# A molecularly distinct accumbal-to-lateral hypothalamic circuit modulates food seeking and consumption

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## Abstract

Understanding the mechanism of energy homeostasis is expected to lead to effective treatment to obesity and metabolic diseases[1,2]. However, energy homeostasis is a complicated process largely controlled by neuronal circuits in the hypothalamus and brainstem[3-5], whereas reward and motivation of food intake are mainly controlled by the limbic regions[6] and cerebral cortex[7,8]. Although the limbic and hypothalamus connection like Nucleus Accumbens shell (NAcSh) to the lateral hypothalamus (LH) circuit has been reported to regulate feeding[9,10], the neuron subtypes involved, and how do the humoral/neuronal signals coordinate to direct feeding behavior remain unknown. Here we show that the projection from dopamine receptor D1(Drd1)- and *Serpinb2*-expressing subtype to leptin receptor (LepR) expressing neurons in LH modulates food seeking and consumption. We demonstrate that the *Serpinb2*$^{+}$ neuronal activity is dynamically modulated during feeding. Conversely, chemo/optogenetics-mediated modulation of *Serpinb2*$^{+}$ neurons bidirectionally regulate food seeking and consumption. Importantly, circuitry stimulation revealed the NAcSh$^{Serpinb2}$→LH$^{LepR}$ projection controls refeeding and overcomes leptin-mediated feeding suppression. Ablation of NAcSh$^{Serpinb2}$ neurons could decrease body weight. Together, our study reveals a molecularly distinct accumbal-to-lateral hypothalamic neural circuit that controls internal state-dependent food consumption, which provides a promising therapeutic target for anorexia and obesity.


## Main

The hypothalamus, with highly heterogenous neuronal composition[11], plays a critical role in controlling feeding behavior[12,13], where feeding related hormones, such as ghrelin and leptin, coordinately produce sensations of appetite and satiety leading to behavioral response[14,15]. Traditionally, people regard the arcuate nucleus (Arc) as a major location where LepR performs anorexic function by acting on leptin receptors (LepR) to suppress food intake and bodyweight gain[16]. In addition to the Arc, lateral hypothalamus (LH)[17] also highly expresses LepR to play a similar role. However, the specific neuronal subtype targeting LH$^{LepR}$ neurons beyond hypothalamus to regulate feeding remain to be elucidate. In recent years, several studies have analyzed the role of mediodorsal NAcSh in feeding[10,18,19] and revealed that activation of dopamine receptor 1 expressing medium spiny neurons (D1-MSNs)[20,21] projections to LH[9,10] stops ongoing food consumption. However, other studies showed that D1-MSNs activity was

enhanced during appetitive phase[22] as well as during consumption[23]. Although temporally distinct phases of feeding behavior, such as food seeking, food evaluation and consumption, could potentially account for such discrepancy, the neuronal heterogeneity of NAc[24] could underlie the seemingly conflict feeding behavior as the different studies might have manipulated different neuron subtypes with opposing functions. With the application of single cell RNA-seq and spatial transcriptome techniques to decipher the neuron heterogeneity of different brain regions[25-29], we could focus on different neuron subtypes located in the NAcSh implicated in feeding behavior[9,10,19,22,30,31] to identify the neuron subtype(s) responsible for regulating feeding behavior.

**_Serpinb2_$^+$ neurons are activated in refeeding process**

Using iSpatial[32], an algorithm that integrates single cell transcriptome and spatial transcriptomic information[24], we analyzed NAc MSN subtypes with medial dorsal NAcSh distribution. We found that the *Tac2, serpinb2* and *Upb1* MSN subtypes exhibit distinct distribution patterns in medial dorsal NAcSh (Fig. 1a). Since D1-MSNs, but not D2-MSNs, provide the dominant source of accumbal inhibition to LH with rapid control over feeding via LH GABA neurons[19,22], we focused our effort on the *Tac2* and *Serpinb2* D1-MSN subtypes. tSNE plots of scRNA-seq result indicated that the *Tac2*$^+$ neurons are mainly enriched in the D1-MSN subclusters 6 and 8, while *Serpinb2*$^+$ neurons are enriched in D1-MSN subcluster 2[24] (Fig. 1b). RNA-FISH further confirmed that both *Tac2* and *Serpinb2* subtypes belong to D1-MSN (co-express Drd1) and are mainly localized to medial dorsal NAcSh (Fig. 1c). Consequently, *Tac2*$^+$ and *Serpinb2*$^+$ MSN subtypes are good candidates with potential in mediating feeding behavior.

To determine whether *Tac2*$^+$ and *Serpinb2*$^+$ MSN subtypes can mediate feeding behavior, we first asked whether the activities of these neurons respond to feeding behavior by monitoring the *cFos* expression under three conditions: ad libitum access to food, after 18 hours of fast, and refeeding (Fig. 1d). By counting *cFos*$^+$ neurons that co-express *Serpinb2* or *Tac2* in the medial dorsal NAcSh under Ad libitum, fast and refed conditions (Fig. 1e), we determined whether the *Serpinb2* and *Tac2* neurons respond to the different feeding status. We found fasting and refeeding both increased neuronal activities compared to Ad libitum as indicated by the increased *cFos*$^+$ neuron numbers (Fig. 1f). Importantly, most of the *Serpinb2*$^+$ neurons (~70%)

were activated by refeeding, but not fasting (Fig. 1g). In contrast, the *Tac2*$^{+}$ neurons do not respond to refeeding or fasting (Fig. 1h). Collectively, these data indicate that the majority of *Serpinb2*$^{+}$ neurons respond to refeeding process.

**The *Serpinb2*$^{+}$ neurons respond to eating behavior**

To facilitate studying the role of *Serpinb2*$^{+}$ neurons in the feeding process, we generated a *Serpinb2*-Cre mouse line (Extended Data Fig. 1a,b). We validated this mouse model by injecting a Cre-dependent AAV vector expressing light-gated cation channel channelrhodopsin (ChR2) and observed about 90% colocalization of *Serpinb2*::ChR2-eYFP signal with endogenous *Serpinb2* mRNA signal (Extended Data Fig. 1c), which is consistent with endogenous *Serpinb2* expression in the NAcSh as shown by Allen Brain Atlas. Thus, our *Serpinb2*-Cre mouse line can be used for studying *Serpinb2*$^{+}$ neurons.

To test whether *Serpinb2*$^{+}$ neurons are involved in regulating feeding, we used fiber photometry to monitor *Serpinb2*$^{+}$ neuronal activity during food seeking and consumption. To this end, we inserted an optic cannula into the NAcSh to record the total *Serpinb2*$^{+}$ neuronal activity (reflected by calcium reporter fluorescence intensity) by stereotaxic injection of a Cre-dependent AAV expressing the seventh-generation calcium reporter GCaMP7s into the NAcSh region of the *Serpinb2*-Cre mice. In parallel, we also implanted cannula to the *Drd1*-Cre mice for comparative study (Fig. 2a). Three weeks after the viral injection, we performed fluorescence recording during the feeding process. To monitor the activity dynamics of the *Serpinb2*$^{+}$ neurons, we designed 3-chamber food seeking and food consumption assay (Fig. 2d) and aligned calcium traces of mice with behavioral events that include habituation, food zone approaching, eating, and leaving (Fig. 2b, c). First, we recorded the $Ca^{2+}$ signals during the different feeding phases after the mice were fasted overnight. In habituation phase, mice were allowed to freely travel among the 3 chambers and we detected negligible response when mice approach the two empty food cups (Fig. 2d, first graph). In food approaching phase, when mice enter the food zone to interact with caged food pellets, the activity of the *Serpinb2*$^{+}$ neurons increased immediately when mice entered the food zone and lasted for seconds (Fig. 2d, second graph), indicating that *Serpinb2*$^{+}$ neurons respond to appetitive food seeking. In the eating phase, we observed a significant increase of $Ca^{2+}$ signals when mice start eating the food (Fig. 2d, 3$^{rd}$ graph), and

subsequently, declined to lower than baseline level after eating finished. Usually, mice turned away and left the food zone (Fig. 2d, 4$^{th}$ graphs).

To quantify the total *Serpinb2*$^+$ neuronal activity at different phases, we calculated the area under curve index (AUC) of the calcium signal curve for each trial. We find that the AUC is significantly higher in the food zone approaching and eating phases and lower in the post eating phases compared with that of habituation (Fig. 2d, right panel). In parallel experiments with the *Drd1*-Cre mice, we observed $Ca^{2+}$ signal tended to increase when approach to food zone (Fig. 2e, second graph) and reduce during consumption (Fig. 2e, 3$^{rd}$ graph). After eating, *Drd1*$^+$ neuronal activity increased concomitantly (Fig. 2e, 4$^{th}$ graphs). Similar results were also observed in ad libitum status (Extended Data Fig. 2). These results indicate that *Serpinb2*$^+$ neurons function differently from other *Drd1*$^+$ neurons especially in appetitive and consumption phases.

**_Serpinb2_$^+$ neurons bidirectionally regulate food intake in hungry state**

To determine whether *Serpinb2*$^+$ neuronal activity plays a causal role in regulating feeding behavior, we asked whether the feeding behavior can be changed by manipulating *Serpinb2*$^+$ neurons' activity. To this end, we applied chemogenetic techniques[33] to the *Serpinb2*-Cre mice by injecting the activating AAV encoding a modified human M3 muscarinic receptor (hM3Dq) or the inhibitory vector AAV-DIO-hM4Di-mCherry into the NAcSh region. As controls, we also performed parallel experiments using *Tac2*-Cre and *Drd1*-Cre mice (Extended Data Fig. 3a). We first confirmed the accuracy of the injection site (Extended Data Fig. 3b), then the efficiency of the activation and inhibition with *cFos* expression (Extended Data Fig. 3c).

We then tested whether activation or inhibition of the *Serpinb2*$^+$ neurons affect feeding and reward-related behaviors (Fig. 3a). Interestingly, under Ad libitum conditions, manipulation of the *Serpinb2*$^+$ neuronal activity does not affect food intake (Fig. 3b**,** left panel), indicating that the eating behavior under Ad libitum maybe controlled by other neurons. Next, we performed the same test using fasted mice to analyze total food consumption during refeeding (Fig. 3a-c). We found that activation of the *Serpinb2*$^+$ neurons increased food consumption, while inhibition of the *Serpinb2*$^+$ neurons decreased food consumption (Fig. 3b**,** right panel). However, similar manipulation on *Tac2*-Cre or *Drd1*-Cre mice did not affect food consumption (Fig. 3c). In

addition to the total food consumption, we also analyzed food preference indicated by the time spent in the food zone (Fig. 3c). As expected, fasted mice would significantly increase their time spent in the food zone for the control mouse group (Fig. 3d). Importantly, activation (hM3Dq) or inhibition (hM4Di) the *Serpinb2*$^+$ neurons respectively increased or decreased the time the mice spent in the food zone compared to that of the control mice (Fig. 3e). In contrast, similar manipulation of the *Tac2*-Cre or *Drd1*-Cre mice showed no such effect compared to that in the control mice (Fig. 3e). These results indicate that manipulation the *Serpinb2*$^+$ neuronal activity can regulate food seeking and intaking behavior in hungry state.

To further determine whether the neuronal activity of the *Serpinb2*$^+$ neurons has a causal role in regulating food motivation, we next carried out food operant chamber test (Fig. 3f, g). To maintain a similar food motivation status, all mice were food restricted to reduce body weight to around 90% of their original value. After trained to operantly respond to sweetened chow pellets on fixed ratio (FR) 1, 3, 5 schedule, the animals were then tested for lever pressing upon CNO-induced chemogenetic manipulation. For the FR5 test, *Serpinb2*$^+$ neuron activation significantly increased the active lever pressing and pellet reward (Fig. 3h), while *Serpinb2*$^+$ neuron inhibition elicited the opposite effect (Fig. 3h). For the progressive ratio (PR) 5 test, *Serpinb2*$^+$ neuron activation showed a tendency of increased number of active lever pressing and significantly increased pellet reward (Fig. 3i), while *Serpinb2*$^+$ neuron inhibition decreased both the total active lever pressing and the reward (Fig. 3i). Taken together, these results demonstrate that the *Serpinb2*$^+$ neurons are involved in bidirectional control of goal-direct food seeking behavior.

NAc shell is also known to regulate anxiety[34] and drug reward[35,36]. Thus, we asked whether *Serpinb2*$^+$ neurons also regulate these behaviors under the same chemogenetic manipulation. We found that manipulation of *Serpinb2*$^+$ neuronal activity does not affect locomotion in open field test (Extended Data Fig. 4a,b), or drug reward in cocaine conditioned place preference (CPP) test (Extended Data Fig. 4c), or anxiety in elevated plus maze (EPM) test (Extended Data Fig. 4d,e). Taken together, these results support that *Serpinb2*$^+$ neurons are specifically involved in food refeeding process, but not involved in locomotion, anxiety or drug seeking behaviors.

