

Mapping P2X and P2Y receptor proteins in striatum and substantia nigra: An immunohistological study

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Abstract Our work aimed to provide a topographical analysis of all known ionotropic P2X_{1–7} and metabotropic P2Y_{1,2,4,6,11–14} receptors that are present in vivo at the protein level in the basal ganglia nuclei and particularly in rat brain slices from striatum and substantia nigra. By immunohistochemistry-confocal and Western blotting techniques, we show that, with the exception of P2Y_{11,13} receptors, all other subtypes are specifically expressed in these areas in different amounts, with ratings of low (P2X_{5,6} and P2Y_{1,6,14} in striatum), medium (P2X₃ in striatum and substantia nigra, P2X_{6,7} and P2Y₁ in substantia nigra) and high. Moreover, we describe that P2 receptors are localized on neurons (colocalizing with neurofilament light, medium and heavy chains) with features that are either dopaminergic (colocalizing with tyrosine hydroxylase) or GABAergic (colocalizing with parvalbumin and calbindin), and they are

also present on astrocytes (P2Y_{2,4}, colocalizing with glial fibrillary acidic protein). In addition, we aimed to investigate the expression of P2 receptors after dopamine denervation, obtained by using unilateral injection of 6-hydroxydopamine as an animal model of Parkinson's disease. This generates a rearrangement of P2 proteins: most P2X and P2Y receptors are decreased on GABAergic and dopaminergic neurons, in the lesioned striatum and substantia nigra, respectively, as a consequence of dopaminergic denervation and/or neuronal degeneration. Conversely, P2X_{1,3,4,6} on GABAergic neurons and P2Y₄ on astrocytes augment their expression exclusively in the lesioned substantia nigra reticulata, probably as a compensatory reaction to dopamine shortage. These results disclose the presence of P2 receptors in the normal and lesioned nigro-striatal circuit, and suggest their potential participation in the mechanisms of Parkinson's disease.

Keywords Parkinson's disease · Purinergic receptors · Rat brain · Tyrosine hydroxylase · γ -Aminobutyric acid · 6-Hydroxydopamine

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Abbreviations

BG	Basal ganglia
GABA	γ -Aminobutyric acid
GFAP	Glial fibrillary acidic protein
NF-L	Neurofilament-L protein
6-OHDA	6-Hydroxydopamine
MBP	Myelin basic protein
P2r	P2 receptors
SN	Substantia nigra
SNC	Substantia nigra pars compacta
SNR	Substantia nigra pars reticulata
TH	Tyrosine hydroxylase

Introduction

It is now well established that the arrangement of ionotropic P2X and metabotropic P2Y receptors [1, 2] on a cell membrane is a very dynamic process, often related to developmental or physiopathological conditions. Moreover, it is common knowledge that multiple P2 proteins are simultaneously recruited on a cell membrane for triggering biological functions. As a consequence, P2 receptors are rightly considered more than the sum of their single entities and must be therefore regarded as a complex network of cooperating receptors. Under this perspective, a numerical model was also introduced, the combinatorial receptor web model, which explains the biological efficacy of combining an assorted array of different P2 proteins on a given cell, in order to integrate, upgrade, guarantee and optimize specific receptor-dependent functions [3].

This trend of course applies to the central nervous system (CNS) as well, where *in situ* hybridization of P2 mRNA subtypes and immunohistochemistry of P2 proteins shows, for instance, wide but heterogeneous simultaneous distribution of both P2X [4–11] and P2Y [12–15] classes of receptors. In particular, P2X_{2,4,6} and P2Y₁ subtypes are abundant and widespread approximately in the entire brain, while P2X₁ protein is enriched in the cerebellum, P2X₃ in the brain stem, and P2X₇ is largely prejunctional. The hippocampus concurrently expresses all P2X and, moreover, P2Y_{1,2,4,6,12} receptor subtypes. Particularly in the basal ganglia (BG), neostriatal medium-spiny neurons and cholinergic interneurons highly express P2X₂ and P2Y₁ receptors, but it appears that they become functional only under certain, as yet unknown, conditions [16]. Moreover, P2X₂ receptor protein was described in substantia nigra pars compacta (SNc) [17], whereas both protein and mRNA were described in SNc and striatum [18]. Finally, only very low levels of P2X_{4,6} mRNAs were detected in substantia nigra (SN) and striatum [19].

By functional analysis, ATP release was demonstrated from cultured embryonic neostriatal neurons [20], and ATP-evoked potassium currents in rat striatal neurons were shown to be mediated by P2 receptors [21]. ATP was also proved to increase extracellular dopamine levels in rat striatum through stimulation of P2Y subtypes [22], although it was claimed to inhibit dopamine release in the neostriatum [23]. Extracellular ATP via P2 receptors was finally reported to induce neurotoxicity *in vitro* [24] and *in vivo* [25] in the striatum. Besides P2 receptors on neurons, in BG there is also evidence of P2 receptors on, and release of ATP from, glial cells. P2Y₁₂ subtype is present, for instance, on oligodendrocytes in striatum and SN [26], and P2X₇ receptor is upregulated on microglia in striatum after middle cerebral artery occlusion [27]. In spite of these results, there is a general paucity of studies addressing the cellular distribution of all P2 receptor proteins in BG.

Our work thus aimed to provide the complete topographical analysis of known P2X and P2Y subtypes that are present in rat striatum and SN *in vivo*, and to investigate the dynamic presence of P2 proteins after the induction of experimental parkinsonism by dopamine-denervation achieved by using the unilateral 6-hydroxydopamine (6-OHDA) rat model. By upgrading the current map of P2 receptors expressed in the brain, our study discloses the potential impact of these receptors in the normal and lesioned nigro-striatal circuit.

