different instinctive behaviors might share a common reward mechanism, which exerts broader effects to influence animals' behavioral choices and physiological states.

Feeding provides energy critical for animals' survival and it is not surprising that

feeding is under tight and complex regulations (Atasoy et al., 2012; Morton et al.,

2006; Xu et al., 2008; Yapici et al., 2016; Zhan et al., 2016). While a high volume of

studies focus on mechanisms of how metabolic states modulate feeding behavior

(Ganguly et al., 2021; Inagaki et al., 2012; Marella et al., 2012; Pool & Scott, 2014;

Shiu et al., 2022), our study suggests the motivational drive of feeding, especially that

influenced by mating experience. Previous investigations have shown that males

failed in sexual behaviors increase their ethanol preference (Shohat-Ophir et al., 2012),

have an accelerated aging process (Gendron et al., 2014), and exhibit a sexual

experience-induced preference behavior (Shen et al., 2023). Here we demonstrated

that sexual failure suppressed feeding behavior, one of the most basic homeostatic

systems critical for animal fitness. These results collectively suggest that there may be

a common neural mechanism that underlies the crosslink between mating failure and

other innate behaviors.

Notably, we found that this behavioral coordination persisted for an extended duration

and gradually faded over time, yet could be quickly reversed by successful mating.

This suggests that the regulatory mechanism may modulate the internal state of the

animal. A likely mechanism for this is through the reward system, as both successful

mating and feeding are known to activate the reward system (Burke et al., 2012; Dai

et al., 2022; Schultz, 2010; Stuber & Wise, 2016; Zer-Krispil et al., 2018). To support

this idea, our study demonstrated that dopamine, a key neurotransmitter associated

with reward (Peters et al., 2021; Wise, 2004), plays an integral role in mediating this

behavioral regulation. Dopamine plays a central role in regulating internal

motivational states. It has been shown that factors like social interactions, mating,

hunger, and vigilance influence behavioral output through dopamine signaling (Dai et

al., 2022; de Jong et al., 2019; Inagaki et al., 2012; Marella et al., 2012; Schultz, 2010;

Watabe-Uchida & Uchida, 2018). Dopamine has been implicated in coding the

valence of mating and feeding experiences (Burke et al., 2012; Dai et al., 2022; Stuber

& Wise, 2016; Zhang et al., 2019). Thus, it is possible that dopamine system acts as a

motivation/reward center, allocating limited energy toward different behaviors in an

internal state-dependent manner. This raises a critical question: does dopamine also

mediate interactions among various innate behaviors, such as sleep, aggression,

courtship, fear, and feeding, especially when these behaviors are in conflict?

Additionally, it remains to be determined whether a single group of dopamine neurons

mediates the crosslink amongst these behaviors, or whether distinct neuronal

populations regulate separate behavioral interactions.

Disturbances in innate behaviors have widespread and lasting effects on animals'

overall behavioral patterns and physiological states (Goldstein et al., 2018; Grzelka et

al., 2023; Jourjine et al., 2016; Sun et al., 2023). This cross-behavioral influence is

likely connected to the concept of primal emotions. Emotions are a fundamental

mechanism for behavioral regulations across species, not exclusive to more complex

animals. In *Drosophila*, previous studies have shown that the flies have emotion-like

states such as stress and fear (Anderson & Adolphs, 2014; Gu et al., 2019). Our

research suggests that mating failure may act as a trigger for negative emotions,

affecting other behaviors like feeding and reward responses. This finding provides a

new perspective on how emotions influence behavioral adaptation through neural

mechanisms and offers a model for future research on emotion and behavior.

In summary, this study reveals the neural mechanisms through which mating failure

suppresses sweet taste perception in *Drosophila* and establishes a foundation for

investigating the complex interactions between emotions and the regulation of innate

behaviors. Future research will explore how dopamine system modulates the output of

various innate behaviors under different emotional states and whether such regulatory

mechanism is conserved across species.

