Neuropeptide Y expression defines a novel class of GABAergic projection neurons in the

inferior colliculus

Abbreviated title: NPY neurons and NPY signaling in the IC

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Abstract

Located in the midbrain, the inferior colliculus (IC) integrates information from numerous

auditory nuclei and is an important hub for sound processing. Despite its importance, little is

known about the molecular identity and functional roles of defined neuron types in the IC.

Using a multifaceted approach in mice, we found that neuropeptide Y (NPY) expression

identifies a major class of inhibitory neurons, accounting for approximately one-third of

GABAergic neurons in the IC. Viral tract tracing showed that NPY neurons are principal neurons,

with axons that project locally and to the auditory thalamus and contralateral IC. In brain slice

recordings, many NPY neurons fired spontaneously, suggesting an intrinsic pacemaker ability

that may drive tonic inhibition onto postsynaptic targets. Morphological reconstructions

showed that NPY neurons are stellate cells, and the dendrites of NPY neurons in the

tonotopically-organized central nucleus of the IC cross isofrequency laminae. Immunostaining

confirmed that NPY neurons express NPY, and we therefore hypothesized that NPY signaling

regulates activity in the IC. In crosses between Npy1r^{cre} and Ai14 Cre-reporter mice, we found

that NPY Y$_1$ receptor (Y$_1$R)-expressing neurons were broadly distributed throughout the rostro-

caudal extent of the IC. In whole-cell recordings from Y$_1$R-expressing IC neurons, application of

a high affinity Y_1R agonist led to hyperpolarization of a subset of neurons. Thus, NPY neurons

represent a novel class of inhibitory principal neurons that are well poised to use GABAergic

and NPY signaling to regulate the excitability of circuits in the IC and auditory thalamus.

Introduction

The inferior colliculus (IC) is the hub of the central auditory pathway (Adams, 1979; Cant

and Benson, 2006, 2007), a critical processing center for most aspects of hearing, and an

important site of plasticity after hearing loss (Chambers et al., 2016; Sturm et al., 2017). Despite

its importance in auditory processing, it has proven difficult to identify the neuron classes that

comprise the IC (Peruzzi et al., 2000; Palmer et al., 2013; Beebe et al., 2016). As a result, the

cellular organization and function of neural circuits in the IC remain largely unclear.

The IC contains three main subdivisions: the central nucleus (ICc), dorsal cortex (ICd), and

lateral cortex (IClc) (Morest and Oliver, 1984; Faye-Lund and Osen, 1985). The ICc is

tonotopically organized into isofrequency laminae, and neurons in the ICc are divided in two

broad morphological families: disc-shaped and stellate neurons (Meininger et al., 1986;

Malmierca et al., 1993). Disc-shaped neurons, the majority of neurons, maintain their dendritic

arbors within isofrequency laminae (Oliver and Morest, 1984), while stellate neurons extend

their dendritic arbors across isofrequency laminae and integrate information across sound

frequencies (Oliver et al., 1991). However, there is considerable diversity within the disc-shaped

and stellate groups, suggesting that they contain multiple neuron types (Oliver et al., 1994;

Peruzzi et al., 2000; Ono et al., 2005).

GABAergic neurons represent ~25% of neurons in the IC (Oliver et al., 1994; Merchán et al.,

2005; Beebe et al., 2016). In addition to providing local inhibition, many GABAergic neurons

project to the medial geniculate nucleus (MG, auditory thalamus) (Winer et al., 1996; Peruzzi et

al., 1997). Anatomical studies have subdivided GABAergic neurons into classes based on soma

size and extracellular markers (Ito et al., 2009; Beebe et al., 2016), and neurons in all these

classes can project to the MG (Beebe et al., 2018). However, it is unclear whether these

anatomically defined groups represent neuron classes with distinct functional roles.

In many brain regions, an approach that combines anatomical and physiological analyses

with molecular markers has proven key to identifying functionally distinct neuron types

(Tremblay et al., 2016; Zeng and Sanes, 2017). Using such an approach, we recently identified

vasoactive intestinal peptide (VIP) as a marker for a distinct class of excitatory principal neurons

in the IC (Goyer et al., 2019), but there are currently no known molecular markers for

GABAergic neuron classes in the IC.

NPY is a 36 amino-acid neuropeptide and one of the most abundant peptides in the brain

(Chronwall et al., 1985; O'Donohue et al., 1985). NPY is released from neurons and binds to the

NPY family of $G_{i/o}$-coupled G-protein coupled receptors (Dumont et al., 1998). NPY signaling

modulates synaptic transmission, neuronal excitability, and other physiological processes (Bacci

et al., 2002). Across several brain regions, NPY has proven to be an important molecular marker

for distinct classes of GABAergic neurons (Tong et al., 2008; Polgár et al., 2011). Because

previous studies suggest that NPY (Nakagawa et al., 1995) and NPY Y_1 receptors (Y_1R) are

expressed in the IC (Kishi et al., 2005; Eva et al., 2006), we hypothesized that NPY is a molecular

marker for a class of GABAergic neurons in the IC.

Using a multifaceted approach, we found that NPY is expressed by a distinct population of

GABAergic neurons in the IC. These neurons have a stellate morphology and represent

approximately one-third of the GABAergic neurons in the IC. Among other targets, NPY neurons

project to the ipsilateral and contralateral IC and the MG. In brain slice recordings, NPY neurons

fired spontaneously, suggesting an intrinsic pacemaker ability. Using Npy1r[cre] mice to target

recordings, we found that application of a Y_1R-selective agonist hyperpolarized the membrane

potential of a subset of Y_1R-expressing IC neurons. Thus, our data indicate that NPY neurons

represent a novel class of GABAergic IC principal neurons and that these neurons may use

GABAergic and NPY signaling to regulate the excitability of postsynaptic targets.

Materials and Methods

Animals

All experiments were approved by the University of Michigan Institutional Animal Care

and Use Committee and were in accordance with NIH guidelines for the care and use of

laboratory animals. Animals were kept on a 12-hour day/night cycle with ad libitum access to

food and water. NPY-hrGFP mice were obtained from Jackson Laboratory (stock # 006417; van

den Pol et al., 2009) and were maintained hemizygous for the hrGFP transgene by crossing with

C57BL/6J mice (Jackson Laboratory, stock # 000664). To visualize neurons that express Y_1R,

Npy1r^{cre} mice (B6.Cg-$Npy1r^{tm1.1(cre/GFP)Rpa}$/J, Jackson Laboratory, stock # 030544; Padilla et al.,

2016) were crossed with Ai14 Cre-reporter mice (B6.Cg-$Gt(ROSA)26Sor$ $^{tm14(CAG-tdTomato)Hze}$,

Jackson Laboratory, stock # 007914). All mouse lines were on a C57BL/6J background. Because

C57BL/6J mice are subject to age-related hearing loss due to the $Cdh23^{ahl}$ mutation (Noben-

Trauth et al., 2003), experiments were performed in animals aged P22 – P74, before the onset

of hearing loss symptoms at about 3 months of age.

Analysis of the distribution of NPY neurons

Three NPY-hrGFP mice were used to assess the distribution of NPY neurons in the IC.

Each hrGFP-expressing IC neuron was identified in a series of every third section through the

rostro-caudal extent of one IC. The locations of the labeled neurons were plotted using a

Neurolucida system (MBF Bioscience) attached to a Zeiss Axio Imager.Z2 microscope. IC

subdivisions were determined by examining staining for GAD67 and GlyT2 (Buentello et al.,

2015). Low magnification photomicrographs of the distribution of NPY neurons were collected

using a 5x objective on a Zeiss Axio Imager.Z2 microscope. Neurolucida Explorer was used to

export data to Excel for analysis.

Immunohistochemistry

Mice were deeply anesthetized and perfused transcardially with 0.1 M phosphate-

buffered saline (PBS), pH 7.4, for 1 min and then with a 10% buffered formalin solution

(Millipore Sigma, cat# HT501128) for 15 min. Brains were collected and post-fixed in the same

fixative for 2 hours and cryoprotected overnight at 4°C in 0.1 M PBS containing 20% sucrose.

