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Neurotransmitter Diversity In Pre-Synaptic Terminals Located in the Parvicellular Neuroendocrine Paraventricular Nucleus of the Rat and Mouse Hypothalamus

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Abstract

Virtually all rodent neuroendocrine corticotropin-releasing-hormone (CRH) neurons are in the dorsal medial parvicellular (mpd) part of the paraventricular nucleus of the hypothalamus (PVH). They form the final common pathway for adrenocortical stress responses. Their activity is controlled by sets of GABA-, glutamate-, and catecholamine-containing inputs arranged in an interactive pre-motor network. Defining the nature and arrangement of these inputs can help clarify how stressor type and intensity information is conveyed to neuroendocrine neurons. Here we use immunohistochemistry with high-resolution 3-dimensional image analyses to examine the arrangement of single- and co-occurring GABA, glutamate, and catecholamine markers in synaptophysin-defined pre-synaptic terminals in the PVHmpd of unstressed rats and *Crh-IRES-Cre;Ai14* transgenic mice: respectively, vesicular glutamate transporter 2 (VGluT2), vesicular GABA transporter (VGAT), dopamine β -hydroxylase (DBH), and phenylethanolamine N-methyltransferase (PNMT). Just over half of all PVHmpd pre-synaptic terminals contain VGAT, with slightly less containing VGluT2. The vast majority of terminal appositions with mouse CRH neurons occur non-somatically. However, there are significantly more somatic VGAT than VGluT2 appositions. In the rat PVHmpd, about five times as many pre-synaptic terminals contain PNMT than DBH only. However, because epinephrine release has never been detected in the PVH, PNMT terminals may functionally be noradrenergic not adrenergic. PNMT and VGluT2 co-occur in some pre-synaptic terminals indicating the potential for co-transmission of glutamate and norepinephrine. Collectively, these results provide a structural basis for how GABA/glutamate/catecholamine interactions enable adrenocortical responses to fast-onset interoceptive stimuli, and

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CONFLICT OF INTEREST STATEMENT

All of the authors declare that they have no known or potential conflicts of interest.

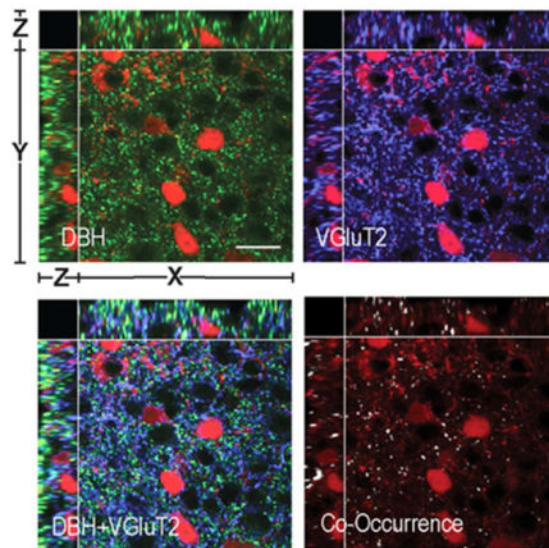
ROLE OF THE AUTHORS

All authors had full access to all the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis. Study concept and design: CSJ, AGW. Acquisition of data: CSJ. Analysis and interpretation of data: CSJ, AGW. Drafting of the manuscript: CSJ, AGW. Critical revision of the manuscript for important intellectual content: CSJ, JSB, AGW. Statistical analysis: CSJ, AGW. Technical and material support: CSJ, AGW, JSB. Obtained funding: AGW, JSB.

more broadly, how combinations of PVH neurotransmitters and neuromodulators interact dynamically to control adrenocortical activity.

Graphical Abstract

The activity of neuroendocrine neurons in the paraventricular nucleus of the hypothalamus (PVH) is controlled by sets of pre-motor inputs. Using immunohistochemistry and confocal image analysis we examine single and co-occurrence of catecholaminergic, glutamatergic, and GABAergic markers in synaptophysin-defined axon terminals in the PVH region that mostly contains CRH neurons. We provide a structural basis for how these transmitter interactions enable adrenocortical responses to fast-onset interoceptive stimuli, and more broadly, how combinations of PVH neurotransmitters and neuromodulators interact dynamically to control adrenocortical activity.



Keywords

Corticotropin-Releasing Hormone; Catecholamine; GABA; Glutamate; Synaptophysin; Pre-motor

INTRODUCTION

The medial parvicellular part, dorsal zone (mpd) of the rat and mouse paraventricular nucleus of the hypothalamus (PVH) mostly contains neuroendocrine motor neurons [Simmons and Swanson, 2009; Biag et al., 2012]. These neurons send axons to the median eminence where their terminals release various neurochemicals into the hypophysial portal vasculature to control hormone release from the anterior pituitary [Lechan et al., 1980; Wiegand & Price, 1980]. Of these, corticotropin-releasing hormone (CRH) neurons are perhaps the most well-studied, primarily because they form the apex of the hypothalamic-pituitary-adrenal (HPA) axis and control its response to all stressors, real or perceived.

CRH neurons are controlled by a complex pre-motor network that uses multiple transmitters and peptidergic modulators to relay information from various parts of the brain to the PVHmpd [Sawchenko and Swanson, 1983; Swanson and Sawchenko, 1983; Sawchenko et al., 1985; Herman and Cullinan, 1997; Herman et al., 2002, 2003; Watts, 2005; Bains et al., 2015]. Substantial evidence supports key roles for GABA and glutamate as primary controllers of neuroendocrine neurons [Boudaba et al., 1996, 1997; Herman et al., 2002; Herman et al., 2004; Kuzmiski et al., 2010; Inoue et al., 2013]. At least half of the pre-synaptic terminals in the medial hypothalamus are GABAergic [Decavel and van den Pol, 1992]. CRH neurons receive direct GABAergic synaptic innervation [Miklós and Kovács, 2002; Flak et al., 2009], and GABA-mediated inhibition is a crucial control element for these neurons [Cole and Sawchenko, 2002; Bali and Kovács, 2003; Wamstecker and Bains, 2010; Bains et al., 2015]. Tonic activation of GABA_A receptors inhibits neuroendocrine CRH activity, while blocking these receptors is sufficient to activate the HPA axis [Cole and Sawchenko, 2002]. In this way, neuroendocrine responses to stressors are dependent on GABA receptor mediated control of Cl⁻ homeostasis [Hewitt et al., 2009; Sarkar et al., 2011]. Similarly, numerous glutamate synapses are found within the PVH [Ziegler and Herman, 2000], and glutamate-containing pre-synaptic terminals appose CRH neurons [Wittmann et al., 2005]. Glutamatergic signaling activates the HPA axis [Ziegler and Herman, 2000; Bains et al., 2015].

A third set of inputs in the neuroendocrine pre-motor network are from hindbrain catecholamine neurons that innervate all subdivisions of the PVH [Swanson et al., 1981; Sawchenko and Swanson, 1982, 1983, Swanson and Sawchenko, 1983; Liposits et al., 1986b; Cunningham and Sawchenko, 1988; Cunningham et al., 1990]. They are defined by the presence or absence of three biosynthetic enzymes: tyrosine hydroxylase (TH), dopamine β-hydroxylase (DBH), and phenylethanolamine N-methyltransferase (PNMT) [Swanson et al., 1981]. PNMT-containing fibers predominate in the parvocellular PVH, with greatest innervation of the PVHmpd [Swanson et al., 1981; Cunningham et al., 1990], while the DBH-only innervation is mostly confined to the periventricular and magnocellular PVH [Swanson et al., 1981; Sawchenko and Swanson, 1982]. Catecholaminergic pre-synaptic terminals appose both magno- and parvocellular PVH neurons [Liposits et al., 1986a], including neuroendocrine CRH neurons [Liposits et al., 1986b].

Catecholaminergic afferents provide stimulus-specific information to the PVHmpd, particularly for acute physiological challenges [Ritter et al., 2011; Guyenet et al., 2013]. These neurons are indispensable for glucocorticoid responses to insulin-induced hypoglycemia, and are major contributors for responses to 2-deoxy-D-glucose (2DG) or GLP-1 agonists [Ritter et al., 2001, 2003; Khan et al., 2007; Lee et al., 2016]. They are also required to maintain the sensitivity of CRH mRNA to elevated levels of corticosterone [Kaminski and Watts, 2012] and to gate metaplasticity at GABA inputs on CRH neurons following an acute stress [Inoue et al., 2013]. However, they are not required for the daily rhythms of corticosterone secretion, or for neuroendocrine responses to psychogenic and other complex stressors [Ritter et al., 2003; Khan et al., 2011; Flak et al., 2014].

PVHmpd afferent systems are highly interactive [Herman et al., 2002, 2003; Watts & Sanchez-Watts, 2002; Watts, 2005; Yang et al., 2007, 2008; Wamstecker and Bains, 2010;

Watts and Khan, 2013]. For example, catecholamine-mediated excitation of neurons within the PVH requires a glutamatergic component located in or around the PVH [Daftary et al., 2000; Herman et al., 2003]. Knowing this, some hindbrain catecholamine neurons contain vesicular glutamate transporter 2 (VGluT2) mRNA [Stornetta et al., 2002] raising the possibility that these neurons have an intrinsic glutamatergic transmitter component [El Mestikawy et al., 2011; DePuy et al., 2013]. It is also important to note that further afferent regulation derives from the many neuropeptides that co-occur in glutamatergic, GABAergic, and catecholaminergic inputs. These include NPY, AgRP, α -MSH, GLP-1, CART, PrRP, and PACAP. [Légrádi et al., 1998; Li et al., 2000; Matsumoto et al., 2000; Khan et al., 2013; Yoon et al., 2013; Zheng et al., 2015].

Our goal here is to elaborate how pre-synaptic terminals in the PVHmpd contribute to the functional anatomy of the pre-motor control network that controls neuroendocrine, and particularly CRH, neuronal activity [Herman et al., 2002, 2003; Watts, 2005; Watts and Khan, 2013]. To do this we analyze the distribution and neurochemical nature of synaptophysin-containing pre-synaptic terminals in the rat and mouse PVHmpd. Synaptophysin is a ubiquitous, vesicular membrane-bound protein [Wiedenmann et al., 1985] and its presence indicates pre-synaptic terminals [Masliah et al., 1990; Thiel, 1993; Flak et al., 2009; Kwon et al., 2011; Chavan et al., 2015]. Our analyses are limited to markers of glutamate (vesicular glutamate transporters 2 and 3; VGluT2, VGluT3) and GABA (vesicular GABA transporter; VGAT) neurotransmission, the catecholamines (PMNT and DBH), and neuropeptide Y (NPY) and agouti-related peptide (AgRP).

