ORIGINAL ARTICLE

Monoclonal antibody Py recognizes neurofilament heavy chain and is a selective marker for large diameter neurons in the brain

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Abstract Almost 30 years ago, the monoclonal antibody Py was developed to detect pyramidal neurons in the CA3 region of the rat hippocampus. The utility of this antibody quickly expanded when several groups discovered that it could be used to identify very specific populations of neurons in the normal, developing, and diseased or injured central nervous system. Despite this body of literature, the identity of the antigen that the Py antibody recognizes remained elusive. Here, immunoprecipitation experiments from the adult rat cortex identified the Py antigen as neurofilament heavy chain (NF-H). Double immunolabeling of sections through the rat brain using Py and NF-H antibodies confirmed the identity of the Py antigen, and reveal that $Py/NF-H+$ neurons appear to share the feature of being particularly large in diameter. These include the neurons of the gigantocellular reticular formation, pyramidal neurons of layers II/III and V of the cortex, cerebellar Purkinje neurons as well as CA3 pyramidal neurons. Taken together, this finding gives clarity to past work using the monoclonal Py antibody, and immediately expands our

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understanding of the importance of NF-H in neural development, functioning, and disease.

Keywords Monoclonal antibody Py - Neurofilament heavy chain - Large diameter neuron - Neurodegenerative disease

Introduction

In 1989, a study by Woodhams et al. [\(1989](#page-12-0)) detailed the production and characterization of a new antibody that labeled a subset of neurons in the hippocampus, cortex, cerebellum and brain stem. Because the antibody was useful for distinguishing pyramidal neurons from the CA3 region of the hippocampus both in vivo and in vitro, they called the monoclonal IgM antibody ''Py''. The usefulness of this antibody was almost immediately realized, as Raisman and colleagues used it to identify and detail CA3 hippocampal transplants to adult rats (Field et al. [1991](#page-11-0)), and Whittemore and colleagues that the Py antigen was expressed by immortalized stem cells transplanted (and incorporating) into the CA3 region of the rat (Shihabuddin et al. [1996](#page-11-0)). Subsequent work by Brook et al. [\(1997a,](#page-11-0) [1998\)](#page-11-0) found that the antibody stained neurons along the spinal column, most notably motor neurons, and the neurons of Clarke's nucleus. Additional work by Houweling et al. ([1999\)](#page-11-0) found that Py labeled neurons in the developing spinal cord at very early stages of development (i.e., embryonic day 15). The antibody further proved useful for highlighting important changes in anatomical structures of the brain and spinal cord in response to gene alterations, injury or disease models. Py, for example, was used to highlight the cortical neurons affected by the deletion of the mouse-enabled gene, which

displays midline crossing defects in the corpus callosum (Lanier et al. [1999](#page-11-0)). In the red nucleus and Clarke's nucleus, Py highlighted a loss of cellular content after axotomy of the rubrospinal and spinocerebellar tracts, respectively (Brook et al. [1997b](#page-11-0), [1998](#page-11-0)). In a transgenic mouse model of amyotrophic lateral sclerosis (G93A), Py was shown useful for detecting early changes in motoneurons before the onset of neuronal cell death in the spinal cord (Joosten et al. [2001\)](#page-11-0).

However, the identity of the antigen that the Py antibody recognizes has, for almost 30 years, remained elusive. Determining the antigen recognized by the Py antibody would be beneficial not only for understanding the findings of past work that has used the antibody, but may also allow for some understanding of what the function of such a protein might be for neurons that express the antigen in the developing and adult central nervous system (CNS). Here, immunoprecipitation experiments from the adult rat cortex have identified the Py antigen as neurofilament heavy chain (NF-H). This finding was confirmed using double-label immunofluorescence analysis in sections through the rat brain, which highlighted its expression in a subset of neurons in the cortex, hippocampus, septum, basal forebrain, brain stem and cerebellum. Coupling findings from studies using the Py antibody with studies using the NF-H antibody should enhance our understanding of the role that NF-H plays in the development, maturation, and diseased states of the mammalian CNS.

