A brainstem to hypothalamic arcuate nucleus GABAergic circuit drives feeding

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Highlights

- Nucleus of the solitary tract (NTS) GABA neurons are responsive to nutritional status.

- Chemogenetic GABA^{NTS} neuron activation reduces food intake and body weight.

- GABA^{NTS} projections to the hypothalamic arcuate nucleus (ARC) promote satiety.

- Optogenetic GABA$^{NTS}\rightarrow$ARC stimulation inhibits orexigenic AgRP/NPY neurons.

In Brief

Martinez de Morentin et al. identify GABAergic neurons in the nucleus of the solitary tract as a new player in the circuit governing feeding behavior and body weight.

Abstract

The obesity epidemic is principally driven by the consumption of more calories than the body

requires. It is therefore essential that the mechanisms underpinning feeding behavior are

defined. The brainstem nucleus of the solitary tract (NTS) receives direct information from

the digestive system and projects to second order regions in the brain. Though γ-

Aminobutyric acid is widely expressed in the NTS (GABA^{NTS}), its function has not been

defined. Characterization of GABA cells using single nucleus RNA sequencing (Nuc-Seq)

identified at least 19 clusters. Here we provide insight into the function of GABA^{NTS} cells,

revealing that selective activation of GABA^{NTS} neurons significantly controls food intake and

body weight. Optogenetic interrogation of GABA^{NTS} circuitry identified GABA^{NTS}→arcuate

nucleus of the hypothalamus (ARC) projections as appetite suppressive without creating

aversion. Electrophysiological analysis revealed GABA^{NTS}→ARC stimulation inhibits hunger

promoting agouti-related protein/neuropeptide Y (AgRP/NPY) neurons via GABA release.

Adopting an intersectional genetics strategy, we clarify that the GABA^{NTS}→ARC circuit

induces satiety. These data identify GABA^{NTS} as a new modulator of feeding behavior, body

weight and controller of orexigenic AgRP/NPY activity, thereby providing insight into the

neural underpinnings of obesity.

Keywords

y-Aminobutyric acid, agouti-related -peptide, Neuropeptide Y, nucleus of the solitary tract,

food intake.

Introduction

Obesity represents a key challenge to human health and is primarily due to the consumption of calories in excess to body's energy requirements. Eating is a complex behavior that not only depends on the basic energy demands at a cellular and organism level, but also the integration of internal and environmental cues, the reward value of food, motivation, and conditioning behavior (Andermann & Lowell, 2017; Campos et al., 2022). The aim of the present study was to probe neurocircuitry regulating feeding and body weight with the objective of uncovering critical energy homeostasis circuitry.

One of the primary nodes for the integration of energy-related information from the periphery to the brain is the nucleus of the solitary tract (NTS) within the brainstem dorsal vagal complex (DVC)(Hyun & Sohn, 2022). The NTS contains a heterogeneous population of energy-related sensitive cells(Cheng et al., 2022; Dowsett et al., 2021; Grill & Hayes, 2012). Recent studies using single-nucleus RNA sequencing provide a detailed expression map of identified cellular populations involved in energy balance, and most were revealed to be glutamatergic(Dowsett et al., 2021; Ludwig et al., 2021). For example, subpopulations of these glutamatergic cells influencing energy balance include leptin receptor (LEPR)(Cheng et al., 2020), calcitonin receptor (CALCR)(J. Chen et al., 2020a), glucagon-like peptide 1 receptor (GLP-1R)(Alhadeff et al., 2017; Fortin et al., 2020), preproglucagon (PPG)(Holt et al., 2019), tyrosine hydroxylase (TH) (Aklan et al., 2020; J. Chen et al., 2020b), cholecystokinin (CCK) (D'Agostino et al., 2016) and proopiomelanocortin (POMC) (Georgescu et al., 2020; Zhan et al., 2013). However, very little is known about the role of NTS inhibitory GABA-releasing clusters (GABA^{NTS}) in energy homeostasis. One report in rats indicated a subpopulation of GLP-1R-expressing neurons releasing GABA are necessary mediators for the anorectic effects of the obesity medication liraglutide (Fortin et al., 2020). Therefore, GABA^{NTS} represents and intriguing and understudied population of NTS cells and is the focus of the present study.

In the regulation of feeding behavior, one of the most widely studied projections from the NTS is to the hypothalamus (Aklan et al., 2020; Blevins et al., 2004; D'Agostino et al., 2016; Liu et al., 2017; Shi et al., 2021; Tsang et al., 2020). However, whether the arcuate nucleus of the hypothalamus (ARC) receives GABAergic inhibitory control from the NTS is not known. Within the ARC is a subpopulation of potent orexigenic neurons expressing both agouti-related peptide and neuropeptide Y (AgRP/NPY) (Betley et al., 2013; Hahn et al., 1998;

Heisler et al., 2006; Heisler & Lam, 2017). We hypothesized that GABA[NTS] neurons
significantly regulate feeding and body weight and project to and inhibit key hunger-
stimulating AgRP/NPY cells.

Results

GABA[NTS] cells are sensitive to energy status and modulate food intake

To characterize GABA[NTS] neurons, we initially used a single nucleus RNA sequencing
(NucSeq) dataset of the mouse dorsal vagal complex (Dowsett et al., 2021). Neuronal cells
expressing transcripts for *Slc32a1* (solute carrier family 32 member 1 or vesicular GABA
transporter, Vgat) were extracted and re-clustered, consisting of 1847 neurons, which formed
19 clusters (**Figure 1A**). Of these clusters, 2-Cacna2d1/Tmem163 expressed Vglut2
transcripts, and lower levels of *Gad1* and *Gad2.* Low expression of classical neurotransmitter
related genes was found in these GABAergic neuronal clusters (**Figure 1B**). Adipocyte
hormone Leptin receptor (*Lepr*) expression was identified in 13-Onecut2/Lepr and 15-
Ebf2/Acly clusters (**Figure 1B**). We identified some expression of incretin receptors including
glucagon-like receptor 1 (GLP-1R) and gastric inhibitory polypeptide receptor (GIPR) in
several clusters (**Figure S1A**). Differential gene expression analysis was performed on each
cluster to identify the effects of an overnight fast on transcript expression in GABAergic
(*Slc32a1*+) neurons (**Figure S1B**). Cluster 3-Dcc/Cdh8 displayed significant upregulation in
transcript expression in response to a fast, however 93.7% of this cluster originates from *ad*
libitum fed animals. Significantly differentially regulated genes in *Slc32a1*^{+} neurons can be
found in **Figure S1C**. These data indicate minimal overlap with specific populations of NTS
neurons that have been previously described with regards to energy balance.

