A neural circuit for female-specific defensive homeostasis in risk assessment

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Highlights

SC CB_1R^+ neurons are necessary for processing visual survival threats in both sexes

CB_1R^+ SC^{GABA} neurons maintain female-specific defensive homeostasis in risk assessment

CB_1R deletion disinhibits GABA release in SC-LHb circuit, impairing female risk assessment

Lacking CB_1R in SC^{GABA} neurons disrupts female stress coping

In Brief

Risk assessment in response to visual threats depends on CB_1R^+ neurons in the SC in both sexes. In females, this process is mediated by a disinhibitory mechanism where eCB from the LHb activates presynaptic CB_1R on SC GABAergic terminals, thereby suppressing GABA release and maintaining defensive homeostasis in risk assessment.

SUMMARY

Risk assessment in defensive behavior is an adaptive mechanism shaped by natural selection, enabling individuals to evaluate potential threats and thereby maintain defensive homeostasis. Although risk assessment is essential for survival, it remains unknown whether specific neural circuits maintain their behavioral homeostasis in a sex-specific manner. Here, we show that visual survival threats activated cannabinoid 1 receptor (CB_1R)-expressing neurons in the superior colliculus (SC), initiating consistent risk assessment in both sexes. Deletion of CB_1R in SC GABAergic neurons female-specifically impairs risk assessment by disinhibiting GABA release in SC-lateral habenula (LHb) projections, resulting in abnormally shortened risk assessment. Furthermore, loss of CB_1R in SC GABAergic neurons increases the occurrence of abnormal spontaneous behavior following chronic stress exclusively in females. Our findings revealed a female-specific CB_1R-modulated SC-LHb GABAergic circuit crucial for maintaining defensive homeostasis in risk assessment, highlighting how disruptions in this circuit underlie deficits in risk assessment and adaptive stress coping.

Key Words: Sex differences; Risk assessment; Defensive homeostasis; Cannabinoid 1 receptor; Superior colliculus; Lateral habenula

INTRODUCTION

Risk assessment during defensive behavior represents an evolutionarily conserved survival mechanism[1-3], refined through natural selection to optimize trade-offs[3,4] between immediate survival threats, such as predation risk, and long-term fitness benefits, such as reproductive success. Such behavioral homeostatic processes to optimize detection and assessment of external stimuli are critical for organism survival. Females exhibit heightened risk-assessment capacities[5,6] to prioritize survival under gestational and lactational constraints[7-9], manifesting as increased risk avoidance[10] and perinatal anxiety[11,12]. Conversely, males display stable risk-taking preferences shaped by selective pressures for resource acquisition[13] and intrasexual competition[14], where consistent behavioral strategies enhance reproductive success. These sexually dimorphic behavioral strategies, driven by asymmetric reproductive investment[15] are evident not only in defensive behaviors[16,17] but also across a range of innate behaviors including mating[18-20] and aggression[21-23] to sex recognition[24-26].

Given that population viability often depends more critically on female survival in many species[27,28], we propose that females evolve more cautious and sensitive defensive homeostasis in risk assessment mechanisms against imminent, life-threatening dangers. These mechanisms prioritize survival certainty over metabolic efficiency, even at the cost of overreaction. Robert Trivers' theory of parental investment[29] suggests that females exhibit risk-averse behaviors to protect their greater reproductive investments, a notion supported by our recent findings of sex-specific defensive strategies in mice facing looming threats[30], suggesting the existence

of sexually dimorphic circuit mechanisms underlying survival behaviors. Neural

circuits modulating sex-dimorphic innate behaviors frequently involve subcortical

hubs that pair conserved anatomy with sex-specific neuromodulation[19,31] to optimize

reproductive success and survival. However, it remains unclear whether subcortical

structures mediate sex-dimorphic innate defensive behavior, particularly those

requiring rapid risk assessment under threat.

The superior colliculus (SC), a conserved hub for processing life-threatening

visual cues across vertebrates[32-36]. Optogenetic activation of SC parvalbumin$^+$ neurons

prolongs escape time in female mice[37], highlighting an evolutionarily driven sexual

dimorphism in defensive control. Cannabinoid 1 receptor (CB_1R), an ancient fear

modulator across vertebrates[38,39], demonstrates sex-specific characteristics in

modulating learned fear and fear-related disorders[40,41], as well as its distribution in the

brain[42,43]. It also contributes to the regulation of behaviors essential for survivals,

including feeding[44], sleep[45], metabolism[46] and social behavior[47,48] as well as

reproductive strategies, such as maternal behavior[49,50] and sexual behavior[51]. It is

noteworthy that endocannabinoid (eCB), as an endogenous ligand of CB_1R, achieves

synthesis and functional action within milliseconds[52], operating orders of magnitude

faster than most documented CB_1R-regulated behaviors. This temporal precision

corresponds to the timescale required for life-or-death defensive responses mediated

by subcortical circuits such as the SC. We therefore hypothesize that the eCB system

evolved specifically to enable precise regulation of survival-critical innate defensive

behavioral homeostasis at subsecond timescales. Thus, investigating the role of CB_1R

in SC-mediated survival threats responses across both sexes is critically important.

Here, we resolve this gap by integrating the hierarchical behavior analysis framework (HBAF)[53] with neural circuit dissection. We demonstrate that risk assessment in response to visual threats is encoded by SC CB_1R^+ neurons in both sexes, while CB_1R in GABAergic SC-lateral habenula (LHb) projections is necessary for female-specific defensive homeostasis in risk assessment. eCB release from the LHb activates the CB_1R on the terminal projecting from the SC, while the deletion of CB_1R in SC GABAergic neurons disinhibits presynaptic GABA release, resulting in a female- specific impairment in risk assessment. Disruption of CB_1R in SC GABAergic neurons induces female-specific stress vulnerability. Importantly, we identify a neural substrate for evolutionarily optimized sex differences in survival strategies, with female-specific defensive homeostasis in risk assessment directly linked to vertebrate-conserved eCB system. These findings establish a novel mechanistic paradigm in which conserved neuromodulatory systems maintain sex-specific defensive homeostasis through subcortical circuit plasticity.

RESULTS

SC CB_1R^+ neurons are necessary for processing visual threats in both sexes

To access CB_1R^+ neurons genetically, we generated a CB_1R^{iCre} knock-in mouse line (Figure S1). *In situ* hybridization confirmed **>** 80% of *CB₁R* mRNA co-localization with Cre protein in both sexes (Figure S1K), exceeded 90% in the superior colliculus (SC) (Figure S1I, K). Quantitative analysis revealed that CB_1R^+ cells make up more than 50% of the SC neuronal populations, independent of sex (Figure S2A-B). To

investigate activation due to threats, we profiled c-Fos mRNA expression patterns

following looming stimuli (Figure S3A). Looming significantly increased c-Fos

mRNA expression and its co-localization with Cre proteins in the SC of both sexes

(Figure S3B-E). Remarkably, **>** 80% of c-Fos^{+} neurons expressed CB_1R (Figure S3F),

indicating predominant engagement of CB_1R^+ neurons in threat processing.

Additionally, we monitored real-time SC CB_1R^+ neuronal activity via calcium

imaging during looming exposure (Figure 1A-B). By injecting either unilateral

AAV2/9-EF1α-DIO-GCaMP6s or a control virus into the SC of CB_1R^{iCre} mice, we

enabled cell-specific recordings (Figure 1A). During looming presentation, we

observed strong calcium transients in SC CB_1R^+ neurons in both male and female

mice (Figure 1B-D), indicating sex-conserved recruitment.

Given the heterogeneity of SC neurons[54], we molecularly characterized the

CB_1R^+ populations using multiplex fluorescent *in situ* hybridization (FISH) (Table

S2). Spatial mapping showed that $CB_1R\square$ neurons predominantly co-expressed

glutamatergic markers (VGLUT2 (*Slc17a6*)$\square$: 28.86±4.83%; CaMKIIα$\square$:

26.72±6.62%) but rarely GABAergic markers (VGAT (*Slc32a1*)$\square$: 21.19±3.89%;

Dlx5/6$\square$ <0.01%) (Figure 1E-I, S2C-D) with no observed sex differences.

CB_1R in SC GABAergic neurons governs female-specific defensive homeostasis

in risk assessment

To dissociate the role of CB_1R in SC GABAergic and glutamatergic neurons, CB_1R-

flox mice were bilateral injected with AAV2/9-Dlx5/6-Cre-GFP (CB_1R cKO) or AAV2/9-

Dlx5/6-GFP (CB_1R Ctrl) (Figure 2A) and AAV-CaMK$\square$α-Cre-GFP (CB_1R cKO) or

AAV-CaMK□α-GFP (CB₁R Ctrl) (Figure S4A). This enabled selective deletion of

CB₁R in GABAergic (Dlx5/6⁺) or glutamatergic (CaMK□α⁺) neurons (Figure 2B,

S4B). Dlx5/6-driven AAVs targeted 94.38±1.94% of GABAergic neurons and

26.52±6.80% of glutamatergic neurons (Figure S5), while CaMK□α-driven AAVs

transduced approximately 71±5.33% of glutamatergic neurons (Figure S5D, S5E).

Using a hierarchical behavioral analysis framework (HBAF)[53], we quantified

defensive responses across groups (Figure 2C, S4C). In Dlx5/6 CB₁R cKO mice, most

escaped to the refuge after looming stimuli (Figure 2D-2E), while some exhibited

alternative strategies like freezing or vigilance (Figure S7, Table S3). Female Dlx5/6

CB₁R cKO mice exhibited accelerated defensive behavior, with reduced response

latency and return time to the refuge (Figure 2E), unlike males (Figure 2D). Refuge

duration remained consistent across groups (Figure 2D-E). In contrast, CaMK□α

CB₁R cKO mice exhibited reduced defensive behavior in both sexes, as evidenced by

fewer flight-to-refuge events (Figure S4D, E). Males additionally showed shortened

refuge time, a phenotype absent in females (Figure S4D).

HBAF-derived 16-movements classification (Table S4, validated in 3D feature

space, Figure 2F-2G, S4F-S4G) identified four behavioral phases: exploration, escape,

hiding, and re-exploration (Figure 2H). Pre-looming movement fractions were

comparable across groups (Figure 2I, J, S6A-D). Female CB₁R cKO mice showed

impaired risk assessment during escape, with decreased looking up (Figure 2L), faster

flight initiation probability (Figure 2P), as well as reduced nose elevation and time-to-

maximum speed (Figure 2R, Movie S1, S2), while males showed no comparable

changes (Figure 2K, 2M, 2O, 2P). These alterations, including a leftward shift in

flight probability distribution (Figure 2P), indicate compromised risk evaluation in

female CB_1R cKO mice. Post-refuge exploration revealed increased sniffing in

females (Figure S6F, S6H) not in males (Figure S6E, S6G), suggesting compensatory

environmental sampling. Principal component analysis (PCA) of the 16-dimensional

movement space detected no behavioral differences in the four phases (Figure S6I-N).