Methods

Py antibody production

A Py IgM hybridoma cell line (2011 clone) was quickly thawed in a 37 \degree C water bath for 1 min. The cell suspension was subsequently centrifuged at 300g for 5 min and the supernatant discarded. The cells were suspended in a hybridoma medium (DMEM, 20 % horse serum, 1 % nonessential amino acids, 1 % glutamax, 1 % penicillin streptomycin fungizone) and centrifuged again at 300g for 5 min and cultured in a T25 flask at 37 °C in 5 % CO_2 . The cells were allowed to expand for 1 week, and were subsequently split 1:3 into T75 flasks filled with hybridoma medium. The cells were grown at 37° C for an additional 2 weeks, when the supernatant was collected for use in immunofluorescence staining. The antibody was stored until use at -80 °C in aliquots to minimize freeze thawing cycles.

Tissue extraction

were carried out under the licensed authority of the UK Home Office (PPL40/3556). Adult, female Sprague–Dawley rats (\sim 230 g weight) were given an overdose of pentobarbitone anesthetic (0.5 mL/100 g) via an intraperitoneal injection, and transcardially perfused with ice-cold sterile 0.9 % sodium chloride. The brains were quickly removed and layers II–V of the cortex (i.e., areas which are known to express the Py antigen, based upon immunofluorescence observations) were dissected from the dorsal region above the striatum. Cortical tissue was placed on a small spatula and excess saline solution removed before placing the sample into a 1.5 mL Eppendorf tube. The dissected tissue was homogenized in approximately four volumes (w/v) of modified RIPA buffer (1 $\%$ NP40, 0.25 % deoxycholate, 1 mM EDTA, 150 mM NaCl, in 50 mM Tris–HCl, pH 7.4) using a small pellet pestle. The extracts were sonicated for 5 s, left on ice for 10 min, followed by centrifugation at 13,000g for 10 min at 4 \degree C to pellet any insoluble material.

Immunoprecipitation

For immunoprecipitation with anti-NF-H, anti-mouse Pan Ig-coated magnetic beads $(50 \mu L)$ (Dynal, Oslo) were washed three times with 4 % BSA in PBS (using a magnetic collector to separate the beads from solution). The beads were then incubated with a monoclonal antibody that recognizes phosphorylated and non-phosphorylated forms of NF-H (Sigma; N0142) (50 μ L of a 1/50 dilution) for 30 min at room temperature with gentle rolling. The beads were then washed three times with PBS (200 μ L), followed by incubation for 1 h at room temperature with the rat cortex RIPA extract $(75 \mu L)$ on a roller. The unbound material was carefully removed from the beads and stored at -80 °C until use. The beads were washed six times with RIPA buffer $(200 \mu L)$ before eluting captured material by heating at 90 °C for 3 min in $2 \times SDS$ sample buffer (75 µL) (2 % sodium dodecyl sulfate–SDS, 5 % 2-mercaptoethanol, 62.5 mM Tris–HCl, pH 6.8). The same method was used for immunoprecipitation with the Py monoclonal antibody $(50 \mu L \text{ neat})$, except that a protein-L matrix $(50 \mu L)$ (CBind L, Sigma) was used for antibody capture instead of Pan Ig magnetic beads because the Py MAb is IgM class.

Western blotting

Protein samples were heated at 90 \degree C in SDS sample buffer (2 % SDS, 5 % 2-mercaptoethanol 62.5 mM Tris–HCl, pH 6.8) for 3 min, separated on 4–12 % SDS-PAGE gradient gel (ThermoFisher Scientific), followed by transfer to nitrocellulose membrane overnight. After blocking nonspecific sites with 4 % powdered milk solution, membranes

were incubated with either Py MAb (1/10), NF-H MAb (Sigma; N0142) (1/1000) or SMI-32 MAb (EMD Millipore; NE1023) (1/1000) diluted in dilution buffer (PBS, 1 % fetal bovine serum, 1 % horse serum and 0.1 % BSA). Antibody reacting bands were visualized by development with peroxidase-labeled rabbit anti-mouse IgG $(1 \mu g/mL)$ in dilution buffer; Dako) (for NF-H and SMI-32) or peroxidase-labeled goat anti-mouse IgM (1/2000; Jackson ImmunoResearch) (for Py), and a chemiluminescent detection system (West Pico, Pierce).