**_Serpinb2_$^+$ neurons mediate food consumption via LH projection**

Thus far, characterization of the *Serpinb2*$^+$ neurons was at the somatic level in NAcSh. Next, we attempt to understand how the *Serpinb2*$^+$ neurons regulate food taking behavior at the circuit level. Previous studies have indicated that NAcSh D1-MSNs could project to multiple brain regions, including ventral tegmental area (VTA)[37], ventral pallidum (VP)[38], and bed nucleus stria terminalis (BNST)[39]. To determine the projection site of *Serpinb2*$^+$ neurons, we performed anterograde tracing by injecting Cre-dependent AAVs expressing ChR2(H134R) into the NAcSh of the *Serpinb2*-Cre mice (Fig. 4a). Analysis of the brain slices after 3 weeks of virus injection revealed that the *Serpinb2*$^+$ neurons only project to lateral hypothalamus (LH) (Fig. 4a). To further validate this projection, we injected the widely used retrograde tracer Cholera toxin subunit B (CTB) [40] into the LH region (Fig. 4b), and observed colocalization of CTB with mCherry in the NAcSh after immunostaining of NAcSh from the *Serpinb2*-Cre mice (Fig. 4b, c), supporting the NAcSh to the LH projection.

To demonstrate that the NAcSh to the LH projection of the *Serpinb2*$^+$ neurons is functionally relevant to food intaking, we asked whether optogenetic manipulation of the *Serpinb2*$^+$ neuron terminals in LH can change the feeding behavior of the *Serpinb2*-Cre mice. To this end, DIO-ChR2-eYFP or DIO-NpHR-eYFP AAV viruses were injected to the NAcSh of the *Serpinb2* -Cre mice with optic cannula implanted into their LH region (Fig. 4d). After fasting the mice for overnight, we activated the *Serpinb2*$^+$ neuron terminals in the LH with blue light on-off stimulation (20 Hz, 2-ms pulses). We found that mice with the *Serpinb2*$^+$ neuron terminal stimulation significantly increased their food intake in 20 mins compared to the control mice that express eYFP (Fig. 4e, left panel). Conversely, *Serpinb2*$^+$ neuron terminal inhibition in the LH with yellow on-off stimulation (20 Hz, 2-ms pulses) decreased total food intake when compared with the control (Fig. 4e, right panel). Collectively, viral tracing and circuit manipulation demonstrate that the NAcSh to LH projecting *Serpinb2*$^+$ neurons have an important function in regulating food intaking behaviors.

## *Serpinb2*$^+$ neurons form a circuit with LH LepR$^+$ GABA$^+$ neurons

Having demonstrated the functional importance of the NAcSh to LH projection, we next attempted to determine the neuron types in LH that receive signals from the *Serpinb2*$^+$ neurons. The LH is a highly heterogeneous brain region controlling food intake, energy expenditure, and

many other physiological functions[41]. Since neural peptides orexin/hypocretin and melanin-concentrating hormone (MCH) are associated with feeding[42,43], and are mainly express in LH, we first asked whether they are the down-stream targets of the NAcSh *Serpinb2*$^+$ neurons. To this end, we injected the AAV-DIO-ChR2 anterograde viruses to the NAcSh of the *Serpinb2* -Cre mice and performed immunostaining of candidate neural peptides or transmitters on slices covering the LH (Fig. 5a). We found very few MCH- or orexin-expressing neurons in LH overlapped with *Serpinb2*$^+$ terminals (Fig. 5b, indicated by arrow heads). Given that GABAergic neuron is the major subtype in the LH and has been reported to be involved in feeding and leptin-regulated energy homogenesis [11,44,45], we next checked whether leptin receptor (LepR) positive GABAergic neurons[17] receive projections from NAcSh *Serpinb2*$^+$ neurons by similar immunostaining. We found over 70% *Serpinb2*$^+$ neuron terminals overlap with LepR$^+$ GABA$^+$ neurons (Fig. 5c). This data indicates that the NAcSh *Serpinb2*$^+$ neurons mainly project to LepR$^+$ GABA$^+$ neurons in LH.

To identify the brain regions that response to refeeding, we performed whole brain *cFos* mapping on the refeeding mice. By postmortem immunostaining, we found *cFos* expression was increased in multiple brain regions compared with that of ad libitum, like NAc, Olfactory tubercle (OT), paraventricular nucleus of the thalamus (PVT), Arc, Dorsomedial nucleus of the hypothalamus (DMH), LH and Zona incerta (ZI) (Fig. 5d, Extended Data Fig. 5). We next attempted to identify inputs for the NAcSh *Serpinb2*$^+$ neurons by performing a modified rabies tracing experiment. To this end, the Cre-inducible avian sarcoma leucosis virus glycoprotein EnvA receptor (TVA) and rabies virus envelope glycoprotein (RG) were injected unilaterally to the NAcSh of *Serpinb2*-Cre mice (Fig. 5e) to allow monosynaptic retrograde transportation and rabies virus infection in the starter neurons, respectively[37,46,47]. Two weeks later, the modified rabies virus SADDG-EGFP (EnvA) was injected unilaterally into the NAcSh and slices of the whole brain were imaged one week later. Confocal imaging results indicated that EGFP-labeled neurons can be found in anterior cingulate area (ACA), PVT, LH, and lateral preoptic area (LPO) (Fig. 5f, Extended Data Fig. 6), indicating that neurons in these regions send monosynaptic projection to NAcSh to form a network regulating food consumption (Fig. 5g).

Combining the major efferent regions with *cFos* mapping (Fig. 5d, f), PVT may serve as a major input for *Serpinb2*$^+$ neurons in NAcSh to regulate food seeking and taking. Collectively, our study uncovered a neuronal network where the NAcSh *Serpinb2*$^+$ neurons may receive signals from PVT neurons to inhibit the LepR$^+$GABA$^+$ neurons in LH, to regulate food seeking and eating behaviors.

**Modulating *Serpinb2*$^+$ neuronal activity can overcome leptin effect and alter bodyweight**

As an adipose-derived hormone, leptin plays a central role in regulating energy homeostasis[48-50]. Leptin performs most of its functions, including suppression of food intaking, by activating the LepR on central nerve system (CNS) neurons[51,52]. Since the NAcSh *Serpinb2*$^+$ neurons are projected to LepR$^+$ GABAergic neurons in LH (Fig. 5c), we anticipate that both leptin and the NAcSh *Serpinb2*$^+$ neurons have shared neuron targets and consequently they should have functional interaction. To analyze their functional interaction in food intaking, we implanted catheter in the LH for leptin delivery (catheter administration) on the *Serpinb2*-Cre mice that were also injected with hM3Dq-mCherry-expressing AAV into the NAcSh so that the NAcSh *Serpinb2*$^+$ neurons can be activated by CNO by i.p. injection. First, we established that 1 μg of bilateral intra-LH leptin cannula delivery[53] significantly decreased food intake in 3 hours compared to the control with saline treatment (Fig. 6a). Then we used 1 μg of leptin for all the following tests. As we have shown previously (Fig. 3b), CNO-induced *Serpinb2*$^+$ neuron activation increased food intake (Fig. 6b). Importantly, although leptin delivery reduced the food intake, the leptin effect can be at least partly overcome by CNO-induced *Serpinb2*$^+$ neuron activation (Fig. 6b). This data indicates that *Serpinb2*$^+$ neurons' innervation to LH can at least partly overcome leptin's inhibitory effect on food intake.

To access whether loss function of the *Serpinb2*$^+$ neurons can exert a long-term effect on energy homeostasis, we selectively ablated NAcSh *Serpinb2*$^+$neurons in *Serpinb2*-Cre mice by injecting a flex-taCasp3-TEVp AAV expressing caspase-3 which eliminates the neurons by inducing cell death (Fig. 6c). NAcSh *Serpinb2*$^+$ neurons ablation decreased food intake (Fig. 6d) as well as reduced body weight gain by 10% in 7 weeks (Fig. 6e). Together, these results demonstrate that transient change in *Serpinb2*$^+$ neuronal activity could alter food seeking and consumption

behaviors. Modulating *Serpinb2*$^+$ neuronal activity could partly overcome leptin effect to maintain energy homeostasis, and ablation of *Serpinb2*$^+$ neurons can lead to bodyweight loss.

## Discussion

Feeding is an essential goal-directed behavior that is heavily influenced by homeostatic state and motivation. The accumbal-to-lateral hypothalamic pathway has been implicated in regulating feeding behavior, but the specific neuron subtypes and precise neuronal circuit in LH are not clear. In this study, we filled in this knowledge gap by delineating a circuit which integrates neuronal and humoral signal to regulate food consumption in an innate energy state-dependent manner. Specifically, we identified a D1-MSN subtype located in the NAcSh and expresses *Serpinb2* to regulate the feeding behavior through a neuronal circuit involving NAcSh$^{Serpinb2}$→LH$^{LepR}$ that is beyond the hypothalamus. We demonstrate that the *Serpinb2*$^+$ neurons bidirectionally modulate food motivation and consumption specifically in hungry state. Importantly, *Serpinb2*$^+$ neurons target the LepR expressing GABAergic neurons in LH and their activation can at least partly overcome the suppressive effect of leptin on food intaking, and whose ablation can chronically cause bodyweight loss.

### The *Serpinb2*$^+$ neurons are functionally distinct from the pan D1-MSNs in NAcSh

Previous studies have observed reduced D1 firing during food consumption, and consistently, suppressing D1-MSNs activity prolonged food intake[19]. Using *Serpinb2*-Cre and *Drd1*-Cre mice, we compared the *Serpinb2*$^+$ MSNs neurons and the D1-MSNs in regulating feeding behaviors (Fig. 2, 3), and found their manipulation have different outcomes. First, *Serpinb2*$^+$ neurons are activated by both food approaching and food consumption, while *D1*$^+$ neurons show bi-phasic response, activated during food approaching but suppressed during food consumption. Second, *Serpinb2*$^+$ neurons bidirectionally regulate food seeking and intake, particularly during refeeding, while *D1*$^+$ neuron manipulation does not significantly alter feeding behavior. On the other hand, a previous study showed that *D1*$^+$ neuron inhibition promoted liquid fat food intake[19]. This difference might be due to the different feeding assays used in the two studies, while we used free-access chow food intake, the previous study used a head-fixed mice licking liquid fat food as the assay[19]. Third, *Serpinb2*$^+$ neuron ablation significantly reduced food intake (Fig. 6), which

is consistent with our finding that *SerpinB2*$^+$ neuron activation positively regulates food intake (Fig. 3). However, a previous study indicated that lesions or inactivation of the NAc neurons do not significantly alter the food consumption[54]. We do not consider these results to be in conflict as NAc is composed of many D1- and D2-MSN neuron subtypes, of which many are not involved in regulating food intake, while others can positively or negatively regulate food intake. Consequently, manipulating *Serpinb2*$^+$ neurons and the entire NAc neurons can have totally different outcomes. This further indicates that finer granularity and cell type-specific approaches are needed to dissect the function of different neuron subtypes in NAc. For example, although the *D2*$^+$ neuronal activity as a whole is not altered during food consumption[19,22], the D2 receptors are indeed downregulated in obese rodent as well as human[55,56], whether certain D2-MSN subtypes are involved in regulating food intake remains to be determined.

**_Serpinb2_$^+$ and _Tac2_$^+$ MSNs respectively modulate food and drug reward**

The NAcSh has long been implicated in regulating reward-related behaviors that are associated with food, social, and drug. Previous studies were mainly based on dichotomous MSNs subtypes that express dopamine receptor 1 or dopamine receptor 2 (D1- or D2-MSNs)[57]. Increasing evidence suggest that the NAcSh is highly heterogeneous in terms of the molecular features and anatomical connections of the neurons located in this region. This raises an interesting question that whether these reward-related behaviors are regulated by distinct or overlap neuron subtypes and/or projections. By combining iSpatial analysis, *cFos* mapping, neuronal activity manipulation of different subtypes, and behavioral tests, we found that the *Serpinb2*$^+$ D1-MSNs of NAcSh specifically regulate food reward, but not drug reward or other emotional and cognitive functions (Fig. 1, 3, Extended Data Fig. 4). On the other hand, we have previously showed that the *Tac2*$^+$ D1-MSNs of NAcSh specifically regulate cocaine reward[58]. These studies indicate that different reward behaviors are at least partly regulated by distinct MSNs subtypes. An important task for future studies will be to identify the relevant neuron subtypes regulating the various reward-related behaviors, and eventually to link the cellular heterogeneity to functional diversity of each brain regions. This way, the cellular and circuit mechanisms underlying the various behaviors can be elucidated.