Materials and methods

Histological procedures

Wistar rats (Harlan, Udine, Italy) were anesthetized by *i.p.* injections of sodium pentobarbital (60 mg/kg), and transcardially perfused with saline (0.9 % NaCl) followed by 4% paraformaldehyde, in phosphate buffer (PB, 0.1 M pH 7.4). Each brain was immediately removed, post-fixed in the same fixative for 2 h, and then transferred to 30% sucrose in PB at 4°C, until it sank. The experimental protocol used in this study was approved by the Italian Ministry of Health and was in agreement with the guidelines of the European Communities Council Directive of November 24, 1986 (86/609/EEC) for the care and use of laboratory animals. All efforts were made to minimize the number of animals used and their suffering.

Double immunofluorescence

Transverse sections (40- μ m thick) were cut on a freezing microtome and were processed for double immunofluorescence studies. Non-specific binding sites were blocked with 10% normal donkey serum in 0.3% Triton X-100, in phosphate buffered saline (PBS) for 30 min at room temperature. The sections were incubated in a mixture of primary antisera for 24–48 h in 0.3% Triton X-100 in PBS. Rabbit anti-P2r (1:300, Alomone, Jerusalem, Israel) was used in combination with either mouse anti-calbindin-D-28K (1:200, Sigma, Mi, Italy), mouse anti-tyrosine hydroxylase (TH, 1:500, Sigma), mouse anti-parvalbumin (1:200, Chemicon International, Temecula, CA, USA), mouse anti-glial fibrillary acidic protein (GFAP) (1:400, Sigma), mouse anti-myelin basic protein (MBP, 1:200, Chemicon International), mouse anti-neurofilament H non-phosphorylated (SMI 32, 1:500, Sternberger Monoclonals, Lutherville, MD, USA), mouse anti-neurofilament H and M non-phosphorylated (SMI 33, 1:500, Sternberger Monoclonals), mouse anti-neurofilament 160 (NF160, 1:500, Sigma) or goat anti-neurofilament-L protein (NF-L, 1:100, Santa Cruz, Mi, Italy). The secondary antibodies used for double labeling

were Cy3-conjugated donkey anti-rabbit IgG (1:100, red immunofluorescence, Jackson ImmunoResearch, West Baltimore Pike, PA, USA), Cy2-conjugated donkey anti-mouse IgG (1:100, green immunofluorescence, Jackson ImmunoResearch) or Cy2-conjugated donkey anti-goat IgG (1:100, green immunofluorescence, Jackson ImmunoResearch).

The sections were washed in PBS three times for 5 min each, and then incubated for 3 h in a solution containing a mixture of the secondary antibodies in 1% normal donkey serum in PBS. After rinsing, the sections were mounted on slide glasses, allowed to air dry and coverslipped with gel/mount anti-fading medium (Biomedica, Foster City, CA, USA).

Confocal microscopy

Double- or triple-label immunofluorescence was analyzed by means of a confocal laser scanning microscope (CLSM) (LSM 510, Zeiss, Arese, Mi, Italy) equipped with argon laser emitting at 488 nm, helium/neon laser emitting at 543 nm, and helium/neon laser emitting at 633 nm. Specificity of the antibodies was positively proved by performing confocal analysis in the absence of the primary antibodies, but in the presence of either anti-rabbit or anti-mouse secondary antibodies. Specificity was further confirmed for the P2r antiserum by performing immunoreactions in the simultaneous presence of the P2r neutralizing immunogenic peptides.

Isolation of cerebral areas and protein extraction

Wistar rats were anesthetized by i.p. injections of sodium pentobarbital (60 mg/kg) and, after decapitation, brains were removed. Each brain was transversally cut on a vibratome (300 μ m). The specific cerebral areas were isolated with the aid of a dissection microscope and homogenized in RIPA buffer (1% Nonidet P-40, 0.5% sodium deoxycholate, 0.1% SDS in PBS containing protease inhibitors). After short sonication, the homogenates were incubated on ice for 1 h and centrifuged at 14,000 r.p.m. for 10 min at 4°C. Protein quantification was performed in the supernatants by Bradford colorimetric assay (Biorad, Milan, Italy).

Western blot analysis

Equal amounts of cell lysate (20–30 μ g of protein from each cerebral area) were separated by electrophoresis on 10–12% SDS-PAGE and transferred to nitrocellulose membranes Hybond-C extra (Amersham Biosciences, Cologno Monzese, Italy). The filters were pre-wetted in 5% non-fat milk in TBS-T (10 mM Tris pH 8, 150 mM NaCl, 0.1% Tween 20) and hybridized overnight with P2X_{1,2,4} antisera (Alomone, 1:500), P2X₅ and P2Y_{4/14}

(1:200), P2Y₆ (1:300) or P2Y₂ (1:400). The antisera were immunodetected with an anti-rabbit HRP-conjugated antibody (1:5,000) and developed by ECL chemiluminescence (Amersham Biosciences), using Kodak Image Station (KDS IS440CF).

Anti-P2r specificity

The polyclonal P2r antisera used in this study were raised against P2r highly purified peptides (identity confirmed by mass spectrography and amino acid analysis, as indicated in the certificate of analysis provided by the manufacturer), corresponding to specific epitopes not present in any other known protein. The specificity of the P2r signals was moreover assessed by incubating Western blots either in the absence of the primary antiserum, or in the presence of the primary antiserum together with the neutralizing P2r immunogenic peptides (μ g protein ratio 1:1 between peptide and antiserum).