To determine whether the epitope recognized by the Py antibody is glycosylated, electro-blotted membranes were subject to sodium metaperiodate treatment to destroy pentose and hexose carbohydrate epitopes by oxidation, as described by Woodward et al. [\(1985](#page-12-0)). To ensure that the technique did not result in denaturation of protein epitopes, an additional electro-blotted membrane was subject to the same conditions but without addition of metaperiodate. Briefly, electro-blotted membranes were incubated in either 10 mM sodium metaperiodate (in 50 mM sodium acetate buffer, pH 4.5) or just 50 mM sodium acetate buffer, pH 4.5 (for the controls), for 1 h in the dark, followed by incubation with 50 mM sodium borohydride in PBS for 30 min. After rinsing several times with PBS, the blots were blocked and developed in the same way as described above.

Immunofluorescence

Adult, female Sprague–Dawley rats (\sim 230 g in weight) were given an overdose of pentobarbitone anesthetic (0.5 mL/100 g) via an intraperitoneal injection, and cardiac perfused with a 4.0 % solution of paraformaldehyde (PFA) in Tris-buffered saline (TBS). The brain was removed and placed in PFA overnight. The following day specimens were transferred to a 30 % sucrose solution in TBS, and allowed to sink fully in the solution at room temperature (RT). Subsequently, specimens were placed on a sliding, freezing microtome and 40 µm coronal or sagittal sections cut through the brain from the most anterior or rostral section in the coronal plane in which the striatum can be recognized, to the cerebellum. Sections were placed in 24 well plates containing TBS with sodium azide, and 1 in 6 sections were taken for immunofluorescence using Py and NF-H antibodies.

For immunostaining, sections were rinsed 3 times in TBS, and subsequently incubated in a 0.5 % Triton X-100/ TBS solution (TxTBS) for 1 h at RT. Double-label antibody staining was conducted sequentially to avoid crossreaction with secondary antibodies. Control sections were processed at the same time in the absence of either the Py or NF-H primary antibody, but included all steps containing secondary antibodies. Sections were first incubated overnight in a 1:500 dilution of NF-H (Sigma; N0142) in TxTBS. The following day, the sections were rinsed three times for 5 min in TBS, and subsequently placed in a 1:500 dilution of Alexa Fluor[®] 594 goat anti-mouse IgG secondary (Life Technologies) in TxTBS for 2 h. Sections were subsequently rinsed three times for 20 min in TBS, and placed in a 1:50 dilution of Py antibody in TxTBS overnight. The following day sections were rinsed three times for 5 min, and subsequently incubated in a 1:500 dilution of Alexa Fluor[®] 488 goat anti-mouse IgM (Life technology) and $0.5 \mu g/mL$ DAPI in TxTBS for 2 h. Finally, the sections were rinsed three times for 20 min in TBS, and then mounted and coverslipped for microscopy. Images were captured using a Nikon Eclipse 80i microscope fitted with a Hamamatsu fluorescent camera operated with the NiS Elements computer software.

Results

Monoclonal antibody Py recognizes neurofilament heavy chain

Immunoprecipitation from a rat cortex extract using Py MAb attached to protein-L, followed by mass spectrometry analysis of the captured material, resulted in the identification of several Py antigen candidates (data not shown). Whilst some of these presumably bound to the protein-L nonspecifically, only one candidate, neurofilament heavy chain (NF-H), corresponded to the approximate molecular weight of the major band recognized by Py MAb on a western blot (\sim 200 kDa). Immunoprecipitations from a fresh extract of rat cortex using either Py MAb or an anti-NF-H MAb, followed by western blot analysis of the captured material, confirmed that immunoprecipitation and detection of the \sim 200 kDa antigen was reciprocal (Fig. [1a](#page-3-0)). Control pull-downs, using cortex extract added to Dynabeads without antibody (for NF-H) and protein-L without antibody (for Py), verified that the antigen did not bind to either matrix nonspecifically (Fig. [1](#page-3-0)a).

Py has historically been referred to as a ''146 kDa glycoprotein'' (Woodhams et al. [1989;](#page-12-0) Joosten et al. [2001](#page-11-0); Houweling et al. [1999](#page-11-0); Brook et al. [1997a,](#page-11-0) [b](#page-11-0)). To determine whether the epitope recognized by the Py antibody is glycosylated, electro-blotted membranes were subject to sodium metaperiodate treatment to destroy pentose and hexose carbohydrate epitopes by oxidation (Woodward et al. [1985](#page-12-0)). This treatment did not abolish immunoreactivity of the Py MAb, suggesting that the Py antibody epitope is unlikely to be glycosylated (Fig. [1](#page-3-0)b).