We therefore, examined whether GABA[NTS] cells are responsive to energy status. We
facilitated the visualization of GABA[NTS] neurons by crossing *Vgat-ires-Cre* mice with
tdTomato^{fl/fl} reporter mice (*Vgat[tdTom]*, **Figure 1D and 1E**). We then examined the expression
levels of the neuronal activity marker c-Fos (Olson et al., 1993) in mice that were overnight
fasted and in mice that were refed for 2 hours after overnight fasting. Re-fed mice showed
significantly more c-Fos in the NTS (**Figure S1D**) and specifically an increased number of
GABAergic cells activated (**Figure 1F and 1G**) when compared to fasted mice. This suggests
that GABA[NTS] cells are responsive to positive energy status.

To determine whether GABA^{NTS} neurons have a role in the control of food intake, we used

chemogenetic Designer Receptors Exclusively Activated by Designer Drugs (DREADD) to

manipulate GABA^{NTS} cellular activity (Alexander et al., 2009). Specifically, AAVs expressing

hM3Dq, hM4Di or control mCherry were bilaterally injected into the NTS of *Vgat^{Cre}* mice

(**Figure 1H**). This allowed to modulate GABA^{NTS} neuron activity with the administration of the

designer drug clozapine-N-oxide (CNO). Administration of CNO in hM3Dq-injected mice

induced strong c-Fos expression in the NTS (**Figure 1I**). We next assessed the effect of the

modulation of GABA^{NTS} on food intake in mice under different scenarios. In ad libitum fed

mice, hM3Dq chemogenetic activation of GABA^{NTS} cells significantly reduced overall food

intake during the active dark cycle when compared to control (mCherry) siblings (**Figure 1J**

and S1E). Similarly, hM3Dq GABA^{NTS} cell activation reduced food intake in hunger-induced

fasted mice when compared to control mice (**Figure S1F and S1G**). Conversely, hM4Di

chemogenetic inhibition of GABA^{NTS} cell in satiated mice during the light cycle significantly

increased food intake when compared to satiated littermate mCherry controls (**Figure 1K**).

Given the potent reduction of food intake produced by the activation of GABA^{NTS} neurons,

we examined whether prolonged activation impacted body weight. Twice-daily administration

of CNO (**Figure 1L**) in hM3Dq-expressing mice induced a significant reduction in daily food

intake (**Figure 1M**) and a progressive reduction in body weight (**Figure 1N and 1O**). Although

feeding returned to baseline levels 72 hours after the final CNO administration, body weight

remained significantly lower in hM3Dq-expressing mice (**Figure 1N**). In contrast, prolonged

GABA^{NTS} neuron inhibition using hM4Di did not induce greater food intake (**Figure S1H**) nor

changes in body weight (**Figure S1I**) when compared to their littermate mCherry controls.

These results indicate that the selective activation of GABA^{NTS} neurons is sufficient to result

in a sustained reduction in energy intake and hence body weight.

Recently, it has been reported that the inhibition of neurons expressing the GABA-producing

enzyme, glutamate decarboxylase (GAD), in the NTS of rats partially blunted the anorectic

effect of the GLP1-R agonist liraglutide (Fortin et al., 2020). These data, suggest that

GABA^{NTS} contributes to the therapeutic effects of GLP1-R agonists in rats. However, GLP-

1R expression in rats and mice differs, especially its receptor density (Cork et al., 2015;

Graham et al., 2020; Jensen et al., 2018). We therefore assessed the feeding response of

liraglutide in GABA^{NTS}-hM4Di/mCherry-expressing mice. Chemogenetic inhibition of

GABA^{NTS} partially blunted the acute anorectic effects of liraglutide (**Figure S1J and S1K**).

This provides support that GABA^{NTS} participates in the anorectic effects of liraglutide in mice.

Optogenetic stimulation of GABA[NTS] terminals inhibits AgRP/NPY[ARC] cells

To clarify the circuitry through which GABA[NTS] neurons influence feeding and body weight, we used Channelrhodopsin 2 (ChR2)-assisted circuit mapping (CRACM) (González et al., 2016; Petreanu et al., 2007). Specifically, an AAV expressing ChR2-mCherry was injected into the NTS of *Vgat[Cre]* mice (**Figure 2A**) and projection patterns were analyzed. GABA[NTS] cells project to hypothalamic subregions including the ARC, paraventricular nucleus (PVH) and dorsomedial nucleus (DMH) and various other extra-hypothalamic regions (**Figure S2A**). Within the ARC, we observed a dense array of projections (**Figure 2B**). This identified the ARC as a candidate second order region involved in the GABA[NTS] control of food intake and body weight.

Fasting induces activation of AgRP/NPY-expressing neurons (Hahn et al., 1998) and direct AgRP/NPY neuron activation induces robust feeding (Aponte et al., 2011; Krashes et al., 2011). Since GABA-releasing neurons are the main inhibitory network in the brain (Krnjević & Schwartz, 1967), we hypothesized that GABA[NTS] cells projecting to the ARC would target AgRP/NPY neurons to decrease food intake. To interrogate this potential GABA[NTS]→AgRP/NPY[ARC] circuit, *Vgat[Cre]* were crossed with *Npy[hrGFP]* mice (*Vgat[Cre]::Npy[hrGFP]*) and bilaterally injected with AAV-ChR2-mCherry into the NTS. We observed that a subset of NPY[hrGFP] cell bodies were surrounded by mCherry-containing fibers (**Figure 2C**). In contrast, terminals were not found surrounding neurons expressing other neuropeptides involved in the regulation of food intake such as POMC neurons (Williams & Schwartz, 2005) (**Figure 2D**).

We next investigated whether this anatomical connectivity produced functional interactions between GABA[NTS] terminals and AgRP/NPY[ARC] and POMC neurons. Specifically, photo-stimulation of ChR2-containing axon terminals from GABA[NTS] neurons produced robust synaptic responses in 14% of AgRP/NPY cells in the ARC but not in POMC cells (**Figure 2E**). The rapid synaptic currents triggered in AgRP/NPY cells by the optical stimulation changed polarity near the equilibrium potential for chloride (**Figure 2F and 2G**), as expected from ionotropic GABA receptors. Light-induced currents were unexpectedly small, and their reversal potential was more positive than that expected for GABA-activated currents (about -60 mV) (**Figure 2H-J**). These effects could be explained by voltage- and space-clamp errors (Spruston et al., 1993) and in turn suggest that perhaps GABA[NTS] terminals reach ARC NPY

cells at dendrites distant from the soma. While it is not possible to directly demonstrate that

this is the case, we tested whether this reasoning was justified by simulating the optogenetic

activation of GABA synaptic events at distal vs proximal dendrites in a model neuron. Using

a predictive neuronal model (Lindroos & Hellgren Kotaleski, 2021), we found that GABAergic

post-synaptic currents become progressively smaller with their reversal potential

progressively more positive, the further the GABA inputs are from the soma (**Figure S2 B-**

E).