Brains were cut into 40 µm sections on a vibratome or freezing microtome. Sections were

washed in 0.1 M PBS, and then treated with 10% normal donkey serum (Jackson

ImmunoResearch Laboratories, cat# 017-000-121, West Grove, PA) and 0.3% Triton X-100 for 2

hours. Sections were then incubated for 24 hours at 4 °C in mouse anti-GAD67 (1:1000;

Millipore Sigma, cat# MAB5406), rabbit anti-NeuN (1:500; Millipore Sigma, cat# ABN78), rabbit

Neuro-Chrom pan neuronal marker (1:1000; Millipore-Sigma, cat# ABN2300), rabbit anti-NPY

(1:1000; Peninsula Labs, cat# T-4070), guinea pig anti-GlyT2 (1:2000; Synaptic Systems cat#

272-004) or mouse anti-bNOS (1:1000, Millipore Sigma, cat#: N2280). On the following day,

sections were rinsed in 0.1 M PBS and incubated in Alexa Fluor 647-tagged goat anti-mouse IgG

or Alexa Fluor 750-tagged goat anti-rabbit IgG (1:100, Thermo Fisher, cat# A21235 and A21039),

or Alexa Fluor 647-tagged donkey anti-mouse IgG, Alexa Fluor 647-tagged donkey anti-rabbit

IgG, or Alexa Fluor 594-tagged goat anti-guinea pig IgG (1:500, Thermo Fisher, cat# A-21202, A-

21206 and A-11076) for 1.0 – 1.5 hours at room temperature. Sections were then mounted on

Superfrost Plus microscope slides (Thermo Fisher, cat# 12-550-15) and coverslipped using

Fluoromount-G (SouthernBiotech, cat# 0100–01) or DPX (Sigma-Aldrich cat# 06522). Images

were collected using a 1.30 NA 40x oil-immersion objective or a 1.40 NA 63x oil-immersion

objective on a Leica TCS SP8 laser scanning confocal microscope.

Antibody validation

All antibodies used in this study, except anti-NPY, have been previously validated in the

IC. To label GABAergic neurons, we used the mouse monoclonal anti-GAD67 antibody (Millipore

Sigma, cat# MAB5406). This antibody was raised against the 67 kDa isoform of glutamic acid

decarboxylase (GAD). The manufacturer reports that western blot analysis showed no cross-

reactivity with the 65 kDa isoform of GAD (GAD65). This antibody has been used in several

studies to identify GABAergic neurons in the IC (Ito et al., 2009; Mellott et al., 2014; Beebe et

al., 2016; Goyer et al., 2019). To identify neurons, we performed anti-NeuN staining with a

rabbit polyclonal antibody (Millipore Sigma, cat# ABN78). The manufacturer reports that anti-

NeuN specifically recognizes the DNA-binding, neuron-specific protein NeuN. Previous studies

used this antibody to identify neurons in the IC (Foster et al., 2014; Mellott et al., 2014; Beebe

et al., 2016; Goyer et al., 2019). To define the borders of IC subdivisions, we performed a

double staining using anti-GAD67 and anti-GlyT2 antibodies (Buentello et al., 2015). Guinea pig

anti-GlyT2 antibody (Synaptic Systems, cat# 272 004) was used to identify glycine transporter 2,

which is localized in axons. We also immunostained with anti-brain nitric oxide synthase (bNOS)

to define the borders of the IC subdivisions (Millipore Sigma, cat# N2280). This antibody was

raised against the IgG1 isotype from the NOS-B1 hybridoma. The manufacturer reports that

anti-bNOS reacts specifically with nitric oxide synthase (NOS), derived from brain (bNOS, 150 –

160 kDa). Previous studies used anti-bNOS to delineate the borders of IC subdivisions in guinea

pig and mouse (Coote and Rees, 2008; Keesom et al., 2018). Finally, to label NPY, we used the

polyclonal rabbit anti-NPY antibody (Peninsula Laboratories, cat# T-4070). Pretreatment with

antiserum containing NPY peptide abolished all staining, reinforcing the high specificity of this

antibody (Rowan et al., 1993). This antibody has been successfully used in cortex (Rose et al.,

2009; Milstein et al., 2015) and hippocampus (Ledoux et al., 2009).

156   Analysis of GAD67 staining

Analysis of GAD67 immunostaining was performed as previously reported (Goyer et al.,

2019). Images from representative sections of the IC (n = 2 male P74 mice, three sections per

mouse, one caudal, one medial and one rostral) were collected at 2 μm depth intervals with a

1.30 NA 40x oil-immersion objective on a Leica TCS SP8 laser scanning confocal microscope.

Images were analyzed using Leica LAS X software (version 3.3.0). As previously reported, anti-

GAD67 antibody did not penetrate the entire depth of the tissue sections (Beebe et al., 2016;

Goyer et al., 2019). For this reason, the analysis was restricted to the top 10 – 12 μm of each

section, where the antibody fully penetrated. Within this region, we manually marked every

hrGFP[+] cell body in the left IC. We analyzed the green (hrGFP) and red (Alexa 647) channels

separately, so that labeling in one channel did not influence analysis of the other channel. After

the hrGFP cells were marked, the red and green color channels were merged, and in every

instance where a cell body contained markers for hrGFP, we evaluated if GAD67 was also

present. The number of double-labeled cells was compared to the total number of hrGFP$^+$

neurons to determine the percentage of hrGFP$^+$ neurons that were GAD67$^+$ as well.

Analysis of NeuN staining with design-based stereology

We used a design-based stereology approach to estimate the numbers of hrGFP$^+$ and

NeuN$^+$ neurons in brain sections that were stained with anti-NeuN antibody. This approach has

been previously described and validated in the IC (Goyer et al., 2019). In brief, we collected

systematic random samples using a virtual grid of 370 µm x 370 µm squares that was overlaid

on the IC section. The starting coordinates for the grid were set using the Mersenne Twister

random number generator in Igor Pro 7 or 8 (WaveMetrics Inc). We then collected images of

the coordinates determined by the upper-left intersection of each grid-square that fell over the

IC. Each image consisted of a 184 µm x 184 µm Z-stack collected at 1 µm depth intervals with a

1.40 NA 63x oil immersion objective on a Leica TCS SP8 confocal microscope. Eight to twenty

images were collected per slice. Three slices (caudal, middle and rostral) were analyzed per

mouse (n = 2). Using Neurolucida 360 (MBF Bioscience), we counted neurons using the optical

fractionator approach (West et al., 1991). In this approach, we defined guard zones as ≥2 µm-

thick regions at the top and bottom of the slice and excluded these from subsequent analysis.

Neurons between the guard zones, within a 15 µm-thick region at the center of the slice, were

counted by making a single mark at the top of each cell. Cells crossing the right and top borders

of the image stack were counted, whereas those crossing the left and bottom borders were

not. The red (NeuN) and green (hrGFP) color channels were analyzed separately, so that

labeling in one channel did not influence analysis of the other. In all neurons counted, hrGFP$^+$

cells were also NeuN$^+$ (1097/1097 cells). The total number of double-labeled (hrGFP$^+$/NeuN$^+$)

cells was then compared to the total number of NeuN$^+$ cells. We also collected tile scan images

of each IC section analyzed using a 10x objective and used GAD67 and GlyT2 staining to

determine the border separating the ICc from the IC shell (ICs) regions. The 63x Z-stacks were

aligned to these tile scans using Adobe Photoshop, and counted neurons were assigned to the

ICc or IC shell.