**The PVT- NAcSh $^{Serpinb2+}$ -LH $^{LepR+}$ circuit controls feeding in hungry state**

Previous studies have shown that either LH GABA or Vglut2 neuronal subpopulation can receive NAc innervation[19,59]. However, the LH GABA and Vglut2 neurons are extremely heterogeneous, and can be further divided into 15 distinct populations, respectively[60]. Thus, the specific cell types that receive NAc innervation were unknown. Previous studies also showed that NAcSh D1- MSN to LH inhibitory transmission stops eating, and endocannabinoids mediated suppression of this projection promotes excessive eating of highly palatable chow[61], but the D1 subtype involved in this projection was not known. Using viral tracing, we discovered that the NAcSh *Serpinb2*$^+$ D1-MSNs project to *LepR*$^+$ neurons in LH underlying the *Serpinb2*$^+$ neuron function in food intake (Fig. 6c). Distributed in numerous regions involved in the regulation of energy balance, the *LepR*$^+$ neurons lie in the mediobasal hypothalamic (MBH) "satiety centers" and in LH that is regarded as the "feeding center"[50,62]. Leptin treatment induced *cFos* expression and 100 nM of leptin depolarized 34% of LepR-expressing neurons in LH[17]. Unilateral intra-LH leptin decreased food intake and bodyweight[17]. In our study, we found activation of the *Serpinb2*$^+$ neurons increased the inhibition of the *LepR*$^+$ neurons excitability, resulting in increased food consumption; while inhibition of *Serpinb2*$^+$ neurons decreased the inhibition of *LepR*$^+$ neurons excitability, leading to decreased food consumption even after fasting (Fig. 3b, 4e). Our results are consistent with previous reports demonstrating that LH LepR neuron activation decreased chow intake[63]. Importantly, manipulating *Serpinb2*$^+$ neuronal activity could override leptin's effect in LH to modulate food consumption (Fig. 6b). It is not clear whether the lack of effect of the *Serpinb2*$^+$ neurons on feeding behavior in ad libitum is due to the lack of food taking motivation or the relatively high leptin level masked the *Serpinb2*$^+$ neuron effect. Alternatively, it is possible that the endocannabinoid and leptin signaling may interact in LH, where activation of NAcSh *Serpinb2*$^+$ neurons suppresses LepR neurons in the LH, which may increase the synthesis and release of endocannabinoids[64] and thus promote feeding. Our study thus reveals a parallel and compensatory circuit from NAcSh to LH$^{LepR}$, which is beyond the hypothalamus circuit that directly modulates food intake, to maintain energy homeostasis.

For the efferent site, *Serpinb2*$^+$ neurons majorly receive input from PVT based on the GFP$^+$ cell numbers (Fig. 5f), which is believed to be an integration hub processing information and sending "command" to the downstream targets[65-67]. Previous studies revealed that the *Slc2a2*$^+$ neurons *in*

*PVT* are activated by hypoglycemia and their activation by optogenetics increases motivated sucrose seeking behavior[68]. On the other hand, the *Gck*$^+$ neurons in PVT have the opposite glucose sensing property as their optogenetic activation decreased sucrose seeking behavior[69]. Taken together, we believe that in hungry state, PVT receives the "hungry signal" and send it to *Serpinb2*$^+$ neurons. The activated *Serpinb2*$^+$ neurons then instruct the LH *LepR*$^+$ neurons to promote eating. This PVT- NAcSh $^{Serpinb2+}$-LH $^{LepR+}$ circuit controls feeding in hungry stage and strengthens the sentinel role of NAcSh[19,70].

**_Serpinb2_$^+$ neurons connect energy homeostasis and motivation**

Feeding is an essential goal-directed behavior that is influenced by cellular homeostasis state and appetitive motivation. Interestingly, we found that *Serpinb2*$^+$ neurons promote feeding in hunger state, rather than in normal feeding state, suggesting that *Serpinb2*$^+$ neuron function is regulated in an internal metabolic state-dependent manner. Consistently, ablation of *Serpinb2*$^+$ neurons significantly reduced food intake and leading to bodyweight loss (Fig. 6e,f). The *Serpinb2*$^+$ neurons are activated by both appetitive food approaching and food consumption. To measure appetitive food motivation, we conducted operant food intake assay, where mice need to press levers to earn food pellets. We found that *Serpinb2*$^+$ neuronal activity bi-directionally regulates active lever presses and the earned reward (Fig. 3h,i). These studies suggested that *Serpinb2*$^+$ neurons regulate both energy homeostasis and appetitive motivation which is consistent with the demonstrated function of NAc in integrating descending signals pertaining to homeostatic needs and goal-related behaviors[3,71]. Collectively, these data indicate that lose function of the NAcSh *Serpinb2*$^+$ neurons can disrupt energy homeostasis and appetitive motivation, which provides a potential therapeutic target for obese treatment. Conversely, activation of the NAcSh *Serpinb2*$^+$ neurons can be a potential strategy for anorexia treatment.

In conclusion, we identified a molecularly defined neuron population with crucial functions in regulating food intake via neuron-hormone axis. From a therapeutic point of view, our findings are highly relevant because activating or ablating a small population of molecularly defined neurons could respectively rescue food intake at low energy status or lead to long-term bodyweight loss. Given its function, we believe the small population of NAcSh *Serpinb2*$^+$

398 neurons is an ideal entry point for understanding the complex brain-metabolism regulatory
399 network underlying eating and bodyweight control.
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## Methods

### Mice

All experiments were conducted in accordance with the National Institute of Health Guide for Care and Use of Laboratory Animals and approved by the Institutional Animal Care and Use Committee (IACUC) of Boston Children's Hospital and Harvard Medical School. The *Serpinb2*-Cre mice were generated as described below. For behavioral assays, 12 -16 weeks old male mice were used. The mice were housed in groups (3-5 mice/cage) in a 12-hr light/dark cycle, with food and water ad libitum unless otherwise specified. The Tac2-Cre knock-in mouse line was a gift from Q. Ma at Dana-Farber Cancer Institute and Harvard Medical School. The D1-cre mouse line was obtained from Jackson Laboratory (JAX: 037156).

### Fluorescence *in situ* hybridization (FISH) and immunofluorescence (IF) staining

Mice were transcardially perfused with PBS followed by 4% paraformaldehyde. Brains were then placed in a 30% sucrose solution for 2 days. The brains were frozen in Optimal Cutting Temperature (OCT) embedding media and 16 μm (for FISH) or 40 μm (for IF) coronal sections were cut with vibratome (Leica, no. CM3050 S). For FISH experiments, the slices were mounted on SuperFrost Plus slides, and air dried. The multi-color FISH experiments were performed following the instructions of RNAscope Fluorescent Multiplex Assay (ACD Bioscience). For IF, cryostat sections were collected and incubated overnight with blocking solution (1×PBS containing 5% goat serum, 5% BSA, and 0.1% Triton X-100), and then incubated with the following primary antibodies, diluted with blocking solution, for 1 day at 4 °C. Samples were then washed three times with washing buffer (1×PBS containing 0.1% Tween-20) and incubated with the Alexa Fluor conjugated secondary antibodies for 2 h at room temperature. The sections were mounted and imaged using a Zeiss LSM800 confocal microscope or an Olympus VS120 Slide Scanning System. Antibodies used for staining were as follows: rabbit anti-cFos (1:2000, Synaptic systems, #226003), chicken anti-GFP (1:2000, Aves Labs, no. GFP-1010), chicken anti-mCherry (1:2000, Novus Biologicals, no. NBP2-25158), Orexin-A (KK09) antibody, (1:500, Santa Cruz Biotechnology, Cat# sc-80263 ), anti-MCH, Ab1-pmch antibody (1:2000, Phoenix Pharmaceuticals, Cat# H-070-47 ), Anti-GABA antibody (1:1000, Sigma, Cat# A2052) , Leptin Receptor antibody (Abcam, Cat# ab104403), Goat anti-chicken Alexa Fluor 488(Thermo Fisher

Scientific, Cat# A11039), Donkey anti-rabbit Alexa Fluor 488(Thermo Fisher Scientific, Cat# A21206), Donkey anti-rabbit Alexa Fluor 568(Thermo Fisher Scientific, Cat# A10042).

**AAV vectors**

The following AAV vectors (with a titer of $>10^{12}$) were purchased from UNC Vector Core: AAV5-EF1a-DIO-hChR2(H134R)-EYFP, AAV5-EF1a-DIO-EYFP, AAV-DJ-EF1a-DIO-GCaMP6m. The following AAV vectors were purchased from Addgene: AAV5-hSyn-DIO-hM3D(Gq)-mCherry (#44361), AAV5-hSyn-DIO-hM4D(Gi)-mCherry (#44362), AAV5-hSyn-DIO-mCherry (#50459), pAAV-flex-taCasp3-TEVp (#45580), pAAV-Ef1α-DIO eNpHR 3.0-EYFP (#26966), EnvA G-deleted Rabies-EGFP (Salk Institute).

**Stereotaxic brain surgeries**

The AAV vectors were injected through a pulled-glass pipette and the nanoliter injector (Nanoject III, Drummond Scientific - 3-000-207). The injection was performed using a small-animal stereotaxic instrument (David Kopf Instruments, model 940) under general anesthesia by isoflurane (0.8 liter/min, isoflurane concentration 1.5%) in oxygen. A feedback heater was used to keep mice warm during surgeries. Mice were allowed to recover in a warm blanket before they were transferred to housing cages for 2–4 weeks before behavioral evaluation was performed. For chemogenetics experiments, 0.1~0.15 μl of AAV vector was bilaterally delivered into target regions. For optogenetics experiments, following viral injection, the fiber optic cannula (200 μm in diameter, Inper Inc.) were implanted 0.1 mm above viral injection site and were secured with dental cement (Parkell, #S380). For the drug delivery cannula implantation, the cannula (Guide cannula: C.C 2.0mm, C=4.5mm; Injector: G1=0.5mm; Dummy cannula: G2=0, RWD Life Science) were directly implanted 0.5 mm above the LH and were secured with dental cement (Parkell, #S380). The coordinates of viral injection sites are based on previous literature as follow: NAc (anterior-posterior [AP] +1.2, medial-lateral [ML] ± 0.6, dorsal-ventral [DV] -4.5 mm) and LH (AP -1.3, ML ± 1.2, DV -5.0 mm).

**Neuronal Tracing**

For CTB tracing, mice were injected with 100–200 μl CTB-647 (AF-CTB, all from Life Technologies) unilaterally into the LH (AP –1.2, ML +1.2, DV –4.75 mm). For rabies tracing,

*Serpinb2*-cre mice were first unilaterally injected in the NAcSh with the starter AAV. After 14 d, the same mice were injected with the rabies virus. Then, 11 days after CTB injections and 7 days after rabies injections, brains tissue was collected and processed for confocal imaging. To aid visualization, images were adjusted for brightness and contrast using ImageJ, but alterations always were applied to the entire image.

**Behavioral assays**

**Open-field tests (OFT).** A clear box (square 27.3 cm x 27.3 cm square base with 20.3 cm high walls) used for the open field test, and the center zone was 50% of the total area. Prior to testing, mice were habituated to the test room for at least 20 minutes. Mice were placed in the center of the box at the start of the assay. Movement was recorded using a measurement (Med Associates, St. Albans, VT, ENV-510) 1 hour in 5 mins bins. In addition to regular parameters related to locomotor activity (such as total travel distance, velocity, ambulatory time, resting time), time spent, and distance travelled in the center area of the testing arena were also recorded and analyzed.

**Elevated plus maze (EPM).** EPM was used to measure anxiety effect. Before EPM test, mice were brought to the testing room for environmental habituation for at least 30 min. The EPM apparatus is consisted of an elevated platform (80 cm above the floor), with four arms (each arm is 30 cm in length and 5 cm in width), two opposing closed arms with 14 cm walls and two opposing open arms. Mice were attached into the fiber-optic patch cord and were individually placed in the center of the EPM apparatus, towards one of the open arms. The mice trajectories were tracked for 5 minutes, and the time spent in the open arms was analyzed using Ethovision XT11 (Noldus).

**Cocaine conditioned place preference (cocaine-CPP).** Mice were allowed to freely explore both sides of a custom-made (Med Associates) CPP training apparatus (25 × 19 × 17 cm for L × D × H) for 30 min. Trajectories were tracked by infra-red photobeam detectors, and the travel distance and the duration were recorded to assess their baseline place preference. Mice that showed strong bias (< 25% preference) were excluded from the experiments. Then, for

chemogenetic activation or inhibition during CPP formation, these mice were injected with saline (i.p.) and confined to their preferred side of the chamber for 30 min before returned to their home cage. At least four hours later, the same mice received CNO at least 15 min before an i.p. injection of 15 mg/Kg cocaine and were confined to their non-preferred side of the chamber for 30 minutes. They were then returned to their home cage. The same training with saline and CNO injection were performed for three consecutive days. Twenty-four hours after the final training session, mice were re-exposed to the CPP chamber and allowed to explore both sides of the chamber for 30 min.