6-OHDA lesion and Nissl staining

Deeply anesthetized rats (45 days old, about 150 g body weight) were injected with 8 μ g/4 μ l 6-OHDA in saline 0.1% ascorbic acid in the medial forebrain bundle (stereotaxic coordinates ap=-4.4; l=+1.2; vd=-7.8, see also Paxinos et al. [28]) at a rate of 0.38 μ l/min. Fifteen days later, the lesioned rats were tested with 0.05 mg/kg s.c. of the D1/D2 dopamine agonist apomorphine, in order to verify the efficacy of the 6-OHDA lesion, and contralateral turns to the lesion were counted for 40 min. Only those rats that made at least 200 contralateral turns were used for the study. It has been previously demonstrated that rats meeting this screening criterion have greater than 95% depletion of striatal dopamine [29]. At 1.5 months after the 6-OHDA lesion, rats were used for immunohistological experiments ($n=3$). In order to evaluate cell damage, 40- μ m rat brain sections were mounted onto gelatinized slides. They were dehydrated through alcohols, and then rehydrated and stained in 2% cresyl violet for 45 min. Following deionized water rinses, the slides were dehydrated in a standard alcohol series, cleared in xylene, and coverslipped.

Results

P2X and P2Y receptor proteins in rat striatum

We describe in this work the cellular and subcellular in vivo distribution of P2X and P2Y receptors in transverse sections of adult rat striatum, showing by double immunofluorescence confocal analysis that the various P2 subtype proteins are distinguished by different degrees of expres-