We used double- and triple-label immunohistochemistry (IHC) followed by high resolution three dimensional (3D) image analyses of confocal photomicrographs to determine three parameters: the relative abundance of different pre-synaptic terminal phenotypes within the PVHmpd of unstressed rats and mice; any co-occurrence of neurochemical markers within these terminals; and in mice, their appositions with CRH neurons.

MATERIALS AND METHODS

Animals

Nine adult male Sprague Dawley rats (~300g body weight) (Harlan) were housed two- or three-to-a-cage in a climate-controlled room (20–22°C), on a 12h light/dark cycle (lights on 0600h) with unrestricted access to food and water. Animals were acclimated and handled daily for at least five days before any procedures. The brains of six adult female *Crh-IRES-Cre;Ai14* mice [Wamsteeker-Cusulin et al., 2013] were shipped from the University of Calgary and immediately stored at –80°C until further processing. All procedures were approved by the local Institutional Care and Use Committees.

Perfusion and Sectioning

Rats were transcardially perfused with 100mL of cold 0.9% saline, followed by 500mL of cold 4% paraformaldehyde in a sodium borate solution (pH 9.5). Brains were removed and post-fixed overnight at 4°C in a solution of the same fixative, with the addition of 12% (w/v) sucrose. Brains were frozen in hexanes cooled by dry ice before being sectioned coronally at

20 μ m through the rostrocaudal extent of the PVH using a sliding microtome, then stored in a cryoprotectant solution (50% 0.05M sodium-phosphate buffer, 30% ethylene glycol, 20% glycerol) at -20°C until further processing. Seven series were reserved for IHC and one series was reserved for thionin staining to verify cytoarchitecture. Mouse brains were perfused and processed in the same manner but with a lower volume of fixative [Wamsteeker et al., 2010, 2013].

Immunohistochemistry

Sections were washed in tris-buffered saline (TBS, pH 7.4, six rinses for five minutes each at room temperature (RT)), and allowed to incubate for two hours (RT) in a blocking solution of TBS containing 2% (v/v) normal donkey serum (Chemicon, Temecula, CA, S30-100ML) and 0.03% (v/v) Triton X-100 (Sigma Aldrich, 9002-93-1). Sections were incubated in the same blocking solution containing various combinations of primary antibodies (Table 1) for 72 hours at 4°C . Sections were again washed in TBS and incubated in a solution of appropriate and corresponding species-sourced secondary antibodies (Table 2) overnight at 4°C . The sections were washed a final time as before then mounted onto Superfrost slides and allowed to air dry in the dark at RT. Once dry the slides were coverslipped with a solution of 50% TBS/50% glycerol, sealed with clear nail polish, and stored in the dark at 4°C .

The Characterization and Specificity of the Primary Antibodies

Rabbit anti-Agouti-Related Peptide (AgRP) antibody—Specificity was determined through preabsorption of the antiserum with the peptide antigen (Légrádi et al., 1999; Nilsson et al., 2008), following which the authors found no immunolabeling of AgRP in any sections. A single band between 14–21kDa was recognized by this antibody with western blot analysis of total duck brain or hypothalamic tissue extracts (Mirabella, Esposito, Squillacioti, De Luca, & Paino, 2004).

Mouse anti-Dopamine β Hydroxylase (DBH) monoclonal antibody—Western blot analysis recognizes a band of $\sim 70\text{kDa}$ in rat tissue (Bruinstroop et al., 2012). Additionally, immunolabeling is abolished following preadsorption with the immunogen (Rinaman, 2001).

Rabbit anti-DBH antibody—This antibody was characterized using Western blot of purified DBH, and recognizes a triplet at approximately 72–74kDa. The manufacturers tested the antibody in the rat brainstem, cerebellum, and adrenal medulla. The antibody also replicates the known pattern of distribution when used simultaneously with another antibody against DBH (Halliday and McLachlan, 1991).

Sheep anti-Neuropeptide Y (NPY) antibody—Specificity was determined by the manufacturers through the use of radioimmunoassay, and immunolabeling was completely abolished following preabsorption with the immunogen. The antibody showed $\sim 36\%$ cross-reactivity with Peptide YY, though through the use of NPY knockout mice there is no immunolabeling by the antibody in either the forebrain or brainstem, indicating that there is no substantial cross-labeling with Peptide YY (Lee et al., 2013).

Rabbit anti-Phenylethanolamine N-Methyltransferase (PNMT) antibody—This antibody only labels the adrenergic cell groups (C1-3) of the hindbrain (Bohn et al., 1987; Rosin et al., 2006). Additional specificity controls were established through Western blot analysis, where the antibody was shown to recognize a single ~35kDa band, while labeling no bands from the liver, spleen, or cerebral cortex (Bohn et al., 1987).

Goat anti-Synaptophysin antibody—Specificity was verified by the manufacturers through Western blot analysis, where a single band was detected at ~38kDa in human, mouse, and rat brain tissue. Additionally, the generation of transgenic mice wherein the presumed presynaptic B-secretase enzyme is fused with yellow fluorescent protein showed colocalization with the synaptophysin antibody (Buggia-Prévoit et al., 2014).

Rabbit anti-VGAT antibody—Western blot analysis show a single ~60kDa band in rat brain tissue (Chaudhry et al., 1998). Additionally, Western blot analyses of brain tissue from VGAT^{-/-} knockout mice is completely devoid of this band (Martens et al., 2002).

Guinea pig anti-VGluT2 antibody—The manufacturer's Western blot analysis shows a single band at ~52kD. Additionally, when the tissue was incubated with the immunogen, immunolabeling was abolished (Bochorishvili et al., 2012).

Guinea pig anti-VGluT3 antibody—Western blot analysis recognizes a single ~65kDa band (from the manufacturers). Additionally, as reported by both the manufacturer and in the literature (Gabellec et al., 2007), preabsorption with the immunogen peptide abolishes all immunolabeling. Finally, specificity was further tested by the manufacturers on tissue from the rat central nervous system through the use of immunofluorescence histochemistry, where the staining pattern of the antibody was found to correspond with the known expression of VGluT3 mRNA in cell bodies (Fremeau et al., 2002; Gras et al., 2002; Soiza-Reilly and Commons, 2012).

Confocal Microscopy

Images were captured using a Zeiss 700 Laser Scanning Confocal Microscope equipped with a Zeiss AxioImager Z1 camera (Carl Zeiss MicroImaging, Inc., Thorwood, NY), using a 40x (numerical aperture 1.3) oil-corrected objective. An optical zoom of 1.6 was applied using the Zeiss Zen software (Zeiss Zen Black) to result in an image magnified to 64x. Images were taken using a unidirectional line scan at 1024 x 1024 pixels captured on an 8 bit gray scale. Each pixel was scanned twice and the result was averaged to reduce optical noise.

On this scale the intensity of a pixel is given a value ranging from 0 (black, no fluorescent labeling) to 255 (white, maximal fluorescent labeling). To ensure we captured the full range of fluorescent intensities, as well as to ensure proper alignment of each optical channel, images were first taken using fluorescent beads (15µm in diameter) (FocalCheck Fluorescence Microscope Test Slide #1, Invitrogen, F36909) expressing fluorophores for each laser line used. The lower limit (a black value of 0) was set by blocking the light path completely and verifying that all pixels registered a value of 0. This was adjusted using the *Offset* function in the Zeiss Zen software, and was done for each fluorophore channel used.

To set the upper limit, or saturated value of 255, we used the fluorescent beads. For each fluorophore channel the image was adjusted using the *Master Gain* function to ensure that all labeled pixels registered a value of 255, as the concentrated fluorescent dye in the beads is presumably more intense than labeling in physiological tissue. By ensuring that the camera is capable of capturing every possible value assigned to a pixel, we can analyze and compare different levels of fluorophore intensity.

The Airy unit was set to ~1.0 for all laser channels, allowing for an optimal signal to noise ratio based on the properties of each laser channel. Appropriate emission filters were applied to prevent bleed-through between optical channels. 3D images were constructed by taking images of planes (~1 μm) through the Z axis, with a frequency of ~0.4 μm , throughout the entire depth of the tissue section. Images were analyzed using Volocity software (PerkinElmer, version 6.3, Waltham, MA), operating on a MacBook Pro (Apple, Inc.).

The images for the rat PVHmpd were captured from sections corresponding to Levels 25–26 of the Swanson Rat Brain Atlas [Swanson, 2004]. Images of the mouse PVHmpd corresponded to Levels 61–62 of the Allen Reference Atlas [Dong, 2007; Biag et al., 2012]. Anatomical regions of interest were identified using adjacent Nissl-stained sections (Fig. 1).

Image Adjustments

A number of image corrections were employed before analysis. Images of the fluorescent beads were analyzed to ensure that all channels were properly aligned. At the scale of pre-synaptic terminals, any misalignment of images caused by light properties, external vibrations, or other sources of interference has serious consequences on accurate analysis. If all labeled fluorophores did not align in each dimension (X, Y, and Z), the alignment was corrected using the *Registration Correction* feature in Volocity. Subsequent experimental images were corrected using the same settings. To reduce optical aberrations caused by diffraction, particularly along the Z axis, image deconvolution was achieved through the *Iterative Restoration* feature using the *Measured Point Spread Function* in Volocity. One caveat to these methods is that our ability to resolve smaller appositions in the Z-axis is reduced compared to the X-Y axes because of spherical aberration. However, because our results are reported as relative comparisons and these differences are equally distributed across all samples, we believe they are adequately controlled.

Image Analysis

The Rat and Mouse PVH—Although the relative locations of the functional compartments of the rat and mouse PVH are comparable, the distribution of chemically defined neurons within them is somewhat more homogenous in the rat than the mouse [Simmons and Swanson, 2009; Biag et al., 2012]. In the rat most neuroendocrine CRH neurons are clustered in the PVHmpd between levels 25 and 26 [Swanson, 2004], where they are by far the most predominant peptide phenotype [Simmons and Swanson, 2009; Watts and Khan, 2013]. We therefore limited our analyses to these two levels of the rat PVHmpd (Fig. 1).