**Post-fasted food intake.** Mice were individually placed in the home cage and fasted overnight (18 hours). Mice received N-clozapine (CNO, 2mg/mL for hM3Dq group and 5mg/mL for hM4Di group) via i.p. injection and then regular chow pallets (3 g per pellet) were put in the hopper. Three hours later, the remaining food pallets were collected and measured to calculate total amount of food consumed (g). For the leptin treatment test, 15 mins after CNO injection, 1 μg of leptin (R&D, Cat# 498-OB) was delivered through cannula by pump for 5 mins and waited for another 5 mins before adding pellets. For the Ta-Casp3 treatment group, the test was carried out 3 weeks after virus injection.

**Food place preference.** Animals were placed in a custom three-chamber, 45 × 60 × 35 cm arena to assess the amount of food consumed and time spent in a designated food zone area. The arena contained two 64-$cm^2$ food cups in two outer corners of separate chambers. One cup contained standard grain-based chow (Harlan, Indianapolis, IN), while the other cup remained empty. Mice were allowed to explore the arena freely, and spatial locations were tracked using EthoVision XT 10 (Noldus, Leesburg, VA) and CCD cameras (SuperCircuits, Austin, TX).

**Operant behavior.** Animals were first given access to 20 mg sweetened chow pellets in their home cage before testing. Animals were then trained to enter the chamber to retrieve a pellet. Each pellet was delivered 10 s after the prior pellet retrieval. After at least 2 days training and until >30 pellets earned in a single session, animals were trained for the fixed ration 1 (FR1) task, in which each active lever pressing was rewarded with a pellet. A new trial does not begin until animals entered the magazine to retrieve the pellet. Retrieval was followed by a 5 s intertrial

interval, after which the levers were reactivated, indicated by a cue light. Training continued until >40 pellets were earned in a single 60 mins session.

**Progressive ratio**. After FR1, FR3, FR5 training sessions, all mice were tested with CNO treatment. For progressive ratio (PR) task, a schedule of reinforcement, each subsequent reward required exponentially more lever pressing based on the formula $(5 \times e^{0.2n}) - 5$, rounded to the nearest integer, where $n$ = number of rewards earned[72]. 60 mins per session.

**Optogenetic modulations of post-fasted food intake**

Mice received 20 min laser stimulation (4 × 5 min, On-Off-On-Off), and then the remaining food pallets were collected and food intake was measured. For photostimulating ChR2, a 473-nm laser (OEM Lasers/OptoEngine) was used to generate laser pulses (10-15 mW at the tip of the fiber, 5 ms, 20 Hz) throughout the behavioral session, except when noted otherwise, controlled by a waveform generator (Keysight). For NpHR photostimulation, a 532-nm laser (OEM Lasers/OptoEngine) generated constant light of 8-10 mW power at each fiber tip.

**Fiber photometry during feeding**

The *Serpinb2*$^{+}$ neuronal dynamics during feeding was measured using fiber photometry. Following injection of an AAV1-hSyn-FLEX-GCaMP7s vector into NAcSh of *Serpinb2*-Cre mice, an optical cannula (Ø200 μm core, 0.37 numerical aperture) was implanted 100 μm above the viral injection site. Mice were allowed to recover for 3 weeks and then subjected to behavioral test. GCaMP fluorochrome was excited, and emission fluorescence was acquired with the RZ10X fiber photometry system, which has built-in LED drivers, LEDs, and photosensors (Tucker-Davis Technologies). The LEDs include 405 nm (as isosbestic control) and 465 nm (for GCaMP excitation). Emitted light was received through the Mini Cube (Doric Lenses) and split into two bands, 420 to 450 nm (autofluorescence) and 500 to 550 nm (GCaMP7 signal). Mice with optical cannula were attached to recording optic cables, and the LED power at the tip of the optical cables was adjusted to the lowest possible (~20 μW) to minimize bleaching. Mice were placed in the 3- chamber for food preference and food consumption test. Mice behaviors were recorded using EthoVision XT 10 (Noldus, Leesburg, VA) and CCD cameras (Super Circuits, Austin, TX).

For the 3-chamber food seeking and food consumption, mice were fasted overnight, and were then habituated for 10-min in the chamber that contain 2 empty food cups, after that, we put a non-eatable object and food pellets in the two food cups, during this phase, food pellets are caged so that mice can sense the food but are unable to eat them. After 10-min recording, we then removed the barrier of food and mice have free-access to the food, mice eating events were then recorded. Behavioral events, such as baseline of free moving, entering food zone, food consumption and post eating were scored manually and synchronized with fluorescence signal based on recorded videos. The voltage signal data stream was acquired with Synapse software (Tucker-Davis Technologies) and were exported, filtered, and analyzed with custom-written Matlab code. To calculate ΔF/F, a polynomial linear fitting was applied to isosbestic signal to align it to the GCaMP7 signal, producing a fitted isosbestic signal that was used to normalize the GCaMP7 as follows: ΔF/F = (GCaMP7signal − fitted isosbestic)/fitted isosbestic signal.

**Lead Contact**

Further information and requests for reagents should be directed to and will be fulfilled by the Lead Contact, Yi Zhang (yzhang@genetics.med.harvard.edu).

**Data and Code Availability**

The custom code that supports the findings from this study are available from the Lead Contact upon request.

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### Contributions

Y.Z. conceived the project; Y.L., and Y.Z. designed the experiments; Y.L. performed most of the experiments. Z.-D.Z. helped with the fiber photometry. G.X. helped with the catheter administration. R.C. initiated the Serpinb2-Cre mouse generation. Y.L., Z.-D.Z., R.C. and Y.Z. interpreted the data; Y.L. and Y.Z. wrote the manuscript with input from Z.-D.Z and R. C.

### Ethics declarations

### Competing interests

The authors declare no competing interests.

802 **Figure Legends**

803

804 **Fig. 1: NAcSh *Serpinb2*$^+$ neurons and NAc *Drd1*$^+$-MSNs are activated in refeeding process**

805 **a,** Inferred spatial expression patterns from MERFISH database of *Tac2*, *Serpinb2* and *Ubp1*
806 whose expression is highly enriched in medial dorsal NAc shell. Expression level is color-
807 coded. Doted line circle anterior commissure olfactory limb (aco) and NAc core. The dorsal-
808 ventral (DV) and medial-lateral (ML) axes are indicated.

809 **b,** Left, tSNE plot showing the 8 NAc D1-MSN subtypes. Middle and right panels indicate the
810 expression of *Tac2* and *Serpinb2* in the NAc D1-MSNs, respectively. Expression level is
811 color-coded.

812 **c,** RNA *in situ* hybridization showing *Tac2* and *Seprinb2* expression in the medial part of the
813 NAc shell. Scale bar: 500 μm (left), 20 μm (right).

814 **d,** Schematic representation of the experimental design for the ad libitum, fast and refed groups.

815 **e,** Coexpression of *Serpinb2* mRNA with *cFos* mRNA in NAc of ad libitum, fasted and refed
816 states. Representative images showing the colocalization of *cFos* (green), *Serpinb2* (red) and
817 *Tac2* (magenta) expressing neurons. Scale bar: 50 μm.

818 **f,** The average number of *cFos*$^+$ neurons in the dorsal medial NAcSh at ad libitum, fasted and
819 refed states. (n = 5 sections from three mice of each group, one-way ANOVA, with Tukey's
820 multiple comparisons).

821 **g,** The percentages of activated *Serpinb2*-expressing neurons in the total *Serpinb2* neurons under
822 different feeding states (n = 5 sections from three mice of each group, one-way ANOVA,
823 with Tukey's multiple comparisons).

824 **h,** The percentages of activated *Tac2*-expressing neurons in the total *Tac2* neurons under
825 different feeding states (n = 5 sections from three mice of each group, one-way ANOVA,
826 with Tukey's multiple comparisons). All error bars represent mean ± SEM. ns, not significant,
827 *P < 0.05.

828

829 **Fig. 2: The activity of *Serpinb2*$^+$ neurons respond to feeding states**

830 **a,** Left, illustration of light pathways of fiber photometry. Right, schematic illustration of the
831 GCaMP7s injection and optic cannula implantation.

**b, c,** Validation of GCaMP7s expression and implantation of optic cannula in *Serpinb2*-Cre mice (b) or *Drd1*-Cre mice (c) (left). Representative trace of real-time monitoring of *Serpinb2*$^+$ neurons (b) or *Drd1*$^+$ neurons (c) (right) during feeding process. Scale bar: 200 μm.

**d,** $Ca^{2+}$ signals at different phases of feeding process of fasted mice (top). Average $Ca^{2+}$ signal at different feeding phases of the *Serpinb2*-Cre mice. Elevated $Ca^{2+}$ signals were observed after entering food zone in approaching and eating phases. Declined $Ca^{2+}$ signals were observed post eating. Quantification of AUC in the four phases is shown in bar graph (right, n=7).

**e,** The same as in panel D except *Drd1*-Cre mice were used. ***P ≤ 0.001, **P ≤ 0.01, *P ≤ 0.05; ns, P > 0.05, one-way ANOVA test. Data are represented as mean ± SEM.

**Fig. 3: *Serpinb2*$^+$ neurons bidirectionally regulate food seek and intake in hungry state**

**a,** Experimental scheme of the food consumption and food preference assays.

**b,** Total food consumption during the 3-hour test. Chemogenetic activation (hM3Dq) or chemogenetic inhibition (hM4Di) of *Serpinb2*$^+$ neurons at ad libitum (left) or fasting state (right). **, p<0.01; ns, p>0.05, unpaired t-test.

**c,** The same as in panel B except the *Tac2*-Cre (left) or *Drd1*-Cre (right) mice were used. ns, p>0.05, unpaired t-test.

**d,** Color map encoding spatial location of a fasted mouse using the free access feeding paradigm.

**e,** Percentage of time that mice spent in food zone. Chemogenetic activation (hM3Dq) or chemogenetic inhibition (hM4Di) of *Serpinb2*$^+$ neurons (left), *Tac2*$^+$ neurons (middle), and *Drd1*$^+$ neurons (right). *, p<0.05; ns, p>0.05, unpaired t-test.

**f,** Experimental timeline of the food operant chamber assay.

**g,** Diagrammatic illustration of the food operant chamber paradigm. Mice were trained to press the lever to get food; pressing the active lever is followed by the delivery of food pellet, while pressing the inactive lever yields no outcome. The behavioral training includes habituation phase and fixed ratio (FR) training phase. Mice received CNO injection (2 mg/kg for the hM3Dq group and 5 mg/kg for the hM4Di group) 15 mins before they were placed into the operant chamber to start the FR and progressive ratio (PR) tests.

**h,** Results of FR=5 test. The total number of active lever pressing (left) and total number of reward (right) after chemogenetic manipulations. **, p<0.01; *, p<0.05; ns, p>0.05; one-way ANOVA test.

**i,** Results of PR=5 test. The total number of active lever pressing (left) and total number of reward (right) after chemogenetic manipulations. *, $p<0.05$; ns, $p>0.05$; one-way ANOVA test.

Data are represented as mean ± SEM.

**Fig. 4: *Serpinb2*$^+$ neurons mediate food intake via LH projection**

**a,** Diagram of antegrade tracing of the NAc *Serpinb2*$^+$ neurons with ChR2-EYFP (left), the expression of DIO-ChR2-EYFP in the NAcSh (2$^{nd}$ panel), and *Serpinb2*$^+$ neuron projection to LH (the right 3 panels). Scale bar: 100 μm.

**b,** Diagram (left) and image (right) of retrograde tracing of LH neurons with CTB 647, with the *Serpinb2*$^+$ neurons labeled by DIO-mCherry. Scale bar: 50 μm.

**c,** Percentage of *Serpinb2*$^{+\ LH}$ projecting neurons over the total mCherry labeled *Serpinb2*$^+$ neurons.

**d,** Diagram illustrate the indicated AAV injection into NAc and optic cannulas implantation in LH area (left), and histology validating virus expression and cannula implantation site (right). Scale bar: 200 μm.

**e,** Optogenetic activation (left) or inhibition (right) of *Serpinb2*$^+$ NAc→LH neurons respectively increased or decreased food intake.

Data in (e) is presented as mean ± SEM. TH, thalamus, CP, caudoputamen, AHNc: anterior hypothalamic nucleus, central part, HPF, hippocampal formation, VL, lateral ventricle. ***P ≤ 0.001; *P ≤ 0.05; ns, P > 0.05, unpaired t-test.