The western blot band recognized by the Py and the NF-H MAb on eluates from both antibodies appeared to have a slightly slower electrophoretic mobility, compared to the

Fig. 1 Reciprocal immunoprecipitation and western blot detection of Py and NF-H. a Western blot analysis of immunoprecipitates from a RIPA extract of rat cortex using either Py MAb or an anti-NF-H MAb confirmed that immunoprecipitation and detection of the \sim 200 kDa antigen was reciprocal. The control pull-down lanes refer to cortex extract incubated with Dynabeads without antibody (for NF-H) and Protein-L without antibody (for Py). The band identified by "asterisk" is presumed to be a cross-reaction of the anti-IgM secondary antibody with the IgM heavy chain that eluted from the Protein-L. The bands identified by "greater than symbol" are presumed to be a minor crossreaction of the Py or anti-IgM antibody with a component of the Protein-L that detached during elution of the immunoprecipitate in SDS sample buffer. b Sodium metaperiodate treatment of electro-blotted membranes did not abolish immunoreactivity of the Py MAb. c Western blot analysis of the cortex extract, unbound extract (after pulldown) and eluates, following extended electrophoresis to separate higher molecular weight forms, is shown for the Py MAb immunoprecipitation and in d for the NF-H immunoprecipitation. Both sets of immunoprecipitates (c, d) were also analyzed using another commercially available anti-neurofilament heavy antibody, SMI-32

unbound material following immunoprecipitation (Fig. 1a). To confirm this, and to determine whether the antibodies immunoprecipitate a particular subset of the NF-H in the cortex extract, the eluates were subject to much longer separation on an SDS-PAGE gel, to better resolve higher molecular weight proteins before western blotting. Comparison of the material eluted from the Py and NF-H antibodies against the cortex extract before pull-down (i.e.,

"input") and the unbound material after pull-down revealed some interesting insights (Fig. 1c, d). While the Py and the NF-H MAb both detected a sharp, well-resolved band of the same apparent molecular weight in the eluates, the NF-H MAb detected a band spanning a greater molecular weight range in the input and unbound material, compared to the Py MAb. Analysis of the same samples with an additional, commercially available antibody

against NF-H, SMI-32, revealed the same pattern of immunoreactivity as seen with the NF-H antibody. Of note is the more discrete-sized band recognized by the Py MAb in the input and unbound (Fig. [1](#page-3-0)c, d), and the clear distinction that the Py MAb makes between the faster electrophoretic mobility of the unbound material compared to the slower electrophoretic mobility of the protein that was immunoprecipitated (Fig. [1c](#page-3-0), d).

Taken together, these findings demonstrate that the Py and NF-H antibodies recognize the same sized isoform of NF-H in rat cortex RIPA extracts before protein denaturation, but following denaturation in SDS sample buffer, a wider range of NF-H molecular weight forms—presumably varying in the degree of phosphorylation or other posttranslational modification—are detectable by the NF-H MAb, compared to the Py MAb.

Py and NF-H colocalize in the brain

Immunofluorescence double-labeling with Py and NF-H antibodies yielded an overlapping staining pattern throughout the rat brain (Figs. [2,](#page-5-0) [3,](#page-6-0) [4,](#page-7-0) [5\)](#page-8-0), while control sections (i.e., sections immuno-processed with both secondary antibodies, but excluding either of the primary antibodies) yielded labeling in only a single channel (Fig. [6](#page-9-0)). DAPI labeling in each experimental condition indicates the presence of cells and confirms the integrity of the tissue. Colocalization of Py and NF-H in coronal sections from rostral regions of the brain (approximately -0.12 mm from Bregma; Paxinos and Watson [2005\)](#page-11-0) was notable throughout the cortex, septal nuclei, and basal forebrain (Fig. [2](#page-5-0)). While layer II/III and V of the cortex in this rostral plane showed obvious colocalization of Py and NF-H in neurons with large apical dendrites (Fig. [2a](#page-5-0)–h), staining in the septal region was exclusively fibrous, and the basal forebrain a mixture of cellular and fibrous staining (Fig. [2i](#page-5-0)–p). In the cortex, pyramidal cells in both layers II/III and V displayed intense staining of the cell body, proximal basal dendrites, as well as the full length of the apical process (Figs. [2](#page-5-0)a–h, [7a](#page-10-0)–d). Branching of the apical process in the most superficial regions of the cortex could be seen emanating from neurons in layer II/III. Fibers stained in the septal region (Fig. [2](#page-5-0)i–l) were thick, though there was no obvious cell body staining to conclusively indicate that they were dendrites. In contrast, a small region of the basal forebrain showed a mixture of fiber (dendritic) and somatic staining (Fig. [2m](#page-5-0)–p).