The distinct compartmental organization of the rat PVH is much less apparent in the mouse where the different chemical phenotypes are distributed more heterogeneously [Biag et al., 2012]. Most CRH neurons are located between levels 61–62 of the Allen mouse brain atlas [Dong, 2007]. To analyze these neurons in the mouse we used a transgenic line, *Crh-IRES-Cre;Ai14*, wherein all CRH neurons express the tdTomato fluorophore throughout their entirety [Wamsteeker et al., 2013] (Fig. 1). This endogenous labeling enables identification of pre-synaptic terminals that are immediately apposed to CRH neurons.

Fluorophore Colocalization—Dunn et al. [2011] have noted the importance of two components when assessing the colocalization of structures labeled with different fluorescent labels: co-occurrence, where two or more markers label the same physical structure; and correlation, wherein the fluorescent intensities of those co-occurring labels vary in a significantly correlated manner.

To determine the co-occurrence of two labels, the total number of each type of fluorescently-labeled structure was quantified within a defined region of interest in the PVHmpd. We then quantified those structures that were double- and/or triple-labeled, using Volocity software to build a protocol that identified each population of structures based on fluorophore expression. Thresholds were set to reduce aberrant signaling in both the volume and fluorescent-intensity parameters. To be considered a labeled structure a minimum volume threshold of $0.5 \mu\text{m}^3$ was applied, along with a maximum threshold of $5\mu\text{m}^3$ for pre-synaptic terminals. Thresholds for volume were determined through comparisons with previously published methods [Bouyer and Simerly, 2013]. Furthermore, a minimum threshold of two standard deviations above the mean background fluorescent intensity was applied. Soma were defined as appropriately-labeled structures with a volume $100 \mu\text{m}^3$, but $500 \mu\text{m}^3$ (mouse) and quantified through the Volocity software throughout the depth of the tissue. These values were determined to be the most appropriate thresholds to allow for the identification of single soma excluding axons and dendrites.

The *Intersect* function in Volocity was applied to determine the co-occurrence of overlapping populations, wherein structures containing both of the indicated fluorophores were counted as a separate population. To find those structures containing DBH but not PNMT the *Intersect* function was applied followed by the *Exclude Touching* function. This prohibited the quantification of DBH-labeled structures that also contained PNMT, leaving only single-labeled structures in the analysis.

To determine the locations of synaptophysin-labeled axon terminals that were apposed to CRH-positive structures in the mouse PVH we used the subtractive methods developed by Bouyer and Simerly [2013] adjusted to measure appositions per unit of surface area, rather than per unit of volume. We calculated the number appositions made to CRH soma by applying appropriate volume thresholds: $>100 \mu\text{m}^3$ for somatic regions; $99.9 \mu\text{m}^3$ for non-somatic regions. We first determined the number of triple-labeled structures (synaptophysin/VGluT2 or VGAT/CRH) within the entire image. We then calculated appositions with non-somatic regions of CRH neurons by dividing the number of somatic appositions (ie. those made with td-Tom labeled structures) by the total number of appositions from the entire image for each terminal marker. The residual structures were designated non-somatic.

We were unable to identify DBH-immunoreactive (ir) appositions to non-somatic regions because of DBH and synaptophysin primary antibody species incompatibility. However, we used the voxel-to-voxel ('touching structures') approach described by Bouyer & Simerly [2013], to identify DBH-ir appositions with CRH neuronal soma.

Excel Analysis

Measurements from Volocity were transferred to Microsoft Excel for further analysis. The number of quantified structures (single-, double-, and triple-labeled) in each image was normalized to the total number of structures per 1000 μm^3 , using the total measured volume of the tissue. Appositions were normalized to 100 μm^2 of measured CRH neuronal surface area. To determine the percent of co-occurrence of a single population with another population the total number of double-labeled structures was divided by the total number of single-labeled structures measured from each of the populations.

Validation Controls for the Methods of Assessing Co-Occurrence

We performed two sets of controls for the rat PVH to validate the co-occurrence methods.

1. *Theoretical Controls.* Sections containing the rat PVHmpd were co-labeled using two different antibodies against DBH: a mouse anti-DBH antibody and a rabbit anti-DBH antibody. Because these antibodies should theoretically identify the same structures, there should be no significant difference between the percent of mouse anti-DBH-labeled structures that express rabbit anti-DBH and the percent of rabbit anti-DBH-labeled structures that express mouse DBH.
2. *Technical Controls.* Sections containing the rat PVHmpd were co-labeled using a mouse anti-DBH antibody and a rabbit anti-PNMT antibody. The percent co-occurrence of the labeling with each antibody was determined. The organization of catecholamine biosynthetic pathways means that all structures containing PNMT must contain DBH, but not all DBH-labeled structures will contain PNMT [Swanson et al., 1981].

Statistics

Significant differences between group means were determined using Student's t test or a one-way analysis of variance (ANOVA) followed by Tukey's multiple comparison post hoc test, as appropriate (JMP Pro, version 12.1, SAS Institute Inc., Cary, NC). All data are expressed as the mean \pm the standard error of the mean (S.E.M.). A *p* value of 0.05 or less was considered statistically significant for all tests.

RESULTS

Establishing the Validity of the Immunohistochemical and Quantitative Analyses

Theoretical Controls—Figure 2a)–d) shows structures in the rat PVHmpd labeled by the mouse or rabbit antibodies against DBH. We found that over 80% of the structures labeled by the mouse anti-DBH antibody were also labeled by the rabbit anti-DBH antibody (Fig. 2e). A similar percentage of the structures labeled by the rabbit anti-DBH antibody were also labeled by the mouse anti-DBH antibody (Fig. 2e). There was no statistical difference in

the percent of co-occurrence between these two measurements (Fig. 2e), indicating that the methods of calculating percent co-occurrence are theoretically valid.

Technical Controls—‘Adrenergic’ structures contain both DBH and PNMT, while ‘noradrenergic’ structures only contain DBH [Goldstein et al., 1972; Swanson et al., 1981]. When we double-labeled sections with DBH and PNMT antibodies (Fig. 2f–i) we found virtually all PNMT-labeled structures contain DBH (Fig. 2j), but significantly fewer DBH-labeled structures contain PNMT (Fig. 2j). These findings validate our ability to use these antibodies to differentiate between structures with these two different catecholaminergic phenotypes.

Vesicular Glutamate and GABA Transporter Distributions in the Rat PVH Region and their Organization within Terminal Appositions

Figure 3 shows the distribution of synaptophysin (Fig. 3a) VGLuT2-ir (Fig. 3b), VGLuT3-ir (Fig. 3c), and VGAT-ir (Fig. 3d) elements in the rat PVH. To determine if vesicular glutamate and GABA transporters were found only in pre-synaptic terminals in the rat and mouse PVHmpd, we assessed the percent co-occurrence of synaptophysin with VGLuT2 or VGAT (Fig. 4).

Virtually every VGLuT2-ir structure in the rat and mouse PVHmpd also contained synaptophysin (Figs. 4d,h). We also found that approximately half of all synaptophysin-ir terminals in the PVHmpd of both species contained VGLuT2 (Fig. 4j). Only about one third of the VGLuT3-ir structures in the rat PVHmpd co-occurred with synaptophysin, and about 10% of the synaptophysin co-occurred with VGLuT3 (Fig. 4j), indicating that most VGLuT3 is not situated in pre-synaptic terminals (Fig. 4j), which is consistent with previous findings [Seal and Edwards, 2006; Santos, 2009]. We also observed apparent VGLuT3 in some axons and in the soma of some PVHpm magnocellular neurons (Fig. 3b), again consistent with the non-terminal expression of this transporter.

Like VGLuT2, the vast majority of VGAT-ir co-occurred with synaptophysin in both rat and mouse PVHmpd (Fig. 4j) indicating that virtually all VGAT-ir structures are pre-synaptic terminals. Furthermore, over half of all synaptophysin-ir structures in the rat and mouse PVHmpd were also labeled for VGAT (Fig. 4j) indicating substantial GABAergic innervation in both species.

VGLuT2 and VGAT Co-Occur in a Population of Terminals in the PVHmpd

When we analyzed the co-occurrence of VGLuT2 and VGAT in the rat PVHmpd (Fig. 5) we found approximately 8% of synaptophysin-labeled pre-synaptic terminals contained both vesicular transporters (Fig. 5e). The number of these double-labeled pre-synaptic terminals was considerably less than the pre-synaptic terminals single-labeled for either VGLuT2 or VGAT only (Fig. 5e). VGLuT2 and VGAT co-occurrence could not be assessed in the mouse because of species incompatibility with the appropriate antibodies.

Catecholaminergic Innervation Patterns in the Rat PVH

DBH (Fig. 6a) and PNMT (Fig. 6b) are differentially distributed in the rat PVH. PNMT was most abundant in the PVHmpd (Fig. 6c, yellow only), while DBH only (Fig. 6c, blue only) was most concentrated in the PVHpm and PVHmpv. The remaining compartments (PVHdp and PVHpv) contained a mixture of PNMT and DBH only. When we analyzed the number of terminals containing synaptophysin and either DBH-only or PNMT in the PVHmpd we found four times more PNMT than DBH-only terminals (Fig. 6d), which is consistent with previous reports showing that PNMT-containing afferents preferentially innervate the parvicellular parts of the PVH that mediate neuroendocrine responses [Swanson et al., 1981; Cunningham et al., 1990].

Catecholaminergic Axon Terminals Express VGluT2 in the PVHmpd of the Rat and the Mouse

In the rat PVHmpd, approximately 25% of all VGluT2-containing pre-synaptic terminals also expressed catecholaminergic markers (Fig. 7a–d, e–h). However, Figure 7i) shows that there was a much higher incidence of VGluT2 co-occurrence in PNMT-containing (21%) than in DBH-only-containing terminals (4%).