**Fig. 5: *Serpinb2*$^+$ neurons project to LH LepR$^+$ GABA$^+$ neurons and receive input related to energy homeostasis**

**a,** Diagram indicating the MCH$^+$, Orexin-A$^+$, GABA$^+$ and LepR$^+$ neurons in LH.

**b,** Images (left two panels) showing colocalization of *Serpinb2*$^+$ neuron terminals (green) with MCH$^+$, Orexin-A$^+$ neurons in LH, and their quantifications (right panel). Scale bar, 50 μm. Percentage = eYFP$^+$MCH$^+$/eYFP$^+$DAPI$^+$ or eYFP$^+$Orexin-A$^+$/eYFP$^+$DAPI$^+$

**c,** Images showing the colocalization of *Serpinb2*$^+$ neuron terminals (green) with GABA$^+$ (red) or LepR$^+$ neurons (red) as indicated, as well as their quantifications (right panel). Scale bar, 50 μm. Percentage = eYFP$^+$GABA$^+$/eYFP$^+$DAPI$^+$ or eYFP$^+$LepR$^+$/eYFP$^+$DAPI$^+$

894

895 **d,** Quantification of cFos$^+$ cells of different brain regions in ad libitum and refeed status. (n = 3
896 sections from three mice of each group, unpaired t-test)
897 **e,** Schematic presentation of modified rabies tracing (left) and representative image confirming
898 the expression of the indicated proteins at the injection site (right); scale bar, 100 μm
899 (left); 20μm (right).
900 **f,** Representative images showing the brain areas with positive signals indicating these regions
901 have neurons projecting to *Serpinb2*$^+$ neurons. Scale bar: 100 μm, arrow heads indicate
902 neurons.
903 **g,** Diagram illustrating brain regions upstream and downstream of *Serpinb2*$^+$ neurons.
904 ACAd: Anterior cingulate area, dorsal part; fa, corpus callosum, anterior forceps; LPO: lateral
905 preoptic area; SI: substantia innominata; PVT: paraventricular nucleus of the thalamus; VMH:
906 ventromedial hypothalamic nucleus; LH: lateral hypothalamus; V3: third ventricle.
907
908 **Fig. 6: Modulating *Serpinb2*$^+$ neuron activity can overcome leptin effect and alter**
909 **bodyweight**
910 **a,** Diagram showing bilateral cannula implantation in LH for leptin delivery (left). CNO delivery
911 was achieved via i.p. injection. Results of total food consumption in 3 hours by fasted mice
912 with different doses of leptin administration. Scale bar: 500 μm.
913 **b,** Same as panel A except food consumption is quantified under different conditions with or
914 without *Serpinb2*$^+$ neuron activation in the presence or absence of 1 μg of leptin delivery.
915 ***P ≤ 0.001, **P ≤ 0.01, *P ≤ 0.05; ns, P > 0.05, unpaired t-test.
916 **c,** FISH and quantification verify *Serpinb2*$^+$ neuron ablation after AAV-DIO-taCasp3 injection.
917 Scale bar: 100 μm. ***P ≤ 0.001, unpaired t-test.
918 **d,** *Serpinb2*$^+$ neuron ablation decreased 3 hours total food consumption by mice. ***P ≤ 0.001,
919 unpaired t-test.
920 **e,** *Serpinb2*$^+$ neuron ablation has a long-time effect on bodyweight loss. ***P ≤ 0.001, **P ≤
921 0.01, *P ≤ 0.05; ns, P > 0.05, unpaired t-test.
922 Data are presented as means ± SEM.

## Extended Data Figure Legends

### Extended Data Fig. 1: Generation and validation of *Serpinb2*-Cre line

**a,** Diagrams showing the targeting strategy.

**b,** Genotyping by PCR. Homozygotes: 413 bp. Heterozygotes: 413 bp/772 bp.

**c,** Left, in situ hybridization (ISH) data of *Serpinb2* from Allen Brain Atlas. Scale bar: 1000 μm. Middle, colocalization of *Serpinb2* RNA (red), AAV-DIO-ChR2-eYFP (green) and DAPI (blue). Scale bar: 50 μm. Right, quantification of $Serpinb2^+$ and $eYFP^+$ neurons among all $Serpinb2^+$ neurons.

### Extended Data Fig. 2: $Serpinb2^+$ neuron activity during different phases of feeding in ad libitum mice

**a,** $Ca^{2+}$ signals at different phases of feeding process of ad libitum mice (top). Average $Ca^{2+}$ signal at different feeding phases of the *Serpinb2*-Cre mice. Elevated $Ca^{2+}$ signals were observed after entering food zone in the approaching and eating phases. Declined $Ca^{2+}$ signals were observed post eating. Quantification of AUC in the four phases is shown in bar graph (right, n=7 mice).

**b,** The same as in panel D except *Drd1*-Cre mice were used.

$^{**}P \leq 0.01$, $^{*}P \leq 0.05$; ns, $P > 0.05$, one-way ANOVA test. Data are represented as mean ± SEM.

### Extended Data Fig. 3: Validation of chemogenetic manipulation

**a,** Experimental scheme of chemogenetic manipulation.

**b,** Validation of virus expression in *Serpinb2*-Cre, *Tac2*-Cre and *Drd1*-Cre mice. Scale bar: 500 μm.

**c,** *cFos* induction after intraperitoneal injection of ligand CNO in mCherry-expressing, hM3Dq-mCherry-expressing and hM4Di-mCherry-expressing mice. The ratio of $cFos^+/mCherry^+$ cells in all $mCherry^+$ cells was calculated and shown on the right panel. Scale bar: 100 μm. *, $p<0.05$; **, $p<0.01$, one-way ANOVA test. Data are represented as mean ± SEM.

### Extended Data Fig. 4: $Serpinb2^+$ neuronal activity does not affect anxiety or drug seeking behavior

**a, b,** Open field test for the effect of *Sepinb2*$^+$ neuronal activation (hM3Dq) (a) or inhibition (hM4Di) (b) on the total distance traveled in the 1-hour post-treatment period after chemogenetic manipulation of *Sepinb2*$^+$ neurons (left) or the distance traveled in 5-min time bin (right).
ns, p>0.05, left, unpaired t-test; right, two-way ANOVA.

**c,** Left, illustration of the two-chamber cocaine-CPP paradigm. Right, cocaine-CPP with chemogenetic activation (hM3Dq) or inhibition (hM4Di) of *Sepinb2*$^+$ neurons. CPP scores were calculated by subtracting the time spent in the preconditioning phase from the time spent in the postconditioning phase. ns, p>0.05, unpaired t-test.

**d, e,** Elevated plus maze test for the effect of *Sepinb2*$^+$ neuronal activation (hM3Dq) (d) or inhibition (hM4Di) (e) on the time spent (left) or distance traveled (right) in open arm and closed arm of the 5-min post-treatment period after chemogenetic manipulation of *Sepinb2*$^+$ neurons. ns, p>0.05, unpaired t-test.

Data are represented as mean ± SEM.

**Extended Data Fig. 5: cFos staining of different brain regions from mice under Ad libitum and refeed states**

Shown are cFos FISH in different brain regions listed below:
ACB: Nucleus accumbens; aco: anterior commissure, olfactory limb; OT:
Olfactory tubercle; V3: third ventricle; ARH: Arcuate hypothalamic nucleus; AHNp: Anterior hypothalamic nucleus, posterior part; VMH: ventromedial hypothalamic nucleus; fx: columns of the fornix; LH: lateral hypothalamus; PVT: paraventricular nucleus of the thalamus; sm: stria medullaris; BLA: Basolateral amygdala; LA: Lateral amygdala; RE: Nucleus of reuniens; DMH: Dorsomedial nucleus of the hypothalamus; ZI: Zona incerta. Scale bar: 100 μm.

**Extended Data Fig. 6: Brain regions that do not innervate *Serpinb2*$^+$ neurons**

Shown are immunostaining of the various brain regions listed below:
PL: Prelimbic area; IL: Infralimibic area; fa: corpus callosum, anterior forceps; ccg: genu of corpus
callosum; LSr: Lateral septal nucleus, rostral part; VL: lateral ventricle; ACB: Nucleus accumbens; aco: anterior commissure, olfactory limb; OT: Olfactory tubercle; ADP:

985 Anterodorsal preoptic nucleus; BST: Bed nuclei of the stria terminalis; V3: third ventricle; ARH:
986 Arcuate hypothalamic nucleus; AHNp: Anterior hypothalamic nucleus, posterior part; VMH:
987 ventromedial hypothalamic nucleus; fx: columns of the fornix; LH: lateral hypothalamus; PVT:
988 paraventricular nucleus of the thalamus; sm: stria medullaris; BLA: Basolateral amygdala; LA:
989 Lateral amygdala; RE: Nucleus of reuniens; DMH: Dorsomedial nucleus of the hypothalamus;
990 ZI: Zona incerta; DG: Dentate gyrus; CA1: field CA1; CA3: field CA3; PAG: Periaqueductal
991 gray ; APN: Anterior pretectal nucleus; HPF: Hippocampal formation. Scale bar: 100 μm