Unlike Dlx5/6 CB_1R cKO mice, CaMK□α CB_1R cKO showed no sex-dependent

phenotypes during escaping (Figure S4K, S4L and Movie S3, S4, Table S5) and

hiding (Figure S4M, S4N), likely due to the low specificity (~70%) of CaMKIIα for

glutamatergic neurons. Subsequent experiments focused on SC GABAergic neurons,

where CB_1R exhibits sexually dimorphic regulatory functions.

CB_1R^+ SC-LHb GABAergic circuit governs female-specific defensive homeostasis in risk assessment

To identify the specific circuits through which downstream SC $CB_1R^+Dlx5/6^+$ neurons

mediate female risk assessment during defensive behavior, we conducted anterograde

neuronal tracing from $CB_1R^+Dlx5/6^+$ neurons in the SC. This revealed over 30

projection targets without sex differences in fiber density (Figure 3). Parallel cell-

specific labeling showed comparable $CB_1R^+ Dlx5/6^+$ neurons density between sexes

(Figure S8), highlighting conserved circuit architecture. c-Fos screening revealed that

LHb neurons were selectively activated by looming stimuli in females following SC

CB_1R ablation in Dlx5/6 neurons (Figure 4A-4D). Given the LHb's established role in

aversive processing[55-62] and eCB-mediated coping[63], we hypothesized that the CB_1R^+

SC-LHb GABAergic circuit regulates female risk assessment. Retrograde tracing

confirmed dense SC CB_1R^+ $Dlx5/6^+$ axons innervation of the LHb (Figure S9). Fiber

photometry recordings used to detect eCB dynamics via the eCB sensor[64-66] (Figure

4E) showed looming stimuli increased eCB and calcium signals in the unilateral LHb

of flight or non-flight-responding mice (Figure 4F-4I), with no sex differences in

signal dynamics (Figure 4J). These results suggest that looming induces similar levels

of post-synaptic neuronal activity and eCB release in downstream nuclei, regardless

of the behavioral response (Flight vs. Non-flight). LHb-specific CB_1R antagonism

female-specifically impaired risk assessment (Figure 4K-4N), unlike interventions in

reuniens thalamic nucleus or zona incerta (ZI) (Figure S10).

Optogenetic activation of CB_1R^+ SC-LHb GABAergic projections (Figure 5A-

5D) disrupted female risk assessment (Figure 5F) but not males (Figure 5E), while

chemogenetic inhibition (Figure 5G-5I) prolonged female risk assessment (Figure 5K),

evidenced by increased latency to initiate flight and return to the refuge. These effects

were not observed in CB_1R^+ GABAergic SC-ZI (Figure S11A-S11D) or SC-lateral

posterior thalamus (LP) (Figure S11E-S11H) protections, confirming the circuit's

necessity. Male defensive behavior (Figure 5J) and spontaneous behavior in both

sexes remained unaffected (Figure 5L-O, S12, Table S6). In summary, CB_1R in SC

GABAergic projections to the LHb are necessary for female to exhibit full,

deliberative risk assessment comparable to males. Their genetic deletion

compromised defensive homeostasis in risk assessment, manifesting shortened and

inadequate threat evaluation prior to flight.

CB$_1$R-mediated suppression of GABA release from SC terminal onto the LHb underlies female-specific phenotypes

Given that presynaptic CB$_1$R inhibits neurotransmitter release[67,68], we measured GABA dynamics in the LHb using a genetically encoded fluorescent sensor for *in vivo* imaging (Figure 6A-6C). Female Dlx5/6 CB$_1$R cKO mice exhibited elevated GABA release during looming (Figure 6E), unlike males (Figure 6D), indicating disinhibited transmission from CB$_1$R$^+$ SC GABAergic projections to the LHb. The LHb expressed high levels of diacylglycerol lipase alpha (Dagla) (Figure 6G), essential for synthesizing 2-arachidonoylglycerol (2-AG)[69,70]. Locally deletion of *Dagla* by bilaterally injecting AAV-EF1α-Cre-EGFP virus into the LHb of Daglα-floxed mice (Figure 6F-6H) induced female-specific risk assessment deficits (Figure 6K), similar to those seen in Dlx5/6 CB$_1$R cKO mice, with no equivalent effects in males (Figure 6J). These findings suggest that 2-AG retrograde signaling in the LHb gates presynaptic GABA release from CB$_1$R$^+$ SC GABAergic terminals, thereby ensuring that females maintain defensive homeostasis in risk assessment comparable to males. Disruption of this circuit through CB$_1$R knockout, circuit modulation, or 2-AG depletion consistently impairs female-specific defensive homeostasis in threat evaluation.

The loss of CB$_1$R in SC GABAergic neurons underlies a female-specific vulnerability to stress

Considering that anxiety can accelerate defensive responses to looming stimuli[71], we assessed stress susceptibility in SC GABAergic CB$_1$R cKO and Ctrl mice (Figure 7A).

Chronic stress resulted in comparable weight stability across groups (Figure 7B), but female Dlx5/6 CB_1R cKO mice showed increased anxiety-like behavior, evidenced by reduced open-arms time in elevated plus maze post-stress (Figure 7C). Stress-induced behavioral reorganization included reduced locomotion (Figure 7G, 7H, 7J, 7K), increased vertical exploration, such as rearing, with decreased nose height (Figure 7I, 7L, 7O, 7P), and decreased horizontal exploration, such as sniffing, with increased nose height (Figure 7I, 7L, 7O, 7P) in male and female CB_1R Ctrl and cKO groups. The stressor elicits a strategic behavioral shift from active exploration to defensive vigilance, manifesting as reduced locomotion and heightened vertical investigation, which potentiates visual information acquisition for environmental surveillance. Female CB_1R cKO mice exhibited a pathological triad post-stress, characterized by anhedonia-like reduced sniffing (10–20 min) (Figure 7L), hypervigilance associated with increased corner occupancy (40–50 min) (Figure 7N), and compulsion-like elevated grooming (50–60 min) (Figure 7L). This progression resembles human anxiety disorders (DSM-5), confirming female-specific vulnerability. Machine learning analysis quantified significant behavioral dissimilarity induced by stress (Figure S13, S14). Collectively, CB_1R deletion in SC Dlx5/6^{+} neurons triggers a female-specific pathogenic cascade characterized by elevated GABA release in the LHb, impaired risk assessment, and heightened vulnerability to stress-induced psychopathology. This defines an evolutionarily conserved circuit that governs sex-specific survival plasticity through a vertebrate-conserved neuromodulatory system.

DISCUSSION

This study identifies a female-specific CB_1R^+ SC-LHb GABAergic circuit that governs defensive homeostasis in risk assessment, suggesting its potential role as a "threat-evaluation switch" to help balance evolutionary trade-offs between predation avoidance and reproductive investment. We demonstrate that the evolutionarily conserved eCB system enables subsecond-scale precision in risk assessment to visual threats, which may represent a critical life-and-death mechanism across mammals. Specifically, GABAergic CB_1R^+ neurons in the SC serves as a neuromodulatory switch optimizing female threat evaluation, consistent with life-history theory predictions regarding sex-specific survival strategies. Furthermore, chronic stress exposure heightens vulnerability in females with circuit dysfunction, indicating that disruption of the CB_1R^+ SC-LHb GABAergic circuit not only impairs defensive homeostasis in risk assessment during acute threats but may also promote persistent stress susceptibility. Thus, despite conserved anatomical connectivity between sexes, neuromodulation via CB_1R may emerge as a central mechanism driving female-specific defensive behaviors and their associated vulnerabilities.

CB_1R as a sexually dimorphic neuromodulator in the homeostatic maintenance of risk assessment

CB_1R, a key component of the evolutionarily conserved eCB system across vertebrates[39], modulates various innate behaviors[72,73] and exhibits sexual differences in neural function[74]. Although associated with locomotor activity[65,75,76], nociceptive processing[77] and affective states[47,78], here we reveal that CB_1R in the SC are necessary for visual threat-induced defensive behavior in both sexes, expanding previous studies upon its role in defensive behavior[79,80]. Notably, although visual threats activate SC

CB_1R^+ neurons in both males and females, the CB_1R^+ SC-LHb GABAergic circuit

female-specifically governs risk computation during defensive behavior, as evidenced

by reduced risk-assessment movements such as looking up. This highlights that

conserved neural activation does not preclude sex-specific neuromodulatory control.

Consistent with prior studies demonstrating that CB_1R modulates behavior in a

highly nucleus-[81-83] and cell type-specific[84-89] manner, our findings further highlight

this functional compartmentalization of eCB signaling. Specifically, despite conserved

anatomical projections between the SC and LHb, CB_1R enhances GABAergic

transmission within this pathway exclusively in females, facilitating more deliberate

threat evaluation, an adaptation likely driven by evolutionary pressures that prioritize

female survival. In contrast, deletion of CB_1R in glutamatergic neurons reduced

escape behavior in both sexes, suggesting a sexually non-dimorphic role in globally

regulating response thresholds to balance defensive and exploratory motivations. This

aligns with the established role of glutamatergic neurons in the SC promoting defense

initiation[33,35,90-92], whereas GABAergic neurons provide inhibitory control over these

responses[93].

GABAergic neurons orchestrate behaviors by regulating excitation-inhibition

balance[94], gating signal throughput[95], and facilitating disinhibition[96], which allows for

precise control over the intensity, direction, and timing. Different GABAergic

subtypes across brain regions encode specific fear behavior[97], regulate defensive

outputs[33], and coordinate vigilance[98,99]. Critically, these GABAergic functions

themselves are often sexually dimorphic[100], contributing to sex differences in

defensive behavior. Thus, these findings suggest that CB_1R modulates defensive behavior through complementary mechanisms, orchestrating behavioral initiation via glutamatergic regulation in both sexes, while maintaining defensive homeostasis through sexually dimorphic modulation via GABAergic circuits. This dual arrangement may support immediate individual survival and long-term species-level reproductive strategies.

This functional dimorphism mediated via CB_1R parallels mechanisms such as tachykinin receptor 1 activation in preoptic area of the hypothalamus neurons, which drives male-specific mating behavior[19]. Both findings highlight how evolutionarily distinct neuromodulatory systems, the endocannabinoid and substance P systems, can be deployed within specialized circuits to promote sex-specific behaviors. We propose that such dimorphism may originate from early developmental programming, as eCB signaling has been shown to regulate microglia-dependent synaptic pruning and astrogenesis, thereby shaping sex differences in juvenile social behavior[101]. Further supporting a developmental basis, endocrine signals have been shown to influence the connectivity and gene expression profiles of hormone-responsive neurons[31], creating latent sex biases that are selectively activated by specific threat contexts in adulthood.