Brain slice preparation

To characterize the intrinsic physiology of NPY neurons and the effect of NPY on

neurons in the IC, whole-cell patch-clamp recordings were performed in NPY-hrGFP (van den

Pol et al., 2009) and Npy1r[cre] mice (Padilla et al., 2016). Both males (n = 22) and females (n = 33)

aged P22 – P50 were used. No differences were observed between animals from different sexes

or ages (linear regression, p>0.05). Mice were deeply anesthetized with isoflurane and then

rapidly decapitated. The brain was removed, and the IC was dissected in ~34 °C artificial

cerebrospinal fluid (ACSF) containing (in mM): 125 NaCl, 12.5 glucose, 25 $NaHCO_3$, 3 KCl, 1.25

NaH_2PO_4, 1.5 $CaCl_2$, 1 $MgSO_4$, 3 sodium pyruvate and 0.40 L-ascorbic acid, bubbled to a pH of

7.4 with 5% CO_2 in 95% O_2. Coronal sections of the IC (200 μm) were cut in ~34 °C ACSF with a

vibrating microtome (VT1200S, Leica Biosystems) and incubated at 34 °C for 30 min in ACSF

bubbled with 5% CO_2 in 95% O_2. After incubation, slices were placed at room temperature for

at least 30 min before being transferred to the recording chamber. All chemicals were obtained

from Thermo Fisher Scientific unless otherwise noted.

Electrophysiological recordings

For recordings, slices were placed into a chamber that was continuously perfused at ~2

ml/min with 34°C oxygenated ACSF. NPY-hrGFP neurons and Y_1R-tdTomato neurons in IC slices

were identified with epifluorescence using a Nikon FN1 microscope. Whole-cell current-clamp

recordings were performed with a BVC-700A patch clamp amplifier (Dagan Corporation). Data

were low-pass filtered at 10 kHz, sampled at 50 kHz with a National Instruments PCIe-6343 data

acquisition board, and acquired using custom software written in IgorPro.

Recording pipettes were pulled from borosilicate glass pipettes (outer diameter 1.5 mm,

inner diameter 0.86 mm, cat# BF150-86-10, Sutter Instrument) using a P-1000 microelectrode

puller (Sutter Instrument). Pipettes were filled with an internal solution containing (in mM): 115

K-gluconate, 7.73 KCl, 0.5 EGTA, 10 HEPES, 10 Na_2phosphocreatine, 4 MgATP, 0.3 NaGTP,

supplemented with 0.1% biocytin (w/v), pH adjusted to 7.3 with KOH and osmolality to 290

mmol/kg with sucrose. Pipettes with resistances of 3.0 - 4.8 MΩ when filled with internal

solution were used for recordings.

Input resistance was assessed by applying a series of 100 ms current steps that

hyperpolarized the membrane potential from just below rest to < −100 mV. For each step

response, the peak and steady-state voltage changes were measured, and these values were

used to generate voltage versus current plots from which the peak (R_{pk}) and steady-state (R_{ss})

input resistances were calculated based on the slope of a linear regression. Membrane time

constant was determined by applying fifty, 300 ms current steps that hyperpolarized the

membrane potential by 2 - 4 mV and taking the median of the time constants obtained by

fitting an exponential function to each response. All membrane potential values were corrected

for the liquid junction potential (-11 mV). Capacitance and series resistance were compensated

using bridge balance. Recordings with series resistance >20 MΩ or changes in series resistance

>15% during the recording period were discarded.

To test if NPY neurons are pacemaker neurons (n = 8), we blocked synaptic inputs using

50 µM D-AP5 (NMDA receptor antagonist, Hello Bio, cat# HB0225) and 10 µM NBQX disodium

salt (AMPA receptor antagonist, Hello Bio, cat# HB0443). All drugs were diluted in standard

ACSF.

To evaluate the effects of Y_1R activation, Y_1R-tdTomato neurons were targeted for

recordings. Since most NPY neurons spontaneously fire (n = 87 out of 121 cells), we included 1

µM tetrodotoxin (TTX, Hello Bio, cat# HB1035) in the ACSF in this experiment to block action

potential mediated release of NPY. In current clamp mode, Y_1R neurons were held for 10 min

before starting the control period recording to limit possible effects of intracellular dialysis.

During these 10 min, R_{pk}, R_{ss} and membrane time constant were collected to monitor the health

of the neuron. At the start of the 5 min control period, we initiated a protocol that recorded the

resting membrane potential (V_{rest}) every 10 seconds. Only neurons with a stable V_{rest} , defined

as V_{rest} drift <3 mV, were included in the analysis. We then applied 500 nM [Leu^{31}, Pro^{34}]-NPY

(Tocris, cat # 1176), a high affinity Y_1R agonist. We used 500 nM [Leu^{31}, Pro^{34}]-NPY based on

previous studies performed in the hypothalamus (Roa and Herbison, 2012) and hippocampus

(Giesbrecht et al., 2010). [Leu^{31}, Pro^{34}]-NPY was diluted in 10 ml of ACSF and bath applied for ~5

min. The recording was then continued for a 20 - 50 min washout period. Only neurons that

had a washout period over 20 min were included in the analysis. Control experiments followed

the same approach, using a vehicle solution (normal ACSF).

To analyze the effect of [Leu^{31}, Pro^{34}]-NPY on the V_{rest} of IC neurons, we compared V_{rest}

during the final two min of the baseline period with the initial 2 min of the drug application

period (see Figure 7B, C, gray bars). The drug application period started 90 s after the end of the

control period to allow time for the drug solution to reach the recording chamber. A cell was

considered sensitive to [Leu^{31}, Pro^{34}]-NPY if it exhibited a >2 mV change in V_{rest} compared to

baseline.

Post hoc reconstructions of morphology and morphology analysis

Neurons were filled with biocytin via the recording pipette. After the recording, an

outside-out patch was formed, by slowly removing the pipette, to allow the cell membrane to

reseal. Brain slices were then fixed overnight in 4% paraformaldehyde (PFA) in 0.1 M phosphate

buffer (PB, pH 7.4) and moved to 0.1 M PBS the next day. Slices were stored at 4 °C for up to

three weeks and then stained using fluorescent biocytin-streptavidin histochemistry. For this,

slices were washed in 0.1 M PBS three times for 10 min (3 × 10 min in PBS), the membrane was

permeabilized in 0.2% Triton X-100 in 0.1 M PBS for 2 hours, washed 3 × 10 min in PBS, and

stained at 4 °C for 24 hours with streptavidin-Alexa Fluor 647 (1:1000). The next day, slices

were washed 3 × 10 min with PBS, post fixed in 4% paraformaldehyde (PFA) in 0.1 M phosphate

buffer (PB, pH 7.4) for 1 hour and then washed 3 × 10 min with PBS. Slices were then mounted

on Superfrost Plus microscope slides in anti-fade media (Fluoromount-G). Images were

obtained with a Leica TCS SP8 laser scanning confocal microscope using a 1.40 NA 63x oil-

immersion objective. Z-stack tile scans were performed to image the soma and dendritic arbor.

Image stacks were imported into Neurolucida 360 (MBF Bioscience), where three-dimensional

reconstructions and quantitative analyses of neuronal morphology were performed.

Intracranial virus injections

To evaluate the axonal projections of NPY neurons, we used the "Cre recombinase dependent on GFP" (Cre-DOG) system to selectively express Cre recombinase in GFP expressing neurons (Tang et al., 2015) in the IC of NPY-hrGFP mice. For this, NPY-hrGFP mice were injected in the right IC with a mixture of three viruses: a Cre-dependent tdTomato virus, rAAV2/1.CAG.LSL.tdTomato (Addgene # 100048; titer = 1.10 x 10^{13} gc/ml), and the two AAVs needed for Cre-DOG, rAAV2/1-EF1a-N-CretrcintG (Addgene # 69570; titer = 8.223 x 10^{13} gc/ml) and rAAV2/1-EF1a-C-CreintG (Addgene # 69571; titer = 8.227 x 10^{13} gc/ml). The latter two of these viruses produce two halves of a split Cre-GFP binding protein that assemble in the presence of GFP to produce functional Cre recombinase.