Next, we tested whether activation of GABA receptors at NPY cells is accompanied by

release of GABA from GABA^{NTS} ChR2-containing terminals. We first stimulated ChR2-

mCherry expressing GABA^{NTS} fibers in the ARC and performed a "sniffer patch" experiment

registering GABA$_A$R single-channel openings above the NPY^{hrGFP} cells contacted by

GABA^{NTS} fibers (**Figure 2K**). In this experiment, the GABA$_A$R response was isolated with a

specific cocktail of antagonists (see Methods). This provides semi-quantitative monitoring of

extracellular levels of GABA (Sylantyev et al., 2020). A burst of light directed to GABA^{NTS}

ChR2-containing terminals in the ARC induced single-channel openings in membrane patch

in control conditions, and the currents had amplitudes comparable to GABAergic inhibitory

currents (Sylantyev et al., 2020) (**Figure 2L, *left***). Application of GABA$_A$R competitive

antagonist gabazine reversibly blocked the single-channel openings (**Figure 2L, *middle and***

right). The same pattern of receptor opening time was observed in all assessed patches

(**Figure 2M**). This suggested that postsynaptic NPY cells are sensitive to GABA^{NTS}

presynaptic release of GABA.

To test whether NPY cells were inhibited by the release of GABA from GABA^{NTS} terminals,

we designed a protocol of additive subthreshold electrical stimuli. This method is designed to

evoke an action potential after 5 stimuli. In addition, we coupled a 470 nm light burst to the

same trigger (**Figure S2F**). We then alternated a sequence of electrical stimuli and electrical

stimuli with light burst. The electrical stimulation evoked an action potential(s) (**Figure S2G,**

left) followed by low-frequency or no single-channel openings in the sniffer patch. When we

coupled the electrical stimulation with the light burst, the occurrence of actions potentials was

blocked and accompanied with single-channel openings in a sniffer patch (**Figure S2G,**

right). This happened in all cells patched (**Figure S2H**). These openings resembled GABA$_A$R

openings illustrated in Figure 2L. Subsequent stimulations showed a decrease in channel

opening intensity, suggesting a depletion of GABA stores from the presynaptic terminal,

which was eventually insufficient to prevent the action potential (**Figure S2I**). These results

provide strong evidence that GABA[NTS] fibers inhibit AgRP/NPY cells due to the release of the
fast neurotransmitter GABA.

GABA[NTS]→ARC optogenetic stimulation reduces feeding and is not aversive

Given the dense fiber projection pattern of NTS-GABAergic cells to the ARC, and that its
activation induced a strong inhibition of AgRP/NPY neurons, we interrogated whether this
circuit is sufficient to influence food intake. To investigate this, AAVs expressing ChR2-
mCherry were infused into the NTS of *Vgat[Cre]* mice and an optic fiber was placed above the
ARC (**Figure 3A**). Food intake was measured both without and with ARC photo-stimulation
prior to the onset of the dark cycle (**Figure 3B and 3C**). Light stimulation of GABA[NTS]→ARC
terminals induced a strong acute inhibition of food intake that lasted 60 min from food
presentation, as compared to the same mice without light stimulation (**Figure 3D and 3E**).

The DVC has been proposed to be a key region modulating food intake reduction associated
with aversive states (D'Agostino & Luckman, 2022). To assess whether GABA[NTS]→ARC
activation produces aversion or negative valence (Berridge, 2004; Betley et al., 2015), we
evaluated the existence of passive avoidance behavior using an adapted real-time place
preference (RTPP) task (D'Agostino et al., 2016; Kim et al., 2013; Stamatakis & Stuber,
2012). Specifically, in a two-sided open arena, light stimulus was coupled to one of the sides
(**Figure 3F**). GABA[NTS]→ARC stimulation or lack of stimulation did not produce a place
preference (**Figure 3G**) and mice travelled a similar distance in both sides of the arena
(**Figure 3H**). These findings indicate that GABA[NTS]→ARC stimulation does not produce
negative valence or aversion.

In addition to influencing homeostatic feeding, modulation of AgRP/NPY cells has also been
reported to elicit anxiety-like behaviors influencing exploration and foraging which impacts
food consumption (Dietrich et al., 2015; Heinz et al., 2021; Li et al., 2019). To test whether
the activation of the GABA[NTS]→ARC circuit produced anxiety-related behavior, mice were
assessed in an open-field arena (OFA) and an elevated zero-maze task (EZM) (**Figure S2A
and S2D**). GABA[NTS]→ARC stimulation did not alter the time mice spent in the center of the
OFA (**Figure S2B**). Likewise, mice displayed similar ambulatory patterns and travelled
comparable distance during the OFA test with and without GABA[NTS]→ARC stimulation
(**Figure S2C**). Consistent with the OFA data, GABA[NTS]→ARC stimulation did not alter either
the time (**Figure S2E**) or distance travelled in the exposed zones of the EZM (**Figure S2F**).

These results provide evidence that stimulation of the GABA^{NTS}→ARC circuit impacts feeding
without altering other behavioral states such as anxiety-like behavior.

We postulated that since our model involved fast neurotransmission rather than long term
release of neuropeptides, the reduction in food intake that we observed could begin by
reducing the interaction with the nutritional cue at first instance. To investigate this, we
designed a task where mice would first be allowed to explore an empty arena and then three
items would be presented at the same time, an inedible object, a novel palatable food item
and a known nutritional food item (**Figure 3I**). The number of interactions with each item was
measured with and without GABA^{NTS}→ARC stimulation. GABA^{NTS}→ARC stimulation did not
induce changes in the interaction with the novel object (**Figure 3J**), further suggesting that
GABA^{NTS}→ARC does not induce anxiety-like behavior. The number of interactions with the
novel palatable food item was increased under both non-stimulation and stimulation trials
(**Figure S2F and S2G**). However, we found that under optical stimulation mice had less
interactions with the known nutritional item (**Figure 3J and S2G**), suggesting that
GABA^{NTS}→ARC activation reduces hunger. Taken together, these findings indicate that
GABA^{NTS}→ARC stimulation decreases feeding and hunger without inducing aversion or
anxiety.

GABA^{NTS}→ARC neuron activation promotes satiety

We next used a two-virus intersectional approach (Fenno et al., 2014, 2017) to provide a
more detailed characterization of GABA^{NTS}→ARC activation in energy homeostasis. *Vgat^{Cre}*
mice were bilaterally injected with AAVs expressing Flipase recombinase (FlpO) (AAV-DIO-
FlpO) under the control of Cre into the NTS of *Vgat^{Cre}* mice, allowing us to express a second
recombinase only in GABA^{NTS} cells. After surgery recovery, a retrograde AAV encoding for a
FlpO dependent hM3Dq-mCherry (rgAAV-fDIO-hM3Dq-mCherry) was bilaterally injected into
the ARC to retrogradely deliver hM3Dq in a FlpO-dependent manner to GABA^{NTS} cells.
Therefore, hM3Dq was expressed only in GABA^{NTS} cells projecting to the ARC in *Vgat^{Cre}*
mice (**Figure 4A**).