Materials and methods

Flies

Flies were maintained in vials with standard cornmeal food at 25°C and 60% humidity in a 12 h:12 h light: dark cycle unless otherwise indicated. For temperature-sensitive flies (Shibire^{ts} and dTRPA1), male flies were kept at 20°C before the behavioral tests. Fly strains used in the manuscript:

elav-GAL4 (#25750), *UAS-mCD8::GFP* (#5137), *LexAop-rCD2RFP*, *UAS-mCD8GFP* (#67093), *10×UAS-IVS-mCD8::RFP, 13×LexAop2-mCD8::GFP* (#32229), *UAS-GCaMP6m* (#42748), *UAS-CalexA* (#66542), *UAS-NaChBac* (#9469), *UAS-Kir2.1*(#6595), *Gr5a-GAL4* (#57592), *Gr5a-LexA* (#93014) were obtained from the Bloomington *Drosophila* Stock Center at Indiana University. *UAS-Shi^{ts}* and *UAS-dTRPA1* were kindly shared by David Anderson (California Institute of Technology). *Dop1R1$^{-/-}$* mutant, *Dop1R2$^{-/-}$* mutant, *Dop2R$^{-/-}$* mutant, *DopEcR$^{-/-}$* mutant, *Th$^{-/-}$* mutant, *Dop1R1-GAL4*, *Dop1R2-GAL4*, *Dop2R-GAL4*, *DopEcR-GAL4*, *TH-GAL4* were kindly shared by Yi Rao (Peking University). *LexAop-GCaMP6m* and *UAS-CsChrimson* were kindly shared by Yufeng Pan (Southeast University). *UAS-Dop1R1 RNAi* (#4278), *UAS-Dop1R2 RNAi* (#5285), *UAS-Dop2R RNAi* (#2141), and *UAS-DopEcR RNAi* (#3275) were obtained from the Tsinghua Fly Center.

Conditioning protocols

We used a modified courtship conditioning protocol (McBride et al., 1999;

Shohat-Ophir et al., 2012) to generate Naïve, Failed, and Satisfied males. The Failed

group was generated by exposing virgin males to either mated females or decapitated

virgin females. Individual virgin males were placed with either 25 mated females or

25 decapitated virgin females in a food vial for 4.5 h. To generate mated females for

the conditioning, 50 males were added to the 25 virgin females for 16 h before the

conditioning. The decapitated female flies were prepared freshly before assays. The

Satisfied group was generated by exposing virgin males to virgin females. Individual

virgin males were placed with 25 virgin females in a food vial for 4.5 h. The Naïve

group consisted of the age-matched virgin males kept isolated from females until the

assays.

The MAFE assay

The MAFE assay was carried out as described previously (Qi et al., 2015). Briefly,

individual male flies were gently introduced and immobilized in a 200 μl pipette tip.

The end of each pipette tip was shaped to expose the fly's proboscis. Flies were first

sated with sterile water and then presented with 400 mM sucrose filled in a graduated

glass capillary (VWR, #53432-604). The sucrose was presented until the flies no

longer responded to a series of 10 sucrose stimuli. The ingested volume was

calculated based on the volume change before and after the feeding.

PER assay

PER was assayed as described previously (Qi et al., 2015). Individual male flies were mounted into a 200 μl pipette tip. Flies were first sated with sterile water and then subjected to different sugar solutions, each tested 3 times. Flies showing PER responses to at least 1 of 3 trials were considered positive to that sugar concentration. For drug treatment experiments, flies were maintained on food containing a drug for 2 days. Drugs used for treatment included 3-Iodo-L-tyrosine (3IY) (Sigma, 10 mg/ml), DL-p-chlorophenylalanine (pCPA) (Sigma, 10 mg/ml), Methysergide maleate (abcam, 25mg/ml). For the dopamine injection experiment, dopamine (Aladdin 0.1-10 ng/μl) was injected into the proboscis of each male fly using a glass micropipette. The standard PER assay was carried out at 25°C. For Shibire[ts] and dTRPA1 experiments, flies were raised at 20°C, and assays were performed at either 20°C or 30°C. For PER assays performed at 24, 48, and 72 hours post-treatment, male flies were conditioned and then transferred into vials containing standard food before the behavioral assays. For Failed-re-copulated experiments, individual Failed male flies were allowed to mate with virgin females in a food vial for 4.5 h. All PER assays had n>4 replicates with 10–15 flies per replicate unless otherwise stated.