Just caudal to this (approximately -0.48 mm from Bregma; Paxinos and Watson), coronal sections show Py and NF-H colocalizing in the soma and apical dendrites of neurons in Layers II/III and V of the cortex (Fig. [3a](#page-6-0)–h). However, though pyramidal neurons in the cingulate and somatosensory regions of the cortex were intensely immunopositive, the primary and/or secondary motor cortex appeared to have little or no immunoreactivity (Fig. [3a](#page-6-0)–h). In this same plane of section, thick fiber staining highlighted the globus pallidus (Fig. [3](#page-6-0)i–l) with both Py and NF-H antibodies, though it was not obvious whether these were dendritic or axonal. There was, however, a complete absence of fiber or cell body staining in the neighboring striatum.

Even more caudally (approximately -3.48 mm from Bregma; Paxinos and Watson [2005](#page-11-0)), coronal sections showed somatic and dendritic staining throughout the tightly packed CA3 region of the hippocampus (Figs. [4](#page-7-0)a–h, [7](#page-10-0)e–h), and exclusively fibrous staining in the thalamus (Figs. [4](#page-7-0)i–l). In the hippocampus, the soma of pyramidal neurons along with a short portion of their apical process was intensely Py/NF-H+ (Fig. [7e](#page-10-0)–h). In both the thalamic and hippocampal region, staining in the red (NF-H) channel was occasionally more obvious (particularly in white matter tracts), compared to the staining seen with the Py antibody (green). This slight difference in antibody specificity may bear some relationship to the slightly narrower range of molecular weight forms that the Py antibody recognizes after protein denaturation by western blot, compared to the NF-H antibodies (Fig. [1](#page-3-0)).

Caudally, in the cerebellum and brain stem (Fig. [5](#page-8-0)), double-labeling of Py and NF-H in sagittal sections showed staining in Purkinje cell soma and dendrites (Fig. [5a](#page-8-0)–d), and in the soma of cells in the gigantocellular reticular formation (Fig. [5e](#page-8-0)–l). Within the cerebellum, there was a notable absence of staining of any of the numerous neurons of the molecular and granule cell layer (Fig. [5a](#page-8-0)–d). Both the soma and proximal (thick) dendritic process of Purkinje cells were clearly $Py/NF-H+$, though the more fine structures of the dendritic tree were not obvious (Fig. [5](#page-8-0)a–d; [7](#page-10-0)i– l). Again, staining with the NF-H antibody was more apparent in the white matter tract of the cerebellum in comparison to Py immunolabeling (Fig. [5b](#page-8-0)–d). In the brain stem, the loosely packed group of neurons in the region of the gigantocellular reticular formation were intensely Py/ $NF-H+$ (Fig. [5e](#page-8-0)–l). Though fine tubular staining could be seen throughout the soma of the cells, there was little obvious neurite staining surrounding them (Fig. [7m](#page-10-0)–p).

Discussion

Previously, the monoclonal antibody Py was shown to be useful for revealing intricate details of subsets of neurons within the brain and spinal cord. However, the identity of the antigen the antibody recognizes remained a mystery. Here, immunoprecipitation from extracts of rat cortex, and subsequent immunofluorescence characterization shows that Py recognizes the neurofilament heavy chain subunit.