Corroborating the optogenetic data presented above, the selective chemogenetic stimulation
of the GABA^{NTS}→ARC circuit significantly reduced acute food intake (**Figure 4B** and **4C**) in
ad libitum fed mice. GABA^{NTS}→ARC circuit activation did not alter overall locomotor activity

(**Figure 4D** and **4E**), respiratory quotient (**Figure 4F** and **4G**) or heat production (**Figure 4H**
and **4I**). GABA^{NTS}→ARC neuron activation also reduced food intake in refed mice after
overnight fasting (**Figure 4J**). Fasting is associated with a rise in systemic levels of ghrelin
(Cummings et al., 2001) which acts as a pre-prandial effector stimulating AgRP/NPY neurons
to initiate a feeding response (H. Y. Chen et al., 2004; Luquet et al., 2007). To examine
whether GABA^{NTS}→ARC neuron activation is sufficient to dampen a hunger cue,
GABA^{NTS}:hM3Dq-expressing mice were pre-treated with CNO prior to ghrelin. Activation of
the GABA^{NTS}→ARC neurons with CNO prevented the feeding induced by an orexigenic dose
of ghrelin (**Figure 4L and 4M**). We next performed an analysis of the microstructure of the
feeding event during the anorectic episode produced by CNO (Clifton, 2000; Richard et al.,
2011; Zorrilla et al., 2005). GABA^{NTS}→ARC neuron activation reduced meal size (**Figure 4N**)
and significantly reduced the number of meal events in the earliest interval compared to saline
treatment (**Figure 4O** and **4P**). Taken together, these findings indicate that GABA^{NTS}→ARC
neuron activation is sufficient to blunt fasting and ghrelin-induced hunger and significantly
reduces food intake by promoting satiety.

Discussion

Here we identify a critical new brain circuit modulating appetite and body weight. We focused
on the NTS because it is a brain region positioned to receive and integrate energy-related
information from the periphery and relay it within the CNS to promote energy homeostasis.
However, the NTS is neurochemically heterogeneous and key neurons within it performing
this function have not been fully defined. Using multi-methodological approach, here we
identify GABA^{NTS} neurons as sufficient to control feeding behavior and body weight in mice.

Recent efforts to decode the function of specific chemically defined neurons within the NTS
have revealed that distinct subpopulations of glutamatergic cells play a role in energy
homeostasis (Dowsett et al., 2021). However, NTS inhibitory GABA-releasing neurons have
not been studied in detail and this is necessary to clarify the role of both excitatory and
inhibitory NTS signals in the regulation of energy balance (Cheng et al., 2022). Here we
provide a detailed characterization of the effect of GABA^{NTS} in the regulation of energy
homeostasis and body weight. A recent report provided evidence that obesity medication
liraglutide engages GABA^{NTS} neurons to reduce food intake in rats, providing a rationale that

activating GABA[NTS] cells may have translational relevance for the treatment of human obesity
(Fortin et al., 2020).

The NTS is involved in satiety and satiation, and refeeding induces a strong neuronal
activation in this region (D'Agostino et al., 2016; Wu et al., 2014). We discovered that
refeeding significantly activates a subset of GABA[NTS] cells. Further, we found that activation
of GABA[NTS] neurons with chemogenetics was anorectic, and controlled food intake and body
weight. Consistently, when we inhibited these neurons, we observed a potent induction of
feeding in satiated mice, suggesting a crucial role in the induction of early eating. However,
when the inhibition occurred in ad libitum mice entering the dark cycle, we didn't observe an
increased food intake. This could be explained by two reasons. First, despite of mice being
kept in ad libitum conditions, they are not satiated at the onset of the dark cycle, meaning
GABA[NTS] neuros are likely to not be active at this time, therefore an inhibition would not
meaningful. Second, during the active dark cycle there are multiple feeding cues not only
vagal sensory afferents but afferent from the sensory system that would modulate feeding
and could counteract GABA[NTS] lack of inhibition in second order neurons. Our data illustrates
that GABA[NTS] neurons project widely within the brain, with particularly dense innervation of
the hypothalamus. Several studies indicate that there is a coordination between the NTS and
the hypothalamus to orchestrate the meal event (Aklan et al., 2020; Blevins et al., 2004;
D'Agostino et al., 2018; Liu et al., 2017; Tsang et al., 2020). However, whether the ARC
receives inhibitory control from the NTS is not known and was examined here.

Given that GABA is an inhibitory neurotransmitter, we hypothesized that GABA[NTS] neurons
inhibit appetite stimulating neurons. We focused on AgRP/NPY[ARC] neurons because of the
dense GABA[NTS] innervation that we found and the potent orexigenic properties of AgRP/NPY
(Aponte et al., 2011; Krashes et al., 2011; Luquet et al., 2005). In addition, AgRP/NPY
neurons display a multiple timescale activation (Mandelblat-Cerf et al., 2015) that could
explain the lack of feeding effect during the chemogenetic inhibition of GABA[NTS]. We
demonstrated that GABA[NTS] terminals in the ARC release GABA and that this is synchronized
with suppression of the action potential propagation postsynaptic AgRP/NPY[ARC] cells. The
timescale of GABA$_A$R openings after optical stimulation in a membrane patch placed over the
GABA[NTS] terminal suggests the release of GABA from this terminal rather than from another
inhibitory neurons. The number of AgRP/NPY neurons responding is in line with the 20%
ARC responders to NTS innervation published previously (Aklan et al., 2020). However,
despite the relatively small number of AgRP/NPY[ARC] neurons inhibited by GABA[NTS] terminals,

it was sufficient to significantly reduce feeding in freely behaving mice. AgRP/NPY cells are

poised to orchestrate the integration of homeostatic, reward and sensory cues as well as

learning and conditioned behaviors (Aponte et al., 2011; Berrios et al., 2021; Deem et al.,

2022; Dietrich et al., 2012, 2015; Garau et al., 2020; Han et al., 2021; Jikomes et al., 2016;

Krashes et al., 2011, 2014; Wang et al., 2021). A detailed analysis of feeding behavior

revealed that activation of the $GABA^{NTS} \rightarrow ARC$ pathway specifically reduces hunger and

promotes satiety and does not induce negative valence, aversion or anxiety.

Obesity is an international health concern that is primarily the consequence of over-eating.

Defining the mechanisms governing hunger and food intake is therefore of paramount

importance. Here we identify a new player that controls appetite and body weight, $GABA^{NTS}$.