MAT calculation

The method to calculate MAT has been thoroughly described in previous papers (Inagaki et al., 2012; Inagaki et al., 2014). In general, sigmoid interpolation was

447   performed to fit the results into sigmoidal curves. The sigmoid curves for sugar were

448   defined as follows:

$$F_S = \frac{1}{1 + e\left(-\alpha_s \, log_2 \, \frac{S_{con}}{S_{50}}\right)}$$

F_S: Fraction of flies showing the PER on different sugar concentrations.

Scon: Sugar concentration.

S_{50}: Sucrose concentration where 50% of flies show PER

α_S: Slope of the sigmoid curve

Based on the experimentally obtained results (F_S, Scon), S_{50} and α_S were chosen to fit

the results. For all experimental results, fitting based on nonlinear regression was

calculated with Python.

Immunohistochemistry

Fly brains or labella were dissected in PBS on ice and fixed in 4% paraformaldehyde

at room temperature for 60 min. After rinsing with PBS for 3 times, the fixed brains or

labella were blocked in Penetration/Blocking Buffer (10% Calf Serum and 0.5%

Triton X-100 in PBS) at room temperature for 90 min and incubated with primary

antibodies at 4°C for 24 h. The samples were then rinsed with Washing Buffer (0.5%

Triton X-100 in PBS) at room temperature for 3×30 min and incubated with

secondary antibodies at 4°C for 24 h. After another 3 rinses with Washing Buffer, the

brains or labella were mounted in Fluoroshield (Sigma-Aldrich) for imaging. Images

were acquired with Nikon C2 confocal microscope ($40 \times /0.80$w DIC N2).

The antibodies were used as follows: rabbit anti-GFP (1:1000, Life Technologies),

mouse anti-GFP (1: 1000, Abcam), rabbit anti-DsRed (1:1000, ClonTech), Alexa

Fluor 488 goat anti-rabbit (1:1000, Life Technologies), Alexa Fluor 488 goat

anti-mouse (1:1000, Life Technologies), Alexa Fluor 546 goat anti-rabbit (1:1000,

Life Technologies).

Calcium imaging

The in vivo calcium imaging experiment was performed as previously described (Qi

et al., 2021). Flies were briefly anesthetized on ice and mounted onto transparent

plastic tape. The cuticle part around the Gr5a$^+$ region of the brain was gently removed

with forceps and the brain was bathed in sugar-free adult hemolymph-like solution

(AHL, 108 mM NaCl, 8.2 mM $MgCl_2$, 26 mM $NaHCO_3$, 1 mM NaH_2PO_4, 5 mM

KCl, 2 mM $CaCl_2$, 5 mM HEPES, pH7.3). A micro manipulator delivered sucrose to

the proboscis of the fly and the actual feeding bouts were imaged by a digital camera

(FLIR USB 2.0, Point Gray). The calcium signals of Gr5a$^+$ neurons were collected by

a Nikon C2 confocal microscope, with a water immersion objective lens ($40 \times /0.80$w

DIC N2).

For CsChrimson experiments, newly enclosed flies were maintained in vials with

standard cornmeal food for 3 days and transferred to standard cornmeal food

supplemented with 400 μM all-trans-retinal (Sigma R2500) for 2 days before

experiments. CsChrimson activation was provided by a fiber-coupled 625 nm LED

(M625L3, Thorlabs). Image analyses were performed in ImageJ and plotted in

GraphPad Prism6 or MATLAB. The ratio changes were calculated using the

following formula: $\Delta F/F = [F - F_0]/F_0$, where F was the peak intensity, and F_0 was

the average intensity at baseline.

Statistical analysis

Data presented in this study were all verified for normal distribution by

D'Agostino-Pearson omnibus test. In one-way ANOVA (for comparisons among three

or more groups) tests, statistical significance was corrected using post hoc Tukey's

test. The two-way ANOVA (for comparisons with more than one variant) tests use

post hoc Bonferroni correction.

Competing interest

The authors declare that they have no competing interests exist.