We also analyzed the number of pre-synaptic terminals and the co-occurrence of VGluT2 and DBH in the mouse (Fig. 7j–m). Because the PNMT antibody did not produce specific staining in mouse brain sections, we were unable to differentiate PNMT-ir structures from DBH-only-ir structures. Like the rat PVHmpd, there were significantly more VGluT2-containing pre-synaptic terminals than catecholaminergic pre-synaptic terminals in the female mouse PVHmpd (Fig. 7n). VGluT2 and DBH co-occur in the mouse, but there are significantly fewer double-labeled pre-synaptic terminals than either VGluT2-only or catecholaminergic-only pre-synaptic terminals (Fig. 7n).

Catecholamines and NPY Co-Occur in Axon Terminals in the PVHmpd of the Rat

Figure 8 shows that almost two thirds of the NPY-ir structures co-occur with either DBH or PNMT or both, while the remaining third co-occurs with AgRP. Furthermore, almost all of the structures labeled with AgRP also contained NPY.

Appositions to CRH Neurons in the Mouse PVHmpd

There were large numbers of synaptophysin-labeled axon terminal in the mouse PVHmpd (Fig. 9a–c,m). We found approximately twice the number of somatic VGAT-containing appositions (Fig. 9d–f,n) than either VGluT2- (Fig. 9g–i,n) or DBH-containing (Fig. 9j–l,n) appositions.

We found that approximately 90% of synaptophysin-, VGluT2-, and VGAT-containing appositions were with non-somatic regions rather than somatic regions (Fig. 9o). Although we could not account for the cellular locations of all non-somatic appositions, many VGAT terminals were clearly apposed to what appeared to be dendrites of CRH neurons (Fig. 10).

DISCUSSION

Our goal was to determine the configuration and the degree of co-occurrence of protein markers for glutamate-, GABA-, and catecholamine-containing presynaptic terminals in the rat and mouse PVHmpd. Five main findings are evident (Fig. 11). First, there are somewhat more VGAT- (GABAergic) than VGluT2-containing (glutamatergic) pre-synaptic terminals. Second, about 80% of catecholamine pre-synaptic terminals contain PNMT, the remainder are DBH-only. Third, almost 20% of all VGluT2-containing (glutamatergic) terminals also contain PNMT/DBH or DBH-only, with VGluT2 being five times more prevalent in PNMT- than DBH-only-containing terminals. Fourth, while there are substantially more terminal appositions of all types to non-somatic regions of mouse CRH neurons, there are significantly more somatic GABAergic appositions than either glutamatergic or catecholaminergic appositions. Finally, our results are consistent with the finding that approximately two-thirds of the NPY innervation to neuroendocrine CRH neurons originates from the catecholaminergic cell groups of the hindbrain [Bai et al., 1985; Sawchenko et al., 1985; Füzesi et al., 2007], with the remainder originating from other sources, for example the arcuate and dorsomedial nuclei of the hypothalamus [Bai et al., 1985; Füzesi et al., 2007; Lee et al., 2013]. The NPY innervation originating from the arcuate nucleus also contains AgRP [Hahn et al., 1998] and is GABAergic [Tong et al., 2008].

Glutamate/GABA

Figure 11 shows that presynaptic terminals containing VGluT2 and VGAT account for at least 85% of the total number of synaptophysin-labeled terminals in the rat and mouse PVHmpd. Of these, just over half contain VGAT. These numbers agree with previous electron microscopic (EM) surveys of GABAergic synapses [Decavel and van den Pol, 1990; Miklós and Kovács, 2003, 2011].

GABAergic terminals make direct contact with CRH neurons and use GABA_A receptor signaling to exert strong inhibitory influence over them and the HPA axis [Cole and Sawchenko, 2002; Miklós and Kovács, 2002, 2011; Bali and Kovács, 2003; Cullinan, 2000; Cullinan et al., 2008]. We found twice as many VGAT-containing terminals apposed to CRH neuronal soma than VGluT2- or DBH-containing terminals to help enable this strong inhibitory control.

The second largest group of synaptophysin-labeled terminals contain VGluT2 (Fig. 11). The fact that afferent activation elicits glutamatergic excitatory post-synaptic potentials in parvicellular PVH neurons [Wuarin and Dudek, 1991; Daftary et al., 2000], is consistent with the likelihood that VGluT2 expressed in PVHmpd terminals is associated with pre-synaptic glutamate release [Bellocchio et al., 1998]. The lack of suitable markers has made EM examinations of glutamatergic synapses in the PVHmpd more challenging than for GABA, but the consensus is that glutamate provides a significant innervation to the PVH including CRH neurons [van den Pol et al., 1990; Wittmann et al., 2005]. Our results are consistent with these findings.

To understand how afferent networks control neuroendocrine neurons it is important to know on which parts of these neurons particular inputs form synapses. Electron microscopy (EM)

provides the most detailed level of analysis. These EM studies consistently indicate that the great majority of PVH pre-synaptic terminals are found on dendrites [Decavel and van den Pol, 1990]. Using a CRH reporter mouse [Wamsteeker et al., 2013] we found only about 10% of all synaptophysin-defined terminals apposed to CRH neuronal soma. These findings are consistent with previous EM distribution analyses of somatic versus non-somatic synapses in the rat medial parvocellular PVH [Decavel and van den Pol, 1990; Miklós and Kovács, 2011]. Our percentages are somewhat higher than those reported by Miklós and Kovács (72%) [2011] perhaps because they used colchicine to help visualize CRH neurons. Colchicine can dramatically alter CRH neuronal morphology, including somatic area, and dendritic spine number and density [Rho and Swanson, 1989]. We found that somatically located GABAergic terminals are significantly more abundant than glutamatergic and DBH-containing terminals. The remaining synaptophysin-defined terminals are most likely axo-dendritic, but with no apparent differences between the proportions of GABAergic and glutamatergic terminals. These results are again consistent with GABAergic inhibition being the predominant control component for these neurons.

Although we found some VGluT3/synaptophysin co-occurrence that may correspond to synaptic glutamate release, it was much less frequent than with VGluT2 or VGAT, meaning that VGluT3 has a more widespread non-terminal distribution within PVHmpd neurons than the other two transporters. The apparent expression of VGluT3 in the soma of magnocellular neurons is consistent with this view. VGluT3 is found in pre-synaptic terminals, cell bodies, and dendrites of neurons not traditionally considered glutamatergic, particularly serotonergic and cholinergic neurons [Fremeau et al., 2002; Gras et al., 2002; Herzog et al., 2004; Kljakic et al., 2017; Somogyi et al., 2004; Commons, 2009]. Therefore VGluT3/synaptophysin co-occurrence may be in serotonergic or possibly cholinergic terminals rather than glutamatergic (Fig. 11).

VGluT2 and VGAT Co-occurrence

We found a relatively small number of pre-synaptic terminals containing both VGluT2 and VGAT. Although these populations have generally been considered mutually exclusive it is evident that this is not always the case [Boudaba et al., 1996; Ottem et al., 2004; Zander et al., 2010]. While the primary function of vesicular transporters is to package fast-acting neurotransmitters into synaptic vesicles [McIntire et al., 1997; Chaudhry et al., 1998; Fremeau et al., 2002; Herzog et al., 2001], the presence of the VGluTs does not unambiguously mark synaptically releasable glutamate. When VGluTs are found in the same terminals as other transporters, including VGAT, they can augment the packaging ability of the principal transmitter transporter, a process called vesicular synergy [El Mestikawy et al., 2011; Münster-Wandowski et al., 2016]. Additionally, VGluTs may be involved in the transport of inorganic phosphate into vesicles, independent of glutamate transport [Ni et al., 1994; Aihara et al., 2000; Juge et al., 2006]. Collectively these results mean that PVHmpd pre-synaptic terminals with VGluT2/VGAT co-occurrence are more likely GABAergic than glutamatergic (Fig. 11).

Catecholamines

The locations of catecholaminergic neurons have traditionally been defined by the expression of their biosynthetic enzymes: dopaminergic (A9 etc. groups; TH, but not DBH or PNMT); noradrenergic (A1–A8 groups; TH and DBH, but not PNMT); and adrenergic (C1–C3 groups; all three enzymes). Using this nomenclature, virtually all the catecholaminergic innervation of the PVH originates from two medullary locations: the ventrolateral medulla (VLM) (A1, A1/C1, and C1) and the nucleus of the solitary tract (NTS) (A2 and C2), although some periventricular noradrenergic inputs come from the locus coeruleus (A6) [Cunningham and Sawchenko, 1988]. The A1 group primarily innervates magnocellular neurons; the A2 group innervates both parvi- and magnocellular parts [Swanson et al., 1981; Cunningham and Sawchenko, 1988]; while the C1 and C2 groups target its parvicellular parts [Swanson et al., 1981; Cunningham et al., 1990].

VGLUT2 mRNA is found in the majority of PNMT-expressing cell groups, as well as in some A2 neurons, but its expression appears to be most widespread in C1 neurons [Stornetta et al., 2002]. Guyenet and colleagues have provided compelling evidence that glutamate is the primary ionotropic transmitter in those C1 neurons that project to the spinal cord and hindbrain to control cardiovascular processes [DePuy et al., 2013].

C1 neurons that project to the hypothalamus are found at somewhat different locations in the VLM from those with hindbrain or spinal cord projections [Tucker et al., 1987; Guyenet et al., 2013]. These C1 neurons project heavily to the PVHmpd [Swanson et al., 1981; Cunningham et al., 1990]. They make asymmetric synapses with their targets [Liposits et al., 1986ab], which are structurally similar to the glutamatergic synapses made by C1 neurons in the hindbrain [Sheng and Kim, 2011; DePuy et al., 2013]. We now provide the first evidence that a significant number of DBH/PNMT-containing axon terminals in the PVHmpd also contain VGLUT2, meaning that—like caudally directed C1 neurons—glutamate is probably their primary ionotropic transmitter. Previous reports suggest that catecholaminergic neurons engage glutamatergic interneurons to alter parvicellular neuronal function in the PVH [Daftary et al., 2000; Herman et al., 2003], but our results now indicate that at least a subset of these catecholaminergic inputs can themselves release glutamate.