Fig. 2 Immunofluorescence of coronal sections through the brain reveals colocalization of Py and NF-H in the cortex, septum, and basal forebrain. At approximately -0.12 mm from bregma, Py and NF-H staining was evident in layers II/III and V of the cingulate (a– d) and dorsal neocortex (e–h), as well as the fimbria (FI) and lateral septal (LS) regions (i-l), and the basal forebrain (m-p). While the

soma (arrows b and f) and apical process (arrowheads b and f) of neurons in the cortex showed evident staining in layers II/III and V, there was only fibrous staining in the septal region, and a mixture of cell body and fiber staining in the basal forebrain. Scale bars in a and $m = 200 \text{ }\mu\text{m}; \text{ } e = 100 \text{ }\mu\text{m}; \text{ } i = 400 \text{ }\mu\text{m}.$ Diagram adapted from Paxinos and Watson [\(2005](#page-11-0))

Fig. 3 Immunofluorescence of coronal sections through the brain reveals an absence of Py and NF-H staining in the motor cortex, and only fibrous staining in the globus pallidus. While the neurons in layers II/III and V of the somatosensory (arrowhead, b) and cingulate cortex (arrow, **b**) stained intensely for Py and NF-H (a–h), there was a virtual absence of staining in much of the primary (M1) and

In the original article detailing the production of the Py antibody, Woodhams et al. ([1989\)](#page-12-0) describe how the antibody recognized a major band at 146 kDa and a fainter band at 166 kDa in various rat brain homogenates. However, in our hands, the major band recognized by the Py antibody has an apparent molecular weight of approximately 200 kDa, with some much fainter bands present at lower molecular weights (see Fig. [1a](#page-3-0)). In contrast to the western blots shown here that were developed with an anti-IgM secondary

secondary (M2) motor cortex (*b), particularly in layers II/III (a-h). Ventral to this, fibrous staining could be seen throughout the globus pallidus (GP), though the whole of the striatum (CPu) was immunonegative for Py and NF-H (i-l). Scale bars in a and $i = 400 \text{ µm}; e = 200 \text{ µm}.$ m motor cortex, C_g cingulate cortex. Diagram adapted from Paxinos and Watson ([2005\)](#page-11-0)

antibody, Woodhams et al. [\(1989](#page-12-0)) developed their Py blot using an anti-IgG secondary antibody that potentially lacked the required specificity to detect the (IgM class) Py MAb. It is also possible that in previous attempts to resolve the Py antigen on an SDS-PAGE gel, the length of time the extracts were subject to electrophoresis was insufficient to enable the 200 kDa protein to enter the resolving gel.

Py has historically been referred to as a ''146 kDa glycoprotein'' (Woodhams et al. [1989;](#page-12-0) Joosten et al. [2001](#page-11-0);

Fig. 4 Immunofluorescence of coronal sections through the brain reveals overlapping staining for Py and NF-H in the hippocampus and thalamus. In caudal regions of the brain, Py and NF-H had completely overlapping staining in the CA3 region of the hippocampus (a–h), and reticular (rt) and ventral posteromedial (VPM) and ventral posterolateral (VPL) thalamic nuclei (i-l). Note that although the cell soma

and apical processes of neurons in the stratum lucidum (SLu) and CA3 regions of the hippocampus are labeled for Py and NF-H (e–h), only fibrous staining could be seen in nuclei of the thalamus (i–l). Scale bars in **a** and $i = 400 \mu m$; $e = 200 \mu m$. Diagram adapted from Paxinos and Watson [\(2005](#page-11-0))

Houweling et al. [1999;](#page-11-0) Brook et al. [1997a](#page-11-0), [b\)](#page-11-0), presumably because it was a glycoprotein enriched fraction, isolated from a lentil lectin affinity column that was used for immunization (Woodhams et al. [1989](#page-12-0)). Although a glycosylated form of NF-H has been described previously (Cheung and Hart, [2008\)](#page-11-0), our data suggest that the epitope that is recognized by the Py MAb is not glycosylated, since periodate treatment to destroy pentose and hexose carbohydrate epitopes by oxidation (Woodward et al. [1985\)](#page-12-0) did not abolish immunoreactivity of the Py MAb (Fig. [1](#page-3-0)). In the future, it will be interesting to determine whether differences in posttranslational modification (e.g., phosphorylation and/or glycosylation) explain the narrower molecular weight range of NF-H that is recognized by the Py MAb on western blots and whether this has any functional relevance.