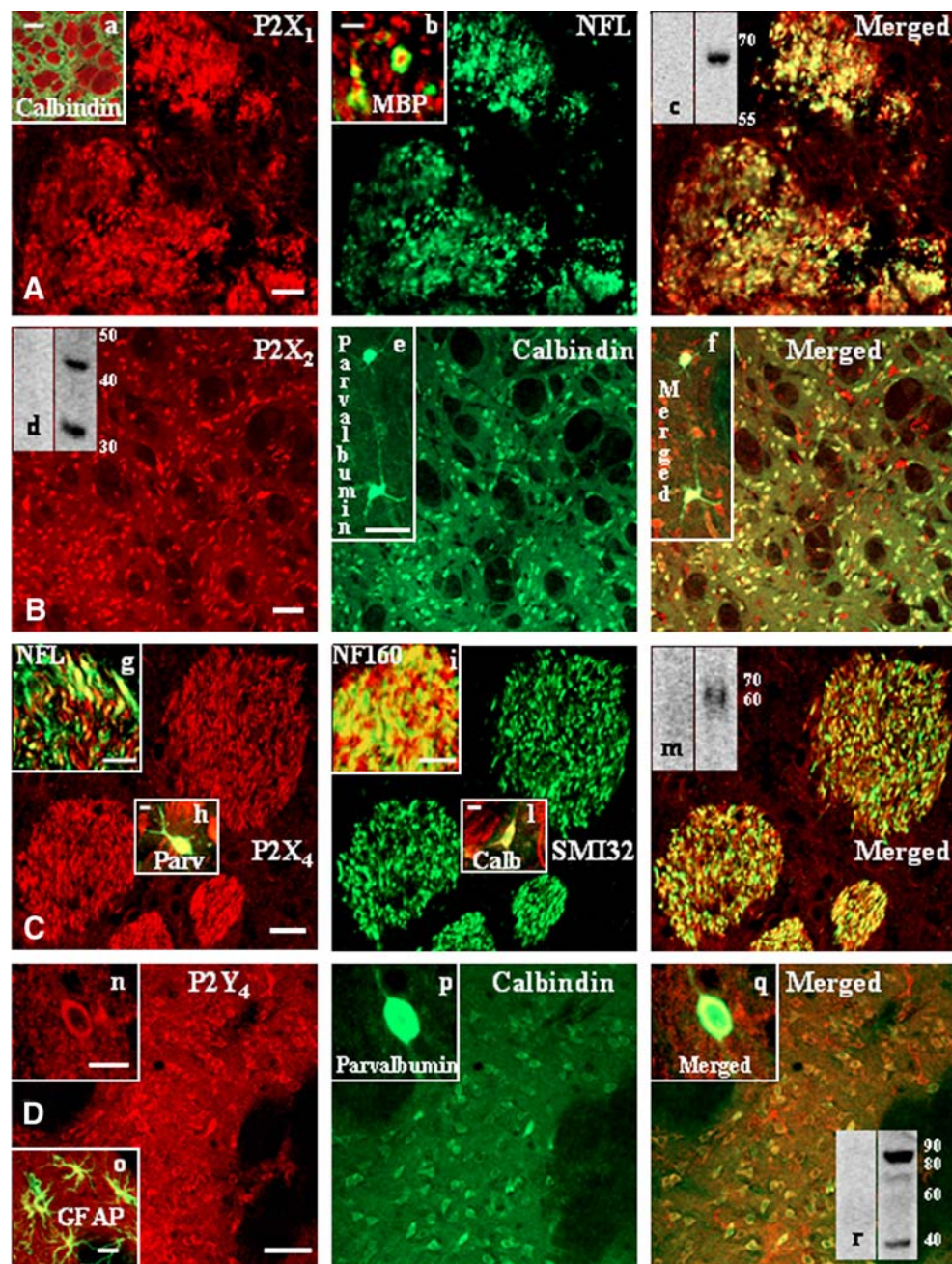


Fig. 1 P2X and P2Y receptor proteins in rat striatum. Transverse sections through the striatum of adult rats were processed for double immunofluorescence studies. Rabbit polyclonal antisera against P2X_{1,2,4} and P2Y₄ receptors (red Cy3 immunofluorescence) were used in combination with antibodies against neuronal or glial markers (green Cy2 immunofluorescence). **Panel A** P2X₁: confocal images illustrate clear colocalization of P2X₁ receptor with neurofilament-light protein (NF-L). The merged field of inset *a* shows absence of colocalization between the neuronal GABAergic marker calbindin (green) (a calcium-binding protein expressed mainly in medium spiny neurons of the striatum) and P2X₁ receptor (red). The merged field of inset *b* shows the merged field of P2X₁ (red) and MBP (green) overlapping immunoreactivities at higher magnification. **Panel B** P2X₂: double immunofluorescence demonstrates that P2X₂ receptor immunoreactivity (red) colocalizes with calbindin protein (green). The insets *e* and *f* show colocalization with the neuronal GABAergic marker parvalbumin (green) (a calcium-binding protein that is expressed in interneurons of

the striatum). **Panel C** P2X₄: red immunofluorescence for P2X₄ protein merges with the green signals of the three types of neurofilament proteins: NF-L (inset *g*, merged field), NF160 (inset *i*, merged field) and SMI 32 and, moreover, with parvalbumin (inset *h*, merged field) and calbindin (inset *i*, merged field). **Panel D** P2Y₄: red P2Y₄ immunoreactivity is present on calbindin-positive neurons (green), on parvalbumin-positive neurons (green) (insets *n*–*q*), and on GFAP-positive astrocytes (inset *o*, merged field). Western blot analysis also confirms the presence of receptor proteins P2X_{1,2,4} (insets *c* in panel *A*, *d* in panel *B*, *m* in panel *C*, respectively) and P2Y₄ (inset *r* in panel *D*) in striatum. Specificity of the P2 receptor signals was assessed by incubations of the primary antisera with the corresponding neutralizing immunogenic peptides (μg protein ratio 1:1 between peptide and antiserum). *Scale bars* are 10 μm in *A*; 100 μm in inset *a*; 2 μm in inset *b*; 50 μm in *B* and in insets *e* and *f*; 20 μm in *C*; 10 μm in insets *h*, *i*, *l*; 5 μm in inset *g*; 50 μm in *D*; 5 μm in inset *n*; and 20 μm in inset *o*. Similar results were obtained in at least four independent experiments

sion and are not uniformly distributed throughout the entire tissue (Fig. 1).

In particular, a strong P2X₁ receptor immunoreactivity (red) confers a patchy appearance to the striatum, being localized mainly in white matter, while sparing the projecting calbindin-positive GABAergic neurons that are highly enriched in gray matter (Fig. 1A, inset a, green). Moreover, P2X₁ protein immunofluorescence is present on NF-L positive, transversally oriented neuronal fibers, although the merged field provides only partial colocalization between the two signals (Fig. 1A). In addition, the high magnification analysis (Fig. 1A, inset b) of P2X₁ (red) and MBP (green) immunoreactive signals shows that P2X₁ receptor is surrounded by MBP, proving the presence of P2X₁ protein on myelinated fibers. Due to the close vicinity of the two signals, overlapping yellow immunofluorescence is also observed. Finally, P2X₁ receptor in striatum is recognized by Western blot analysis as a single protein band of 60–65 kDa, additionally abolished in the presence of the P2X₁ receptor–neutralizing immunogenic peptide (Fig. 1A, inset c).

Conversely, an abundant P2X₂ receptor immunoreactivity (red) is found in gray matter of striatum (Fig. 1B), while sparing the bundles of white matter. Specific receptor immunolabeling is present not only on the highly expressed calbindin-positive projecting GABAergic neurons, but also on the fewer parvalbumin-positive GABAergic interneurons (Fig. 1B, insets e, f). By Western blot analysis, we show that P2X₂ receptor is present in striatum under two isoforms of about 45 and 32 kDa, furthermore, it is abolished in the presence of the P2X₂ receptor–neutralizing immunogenic peptide (Fig. 1B, inset d).

P2X₃ receptor immunostaining in striatum is of medium intensity (Table 1), and mainly localizes on GABAergic neurons of gray matter (data not shown).

P2X₄ receptor signal is instead very copious in white matter, although present on a few fibers of gray matter as well (Fig. 1C, red). It partially colocalizes with all types of heavy-, light- and medium-chain neurofilament proteins (merged fields): SMI 32 (green), NF-L (inset g), and NF160 (inset i). Moreover, we find P2X₄ protein also on GABAergic interneurons (inset h) and GABAergic spiny neurons (inset l). By Western blot analysis, we demonstrate that P2X₄ receptor is present in striatum as a single band of about 60 kDa, moreover, it is abolished in the presence of the P2X₄ receptor–neutralizing immunogenic peptide (Fig. 1C, inset m).

P2X_{5,6,7} and P2Y₁ receptor immunoreactivities in striatum are very weak (Table 1) in gray matter, although totally absent from white matter under our experimental conditions (data not shown).

The P2Y₂ receptor is highly expressed in striatum on axons of white matter and astrocytes of gray matter

(Table 1). Moreover, it is detected as a double protein band in the 55–65 kDa range (data not shown).

A strong P2Y₄ receptor immunoreactivity is present only in gray matter of striatum, localized on both types of GABAergic neurons: calbindin-positive (Fig. 1D) and parvalbumin-positive (insets n–q). Nevertheless, the receptors are also widespread throughout the striatum on astrocytes, as shown by colocalization with the GFAP marker (inset o). By Western blot analysis, we prove that P2Y₄ receptor is present in striatum as a double band of about 42 and 85 kDa (inset r), likely corresponding to the monomeric and dimeric aggregation states of the receptor [30, 31].