**Fig. 1: NAcSh *Serpinb2*$^{+}$ neurons and NAc *Drd1*$^{+}$-MSNs are activated in refeeding process.**

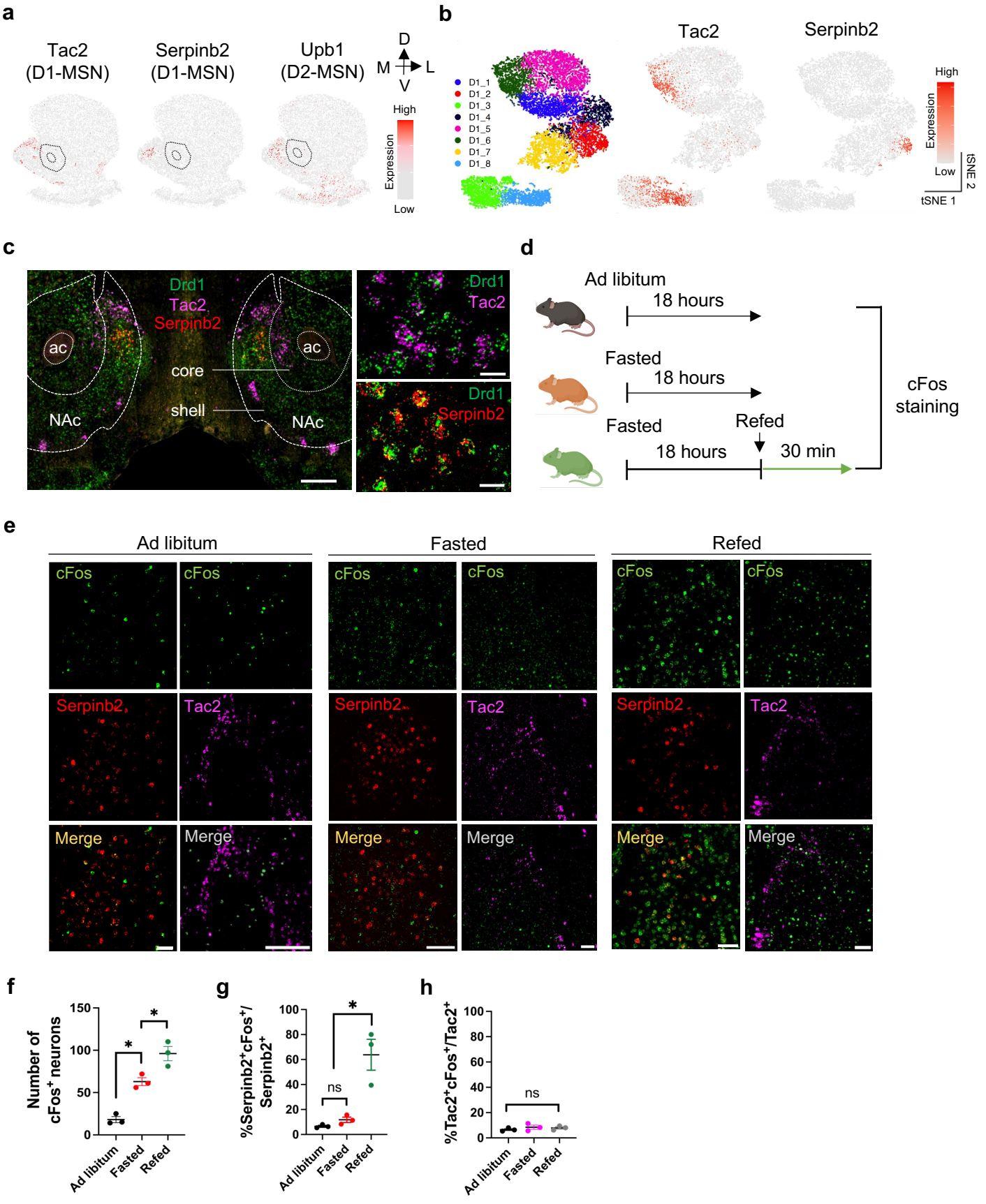


**Fig. 2: The activity of *Serpinb2*$^{+}$ neurons respond to feeding states.**

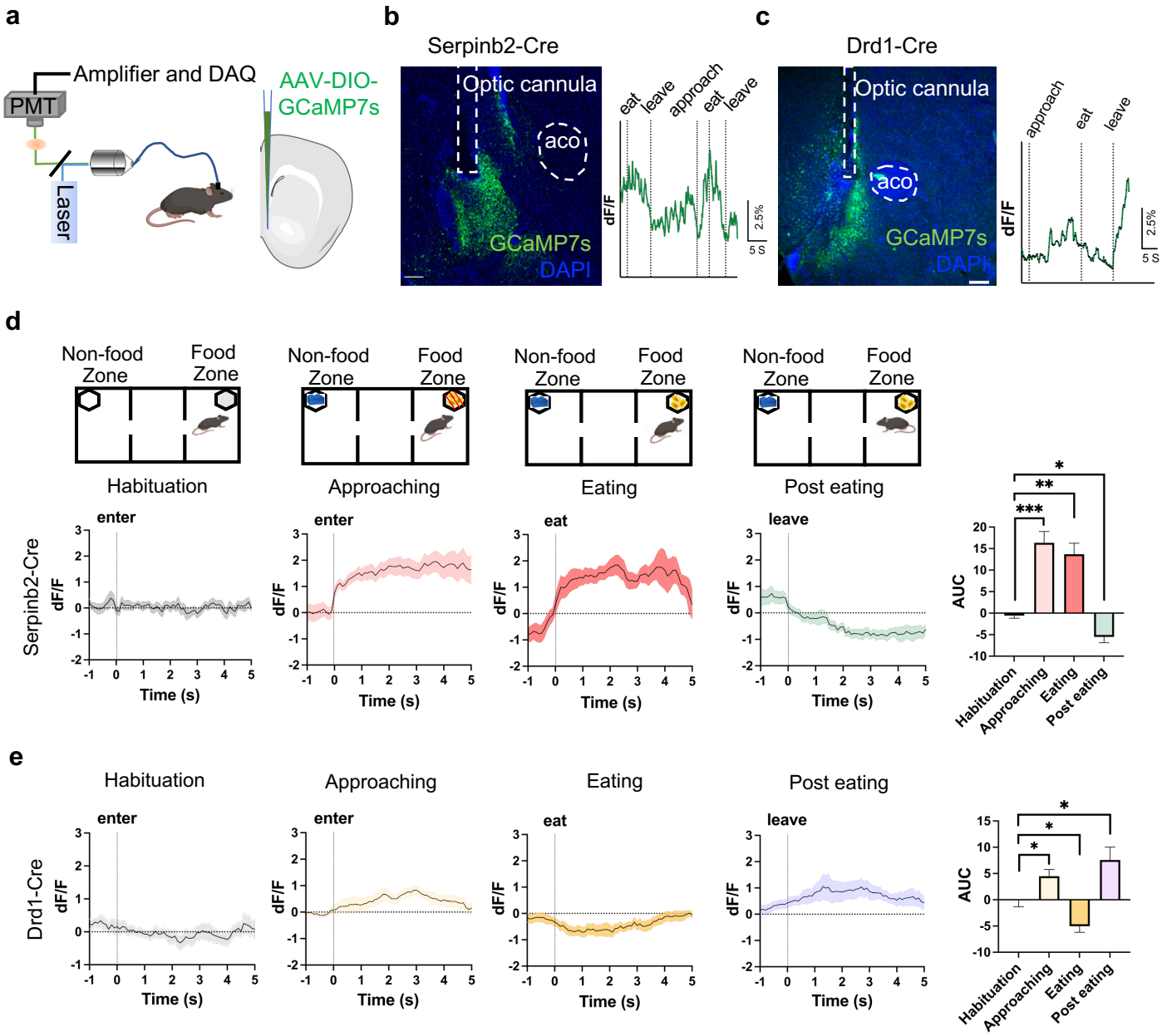


**Fig. 3: *Serpinb2*$^{+}$ neurons bidirectionally regulate food seek and intake in hungry state.**

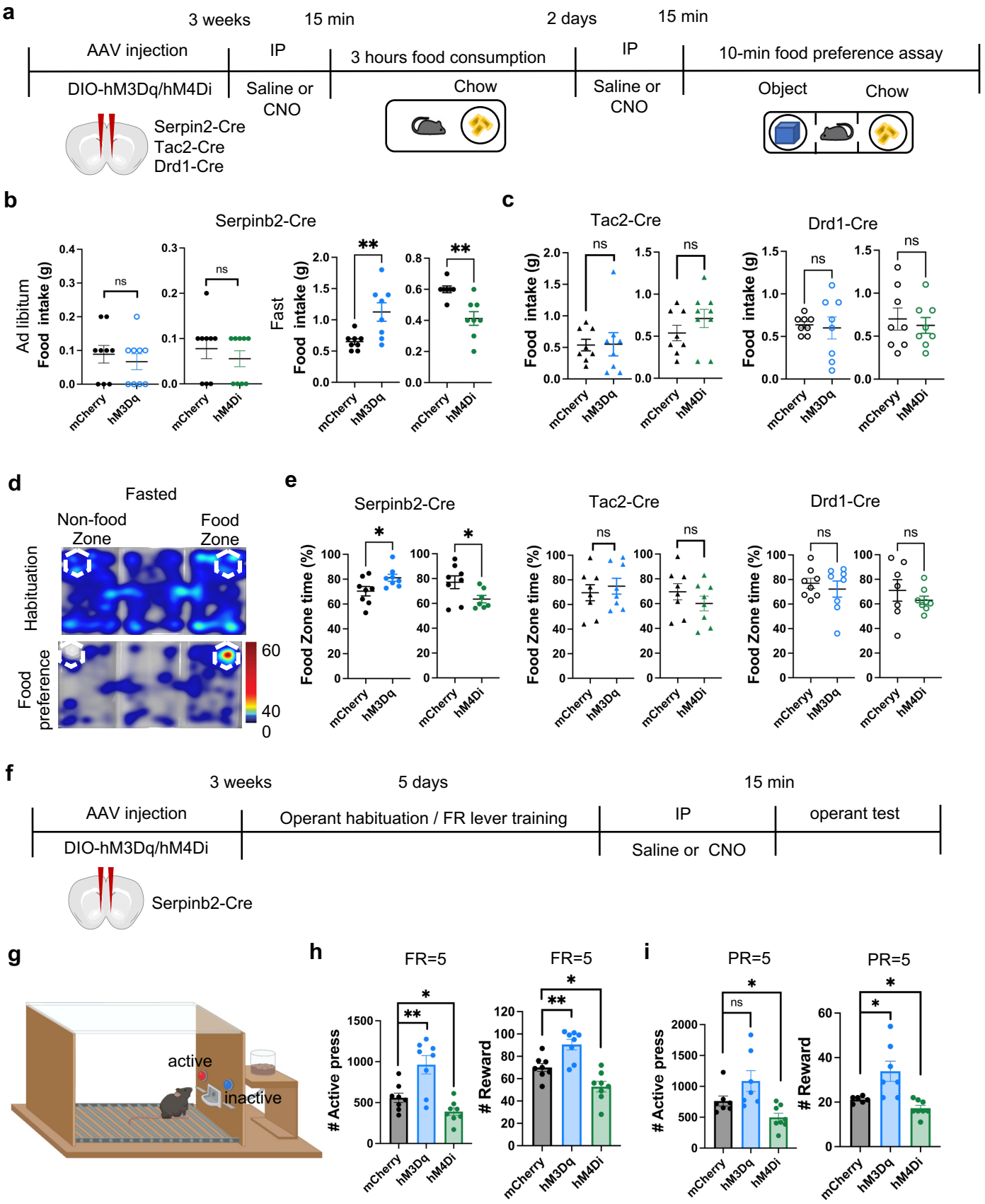


**Fig. 4: *Serpinb2*$^{+}$ neurons mediate food intake via LH projection.**

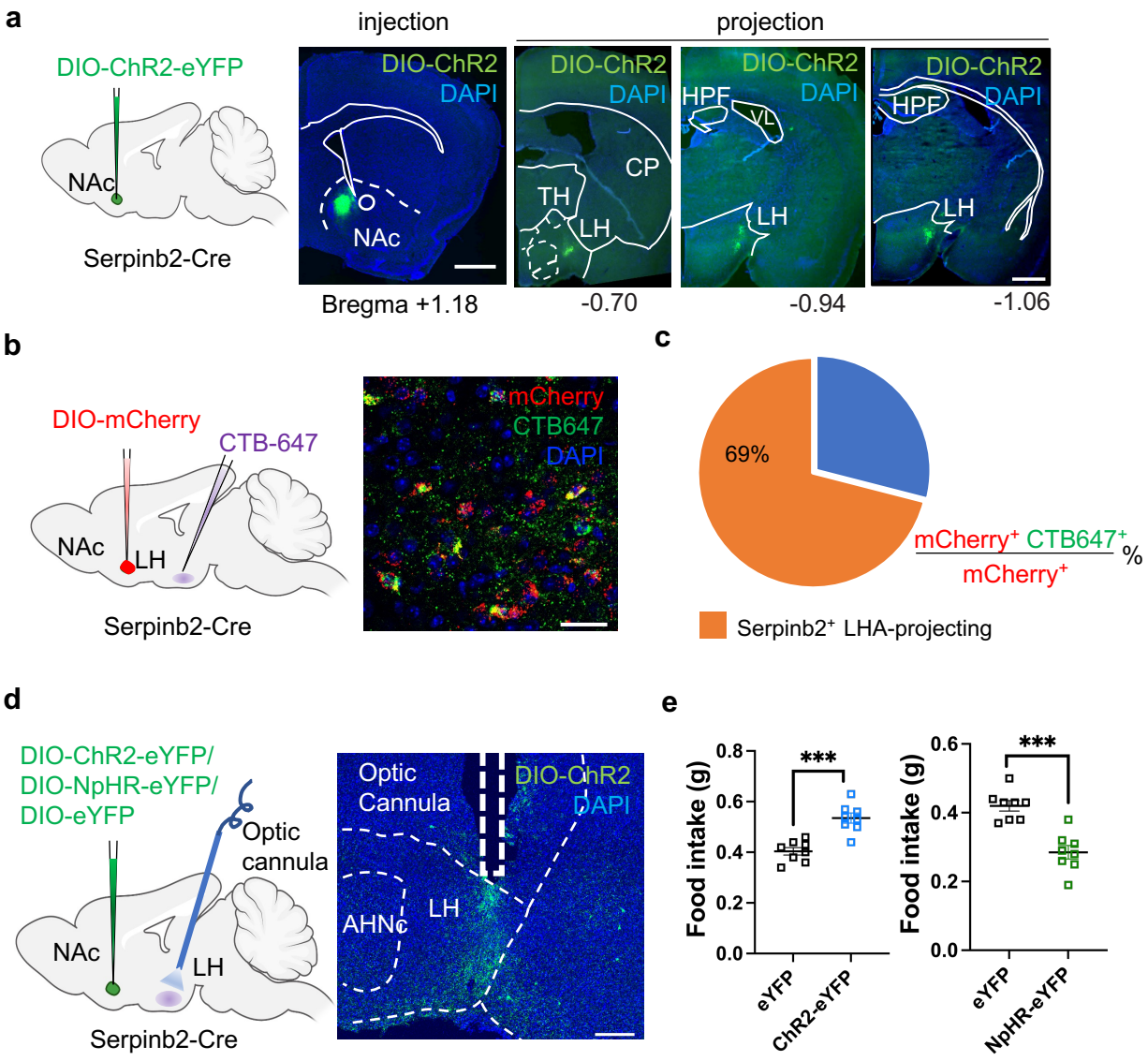


**Fig. 5: *Serpinb2*$^{+}$ neurons project to LH LepR$^{+}$ GABA$^{+}$ neurons and receive input related to energy homeostasis.**