Subcortical GABAergic circuit architecture governs sex-specific threats evaluation

Subcortical circuit governing sex-specific survival strategies exhibit conserved anatomy alongside different neuromodulation. While sexually dimorphic neural architectures are well-documented in hypothalamic[20,102] and limbic hubs[103], manifesting as sex differences in gene expression[104], synaptic activity[105,106], and cell

density[102], the SC-LHb circuit demonstrates functional sexual dimorphism without structural differences, as previously reported[54] and confirmed here. The SC, a conserved detector of survival-relevant visual threats across vertebrates, processes threat information through glutamatergic neurons[33,35,90-92,107] and GABAergic neurons[93]. Our findings reveal a specific sexually dimorphic mechanism within the CB_1R^+ SC-LHb GABAergic circuit. We show that female-specific activation of the LHb depends on CB_1R-mediated retrograde 2-AG signaling that selectively enhances presynaptic inhibition of GABA release. This neuromodulatory specialization aligns with life-history trade-offs that prioritize female survival due to greater reproductive investment.

The LHb serves as a key node encoding aversion, anxiety, and stress, integrating negative valence signals and coordinating adaptive physiological responses[56,108,109]. Our work identifies that the LHb integrates female-specific survival behaviors. Its broad involvement across these behaviors suggests that the LHb may function as a central integrator of survival-stress information, with sex differences in connectivity[55,110] potentially underlying differential susceptibility to stress-related disorders. This integrative function may also account for the heightened threat reactivity often seen in individuals with emotional disorders. Beyond risk assessment, the LHb also contributes to risk preference evaluation[111] and parental behavior[112,113] via distinct afferent pathways, highlighting its broader function in balancing survival-reproduction trade-offs. The engagement of these functions in the LHb highlights its central role in female-specific adaptive behaviors, with eCB signaling providing

essential neuromodulatory flexibility to dynamically prioritize competing demands, based on internal and external contexts.

More broadly, our results align with growing evidence that subcortical structures, including the BNST[19,24,114] and VMH[18,21,31], encode sexually dimorphic behaviors in a cell-type-specific mechanisms involving neuronal populations expressing Tac1[19], serotonin[114], aromatase[24], or progesterone-receptor[18,21]. This supports a unifying paradigm in which evolutionarily conserved subcortical circuits use genetically specified neuronal subpopulations to regulate innate behaviors in a sex-dependent manner. Such functional specialization, often mediated by inhibitory signaling, is conserved across species[115-118] while allowing for species-specific adaptations. Importantly, dysregulation of this circuits can disrupt defensive homeostasis, as evidenced by the LHb's involvement in female-biased disorders such as PTSD[119,120] and postpartum anxiety[121,122], highlighting the clinical relevance of these evolutionarily tuned circuits.

Defensive homeostasis, evolutionary trade-offs and therapeutic translation for sex-biased disorders

The behavioral homeostasis theory suggests that the evolutionary function of defensive behavior is to rapidly maximize an organism's overall readiness to cope with new survival threats and to minimize unnecessary energy expenditure[123-125]. This concept aligns with recently identified homeostatic systems regulating fundamental needs such as sleep[126] and social interaction[127]. Building on our previous findings that complete defensive behavior includes not only threat detection and escape to shelter,

but also the critical process of re-emergence for risk reassessment and environmental exploration[30], we introduce the concept of defensive homeostasis. This homeostatic process involves the entire sequence of responses to life-threatening stimuli, emphasizing the optimization of an organism's ability to rapidly evaluate whether external stimuli present imminent survival risks. Notably, such homeostatic systems often exhibit inherent sexual dimorphism, likely rooted in genetic and epigenetic mechanisms—as seen in the sex-specific imprinting of hypothalamic genes[128] which may predispose differential defensive strategies between males and females. Although both sexes showed consistent escape responses to immediate threats, females displayed quicker re-exploration and more thorough risk reassessment after reaching safety[30]. This enhanced defensive homeostasis in females can be understood through the life-history trade-off theory proposed by William Williams in 1957. Due to asymmetric parental investment, with females bearing the costs of gestation and lactation, female behavioral strategies prioritize survival through more cautious and comprehensive risk assessment. We propose that the comparable assessment duration, between sexes served in females as a proactive phase for gathering and processing more detailed information, thereby \ qualitatively enhancing their defensive homeostasis. The reduction of this duration following CB_1R deletion in SC GABAergic neurons highlights the role of CB_1R in facilitating the deliberative processing characteristic of female defensive strategy.

Chronic stress exposure further highlights these evolutionary trade-offs, where CB_1R dysfunction transforms adaptive vigilance into female-specific psychopathology.

Consistent with the role of CB_1R in managing acute[129] and chronic stress[47,130], Dlx5/6

CB_1R-KO mice exhibit maladaptive behaviors exclusively in females, including

reduced exploration, heightened thigmotaxis, and compulsive grooming. These

phenotypes closely align with the core symptoms of anxiety and depression as

outlined in the DSM-5. This maladaptation may arise from disrupted threat calibration

within the CB_1R^+ SC-LHb GABAergic circuit. From an evolutionary perspective, this

vulnerability reflects the costs of optimizing female survival strategies. Females with

CB_1R dysfunction may overestimate risks, a strategy that could theoretically enhance

offspring protection during maternal investment[131] but imposing significant energetic

costs. Conversely, males maintain more accurate defensive homeostasis in risk

assessment, supporting competitive resource acquisition and reproductive success[13].

These sexually dimorphic behavioral strategies collectively enhance species survival

and reproductive adaptability. Clinically, aberrant risk calibration is mechanistically

linked to disorders predominantly affecting females, such as pathological

hypervigilance in post-traumatic stress disorder (PTSD)[132], which correlated with

habenular hyperactivity[133,134]. Thus, enhancing CB_1R signaling in this circuit presents

a promising therapeutic approach for addressing sex-specific psychopathologies.

Limitations of the study

The Dlx5/6 promotor exhibits approximately 20% glutamatergic specificity, yet fails

to fully resolve the heterogeneity among GABAergic subpopulations in the SC.

Future studies should employ intersectional genetics, such as Cre/Flp-dependent

transgenic mice with viral tools, for subpopulation-specific manipulation. Additionally,

the continuous stress modeling protocol limited the dissection of stage-specific

adaptations. Future designs should incorporate *in vivo* recordings during chronic stress

exposure to characterize temporal dynamics. Finally, therapeutic translation warrants

testing CB_1R-positive allosteric modulators[135] for rescuing female-specific deficits.

FIGURE LEGENDS

Figure 1. SC CB$_1$R$^+$ neurons are activated during visually evoked defensive behavior

(A) Left: Schematic representing viral injection of AAV2/9-EF1α-DIO-GCaMP6s or AAV2/9-EF1α-DIO-EGFP into the unilateral SC of CB$_1$R-iCre-EGFP mice. Right: Representative image showing the recording site within the SC. Scale bar, 500 μm and 50 μm (enlarged view).

(B) Experimental timeline and schematic of the looming stimuli paradigm used to record calcium signals via fiber photometry in a refuge-containing open field apparatus.

(C and D) Left: Average calcium transients for the entire test group in males (C) and females (D). Shaded areas represent error bars, while the gray shadow indicates the looming stimuli event. Right: Plots showing calcium transients during looming stimuli in males (C) and females (D). Unpaired Student's t test with Mann-Whitney test was performed.

(E) Representative sections showing gene expression in the superior colliculus (SC). *Cnr1*, cannabinoid receptor 1; *Slc32a1*, solute carrier family 32 (GABA vesicular transporter), member 1; *Slc17a6*, solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 6; *CaMKIIα*, calcium/calmodulin-dependent protein kinase II alpha; *Dlx5*, distal-less homeobox 5; *Dlx6*, distal-less homeobox 6, displayed from left to right.

(F) Neural subtypes cell plots color-coded by gene expression. magenta, *Cnr1*; red,

Slc32a1; green, *Slc17a6*; cyan, *CaMKIIα*; yellow, *Dlx5*; and gray, *Dlx6*.

(G) Multiplexed fluorescent *in situ* hybridization of *Cnr1* with *Slc32a1, Slc17a6,*

CaMKIIα, Dlx5, Dlx6 (from left to right) in the SC within a single representative

section. Scale bar, 500 μm and 50 μm (enlarged view).

(H and I) Quantification of neural subtypes (I) and double-positive types, expressed as

a percentage of DAPI. Two-Way ANOVA with Sidak's multiple comparisons post hoc

test was performed.

Data are presented as mean ± SEM. n=7 male Ctrl group (C), 6 male GCaMP6s group;

7 female Ctrl group, 6 female GCaMP6s group (D); 13 sections from 3 males and 3

females (H, I). ns (not significant), *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p <$

0.0001. For detailed p values, refer to Table S1.

Figure 2. Deletion of CB_1R in SC Dlx5/6^{+} neurons female-specifically impairs

defensive homeostasis in risk assessment

(A) Schematic representation of viral injection of AAV-Dlx5/6-GFP or AAV-Dlx5/6-

Cre-GFP into the bilateral SC of CB_1R-floxed mice.

(B) Left: Representative images showing AAV-Dlx5/6-GFP (top) and AAV-Dlx5/6-

Cre-GFP (bottom) expression in the SC, alongside CB_1R mRNA (red) and Dlx5/6[+]

cells bodies (green) in CB_1R Ctrl and CB_1R cKO group. Scale bar, 500 μm and 20 μm

(enlarged view). White arrows indicate CB_1R mRNA fluorescent signals colocalized

with GFP positive signals. Right: Quantification of CB_1R^+ cells in Dlx5/6^{+} neurons in

CB_1R Ctrl and CB_1R cKO groups for both sexes.

(C) Experimental timeline and schematic paradigm of virus injection and the looming

test in a refuge-containing apparatus.

(D and E) Defensive strategy (left) and flight-to-refuge behaviors (from left to right: response latency, return time, time spent in the refuge) in male (D) and female mice (E) across CB_1R Ctrl and CB_1R cKO group.

(F) Spatiotemporal feature space of behavioral components.

(G) Sixteen movement types during defensive behavior, represented by corresponding colors.

(H) Schematic paradigm of the different stages of defensive behavior: exploring, escaping, hiding and re-exploring.

(I and J) Ethograms showing the 16 movement types in males (I) and females (J) across groups (CB_1R Ctrl group on the left, CB_1R cKO group on the right). Scale bar, 10 s.