While P2Y₆ receptor is barely detectable (Table 1) on GABAergic neurons in striatum (data not shown), P2Y_{11,13,14} receptor proteins were not identified by any means under our experimental conditions (Table 1). Finally, P2Y₁₂ receptor in striatum (Table 1) is abundantly expressed only on oligodendrocytes and myelin sheets, as previously shown [26].

P2X and P2Y receptor proteins in substantia nigra

We conducted a parallel analysis on the cellular and subcellular in vivo distribution of P2X and P2Y receptors in transverse sections of adult rat SN. We showed by double immunofluorescence confocal analysis that the different P2 receptor proteins possess more comparable levels of expression with respect to the striatum, and are

Table 1 Map of P2 receptor proteins in striatum and substantia nigra

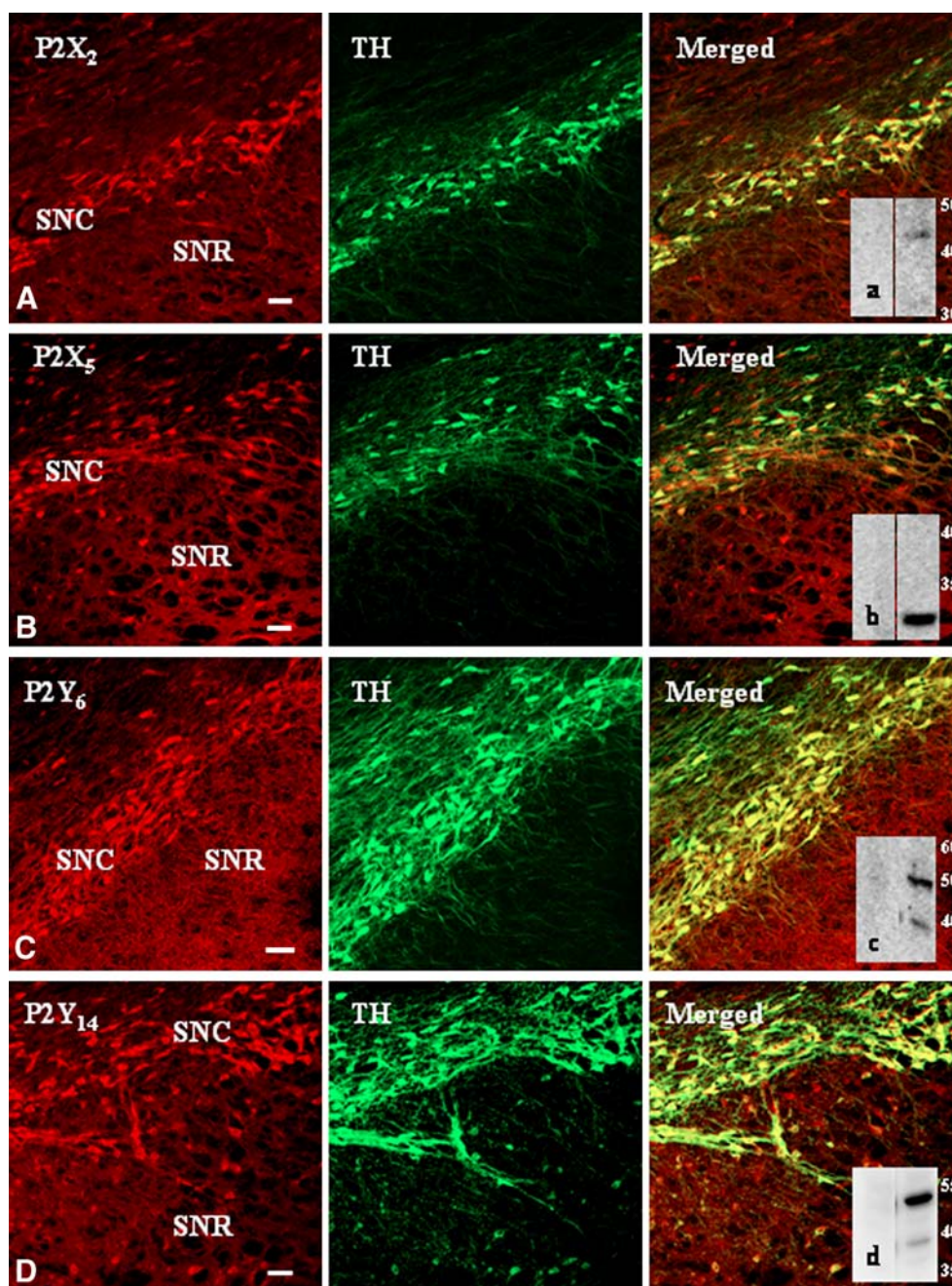
	Striatum	Substantia nigra
P2X ₁	+++	+++
P2X ₂	+++	+++
P2X ₃	++	++
P2X ₄	+++	+++
P2X ₅	+	+++
P2X ₆	+	++
P2X ₇	++	++
P2X ₁₁	–	–
P2X ₁₂	+++	+++
P2X ₁₃	–	–
P2X ₁₄	–	+++