Morphologically, the cells that stain positive for Py/ NF-H appear to have several characteristics in common.

Fig. 5 Immunofluorescence of sagittal sections through the brain reveals overlapping staining for Py and NF-H in the cerebellum and brain stem. In caudal regions of the brain, Py and NF-H stained large cells of the Purkinje cell layer (PCL) of the cerebellum (a–d) while the much smaller, but numerous, neurons of the internal granule cell (IGL) and molecular layer (ML) were negative. Similarly, in the brain

Neurons that express the antigen are large in size and many support large (typically apical) dendritic processes. This is illustrated well in layers II/III and V of the cortex (Figs. [2](#page-5-0), [3](#page-6-0)), and in the CA3 region of the hippocampus (Fig. [4](#page-7-0)). Within the cerebellum, there is a striking contrast between the immunoreactivity of the large cells in the Purkinje cell layer, and the lack of any detectable Py/NF-H staining in the adjacent internal granule cell and molecular layers (Fig. 5). In addition, the particularly large neurons in the gigantocellular reticular formation (Fig. 5) are similarly immunopositive for Py/NF-H, while many brain stem neurons remain negative. Such observations are similar to work in the spinal cord by Brook et al. ([1997a](#page-11-0)) who showed that only large diameter alpha motoneurons and the large neurons in Clarke's nuclei are positive for Py, while the small diameter neurons in the substantia gelatinosa are negative. This observation suggests that one possible explanation for the specific staining pattern of a subset of cells could be structural, in that the larger subunit of neurofilament may be important to the maintenance of the cytoskeletal framework of particularly large neurons. Indeed, work by Elder et al. ([1998\)](#page-11-0)

stem, low (e–h) and high (i–l) powered images show that the very large cells of the gigantocellular group of the reticular formation (arrowheads j) are strongly positive for Py and NF-H staining. Boxed area in f, shown at higher magnification in i, j. Scale bars in a and $e = 200 \text{ µm}; i = 100 \text{ µm}$

have shown that mice with NF-H knocked out fail to develop large diameter axons in both the central and peripheral nervous system.

A notable exception to large diameter neurons staining positive for $Py/NF-H+$ are the very large pyramidal neurons (particularly, Betz cells) in the motor cortex. Betz cells are among the largest neurons in the mammalian brain, yet they do not appear to stain positive for Py/NF-H. In an ultrastructural analysis of Betz cells in the human brain, Sasaki and Iwata ([2001\)](#page-11-0) have shown that neurofilament staining of these large neurons is rare, but that neurofilament accumulation is increased in Bunia-like bodies in Betz cells of human brains >65 years of age and in neurons of the anterior horn of the spinal cord of ALS patients (Sasaki et al. [1989\)](#page-11-0). Also, they observed that Betz cells appear to contain filaments that are much thicker than neurofilaments (i.e., 20–25 nm in diameter). Such observations suggest that Betz cells have a particular cytoskeleton that is not dominated by a typical neurofilament structure, and that neurofilaments may mainly occur in these cells with increasing age or the presence of unusual cytoplasmic inclusions.

Fig. 6 Control staining for Py and NF-H show no cross reactivity with secondary antibodies. Low magnification $(1\times)$ image of coronal section of the rat brain double-labeled using Py (IgM) and NF-H (IgG) primary antibodies and corresponding anti-IgM (green) and anti-IgG (red) secondary antibodies (a–d). Inclusion of both primary and corresponding secondary antibodies reveal overlapping staining in the brain (d). In control sections, where the NF-H antibody was

A second commonality among many of the cells and processes that stain positive for Py/NF-H, is that they reside in regions of the brain that appear to be particularly vulnerable to dementia or age-related degeneration. The high expression of Py/NF-H in the cortex and CA3 hippocampal region, for example, correlates well with patterns of abnormal tau phosphorylation in Alzheimer's disease (Blazquez-Llorca et al. [2011](#page-11-0)). Furthermore, staining in the basal forebrain's substantia innominata likely include cells of the nucleus basalis of Meynert, which is known to be significantly affect by both Alzheimer's disease (AD) and Parkinson's disease (PD) (Liu et al. [2015](#page-11-0)). Similarly, connections to and from the cerebellum (Guo et al. [2016\)](#page-11-0) and the septal region of the brain (Stroessner-Johnson et al. [1992\)](#page-12-0) are known to be affected by neurodegenerative disease or age-related cell neuronal loss. Interestingly, seminal work by Chapman et al. ([1989\)](#page-11-0) showed that sera from AD patients showed a high content of antibodies directed again NF-H, and Roder and Ingram ([1991\)](#page-11-0) that age and AD-related changes in ATP may affect neurofilament/tau kinase activity.