(K-N) Trajectories (left and middle) and quantification of fractions (right) of mice during the escaping stage (from the onset of looming to the end of flight, K, L) and hiding in the refuge (M, N) in males (K, M) and females (L, N) across CB_1R Ctrl and CB_1R cKO groups. Histograms show the percentage of mice occupying horizontal and vertical positions in the apparatus. Symbols indicate mouse locations: five-pointed star for looming onset, triangle for flight initiation, and circle for flight termination.

(O and P) Distribution and positions of mice during flight, looking up, and freezing from 5 s pre-looming to 15 s post-looming in males (O) and females (P) across groups (CB_1R Ctrl on the left, CB_1R cKO on the right).

(Q and R) Left: Real-time speed and nose height from 5 s pre-looming to 15 s post-looming in males (Q) and females (R) across groups. Right: Quantification of the time taken to reach maximum speed and mean nose height during looming stimuli in males (Q) and females (R).

Data are presented as mean ± SEM (B, D, E, Q, R) or mean (K-N). n=5 male CB_1R Ctrl, 4 male CB_1R cKO, 4 female CB_1R Ctrl, 5 female CB_1R cKO (B); 28 male CB_1R Ctrl, 24 male CB_1R cKO (D), 20 female CB_1R Ctrl, 21 female CB_1R cKO (E); 22 male CB_1R Ctrl and CB_1R cKO, 16 female CB_1R Ctrl and CB_1R cKO (K-R). Two-Way ANOVA with Sidak's multiple comparisons post hoc test (B, K-N) and Unpaired Student's t test with Mann-Whitney test (D, E, Q, R) were performed. ns (not significant), $*p < 0.05$, $***p < 0.001$. For detailed p values, refer to Table S1.

Figure 3. Comparable downstream targeting of SC $CB_1R^+Dlx5/6^+$ neurons in males and females

(A) Schematic illustration of the surgical procedure for anterograde tracing of SC $CB_1R^+ Dlx5/6^+$ neurons.

(B) Representative image showing viral infection in SC $CB_1R^+Dlx5/6^+$ neurons in a female mouse. Red, $CB_1R^+Dlx5/6^+$ starting cells; green, terminals of $CB_1R^+Dlx5/6^+$ cells. Scale bar, 500 μm.

(C-N) Representative image of various brain regions targeted by SC $CB_1R^+Dlx5/6^+$ neurons in a female mouse: Re (C), LH (C), LD (D), MD (D), PVT (E), ZI (F), PH (F), opt (F), LHb (G), PVG (H), PaF (H), APT (H), LP (H), LGN (H), PT (I), RPC (J), PP (J), CnF (K), MiTg (K), NLL (K), IC (L), GiA (L), CnF (M), mRt (M), and IOC

1 (N).

2 Abbreviations: Re, reuniens thalamic nucleus; LH, lateral hypothalamus; LD,

3 laterodorsal thalamus; MD, mediodorsal thalamus, PVT, paraventricular thalamus,

4 LHb, lateral habenula; PVG, periventricular gray; PaF, parafascicular thalamic

5 nucleus; APT, anterior pretectal nucleus, LP, lateral posterior thalamus; LGN, lateral

6 geniculate nucleus; PH, posterior hypothalamus; ZI, zona incerta, opt, optic fiber; PT,

7 pretectal nucleus (PT); RPC, red nucleus, parvicellular part; PP, peripeduncular

8 nucleus; CnF, cuneiform nucleus; MiTg, microcellular tegmental nucleus; NLL,

9 nucleus of the lateral lemniscus pontine reticular nucleus; IC, inferior colliculus, mRt,

10 mesencephalic reticular format; GiA, gigantocellular reticular nucleus, alpha part; and

11 IOC, inferior olive, subnucleus C of medial nucleus.

12 (O) Schematic representation showing outputs of SC $CB_1R^+Dlx5/6^+$ neurons.

13 (P) Sankey network showing projections of SC $CB_1R^+Dlx5/6^+$ neurons in male and

14 female mice, with curve width representing connection strength.

15 (Q) Quantification of normalized fluorescence density in ipsilateral downstream brain

16 subregions targeted by SC $CB_1R^+Dlx5/6$ neurons in male and female mice.

17 Data are presented as mean ± SEM. n=3 each group. Two-Way ANOVA with Sidak's

18 multiple comparisons post hoc test was performed. For detailed *p* values, refer to

19 Table S1.

20 **Figure 4. LHb CB_1R antagonism female-specifically impairs defensive**

21 **homeostasis in risk assessment**

22 (A) Left: Schematic representation of viral injections of AAV-Dlx5/6-GFP or AAV-

Dlx5/6-Cre-GFP into the bilateral SC of CB$_1$R-floxed mice. Right: Representative

images showing Dlx5/6$^+$ GFP signals expression in the SC. Scale bar, 500 μm.

(B) Experimental timeline and schematic paradigm of the looming test.

(C) Representative images of c-Fos expression in the LHb in response to looming

stimuli in female CB$_1$R Ctrl and CB$_1$R cKO mice. Scale bar, 200 μm.

(D) Quantification of c-Fos expression in different brain regions across sexes.

Abbreviations: LHb, lateral habenula; LP, lateral posterior thalamus; ZI, zona incerta,

PVT, paraventricular thalamic nucleus; Re, reuniens thalamic nucleus.

(E) Experimental timeline (top), schematic paradigm (middle) and representative

image (bottom) of the viral infection in LHb neurons. Scale bar, 500 μm

(F, G) Heatmap visualization of endocannabinoid (eCB) and calcium (lower panel)

dynamics (left), signals (middle), and peak signals (right) in male (F) and female (G)

mice during flight response to looming stimuli. Shaded areas around the means

indicate error bars; the gray shadow marks the looming stimuli event.

(H, I) eCB and calcium fluorescence changes in male (F) and female (G) mice during

non-flight response to looming stimuli.

(J) Statistical graph showing the area under the curve (AUC) of eCB and calcium

signaling in male and female mice with flight (left) or non-flight (right) response

under looming stimulus.

(K) Timeline and schematic representation of pharmacological experiment during the

looming test.

(L) Representative image showing the placement of track cannula in the bilateral LHb.

Scale bar, 500 μm.

(M and N) Metrics of defensive behavior (from left to right: response latency, return

time, time spend in the refuge and freezing time in the refuge after looming stimuli)

for male (M) and female mice (N) in vehicle (Veh) and NESS 0327 (NESS) groups.

Data are presented as mean ± SEM. n=5 per group (D), 4 per group (H), 5 per group

(F-J), 8 male Veh, 10 male NESS (K), 9 female Veh, 10 female NESS (L). Unpaired

Student's t test with Mann-Whitney test (D, K, L) and two-way ANOVA with Sidak's

multiple comparisons post hoc test (H) were performed. ns (not significant), $*p < 0.05$,

$**p < 0.01$, $***p < 0.001$, $****p < 0.0001$. For detailed p values, refer to Table S1.

Figure 5. Modulating SC-LHb $CB_1R^+Dlx5/6^+$ projections female-specifically

affects defensive homeostasis in risk assessment

(A) Schematic representation of unilateral blue light activation targeting CB_1R^+ SC-

LHb $Dlx5/6^+$ projections.

(B) Protocol for optogenetic stimulation.

(C) Representative image showing ChR2 virus expression in the SC of a CB_1R-iCre-

EGFP mouse and the location of the fiber track (blue, DAPI; red, ChR2-mCherry;

dotted border line, fiber track). Scale bars, 500 μm.

(D) Left: Schematics of patch-clamp slice recording during optogenetic stimulation of

SC $CB_1R^+Dlx5/6^+$ neurons. Right: Light-pulse train triggered spikes (red) in ChR2-

mCherry–positive neurons.

(E and F) Metrics of defensive behavior in males (E) and females (F) (from left to

right: response latency, return time and time spend in the refuge after looming) under

optogenetic activation of CB_1R^+ SC-LHb Dlx5/6$^+$ terminals.

(G) Schematic representation of bilateral chemogenetic inhibition targeting CB_1R^+ SC-LHb Dlx5/6$^+$ projections.

(H) Representative image showing CB_1R^+Dlx5/6$^+$ neurons in the SC and the fiber track location (blue, DAPI; red, ChR2-mCherry; dotted border line, fiber track). Scale bars, 500 μm.

(I) Defensive strategy in males (left) and females (right).

(J and K) Metrics of defensive behavior (from left to right: response latency, return time, time spent in the refuge, and freezing time in the refuge after looming stimuli) in male (J) and female mice (K) in mCherry and hM4Di group.

(L, M) Comparison of the movement fraction between male (L) and female (M) mice in the mCherry and hM4Di groups.

(N, O) Proportion of clusters (left), a random forest classifier trained on 14-dimensional movement fractions to predict group labels (middle), and PCA visualization of behavioral features (right) for male (N) and female (O) in the mCherry and hM4Di groups.

Data are presented as mean ± SEM. n=20 male mCherry, 18 male ChR2 (E), 18 female mCherry and ChR2 (F),11 male mCherry and hM4Di (J, L, N), 9 female mCherry, 12 female hM4Di (K, M, O). Unpaired Student's t test with Mann-Whitney test was performed. ns (not significant), *$p < 0.05$, **$p < 0.01$. For detailed p values, refer to Table S1.

Figure 6. SC-LHb GABA transmission is required for risk assessment in a

female-specific manner

(A) Schematic representation of viral injection: AAV-Dlx5/6-GFP or AAV-Dlx5/6-Cre-GFP into the bilateral SC (left) and AAV-EF1α-iGABASnFR into the unilateral LHb (right) of CB$_1$R-floxed mice.

(B) Representative image showing Dlx5/6$^+$ GFP signals in the SC and iGABASnFR signals in the LHb, and the position of the fiber track (blue, DAPI; green, Dlx5/6$^+$ GFP or iGABASnFR signals; dotted border line, fiber track). Scale bars, 500 μm.

(C) Apparatus for looming test and schematic of the recording system to measure GABA signals using fiber photometry during looming stimuli.

(D and E) Heatmap of iGABASnFR fluorescence changes (left), signals (middle), and peak signals (right) in male (D) and female (E) mice. Shaded areas around the means indicate error bars; the gray shadow marks the looming stimuli event.

(F) Schematic representation of viral injection: AAV-EF1α-EGFP or AAV-EF1α-Cre-EGFP into the unilateral LHb (left) of Daglα-floxed mice.

(G) Representative images showing AAV-EF1α-EGFP (left) and AAV-EF1α-Cre-EGFP (right) expression in the LHb (red, Daglα mRNA; green, EGFP signals). Scale bar, 200 μm and 20 μm (enlarged view).