Surgeries were performed on males and females, aged P21 – P24, using standard aseptic techniques. Mice were deeply anesthetized with isoflurane (2 – 3%) and their body temperature maintained with a homeothermic heating pad. The analgesic carprofen (5 mg/kg, CarproJect, Henry Schein Animal Health) was injected subcutaneously. The scalp was shaved and a rostro-caudal incision was made along the midline to expose the skull. A unilateral craniotomy was performed over the right IC, using a micromotor drill (K.1050, Foredom Electric Co.) with a 0.5 mm burr (Fine Science Tools). Injection sites were defined using stereotaxic coordinates relative to the lambda suture, with injection depths relative to the surface of the skull at lambda. Injections were made in two penetrations: penetration 1 – 900 μm caudal, 1000 μm lateral, and 2000 μm deep; penetration 2 – 900 μm caudal, 1250 μm lateral, and 2000 μm deep. One injection of 50 - 60 nl of virus was performed in each penetration, resulting in two virus deposits with a total volume of 100 -120 nl. Viral constructs were injected with a NanoJect III nanoliter injector (Drummond Scientific Company) connected to a MP-285 micromanipulator (Sutter Instruments). Glass injection pipettes were pulled using 1.14 mm outer diameter, 0.53 mm inner diameter capillary glass (cat# 3-000-203-G/X, Drummond Scientific Company) with a P-1000 microelectrode puller (Sutter Instrument). The injector tip was cut and then beveled at 30° with a BV-10 pipette beveller (Sutter Instrument). Injectors

were back-filled with mineral oil and then front-filled with virus. After the injections were

complete, the scalp was sutured, using Ethilon 6–0 (0.7 metric) nylon sutures (Ethicon USA LLC),

and the incision was treated with 0.5 ml 2% Lidocaine hydrochloride jelly (Akorn Inc.). After

recovery from anesthesia, mice were returned to the vivarium and were monitored daily until

the wound healed. After allowing 3 – 4 weeks for protein expression, mice were perfused and

axonal projections were analyzed.

Subdivisions of the IC

We used two different approaches to define the borders between the ICc, ICd, and IClc.

For the stereology experiments and for the analysis of the distribution of NPY neurons,

subdivision borders were defined using a combination of GlyT2 and GAD67 staining (Buentello

et al., 2015). To identify the subdivisions in which reconstructed neurons were located, we used

bNOS staining (Coote and Rees, 2008). Both methods have been previously validated in the IC,

and we did not observe any obvious difference between the subdivision borders obtained with

these approaches.

Analysis of axonal projections

Axonal projections of NPY neurons were analyzed in 11 NPY-hrGFP mice that received

intracranial injections of Cre-DOG and FLEX-tdTomato viruses. Neurons expressing tdTomato

were present in one IC in all but one case. Injection sites were characterized by neurons filled

with tdTomato that also expressed hrGFP. In each case, the brain was examined for tdTomato-

expressing axons and boutons, which were assumed to originate from NPY neurons in the

injected IC. In some cases, the injection site encroached on the ventral IC border, but

tdTomato-expressing axons and boutons were present in substantial numbers in each of the

major projection targets reported here in every case, including those where the injection site

was restricted to the IC. Staining for NeuN and Neuro-Chrom was used to identify target nuclei.

Low magnification images of the injection site were taken using a 5x objective on a Zeiss

AxioImager.Z2 microscope. High magnification images of neurons at the injection site and of

tdTomato-labeled axons and boutons were collected as z-stacks using a 63x oil-immersion

objective (NA = 1.4) on a Zeiss Axio Imager.Z2 microscope using structured illumination

(Apotome 2, with 0.5 µm z-step interval) to provide optical sectioning. Presented images are

maximum projections of collected stacks.

Statistics

All statistical analyses were performed in Igor Pro. Data are presented as mean ± SD.

The statistical tests used for each experiment are indicated in the results. Results were

considered significant when P < 0.05.

Results

NPY neurons are distributed throughout the main subdivisions of the IC

To target experiments to NPY-expressing neurons, we obtained the NPY-hrGFP mouse

line, in which hrGFP expression is driven by the NPY promoter (van den Pol et al., 2009). Figure

1 shows that hrGFP-labeled cells are present throughout the rostro-caudal extent of the IC in

these mice. We performed immunostaining against the NPY peptide to determine whether this

mouse line selectively labels NPY-expressing neurons in the IC. In brain slices from two P58

males, we counted hrGFP$^+$ neurons in one caudal, one middle and one rostral section of the IC

per animal (n = 3205 hrGFP$^+$ neurons, Figure 2A-C). We then evaluated whether each hrGFP$^+$

neuron co-labeled with anti-NPY. We found that 94.7% of hrGFP$^+$ neurons were NPY$^+$, indicating

that the NPY-hrGFP mouse line provides highly selective labeling of NPY-expressing neurons (n

= 3035 hrGFP$^+$/NPY$^+$ neurons and 170 hrGFP$^+$/NPY$^-$ neurons, Table 1).

Figure 1. NPY neurons are distributed throughout the IC. Micrographs of a series of coronal sections through the IC of an NPY-hrGFP mouse. NPY^{+} cells (green) were observed in all sections through the IC, but were more numerous caudally, especially in the dorsomedial aspects of these sections. White arrows indicate NPY[+] axons crossing the IC commissure. Aq- cerebral aqueduct, Cb- cerebellum, SC- superior colliculus. Scale bar = 500 μm.

NPY neurons represent one-third of the GABAergic neurons in the IC

NPY neurons are nearly exclusively GABAergic in the cerebral cortex (Karagiannis et al., 2009), hippocampus (Milner et al., 1997) and hypothalamus (Henry et al., 2015; Marshall et al., 2017). To investigate the neurotransmitter content of NPY neurons in the IC, we performed immunostaining against GAD67, an enzyme essential for GABA synthesis. Using brain slices from two P74 males (Figure 2D-F), we counted hrGFP^{+} neurons in one caudal, one middle and one rostral section of the IC per animal (n = 2673 hrGFP^{+} neurons). We then determined whether each hrGFP[+] neuron co-labeled with anti-GAD67. Out of the 2673 hrGFP^{+} neurons counted, 98.5% were GAD67^{+} (n = 2633 hrGFP[+]/GAD67^{+} neurons and 40 hrGFP[+]/GAD67^{-} neurons, Table 1). We suspect that the absence of GAD67 immunostaining in 40 hrGFP^{+} neurons was due to rare cases of poor antibody penetration or nonspecific expression of hrGFP. Thus, these data indicate that NPY neurons in the IC are GABAergic.

Next, we used design-based stereology to determine the percentage of neurons in the IC that express NPY. For this experiment, we analyzed the ICc and IC shell (ICs = ICd plus IClc) separately, defining the borders between these subdivisions according to the pattern of GAD67 and GlyT2 immunoreactivity (Buentello et al., 2015). Sections from two P54 male NPY-hrGFP

Figure 2. NPY neurons express NPY, are GABAergic, and represent 7.6% of neurons in the ICc. *A, D, G,*
Confocal images of NPY neurons expressing hrGFP in coronal sections of the ICc from NPY-hrGFP mice. ***B,***
Confocal images showing immunostaining for NPY (magenta). ***C,*** Nearly all hrGFP^{+} neurons colocalized
with anti-NPY. ***E,*** Confocal images showing immunostaining for GAD67 (magenta). ***F,*** Nearly all hrGFP[+]
neurons were positive for GAD67 (magenta). ***H,*** Confocal images showing immunostaining for NeuN
(magenta). ***I,*** All NPY neurons were positive for NeuN. White arrowheads indicate NPY neurons labeled by
NPY, GAD67 and NeuN. Scale bar = 30 µm, applies to all images.

mice were stained with anti-NeuN, a neuron-selective antibody previously shown to label most

or all neurons in the IC (Mellott et al., 2014; Beebe et al., 2016). Three sections per mouse were

analyzed: one caudal, one middle, and one rostral. To ensure unbiased counting of neurons, we

applied the optical fractionator method (see Materials and Methods). Sections adjacent to the

Table 1. NPY neurons express NPY and are GABAergic.