Are catecholamines released by the terminals of C1 or C2 neurons projecting into the PVH? There is a notable lack of direct evidence for epinephrine and/or norepinephrine release from the axon terminals, dendrites, or other parts of caudally projecting C1 neurons [Guyenet et al., 2013]. The fact that PNMT in the hindbrain and spinal cord does not reliably indicate the presence of releasable epinephrine is consistent with this view [Sved, 1989]. These neurons synthesize but do not store epinephrine in vesicles, most likely because elevated levels of monoamine oxidase quickly degrades non-vesicular epinephrine [Sved, 1989; 1990]. We find approximately five times as many PNMT-containing pre-synaptic terminals in the PVHmpd as those containing only DBH. Conventionally this should mean that these neurons will release epinephrine and norepinephrine, respectively. Therefore, it is very surprising that epinephrine release in the PVH has never been detected by microdialysis or fast-scan cyclic voltammetry after various challenges [Pacak et al., 1995, 1999, 2001; Gerth and Roitman, 2016]. The difference between C1 neurons projecting to the hindbrain/spinal cord and those that project to the hypothalamus may result from how hypothalamic monoamine

oxidase acts on epinephrine and norepinephrine, the different cellular locations of DBH (vesicular) and PNMT (cytoplasmic) [Sved, 1990], and the reported absence of a membrane transporter in most C1 neurons [Lorang et al., 1994; Comer et al., 1998].

A lack of measurable epinephrine release in the PVH has two interpretational consequences. First, DBH- and PNMT-containing terminals in the PVH both release norepinephrine, and therefore both should be considered noradrenergic (Fig. 11). Second, the presence or absence of PNMT may not coincide with different releasable catecholamines. Instead it identifies two anatomically distinct neuronal populations with different neuropeptides, afferents, and functional consequences for post-synaptic PVH neurons. For example, PNMT-containing neurons are preferentially recruited by 2DG-induced glucoprivation [Ritter et al., 1998], and particularly by insulin-induced hypoglycemia, which activates more neurons in the C1 region than in A1, A2, and C2 [Jokiaho et al., 2014].

The post-synaptic effects of norepinephrine on PVH parvicellular neurons are varied and complex. Applying norepinephrine to hypothalamic slices containing the PVH increases the firing rates of most parvicellular neurons using ionotropic glutamate receptor-, $\alpha 1$ adrenoreceptor-, and mitogen-activated protein kinase kinase (MEK)-dependent mechanisms [Daftary et al., 2000; Khan et al., 2011]. Conversely, a minority of neurons are inhibited by norepinephrine using β -adrenoreceptor-dependent mechanism [Daftary et al., 2000]. Our structural results now show that at least part of the glutamatergic component in this effect likely originates directly from mostly C1-originating and PNMT-containing terminals in the PVHmpd.

While the relationship between the mechanisms used by norepinephrine to alter parvicellular neuron excitation and the various neuronal components in the PVHmpd are unknown, one key synaptic mechanism is worth noting. Norepinephrine can regulate GABA synaptic plasticity using a mGluR1-dependent metaplastic effect on the ability of GABA synapses to express LTP [Inoue et al., 2013]. This occurs because the norepinephrine released by a stressor into the PVHmpd increases post-synaptic GABA_A receptor membrane insertion, thereby increasing GABA LTP, and consequently post-synaptic responsiveness to subsequent stressors [Bains et al., 2015].

Conclusions

Our results reveal complex relationships between glutamate, GABA, and norepinephrine in pre-synaptic terminals of the neuroendocrine pre-motor networks in the PVHmpd. Complexity is further increased when we consider the other co-occurring peptides and other signaling systems that are known to be present in PVHmpd afferents. An important functional consideration is that the inability of other studies to detect epinephrine release in the PVH suggests that all catecholamine terminals in the PVHmpd, including those containing PNMT, release norepinephrine. For CRH neurons, the dynamic regulation of their pre-synaptic signaling molecules should significantly increase synaptic flexibility in a way that is well-placed to mediate adaptive CRH neuronal responses to different glucocorticoid environments and stressor exposures [Watts, 2005]. Further investigation is needed to determine how and when this dynamic regulation is mediated—including for

example, whether the various signaling molecules are arranged in a co-release or a co-transmission configuration [Vaaga et al., 2014]—and under what circumstances the various signals are released.

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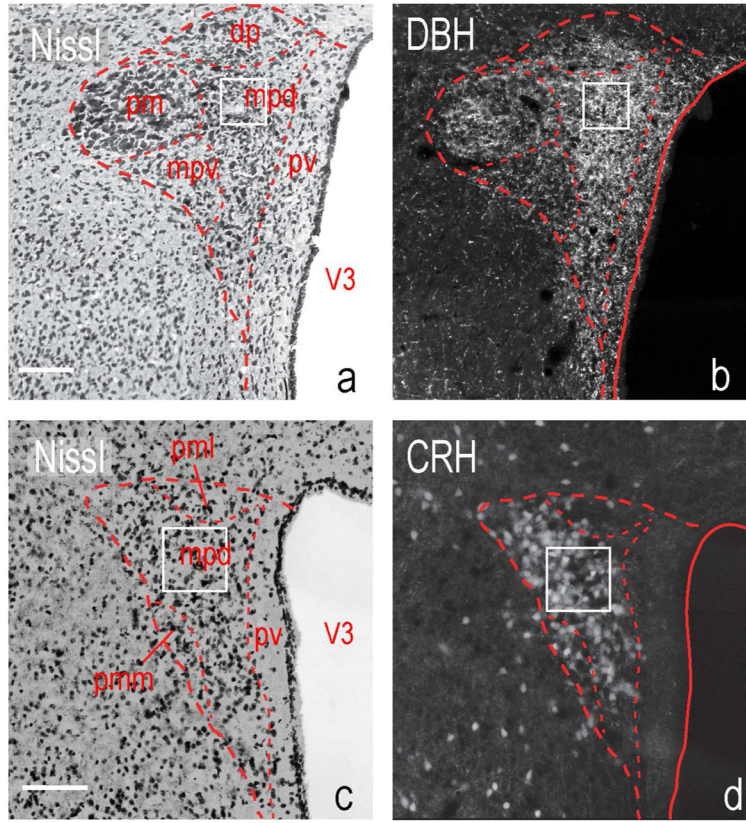


Figure 1. The rat and mouse paraventricular nucleus of the hypothalamus (PVH)
 Thionin-stained sections containing (a) the rat PVH at level 26 of Swanson [2004], and (c) mouse PVH at level P56, Image 61 of Dong [2007] (Also see Biag et al., [2012]). Appropriate serial sections show either immunostaining for DBH (b, rat) or endogenous td-Tomato fluorescence in a *Crh-IRES-Cre;Ai14* mouse (d). White boxes indicate the sampling region for all images. Scale bars indicate 100µm. Abbreviations: dp, dorsal parvicellular part; mpd, medial parvicellular part, dorsal zone; mpv, medial parvicellular part, ventral zone; pm, posterior magnocellular part; pv, periventricular part; V3, third ventricle.

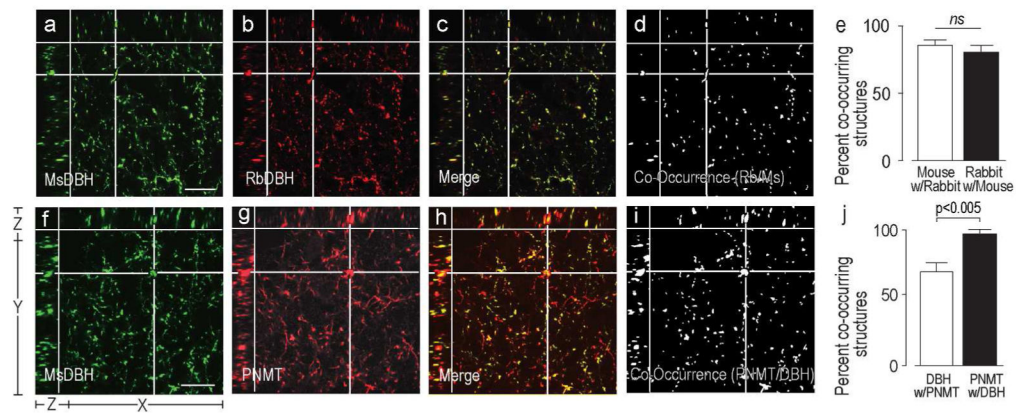


Figure 2. Validation of Image Analytical Methods

Three plane orthogonal views obtained from confocal Z-stacks of part of the rat PVHmpd. The arrangement of the X, Y, and Z planes are shown on panel a. The XY part of each panel shows the uppermost image of each Z-stack. The images show the degree of co-occurrence of structures immunostained with either a mouse monoclonal DBH antibody and a rabbit polyclonal DBH antibody (a–d), or a mouse monoclonal DBH antibody and a rabbit polyclonal PNMT antibody (f–i). Panels a), b), f), and g) show structures stained with the individual antibodies. Panels c) and h) are the respective merged images where single labeled structures are in red or green and co-occurring structures in yellow. Panels d) and i) show any structure that contained labeling from both antibodies (white) extracted from panels c) and h), respectively. The horizontal and vertical white lines in panels a)–d) and f)–i) target a representative double-labeled structures in the X, Y, and Z planes. There is no significant difference between the percent co-occurrence of the structures labeled with the two DBH antibodies (e). However, because all PNMT always co-occurs with DBH but not all DBH occurs with PNMT there was a significant difference between the percent co-occurrence of the two antigens depending on which one was considered the reference antigen (j). Error bars in e) and j) indicate mean + S.E.M. Scale bar represents 20 μ m.