Acknowledgements

We thank all Wang Lab members for helpful discussions and technical assistance. Ye Wu and Tingting Song provided scientific and administrative support in the laboratory. We thank the Bloomington Drosophila Stock Center at Indiana University and the Tsinghua Fly Center for fly stocks and reagents. This study was funded by the National Key R&D Program of China (2019YFA0801900 and 2019YFA0802400), the National Natural Science Foundation of China (32071006), startup funds from Shenzhen Bay Laboratory and Chinese Institutes for Medical Research.

Author contributions

Gaohang Wang, Resources, Data curation, Formal analysis, Validation, Investigation, Methodology, Writing -original draft; Wei Qi, Investigation; Rui Huang, Resources, Formal analysis, Supervision, Funding acquisition, Investigation, Methodology, Writing - original draft, Project administration, Writing - review and editing; Liming Wang, Conceptualization, Resources, Supervision, Funding acquisition, Investigation, Writing - original draft, Project administration, Writing - review and Editing.

 Figure Legends

 Figure1 Sexual failure decreased sweet sensitivity.

(A) Schematic illustrating courtship conditioning strategy. (B) Percentage of males

that performed feeding to 400 mM sucrose. (C) Fraction of Naïve, Failed and

Satisfied male flies showing PER responses to different concentrations of sucrose.

The PER experiment was performed immediately after conditioning unless otherwise

indicated. Left: average PER responses of Naïve (blue), Failed (black), and Satisfied

(red) flies (n=60-90). Right: MAT (mean acceptance threshold; the sucrose

concentration where 50% of the flies show PER), plotted as a function of sexual state.

(D) Fraction of male flies showing PER responses to different concentrations of

sucrose. The PER experiment was performed immediately, 24 h, 48 h, and 72 h after

conditioning (n=48-60). (E) Fraction of male flies showing PER responses to different

concentrations of sucrose. Failed male flies were generated by two methods: one

group was subjected to mated females (with cVA), and the other was subject to

decapitated virgin females (without cVA) (n=48-60). (F) Fraction of male flies

showing PER responses to different concentrations of sucrose. Failed-re-copulated

males were generated by placing individual failed males with 25 virgin females in a

food vial for 4.5 h. Data are shown as means ± SEM. Error bars represent SEM. n.s.,

$p > 0.05$; *$p < 0.05$; **$p < 0.01$; ***$p < 0.001$, ****$p < 0.0001$.

Figure 2 Dopamine signaling modulated the effect of sexual experience

on sweet sensation.

(A) PER responses in male flies fed with 3IY, compared to male flies fed with vehicle

(n=48-60). (B) PER responses in male flies with dopamine injection, compared to

male flies injected with vehicle (n=48-60). (C-D) PER responses in male flies with

indicated genotypes at 30°C (n=48-60). Data are shown as means ± SEM. Error bars

represent SEM. n.s., $p > 0.05$; *$p < 0.05$; **$p < 0.01$; ***$p < 0.001$, ****$p < 0.0001$.

Figure 3 Dopamine neurons modulated the activity of Gr5a$^+$ neurons

upon sexual failure.

(A) Schematic diagram of *in vivo* calcium imaging paradigm. (B) Quantification of

the *in vivo* calcium responses of Gr5a$^+$ neurons to the indicated concentrations of

sucrose (n=6-22). (C) Averaged traces of the calcium responses to 200 mM sucrose.

The horizontal black bar represents sucrose application. The solid lines represent

average trace, and shaded regions represent ±SEM. (D) Representative pseudocolor

images of the calcium responses to 200mM sucrose. The dashed area was quantified

to measure the response. (E) Top: Representative images from GRASP experiment

between dopaminergic neurons and sweet GRNs. Bottom: Enlarged images of the

SEZ region seen on the top. Scale bars, 20 μm. (F) Average fluorescence reported by

TH>nsyb-spGFP1-10, Gr5a>CD4-spGFP11 (normalized to Naive) in the SEZ. Lines

represent mean ± SEM, and dots indicate values for individual flies. (G) Top:

Representative images of CaLexA-induced GFP reporter in dopaminergic neurons.