Animal	Slice #	# hrGFP$^+$	# NPY$^+$	# hrGFP$^+$ NPY$^-$	% hrGFP$^+$ NPY$^+$
P58 male #1	1 (caudal)	464	436	28	94.0
	2 (middle)	563	537	26	95.4
	3 (rostral)	567	537	30	94.5
	Total	**1594**	**1510**	**84**	**94.7**
P58 male #2	1 (caudal)	496	485	11	97.8
	2 (middle)	602	555	47	92.2
	3 (rostral)	514	486	28	94.6
	Total	**1612**	**1526**	**86**	**94.7**
Grand total		**3206**	**3036**	**170**	**94.7**
Average across two mice					**94.7**

Animal	Slice #	# hrGFP+	# GAD67+	# hrGFP$^+$ GAD67-	% hrGFP$^+$ GAD67$^+$
P74 male #1	1 (caudal)	456	450	6	98.7
	2 (middle)	387	380	7	98.2
	3 (rostral)	328	322	6	98.2
	Total	**1171**	**1152**	**19**	**98.4**
P74 male #2	1 (caudal)	468	462	6	98.7
	2 (middle)	346	337	9	97.4
	3 (rostral)	688	682	6	99.1
	Total	**1502**	**1481**	**21**	**98.6**
Grand total		**2673**	**2633**	**40**	**98.5**
Average across two mice					**98.5**

anti-NeuN sections were immunostained with anti-GAD67 and anti-GlyT2, and IC subdivision

boundaries were determined by comparing these to the adjacent anti-NeuN stained sections.

This analysis showed that NPY neurons represented 7.6% of neurons in the ICc and 7.8% in the

ICs (Table 2). Since previous studies indicate that 20 – 27% of IC neurons are GABAergic (Oliver

et al., 1994; Merchán et al., 2005; Beebe et al., 2016), our results suggest that NPY-expressing

neurons account for approximately one-third (27 – 39%) of GABAergic neurons in the IC.

Table 2. NPY neurons represent 7.6% of neurons in the ICc and 7.8% of neurons in the ICs.

ICc				
Coronal Slice	**P58 Male #1**	**P58 Male #2**	**Per slice pane**	**Grand average**
Caudal	8.0% (60/755) 7 samples	9.4% (15/159) 4 samples	8.2% (75/914) 11 samples	
Medial	7.9% (84/1070) 11 samples	7.0% (90/1278) 9 samples	7.4% (174/2348) 20 samples	
Rostral	7.7% (77/999) 10 samples	7.2% (68/942) 10 samples	7.5% (145/1941) 20 samples	
Per mouse	7.8% (221/2824)	7.3% (173/2379)		7.6% (394/5203)

ICs				
Coronal Slice	**P58 Male #1**	**P58 Male #2**	**Per slice pane**	**Grand average**
Caudal	8.6% (140/1620) 12 samples	6.3% (29/460) 7 samples	8.1% (169/2080) 19 samples	
Medial	6.9% (128/1849) 11 samples	10.4% (76/728) 8 samples	7.9% (204/2577) 19 samples	
Rostral	8.3% (55/668) 12 samples	6.7% (77/1155) 11 samples	7.2% (132/1823) 23 samples	
Per mouse	7.8% (323/4137)	7.8% (182/2343)		7.8% (505/6480)

Together, the above data show that NPY is expressed in a population of GABAergic neurons in the IC, but do not indicate whether this population represents one or several neuron types. To test this, we next analyzed the physiological and morphological properties of NPY neurons.

Most NPY neurons exhibit a sustained firing pattern and have a propensity to spontaneously fire

In acute brain slices, we targeted whole-cell current clamp recordings to NPY-hrGFP

neurons in the Icc, Icd, and Iclc. The location of the recording was noted at the end of the

recording and confirmed during retrieval of neuronal morphology (morphology was recovered

from 50 out of 146 recorded neurons). Because the intrinsic physiology of NPY neurons did not

differ across IC subdivisions, neurons from Icc, Icd, and Iclc are lumped together here. Most NPY

neurons presented moderate voltage sag in response to hyperpolarizing current steps (Figure

3A, left) and a sustained firing pattern in response to depolarizing current steps (Figure 3A,

right). Firing versus current functions were determined with 1 s depolarizing current steps and

showed that NPY neurons had relatively linear input-output functions (Figure 3B). NPY neurons

had an average resting membrane potential of -63.4 ± 3.8 mV (Figure $3C_1$) and exhibited

relatively low expression of I_h current, with voltage sag ratios (steady-state/peak) of 0.89 ± 0.09

and 0.90 ± 0.09 measured from current steps that elicited peak hyperpolarization of -91.1 ± 1.8

mV and -111.0 ± 0.7 mV, respectively (Figure C_2). In addition, NPY neurons exhibited moderate

membrane time constants (15.9 ± 8.6 ms, Figure $3C_3$) and input resistances (R_{pk} = 238.7 ± 124.0

MΩ, R_{ss} = 249.5 ± 167.0 MΩ, Figure $3C_4$) and cells that did not spontaneously fire had low

rheobase values (46.7 ± 25.1 pA, Figure $3C_5$).

Neurons were classified as having a sustained firing pattern if their spike frequency

adaptation ratio (SFA ratio, last interspike interval/first interspike interval) was less than 2 and

as adapting if their SFA ratio was greater than 2 (Peruzzi et al., 2000). We found that 92.4% of

NPY neurons (n = 135 out of 146) had a sustained firing pattern in response to current steps

that elicited ~5 spikes and 89.7% (n = 96 out of 107) had a sustained firing pattern in response

to current steps that elicited ~10 spikes (Figure $3C_6$). Because neurons in the IC exhibit diverse

firing patterns, the finding that nearly all NPY neurons had sustained firing patterns supports

the hypothesis that NPY neurons constitute a distinct neuron class.

Interestingly, 71.9% of NPY neurons fired spontaneously at rest, with a median firing

rate of 0.22 Hz (n = 87 out of 121 neurons, range: 0.01 – 28.06 Hz, mean = 2.70 Hz; Figure 3D).

Figure 3. NPY neurons have a sustained firing pattern and most fire spontaneously. *A,* NPY neurons
exhibited a sustained firing pattern in response to depolarizing current steps and a moderate voltage sag
in response to hyperpolarizing current steps. ***B,*** Firing vs input analysis showed a linear input-output
response, with increasing firing rates in response to increasingly large depolarizing current steps. Example
FI curves from 15 neurons are shown. ***C,*** NPY neurons exhibited moderate intrinsic physiological
properties. ***C₁,*** resting membrane potential, ***C₂,*** voltage sag ratio, ***C₃,*** membrane time constant, ***C₄,*** input
resistance, ***C₅,*** rheobase, and ***C₆,*** spike frequency adaptation ratio. Boxplots show median, 25th and 75th
percentile (box), and 9th and 91st percentile (whiskers). ***D,*** Most NPY neurons spontaneously fired at rest,
and spontaneous firing persisted in 7 out of 8 cells in the presence of synaptic blockers (50 µM D-AP5 and
10 µM NBQX).

To determine whether spontaneous firing was driven by spontaneous synaptic input, we

applied 50 µM D-AP5 and 10 µM NBQX to block AMPA and NMDA receptors and found that

spontaneous firing remained in 7 out of 8 neurons (Figure 3D). These results suggest that NPY

neurons possess an intrinsic pacemaker ability, which may enable NPY neurons to provide tonic

GABAergic and NPY tone in the IC.

NPY neurons have stellate morphology

Using biocytin-streptavidin staining, we reconstructed the morphology of 49 NPY

neurons. Twenty-four of these neurons were located in the ICc, 18 in ICd and 7 in IClc. Figure 4

shows examples of the morphology of NPY neurons in the ICc, ICd and IClc. For comparison

purposes, neurons are oriented as they would appear in a coronal section of the left IC viewed

from a caudal perspective. Visual inspection suggested that NPY neurons had a stellate

morphology in all subdivisions. This is expected for ICd and IClc, where most neurons are

stellate, but in the ICc, only 15 - 20% of neurons have stellate morphology (Oliver and Morest,

1984; Oliver et al., 1991).