Relative abundance of all P2X and P2Y receptor proteins was analyzed by confocal immunofluorescence microscopy, as described in “Materials and methods”. The intensity of the specific immunostaining was scored as follow: – = not detected; + = just sufficient to evaluate presence and outline of positive cells; ++ = adequate to assess morphological features of cell bodies and/or cellular processes; +++ = very bright

also more uniformly, although differently, distributed throughout the entire SNC and SNR (Fig. 2). In particular, strong signals for ionotropic P2X_{2,5} (red, Fig. 2A,B), P2X_{1,4} (Table 1), metabotropic P2Y_{6,14} (red, Fig. 2C,D) and P2Y₄ (Table 1), or moderate signals for P2X_{3,6} and P2Y₁ receptors (Table 1) are present on dopaminergic neurons (TH-positive) of SNC. Moreover, P2Y₂ and P2Y₁₂ receptors are abundantly expressed in SN (Table 1), but P2Y₂ is expressed on axons and astrocytes, and P2Y₁₂ only on oligodendrocytes and myelin sheets [26]. Conversely, in SNR, a weak P2X/Y receptor immunoreactivity is limited

to sparse neuronal bodies, likely identified as GABAergic neurons by colocalization with parvalbumin (data not shown). The presence at the tissue level in SN of ionotropic P2X_{2,5} (insets *a* in panel A, and *b* in panel B of Fig. 2, respectively) and metabotropic P2Y_{6,14} (insets *c* in panel C, and *d* in panel D of Fig. 2, respectively) proteins is confirmed by Western blot analysis performed in all cases in the presence of specific receptor-neutralizing immunogenic peptides. Similarly to the striatum, immunoreactive signals for P2Y_{11,13} receptors were not identified by any means under our experimental conditions (Table 1).

Fig. 2 P2X and P2Y receptor proteins in rat substantia nigra. Double immunofluorescence visualized by confocal analysis was performed in transverse sections through the substantia nigra of adult rats. Strong signals for ionotropic P2X_{2,5} and metabotropic P2Y_{6,14} (red Cy3 immunofluorescence) are present on dopaminergic neurons (TH-positive, green Cy2 immunofluorescence) of substantia nigra pars compacta (SNC), whereas in substantia nigra pars reticulata (SNR) P2X/Y immunoreactivity is limited to sparse neuronal bodies. Western blot analysis confirms the presence in substantia nigra of receptor proteins P2X_{2,5} (insets *a* in panel A, and *b* in panel B, respectively) and P2Y_{6,14} (insets *c* in panel C, and *d* in panel D, respectively). Specificity of the P2 receptor signals was assessed by incubation of the primary antisera with the corresponding neutralizing immunogenic peptides (μg protein ratio 1:1 between peptide and antiserum). Scale bars in all panels are 50 μm . Similar results were obtained in at least four independent experiments



6-Hydroxydopamine modulates the expression of selected P2 receptors in striatum and substantia nigra

No contralateral rotation as a sign of motor deficit was reported in rats before being 6-OHDA-lesioned, but was instead detected after the lesion rotation (data not shown), together with loss of dopaminergic TH-positive neurons only from the ipsilateral hemisphere of SNC (Fig. 3A and insets *a*, *b*).

Concomitantly, we prove that dopamine denervation in the 6-OHDA-lesioned rat generates a significant and selective rearrangement of P2 receptor proteins. Whereas the expression pattern and immunofluorescence intensities of P2X_{1,4}, P2Y₂ (colocalizing with all neurofilaments and present in white matter on fibers projecting from the

cortex), and P2Y₁₂ (present on oligodendrocytes of white matter) remain constant in both ipsi- and contralateral hemispheres after 6-OHDA treatment (as well as in control animals), all other P2X and P2Y receptors are decreased on parvalbumin- and calbindin-positive GABAergic neurons of deafferented ipsilateral striatum (but not contralateral and in control animals), as measured by semiquantitative analysis (Table 2) ($n=3$).

Similarly, all P2X and P2Y receptors are lost in the lesioned (but not contralateral) substantia nigra pars compacta, consequent to the degeneration of the majority of TH-positive dopaminergic neurons (Table 2). Conversely, P2X₁ (Fig. 3B) and P2X_{3,4,6} (Table 2) receptors present on GABAergic neurons, and P2Y₄ receptors on astrocytes augment their expression only in ipsilateral substantia nigra

Fig. 3 6-Hydroxydopamine modulates the expression of selected P2 receptor proteins in striatum and substantia nigra. Staining of rat substantia nigra after 6-hydroxydopamine treatment. **Panel A** Conventional microscopy images of Nissl staining shows several dopaminergic neurons (arrows) in the contralateral control hemisphere, which are lost (asterisks) in the ipsilateral lesioned hemisphere. Specific ipsilateral dopaminergic lesion of substantia nigra pars compacta (SNC) was also visualized by confocal TH-immunostaining (green) (insets *a*, *b*). **Panel B** Confocal merged yellow images show upregulation of P2X₁ receptor protein (red) in parvalbumin-positive GABAergic neurons (green) in the lesioned side of substantia nigra pars reticulata (SNR) of 6-hydroxydopamine-treated rats. **Panel C** Confocal merged yellow images show a drastic increase in GFAP-positive astrocytes (green) in the lesioned side of 6-hydroxydopamine-treated rats and, correspondingly, an augment of P2Y₄ signal (red) (inset *c*). Scale bars are 100 μ m in *A*, *B* and in insets *a*, *b*; 20 μ m in *C*; and 10 μ m in inset *c*. Similar results were obtained in at least three independent experiments