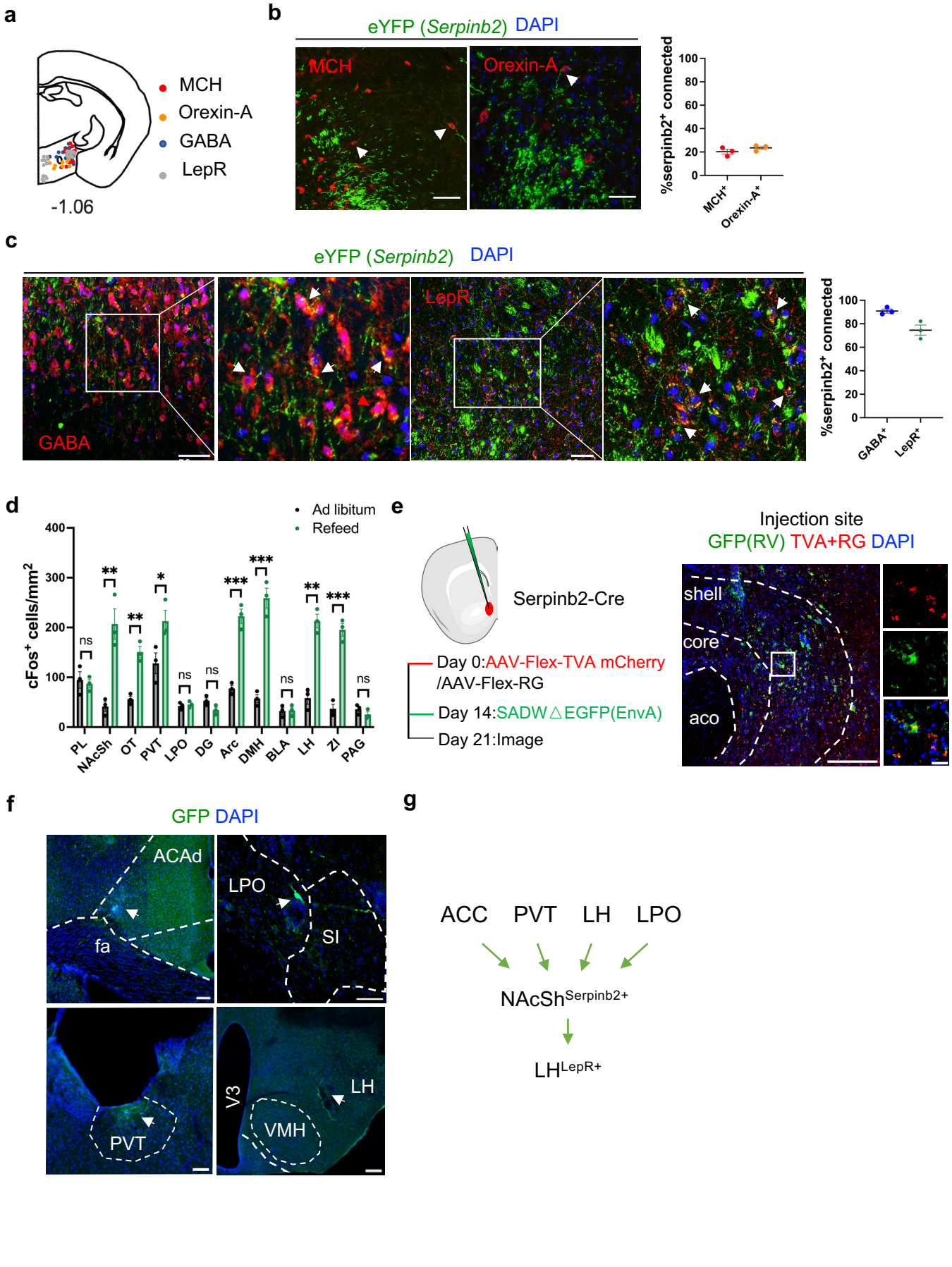
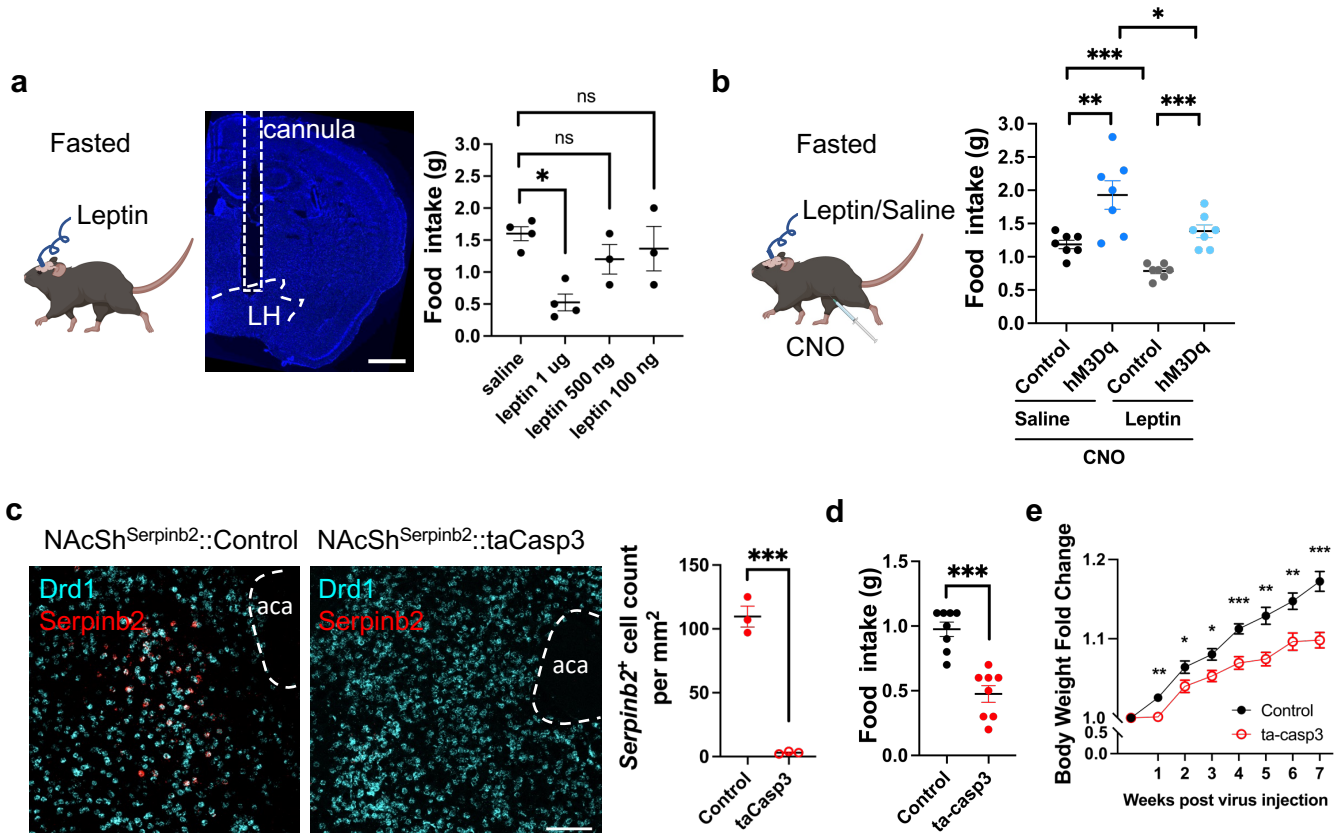
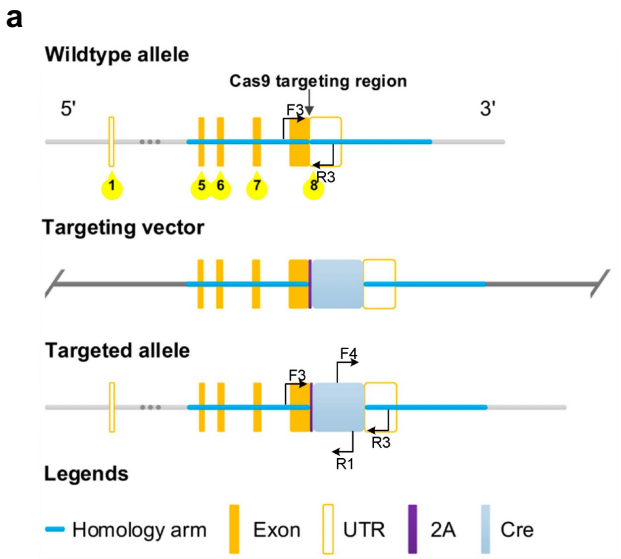
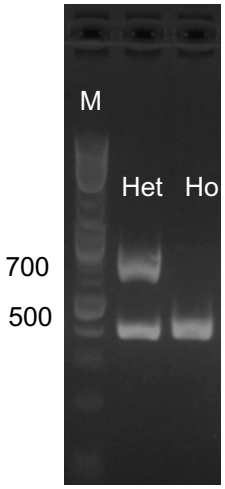


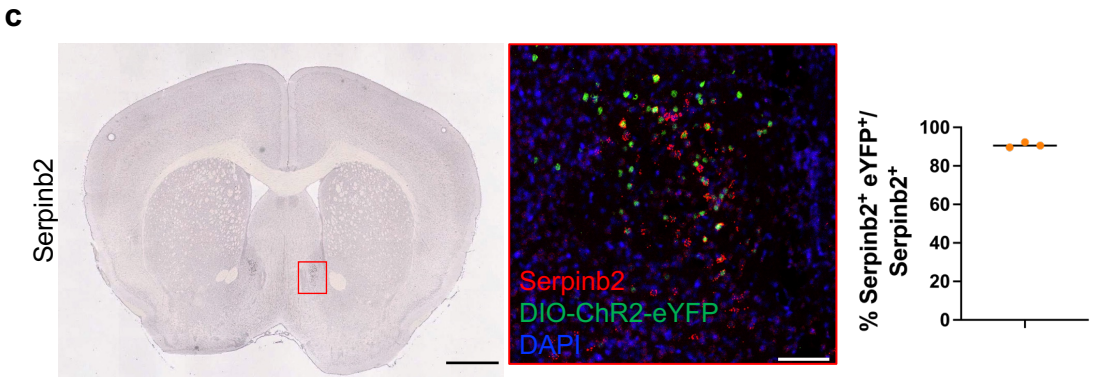
**Fig. 6: Modulating *Serpinb2*$^{+}$ neuron activity can overcome leptin effect and alter bodyweight.**



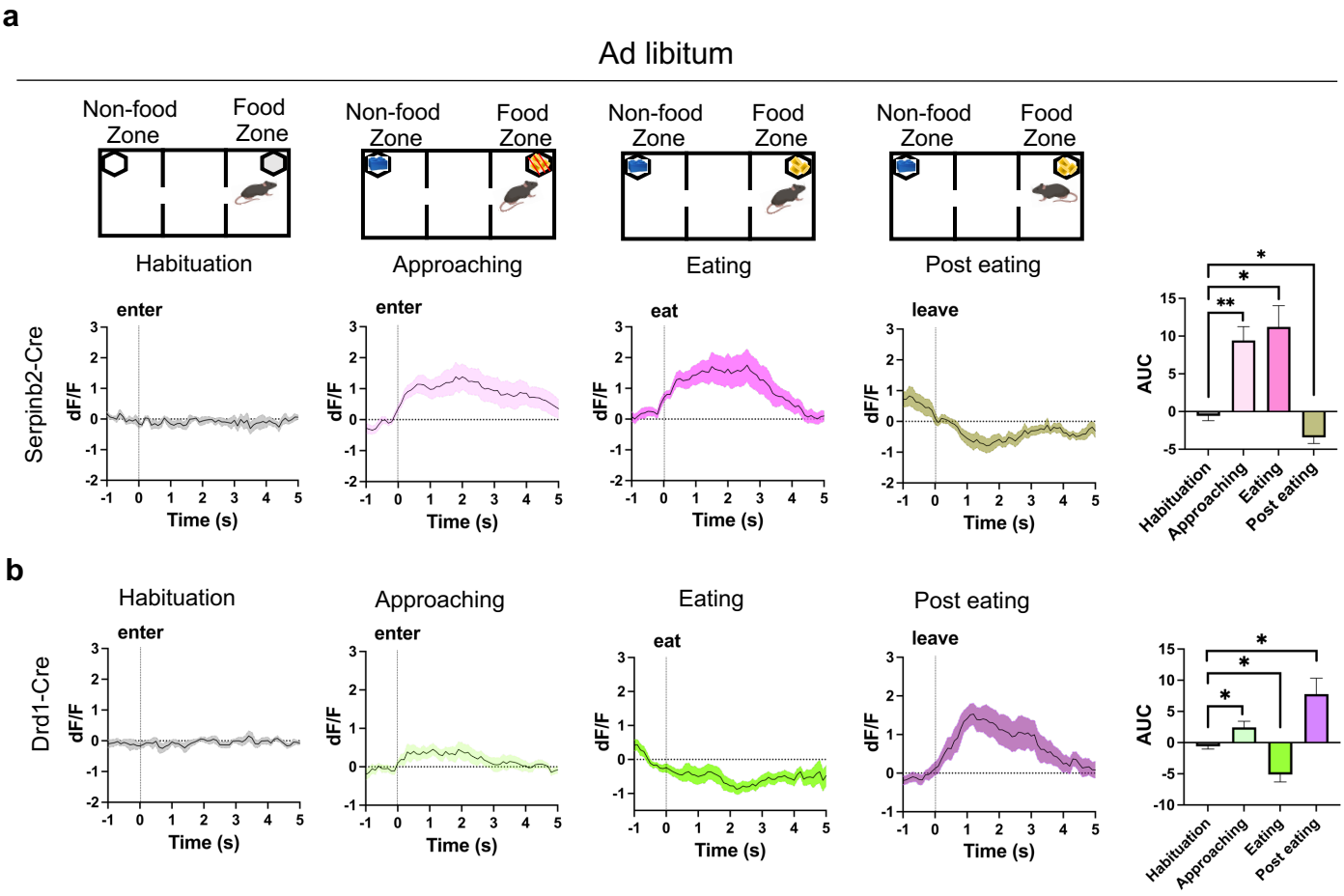
**Extended Data Fig. 1: Generation and validation of *Serpinb2*-Cre line**



