

Plasticity of Subventricular Zone Neuroprogenitors in MPTP (1-Methyl-4-Phenyl-1,2,3,6-Tetrahydropyridine) Mouse Model of Parkinson's Disease Involves Cross Talk between Inflammatory and Wnt/ β -Catenin Signaling Pathways: Functional Consequences for Neuroprotection and Repair

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In Parkinson's disease (PD), neurogenesis is impaired in the subventricular zone (SVZ) of postmortem human PD brains, in primate nonhuman and rodent models of PD. The vital role of *Wingless-type MMTV integration site (Wnt)/ β -catenin* signaling in the modulation of neurogenesis, neuroprotection, and synaptic plasticity coupled to our recent findings uncovering an active role for inflammation and *Wnt/ β -catenin* signaling in MPTP-induced loss and repair of nigrostriatal dopaminergic (DAergic) neurons prompted us to study the impact of neuroinflammation and the *Wnt/ β -catenin* pathway in the response of SVZ neuroprogenitors (NPCs) in MPTP-treated mice. *In vivo* experiments, using bromodeoxyuridine and cell-specific markers, and *ex vivo* time course analyses documented an inverse correlation between the reduced proliferation of NPCs and the generation of new neuroblasts with the phase of maximal exacerbation of microglia reaction, whereas a shift in the microglia proinflammatory phenotype correlated with a progressive NPC recovery. *Ex vivo* and *in vitro* experiments using microglia–NPC coculture paradigms pointed to NADPH-oxidase (gpPHOX⁹¹), a major source of microglial ROS, and reactive nitrogen species as candidate inhibitors of NPC neurogenic potential via the activation of glycogen synthase 3 (pGSK-3 β ^{Tyr216}), leading to loss of β -catenin, a chief downstream transcriptional effector. Accordingly, MPTP/MPP⁺ (1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine) caused β -catenin downregulation and pGSK-3 β ^{Tyr216} overexpression, whereas manipulation of *Wnt/ β -catenin* signaling with RNA interference-mediated GSK-3 β knockdown or GSK-3 β antagonism reversed MPTP-induced neurogenic impairment *ex vivo/in vitro* or *in vivo*. Reciprocally, pharmacological modulation of inflammation prevented β -catenin downregulation and restored neurogenesis, suggesting the possibility to modulate this endogenous system with potential consequences for DAergic neuroprotection and self-repair.

Introduction

The generation of new neurons in the subventricular zone (SVZ) of the brain continues throughout adult life and may contribute to

endogenous repair mechanisms after brain damage and/or disease (Curtis et al., 2007). In Parkinson's disease (PD), a neurodegenerative disorder characterized by the selective degeneration of dopaminergic (DAergic) neurons in the substantia nigra pars compacta, neurogenesis is impaired. The decreased proliferation of neural stem/progenitor cells (NPCs) in the SVZ of human PD brains, in primate nonhuman, rodent 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP), and 6-hydroxydopamine models, has been attributed to the loss of the neurotransmitter dopamine (Baker et al., 2004; Hoglinger et al., 2004; Freundlieb et al., 2006; O'Keefe et al., 2009a; Lenington et al., 2011). Other studies have proposed dopamine-independent effects of MPTP, but the cellular contributors and signaling pathways through which MPTP may exert its direct action on SVZ cells are currently unknown (He et al., 2006; Shibui et al., 2009). Here, we focused on glia and dissected the influence of MPTP, astrocytes, and microglia on the SVZ response to PD.

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Indeed, NPCs are in intimate contact with surrounding glia, forming the so-called SVZ “stem cell niche” that can influence NPC proliferation and differentiation (Lim and Alvarez-Buylla, 1999; Alvarez-Buylla et al., 2001; Song et al., 2002; Kazanis, 2009). Importantly, in the striatum bordering the SVZ of PD experimental models, astrocytes and microglia exhibit remarkable morphological and functional changes, including expression of an array of proinflammatory and anti-inflammatory cytokines and chemokines, as well as the production of reactive oxygen species (ROS) and reactive nitrogen species (RNS) (Marchetti and Abbraccio, 2005; Gao and Hong, 2008; Hirsch and Hunot, 2009). Of special interest, MPTP-dependent inflammatory mechanisms are recognized to contribute to nigrostriatal DAergic degeneration and self-repair (L'Episcopo et al., 2010a). However, the potential signaling pathways involved in the neuroinflammatory regulation of neurogenesis (Ekdahl et al., 2003, 2009; Monje et al., 2003; Butovsky et al., 2006; Pluchino et al., 2008) in PD are ill defined.

One attractive pathway chiefly involved in the modulation of neurogenesis, neuroprotection, and synaptic plasticity is the *Wingless-type MMTV integration site (Wnt)/β-catenin* signaling system (Kalani et al., 2008; Maiese et al., 2008; Toledo et al., 2008; Munji et al., 2011; Zhang et al., 2011). The *Wnt/β-catenin* pathway plays a central role in the generation, survival, and protection of midbrain DAergic neurons (Prakash et al., 2006; Inestrosa and Arenas, 2010; L'Episcopo et al., 2011a,b). Interestingly, in response to nigrostriatal injury, reactive astrocytes express *Wnt1* and protect DAergic neurons against different neurotoxic insults via the activation of *Wnt/β-catenin* signaling (L'Episcopo et al., 2011a,b). In addition, reactive astrocytes and *Wnt/β-catenin* signaling activation promote neurogenesis from adult NPCs (L'Episcopo et al., 2011a).

Here, using *in vivo*, *ex vivo*, and *in vitro* experiments; different glial–NPC coculture paradigms; along with pharmacological antagonism/RNA silencing experiments coupled to functional studies, we provide evidence supporting an active and concerted role of reactive astrocytes and microglia in the remodeling of the SVZ niche after MPTP/1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPP⁺) injury, at least in part regulated by cross talk between inflammation and *Wnt/β-catenin* signaling cascades, with potential consequences for DAergic neuroprotection and self-repair.

Materials and Methods

Mice and treatments

Eight- to ten-week-old male C57BL/6 mice (body weight, 24–26 g; Charles River, Calco) received $n = 4$ intraperitoneal injections of vehicle (saline, 10 ml/kg) or MPTP-HCl (20 mg kg⁻¹ free base; Sigma-Aldrich) dissolved in saline, 2 h apart in 1 d, according to the acute MPTP injection paradigm (Jackson-Lewis and Przedborski, 2007). Mice were killed at different (1–42 d) time points after MPTP administration (see Fig. 1A). Studies were conducted in accordance with the *Guide for the Care and Use of Laboratory Animals* (NIH) and approved by the Institutional Animal Care and Use Committee.

At the indicated days after MPTP treatment, mice were given bromodeoxyuridine (BrdU; 50 mg kg⁻¹, injected four times, 2 h apart) according to Ahlenius et al. (2009) and killed 2 h after the last injection.

Immunohistochemistry

Mice were deeply anesthetized and transcardially perfused, and the brains were carefully removed, postfixed overnight, placed in 20% sucrose in 0.1 M phosphate buffer for 24 h, frozen (–80°C), and stored until further analyses (Morale et al., 2004). Criostatic coronal sections (14 μm thick) containing the SVZ (0.74, 0.5, 0.14, and 0.02 mm anterior to bregma) were collected according to Paxinos and Watson (1997) and mounted on poly-L-lysine-coated slides. The following preabsorbed pri-

mary antibodies were used: rat anti-BrdU (1:200; Sigma-Aldrich), goat anti-doublecortin (DCX; 1:400; Santa Cruz Biotechnology), mouse anti-proliferating cell nuclear antigen (PCNA) (1:500; Dako), mouse anti-glial fibrillary acidic protein (GFAP; 1:500; Sigma-Aldrich), rabbit anti-GFAP (Dako), sheep anti-epidermal growth factor receptor (EGF-R; 1:50; Millipore), goat anti-ionized calcium-binding adapter molecule 1 (IBA1; 1:200; Novus Biologicals), rabbit polyclonal, anti-inducible nitric oxide synthase (iNOS; 1:200; Santa Cruz Biotechnology), rabbit anti-3-nitrotyrosine (3-NT; 1:200; Millipore), rat anti-dopamine transporter (DAT; 1:500; Millipore), mouse anti-neuron-specific nuclear protein (NeuN; 1:500; US Biologicals), and rabbit anti-cleaved Caspase3 (1:200; Cell Signaling Technology). Immunohistochemistry for PCNA required antigen retrieval performed by incubating the sections in antigen retrieval solution (10× Target Retrieval Solution; Dako) diluted to 1× in distilled water. The antigen retrieval solution was preheated in a water bath at 80°C before the sections were added for 30 min. The sections were subsequently washed three times in PBS at room temperature before immunohistochemistry was performed. Visualization of incorporated BrdU requires DNA denaturation performed by incubating the sections in HCl for 30 min at 65°C. After overnight incubation, sections were rinsed and incubated in darkness for 2 h with CY3-conjugated donkey anti-rat, donkey anti-rabbit, and donkey anti-goat antibodies (1:200; Jackson ImmunoResearch Laboratories), mounted on glass slides, and coverslipped with glycerol-based mounting medium. Nuclei were counterstained with 4',6-diamidino-2-phenylindole (DAPI) in mounting medium (Vector Laboratories) or propidium iodide (PI). In all of these protocols, blanks were processed as for experimental samples except that the primary antibodies were replaced with PBS.

Microscopical analysis

Counting was performed blind to treatment condition. Immunostaining was examined using a Leica LCS-SPE confocal microscope. Briefly, four coronal sections through the SVZ, located at 0.74, 0.5, 0.14, and 0.02 mm anterior to bregma (Paxinos and Watson, 1997), were analyzed bilaterally. Stereological estimations of BrdU⁺, PCNA⁺, DCX⁺, EGF-R⁺, GFAP⁺ (cell numbers per cubed millimeter of SVZ; Nv), were performed using a semiautomatic stereology system (ExploraNova Mercator), using conventional stereological equations (Gundersen and Jensen, 1987), with the formula: $Nv = NC/(t + D)$, where NC is the number of cells/mm², t is the thickness of the section, and D is the cell diameter. Quantification of the percentage of each phenotype marker expressing BrdU (i.e., DCX⁺/BrdU⁺, EGF-R⁺/BrdU⁺, and GFAP⁺/BrdU⁺ cells) was performed in 100 cells in high-magnification fields in the SVZ in four sections per brain, with LAS-SPE imaging software (Leica). 3D reconstruction from z-series was used to verify colocalization in the x – y , y – z , and x – z planes (see Fig. 1C).