(H) Percentage of DAGLα$^+$ cells in LHb neurons in DAGLα Ctrl and DAGLα cKO group.

(I) Schematic paradigm for the looming test, open field test and elevated plus maze test.

(J and K) Metrics of defensive behavior (from left to right: response latency, return

time, time spent in the refuge and freezing time in the refuge after looming stimuli) of

male (J) and female mice (K) in DAGLα Ctrl and DAGLα cKO group.

Data are presented as mean ± SEM. n = 8 male CB$_1$R Ctrl, 9 male CB$_1$R cKO (D), 14

female CB$_1$R Ctrl, 8 female CB$_1$R cKO (E), 4 per group (H), 16 male DAGLα Ctrl

and DAGLα cKO (J), 18 female DAGLα Ctrl, 19 female DAGLα cKO (K). Unpaired

Student's t test with Mann-Whitney test. ns (not significant), *$p < 0.05$. For detailed p

values, refer to Table S1.

Figure 7. CB$_1$R deletion in SC GABAergic neurons increases abnormal behavior

patterns specifically in females after stress

(A) Schematic representation of viral injection of AAV-Dlx5/6-GFP or AAV-Dlx5/6-

Cre-GFP into the bilateral SC of CB$_1$R-floxed mice and the timeline of behavioral

teats conducted pre- and post- chronic stress.

(B, C) Changes in body weight and percentage of time spent in open arms during

elevated plus maze test of mice pre- and post- chronic stress across different groups.

#$p < 0.05$.

(D) Trajectories showing speed during 60-min of open field activity.

(E) Representative trajectories illustrating different movements in the open field

apparatus. Each of the 14 colors corresponds to a specific movement type.

(F) Density map showing the spatial distribution of dorsal points during 60-min of

open field (left) and representative trajectories in the center, periphery, and corner

(right) of the apparatus.

(G, J) Travel distance per minute throughout the experiment for males (G) and

females (J).

(H, K) Total travel distance and average speed (10-min epochs) throughout the experiment for males (H) and females (K).

(I, L) Comparison of the fraction (10-min epochs) between male (I) and female (L) mice in the CB_1R Ctrl and CB_1R cKO groups.

(M, N) Pie charts showing relative percentage of time spent in different areas for males (M) and females (N). Inner and outer rings represent pre- and post-stress percentages, respectively, while left and right halves correspond to CB_1R Ctrl (left) and CB_1R cKO (right) groups.

(O, P) Nose height during different movements in the 60-min open field apparatus (bottom) for males (O) and females (P).

Data are presented as mean ± SEM (B, C, G, H, J, K, O, P) and as mean (I, L). n=18 pre-stress, 17 post-stress male CB_1R Ctrl; 18 pre-stress, 16 post-stress male CB_1R cKO; 17 pre-stress, 20 post-stress female CB_1R Ctrl; 17 pre-stress, 18 post-stress female CB_1R cKO. Two-Way ANOVA with Sidak's multiple comparisons post hoc test was performed. #$p < 0.05$; ##$p < 0.01$; ###$p < 0.001$; ####$p < 0.001$ pre-stress versus post-stress; *$p < 0.05$; **$p < 0.01$ post-stress CB_1R Ctrl versus CB_1R cKO For detailed p values, refer to Table S1.

STAR ★ METHODS

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- ● **QUANTIFICATION AND STATISTICAL ANALYSIS**

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AUTHOR CONTRIBUTIONS

F.W. and L.W. conceived the project and designed the experiments. X.L., H.H., X.F., X.L., H. Y., Z.G., Z.L. Z.W and J.L. conducted the experiments and analyzed data.

1 J.Y., H.H. and X.L. contributed to 3D behavioral analysis. R.C., S.C., and Y.L.

2 developed and provided eCB3.0 sensor. F.W., X.L. and H.H. wrote the manuscript

3 with input from all authors. L.W. and F.W. supported all aspects of this project.

4 **DECLARATION OF INTERESTS**

5 The authors declare no competing interests.

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1 **STAR ⬚ METHODS**

2 Detailed methods are provided in the online version of this paper and include the

3 following:

4 **KEY RESOURCES TABLE**

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Antibodies		
Rabbit anti-Cre	Abcam	Cat #: ab190177, RRID: AB_2860024
Goat anti-GFP	Rockland	Cat #: 600-101-215, RRID: AB_218182
Rabbit anti-RFP	Rockland	Cat #: 600-401-379, RRID: AB_2209751
Sheep antibody to digoxigenin peroxidase	Roche	Cat #: 11207733910, RRID: AB_514500
Donkey anti-Goat, Alexa Fluor 488 conjugate	Jackson ImmunoResearch	Cat #: 705-545-003, RRID: AB_2340428
Donkey anti-Rabbit, Alexa Fluor 488 conjugate	Jackson ImmunoResearch	Cat #: 711-547-003, RRID: AB_2340620
Donkey anti-Rabbit, Alexa Fluor 594 conjugate	Jackson ImmunoResearch	Cat #: 705-585-003, RRID: AB_2340432
Bacterial and virus strains		
AAV-hEF1α-DIO-GCaMP6s-WPRE-pA	Taitool Bioscience Co., Ltd., Shanghai	Cat #: S0351-9
AAV-EF1α-DIO-EGFP	Fuqiang Xu at the SIAT	N/A
AAV2/9-Dlx5/6-Cre-GFP	Liping Wang's Lab at the SIAT	N/A
AAV2/9-Dlx5/6-GFP	Liping Wang's Lab at the SIAT	N/A
AAV2/9-EF1α-EGFP	Liping Wang's Lab at the SIAT	N/A
AAV2/9-hSyn -Cre-WPRE-pA	Taitool Bioscience Co., Ltd., Shanghai	S0278-9
rAAV-CaMKIIα-EGFP	Brain Case	Cat #: BC-0027
rAAV-CaMKIIα-Cre-EGFP	Brain Case	Cat #: BC-0165
AAV2/9-Dlx5/6-DIO-mCherry-WPRE-hGH pA	BrainVTA Co., Ltd., Wuhan	Cat #: PT-0850
AAV2/9-mDLX-DIO-hChR2(H134R)-mCherry-WPRE-hGH pA	BrainVTA Co., Ltd., Wuhan	Cat #: PT-0970
rAAV-mDlx5/6-DIO-hM4D(Gi)-mCherry-WPRE-hGH pA	BrainVTA Co., Ltd., Wuhan	Cat #: PT-1191
rAAV-mDlx-FLEX-tdTomato-T2A-Synaptophysin-EGFP	Brain Case	Cat #: BC-0659
AAV2/2Rretro-hSyn-Cre-WPRE-pA	Taitool Bioscience Co., Ltd., Shanghai	Cat #: S0278-2R
AAV9-GfaABC1D-eCB3.0	Yulong Li's Lab at the Peking	N/A

	University	
AAV9-hSyn-RCaMP3	Vigene Biosciences (this paper)	N/A
AAV2-EF1α-iGABASnFR	Fan Jia at the SIAT	N/A
AAV2/9-Dlx5/6-Cre on-ChR2-mCherry	Shanghai Sunbio Medical Biotechnology., Shanghai	N/A
AAV2/9-Dlx5/6-Cre out-ChR2-YFP	Shanghai Sunbio Medical Biotechnology., Shanghai	N/A
Chemicals, peptides, and Recombinant proteins		
NESS 0327	Cayman	Cat #: 10004148
DAPI	Sigma-Aldrich	Cat #: D9542
Clozapine N-oxide	Tocris	Cat #: 4936
Proteinase K	Roche	Cat #: 03654672103
Formamide	Sigma-Aldrich	Cat #: F9037
20×standard saline citrate (SSC)	Thermo Fisher Scientific	Cat #: AM9765
Ribonucleic acid from baker's yeast (S. cerevisiae)	Sigma-Aldrich	Cat #: R6750
Deoxyribonucleic acid from heparin sperm	Sigma-Aldrich	Cat #: D-7290
1× Amplification Diluent	APExBIO	CAT#: H1001
Protector RNase Inhibitor	Roche	Cat #: 3335399001
5×Denhardt's solution	Sigma-Aldrich	Cat #: D2532
DL-Dithiothreitol	Sigma-Aldrich	Cat #: 43816
diethyl pyrocarbonate (DEPC)	Sigma-Aldrich	Cat #: D5758
Fluoromount-G	Southern Biotech	Cat #: 0100-01
Fluorescein Tyramide TSA plus fluorescein kit	APExBIO	Cat#: K1084
Cy3 TSA Fluorescence System Kit	APExBIO	Cat#: K1051
Experimental models: Organisms/strains		
C57BL/6JNifdc mice	Zhejiang Vital River Laboratory Animal Technology Co., Ltd., Zhejiang	RRID: IMSR_JAX:000664
CB$_1$R-floxed	Biocytogen Pharmaceuticals Co., Ltd., Beijing	N/A
DAGLα-floxed	Biocytogen Pharmaceuticals Co., Ltd., Beijing	N/A
CB$_1$R-iCre-EGFP	Biocytogen Pharmaceuticals Co., Ltd., Beijing	N/A
Sprague-Dawley	Zhejiang Vital River Laboratory Animal Technology Co., Ltd., Zhejiang	RRID: RRRC_00239
Oligonucleotides		
Primers for *Cnr1* FISH:	This study	N/A

GTTCAAGGAGAACGAGGA CAAC CAGCAGGCAGAGCATACTA CAG		
Primers for *VGLUT2* FISH: CCAAATCTTACGGTGCTAC CTC TAGCCATCTTTCCTGTTCC ACT	This study	N/A
Primers for *GAD2* FISH: TCTTTTCTCCTGGTGGCG TTGAGAGGCGGCTCATTC	This study	N/A
Primers for *c-Fos* FISH: GCCCACCGACCTGCCTGCA A CTTTCGGTGGGCAGCGGC AC	This study	N/A
Primers for Genotyping of CB₁R-iCre-EGFP: GCCTGCACAAGCACGCCAA T TGAAGGATCGCCGAGCAA CTG GGCACAAGCTGGAGTACA AC	This study	N/A
Primers for Genotyping of CB₁R-floxed: AGGCAACACTCAGTGACA GC TAGGTGGATCTCAGGAATC TTGTAG	This study	N/A
Primers for Genotyping of DAGLα-floxed: GGTGCTGATTGAGGCCTAC CTGT AGCCCTGGCTTGTTTACCC AA	This study	N/A
Software and Algorithms		
Python v3.8	Python Software Foundation	https://www.python.org/, RRID: SCR_008394
R studio Desktop	Posit Software, PBC	https://posit.co/download/rstudi o-desktop/
R v4.3.1	R Core Team	https://www.r-project.org/, RRID: SCR_001905
R package 'networkD3' v0.4 [136]	Allaire J et al., 2017	https://CRAN.R-project.org/package=networkD3
R package 'tidyverse' v2.0.0 [137]	Wickham H et al., 2019	https://doi.org/10.21105/joss.01 686
TissueFAXS Viewer	TissueGnostics	https://www.tissuegnostics.com /cn
Adobe Photoshop	Adobe Systems Inc	https://www.adobe.com/, RRID:SCR_014199