We used several quantitative approaches to test whether the dendritic arbors of NPY

neurons in the ICc were consistent with the stellate morphology predicted by visual inspection.

First, we used two-dimensional principal component analysis (PCA) to determine the

orientation and extent of the "length" (first principal direction) and "width" (second principal

direction) axes of each NPY neuron within the coronal plane (Figure 5A). In the mouse, the

isofrequency laminae extend at a ~40° - 50° angle relative to the horizontal plane, and it is

expected that the long axis of disc-shaped neurons extends parallel to the isofrequency laminae

(Stiebler and Ehret, 1985). Our data show that the long axis of NPY neurons in the ICc did not

have a preferred orientation, and only 3 of 24 neurons had a long axis oriented between 30°

and 60° (Figure 5B).

Second, because the dendrites of ICc stellate neurons typically extend across more than

one isofrequency lamina, we measured how far ICc NPY neurons extended their dendrites

perpendicular to a 45° laminar plane. Although the width of isofrequency laminae has not been

Figure 4. Morphology of NPY neurons in the ICc, ICd, and IClc. Representative reconstructions of the morphology of 6 NPY neurons from the ICc (***A***), 6 NPY neurons from the ICd (***B***), and 3 NPY neurons from the IClc (***C***). The gray lines in the ICc representation were drawn to illustrate the approximate orientation and spacing of the isofrequency laminae (45° angle). Solid gray lines are spaced 200 µm apart, dashed lines and solid lines are spaced 100 µm apart.

determined in mice, the distance between laminae in rats presumably provides a conservative

estimate for mice and ranges between 90 µm and 150 µm (Malmierca et al., 1993). We found

that only 4 out of 24 ICc NPY neurons had dendritic arbors that extended <90 µm perpendicular

to the laminar plane, and only 9 out of 24 had dendrites that extended <150 µm perpendicular

Figure 5. NPY neurons are a class of stellate cells. A, Orientation of the dendritic arbors of NPY neurons from the ICc. Combined: all reconstructed neurons from the ICc (n = 24). Blue lines indicate the orientation and length of the longest axis (first principal direction) of each neuron determined using 2D PCA. Caudal, middle and rostral: orientation of dendritic fields separated according to position along the rostro-caudal axis of the ICc. Blue lines indicate longest axis, perpendicular red lines indicate second longest axis (second principal direction) as determined by 2D PCA. **B,** Angular orientation of the long axis for every reconstructed NPY neuron located in the ICc. Angles indicate counter-clockwise rotation relative to the medial-lateral (horizontal) axis. **C,** Extent of dendritic arbors of ICc NPY neurons perpendicular to the plane of the isofrequency laminae (45°). **D,** 3D PCA was used to determine the length and width axes of NPY neurons in the ICc. 92% (22 out of 24) of NPY neurons had a length to

width ratio <3, indicating that they are stellate cells. **E,** Orientation of the dendritic fields of NPY neurons
from the ICd and IClc (combined as ICs). Colors of lines and arrangement of sections are the same as in
A. **F,** Angular orientation of the long axis for every reconstructed NPY neuron located in the ICs (n = 25).
Angles indicate counter-clockwise rotation relative to the medial-lateral (horizontal) axis. **G,** Comparison
of the mean dendritic branch order between NPY neurons in the ICc and ICs. **H,** Comparison of the
length/width ratio between NPY neurons in the ICc and ICs. The dashed line indicates length/width ratio
= 3.

to the laminar plane (Figure 5C). This suggests that the majority of ICc NPY neurons have

dendritic arbors that extend across two or more isofrequency laminae.

Third, we performed three-dimensional principal component analysis on the x, y, z

coordinate set for each ICc NPY neuron to determine the length (extent along the first principal

direction) and width (extent along the second principal direction) of each neuron in three

dimensions. We then calculated the length-to-width ratio of the dendritic arbor. Oliver and

colleagues showed that stellate neurons have a length-to-width ratio <3 and disc-shaped

neurons have a ratio $\geq$ 3 (Oliver et al., 1991). We found that 22 out of 24 NPY neurons (92%) in

the ICc had a length to width ratio < 3, consistent with the hypothesis that NPY neurons are

stellate (Figure 5D). Combined, the above quantitative analyses support the conclusion that ICc

NPY neurons are a class of stellate cells. Because stellate cells represent 15 – 20% of ICc

neurons (Oliver and Morest, 1984; Oliver et al., 1991) and our stereology data (Figure 2 G-I)

indicate that NPY neurons represent 7.6% of neurons in the ICc, NPY neurons represent 38 -

50% of stellate neurons in the ICc.

We also performed two-dimensional principal component analysis on NPY neurons

located in the ICd and IClc (combined as ICs; Figure 5E). As in the ICc, the longest axis of NPY

neurons in the ICs did not exhibit a preferred orientation (Figure 5F). Instead, the long axes of

ICs NPY neurons appeared to run parallel to the outer edge of the IC. Next, we compared the

morphology of NPY neurons in the ICc versus those in the ICs. Interestingly, no difference was

observed in the mean branch order number (ICc: 3.11 ± 1.54; ICs: 3.11 ± 1.04; two-tailed Mann-

Whitney-Wilcoxon test, p = 0.713; Figure 5G) or the dendritic length/width ratio (ICc: 2.16 ±

0.87; ICs: 1.74 ± 0.55; two-tailed Mann-Whitney-Wilcoxon test, p = 0.091; Figure 5H). Thus,

there were no gross morphological differences between NPY neurons in the ICc and ICs.

NPY neurons project to auditory thalamus and other targets

To examine the projections of NPY neurons in the IC, we made injections of AAV-Cre-

DOG and AAV-FLEX-tdTomato into the IC unilaterally in NPY-hrGFP mice. The combination of

these viruses allows Cre expression in cells expressing GFP (here, NPY cells), and subsequent

tdTomato expression in those Cre-expressing cells. In this way, we were able to use tdTomato

expression to identify axons and boutons throughout the brainstem that originate in NPY

neurons of the IC. An example of an injection site is shown in Figure 6A. Note that hrGFP

Figure 6. NPY neurons send GABAergic projections to several long-range targets. *A,* Micrograph of an
rAAV injection site in the IC of an NPY-hrGFP mouse. The IC was injected with rAAVs for the Cre-DOG
system and Cre-dependent tdTomato, resulting in tdTomato expression in transfected NPY neurons
(magenta). hrGFP expression is shown in green. Scale bar = 250 μm. ***B-E,*** Higher magnification micrographs
show tdTomato[+] axons and boutons from IC NPY neurons (magenta) terminating in four brain regions:
ipsilateral IC, contralateral IC, ipsilateral nucleus of the brachium of the IC (NBIC), and ipsilateral medial
geniculate nucleus (MG). Staining for the pan-neuronal markers NeuN and Neuro-Chrom is shown in
yellow; scale bar = 20 μm.

expressing cells are present throughout the IC, but tdTomato expressing cells are present only

at the AAV injection site.

In every case that exhibited tdTomato expressing cells in the IC, clearly labeled axons

with terminal and en passant boutons could be seen in four areas: locally within the ipsilateral

IC (Figure 6B), in the contralateral IC (Figure 6C; this is consistent with the labeling of

commissural axons with hrGFP in Figure 1), in the ipsilateral nucleus of the brachium of the IC

(NBIC; Figure 6D), and in the ipsilateral medial geniculate nucleus (MG; Figure 6E). Fields of

axons and terminals in the IC and the MG were spread widely, without any apparent preference

for a specific subdivision. Less frequently, fewer axons with boutons were observed in other

targets, for example the contralateral NBIC and MG, the ipsilateral cuneiform nucleus, the

ipsilateral superior olivary complex, and the periaqueductal gray. These targets are largely

consistent with previous reports of projections of GABAergic IC cells.