Determination of dopaminergic end points in striatum

Measurements of fluorescence intensity (FI) of DAT-immunoreactive (IR) fibers in striatum (Str) were performed by computer-assisted image analysis software (Leica), and changes in average FI (mean ± SD) were expressed as percentage of saline-injected controls. Determination of high-affinity synaptosomal dopamine uptake were performed in Str of saline and MPTP mice, and dopamine uptake levels (total high affinity and mazindol noninhibitable) were expressed as percentage of control as described (Morale et al., 2004; L'Episcopo et al., 2010b, 2011a,b,c).

SVZ dissection and cell isolation

Animals were deeply anesthetized with halothane and killed by cervical dislocation. Brain coronal sections from vehicle- and MPTP-treated mice were taken from mice 2 mm from the anterior pole of the brain, excluding the optic tracts, and 3 mm posterior to the previous cut, at the indicated time intervals [1–35 d posttreatment (dpt); see Fig. 2]. The SVZs were then dissected out under a microscope. Dissected tissue was transferred to Earl's balanced salt solution (Invitrogen) containing 1 mg/ml papain (27 U/mg; Sigma-Aldrich), 0.2 mg/ml cysteine (Sigma-Aldrich), and 0.2 mg/ml EDTA (Sigma-Aldrich) and incubated for 45 min at 37°C on a rocking platform. Tissues were then transferred to DMEM/F-12 medium (1:1 v/v; Invitrogen) containing 0.7 mg/ml ovomucoid (Sigma-Aldrich) and mechanically dissociated. After digestion,

the number of viable cells was determined by trypan blue (Sigma-Aldrich) exclusion. The green fluorescent protein-positive (GFP⁺) cells were isolated from the SVZ and established as described in detail previously (Pluchino et al., 2003, 2005; L'Episcopo et al., 2011a).

Primary neural stem/precursor cell cultures

To analyze self-renewal, continuous *in vitro* propagation was performed in neurosphere growth medium (DMEM/F-12 containing 2 mM L-glutamine, 0.6% glucose, 0.1 mg/ml apo-transferrin, 0.025 mg/ml insulin, 9.6 μ g/ml putrescine, 6.3 ng/ml progesterone, 5.2 ng/ml Na selenite, 2 μ g/ml heparin) supplemented with EGF (20 ng/ml) and FGF2 (10 ng/ml), and the estimated total number of cells was calculated by multiplying the amplification rate (total number of cells obtained at a given subculture passage per 250,000) for the total number of cells obtained at the previous passage, as described (Pluchino et al., 2008). Data from continuous *in vitro* propagation experiments were expressed as absolute number of viable cells \pm SEM from a total of $n \geq 3$ independent experiments.

In vitro neurosphere formation assay

Primary cells isolated from the SVZs of either vehicle- or MPTP-treated mice ($n \geq 4$ per group) were plated in 24-well uncoated plates (0.5 ml/well; Corning) at a density of 8000 cells/cm² in growth medium, as described (Gritti et al., 2002; Pluchino et al., 2008). Under these conditions, neurospheres are derived from single cells and serve as an index of the number of *in vivo* neural stem cells (Morshead and van der Kooy, 1992; Morshead et al., 1994). The number of primary neurospheres with a diameter (ϕ) $\geq 100 \mu$ m in each well was counted 7 d after plating. To determine the self-renewal capacity, we mechanically dissociated primary spheres into single-cell suspensions and recultured them under the same conditions as primary cultures. Again, we determined the number of secondary neurospheres after 7 d. The diameter of the single primary and secondary neurospheres was measured under an inverted microscope (Axiovert S100TV). Data were expressed as either mean clonal efficiency (namely, percentage of colony-forming cells, over total plated cells) or mean diameter (in micrometers), both \pm SEM from a total of $n \geq 3$ independent experiments.

In vitro differentiation conditions

To analyze proliferation, individual spheres (e.g., primary, secondary, established from either saline or MPTP mice; GFP⁺-expressing cells) were mechanically dissociated, and single cells were plated at a final density of 1×10^5 cells/cm on poly-D-lysine-coated 24-well plates in differentiation medium consisting of a mixture of a 1:1 of F-12 and MEM, containing HEPES and glutamine, with N2 supplements (Invitrogen), with 1% fetal calf serum (FCS). Proliferation was studied by addition of the nucleotide analog BrdU (5 μ M) at 2 DIV, in cell cultures exposed to the different treatment/coculture paradigms, and the cells were fixed after 24 h (i.e., at 3 DIV). For neuronal differentiation studies, neurospheres plated in differentiation medium were allowed to differentiate for 5–10 DIV in the absence or presence of the different treatments/coculture paradigms, as described. The direct effect of the neurotoxin MPP⁺ was studied in NPCs grown with or without glia, and in the absence or presence of activator/inhibitors of Wnt/ β -catenin signaling (see Figs. 4, 6, 7), as described below.

Isolation and culture of astrocyte and microglia

Mixed glial cell cultures were obtained from mouse brain, at postnatal days 2–3, as described in detail (Gallo et al., 2000a; Gennuso et al., 2004). The cultures were allowed to grow and differentiate until they reached confluency (14–16 DIV), and the resulting mixed cultures (65 \pm 6% GFAP⁺ astrocytes, 24 \pm 4% IBA-1⁺ microglia, and 5 \pm 2% oligodendrocytes, by immunocytochemistry) were used for coculture experiment (NPC–glia). For purified microglia and purified astrocyte cell preparations, the mixed glial cultures established as described were allowed to grow and differentiate until confluence (14–16 DIV), when the loosely adherent microglial cells were separated by shaking for 5 h at 37°C and 150 rpm, counted, and cultured in microglia medium (RPMI-1640 supplemented with 10% FCS, 1 mM L-glutamine, 1 mM sodium pyruvate, 50 mM β -mercaptoethanol, 100 U/ml penicillin, and 100 mg/ml streptomycin).

The primary astrocytes, obtained after separation of microglia (>95% of the cells were GFAP-IR astrocytes), and the enriched microglial (>95% of the cells were IBA1-IR microglia) monolayers were rinsed with sterile PBS and replated at a final density of $0.4\text{--}0.6 \times 10^5$ cells/cm² in poly-D-lysine (10 μ g/ml)-coated 6-, 12-, or 24-well plates, or in insert membranes (0.4 μ m of polyethylene terephthalate) for indirect coculture (BD Biosciences).

Glial–NPC cocultures and treatments

The direct effect of MPP⁺ was studied in the absence (monotypic cultures) or presence (coculture) of glia, using GFP⁺ cells (L'Episcopo et al., 2011). GFP⁺ cells at passages 8–12 were used in all *in vitro* experiments. Astrocyte, microglial, or mixed astrocyte–microglial monolayers were freshly cocultured with GFP⁺ cells for different time intervals, according to proliferation (3 DIV) and differentiation (5–10 DIV) studies (L'Episcopo et al., 2011a). Both NPC monotypic and glial–NPC cocultures received increasing doses (5–25 μ M) of MPP⁺ (see Fig. 4). These doses were selected based on our previous studies on primary mesencephalic DAergic neurons in culture (L'Episcopo et al., 2011a;b). For proliferation studies, MPP⁺ was applied at 2 DIV, the nucleotide analog BrdU (5 μ M) was added, and the cells were fixed after 24 h, i.e., at 3 DIV. For differentiation studies, GFP⁺ cells grown alone, or layered on top of astrocyte (NPC–Astro), microglia (NPC–Micro), or mixed astrocyte–microglial (NPC–glia) preparations, shifted in differentiation medium, and cells were allowed to differentiate. MPP⁺ was applied at 5 and 9 DIV, and the cells were fixed after 24 h (see Fig. 4). Additionally, we studied the effect of coculture of GFP⁺ cells with microglia isolated “*ex vivo*” from naive or MPTP mice (see Fig. 5). Reciprocally, the effect of coculture of astrocytes with NPC isolated *ex vivo* from naive and MPTP mice, both in the absence or presence of Wnt/ β -catenin signaling activation/antagonism, was also studied as described (see Fig. 7).

For immunocytochemistry, cell cultures were fixed in 4% paraformaldehyde in PBS or with paraformaldehyde/PBS followed by ice-cold acidic ethanol and HCl for BrdU staining (L'Episcopo et al., 2011a). The following markers were used: mouse anti-Tuj1 (Covance Research Products) and rabbit anti-microtubule-associated protein 2abc (Map2a; Abcam) as neuronal markers; rabbit anti-GFAP (Millipore), rat anti-BrdU (Abcam), mouse anti-BrdU (Dako), goat anti-IBA1 (Novus Biologicals), rabbit polyclonal, and anti-iNOS (Santa Cruz Biotechnology). Nuclei were counterstained with DAPI. Analyses were performed using a confocal laser microscope and computer-assisted image analysis (Leica). For quantification of the amount of cells expressing a given marker or marker combinations, the number of Map2⁺ cells was determined relative to the total number of DAPI-labeled nuclei or relative to GFP⁺ cells; the number of BrdU⁺ cells was determined relative to the total number of DAPI⁺ cells or GFP⁺ cells, using the Leica Lite software and three-dimensional overlay to avoid false-positive/negative overlay and double counting.

Ex vivo isolation and culture of macrophage/microglia, measurement of RNS, and intracellular ROS

Isolation of brain macrophage/microglia from saline and MPTP mice was performed in $n = 8$ mice per group as described in detail (Marchetti et al., 2002; Morale et al., 2004; L'Episcopo et al., 2011c), 3 d after saline or MPTP (i.e., at the peak of microglial activation; see Fig. 3B). Cells were immunoreacted for activated macrophage/microglial-specific [IBA1; macrophage antigen-1 (Mac-1)/CD11b] and nonspecific (GFAP for astrocyte, galactocerebroside for oligodendrocyte, or neurofilament for neuron) markers. The isolated cells (>95% IBA1⁺ cells) were counted and plated at a final density of $0.4\text{--}0.6 \times 10^5$ cells/cm² in poly-D-lysine (10 μ g/ml)-coated 6-, 12-, or 24-well plates, or in insert membranes (0.4 μ m of polyethylene terephthalate) for indirect coculture (BD Biosciences). The cells were cultured in microglia medium (RPMI-1640 supplemented with 10% FCS, 1 mM L-glutamine, 1 mM sodium pyruvate, 50 mM β -mercaptoethanol, 100 U/ml penicillin, and 100 mg/ml streptomycin) and used as follows. Part of the cells were processed 24–48 h after plating for ROS measurement using the redox membrane-permeant probe 2',7'-dichlorofluorescein diacetate (DCF-DA; 50 μ M, added for 1 h at 37°C), and cells were viewed under the confocal microscope

(Gennuso et al., 2004). Measurement of iNOS-derived NO was performed in cell-free supernatant using Griess reagent (Marchetti et al., 2002; Morale et al., 2004; L'Episcopo et al., 2011c). To study the effect of inhibition of oxidative and nitrosative stress mediators, the freshly prepared IBA1⁺ cells were cultured in the absence or presence of the ROS antagonist apocynin (Apo; 0.5 mM) (Gao et al., 2003) and the specific iNOS inhibitor L-Nil [L-N⁶-(1-iminoethyl)-lysine, 50 μ M; Sigma-Aldrich] (Marchetti et al., 2002; Morale et al., 2004) applied after plating, and determinations were performed 24–48 h after treatment (see Fig. 5).