Specifically, we show that NTS GABA-releasing neurons are active during satiety and reduce

food intake and body weight without causing aversion or anxiety. We demonstrate that

$GABA^{NTS}$ cells directly activate $GABA_A$Rs on the surface of AgRP/NPY cells which inhibits

neuron activity. These studies reveal for the first time the effect of $GABA^{NTS}$ neurons on

feeding and body weight, and identify a fast inhibitory circuit between the NTS and the ARC

in the control of food intake. These results thereby provide significant insight into the brain

circuits governing appetite and body weight, findings of relevance to the global obesity crisis.

Methods

RNA-Seq

Single nucleus RNA-sequencing data from the mouse hindbrain in the fed and fasted state was taken from (Dowsett et al., 2021). Neuronal nuclei expressing at least 1 UMI count for *Slc32a1* were identified as GABAergic neurons, subsetted and reclustered using Seurat package version 4.3 (Hao et al., 2021). Marker genes for each cluster were calculated using Wilcoxon's rank-sum test. Each cluster was named with 2 marker genes that were expressed in >60% of the cluster, <30% of the rest of the data and had an average log fold change >0.5. If no genes fit these criteria, then the two genes with the lowest p-values were used. Differential gene expression analysis between ad libitum fed and overnight fasted cells was performed using the Wilcoxon's rank sum test. Feature plots were drawn using the Seurat package and ggplot2.

Animals

Vgat-ires-Cre (Vong et al., 2011) (Slc32a1tm2(cre)Lowl; #016962), *NPY-hrGFP* (van den Pol et al., 2009)(van den Pol et al., 2009)(van den Pol et al., 2009)(van den Pol et al., 2009) (B6.FVB-Tg(Npy-hrGFP)1Lowl/J; #006417), POMC-dsRed (Hentges et al., 2009) (Tg(Pomc-DsRed)18Low) and *Rosa26tdTomato-LoxP* (Madisen et al., 2009) (B6.Cg-Gt(ROSA)26Sortm9(CAG-tdTomato)Hze/J, #007909) mice were obtained from The Jackson Laboratory (Bar Harbor, USA) and bred in a C57Bl/6J background. Mice were fed with standard laboratory chow (Standard CRM (P) 801722, Special diets, UK) and provided with water ad libitum, unless otherwise stated. Mice were kept in a 12-hours light:dark cycle (7am-7pm) in environmental controlled conditions (20-22°C and 40-60% RH). All experimental procedures were performed in accordance with the UK Animal (Scientific Procedures) Act 1986.

Viral vectors

Cre-dependent viral vectors were purchased from Addgene, AAV8-hSyn-DIO-hM3D(Gq)-mCherry (1.83×10^{12} gc/ml) was a gift from Bryan Roth (Addgene plasmid # 44361) (Krashes et al., 2011), AAVrg-hSyn-fDIO-hM3D(Gq)-mCherry-WPREpA (1.8×10^{12} gc/ml) was a gift from Ulrik Gether (Addgene plasmid # 154868); AAV8-hSyn-DIO-mCherry (3.6×10^{12} gc/ml) was a gift from Bryan Roth (Addgene plasmid # 50459); AAV8-pEF1a-DIO-FLPo-WPRE-

hGHpA (2×10^{12} gc/ml) was a gift from Li Zhang (Addgene plasmid # 87306) (Zingg et al.,
2017). AAV2-EF1a-DIO-ChR2(E123T/T159C)-mCherry and AAV2-EF1a-DIO-
ChR2(E123T/T159C)-YFP (7.3×10^{12} vp/ml) were a gift from Karl Deisseroth and were
obtained from University of North Carolina Vector Core (Chapel Hil, NC, USA). All viral
particles were delivered into nuclei-specific regions through stereotaxic injections.

Stereotaxic surgeries

For viral delivering in the NTS, stereotaxic surgery was adapted from previous studies
(D'Agostino et al., 2016). Briefly, 12-20 weeks old mice were anaesthetized with isoflurane,
back region of the neck shaved and placed in a stereotaxic instrument (David Kopf
instruments, CA, USA) with a face mask (World Precision Instruments, FL, USA). Head was
inclined ~70 degrees forward and a longitudinal incision was made in the skin at the level of
the C1; neck muscles were retracted to expose the atlanto-occipital membrane. This was
carefully dissected allowing access to the dorsal brainstem and visualization of the obex.
Using a pulled glass capillary (40μ tip diameter) (G1, Narishige, UK) and a pneumatic
microinjector (IM-11-2, Narishige, UK) 200-300 nl of viral preparation was bilaterally injected
into the NTS (obex: AP:0.25 mm AP, L:± 0.25 mm and DV:-0.25mm) at a flow of 50nl/min.
Capillary was left in the injection place for 5 min to allow diffusion and it was removed slowly
to avoid dispersion to neighbor brainstem regions. Viral delivery into the ARC was performed
as previously described (Wagner et al., 2022) at coordinates bregma: AP:1.58 mm AP,
L:±0.2mm and DV:5.90 mm. For optical fiber cannula placement, mice were allowed 4 weeks
recovery form the NTS surgery and a 200 μm core diameter, 0.39NA (CFMLC, Thorlabs, UK)
optical fiber implants were placed in the third ventricle above the ARC. Mice were allowed 3
weeks before any study to allow surgery recovery and maximal viral expression. Post hoc
analysis of injection site, viral expression and canula placement were used as exclusion
criteria for data analysis.

In vivo photo-stimulation protocol

Optical fiber implants were attached to optogenetics patch cables (M83L1, Thorlabs, UK)
connected to a rotary joint (Doric lenses) coupled to a 473-nm laser (Laserglow, Toronto,
Canada) controlled via TTL-USB interface with Arduino board. For feeding experiments, the

stimulation protocol was 1 s followed by 4 s break with 10ms light pulses with a frequency of 30Hz. For behavioral experiments, the stimulation protocol was 1 s followed by 0.5 s break with 10ms light pulses with a frequency of 30Hz. We used 15mW of laser power to achieve an irradiance of 5-10mW/mm^2 (PM100D, Thorlabs) on the target area following https://web.stanford.edu/group/dlab/cgi-bin/graph/chart.php, above ChR2 threshold activation (Lin et al., 2009).

Food intake and body weight studies

For food intake, body weight and metabolic parameters measurements, mice were single housed and habituated in indirect calorimetry system cages for one week (Phenomaster, TSE Systems, Germany). For acute ad libitum studies, access to food was removed in fed mice 2h before entering the dark cycle and CNO 1 mg/kg was i.p. administered 30 min before the dark cycle onset when food was provided. For re-feeding studies, 12 hours dark cycle-food deprived mice were i.p. injected with CNO at the beginning of the light cycle and 30 min after food was provided. For subchronic studies, mice were i.p. injected twice a day (am and pm) for 5 days with CNO following 5 days with saline.