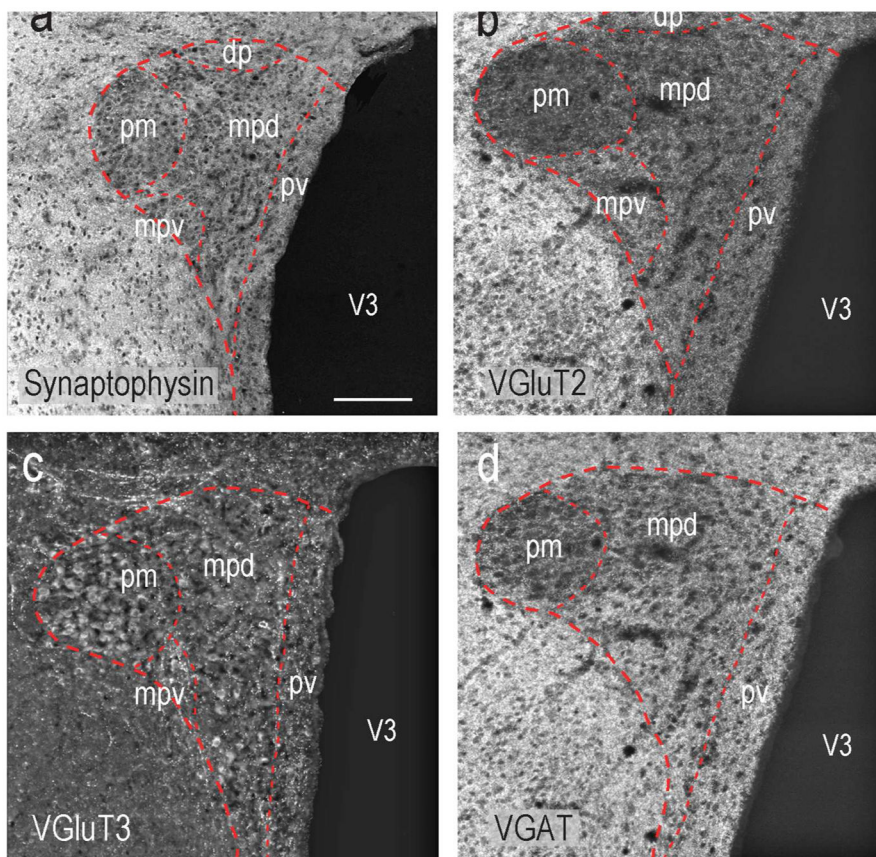


Figure 3. Glutamatergic and GABAergic Innervation Patterns of the rat and mouse PVH
 The locations of synaptophysin (a) VGluT2 (b), VGluT3 (c), and VGAT (d) immunolabeled structures in the rat PVH and adjacent regions. Note that peri-PVH regions have significantly higher levels of VGluT2 and VGAT than the PVH itself. Abbreviations as Figure 2. Scale bar represents 150 μ m.

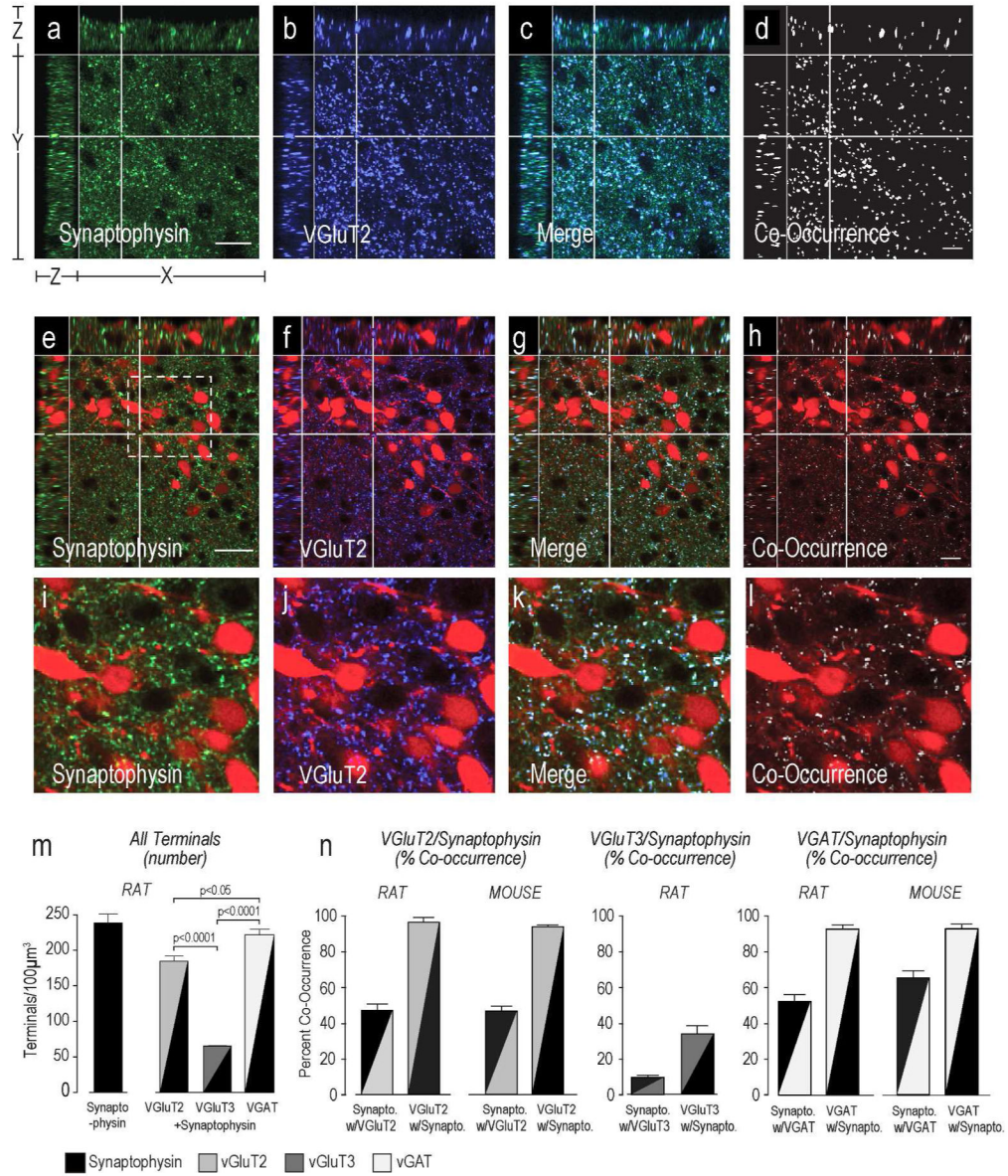


Figure 4. VGLUT2 and VGAT, but not VGLUT3 are Expressed Exclusively in Pre-Synaptic Terminals Synaptophysin (a,e,i), VGLUT2 (b,f,j), and terminals labeled with both molecules (c,d, g,h, and k,l) in the rat (a–d) and mouse (e–l) PVHmpd (see Fig. 1 for orientation). TdTomato-labeled CRH soma are visible in the mouse panels. Equivalent images are not shown for VGAT or VGLUT3. Please see Figure 2 for more information about the general organization of the confocal image panels. Panels i–l show higher magnification images of the area indicated by the dashed line square in e. Scale bar in a) and e) indicates 20µm for panels a – h. Scale bar in i indicates 100µm. All VGLUT2 and VGAT occurs in rat and mouse (female) pre-synaptic terminals (m,n), but not not all VGLUT3 occurs in rat pre-synaptic terminals (m,n). There are significantly more VGAT presynaptic terminals than either Glu transporters, and significantly more VGLUT2 than VGLUT3 terminals (n). Error bars represent the mean + S.E.M.

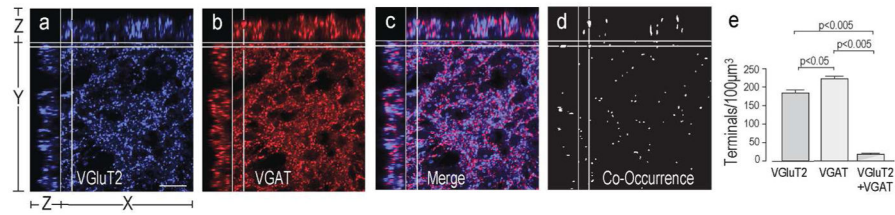


Figure 5. VGlut2 and VGAT Co-Occur in a Population of Terminals in the Rat PVHmpd
 VGlut2 (a) and VGAT (b) co-occur in some terminals (c–e) in the rat PVHmpd. Please see Figure 2 for more information about the general organization of the confocal image panels. There are significantly fewer terminals containing both vesicular transporters than terminals containing only one transporter (e). Scale bar indicates 20μm. Error bars represent mean + S.E.M.

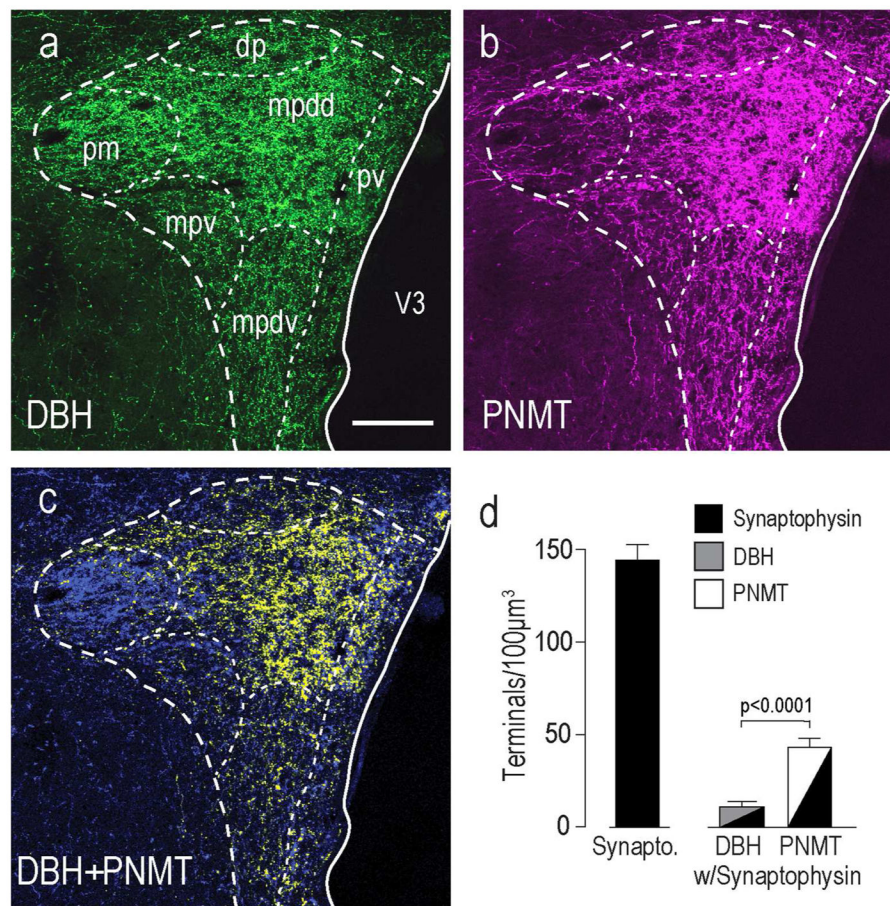


Figure 6. Catecholaminergic Innervation to the PVH

DBH (a) and PNMT (b) immunoreactivity in the rat PVH and its surrounding regions. Panel (c) is a merge of a) and b) to show co-occurring DBH and PNMT structures and DBH-only structures. Within the PVHmpdd there are significantly more PNMT than DBH-only structures (c). Scale bar is 150 μm for all panels. Abbreviations are as Figure 1 except for: mpdd, medial parvicellular part, dorsal zone (dorsal division); mpdv, medial parvicellular part, dorsal zone (ventral division). Panel d) shows that there were significantly more PNMT-containing presynaptic terminals in the PVHmpdv than those containing DBH-only (no confocal images are shown for these results). Error bars indicate mean + S.E.M.