Coculture of microglia isolated ex vivo from saline and MPTP mice with GFP-expressing NPCs

To study the effect of microglia exposure *in vivo* to MPTP (MPTP-microglia) on NPC neurogenic potential, the freshly isolated microglial preparations from saline and MPTP mice, as above, were cocultured with GFP⁺ NPC. As a control (ct-insert), a nonmicroglial preparation (i.e., neuronal GT1–7 cells) was used (Gallo et al., 2000b). For proliferation studies, NPCs were layered on top of the microglial cell monolayers, and after 24 h, the nucleotide analog bromodeoxyuridine (5 μ M) was added and the cells fixed after 24 h (i.e., 48 h after microglial exposure). For differentiation studies, NPCs were grown for 3–4 DIV in differentiation medium as above (N2 medium without growth factors and 1% FCS), the freshly prepared microglial cells were added on top, and the cells were fixed at 5–6 DIV (48 h after the microglial exposure). The cultures were fixed and processed for fluorescent immunocytochemistry as described. Indirect coculture paradigms, i.e., NPCs exposed to saline–microglia, MPTP–microglia, or ct-inserts, were performed as above. The inserts containing the saline, MPTP-microglial monolayers, or ct-inserts were added on top of the NPCs (L'Episcopo et al., 2011a). NPC survival was estimated by counting the number of GFP⁺ cells over the DAPI-positive nuclei and by determination of Caspase3 activity (see Fig. 5).

Caspase3 activity

To verify changes in cell survival, the cells were lysed in ice-cold lysis buffer containing 25 mM HEPES, 5 mM EDTA, 1 mM EGTA, 5 mM MgCl₂, 5 mM dithiothreitol (DTT), 1 mM phenylmethylsulfonyl fluoride, and 10 μ g/ml each of pepstatin and leupeptin, pH 7.5. The cells were left for 20 min on ice and then sonicated. The lysate was centrifuged for 20 min at 10,000 \times g, and the supernatant was quickly frozen in a methanol dry ice bath and stored at -80°C . Lysates (30 μ g protein) were incubated at 37 $^{\circ}\text{C}$ in a buffer containing 25 mM HEPES, pH 7.5, 10% sucrose, 0.1 3-[(3-cholamido propyl) dimethyl ammonio]-1-propanesulphonate, and 10 mM DTT with the fluorogenic substrate DEVD-AFC (15 μ M in dimethylsulfoxide; Calbiochem), and quantification of the DEVD-like fluorescent signal was assessed in a luminescence spectrophotometer (excitation, 400 nm; emission, 505 nm) (L'Episcopo et al., 2011b). Enzymatic activity is expressed as arbitrary fluorescent units.

Western blot analysis

Inflammatory mediators and *Wnt*- β -catenin signaling components were studied by Western blot analysis (L'Episcopo et al., 2010a, 2011a,b,c). Protein extracts were prepared from tissues (Str, SVZ) isolated from saline or MPTP mice ($n = 6$ per group per time point) and from cell cultures within the different experimental groups. Protein samples were diluted to equivalent volumes containing 20 μ g of protein and boiled in an equal volume of Laemmli SDS boiling buffer (Sigma-Aldrich) for 10 min. Samples were loaded into a 9–12% SDS-polyacrylamide gel and separated by electrophoresis for 3 h at 100 V (L'Episcopo et al., 2010b, 2011a,b,c). Proteins were transferred to polyvinylidene difluoride membrane (GE Healthcare) for 1.5 h at 300 mA. The following primary antibodies were used: rabbit anti-Mac1 (Abcam); rabbit anti-p47PHOX (Millipore); mouse anti-gp91PHOX, mouse anti- β -catenin, rabbit anti-3-NT, and mouse anti-glycogen synthase kinase-3 β (GSK-3 β) (all from Transduction Laboratories); and mouse anti-GSK-3 β phospho-Tyr216 (BD Biosciences). Membranes were reprobbed for β -actin (Cell Signaling Technology) or GSK-3 β immunolabeling as internal controls. The bands from the Western blots were densitometrically quantified on x-ray films using ImageQuantity One. Data from experimental bands were normalized to β -actin. Values are expressed as percentage of saline-injected controls (see Figs. 3, 5, 9) or as signal intensity values (fold changes)

compared with saline control (see Fig. 6, 7). Western blot measurements were repeated three times independently.

RNA extraction, reverse transcription, and real-time PCR

Dysregulation of the *Wnt*/ β -catenin signaling pathway in SVZ cells from MPTP mice was also studied measuring the expression level of *Axin2*, a direct *Wnt* target transcriptionally induced after the reception of a *Wnt*/ β -catenin signal (Jho et al., 2002). RNA extraction was performed in cell samples from MPTP- and saline-injected mice homogenized in 1 ml of QIAzol Lysis reagent (79306; QIAGEN) using a rotor-stator homogenizer, as described in detail (L'Episcopo et al., 2011a,b). Total RNA was isolated from homogenized samples using the RNeasy Lipid Tissue kit (74804; QIAGEN) including Dnase digestion. At the end, RNA samples were redissolved in 30 μ l of RNase-free water, and their concentrations were determined spectrophotometrically by A₂₆₀ (Nanodrop-ND 1000); the cDNA was synthesized from 2 μ g of total RNA using the Retroscript kit (Ambion). After purification using the QIAquick PCR Purification kit (QIAGEN), 250 ng of cDNA were used for real-time PCR using predeveloped Taqman Assay reagents (Applied Biosystems). Real-time quantitative PCR was performed with the Step One Detection System (Applied Biosystems) according to the manufacturer's protocol, using the TaqMan Universal PCR master mix (4304437). The assay identifications for *Axin2* (Mm00443610_m1) and β -catenin (Mm00483039_m1) were from Applied Biosystems. For each sample, we designed a duplicate assay, and β -actin (4352341E; Applied Biosystems) was used as the housekeeping gene. Results are expressed as arbitrary units.

Effect of manipulation of *Wnt*/ β -catenin signaling in vitro

Pharmacological activation/antagonism. To study the effect of pharmacological activation of *Wnt*/ β -catenin signaling in NPCs alone, we used the specific GSK-3 β inhibitor AR-AO14418 [N-(4-methoxybenzyl)-N'-(5-nitro-1,3-thiazol-2-yl)urea] (AR; 5 μ M; Osakada et al., 2007; L'Episcopo et al., 2011a,b) or the *Wnt* ligand *Wnt1* (100 ng/ml). AR or *Wnt1* was applied 1 h before MPP⁺/PBS exposure, and NPCs were fixed after 24 h (L'Episcopo et al., 2011a,b). For *Wnt*/ β -catenin antagonism studies in astrocyte–NPC cocultures, we used Dickkopf-1 (*Dkk-1*; 100 ng/ml; R & D Systems). *Dkk-1* is a high-affinity ligand for LRP6 and inhibits *Wnt* signaling by preventing Frizzled (Fz)–lipoprotein receptor-related protein 6 (LRP6) complex formation induced by *Wnt* (Semenov et al., 2001). *Dkk1* was applied to NPCs just before the start of the coculture with astrocytes, and the cell was fixed at the indicated time intervals for proliferation/differentiation studies.

Transient gene silencing with small interfering RNA. To test the effect of GSK-3 β or β -catenin protein depletion, we used targeted mRNA degradation using small interference RNA (siRNA) performed essentially as described (Brazas and Hagstrom, 2005). GSK-3 β siRNA (sc-3525), β -catenin siRNA (sc-29210) (He and Shen, 2009), and control siRNA (sc-37007) were purchased from Santa Cruz Biotechnology, and siRNA introduction was performed according to the protocol provided by Santa Cruz Biotechnology, as reported in detail in our previous study (L'Episcopo et al., 2011b). Briefly, to prepare lipid-siRNA complexes, 80 pmol of the indicated siRNA duplex in 100 μ l of Transfection medium (sc-36868) and 6 μ l of siRNA Transfection reagent (sc-29528) in 100 μ l of Transfection medium were combined, incubated for 30 min at 25 $^{\circ}\text{C}$, and diluted with 800 μ l of prewarmed Transfection medium. NPCs from saline and MPTP mice, dissociated and plated on 24 plate cells in growth medium, were rinsed once with serum-free DMEM, and 1000 μ l of lipid-siRNA mixture described above was applied per well. After incubation for 6 h at 37 $^{\circ}\text{C}$ in a humidified 5% CO₂ cell culture chamber, an additional 1 ml of 20% FBS in DMEM was added per well, and lipofection was allowed to continue overnight. The next morning, the lipofection medium was aspirated, and transfected cells were refed with fresh growth medium. The cells were collected 72 h after transfection for Western blot assays.

Effect of manipulation of *Wnt*/ β -catenin signaling, in vivo

Systemic injections and intracerebroventricular infusions. To link the *Wnt*/ β -catenin signaling pathway to MPTP-induced neurogenic impairment *in vivo*, we addressed (1) the effect of pharmacological activation of *Wnt*/ β -catenin signaling using the specific GSK-3 β inhibitor AR by systemic

or SVZ administration (Adachi et al., 2007; L'Episcopo et al., 2011a,b) and (2) the effect of *Wnt*/ β -catenin signaling antagonism with *Dkk1* intracerebroventricular infusion (Zhang et al., 2008) in the absence or presence of AR systemic administration. Pilot experiments were performed to verify doses and timing capable to decrease active GSK-3 β in the SVZ of both intact and MPTP mice. To test the systemic effect, AR was injected in intraperitoneally at a dose regimen (10 mg/kg twice per day) (L'Episcopo et al., 2011a,b), starting 3 h after MPTP. To kill the mice, we selected 3 d post-MPTP, corresponding to maximal neurogenic impairment and exacerbated microglial reaction (see Fig. 3).