omitted (e–h), but both secondary antibodies were applied, no staining could be seen in the red channel (g). Similarly, in control sections where the Py antibody was omitted (i–l), but both secondary antibodies were applied, no staining could be seen in the green channel (j). M motor cortex, cc corpus callosum, ac anterior commissure, CPu striatum. Scale bar in $a = 1$ mm, $e = 100$ µm, $i = 200 \mu m$

Though these observations should not be overstated (as neurons in many regions of the CNS suffer age-related degeneration or are vulnerable to neurodegenerative disease—and the dopaminergic neurons of the substantia nigra pars compacta are not $Py/NF-H+$) it seems important to continue to explore how large neurons may be more vulnerable to age-related degeneration/disease (Mosconi et al. [2008](#page-11-0)), and whether Py/NF-H may provide a useful marker for these cells. Seminal work by Gou et al. ([1995\)](#page-11-0) and more recently by Veeranna et al. ([2011\)](#page-12-0), in fact, strongly indicate that aging may result in a hyperphosphorylation of NF-H; making cells more susceptible to degeneration. What is almost certain is that NF-H is known to be affected in (and therefore serve as a good marker for) a variety of neurodegenerative conditions. Work by Collard et al. ([1995\)](#page-11-0) provided some of the first evidence for NF-H accumulations being a factor in the degenerative process of amyotrophic lateral sclerosis (ALS). More recent work by Schulz et al. [\(2013](#page-11-0)) have shown that reduced phosphorylation of NF-H may be a contributing factor in neurofibromatosis type 2, and Sellner et al. [\(2014](#page-11-0)) that NF-H may

Fig. 7 High magnification, confocal images, showing cellular detail of Py/NF-H colocalization: In layers II/III and V of the cortex, the soma, proximal basal dendrite (arrow head), and apical process ($arrows$) of pyramidal neurons were Py/NF-H+ $(a-d)$. Similarly, in the tightly packed CA3 region of the hippocampus, the soma of large neurons and a short segment of their dendritic (apical) processes (arrow) could be seen in all pyramidal neurons (e–h). In the cerebellum (i–l), it was notable that none of the vast number of cells

in the granule cell layer (IGL), or the sparse cells of the molecular layer (ML) were immunopositive for Py/NF-H. By contrast, the monolayer of large Purkinje cells and their proximal dendritic tree were intensely Py/NF-H+. In the brain stem $(m-p)$, the sparse neurons of the gigantocellular reticular formation showed intense Py/ NF-H staining of their soma, but little notable staining of neurite processes. Scale bars = $25 \mu m$

be a useful pathological and prognostic marker for acute encephalitis. Interestingly, NF-H has also been shown to be a useful marker for the ganglion cells in the retina whose axons have been damaged (Dräger and Hofbauer [1984](#page-11-0)), as well regenerating fibers of retinal ganglion cells (Bates and Meyer, [1993](#page-11-0)).

In conclusion, the identification of the Py antigen sheds light on past work that has utilized the monoclonal antibody Py, and, when combined with studies using the NF-H antibody, expands our understanding of this protein in the developing, mature and dysfunction CNS. It will be important in the future to gain more understanding of why Py/NF-H expression in the brain appears mostly restricted to large diameter neurons, and whether this feature is relevant to neurodegenerative disorders and CNS injuries. The work presented here, and the collective work with Py and NF-H from previous studies, suggests a particular structure and function to cells which express this antigen in the brain and spinal cord, and illustrates how ultrastructural changes in these cells can be identified (in detail) in both normal and abnormal neural conditions.

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Compliance with ethical standards

All applicable international, national, and/or institutional guidelines for the care and use of animals were followed. All procedures performed in studies involving animals were in accordance with the ethical standards of the institution or practice at which the studies were conducted.

Conflict of interest The authors declare that they have no conflict of interest.

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