NPY hyperpolarizes a subset of Y_1R-expressing neurons in the IC

Previous studies suggest that Y_1Rs are expressed in the IC (Kishi et al., 2005; Eva et al.,

2006), but it is not known whether NPY signaling affects the IC. To test whether NPY signaling

influences the excitability of IC neurons, we used Npy1r^{cre} x Ai14 mice to label Y_1R-expressing

neurons with the fluorescent protein tdTomato (Padilla et al., 2016). We found that Y_1R

neurons are broadly distributed in the IC (Figure 7A). Interestingly, we also observed that blood

vessels in the IC were lined with non-neuronal cells expressing Y_1R (Figure 7A green arrows).

Because NPY induces vasoconstriction in other brain regions (Bao et al., 1997; Cauli et al.,

2004), NPY signaling might regulate blood flow in the IC, an important hypothesis we will

pursue in a future study.

To test how activation of Y_1Rs affects excitability, we targeted current clamp recordings

to Y_1R neurons while including 1 µM tetrodotoxin in the bath to block endogenous release of

NPY from spontaneously active NPY neurons. We found that bath application of [Leu[31], Pro[34]]-

NPY, a high-affinity Y_1R agonist, hyperpolarized V_{rest} in 50% of Y_1R neurons by an average of -6.2

± 4.3 mV (n = 5 out of 10 neurons; two-tailed paired t-test comparing baseline V_{rest} to treatment

Figure 7. [Leu^{31}, Pro^{34}]-NPY hyperpolarized V_{rest} in a subset of Y$_1$R-expressing IC neurons. *A*, Left, tile scan image showing the distribution of tdTomato expression in the IC of an Npy1r^{cre} x Ai14 mouse. Right, higher magnification image. Magenta arrowheads indicate cell bodies of neurons that are tdTomato$^+$, indicating that they express Y$_1$R. Green arrows indicate blood vessels lined with tdTomato$^+$ cells, reflecting the likelihood that IC blood vessels also respond to NPY. ***B, C,*** Graphs showing 15 min of recording from tdTomato$^+$ Y$_1$R neurons. Gray bars indicate periods in which baseline (left bar) and treatment (right bar) measurements were made. ***B,*** Application of 500 nM [Leu^{31}, Pro^{34}]-NPY led to a hyperpolarizing shift in the V_{rest} of 50% of tdTomato$^+$ neurons (-6.2 ± 4.3 mV change; n = 5 out of 10 neurons; two-tailed paired t-test comparing baseline V_{rest} to treatment V_{rest}, p = 0.032). ***C,*** Vehicle had no effect on the membrane potential of tdTomato$^+$ neurons (0.2 ± 1.3 mV change; n = 5; two-tailed paired t-test, p = 0.694). ***D,*** Boxplot showing the change in V_{rest} between vehicle- and [Leu^{31}, Pro^{34}]-NPY-treated neurons (two-tailed Mann-Whitney-Wilcoxon test, p = 0.007). Boxplots show median, 25th and 75th percentile (box), and 9th and 91st percentile (whiskers).

V_{rest}, p = 0.032; Figure 7B). A neuron was considered sensitive to [Leu^{31}, Pro^{34}]-NPY when V_{rest}

hyperpolarized more than 2 mV below baseline in the presence of the drug. Neurons that did

not respond to the drug exhibited a -0.5 ± 0.8 mV change in V_{rest}, which was not significantly

different from the baseline V_{rest} (n= 5 out of 10 neurons; two-tailed paired t-test, p = 0.248). In

control experiments performed with a vehicle solution, there was no significant change in V_{rest}

between the baseline and vehicle periods (n = 5, 0.2 ± 1.3 mV; two-tailed paired t-test, p =

0.694; Figure 7C). Neurons that responded to [Leu^{31}, Pro^{34}]-NPY exhibited a significantly

different change in V_{rest} compared to neurons treated with vehicle (two-tailed Mann-Whitney-

Wilcoxon test, p = 0.007; Figure 7D). After a 30 – 50 minute wash-out period, the V_{rest} of all cells

that responded to [Leu^{31}, Pro^{34}]-NPY returned to baseline (n = 5, two-tailed paired t-test, p =

0.333; data not shown). Together, these results show that NPY signaling provides a novel

mechanism to dampen excitability in the IC. Since NPY neurons express NPY (Figure 2 A-C) and

provide bouton-laden axons to the local and contralateral IC (Figure 6), it is likely that at least

some NPY signaling in the IC originates from local NPY neurons.

Discussion

Using a multifaceted approach, we have identified NPY neurons as a novel class of

inhibitory principal neurons in the IC. NPY neurons are GABAergic, produce NPY, are present in

the three main subdivisions of the IC, and have a stellate morphology, with dendrites that cross

the isofrequency laminae of the ICc. Based on stereological analysis, we conclude that NPY

neurons account for approximately one-third of GABAergic neurons in the IC and 38 – 50% of

stellate neurons in the ICc. NPY neurons exhibited pacemaker activity, suggesting they might

tonically release GABA and NPY onto their postsynaptic targets. Viral tract tracing showed that

NPY neurons project to several targets, including the ipsilateral MG. We tested the effects of

NPY signaling in the IC and found the first evidence that NPY signaling can regulate the

membrane excitability of IC neurons. Thus, NPY neurons represent a major new class of IC

inhibitory neurons that may use both GABAergic and NPY signaling to regulate the activity of

neural circuits in the IC, NBIC and MG.

NPY neurons are a distinct class of IC neurons

The identification of fundamental neuron classes is essential for understanding the

organization and function of neural circuits in the brain. Previous studies showed that the IC

contains a large and diverse population of GABAergic neurons. These efforts relied on neuronal

morphology (Oliver et al., 1994), in vivo physiology (Ono et al., 2017), neurochemical and

extracellular markers (Ito et al., 2009; Beebe et al., 2016) or a combination of morphology,

physiology and neurotransmitter content (Ono et al., 2005). However, these studies yielded

different results, and it is unclear whether the types of GABAergic neurons they defined

represent functionally meaningful groups. In addition, because the neuron groups identified in

these studies lack molecular markers, they are not readily accessible for targeted manipulations

with optogenetics and other molecular tools.

In recent years, it has become clear that combining molecular markers with

physiological and anatomical parameters is key to identifying functionally relevant classes of

neurons (Tremblay et al., 2016; Zeng and Sanes, 2017). Through such an approach, we

identified NPY neurons as a distinct subtype of GABAergic stellate neurons. NPY neurons are

selectively labeled in NPY-hrGFP mice, and they share a common neurotransmitter content,

neuropeptide expression, stellate morphology, and similar physiological features. Ninety-nine

percent of NPY neurons immunostained for GAD67, and 95% immunostained for NPY. Ninety-

two percent of ICc NPY neurons met the criteria for stellate morphology. In our recordings, 90 –

92% of NPY neurons had a sustained firing pattern. Previously, Ono and colleagues reported

that GAD67^{+} neurons exhibited four firing patterns: regular sustained, buildup/pauser, and

transient with or without after-depolarizations (Ono et al., 2005). Our results show that NPY

neurons fit into the regular sustained category, which included 43% of GAD67^{+} neurons in the

Ono study.

We observed that most NPY neurons spontaneously fired at rest. Interestingly,

spontaneous firing has also been reported for NPY neurons in hypothalamus (Top et al., 2004),

striatum (Partridge et al., 2009) and amygdala (Song et al., 2016). This suggests that NPY

neurons in multiple brain regions might preferentially express a common suite of ion channels

that promotes spontaneous firing. The spontaneous firing of NPY neurons in the IC indicates

that NPY neurons in vivo might provide a constant tone of GABA and NPY to the IC, NBIC, and

MG. This inhibitory tone might be rapidly upregulated or downregulated by inputs that

modulate that activity levels of NPY neurons, providing a mechanism for NPY neurons to

increase or decrease the overall excitability in postsynaptic brain regions.