For intracerebroventricular infusion, AR (100 μ M), or the vehicle alone, was infused into the left lateral ventricle of the brain with a miniosmotic pump (model 1007D; flow rate, 0.5 μ l/h; Alzet Osmotic pumps, Cupertino, CA). The cannula was implanted stereotaxically at the following coordinates: anterior, 0 mm; lateral, 1.1 mm; depth, 2.3 mm (relative to bregma and surface of the brain). For *Wnt*/ β -catenin antagonism, we used recombinant human *Dkk1* protein (rh-*Dkk1*; R & D Systems) intracerebroventricular infusion, essentially as reported by Zhang et al. (2009). *Dkk1* was dissolved in sterile physiologic saline (0.9% NaCl) at a final concentration of 1 μ g/ μ l. The infusions of *Dkk1* or the vehicle were performed unilaterally as above using a 2 μ l Hamilton microsyringe. The volume of the solution was infused at a rate of 0.25 μ l/min. The needle was kept in place for 5 min after each infusion before retraction. To address the ability of *Wnt*/ β -catenin activation to reverse *Dkk1* effect, unilateral infusion of *Dkk1* or vehicle was preceded by saline or AR systemic injections as above, starting 48 h before intracerebroventricular administration. The mice were killed 48 h after *Dkk1*/vehicle infusion, and the brains were processed as described for protein expression and neurochemical and histopathological determinations.

Effect of pharmacological modulation of inflammation. To study whether pharmacological modulation of inflammation affects NPC proliferation after MPTP injury via *Wnt*/ β -catenin signaling in SVZ *in vivo*, we addressed the effect of the cyclooxygenase inhibitor, NO-releasing drug (CINOD) 2-fluoro- α -methyl(1,1'-biphenyl)-4-acetic-4-(nitrooxy)butyl ester (HCT1026; kindly provided by Nicox Research Institute, Bresso, Milan, Italy), which belongs to a novel class of nonsteroidal anti-inflammatory drugs endowed with strongly reduced side effects (Keeble and Moore, 2002; Fiorucci et al., 2007; L'Episcopo et al., 2010b, 2011c). HCT1026 was administered in the diet at a dose regimen (HCT1026, 190 ppm in the diet or 30 mg kg⁻¹ per day per animal) shown to mitigate glial neuroinflammatory reaction and afford DAergic neuroprotection, when compared with mice fed with a control diet [plain Teklas 2018 chow (ct diet)] (see L'Episcopo et al., 2010b, 2011c). The HCT1026 diet was started 7 d before MPTP, and mice were killed at 3 d (at the peak of neuroinflammatory microglial reaction) and 14 d after MPTP (corresponding to the phase of stabilization of DAergic degeneration) (Jackson-Lewis and Przedborski, 2007; L'Episcopo et al., 2010b, 2011a).

Data analysis

Statistical significance between means \pm SEM was analyzed by a two-way ANOVA and Student's *t* test for paired or unpaired data. Experimental series performed on different days were compared by the Student–Newman–Keuls *t* test. A value of *p* < 0.05 was considered to be statistically significant.

Results

Loss and recovery of SVZ NPC proliferation and neuroblast formation after MPTP treatment

To assess the contribution of the inflammatory reaction in the SVZ response, we first verified, *in vivo*, spatiotemporal changes in proliferation and neuroblast formation at different time intervals (1–42 d) during loss and recovery of nigrostriatal DAergic neurons, according to the acute MPTP injection paradigm (L'Episcopo et al., 2011a), and correlated the temporal changes in the intrinsic NPC proliferation capacity, *ex vivo* (Fig. 1A). As reported in both rodent and primate models of PD (Baker et al., 2004; Borta and Hoglinger, 2007; O'Keefe et al., 2009a), we found that MPTP induced a significant decrease of PCNA⁺ and

BrdU⁺ (Fig. 1B,D) starting by 1 d after MPTP injury through 14 dpt (from 16.5 ± 2.0 of saline mice to $6.0 \pm 0.5 \times 10^3$ PCNA⁺ cells measured 14 dpt).

The reduced proliferation of MPTP mice was accompanied by a severe reduction of DCX⁺ neuroblasts (Fig. 1B,G) 1–14 dpt (from 870 ± 25 of saline mice to $280 \pm 35 \times 10^3$ DCX⁺ cells measured 14 dpt). The temporal window of neurogenic impairment was associated with the DAergic degeneration phase, as measured by the almost 70–80% depletion of DAT protein in Str, paralleled by a comparable decrease in high-affinity DA uptake by striatal synaptosomes (Table 1). At the midbrain level, the cumulative MPTP dose of 80 mg kg⁻¹ resulted in a severe ($\geq 60\%$) loss of DAergic cell bodies measured 1–21 dpt (L'Episcopo et al., 2011a).

Importantly, between 14 and 28 dpt, a significant recovery of cell proliferation (Fig. 1B,D) was observed in SVZ, reaching pre-MPTP values by 35 and 42 dpt (13.0 ± 1.2 and $15.0 \pm 1.5 \times 10^3$ PCNA cells measured at 35 and 42 dpt, respectively). Likewise, DCX⁺ cells (Fig. 1B,G) exhibited a significant recovery between 14 and 28 dpt, returning back to normal by 35 and 42 dpt (798 ± 35 and $875 \pm 48 \times 10^3$ DCX⁺ cells at 35 and 42 dpt, respectively) and coinciding with progressive striatal DAergic reinnervation (Table 1). At the midbrain level, a reduced ($\leq 38\%$) DAergic cell loss was observed by 35–42 dpt (L'Episcopo et al., 2011a).

As observed previously in nigrostriatal lesioned rodents (Mao et al., 2001; L. W. Chen et al., 2002, 2004; Morale et al., 2004, 2006; Mohapel et al., 2005; L'Episcopo et al., 2010a,b, 2011a,b) and primates (Tandé et al., 2006), astrocyte cell number was significantly increased in the Str of MPTP mice, a phenomenon thought to result from an early reaction of gliogenic precursor cells intrinsic to the striatum (Tandé et al., 2006; Borta and Hoglinger, 2007). Here, the specialized GFAP⁺ astrocytes (B-cells) adjacent to the SVZ, ensheating BrdU⁺ cells in saline-injected mice, were also markedly affected after MPTP injury (Fig. 1B): GFAP⁺ astrocytes were increased (from 120 ± 12 of saline mice to $205 \pm 16 \times 10^3$ GFAP⁺ cells measured 14 dpt) and displayed a highly reactive phenotype, as revealed by hypertrophic cell bodies and increased thickness and length of processes, resulting in an overall disruption of GFAP⁺–BrdU⁺ cell interactions in SVZ (Fig. 1B,E). With time, however, GFAP⁺ astrocytes shifted to a less reactive phenotype, and morphologic interactions with NPCs in SVZ were seemingly restored back to saline-injected controls (Fig. 1B). SVZ astrocytes are infrequently dividing stem cells, which generate frequently dividing, transit amplifying C-cells, which in turn produce restricted neuroblasts, A-cells (Doetsch et al., 1997, 1999). Previous studies have identified epidermal growth factor as a crucial regulator of SVZ expansion, and EGF-R⁺ cells in the SVZ were shown to correspond mainly to the rapidly cycling transit amplifying C-cells (Doetsch et al., 2002). Given the indication that C-cell proliferation is decreased after MPTP (Hoglinger et al., 2004; O'Keefe et al., 2009a,b), we next analyzed sections double stained for BrdU/PCNA and the neuroblast markers DCX, BrdU/PCNA, and GFAP, as well as BrdU/PCNA and EGF-R as C-cell markers (Doetsch et al., 2002; Hoglinger et al., 2004; O'Keefe et al., 2009b) (Fig. 1C,F,H–J). In accordance with previous studies, the percentage of DCX⁺ cells expressing BrdU (Fig. 1H), or the percentage of GFAP⁺ cells expressing BrdU (Fig. 1F), did not change significantly during the window of SVZ impairment, whereas an increase in GFAP⁺ cells expressing PCNA was observed in the Str of MPTP mice, within 24 h of MPTP injury (Fig. 1B). Although, not specifically addressed in the present study, we did not find clear evidence of a cell density gradient from the Str to the SVZ, or vice versa, 1–14