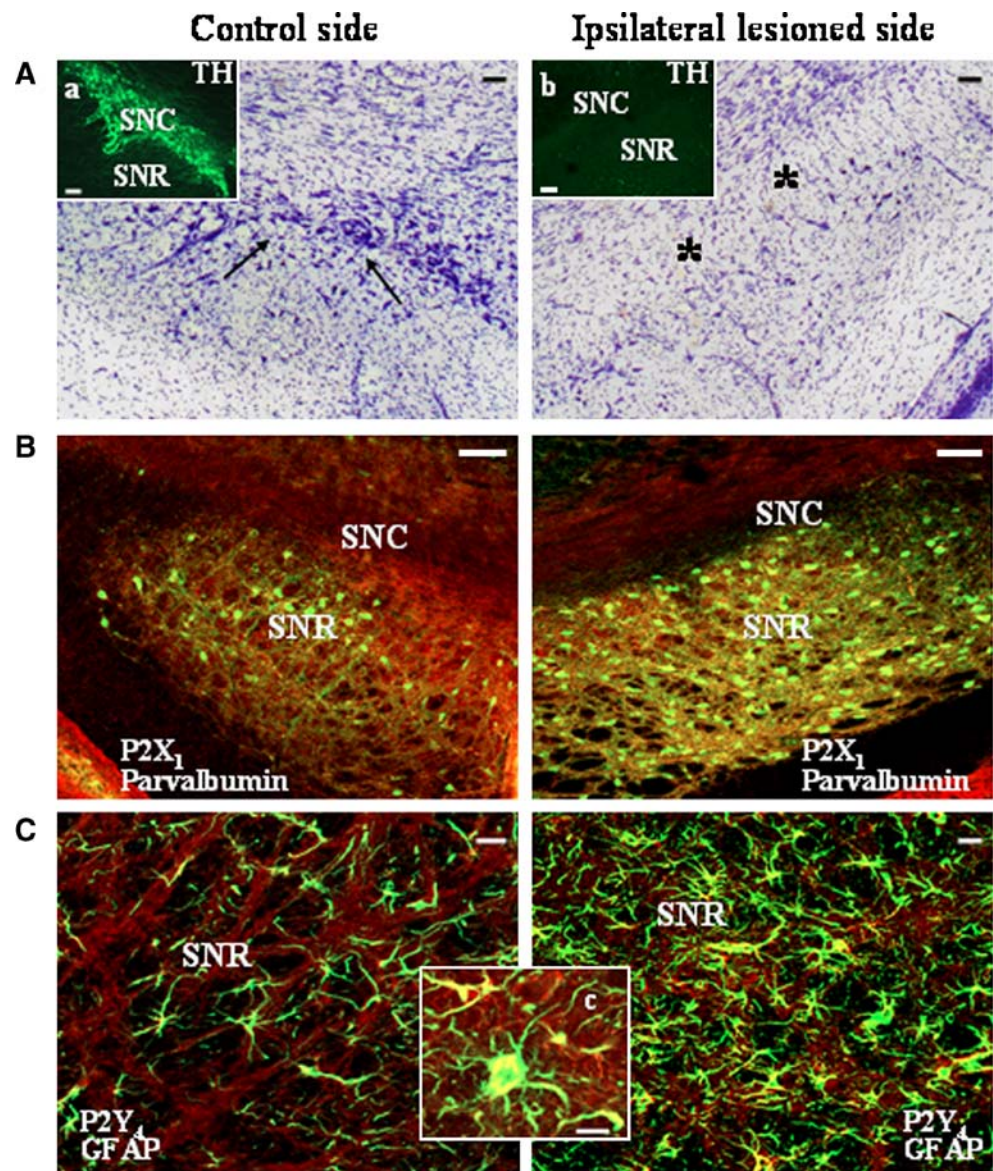


Table 2 Map of P2 receptor modulation after dopamine denervation

	Ipsilateral Striatum	Ipsilateral SN
P2X ₁	=	↑GABA
P2X ₂	↓GABA	↓TH
P2X ₃	↓GABA	↓TH, ↑GABA
P2X ₄	↓GABA	↓TH, ↑GABA
P2X ₅	=	↓TH
P2X ₆	=	↓TH, ↑GABA
P2X ₇	=	=
P2X ₁₁	=	↓TH
P2X ₁₂	=	=
P2X ₁₃	=	=
P2X ₁₄	=	↓TH

Relative increase (↑) or decrease (↓) in P2X and P2Y receptor proteins analyzed by confocal immunofluorescence microscopy in striatum and SN after treatment in rat in vivo with 6-hydroxydopamine (ipsilateral), and in control (not lesioned) brain hemisphere (contralateral). *TH* = presence in dopaminergic neurons, *GABA* = presence in GABAergic neurons, *GFAP* = presence in astrocytes

pars reticulata adjacent to the lesioned pars compacta. In this same area, a phenomenon of astrogliosis is also induced, as detected by more abundant expression of GFAP-positive astrocytes (Fig. 3C).

Discussion

Because the roles of ATP in the CNS have received less attention until recently, often due to lack of appropriate research tools, our knowledge of the functional qualification of P2 receptors in the brain is limited, although rapidly improving. As a group of nuclei interconnected with cerebral cortex, thalamus and brainstem, and associated with a variety of functions, such as motor control, cognition, emotions and learning, the BG [32] is an area that deserves thorough analysis. Our work was aimed at mapping in vivo the presence of P2 receptor subtypes in the BG nuclei of striatum and SN by immunofluorescence-confocal and Western blotting techniques. The specificity of the highly sensitive molecular probes used for the detection of all known P2X and P2Y receptor proteins has been previously validated [33, 34]. In addition, we undertook an analysis that excluded possible cross-reactivity for all antisera used.