Behavioral tests

For valence studies, mice were assessed in an adapted real-time place preference task consisting in an open field arena with two connected identical chambers (30x25cm) (D'Agostino et al., 2016; Kim et al., 2013; Stamatakis & Stuber, 2012), one of them paired with optogenetic stimulation where mice were allowed free movement for 20 min. For anxiety tests, mice were placed in an open arena (50x50cm) with virtual delimited central and peripheral regions and allowed free movement for 10 minutes with and without stimulation in different days. For anxiety and fear assessment, mice were placed in an elevated zero maze (diameter 50cm, elevation 70 cm) with 2 hidden and 2 exposed zones and allowed free movement between zones for 10 min. Tests were performed for each animal with and without stimulation in different days. Time and locomotor parameters for each task and zone were recorded using Any-Maze software (Stoelting, IL, USA).

Immunohistochemistry and imaging

All mice were injected with a terminal dose of anesthesia and transcardially perfused with phosphate-buffered saline (PBS) followed by 10% neutral buffered formalin. Brains were dissected, post-fixed 12 hours in formalin at 4°C, cryoprotected 48 hours with 30% sucrose 4°C and coronally sectioned in 5 series at 25 μm using a freezing microtome (8000, Bright Instruments, UK). Sections were kept in protective anti-freeze solution at 4°C until they were processed for immunohistochemistry as previously described (Yavari et al., 2016). Briefly, NTS sections were washed with PBS-0.2% Tween20 30 min and then PBS (3x10 min), blocked with 1%BSA/5%DS/0.25%Triton X-100 1 hour at room temperature and incubated with primary antibody in blocking solution with anti-c-Fos (1:2500, 2250, CST, USA), anti- c-Fos (for chromogenic) (1:5000, ABE457,Merck, UK), anti-mCherry (1:2000, AB0040-200, Scigen, PT), anti-RFP (1:1000, 600-401-379, Rockland Immunochemicals, USA), anti-POMC (1:3000, H-029-30, Phoenix Pharmaceuticals, USA), anti-hrGFP (1:2000, 240141, Agilent, USA), anti-TH (1:2000, MAB318, Merck, Germany) 16 hours at room temperature. The next day, sections were washed with PBS-Tween and PBS and incubated 1 hour with appropriate secondary antibodies in blocking solution (1:500, AlexaFluor594, AlexaFluor488, Invitrogen, UK) at room temperature. For c-Fos expression quantification in fast vs refed study, chromogenic staining with DAB reagent was performed as previously described (D'Agostino et al., 2016).

Images were acquired using Axioskope2 microscope and Axiovision software (Zeiss, Germany). All images were converted to 8-bit, peudorecolored and cells counted using ImageJ (Fiji).

Electrophysiology

CRACM study

CRACM experiments were performed as previously described (González et al., 2016). Six *Vgat^Cre^:Npy^hrGFP^* mice and six *Vgat^Cre^:Pomc^dsRed^* mice bilaterally injected with AAV-ChR2-mCherry and AAV-ChR2-mCherry respectively into the NTS, aged between 5 and 7 months at the time of the electrophysiology experiments, were used. Expression of mCherry-ChR2 was targeted to NTS^GABA^ cells by stereotaxic injection of AAV as above. Coronal brain sections 180-μm thick were prepared from these mice at least 8 days after virus injections

and were placed in a bath solution consisting of (in mM) 125 NaCl, 2.5 KCl, 1.2 NaH_2PO_4, 21

$NaHCO_3$, 1 glucose, 2 $MgCl_2$, 2 $CaCl_2$. GFP-expressing cells in the ARC were identified using

an upright microscope (Scientifica S-Scope-II) equipped with the appropriate fluorescence

filters. Whole-cell recordings from these cells were obtained with glass pipettes (World

Precision Instruments 1B150F-4) filled with a solution containing (in mM) 120 K-gluconate,

10 HEPES, 10 KCl, 1 EGTA, 2 MgCl2, 4K2ATP, and 1 Na2ATP, tip resistance 3-7 MOhm.

Data was acquired using Axon Instruments hardware (MultiClamp 700B, Digidata 1550). To

test for GABA inputs to AgRP/NPY[ARC] cells, the membrane potential in these cells was

clamped at increasing levels of voltage (from -100 to -10 mV in 10-mV increments), while

ChR2-expressing terminals were stimulated by a single light pulse (CoolLED pE-4000) to

induce post-synaptic currents. Liquid junction potential, estimated to be 10 mV, was

subtracted from the measurements. Chloride equilibrium potential was calculated to be -60.3

mV.

Sniffer-patch recordings

Transverse hypothalamic slices from *Vgat[Cre]:Npy[hrGFP]* mice bilaterally injected AAV-DIO-

ChR2-mCherry were cut at 200-250 using a Leica VT1200S vibratome. Slices were incubated

for one hour in a solution containing (in mM): 124 NaCl, 3 KCl, 1 $CaCl_2$, 3 $MgCl_2$, 26 $NaHCO_3$,

1.25 NaH_2PO_4, 10 D-glucose, and bubbled with 95/5% O_2/CO_2, pH 7.4. After incubation,

slices were transferred to a recording chamber continuously superfused with an external

solution. The external solution composition differed from incubation solution in containing 2

mM $CaCl_2$ and 2 mM $MgCl_2$.

In all experiments the intracellular pipette solution for voltage-clamp recordings contained

(mM): 117.5 Cs-gluconate, 17.5 CsCl, 10 KOH-HEPES, 10 BAPTA, 8 NaCl, 5 QX-314, 2 Mg-

ATP, 0.3 GTP; for current-clamp recordings: 126 K-gluconate, 4 NaCl, 5 HEPES, 15 glucose,

1 $MgSO_4 \cdot 7H_2O$, 2 BAPTA, 3 Mg-ATP (pH 7.2, 295-310 mOsm in both cases); pipette

resistance was 7-9 MOhm; recordings were performed at 33-35ºC using Multiclamp-700B

amplifier with -60 or -70 mV holding current (for voltage-clamp recordings); signals were pre-

filtered and digitized at 10 kHz. In experiments where transmembrane currents were recorded

in outside-out patches only (sniffer-patch recordings), the GABA$_A$ receptors response was

isolated with a ligands cocktail containing 50 µM APV, 20 µM NBQX, 50 nM CGP-55845, 200

µM S-MCPG, 10 µM MDL-72222, and 1 µM strychnine.