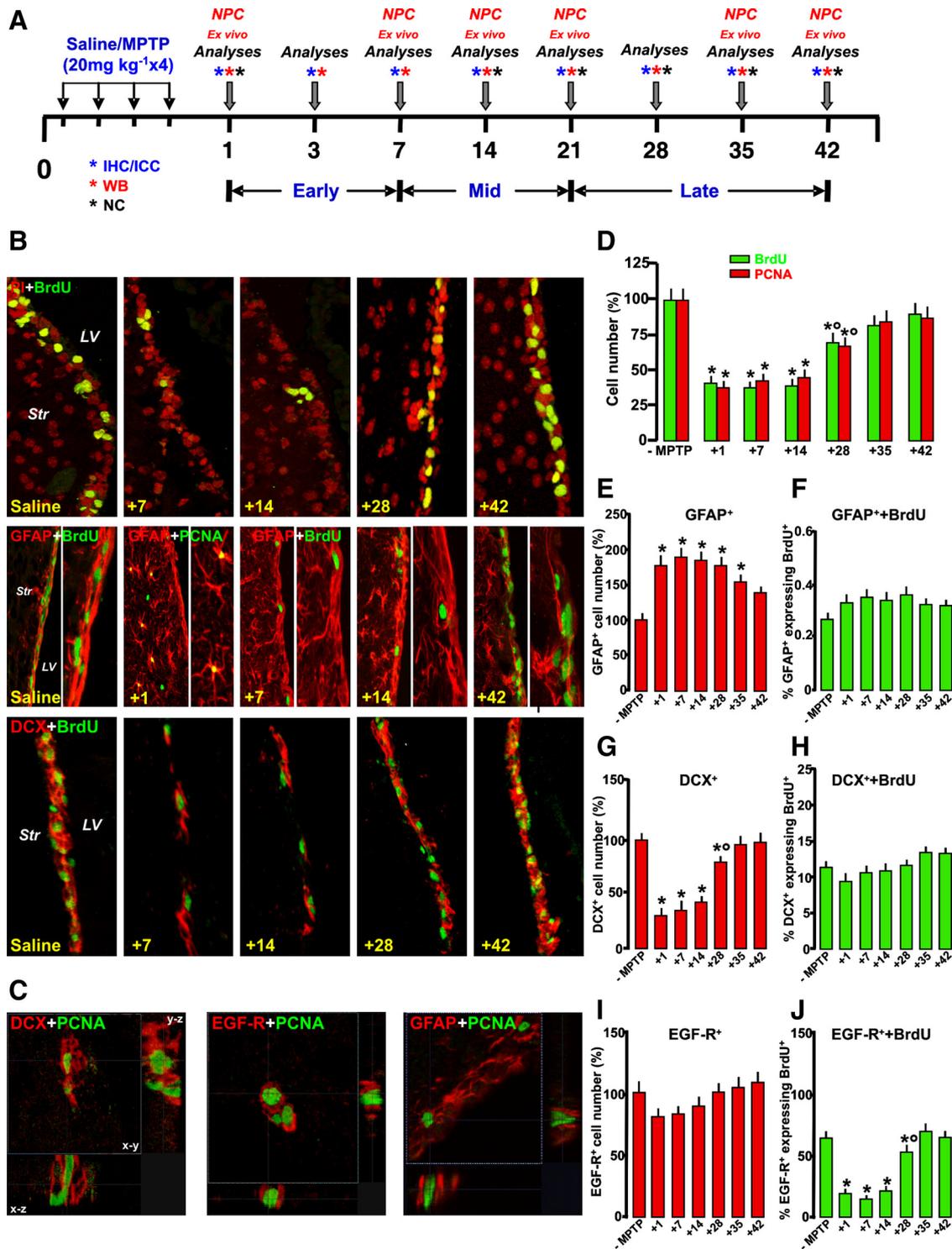


Figure 1. Loss and recovery of SVZ neuroprogenitor proliferation and neuroblast formation after MPTP treatment *in vivo*. **A**, Scheme of experimental design showing time points studied after saline and MPTP treatments (7–8 mice per time point). For *ex vivo* studies of neuroprogenitors, the SVZ region was microdissected and aseptically prepared for isolation and culture of NPCs. For immunohistochemistry (IHC), groups of mice were anesthetized and transcardially perfused (6 mice per time point). For neurochemical (NC) and Western blot (WB) analyses (5 mice per time point), the brains were rapidly dissected and stored at -80°C . ICC, Immunocytochemistry. **B**, Representative confocal images of BrdU⁺ (green) counterstained with the nuclear marker PI (red), and dual staining with GFAP (red) and BrdU (green), with GFAP (red) and PCNA (green), and with DCX⁺ (red) and BrdU (green). **C**, Orthogonal reconstructions of confocal microscopic images of dual staining with DCX and PCNA, EGF-R and PCNA, and GFAP and PCNA in the x–y, x–z, and y–z planes. **D–J**, Quantification of proliferation as assessed by numbers of BrdU⁺ and PCNA⁺ cells (**D**); GFAP⁺ cell numbers (**E**) and percentage of GFAP⁺ expressing BrdU⁺ (**F**); DCX⁺ cell number (**G**) and percentage of DCX⁺ expressing BrdU⁺ (**H**); and EGF-R⁺ cell numbers (**I**) and percentage of EGF-R⁺ expressing BrdU⁺ (**J**). Differences were analyzed by ANOVA, followed by the Newman–Keuls test, and were considered significant when $p < 0.05$. * $p < 0.05$ versus saline; ° $p < 0.05$ versus 1–14 or 1–7 dpt, within each respective group.

Table 1. MPTP-induced loss and recovery of dopaminergic end points in striatum

Analyses	MPTP time course						
	MPTP	+1	+14	+21	+28	+35 d	+42 d
DAT immunoreactivity (FI, % of control)	100 \pm 15	22 \pm 8*	25 \pm 10*	38 \pm 9*	51 \pm 9*	65 \pm 10*	75 \pm 12
[³ H]DA uptake (% of control)	100 \pm 18	31 \pm 10*	29 \pm 8*	35 \pm 10*	45 \pm 10*	58 \pm 12*	68 \pm 12

Male C57BL/6 mice received $n = 4$ intraperitoneal injections of vehicle (saline, 10 ml/kg) or MPTP-HCl (20 mg kg⁻¹ free base; Sigma-Aldrich) dissolved in saline, 2 h apart in 1 d. For each set of analyses, five mice were used for the indicated time points. The brains were processed as indicated for DAT immunohistochemistry. Fluorescence intensity values (means \pm SEM) are expressed as percentage of saline. Synaptosomal high-affinity [³H]DA uptake in striatum (total high-affinity and mazindol noninhibitable) was determined as described, and the values are represented by the changes in DA uptake (expressed as percentage of control). Differences were analyzed by ANOVA, followed by the Newman–Keuls test, and were considered significant when $p < 0.05$. * $p < 0.05$ versus MPTP.

dpt, albeit migration of GFAP⁺ cells from the adjacent Str might explain the observed increased density of astrocytes bordering the SVZ of MPTP mice. On the other hand, the percentage of EGF-R⁺ cells expressing BrdU was significantly decreased during the temporal window of neurogenic impairment (Fig. 1J), whereas by 28 dpt, the percentage of EGF-R⁺/BrdU⁺ cells increased to recover back to normal pre-MPTP levels within 35–42 dpt (Fig. 1J), supporting that the MPTP-induced acute reduction of EGF-R⁺ proliferation might reduce the generation of DCX⁺ cells. In addition, the SVZ recovery phase involved a normalization of GFAP⁺–BrdU⁺ cell interactions and increased EGF-R⁺ cell proliferation, possibly resulting in generation of new DCX⁺ neuroblasts (Fig. 1B).

We next set to identify possible differences in intrinsic properties between NPCs from saline- and MPTP-treated mice at the indicated time intervals, by studying the self-renewal of CNS stem/progenitor cells by using an *ex vivo* assay. It has been shown that neurosphere-forming cells consist mostly of type C (and type B) cells *in vitro* (Morshead et al., 1994). Therefore, we used a well established *in vitro* neurosphere assay to further characterize the SVZ stem cell population of MPTP mice, with the numbers of clonal neurospheres *in vitro* as a measure of the absolute number(s) of putative stem cells *in vivo* (Morshead et al., 1994; Rietze and Reynolds, 2006; Pluchino et al., 2008). We observed a significant reduction of the clonal efficiency of SVZ-derived primary neurospheres at all time points after MPTP ($p < 0.005$, compared with saline-treated control mice) (Fig. 2A). This difference did peak at 14 and 21 d after MPTP, with the clonal efficiency of neurospheres from MPTP-treated mice being threefold lower than that of control neurospheres (Fig. 2A). This decline in clonal efficiency was further confirmed in secondary neurospheres at the very same time points (Fig. 2A; $p < 0.005$, compared with saline-treated control mice), thus further corroborating our own *in vivo* observation that significant impairment of the SVZ NPC compartment occurs at certain time points after MPTP. Interestingly, neurospheres (both primary and secondary) from MPTP-treated mice did not show differences in size at any time point, compared with neurospheres from control mice (Fig. 2B), thus suggesting that SVZ stem cells from MPTP-treated mice do not show intrinsic differences in their mitogenic capacity from cells derived from controls. Finally, we sought to investigate the proliferation capacity of NPCs from MPTP mice under chemically defined serum-free culture conditions *in vitro*. In line with clonal analyses, neurospheres from MPTP-treated mice between 24 h and 14 d did show again significant early impairment in their growth rate that was, however, completely recovered after 12 d *in vitro* in the presence of FGF-II and EGF, thus further confirming that there is not a cell-autonomous process accounting for the impairment in proliferation observed *in vivo* (Fig. 2C).

From both the *in vivo* and *ex vivo* findings, the present time course study indicated that MPTP-induced impairment of neurogenic potential may be attributable to environmental factors rather than to the intrinsic properties studied on NPC progeny.

In addition, the characterized biphasic response of the SVZ after MPTP injury accompanying the early degeneration and the late recovery of nigrostriatal DAergic neurons thus set the temporal window for the study of the influence of neuroinflammation in SVZ plasticity.

MPTP-induced upregulation and downregulation of inflammatory signals in striatum and SVZ is inversely correlated to loss and recovery of NPC proliferation

The well established strong and localized microglial reaction in Str of MPTP-treated mice (see L'Episcopo et al., 2011a) was next temporally correlated with NPC proliferation in SVZ. To this end, we classified the stages of microglia activation according to Kreutzberg (1999) and counted stage 4 IBA1⁺ microglia, exhibiting a round-shaped cell body without processes, as opposed to quiescent stage 1 microglia, with elongated soma and thin/ramified processes (L'Episcopo et al., 2011c). As observed in Figure 3, A and B, an inverse correlation between stage 4 IBA1⁺ microglia and PCNA⁺ cells in the SVZ started as early as after 1 dpt (Fig. 3B), when a sharp increase of activated amoeboid-shaped IBA-1⁺ microglia in both Str and SVZ was detected (Fig. 3A,B), compared with saline-injected mice, where only quiescent stage 1 IBA1⁺ microglia with elongated cell body and ramified processes were seen (Fig. 3A). Here again, numerous PCNA⁺/IBA1⁺ cells were found early (within 24 h) in Str of MPTP mice (Fig. 3A), suggesting their possible migration toward the adjacent SVZ. Correlation of microglia activation with SVZ proliferation indicated that a maximal decrease of PCNA⁺ cells in SVZ correlated with the peak of reactive IBA1⁺ cells, whereas rescue of PCNA⁺ cells coincided with reactive IBA1⁺ cell abatement (Fig. 3B). By this time, IBA-1⁺ microglia with thinner and longer processes were observed in both SVZ and Str for the rest of the experimental period (Fig. 3A). Overactivation of microglia temporally related to NPC impairment was also indicated by the early and sharp increase of Mac-1 (Fig. 3C) and phagocyte oxidase (PHOX) (Fig. 3D) and by increased IBA1⁺ cells expressing iNOS in Str and SVZ (Fig. 3E,F). The severe striatal pro-oxidant/proinflammatory status herein observed is in line with our previous findings revealing upregulation of major oxidative/inflammatory mRNA species (including PHOX, NOs2, NOs3, Hmox1, and UCP2) peaking 3–24 h after MPTP (L'Episcopo et al., 2010b, 2011a). Conversely, PCNA⁺ cell return to pre-MPTP levels (Fig. 3B) was preceded by downregulation of oxidative and proinflammatory mediators in Str (Fig. 3C,D,F). When PHOX- and iNOS-derived NO are active at the same time, then microglia might produce peroxynitrite (ONOO⁻), a potent toxin that may promote nitration of various proteins, and produce hydroxyl radicals that may impair mitochondrial functions (Gao et al., 2003, 2008; Mander and Brown, 2005; Gao and Hong, 2008; Hu et al., 2008). Given the vital role of astrocytes in brain homeostasis, particularly as ROS, peroxynitrite, and glutamate scavengers (Magistretti, 2006; Moncada and Bolanos, 2006; Chen et al., 2009; Sandhu et al., 2009), we then performed dual staining with