Bottom: Enlarged images of the SEZ region shown on the top. Scale bars, 20 μm. (H-I)

Average fluorescence reported by *TH-GAL4/UAS-mLexA-NFAT, LexAop-CD2-GFP*

(normalized to Naive) in the SEZ (H), in the AL (I). Lines represent mean ± SEM, and

dots indicate values for individual flies. Data are shown as means ± SEM. Error bars

represent SEM. n.s., $p > 0.05$; *$p < 0.05$; **$p < 0.01$; ***$p < 0.001$, ****$p < 0.0001$.

Figure 4 The effect of sexual failure was modulated through two

dopamine receptors, Dop1R1 and Dop2R.

(A-E) PER responses in male flies with indicated genotypes (n=48-75). (F) MAT

calculation for the PER data from (A-E), plotted as a function of sexual state and type

of flies. Data are shown as means $\pm$ SEM. Error bars represent SEM. n.s., $p > 0.05$; *p

< 0.05; **$p < 0.01$; ***$p < 0.001$, ****$p < 0.0001$.

Figure 5 Dop1R1^{+} and Dop2R^{+} neurons regulated sexual

experience-dependent sweet sensitivity.

(A-D) PER responses in male flies with indicated genotypes (n=48-60). Data are

shown as means ± SEM. Error bars represent SEM. n.s., $p > 0.05$; *$p < 0.05$; **$p <$

0.01; ***$p < 0.001$, ****$p < 0.0001$.

Figure 6 Gr5a$^+$ neurons received dopaminergic modulation through

Dop1R1 and Dop2R.

(A-B) The expression of membrane-bound GFP (mCD8GFP, Green) in Gr5a$^+$ neurons

driven by *Gr5a-LexA* and membrane-bound RFP (mCD8RFP, Red) in dopamine

receptors neurons driven by *Dop1R1-Gal4* (A), and *Dop2R-Gal4* (B). Scale bars, 20

μm. (C-D) PER responses in male flies with indicated genotypes (n=60-75). (E-F)

Averaged traces of the calcium responses to 200mM sucrose in male flies with

indicated genotypes (n=7-10). The horizontal black bars represent sucrose application.

The solid lines represent average trace, and shaded regions represent ±SEM. Data are

shown as means ± SEM. Error bars represent SEM. n.s., $p > 0.05$; *$p < 0.05$; **$p <$

0.01; ***$p < 0.001$, ****$p < 0.0001$.

Figure 7 Working model for the sexual failure-dependent modulation of

sweet perception.

In the SEZ of the fly brain, dopaminergic neurons form synaptic connections with

sweet-sensing Gr5a$^+$ neurons, which express the dopamine receptors Dop1R1 and

Dop2R. This connectivity enables Gr5a$^+$ neurons to receive dopaminergic modulation,

enhancing the flies' sweet sensitivity. However, after mating failure, the activity of

these SEZ dopaminergic neurons is selectively inhibited, and their synaptic

connections with Gr5a$^+$ neurons are weakened. This reduction in dopaminergic

signaling ultimately decreases the flies' sweet taste perception. This mechanism

highlights a specific pathway by which unsuccessful mating experiences influence

sensory processing through neural plasticity in dopamine-associated circuits.

Figure 1-supplement 1 Sexual failure decreased sweet sensitivity.

(A) Volume of 400mM sucrose consumed by males fed *ad libitum* (n = 35-60). (B)

Fraction of Naïve, Failed and Satisfied male flies showing PER responses to different

concentrations of D-Fructose (n=48-60). (C) Fraction of male flies showing PER

responses to different concentrations of D-Glucose (n=48-60). Data are shown as

means ± SEM. Error bars represent SEM. n.s., $p > 0.05$; $*p < 0.05$; $**p < 0.01$; $***p$

< 0.001, $****p < 0.0001$.

Figure 2-supplement 1 Dopamine but not serotonin signaling modulated

sexual experience-dependent sweet sensitivity.

(A) PER responses in *Th* mutant male flies (n=48-60). (B) PER responses in male

flies injected with various doses of dopamine (n=48-60). (C) PER responses in male

flies fed with DL-p-chlorophenylalanine (pCPA) (n=48-60). (D) PER responses in

male flies fed with methysergide (n=48-60). (E-F) PER responses in male flies with

indicated genotypes at 20°C(n=48-60). Data are shown as means ± SEM. Error bars

represent SEM. n.s., $p > 0.05$; $*p < 0.05$; $**p < 0.01$; $***p < 0.001$, $****p < 0.0001$.