ZEISS Zen Microscopy Software	Zeiss	https://www.micro-shop.zeiss.com/, RRID: SCR_013672
ANY-maze video tracking software	Stoelting Co., IL, USA	https://www.any-maze.com/, RRID: SCR_014289
Graphpad Prism v8.0	GraphPad Software Inc	https://www.graphpad.com/scientificsoftware/prism/, RRID: SCR_002798
Aegisub v3.2.2	Interscape	https://download.html.it/software/aegisub/
OlyVIA v4.1	Olympus	https://www.olympus-lifescience.com.cn/zh/, RRID:SCR_016167
BehaviorAtlas Analyzer v1.0	Bayone BioTech Co., Ltd. Guangdong, China	https://bayonesci.com/
BehaviorAtlas Capture v1.0	Bayone BioTech Co., Ltd. Guangdong, China	https://bayonesci.com/
Automatic infrared behavior-monitor (AIBM) system	Liu Q et al., 2021	https://doi.org/10.1007/s12264-021-00661-4
Fiji	Schindelin J et al., 2012	https://imagej.net/
NetworkX	Python module	https://networkx.org/, RRID：SCR_016864

RESOURCE AVAILABILITY

Lead contact

Further information and requests for resources and reagents should be directed to and

will be fulfilled by the lead contact, Feng Wang (fengwang@siat.ac.cn).

Materials availability

This study did not generate new unique reagents.

Data and code availability

○ The data supporting the current study's findings are available at Zenodo at

https://zenodo.org/uploads/16351971 and are publicly available as of the date of

publication. Original videos will be shared by the lead contact upon request.

○ All original code has been deposited at GitHub at https://github.com/Feng-Wang-Research-Group/A-neural-circuit-for-female-specific-defensive-homeostasis-in-risk-assessment and is publicly available as of the date of publication.

○ Any additional information required to reanalyze the data reported in this paper is available from the lead contact upon request.

EXPERIMENTAL MODEL AND SUBJECT DETAILS

Animals

All experimental procedures adhered to the guidelines established by the Chinese Council on Animal Care and were approved by the Institutional Animal Care and Use Committee (IACUC) of the Shenzhen Institutes of Advanced Technology (SIAT), Chinese Academy of Sciences (No. SIAT-IACUC-200319-NS-WF-A1162 and No. SIAT-BSI-IRB-240613-NS-WF-A0048). The study utilized animals virgin male and female C57BL/6J and mutant mice aged 8~16 weeks and virgin male Sprague-Dawley (SD) rats at an age of 12~20 weeks. All C57BL/6J mice and SD rats were obtained from Zhejiang Vital River Laboratory Animal Technology Co., Ltd (Zhejiang, China). CB_1R-floxed mice were generous gifts from department of Anesthesiology and Perioperative Medicine, Xijing Hospital, Air Force Medical University, Xi'an, China. The CB_1R-iCre-EGFP mouse line was generated as described previously[138]. DAGLα-floxed mice were generated specifically for this study by Biocytogen Pharmaceuticals Co., Ltd., Beijing, China in this study. Genotyping was performed using primers listed in key resources table with correct insertion verified through *in situ* hybridization targeting for gene RNA. Mice were housed in a controlled

environment at 22–25°C on a 12 h light/12 h dark cycle with *ad libitum* access to food

and water. Behavioral tests and recordings were conducted during the 8:00~16:00.

Viruses

(1) *In vivo* **photometry recording experiments**, 300 nL/site AAV-EF1α-DIO-GCaMP6s (2×10^{12} genomic copies/mL; packaged by Taitool Bioscience Co., Ltd., Shanghai) or AAV-EF1α-DIO-EGFP (3×10^{12} genomic copies/mL; packaged by Fuqiang Xu's lab at the SIAT) was injected into the unilateral SC (AP -3.8 mm, ML -0.8 mm, DV -1.8 mm from bregma) of CB_1R-iCre-EGFP mice at a rate of 100 nL/min. 200 nL/site AAV9-GfaABC1D-eCB3.0 (5.46×10^{12} genomic copies/mL; Yulong Li's Lab at the Peking University) and AAV9-hSyn-RCaMP3 (6×10^{14} genomic copies/mL; packaged by Vigene Biosciences Co., Ltd., Shangdong) was infused into the unilateral LHb (AP -1.85 mm, ML -0.55 mm, DV -2.55 mm from bregma) of adult C57BL/6J mice at a rate of 70 nL/min. 50 nL/site AAV-EF1α-iGABASnFR was infused into the unilateral LHb (AP -1.85 mm, ML -0.55 mm, DV -2.55 mm from bregma) of CB_1R-floxed mice at a rate of 70 nL/min. (2) *CB_1R conditional knockout in Dlx5/6$^+$ neurons and control experiments*, 300 nL/site AAV-Dlx5/6-GFP or AAV-Dlx5/6-Cre-GFP (5×10^{12} genomic copies/mL; packaged by Liping Wang's Lab at SIAT) was injected into the bilateral SC (AP -3.8 mm, ML ±0.8 mm, DV -1.8 mm from bregma) at a rate of 100 nL/min. Cre recombination was successful in the target neurons, resulting Cre recombinase expression and the absence of CB_1R mRNA. (3) **Optogenetic manipulation of SC CB$_1$R$^+$ Dlx5/6$^+$ projections to LHb, ZI, or LP**, 300 nL/site Cre dependent recombinant AAV constructs containing hChR2(H134R)-

mCherry (AAV2/9-MDLX-DIO-hChR2(H134R)-mCherry-WPRE-hGH pA; 5×10^{12}

genomic copies/ml; packaged by BrainVTA Co., Ltd., Wuhan) or control viruses

(AAV2/9-Dlx5/6-DIO-mCherry-WPRE-hGH pA; 5×10^{12} genomic copies/ml;

packaged by BrainVTA Co., Ltd., Wuhan) were injected into the unilateral SC (AP -

3.8 mm, ML -0.8 mm, DV -1.8 mm from bregma) of CB_1R-iCre-EGFP mice at a rate

of 100 nL/min. (4) **Confirming SC CB_1R^+ terminals in the LHb**, 300 nL/site AAV-

Dlx5/6-DIO-mCherry   and   AAV-mDlx5/6-FLEX-tdTomato-T2A-Synaptophysin-

EGFP (packaged by Brain Case Co., Ltd., Shenzhen, title: 5×10^{12} genomic copies/mL)

were injected into the unilateral SC of CB_1R-iCre-EGFP mice at a rate of 100 nL/min,

respectively. (5) **Discriminating CB_1R^+ and CB_1R^- projections in SC Dlx5/6$^+$**

neurons, a 1:1 mixture of viruses AAV2/9-Dlx5/6-Cre on-ChR2-mCherry and

AAV2/9-Dlx5/6-Cre out-ChR2-YFP (packaged by Shanghai Sunbio Medical

Biotechnology Co., Ltd., Shanghai, title: 5×10^{12} genomic copies/mL) was injected

into the SC of CB_1R-iCre-EGFP mice at a rate of 100 nL/min. (6) **Chemogenetic**

manipulation of LHb CB_1R^+ GABAergic projections from the SC, 300 nL/site Cre

dependent recombinant AAV constructs containing hM4D(Gi)-mCherry (rAAV-

mDlx5/6-DIO-hM4D(Gi)-mCherry-WPRE-hGH pA; 5×10^{12} genomic copies/ml;

packaged by BrainVTA Co., Ltd., Wuhan) or control viruses (AAV2/9-Dlx5/6-DIO-

mCherry-WPRE-hGH pA; 5×10^{12} genomic copies/ml; packaged by BrainVTA Co.,

Ltd., WuhanLtd., Wuhan) were injected into the bilateral SC of CB_1R-iCre-EGFP

mice at a rate of 100 nL/min. (7) **DAGLα knock-out and control experiments**, 50

nL/site AAV-EF1α-GFP or AAV-EF1α-Cre-GFP (5×10^{12} genomic copies/mL;

packaged by Liping Wang's Lab at SIAT) was injected into the bilateral LHb (AP -

1.85 mm, ML ±0.50 mm, DV -2.75 mm from bregma) at a rate of 70 nL/min.

METHOD DETAILS

Experimental design and randomization

For all between-subject analysis, mice were randomly assigned to experimental

conditions using cage-level randomization (2-3 littermates split between control and

experimental groups) to account for litter effects while minimizing cage-related

variance. Experimental replication was achieved through cohort stratification, with

conditional CB_1R knock out models requiring ≥3 batches (n=3-6/ batches) and other

behavioral and assays experiments requiring ≥2 batches. For 3D behavior analyses,

conditions were randomized, and experimenters remained blinded to group

assignments until statistical was completed. Spatial *FISH* experiments employed

triple-masking protocols during both experimental execution and computational

processing. In neurocircuitry mapping, fiber and cannula placements experiments,

histological verification was performed to assess viral transduction efficiency and

implant localization. Subjects with low transduction efficiency ＜50% or deviation

from target coordinated were systematically excluded prior to data processing.

Stereotaxic surgeries

Stereotaxic injection was performed following the protocol outlined in our published

work[139]. Mice were anesthetized with pentobarbital (i.p., 80 mg/kg) and positioned on

a stereotaxic frame (RWD, Shenzhen, China). Erythromycin eye ointment was applied

to protect the eyes. The scalp was shaved, and a ~1 cm incision was made to expose

the cranium. Connective and muscle tissue were carefully removed, and the skull

surface was leveled to ensure the height difference between bregma and lambda was

< 0.05 mm and aligned in the medial-lateral plane. The skull above the target brain

region was thinned using a dental drill and carefully removed. Viral injections were

performed using a 10-µL Hamilton syringe connected to a 33-gauge needle and a

microsyringe pump (UMP3/Micro4, World Precision Instruments, Sarasota, USA).

The syringe was filled with mineral oil prior to use. At the injection, the needle

remained in place for approximately 5 min to prevent fluid from reflux. Following

surgery and a recovery period of at least 7 days, behavioral tests were conducted.