Statistics and data analysis

Statistical analyses were performed using GraphPad Prism 9 software and are described in the figure legend where 2-tail paired and unpaired Student t-test were used when comparing 2 groups and RM/two-way ANOVA test with Bonferroni post-hoc correction when comparing 4 groups. RNA-Seq data were analyzed as described above. No statistical method was used to predefine sample size, randomization and blinding was performed for histological quantifications. Statistical significance was accepted when the $p \leq 0.05$. Raw data was stored in Excel and figures assembled with CorelVector and Affinity Designer software.

Author Contributions

The project was conceived by PBM and LKH. PBM designed and performed experiments and analyzed data with assistance from YM; GKCD and GSHY performed and analyzed the RNA-Seq data. SS and AG designed and performed electrophysiological recordings. The manuscript was drafted by PBM with input from all other authors.

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Conflict of interest

The authors declare no competing financial interests.

Figure 1

Figure 2

Figure 3

Figure 4

Supplementary Figure 1

Supplementary Figure 2

 Figures Legends

Figure 1. GABA^{NTS} activation reduces food intake and body weight. (A) tSNE plot of
Slc32a1^+ neurons in the DVC colored by cluster, (B)Dot plot showing scaled expression of
marker genes for each of the 19 *Slc32a1^+* clusters (C) tSNE plot of the *Slc32a1^+* nuclei
colored by nutritional status (D) Schematic of the DVC (top) with zoom-in diagram showing
AP, NTS and DMC (bottom). (E) Representative photomicrograph of GABAergic cell
distribution in the medial brainstem (top) and in the DVC (bottom) using *Vgat^{tdTom}* mice. (F)
Quantification GABAergic cells expressing c-Fos following 16h fasting and fasting+2h
refeeding (n=3, Bregma level: two-way ANOVA $F_{(13,34)}$=2.63; p=0.012; Nutritional state: two-
way ANOVA $F_{(1,4)}$=8.69; p=0.042. (E) Representative micrograph of *Vgat^{tdTom}* cells (red)
expressing c-Fos (green) in mice fasted 16h (top) and re-fed for 2h (bottom). (H) Schematic
of AAV-DIO-hM3Dq/hM4Di/mCherry infused into NTS of *Vgat^{Cre}* mice. (I) Representative
photomicrograph of c-Fos (green) expression in hM3Dq-mCherry (red)-expressing cells in
the NTS of VgatCre mice treated with CNO (1 mg/kg, i.p) and (J) 6h cumulative and (J) 2h
and 6h total food intake (n=6, Unpaired t test $t_{(11)}$=2.663, p=0.0238) of GABA^{NTS}:hM3Dq CNO
injected mice compared to mCherry. (K) 2h cumulative and (J) 2h total food intake (n=5,
Unpaired t test $t_{(8)}$=3.205, p=0.0125) GABA^{NTS}:hM4Di CNO injected mice compared to
mCherry. (L) 10-day treatment protocol schematic. Daily CNO treatment significantly reduced
(M) food intake (RM two-way ANOVA $F_{(1,11)}$=20.57; P=0.0008), (N) body weight (n=6/7, RM
two-way ANOVA ($F_{(1,11)}$=21.96; p=0.0007) and (O) body weight change GABA^{NTS:hM3Dq} mice
compared to control GABA^{NTS}:mCherry mice (n=6/7, Unpaired t test $t_{(11)}$=5.252, p=0.0003).
C-D n=4/group, H-K n=6; M-O n=6 mCherry and n=7 hM3Dq. Data represented as
mean±S.E.M. *: p<0.05; **: p<0.01; ***: p<0.001. AP: Area Postrema; CC: Centre canal;
DMV: Doral motor nucleus of the vagus nerve; NTS: Nucleus of the solitary tract.

Figure 2. GABA^{NTS} projections to the ARC release GABA into AgRP/NPY neurons. (A) Illustration depicting details for NTS injection of AAV-DIO-ChR2-mCherry (top) and representative photomicrograph (scale 1mm) of injection (bottom) in *Vgat^{Cre}* mice and (B) ChR2-mcherry containing fibers in the ARC (scale 200um). (B) Representative photomicrographs (scale 100um) and magnification (scale 50 and 10um) ChR2-mCherry fibers in the ARC close to NPY^{hrGFP}-expressing cells and (D) POMC-expressing cells and in *Vgat^{Cre}* mice bilaterally injected AAV-DIO-ChR2-mCherry. (E-J) *Vgat^{Cre}:Npy^{hrGFP}* mice CRACM study. (E) Representative membrane potential response of NPYGFP and POMCDsRed cells after light stimulation of mCherry containing terminals. (F) Representative membrane current in Npy^{hrGFP} cell recorded during a voltage-clamp experiment from -100mV to -10mV in 10-mV steps. (Vertical line: light pulse; green cross: base line; red cross: peak value). (G) Representative light-induced post-synaptic current during voltage clamp experiment showing peak conductance, G_s, and reversal potential, E_{rev}. (H) Individual values and median of (H) E_{rev}, (I) Conductance, G_s and (I) latency of all responsive cells. (K) Diagram of outside-out patch technique in *Vgat^{Cre}:Npy^{hrGFP}* mice bilaterally injected AAV-DIO-ChR2-mCherry. (L) Representative channel opening recordings of control (left) gabazine (middle) and washout (right) recordings. (M) Quantification of receptor opening time (n=4, RM two-way ANOVA $F_{(2,6)}$=16.14; p=0.0039, Bonferroni adjusted p=0.0051 Control vs Gabazine and p=0.0156 Gabazine vs Washout. Data in M are expressed as mean±S.E.M. E-J: 4/27 cells NPYhrGFP and 0/19 cells POMCdsRed; K-M: n=4-5 patches. *: p<0.05; **: p<0.01.

Figure 3. Activation of GABA^{NTS}→ARC projections reduces food intake without inducing aversion. (A) Schematic of viral infection and optic fiber placement for in vivo stimulation of GABA^{NTS}→ARC. (B) Protocol of stimulation. (C) Representative 1min resolution time plot of 3h food intake measurement. GABA^{NTS}→ARC stimulation significantly reduces (D) cumulative 3h food intake and (E) 60min intake (n=7, day 1 (OFF) vs day 2 (ON) paired Student t-test $t_{(6)}$=3.035, p=0.0229; day 2 (ON) vs day 3 (OFF) paired Student t-test $t_{(6)}$=3.283, p=0.0168). (F) Diagram illustrating RTPP task. GABA^{NTS}→ARC does not alter (G) time spent (representative heat map of time) or (H) distance travelled in stimulated and non-stimulated zones. (I) Diagram illustrating an object interaction task. (J) Interactions with novel, known nutritional or novel palatable objects and a representative path track around each object (red circle). GABA^{NTS}→ARC stimulation reduced interaction with known nutritional item (n=6, Student's t test $t_{(5)}$=3.043, p=0.0287). Data are expressed as individual values and as mean±S.E.M, n=6 mice. *: p<0.05; RTPP: real-time place preference.