Our results not only establish that the majority of P2X (P2X_{1–7}) and P2Y (P2Y_{1,2,4,6,11–14}) receptors so far cloned from mammalian tissues are found in striatum and SN, but also prove their distinctive localization on neurons and/or

glial cells. In detail we show that, with the exception of only P2Y₁₁ and P2Y₁₃ receptors (whose immunoreactivity was not identified by any means under our experimental conditions), all other subtypes are specifically localized in striatum and SN (both pars compacta and reticulata), although with different levels of expression, rated as low (P2X_{5,6} and P2Y_{1,6,14} in striatum), medium (P2X₃ in striatum and SN; P2X_{6,7} and P2Y₁ in SN) and high. Moreover, while we show a prevalence of P2 receptors on neurons (P2X_{1,4} and P2Y₂ colocalizing with neurofilament light, medium and heavy chains) with features that are either dopaminergic (P2X_{2–5} and P2Y_{1,4,6,14} colocalizing with TH, in SN) or GABAergic (P2X_{2–4} and P2Y₄ colocalizing with parvalbumin and calbindin, in striatum), we also describe their expression on astrocytes (P2Y_{2,4} in striatum and SN, colocalizing with GFAP), microglia (P2X₇, colocalizing with OX42) [27], and oligodendrocytes (P2Y₁₂, colocalizing with MBP and RIP) [26]. By confirming previous autoradiographic studies [35, 36], our results therefore prove the widespread but diversified P2-receptor protein distribution in striatum and SN, and extend to these nuclei the great level of biological complexity and molecular sophistication pertaining to P2 receptors [3].

Although the configuration of receptor subunits required for assembly into functional cation channels gated by extracellular ATP in different regions of the CNS comprising the BG is not known yet, colocalization of so many different P2X subtypes in striatum and SN is definitely compatible with heteromultimeric assembly of ionotropic subunits. Since a growing body of biochemical and biophysical evidence now indicates that the propensity to form homo- and especially hetero-multimers is frequent also for G protein-coupled receptors [37] comprising the P2Y subtypes [30, 31], the concurrent expression in striatum and SN of as many metabotropic receptors could explain once more a complex hetero-oligomeric architecture. Nevertheless, the biological phenomenon of redundancy could also justify the simultaneous presence of multiple P2 receptor subtypes in these nuclei, with the final outcome of increasing the structural and pharmacological heterogeneity of these brain regions. Finally, the composite architecture of P2 receptors that we depicted in striatum and SN might likely also signify a multipart mechanism of receptor cooperative behavior (Volonté et al., personal communication) that sustains the concomitant level of complexity of this brain area in several tasks, such as planning and modulation of movement pathways, cognitive processes involving executive functions, reward and addiction. These possibilities are, of course, not mutually exclusive.

Striatal neurons, including the most abundant medium spiny neurons, receive convergent synaptic modulation from nigral dopaminergic neurons and from cortical

glutamatergic projections [38]. The present study showing that lesions of nigral dopaminergic neurons do not significantly affect purinergic receptors present on axons of striatum white matter, but do generate a significant overall decrease in P2X and P2Y receptor proteins from striatal spiny neurons and GABAergic interneurons, thus confirms and extends the involvement of P2 receptors and extracellular ATP to the cortex-basal ganglia circuit [21]. Since dopaminergic denervation affects not only the nigrostriatal dopaminergic pathway but, as a consequence, the corticostriatal glutamatergic pathway with an increase in glutamatergic transmission [39–41] and extracellular glutamate levels in the striatum [42], the reduced P2 receptor protein expression that we demonstrate in striatum gray matter could thus not only be a direct effect of the nigrostriatal inhibition, but also a cause of de-inhibitory mechanisms occurring in the corticostriatal circuit. In this regard, it is common knowledge that extracellular ATP participates in excitatory neurotransmission in the CNS [43], that release of extracellular ATP occurs in CNS under both normal and pathological conditions [44] and, not least, that glutamate release is induced by extracellular ATP in CNS glutamatergic neurons [45].

Neurons of the pars compacta responsible for dopamine production in the brain, which we have shown here to completely lose their array of P2 receptors as a consequence of neurodegeneration induced by 6-OHDA treatment, receive inhibiting signals also from neurons of the pars reticulata that produce GABA [46]. Loss of dopamine neurons in the SNC, one of the main pathological features of Parkinson's disease leading to a marked reduction in dopamine function in the brain, thus also impedes the inhibitory pathway of SNR, with a consequent overactivation of GABAergic neurons. Our findings that specific expression of both ionotropic P2X_{1,3,4,6} receptors on GABAergic neurons and metabotropic P2Y₄ receptors on astrocytes is remarkably increased in SNR after dopamine denervation thus probably reflects a parallel compensatory overreaction of GABAergic neurons to dopamine shortage. One possible explanation is that purinergic mechanisms might thus play a crucial role in the fine-tuned regulation not only of dopaminergic and glutamatergic cross-talk in striatum, as it occurs in nucleus accumbens [47], but also of GABAergic and dopaminergic interplay in SN, as it occurs in the mesolimbic neuronal circuit [48]. This is consistent with the overall versatile functions accomplished by P2 receptors in the CNS under both normal and pathological conditions [43, 44, 49] and, in particular, with the intermediary role in oligodendrocyte-to-neuron [26], Bergmann glia-to-neuron, and neuron-to-neuron communication [50] proposed for P2 receptors in various brain regions.

In summary, the importance of our work is twofold. We first provide the complete topographical analysis of all

known P2X and P2Y receptor subtypes expressed in vivo at their protein levels in rat striatum and SN, which, when considered alongside functional studies, supports a key role for extracellular ATP as a cotransmitter/neuromodulator in these brain areas. Then, we prove that dopamine denervation in the 6-OHDA animal model of Parkinson's disease generates a significant rearrangement of P2 receptor proteins in these nuclei, therefore disclosing the participation of P2 receptors in the lesioned nigro-striatal circuit. While requiring further investigation, our findings indicate a potential but noteworthy pharmacological and therapeutic novel outcome for Parkinson's disease.

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