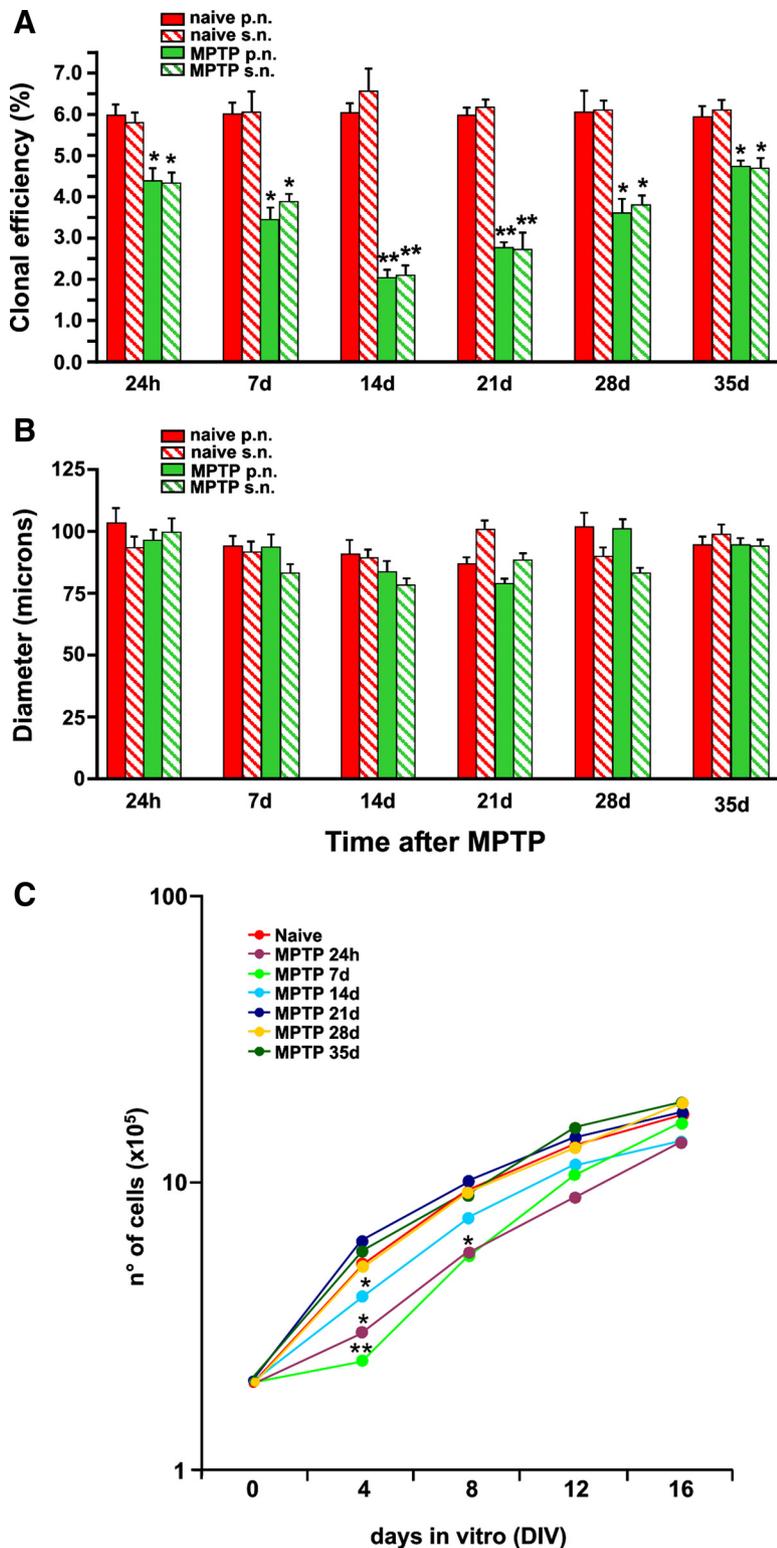


Figure 2. Analyses of intrinsic properties of SVZ neuroprogenitor isolated *ex vivo* from vehicle and MPTP mice. At the indicated time intervals, brain coronal sections were taken from vehicle- and MPTP-treated mice, and the SVZs were rapidly dissected out and processed as described. **A**, Quantitative analysis of the clonal efficiency of neurospheres from naive and MPTP-treated mice. Data are represented as mean percentage of colonies per size over total plated cells (\pm SEM) and have been obtained from a total of $n \geq 3$ independent experiments. **B**, Size of neurospheres from naive and MPTP-treated mice. Data are represented as mean numbers (micrometers) (\pm SEM) and have been obtained from a total of $n \geq 3$ independent experiments. Red in **A** and **B** represents neurospheres from naive mice, whereas green represents neurospheres from MPTP-treated mice. Filled bars in **A** and **B** represent primary neurospheres, whereas wide upward diagonal bars represent secondary neurospheres. **C**, Growth rate analysis of NPC lines established from the SVZ of either naive or MPTP-treated mice. * $p \leq 0.005$ and ** $p \leq 0.0001$, compared with time point-matched controls. p.n., Primary neurosphere; s.n., secondary neurosphere.

GFAP and 3-NT, as a fingerprint of iNOS-derived NO and peroxynitrite generation. Hence, a sharp increase in the 3-NT-immunofluorescent (IF) signal in striatal astrocytes and SVZ was observed shortly after MPTP, but not saline, injection (Fig. 3E). This increase in 3-NT immunofluorescence, supported by Western blot analysis (Fig. 3G), reached a peak 3 dpt and subsided by 21 dpt, indicating astrocyte nitrosylation during the early window of NPC neurogenic impairment. Because previous studies showed MPTP-induced apoptosis in the SVZ (He et al., 2006; Shibui et al., 2009), and given the cytotoxic role of increased peroxynitrite generation, we next localized the death marker cleaved Caspase3. Double localization of the neuronal cell marker NeuN (green) with Caspase3 (red) in the striatal SVZ clearly documented the increased Caspase3-IF signal specifically localized in the SVZ but not in Str of MPTP mice (Fig. 3E) during the peak of microglia activation, while disappearing by 7 dpt (data not shown), thereby indicating that a certain proportion of SVZ cells might be acutely damaged by MPTP challenge.

All together, these data indicated that MPTP-induced NPC impairment, *in vivo* and *ex vivo*, was preceded and accompanied by increased microglial inflammatory mediators and astrocyte nitration both in Str and SVZ, whereas a downregulation of striatal oxidative and inflammatory status preceded the recovery from neurogenic impairment, thus raising the question of the distinct roles played by the neurotoxin MPTP, activated astrocytes, and microglia.

MPP⁺ directly impairs NPC survival, proliferation, and/or differentiation in a dose-dependent fashion

Previous studies proposed a possible dopamine-independent effect of MPTP-induced neuroblast apoptosis in the SVZ, but the cellular contributors and the signaling pathways involved in this process are currently unknown. It is generally accepted that the neurotoxin MPTP, converted into its active metabolite, MPP⁺, in astrocytes, is selectively transported into striatal DAergic terminals via the DAergic transporter DAT, where it induces oxidative stress, the opening of mitochondrial permeability transition pore, the release of cytochrome c, and the activation of caspases (Jackson-Lewis and Przedborski, 2007). In synergy with these early events accounting for ~10% of DAergic neuronal death (Wu et al., 2002), glial inflammatory mechanisms are