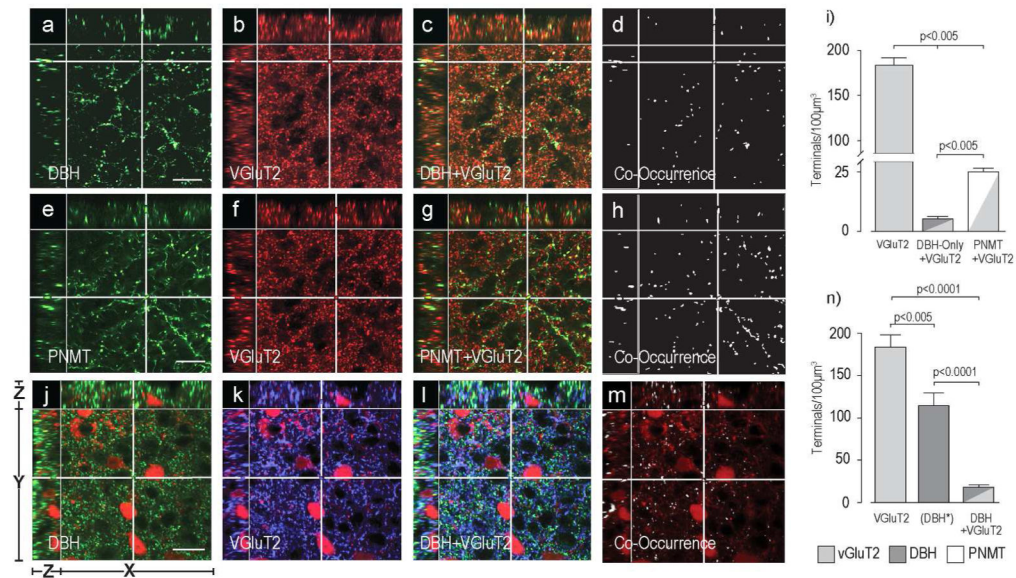


Figure 7. Some Catecholamine Terminals in the PVHmpd of the Rat and Mouse Express VGluT2
 Panels a) – h) show DBH-ir (a–d) or PNMT-ir (e–h) in the rat PVHmpd together with VGluT2-ir (b,f) and its co-occurrence with DBH (c,d) or PNMT (g,h). There are significantly more PNMT than DBH-only terminals that also contain VGluT2 in the rat PVHmpd (i). Please note the change in the scale of the Y axis in panel i). Panels j) – m) show DBH-ir (j) and VGluT2-ir (k) in the female mouse PVHmpd. TdTomato-labeled CRH soma can be seen in panels j) – m). The co-occurrence of DBH and VGluT2 is shown in l) and m). Scale bar indicates 20µm for all panels. Note that in panel n) the (DBH*) data are DBH-immunoreactive structures only. Group data in n) are from female mice. Because of species incompatibility there was no synaptophysin labeling in this experiment to verify that non-VGluT2-ir DBH structures were terminals. Scale bar indicates 20µm for all image panels. Please see Figure 2 for more information about the general organization of the confocal image panels. Error bars indicate mean + S.E.M.

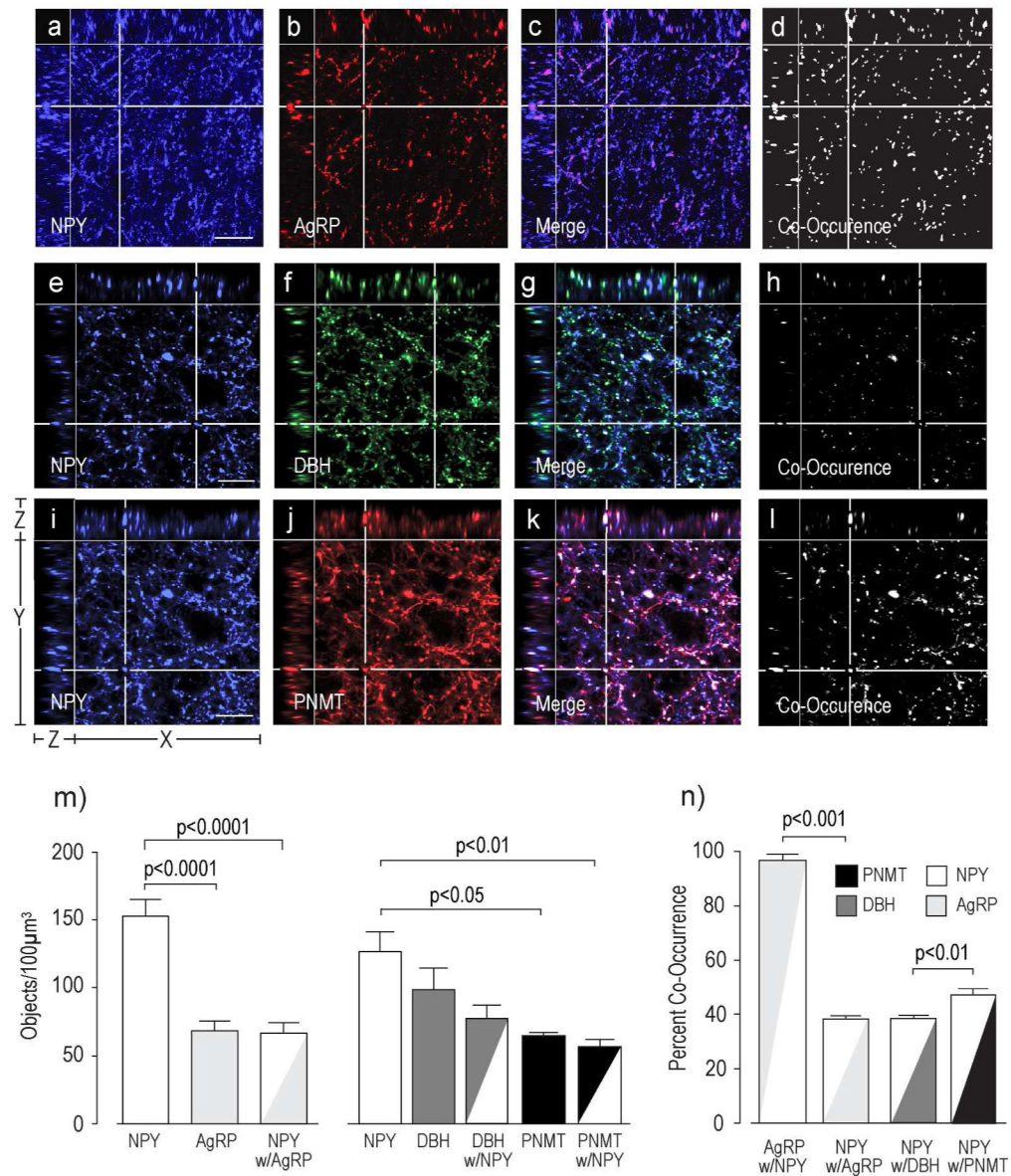


Figure 8. Co-Occurrence of NPY with AgRP, DBH, or PNMT

Panels a) – l) show NPY-ir with AgRP-ir (a–d), NPY with DBH-ir only (e–h), or NPY with PNMT-ir only (i–l) in the rat PVHmpd. Co-occurring structures (c,d, g,h, k,l) were identified and counted (m), together with their percent co-occurrence (n). Please see Figure 2 for more information about the general organization of the confocal image panels. Scale bars indicates 20μm. Error bars indicate mean + S.E.M.