Cannula implantation

For chemogenetics manipulation and pharmacological experiments, a stainless-steel

bilateral cannula (O.D.0.41 mm×I.D.0.25 mm, c.c 1.0 mm, C 2.8 mm, G1 0.5 mm, G2

0.5 mm, RWD Life Science, Shenzhen, China) was implanted into the LHb. For

pharmacological experiment, a bilateral cannula (O.D.0.48 mm×I.D.0.34 mm, c.c 2.4

mm, C 4.5 mm, G1 0.5 mm, G2 0.5 mm, RWD Life Science, Shenzhen, China) was

implanted in the LHb (coordinates of AP: −1.85 mm, ML: ± 0.5 mm, DV:

−2.55 mm), ZI (coordinates of AP: −2.03 mm, ML: ± 1.2 mm, DV: −4.4 mm)

or a singular cannula (0.48 mm in diameter, 5 mm in length, RWD Life Science,

Shenzhen, China) was implanted in the Re (coordinates of AP: −0.59 mm,

ML: 0.1 mm, DV: −4.2 mm). Behavioral tests were performed one week later

after implantation.

Drug delivery

For chemogenetic inhibition of SC-LHb $CB_1R^+Dlx5/6^+$ projections in freely behaving mice, Clozapine N-oxide (CNO, Tocris, Cat #:4936) was prepared by dissolved it in vehicle solution (sterile saline with 0.2% DMSO). Aliquots were frozen and freshly diluted with saline to a final concentration of 3 μg/μL before bilaterally administrated into the LHb via a bilateral guide cannula. The cannula was connected to two Hamilton syringes via a polyethylene catheter. For each side of the LHb, 50 nL of CNO or vehicle solution was infused slowly at a rate of 70 nL/min using two syringe pumps. After infusion, mice were habituated for additional 5 min to allow CNO diffusion before the needle was withdrawn. Animals were observed for 30 min prior to behavioral tests.

For pharmacological experiments, the neutral CB_1R antagonist NESS0327 (Cayman Chemical, 10004184, Michigan, USA) was locally administered to the LHb via a guide cannula. NESS0327 was suspended in saline containing 0.2% Tween 80. On the day of the experiment, a stainless-steel injection cannula (o.d.:0.41; i.d.:0.25, RWD, Shenzhen, China) was inserted into the LHb. The injection cannula, connected to a micro-injection system housing a 10 μL Hamilton syringe via flexible polyethylene tubing, extended 0.4 mm beyond the guide cannula tip. NESS0327 (1 μM/L) was bilateral injected in the LHb at a volume of 50 nL/site. Animals were observed for 3 min before undergoing behavioral tests.

Histology

Fluorescence In Situ Hybridization (FISH)

FISH was carried out following a protocol similar to that previously described[139].

Complementary DNAs (cDNAs) of *CB$_1$R*, *GAD2*, *VGLUT2*, or *c-Fos* (sequences from

Allen Brain Atlas: CB$_1$R, NCBI Accession: NM_007726.2; GAD2, NCBI Accession:

NM_008078.1; VGLUT2, NCBI Accession: NM_080853.2; c-Fos, NCBI Accession:

NM_010234.2) were cloned from a mouse cDNA library into a pBluescript SK

cloning vector. Sequencing was performed to verify the accuracy of the cloned

sequences. Antisense complementary RNA (cRNA) probes were synthesized by *in*

vitro transcription and labeled with digoxigenin. The following primer sequences were

used for cloning of CB$_1$R, GAD2, VGLUT2, and c-Fos:

CB$_1$R (forward/reverse):

GTTCAAGGAGAACGAGGACAAC/CAGCAGGCAGAGCATACTACAG

GAD2 (forward/reverse):

TCTTTTCTCCTGGTGGCG/TTGAGAGGCGGCTCATTC

VGLUT2 (forward/reverse):

CCAAATCTTACGGTGCTACCTC/TAGCCATCTTTCCTGTTCCACT

c-Fos (forward/reverse):

GCCCACCGACCTGCCTGCAA/CTTTCGGTGGGCAGCGGCAC

Spatial FISH

The specific probes for target RNA (Table S2) were designed by Spatial FISH Ltd

(Shenzhen, China). SC sections from adult male (n=3) and female (n=3) C57BL/6J

mice were fixed by 4% paraformaldehyde (PFA) and placed within a reaction

chamber for subsequent processing. After dehydration and denaturation with methanol,

hybridization buffer containing the specific probes were added to the chamber and

incubated overnight at 37°C. The samples were then washed three times with PBST,

followed by ligation of the target probes in a ligation mix at 25°C for 3 h. After

another three washes with PBST, rolling circle amplification was performed using

Phi29 DNA polymerase at 30°C overnight. Fluorescent detection probes in

hybridization buffer were then applied to the samples. Finally, the samples were

dehydrated using an ethanol series and mounted with mounting medium. Images were

captured using Leica THUNDER Imaging Systems, and signal dots were decoded to

analyze RNA spatial position.

Immunohistochemistry

Mice were perfused with 20 ml of 1×PBS followed by 20 ml of 4% PFA. Brains were

harvested, post-fixed in 4% PFA overnight at 4°C, and dehydrated in 30% sucrose for

48 hours. Using a cryostat (CM1950, Leica, Wetzlar, Germany), 40 μm sections were

prepared. Free-floating brain sections were rinsed with PBS (3×5 min), PBST (0.3%

Triton X-100 in PBS, 1×20 minutes), and then blocked with 3% normal donkey serum

(Solarbio, Cat #:SL050) at room temperature. Sections were incubated overnight at

4°C with primary antibody, including rabbit anti-RFP (1:500 dilution, Rockland, Cat

#:600-401-379), rabbit anti-Cre (1:500 dilution, Abcam, Cat #: ab190177), or goat

anti-GFP (1:500 dilution, Rockland, Cat #: 600-101-215). After washing in PBST

(3×5 min), sections were incubated at room temperature for 1 h with secondary

antibodies: donkey anti-rabbit 594 antibody (1:500 dilution, Jackson

ImmunoResearch, Cat #:711-585-152), donkey anti-rabbit 488 antibody (1:500

dilution, Jackson ImmunoResearch, Cat #:711-547-003) or donkey anti-goat 488

antibody (1:500 dilution, Jackson ImmunoResearch, Cat #:705-545-003). This was

followed by three 10-min washes in 0.01 mol/L PBS. Finally, sections were mounted

and covered with the anti-fade reagent DAPI (1:5000 dilution, Sigma-Aldrich, Cat #:

D9542).

Microscopy Imaging and Quantification of Input Neurons

Virus-labeled signals and immunofluorescence-positive cells were traced using an

Olympus VS120 or VS200 microscope equipped with a 10× objective and analyzed

with Adobe Photoshop software (Adobe, San Jose, USA). For classical image

presentation, sections were photographed using a Carl Zeiss ApoTome.2 microscope

(Zeiss, Oberkochen, Germany) and processed with Zen software (Zeiss) or

TissueFAXS Imaging software (TissueGenostics, Austria) using a 20× objective with

tiled imaging (stitched from multiple images). Cell counting was performed manually

on every third 40 μm thick section, ensuring a 120 μm intervals between successive

sections in the coronal plane. Brain regions were identified based on *The Mouse Brain*

(Stereotaxic Coordinates, fourth edition, Franklin, 2013).

Axon projection mapping

To analyze the projection patterns of SC CB_1R^+ $Dlx5/6^+$ neurons, every third brain

sections (40μm thick) spanning the whole brain was collected. Sections were fully

washed with PBS, mounted, and cover-slipped using 50% glycerol. Images were

captured using an Olympus microscopy with a 10× objective (Olympus VS200).

Brain regions were identified using *The Mouse Brain* (Stereotaxic Coordinates, fourth

edition, Franklin, 2013), and projection patterns were analyzed using ImageJ (ImageJ,

RRID: SCR_003070). In detail, for each animal, all images were thresholded to

remove background noise. Then, 2-5 regions of interest (ROIs) were selected for each

brain region, and the average pixel value of each ROI were calculated after

background subtraction (fluorescence density excluding fiber-free regions). For each

animal, the average fluorescence density of a brain region was determined by

averaging the values across all ROIs. Relative density per region was expressed as a

fraction of the total density (summed across all regions). Data processing was

conducted using the tidyverse R package (v2.0.0), while interactive forced-directed

graphs of relative density averages across neuroanatomical regions were generated

using the networkD3 package (v0.4).

Behavioral assays

Looming behavior

To assess the defensive behavior of male and female mice in response to life-

threatening stimuli, we performed 2D looming behavioral tests using an innately

aversive overhead expanding spot (looming task, as previously described[140]). On Day

1, animals were acclimated to the test arena for 10 min. The arena consisted of an

open-top acrylic cylinder ("open field", 50-cm diameter) connected to an adjacent

alley ("refuge", 50 cm×10 cm), allowing free access between compartments. The

entire setup was enclosed by a 30-cm high wall (Figure 1B). On Day 2, mice were

allowed 5 minutes of free exploration in the open-field arena before being subjected to

1-2 looming trials, depending on the timing accuracy of the first-stimulus, with a 3-minute inter-stimulus interval. The looming stimuli was triggered automatically when mice entered a concentric circular zone (central area: 25 cm diameter) within the arena periphery. The looming stimuli consist of a dark disk expanding from 2° to 40° within 300 ms, maintaining this size for 50 ms before disappearing, and repeating 15 times at 30-ms intervals.

For the SC CB$_1$R knock-out experiments, a 3D looming behavioral tests was assessed using multi-view motion-capture device system, as previously described[30]. A transparent looming arena with the same parameter as the 2D behavioral tests was placed at the center of a supporting framework, equipped with four depth cameras (Inter, RealSense D435i) connected to the BehaviorAltas Capture system (version 1.01, Shenzhen Bayone Biotech Co.), ensuring full-view motion capture. Two monitors, mounted on a fixed shelf 45 cm above the bottom of the looming box, delivered uniform and stable white light for behavioral tests. Light intensity was set to 120 lux at the center of the circular trigger zone and 30 lux at the refuge midpoint, matching the brightness parameters of 2D looming behavior setup. Behavioral recordings lasted 15 min, capturing 30 frames per second. matching the brightness parameters of 2D looming behavior setup. The other experimental parameters were consistent with those used in the 2D tests.

Open field test (OFT) in the multi-view motion-capture device

The open field apparatus (50×50×50 cm, matte white polycarbonate) was calibrated to access anxiety-like behavior and locomotor activity. To collect unobstructed

behavioral data for elaborate analysis, a multi-view video capture device was

employed to record spontaneous behavior (Figure S12A, Figure 7A)[53]. Briefly, a

50×50 cm transparent acrylic square enclosure with white ground was placed at the

center of a support framework either 130×130×90 cm (Figure S12) or a 90×90×75 cm

(Figure 7). Four cameras were rigidly mounted on structural support pillars, with pan-

tilt heads finely adjusted to achieve full coverage of the arena. Behavioral recordings

were synchronized (30 Hz, 640×480 or 960×540-pixel resolution) and processed

using BehaviorAtlas Capture Software® v2.0.0.