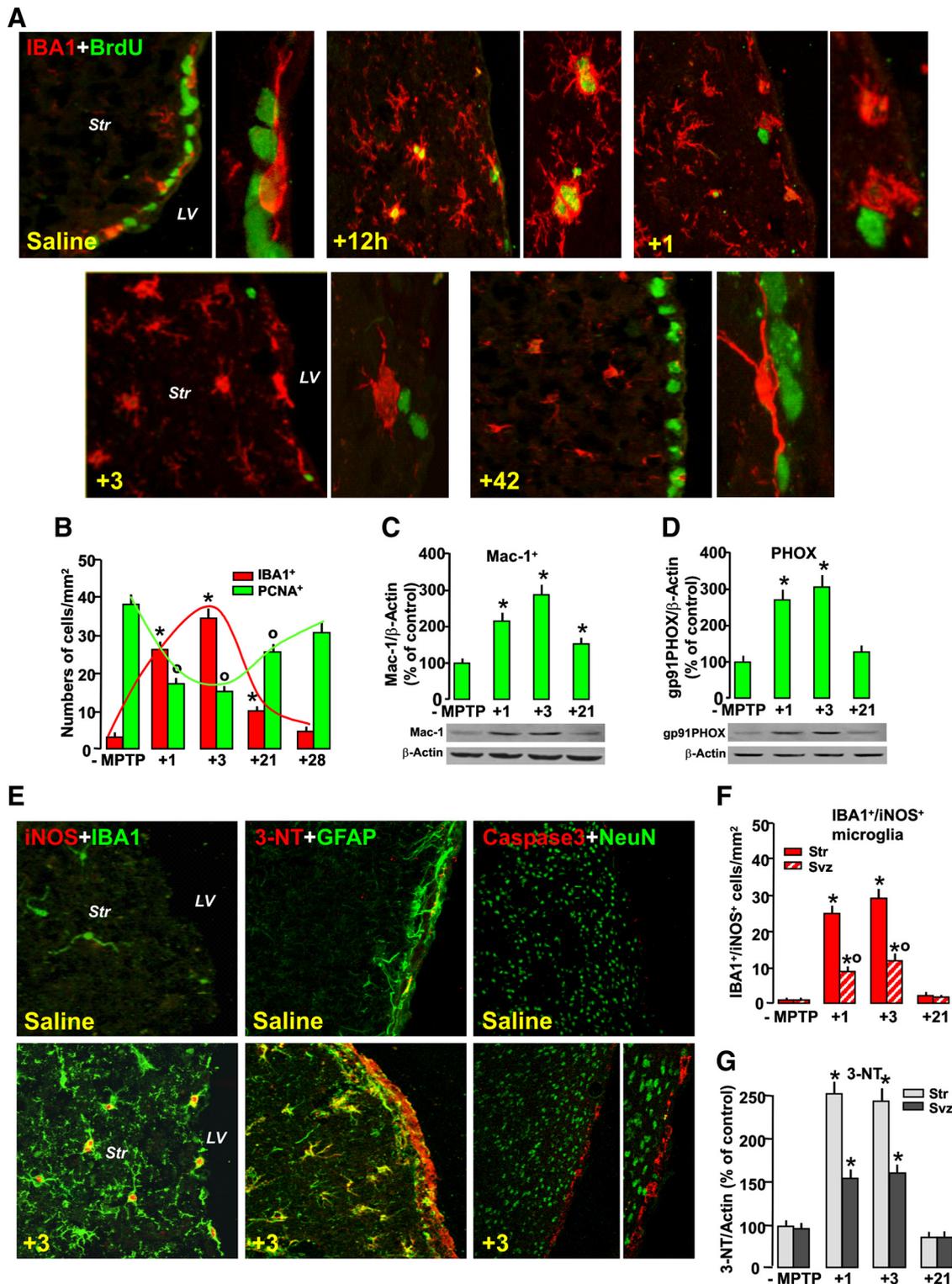


Figure 3. Inverse correlation between MPTP-dependent microglial activation and cell proliferation in SVZ. Mice treated with saline or MPTP were killed, and the brain was processed for immunohistochemical or Western blot analyses. **A**, Representative confocal images showing dual staining with IBA1 (red) and BrdU (green) in saline, 1, 3, and 42 d after MPTP. Note the juxtaposition of stage 1 IBA1⁺ cell soma and process with BrdU⁺ cells in the SVZ of saline mice. Within 12 h from MPTP-activated IBA1⁺, microglia expressing PCNA are observed in the Str. By 24 h and 3 dpt, stage 4 IBA1⁺ microglia are seen both in Str and SVZ. By 42 dpt, BrdU⁺ in the SVZ and IBA1⁺ microglia recovered back to saline-treated controls. **B**, Inverse correlation between NPC proliferation in the SVZ measured with PCNA (numbers of PCNA⁺ cells/10³) and stage 4 IBA1⁺ microglia (cell number per cubed millimeter). Differences were analyzed by ANOVA, followed by the Newman–Keuls test, and were considered significant when $p < 0.05$. * $p < 0.05$ IBA1⁺ cells versus saline and $p < 0.05$ PCNA⁺ cells versus saline, within each experimental group. **C, D**, Western blotting of Mac-1 (**C**) and PHOX (**D**) in Str after saline or MPTP injection. Data from experimental bands were normalized to β -actin, before statistical ANOVA, and values were expressed as percentage of saline-injected controls. Differences were analyzed as above. ^o* $p < 0.05$, compared with saline. **E**, Dual immunostaining with iNOS (red) and IBA1 (green), 3-NT (red) and GFAP (green), and cleaved Caspase3 (red) and NeuN (green) in Str and SVZ in saline and 3 dpt. Note the increased iNOS expression in IBA1⁺ microglia, increased 3-NT colocalization (orange to yellow) in Str astrocytes and SVZ, and increased Caspase3 localization in SVZ, 3 d after MPTP. **F**, IBA1⁺/iNOS⁺ cell counts 1, 3, and 21 dpt. **G**, Immunoblotting shows increased 3-NT in Str and SVZ protein extracts peaking 3 dpt and subsiding by 21 dpt.

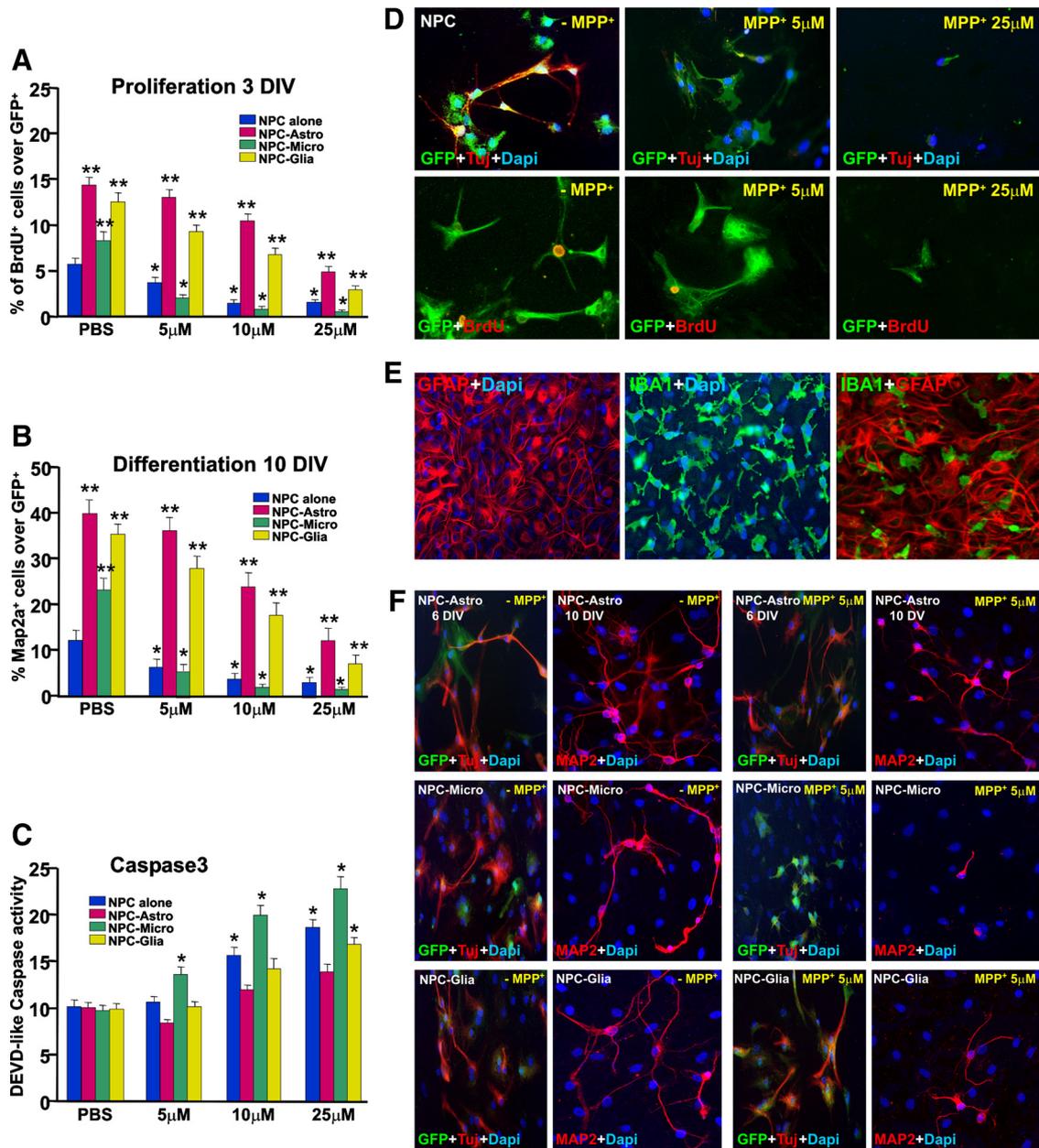


Figure 4. Effect of MPP⁺ with or without glial cocultures on NPC survival, proliferation, and differentiation *in vitro*. The effect of increasing doses of MPP⁺ (5–50 μM) in the proliferation (**A**), differentiation (**B**), and Caspase3 activation (**C**) of adult GFP-expressing NPCs derived from the adult SVZ cultured alone (**D**) or cocultured with purified astrocytes, purified microglia, or mixed astrocyte–microglia preparations (**E, F**). **A**, NPC proliferation was assessed with BrdU as a marker for cell proliferation. **B, F**, Differentiation was measured at 6 DIV using TuJ1⁺ (**B**) and at 10 DIV using Map2a⁺ cells (**B, F**), compared with NPCs cultured alone (**B, D**). Exposure of NPC–Astro cocultures to MPP⁺ efficiently prevented MPP⁺-induced decreased TuJ1⁺ (**F**), BrdU (**A**), and Map2a expression (**B, F**), whereas in NPC–Micro cocultures, MPP⁺ inhibitory effects were not reversed (**A, B, F**). Caspase3 activity was measured using the indirect coculture paradigm (**C**). ***p* < 0.05 versus NPC alone; **p* < 0.05 versus PBS. Astrocyte inserts efficiently reversed MPP⁺-induced increase of DEVD-like fluorescent signal at 10 and 20 μM, whereas microglial inserts did not afford NPC protection (**C**). In mixed astrocyte–microglial cultures (NPC–glia, **F**), increasing astrocyte-to-microglia ratio resulted in astrocyte-dependent reversal of microglia inhibitory effects on NPC proliferation, differentiation, and survival as determined by Caspase3 activity (Fig. **A–C, F**).

thought to contribute to nigrostriatal DAergic degeneration (Hirsch and Hunot, 2009). We thus thought to study a possible direct toxic effect of MPP⁺ on NPC and addressed the contribution of glial-mediated mechanisms, by modeling glia–NPC cross talk *in vitro*. To this end, we used GFP-expressing NPCs derived from the adult SVZ (Pluchino et al., 2008; L'Episcopo et al., 2011a) exposed to increasing doses (5–25 μM) of MPP⁺ both in

the absence or presence of glial cells in different coculture paradigms, as described, and assessed NPC proliferation (Fig. 4A) and differentiation (Fig. 4B). The doses of MPP⁺ were selected according to our *in vitro* studies on primary mesencephalic DAergic neurons in culture (L'Episcopo et al., 2011a,b). In addition, we determined Caspase3 activity, a marker of MPTP/MPP⁺-induced cell death, using the fluorogenic substrate

DEVD-AFC (Fig. 4C). Both NPC monotypic and glial–NPC cocultures received increasing doses (5–25 μ M) of MPP⁺ (L'Episcopo et al., 2001b). For proliferation studies, MPP⁺ was applied at 2 DIV, the nucleotide analog BrdU (5 μ M) was next added, and the cells were fixed after 24 h, i.e., at 3 DIV. For differentiation studies, NPCs grown alone, or layered on top of astrocytes, microglia, or mixed astrocyte–microglial preparations, shifted in differentiation medium (N2 medium and 1% FCS), and cells were allowed to differentiate for 5–10 DIV. MPP⁺ was then applied and cells were fixed after 24 h, with Tuj1 and MAP2a being used as specific neuronal markers. For cell-survival experiments, Caspase3 activity was determined 8 h after MPP⁺.

As observed, GFP⁺ NPCs grown in poly-D-lysine alone in N2 medium and in the presence of 1% of serum express TUJ-1, and some of the GFP⁺ NPCs coexpress BrdU after 3 DIV (Fig. 4A, D). The effect of MPP⁺ was dose dependent: the lowest (\leq 5 μ M) dose did not significantly modify the number of GFP⁺ NPCs over the DAPI-stained nuclei, whereas an almost 30–60% decrease in GFP⁺ NPC number was observed at 10 and 25 μ M doses of MPP⁺, compared with PBS-treated controls. The proliferative potential revealed by the percentage of GFP⁺ NPCs expressing BrdU (Fig. 4A, D) and neuronal differentiation, as determined by the percentage of Tuj1⁺ at 6 DIV (Fig. 4D) and MAP2⁺ neurons at 10 DIV (Fig. 4B) over the GFP⁺ NPCs were dose-dependently reduced by MPP⁺, compared with PBS-treated controls. Finally, the DEVD-like fluorescent signal (Fig. 4C) was dose-dependently increased 8 h after application of 10 and 25 μ M MPP⁺, but not 5 μ M, implicating Caspase3 activation at high MPP⁺ doses. Although different factors play a role in MPTP/MPP⁺-induced neurogenic impairment *in vivo*, including complex cellular contributions and different time courses, the present results demonstrated the ability of the PD neurotoxin to directly impair adult SVZ neuroprogenitor survival, proliferation, and differentiation *in vitro*, in a dose-dependent fashion.