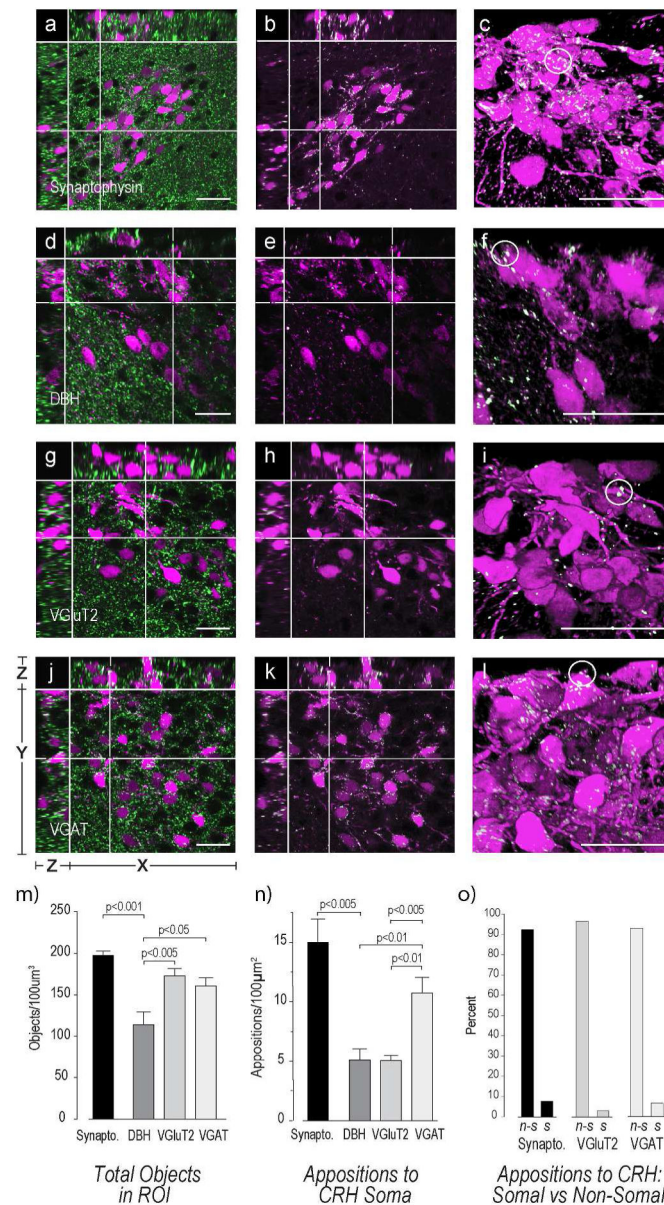


Figure 9. Appositions to CRH Neurons in the Mouse PVHmpd

Female mouse sections containing the PVHmpd were labeled with antibodies against synaptophysin (a–c), DBH (d–f), VGlut2 (g–i), and VGAT (j–l) and analyzed for appositions of each type against the soma of the CRH neuron. Please see Figure 2 for more information about the general organization of the confocal image panels. The white structures in all confocal images except a) d), g), and j) are those that are double-labeled and fulfill the apposition criterion established by Bouyer & Simerly [2013]. See Methods for more details. Images c), f), i), and l) are 3D renderings of the Z-stacks used for images b), e), h), and k), respectively. White circles highlight some somatic appositions. There are significantly more VGAT and VGlut2 terminals than DBH terminals (m). However, VGAT somatic appositions are more numerous than either VGlut2 or DBH (n). There are significantly more appositions to the non-somatic regions of the CRH neuron (o). Group

data in m) – n) are from female mice. Abbreviations for o): n-s, non-somatic; s, somatic.
Scale bars indicate 20µm. Error bars indicate mean + S.E.M.

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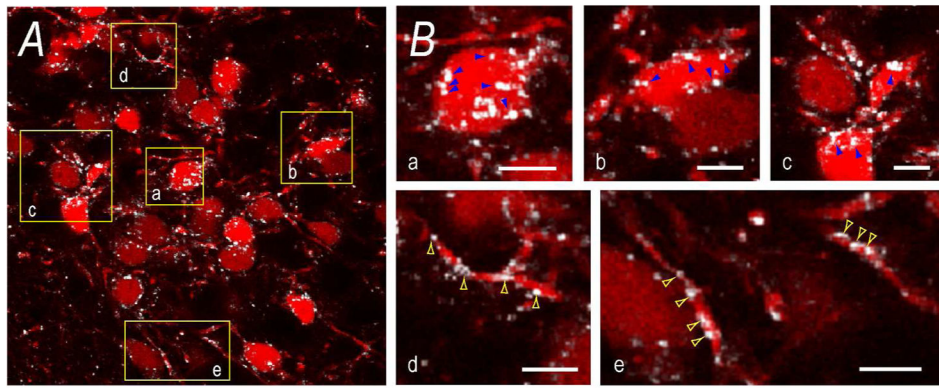


Figure 10. Appositions to CRH Neurons in the Mouse PVHmpd

Panel A) is panel e) from Figure 9. It shows VGAT appositions (white) to TdTomato-labeled CRH soma. To highlight somatic and non-somatic appositions, panel B) shows magnified regions (yellow boxes) from A). Blue arrows in a), b), and c) indicate some VGAT appositions to CRH soma. Yellow arrows in d) and e) indicate some VGAT appositions to non-somatic regions, presumably dendrites. Scale bars represent 20 μ m in A), and 5 μ m in B).

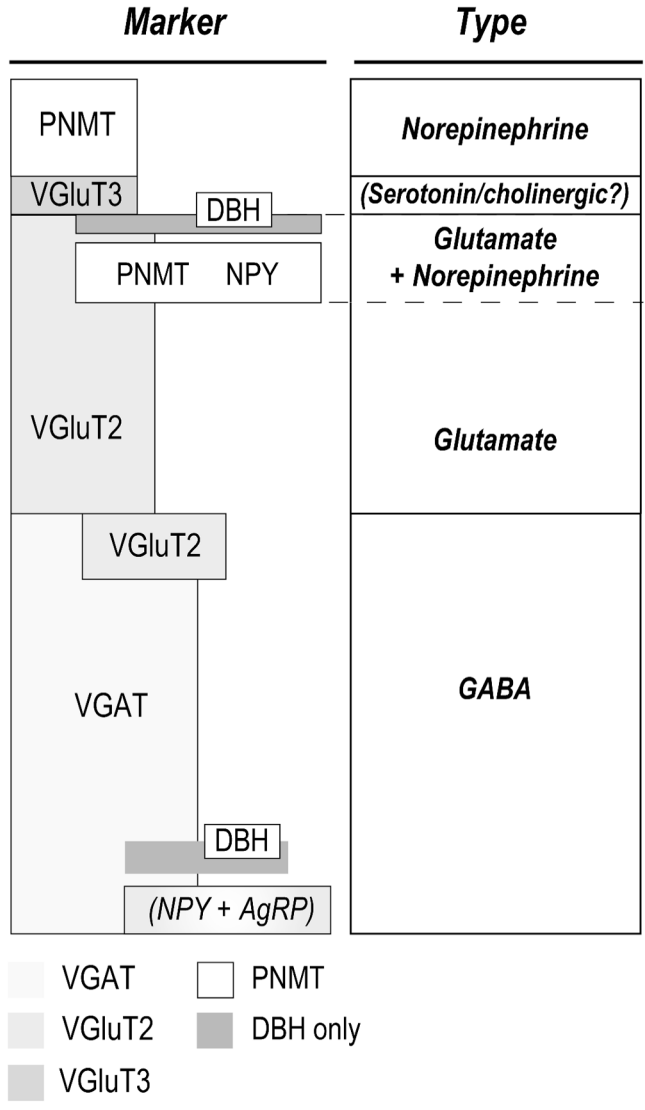


Figure 11. Pre-Synaptic Terminal Diversity in the Rat PVHmpd

Figure 11 summarizes two sets of relationships in pre-synaptic terminals identified in the rat PVHmpd in the absence of stressors:

- 1) **Marker**. The height of each box approximates the proportion of that marker to the total number of synaptophysin-labeled pre-synaptic terminals. Overlapping boxes indicate co-occurrence between two markers. While we did not investigate the co-occurrence of VGAT with NPY, the majority of NPY innervation co-occurs with catecholaminergic (and particularly PNMT-containing) afferents. The remainder originated from hypothalamic sources [Bai et al., 1985; Sawchenko et al., 1985; Füzesi et al., 2007]. NPY originating from the arcuate nucleus co-occurs with AgRP, the majority of which is GABAergic [Tong et al., 2008]. While we did not investigate serotonergic or cholinergic inputs, evidence indicates that serotonergic inputs originate from hindbrain raphe neurons that contain VGlut3 [Gras et al., 2002; Herzog et al., 2004].
- 2) **Type** The presumed major signaling molecule.

Abbreviations: AgRP, agouti-related peptide; DBH, dopamine β -hydroxylase; α MSH, NPY, neuropeptide Y; PNMT, phenylethanolamine N-methyltransferase; VGAT, vesicular GABA transporter; VGlut2, vesicular glutamate transporter 2; VGlut3, vesicular glutamate transporter 3.

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TABLE 1

Primary Antibody	Immunogen	Supplier	Number Catalogue	RRID	Titer
Rabbit anti-Agouti Related Peptide	Synthetic peptide sequence N'- TRSCPMMATGRCYCFANFFR CACPDCCPVQQGSEHLRYCRRSS-C' (Human)	Phoenix Pharmaceuticals	H-003-53	AB_2313908	1:5,000
Mouse anti-Dopamine- β -Hydroxylase	Purified bovine DBH	EMD Millipore	MAB308	AB_2245740	1:20,000
Rabbit anti-Dopamine- β -Hydroxylase	Bovine DBH	ImmunoStar	22806	AB_572229	1:2,000
Sheep anti-Neuropeptide Y	Synthetic NPY peptide conjugated to bovine thyroglobulin	EMD Millipore	AB1583	AB_2236176	1:5,000
Rabbit anti-Phenylethanolamine N-Methyltransferase	Purified PNMT enzyme from rabbit adrenal medulla	Dr. M. Bohn, Northwestern University	N/A	AB_2315181	1:10,000
Goat anti-Synaptophysin	E. coli-derived recombinant human synaptophysin	R&D Systems	AF5555	AB_2198864	1:5,000
Rabbit anti-Vesicular GABA Transporter	Synthetic peptide AEPPVEGDHYQR (aa 75 – 87 in rat) coupled to KLH	Synaptic Systems	131 002	AB_887871	1:20,000
Guinea Pig anti-Vesicular Glutamate Transporter ²	KLH-conjugated linear peptide corresponding to amino acids 501-582 (cytoplasmic C-terminus) of rat VGlut2.	EMD Millipore	AB2251	AB_1587626	1:10,000
Guinea Pig anti-Vesicular Glutamate Transporter ³	Fusion protein amino acids 546-588 (cytoplasmic C-terminus) of rat VGlut3.	EMD Millipore	AB5421	AB_2187832	1:10,000

TABLE 2

Secondary Antibody	Supplier	Catalogue Number	RRID	Titer
AffiniPure Donkey anti-Mouse IgG (H+L) Alexa 488	Jackson ImmunoResearch	715-545-151	AB_2341099	1: 2,000
AffiniPure Donkey anti-Rabbit IgG (H+L) Alexa 488	Jackson ImmunoResearch	711-545-152	AB_2313584	1: 2,000
AffiniPure Donkey anti-Goat IgG (H+L) Cy3	Jackson ImmunoResearch	705-165-147	AB_2307351	1: 2,000
AffiniPure Donkey anti-Rabbit IgG (H+L) Cy3	Jackson ImmunoResearch	711-165-152	AB_2307443	1: 2,000
AffiniPure Donkey anti-Guinea Pig IgG (H+L) Alexa 647	Jackson ImmunoResearch	706-605-148	AB_2340476	1: 2,000
AffiniPure Donkey anti-Sheep IgG (H+L) Alexa 647	Jackson ImmunoResearch	713-605-003	AB_2340750	1: 2,000

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