Figure 4. Chemogenetic activation of GABA^{NTS}→ARC neurons induces satiety. (A) Diagram illustrating the two-virus intersectional strategy to express hM3Dq only in GABA^{NTS} neurons projecting to the ARC. Mice were treated with saline or CNO (1 mg/kg, i.p.). (B) CNO significantly reduced cumulative food intake over 6h (0-6h $F_{(1,15)}$=7.418, p=0.0157), (C) 1h (Unpaired t-test $t_{(15)}$=2.648, p=0.0183) and 3h (Unpaired t-test, $t_{(15)}$=2.568 P=0.0214) food intake compared to saline. (D) CNO did not alter 12h or (E) 1, 3, 6 or 12h quantification of ambulation; (F) 12h RER or (G) AUC quantification; (H) 12h or (I) 1, 3, 6 and 12h heat production per mouse compared to saline. (J-K) CNO significantly reduced 3h food intake following overnight fasting (J, 2-way ANOVA $F_{(1.13)}$=6.139, p=0.0277 and K, unpaired t-test $t_{(13)}$=2.320, p=0.0372). (L-M) CNO attenuated ghrelin hyperphagia over 2h (RM ANOVA $F_{(3,16)}$=11.12; p=0.0003, Bonferroni adjusted p=0.0042 control vs ghrelin, p=0.0003 CNO vs ghrelin and p=0.002 Ghrelin vs Ghrelin+CNO). (N) CNO reduced meal size (Unpaired t-test, $t_{(14)}$=2.256, p=0.0406) and (O-P) decreased meal number (Unpaired t-test, t(14)=2.824, p=0.0135) compared to control. Data are expressed as individual values and as mean±SEM. *: p<0.05; **: p<0.01. ***: p<0.001.

Supplementary Figure 1

(A) Proportion of GLP1-R and GIPR expression in each *Slc32a1*⁺ cluster (B) Number of genes upregulated (red) and downregulated (blue) in response to an overnight fast in each cluster. Genes included were significantly differentially regulated ($P < 0.05$). (C) Top 10 genes differentially expressed in ad libitum vs fasting in *Slc32a1*⁺ neurons (D) Quantification of total NTS c-Fos-expressing cells following 16h fasting and fasting+2h refeeding (n=3, Bregma level: two-way ANOVA $F_{(13,34)}=3.16$; p=0.004; Nutritional state: two-way ANOVA $F_{(1,4)}=18.61$; P=0.012). (E) 12h dark cycle food intake in hM3Dq vs mCherry mice injected with CNO. (F) Food intake following an overnight fast compared to mCherry in GABA^NTS:hM3Dq mice (n=6, RM two-way ANOVA $F_{(1,11)}=10.97$; P=0.0069) and (G) 2h food intake (n=6/7, Unpaired t test, $t_{(11)}=3.392$ p=0.006). (G-I) Food intake and body weight change in hM4Di vs mCherry mice twice daily injected with CNO (2mg/Kg). (J-K) Food intake in hM4Di mice co-administered with CNO and Liraglutide (RM: ANOVA $F_{(3,12)}=15.64$; P=0.002, Bonferroni adjusted p=0.0004 Saline vs Liraglutide, P=0.0004 CNO vs Liraglutide and P=0.0263 Liraglutide vs Liraglutide+CNO).

Figure S2. GABA^NTS projection pattern and CRACM additional analysis. (A) Representative photomicrographs (scale 100µm) of ChR2-mCherry fibers in a serial rostro-caudal distribution of a *Vgat^Cre* mouse brain injected with AAV-DIO-ChR2-mCherry into the NTS. (B-E) A simulated voltage-clamp experiment in an anatomically realistic model of a medium spiny neuron, using the same voltage steps as those described for electrophysiology experiments (Figure 2E-G). The effect of optogenetic stimulation was modelled as a single GABAergic event taking place at each step. (C) Current amplitude was plotted against voltage (as in Figure 2E-G), and a linear fit was used to estimate peak conductance G_s and reversal potential E_{rev}. (D-E) Simulated optogenetic activation of GABA inputs making synaptic contact only on dendrites close to the soma ("prox.", proximal dendrites) was compared to the same type of activation of GABA synapses connected only to dendrites far from the soma ("distal" dendrites) in six different medium spiny neuron models. The reversal potential for GABA in the model was set to -60 mV, and this is close to what was measured at the soma when GABA synaptic inputs were located on dendrites close to the soma. However, when GABA synapses were activated only on dendrites distant from the soma E_{rev}, was more positive than expected and conductance G_s was attenuated. (F) Diagram of

outside-out patch technique coupled to a postsynaptic electrical stimulator. (G, left)

Representative response of Npy^{hrGFP} cell subjected to series of five electrical stimuli, each

evoking excitatory post-synaptic potential (EPSP) of sub-threshold amplitude and (G, right)

same cell with electrical stimuli coupled with 470nm light burst. (M) Quantification of receptor

opening time (n=5, two-way RM ANOVA $F_{(2,8)}$=11.28; p=0.0047, Bonferroni adjusted

P=0.0072 NS vs SL and P=0.0157 S vs SL), L-M n=4-5 cells. (I) Representative response of

a single Npy^{hrGFP} cell subjected to 10 series of five electrical stimuli, each evoking excitatory

post-synaptic potential (EPSP) of sub-threshold amplitude coupled with 470nm light burst in

Vgat^{Cre}::NPY^{hrGFP} mouse injected with AAV-DIO-ChR2-mCherry into the NTS. NS: No

stimulation; S: Stimulation; SL: Stimulation+Light.

Figure S3. Optogenetic activation of GABA^{NTS}→ARC in *Vgat^{Cre}* mice does not induce
anxiety-like behavior. (A) Diagram illustrating open field arena (OFA) task. GABA^{NTS}→ARC
does not alter (B) time spent in the center or (C) distance travelled (representative trace of
the movement) during the test. (D) Diagram illustrating elevated zero maze (EZM) task.
GABA^{NTS}→ARC does not alter (E) time spent in the exposed area or (F) distance travelled in
each zone (representative trace of the movement) during the test. (G-H) Quantification of
number of interactions with a novel object, a known nutritional food item or a novel palatable
food item in mice (G) non-stimulated (RM ANOVA ($F_{(2,10)}$=4.993; p=0.0314, Bonferroni
adjusted p=0.0306 NO vs NP) and (H) stimulated (RM ANOVA $F_{(2,10)}$=10.71; p=0.0033,
Bonferroni adjusted p=0.0051 NO vs NP; p=0.0121 KN vs NP). Data are expressed as
individual values and as mean±S.E.M, n=6 mice. *:p<0.05; **:p<0.01. NO: novel object; NP:
novel palatable; KN: known nutritional.

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