NPY neurons are GABAergic projection neurons

We found that NPY neurons project to several targets including the contralateral IC, the

ipsilateral NBIC and the ipsilateral MG. In most brain regions, GABAergic neurons are

interneurons (Tremblay et al., 2016), but the IC contains a large number of GABAergic neurons

that project to MG and other targets including NBIC and superior colliculus (Winer et al., 1996;

Peruzzi et al., 1997; Ito et al., 2009; Mellott et al., 2014, 2018; Beebe et al., 2018). The

functional roles of these inhibitory projections are largely unknown. The inhibitory projection to

MG is particularly interesting because other regions of primary sensory thalamus do not receive

ascending inhibitory input (Halassa and Acsády, 2016). Since NPY neurons are molecularly

identifiable, it is now possible to use optogenetics and other targeted approaches to test how a

specific inhibitory projection from the IC shapes activity in the MG and NBIC.

NPY neurons likely use both GABA and NPY signaling to regulate activity in the IC and MG

NPY is widely expressed in the central nervous system of mammals and is a potent

modulator of neuronal activity. Previous brain-wide screening studies suggested that NPY and

Y_1Rs are expressed in the IC (Widdowson, 1993; Kishi et al., 2005; Eva et al., 2006), but the

experiments conducted here are the first to characterize IC neurons that synthesize NPY and to

test the effects of NPY signaling on IC neurons. Y_1Rs tend to inhibit neurons when activated

(Giesbrecht et al., 2010). Consistent with this, we found that application of [Leu^{31}, Pro^{34}]-NPY

hyperpolarized V_{rest} in 50% of the tested Y_1R-expressing neurons. It has been shown that Y_1R

activation can induce opening of G protein-coupled inwardly rectifying potassium (G_{IRK})

channels, suggesting that G_{IRK} activation might mediate the effects we observed in the IC (Sun

et al., 2001; Paredes et al., 2003; Fu et al., 2004; Sosulina et al., 2008), although other

mechanisms are possible (Giesbrecht et al., 2010; Villarroel et al., 2018). The fact that not all IC

Y_1R neurons were sensitive to NPY might be due to transient developmental expression of Cre

in some neurons. Alternatively, non-responsive neurons might respond to NPY in ways that we

did not test.

It is likely that IC NPY neurons also use NPY signaling to influence targets beyond the IC.
Y_1R expression has been reported in the MG (Kishi et al., 2005), and we observed tdTomato
expression in a population of MG neurons in Npy1r[cre] x Ai14 mice (data not shown). It is not yet
clear whether NPY receptors are expressed in the NBIC.

Beyond influencing auditory computations, the inhibitory effects of NPY signaling might
serve diverse roles in the IC. In the hippocampus and amygdala, NPY released from GABAergic
NPY neurons acts as an endogenous anticonvulsant (Colmers and Bleakman, 1994; Colmers and
El Bahh, 2003; Noe' et al., 2007). Interestingly, the IC is the main site of initiation of audiogenic
seizures (Ross and Coleman, 2000; Faingold, 2002), and therefore NPY neurons might use NPY
and GABA release to protect against hyperexcitability in the IC. Consistent with this, acoustic
trauma leads to an increase in the expression of brain-derived neurotrophic factor (BDNF) in
the IC (Meltser and Canlon, 2010), and, in other brain regions, it is known that BDNF can
enhance NPY expression (Nawa et al., 1994; Nobuyuki Takei et al., 1996; Wirth et al., 2005).

It has also been shown that the IC plays an important role in auditory task-related
unconditioned fear responses (Muthuraju et al., 2014; Oliveira et al., 2014) and anxiety (Silveira
et al., 1993). In the amygdala (Sajdyk et al., 2008), cortex (Vollmer et al., 2016) and
hippocampus (Li et al., 2017), NPY exhibits anti-anxiolytic properties and regulates behaviors
associated with fear. Thus, NPY neurons in the IC might regulate how IC circuits process
unconditioned fear responses and how they are affected by behavioral states like anxiety.

Interestingly, in the IC of Npy1r[cre] x Ai14 mice we found strong labeling of cells
associated with blood vessels (Figure 7A). Previous studies showed that NPY induces
contraction in cortical microvessels by activating Y_1Rs (Abounader et al., 1999; Cauli et al.,
2004) and plays a similar role in the cochlea (Carlisle et al., 1990). Since the IC is reported to be
the most metabolically active brain region and to have the highest rate of blood flow in the
brain (Sokoloff et al., 1977; Gross et al., 1987; Zeller et al., 1997), IC NPY neurons might play a
critical role regulating energy homeostasis in the IC.

NPY and VIP neurons account for more than half of ICc stellate neurons

ICc stellate neurons have long been a subject of speculation, but their functional roles are almost

completely unknown due to lack of an approach to target or selectively manipulate them during

in vivo recordings or behavioral testing. We showed here that it is possible to selectively express

exogenous proteins in IC NPY neurons (Figure 6). In addition, we recently identified VIP neurons

as a novel class of glutamatergic stellate cells in the IC, and VIP neurons can be selectively

manipulated in VIP-IRES-Cre mice (Goyer et al., 2019). VIP and NPY neurons represent 3.5% and

7.6% of neurons in the ICc, respectively, and combined they account for approximately 55 – 74%

of ICc stellate neurons. Thus, for the first time, we are positioned to determine the functional

roles of excitatory and inhibitory stellate neurons that together represent the majority of ICc

stellate neurons.

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forward. Nature Reviews Neuroscience 18:530–546.

 Key Resources Table

Animals			
Species	**Designation**	**Source**	**Identifiers**
Genetic reagent (*mus musculus*)	NPY-hrGFP	The Jackson Laboratory	JAX:006417
Genetic reagent (*mus musculus*)	Npy1r$^{cre:GFP}$	The Jackson Laboratory	JAX:030544
Genetic reagent (*mus musculus*)	Ai14	The Jackson Laboratory	JAX:007914
Antibodies			
Designation	**Source**	**Identifiers**	**Concentration**
anti-GAD67 (mouse monoclonal)	Millipore	Cat#: MAB5406 RRID: AB_2278725	IHC:1:1000
anti-NeuN (rabbit polyclonal)	Millipore	Cat#: ABN78 RRID:AB_10807945	IHC: 1:500
anti-NPY (rabbit polyclonal)	Peninsula Labs	Cat#: T-4070 RRID: AB_518504	IHC: 1:1000
anti-bNOS (mouse monoclonal)	Sigma-Aldrich	Cat#: N2280 RRID: AB_260754	IHC: 1:1000
anti-GlyT2 (Guinea pig polyclonal)	Synaptic Systems	Cat#: 272 004 RRID: AB_2619998	IHC: 1:2500
anti-rabbit IgG Alexa Fluor 647 (donkey polyclonal)	ThermoFisher	Cat#: A31573 RRID: AB_2536183	IHC: 1:500
anti-mouse IgG Alexa Fluor 647 (donkey polyclonal)	ThermoFisher	Cat#: A-31571 RRID: AB_162542	IHC: 1:500
anti-guinea pig IgG Alexa Fluor 594 (goat polyclonal)	ThermoFisher	Cat#: A-11076 RRID: AB_141930	IHC: 1:500
Chemical compunds/drugs			
Designation	**Source**	**Identifiers**	**Concentration**
D-AP5	Hello Bio	Cat#: HB0225	50 µM
NBQX	Hello Bio	Cat#: HB0443	10 µM
TTX	Hello Bio	Cat#: HB1035	1µM
[Leu31,Pro34] NPY	Tocris	Cat#: 1176	500 nM
Recombinant DNA reagent			
Designation	**Source**		**Identifiers**
rAAV2/1-EF1a-N-CretrcintG	Boston Children's Hospital Viral Core		Addgene # 69570
rAAV2/1-EF1a-C-CreintG	Boston Children's Hospital Viral Core		Addgene # 69571
AAV1.CAG.LSL.tdTomato	University of Pennsylvania Vector Core		Addgene # 100048
Software, algorithm			

Designation	Source	Identifiers
Igor Pro 7 and 8	Wavemetrics	RRID:SCR_000325
Neurolucida	MBF Bioscience	RRID:SCR_001775
Neurolucida 360	MBF Bioscience	RRID:SCR_016788