Figure 3-supplement 1 Dopaminergic neurons had functional connections

with sweet GRNs.

(A) The expression of membrane-bound GFP (mCD8GFP, Green) in dopaminergic

neurons driven by *TH-Gal4* and membrane-bound RFP (rCD2RFP, Red) in Gr5a$^+$

neurons driven by *Gr5a-LexA*. Scale bars, 20 μm. (B) Representative image of

GRASP experiment to show connections between dopaminergic neurons and sweet

GRNs in *TH>nsyb-spGFP1-10, Gr5a>CD4-spGFP11* flies (left). Genetic controls of

GRASP experiments were shown on the center and right. Scale bars, 20 μm. (C-D)

Averaged traces of the calcium responses to 200mM sucrose in male flies with

indicated genotypes (n=6-9). The horizontal black bars represent sucrose application.

The solid lines represent average trace, and shaded regions represent ±SEM.

Figure 4-supplement 1 Dop1R1 and Dop2R were necessary for sexual
failure-induced change in sweet sensitivity.

(A-D) PER responses in male flies with indicated genotypes (n=48-60). Data are

shown as means ± SEM. Error bars represent SEM. n.s., $p > 0.05$; *$p < 0.05$; **$p <$

0.01; ***$p < 0.001$, ****$p < 0.0001$.

Figure 5-supplement 1 Dop1R2$^+$ or DopEcR$^+$ neurons did not affect
sweet sensitivity following sexual failure.

(A-B) PER responses in male flies with indicated genotypes (n=48-60). (C) MAT,

plotted as a function of sexual state and type of flies. Data are shown as means ± SEM.

Error bars represent SEM. n.s., $p > 0.05$; *$p < 0.05$; **$p < 0.01$; ***$p < 0.001$, ****p

< 0.0001.

Figure 6-supplement 1 Dopaminergic modulation of Gr5a$^+$ neurons

following sexual failure did not require DopEcR.

(A) The expression of membrane-bound GFP (mCD8GFP) in dopamine receptors

neurons driven by *Dop1R1-Gal4* (left), *Dop2R-Gal4* (center), and *DopEcR-Gal4*

(right). Scale bars, 20 μm. (B) The expression of membrane-bound GFP (mCD8GFP,

Green) in Gr5a$^+$ neurons driven by *Gr5a-LexA* (left) and membrane-bound RFP

(mCD8RFP, Red) in dopamine receptor neurons driven by *DopEcR-Gal4*. Scale bars,

20 μm. (C) PER responses in male flies with indicated genotypes (n=60-75). Data are

shown as means ± SEM. Error bars represent SEM. n.s., $p > 0.05$; *$p < 0.05$; **$p <$

0.01; ***$p < 0.001$, ****$p < 0.0001$.

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Figure 1

Figure 2

Figure 3

A Gr5a-Gal4 > UAS-GCaMP6m

B ΔF/F (%) vs Sucrose (mM)
- Naive
- Failed
- Satisfied

C 200mM Sucrose
ΔF/F (%) vs (s)
- Naive
- Failed
- Satisfied

D ΔF/F (%) 0 — 1000
Pre-stimulation

200mM Sucrose

Naive   Failed   Satisfied

E TH>nsyb-spGFP1-10, Gr5a>CD4-spGFP11

Naive   Failed   Satisfied

F Relative GRASP signal intensity
n.s.
**   ****
Naive n=23   Failed n=30   Satisfied n=22

G TH-GAL4/UAS-mLexA-NFAT, LexAop-CD2-GFP

Naive   Failed   Satisfied

H SEZ
Relative CalexA signal intensity
*
*   ****
Naive n=13   Failed n=13   Satisfied n=12

I AL
Relative CalexA signal intensity
n.s.
Naive n=13   Failed n=13   Satisfied n=12

Figure 4

Figure 5

Figure 6

Figure 2-supplement 1

Figure 4-supplement 1

Figure 6-supplement 1