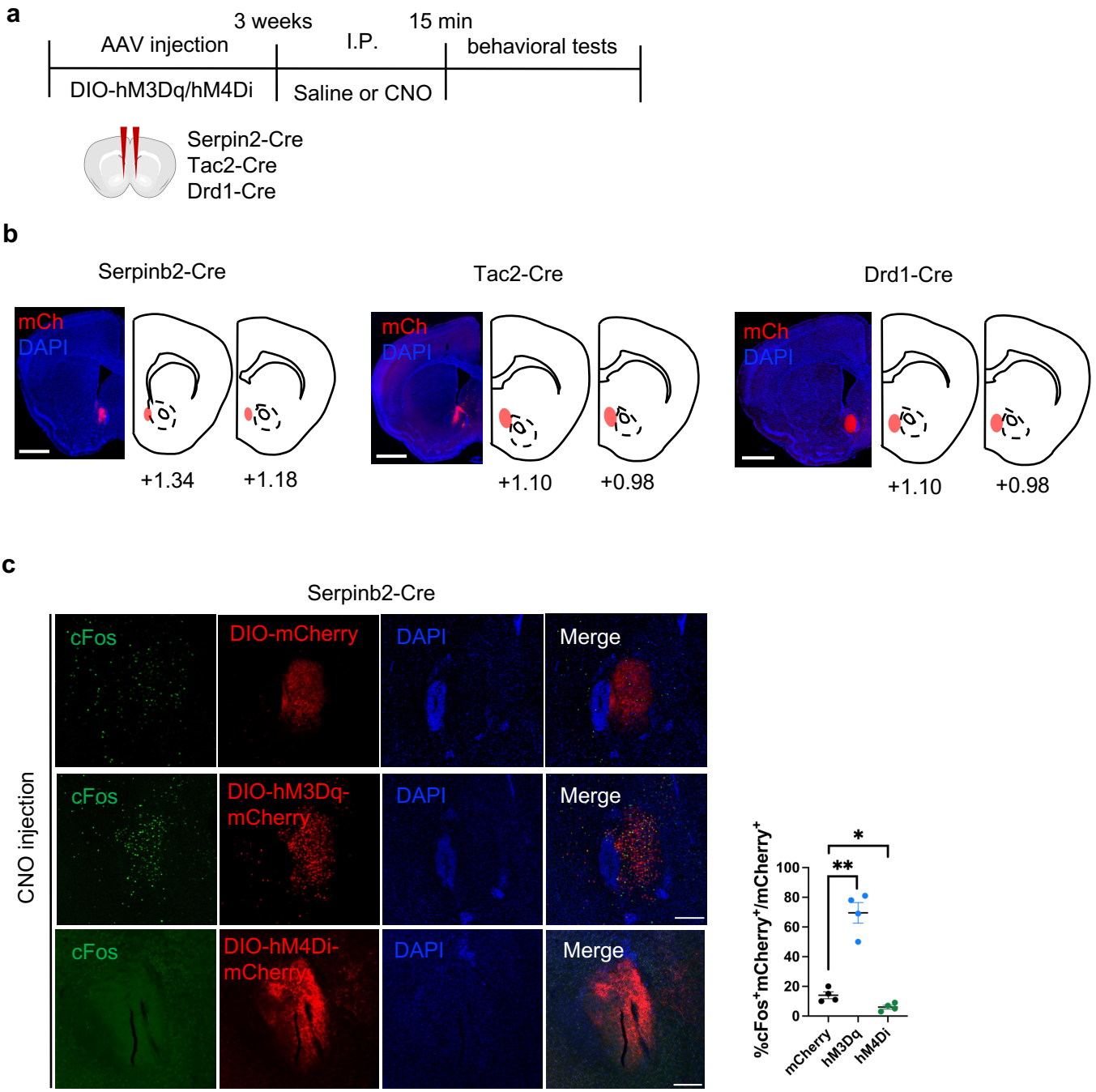



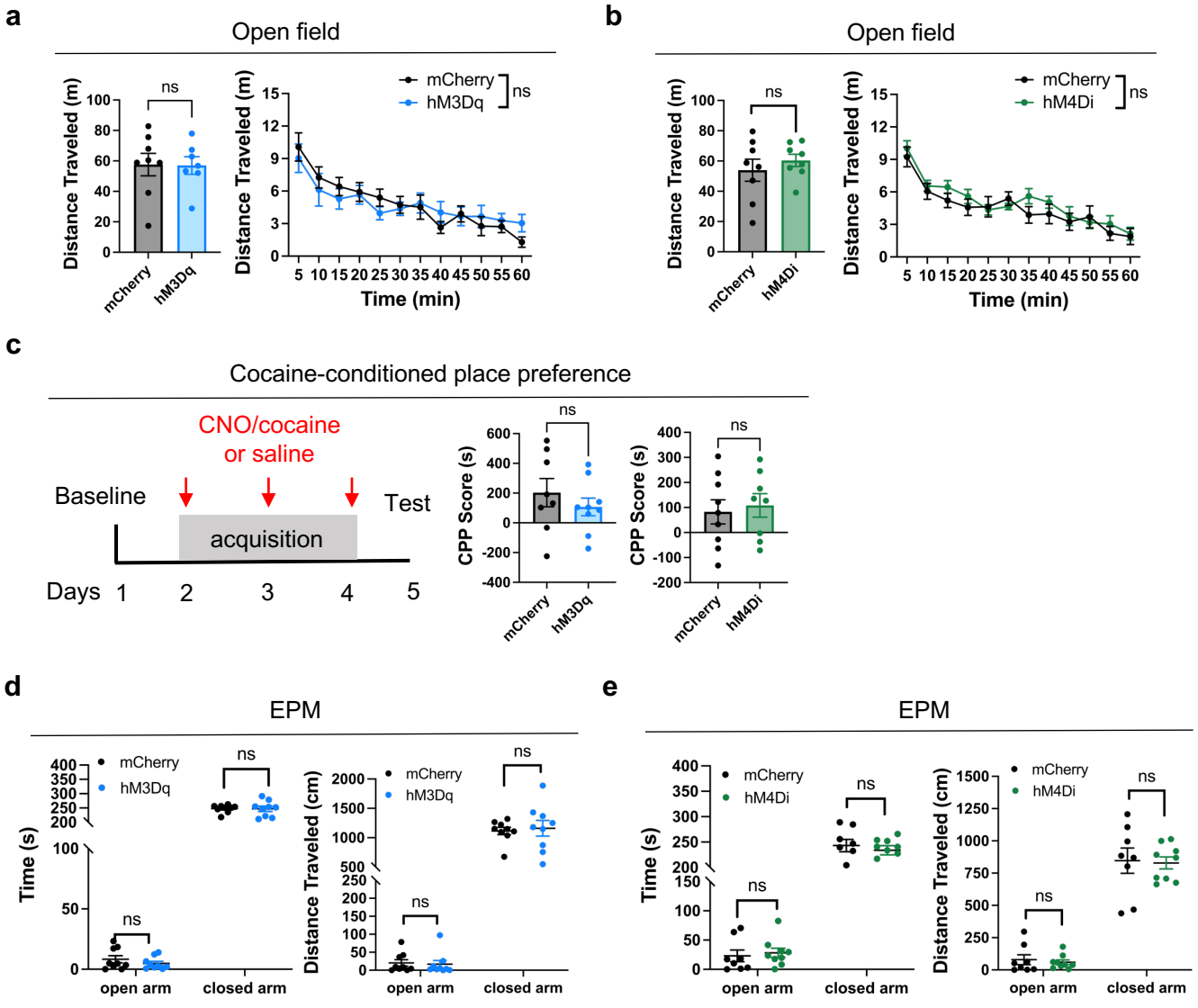
**Extended Data Fig. 2: *Serpinb2*$^{+}$ neuron activity during different phases of feeding in ad libitum mice**



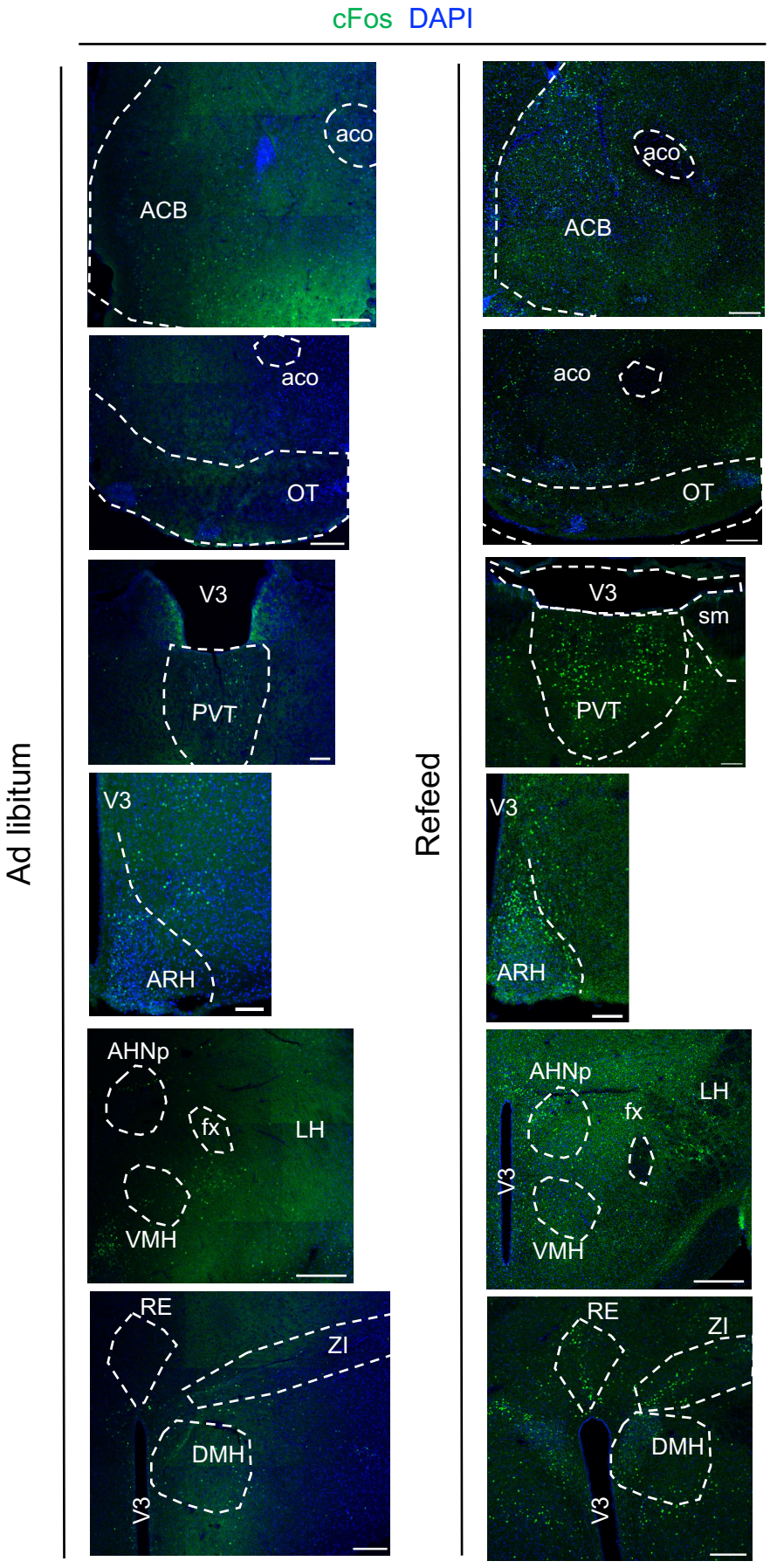
**Extended Data Fig. 3: Validation of chemogenetic manipulation**



**Extended Data Fig. 4: *Serpinb2*$^{+}$ neuronal activity does not affect anxiety or drug seeking behavior**



**Extended Data Fig. 5: cFos staining of different brain regions from mice under Ad libitum and refeed states**



**Extended Data Fig. 6: Brain regions that do not innervate *Serpinb2*$^{+}$ neurons**