3D Behavioral data processing

Data acquired from the BehaviorAtlas Capture System underwent three sequential

stages as previously described[53]: unsupervised movement annotation, supervised label

refinement with manual correction of misclassified movement labels, and ethological

quantification.

Unsupervised movement annotation pipeline

Using pre-training body point data from images, the skeleton of each mouse included

in the study was automatically reconstructed by BehaviorAltas Analyzer software,

incorporating camera calibration information. Movements were classified by

clustering skeleton kinematic features for each frame and assigning digital labels.

Supervised movement labeling

Movements with postural similarities, particularly those occurring in low-light refuge

areas, were subjected to dual-criteria validation with skeleton kinematic parameters

such as speed, head-to-limb angle, body length, and distances between the nose and

limbs. Specific movements like grooming, pausing, and looking up were

systematically reviewed and reclassified when discrepancies arose between 3D

kinematic data and manual annotations (see below). Additionally, movements such as

flight and freezing were manually annotated by experts due to significant inter-

individual variability in kinematic patterns.

Kinematic reclassification protocol

Movement	Threshold
Pausing	Average velocity (based on back coordinate) < 30 mm/s and the movement duration > 2 s
Hunching	Average nose height ranges from 30 to 45 mm and average back height > 25 mm
Looking up	Average nose height > 20mm and average front claw height < 15 mm
Right turning	Average velocity (based on back coordinate) > 50 mm/s and body angle from top-view (neck-back-root tail) >15°
Grooming	Average body angle in 3D space (neck-back-root tail) < 120°
Rotating	Average velocity (based on back coordinate) > 30 mm/s and body angle from top-view (neck-back-root tail) >20°

Ethograms

Four key defensive behavior metrics, flight probability, response latency, return time,

and time in the refuge, were quantified through manual frame-by-frame analysis

(Figure 2D-E). Hierarchical ethograms were constructed by segmenting movement

fragments within specific time windows (Figure 2I-J). Trajectory analysis was

performed to quantify mouse locomotion by tracking the coordinates of the mouse's

back, integrating spatial and temporal movement parameters. Defensive behavior

quantification curves were created by mapping three stereotypic responses, risk-

assessment posturing, flight initiation, freezing duration, against threat-exposure

timeline, with group-averaged movement probability on the y-axis and time on the x-axis. Peak response markers indicated behavior-specific latency (Figure 2O-P). Defensive behavior analysis also incorporated nasal elevation dynamics and locomotor velocity, derived from 3D pose estimation outputs (Figure 2Q-R).

Sample classification and segregation

Movement classification was performed using a random forest classifier with PCA-reduced kinematic features, encompassing all movements and their frequencies. Random forest machine learning algorithms were applied to movement data for group classification, with accuracy evaluated through detailed confusion matrices.

Movement transition and central node definition

Movement transitions were analyzed using probability matrices that captured how often one movement followed another, disregarding the duration of each movement. Line thickness in the diagrams represented the frequency of transitions, while dot size indicated the likelihood of specific movement events (Figure S6K-N). Transitions with probabilities below 0.05, after statistical adjustments, were excluded. Central nodes were identified by calculating the total distance of each movement to all others, with the four movements having the smallest total distances displayed for each group (Figure S12F-I). Network diagrams were generated using Python's NetworkX package (v2.8.4) with spring layout settings.

Elevated plus maze (EPM)

The EPM test was performed a using maze made of white acrylic glass, comprising of a central platform (5×5 cm), two opposing open arms (32.5×5 cm), and two opposing

closed arms (32.5×5×32.5 cm). The maze was elevated 50 cm above the floor. Mice

were placed at the junction of the open and closed arms, facing the closed arms.

Behavioral tracking (5-minute sessions) was conducted using ANY-maze® v7.4

software, with body centroid tracking. Time spent in each area (center, open arms,

closed arms) was quantified.

Between behavior trials, the arena was cleaned by scrubbing with a 25% ethanol

solution.

Chronic rat stress model

The experimental protocol was modified from previous studies[141,142]. In the chronic

stress model, mice were exposed to a rat for 10 consecutive days. Each day, a mouse

was placed in a cage containing a rat for 5 seconds and then co-housed with the rat

using a ventilated plexiglass divider. This apparatus allowed odor transmission while

preventing physical contact, ensuring continuous exposure to predator odors.

Behavioral tests, including the open-field test and elevated-plus maze, were conducted

at two time points: 3 days prior to the establishment of the model (Pre-stress) and on

day 10 post-exposure to rat-induced stress (Post-stress).

Patch-clamp electrophysiology

Coronal brain slices (300 µm thick) containing the SC region (bregma 3.2–4.4 mm)

were prepared from CB_1R-iCre-EGFP mice using standard procedures. Brains were

quickly separated and chilled in ice-cold modified artificial cerebrospinal fluid (ACSF)

containing (in mM): 110 Choline Chloride, 2.5 KCl, 1.3 NaH_2PO_4, 25 $NaHCO_3$, 1.3

Na-Ascorbate, 0.6 Na-Pyruvate, 10 Glucose, 2 $CaCl_2$, 1.3 $MgCl_2$. SC slices were

sectioned in ice-cold modified ACSF using a Leica vibroslicer (VT-1200S). Following

sectioning, slices were allowed to recover for 30 min in a storage chamber containing

regular ACSF at 32~34 °C (in mM): 125 NaCl, 2.5 KCl, 1.3 NaH_2PO_4, 25 NaHCO3,

1.3 Na-Ascorbate, 0.6 Na-Pyruvate, 10 Glucose, 2 $CaCl_2$, 1.3 $MgCl_2$ (pH 7.3~7.4

when saturated with 95% O2/5% CO2). Slices were then maintained at ~25 °C until

transferred to the recording chamber. All solutions had an osmolarity of 300~320

mOsm/kg. For electrophysiological experiments, slices were visualized using infrared

optics on an upright microscope (Eclipse FN1, Nikon Instruments) with a 40× water-

immersion objective. The recording chamber was continuously perfused with

oxygenated ACSF (2 ml/min) at 34 °C. Patch pipettes were pulled by a micropipette

puller (Sutter P-2000 Micropipette Puller) and had a resistance of 3–5MΩ. Electrodes

were filled with intracellular solution containing (in mM): 130 potassium gluconate, 1

EGTA, 10 NaCl, 10 HEPES, 2 $MgCl_2$, 0.133 CaCl2, 3.5Mg-ATP, 1 Na-GTP. In SC

slices, mCherry-positive neurons were patched in current-clamp mode. Action

potentials were elicited by stimulation SC neurons expressing ChR2 with blue light

pulse (10 ms duration, 20 Hz).

Optical fiber implantation

Fifteen min after AAV injection, a ceramic ferrule with an optical fiber (200 μm in

diameter, N.A. 0.37) was implanted, positioning the optical fiber tip above the LHb

(AP: -1.90 mm, ML: ±0.50 mm, DV: -2.55 mm from bregma), ZI (AP: -1.79 mm, ML:

+1.10 mm, DV: -4.3 mm from bregma), or LP (AP: -2.03 mm, ML: +1.20 mm, DV: -

2.2 mm from bregma). Optogenetic stimulation was performed 4 weeks after fiber implantation.

Three weeks after AAV injection, a ceramic ferrule with an optical fiber (200 μm diameter, N.A. 0.37) was implanted, positioning the fiber tip above the SC (AP: -3.8 mm, ML: -0.8 mm, DV: -1.5 mm from bregma), or LHb (AP: -1.90 mm, ML: ±0.50 mm, DV: -2.45 mm from bregma),. Fiber photometry recording was conducted one week after fiber implantation.

Optogenetics

Optogenetic manipulations were performed using laser systems with wavelength-specific settings. Terminal activation in SC-LHb $CB_1R^+Dlx5/6^+$ experiments: a blue laser (473 nm; Aurora-300-470, NEWDOON, Hangzhou) was used to deliver 10 mW/mm² at the fiber tip, with 5-s illumination, 20 Hz frequency, and 10-ms pulse duration.

Fiber photometry

(1) Fluorescence signals of GCaMP6s and GABASnFR were recorded via fiber photometry as previously described[141], using 470-nm LEDs for target excitation and 410-nm LEDs for isosbestic control. The 410-nm light intensity was calibrated to 20 μW/mm² at the fiber tip to match the 470-nm intensity. (2) Fluorescence signals of eCB3.0 and RCaMP3 were recorded via fiber photometry (vista, Inper), using 470-nm, 560 -nm LEDs for target excitation and 410-nm LEDs for isosbestic control. The 410-nm light intensity (0.5 mW/mm²), 470-nm light intensity (0.8 mW/mm²), 561-nm light intensity (1.2 mW/mm²), was calibrated to at the fiber tip. Behavioral video

synchronization was achieved through dedicated system software. Raw data were

smoothed using a 5-25-point moving average variance, with variance analyzed across

different experiments. ΔF/F, areas under the curve, and peak values were calculated

using least-squares regression to scale the 410-nm to 470-nm or 560-nm traces.

QUANTIFICATION AND STATISTICAL ANALYSIS

Statistics

For all experiments, statistical analyses were performed using either unpaired two-

tailed Student's t test or a two-way ANOVA followed by Holm-Sidak's multiple

comparisons test. The statistical significance threshold was set at $\alpha=0.05$, with

significance levels indicated as $*p < 0.05$, $**p < 0.01$, $*** p < 0.001$, $**** p <$

0.0001). Exact p-values are provided in the figure legends and Table S1. The "n" in

these analyses represents the number of mice, cells, or brain sections. Statistical

analyses were carried out using GraphPad Prism 10 (GraphPad Software Inc.).

Additionally, a custom Python script was utilized to evaluate the differences in

movement transition and postures through a permutation test (10,000 iterations),

comparing mean disparities between the two groups. Comprehensive statistical

information is available in the figure legends and Table S1.

Figure 1. SC CB₁R⁺ neurons activation during visually evoked defensive behavior

Figure 2. Deletion of CB₁R in SC Dlx5/6⁺ neurons impairs female risk assessment

Figure 3. Downstream targets of SC CB₁R⁺Dlx5/6⁺ neurons

Figure 4. LHb CB₁R antagonism impairs female risk assessment

Figure 5. Modulating SC-LHb CB₁R⁺Dlx5/6⁺ projections affects female risk assessment

Figure 6. SC-LHb GABA transmission is required for risk assessment in a sex-dependent manner

Figure 7. CB₁R deletion in SC GABAergic neurons increases the abnormal behavior patterns in stressed females