Astrocytes and microglia differentially modulate MPP⁺-induced impairment of NPCs, *in vitro*

Because astrocytes and microglia are recognized to importantly influence MPTP/MPP⁺ toxicity after DAergic neurons, both *in vivo* and *in vitro* (Gao et al., 2003; Hu et al., 2008; Chen et al., 2009; Sandhu et al., 2009; L'Episcopo et al., 2011a,b), we next investigated a possible modulatory effect of glial cells after MPP⁺-induced NPC toxicity. To this end, we studied the effect of astrocyte (>95% GFAP⁺ astrocytes), microglia (>95% IBA1⁺ microglia), or mixed glial (\geq 65 GFAP-IR astrocytes; \leq 24% IBA1-IR microglia) (Fig. 4E) cell preparations cocultured with NPCs in the absence or presence of MPP⁺ (Fig. 4F), as above. In basal conditions, direct coculture of NPCs with purified astrocytes had a robust effect on both proliferation (Fig. 4A) and neuroblast formation at either 6 or 10 DIV (Fig. 4F), as indicated by the almost threefold increase in the proportion of BrdU⁺ (Fig. 4A), Tuj1⁺ (Fig. 4F), and MAP2⁺ (Fig. 4B, F) out of the GFP⁺ cells, compared with NPCs cultured alone. Coculture with purified microglia increased by almost twofold the proportion of MAP2⁺ (Fig. 4B, F) out of the GFP⁺ cells and increased the proportion of BrdU⁺ (Fig. 4A) cells out of the GFP⁺ cells by almost 50%, compared with NPC cultured alone. These findings are in agreement with previous studies, showing the ability of astrocyte and microglial monolayers (Lim and Alvarez-Buylla, 1999; Barkho et al., 2006; Butovsky et al., 2006; L'Episcopo et al., 2011a) to promote neurogenesis from adult NPCs. Interestingly, exposure of NPC–Astro cocultures to MPP⁺ efficiently prevented MPP⁺-induced decreased Tuj1⁺ (Fig. 4F) and MAP2a⁺

(Fig. 4B, F) neuron production and BrdU expression (Fig. 4A), mimicking astrocyte-induced mesencephalic DAergic neuron protection (L'Episcopo et al., 2011a,b), albeit the effect of astrocytes decreased as a function of the concentration of MPP⁺. In sharp contrast, in NPC–Micro cocultures, MPP⁺ inhibitory effects were not reversed (Fig. 4A, B, F). Hence, the decrease in the percentage of Tuj1⁺ and MAP2a⁺ cells observed at both 6 and 10 DIV (Fig. 4B, F), as well as the percentage of GFP⁺ cells labeled with BrdU (Fig. 4A) after MPP⁺, were comparable both in the absence or presence of microglia. These findings suggested that in the presence of MPP⁺, the microglia neurogenesis-promoting effects were inhibited. To verify whether the coculture influenced NPC survival, we measured Caspase3 activity using the indirect coculture paradigm. In this model, the inserts containing the astrocyte or microglia monolayers were added on top of the NPCs. Hence, whereas astrocyte inserts efficiently reversed MPP⁺-induced increase of the DEVD-like fluorescent signal at 10 and 25 μ M, microglial inserts did not afford NPC protection. On the contrary, a greater increase in DEVD-like IF signal (Fig. 4C) was observed in NPC–microglia cocultures at all MPP⁺ doses. The positive effect of reactive astrocytes versus microglia after MPP⁺ exposure was also observed in mixed astrocyte–microglial cultures (NPC–glia), where the increasing astrocyte-to-microglia ratio resulted in a significant reversal of MPP⁺-induced reduced NPC survival (Fig. 4C), proliferation (Fig. 4A), and differentiation (Fig. 4B, F).

These findings indicated that although astrocytes are endowed with protective and neurogenic capabilities both in the absence or presence of MPP⁺, microglia can increase NPC neurogenic potential in basal conditions, whereas in the presence of MPP⁺ they further impair NPC survival, raising the question of the specific factors and the signaling pathways involved in microglial and astrocyte observed effects.

MPTP-induced increased microglial oxidative and nitrosative status impairs NPC proliferation and differentiation *in vitro*

To address the contribution of microglia in MPTP-induced SVZ impairment *in vivo*, and to investigate the nature of the microglial-derived factors involved, we sought to use macrophage/microglia acutely isolated *ex vivo*, from saline or MPTP. We selected 3 dpt since it corresponds to the peak of microglial reaction, *in vivo* (Fig. 3), and used these cell preparations for the determination of PHOX, ROS, and iNOS-derived RNS and for coculture experiments with GFP⁺ cells, as described.

In macrophage/microglia isolated from MPTP mice, PHOX expression was increased by \sim 3.5-fold compared with microglia isolated from saline-treated mice (Fig. 5A). Likewise, in line with previous studies (Morale et al., 2004; Hu et al., 2008; L'Episcopo et al., 2010a,b), the production of ROS and iNOS-derived RNS was significantly increased in macrophage/microglia isolated from MPTP mice, whereas in saline-treated mice very low to undetectable levels of ROS and RNS could be detected (Fig. 5B–D), supporting the *in vivo* data (Fig. 3). The direct implication of these mediators was next studied by exposing NPCs to either microglia (Fig. 5E) or microglial inserts (Fig. 5F) from saline or MPTP mice. A nonmicroglial insert (ct-insert) cultured in comparable conditions was used as a control. After 24 h, we found an almost threefold decrease in GFP⁺ cell incorporation of BrdU in MPTP–microglia–NPC, as opposed to microglia from saline-injected controls (Fig. 5E, F). Likewise, the number of MAP2a⁺ cells was sharply decreased compared with the number of neurons counted in NPCs exposed directly or indirectly to saline–microglia (Fig. 5E–H) or to ct-inserts (data not shown), supporting the

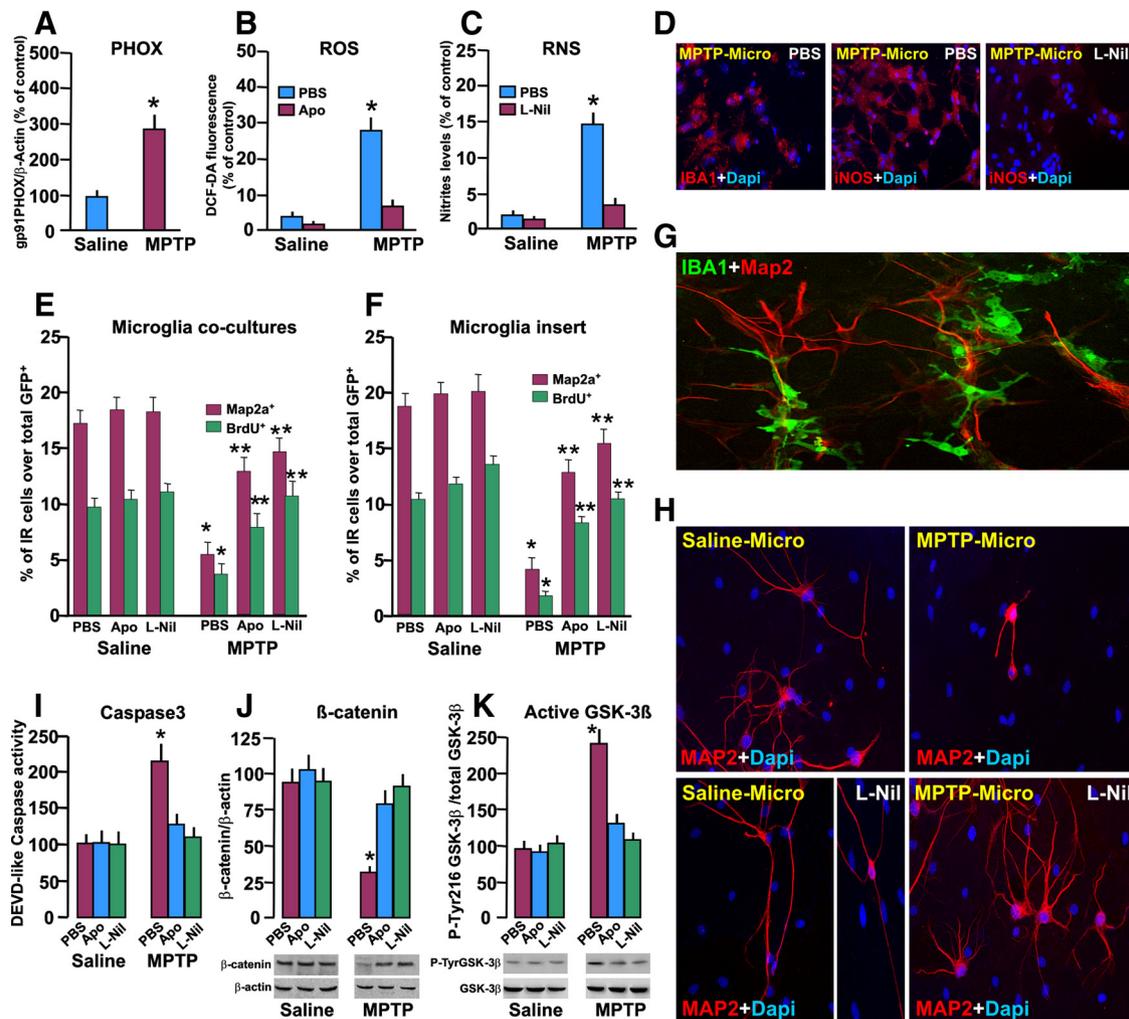


Figure 5. Effect of microglia isolated *ex vivo* from MPTP mice on NPC survival and proliferation/differentiation. Macrophage/microglia acutely isolated *ex vivo*, from saline or MPTP mice at 3 dpt, were processed as described for determination of PHOX, ROS, and RNS and for coculture with GFP⁺ NPCs for determination of proliferation and differentiation potential. Differences analyzed were considered significant when $p < 0.05$. **A–C**, Increased expression of PHOX by Western blotting (**A**), increased production of ROS revealed by the fluorogenic probe dichlorofluorescein diacetate (DCF-DA; **B**), and exacerbated iNOS-derived RNS (**C**) generation at the peak of microglial reaction (3 dpt), compared to macrophage/microglia from naive mice in the absence or presence of Apo (5 mM) or with L-Nil (50 μ M) treatment. * $p < 0.05$ versus saline. **D**, Representative images showing IBA⁺ microglial cells (red) counterstained with DAPI (blue) or iNOS (red) in basal conditions and after L-Nil (**E**, **F**). **E**, **F**, GFP⁺ cells were exposed to either microglia (**E**) or microglial inserts (**F**) from saline or MPTP mice, in coculture experiments for 24 h, and GFP⁺ cell incorporation of BrdU and changes in the number of Map2a⁺ cells were determined. * $p < 0.05$ versus saline; ** $p < 0.05$ versus PBS. **G**, Representative confocal image showing a coculture between microglial cells from saline mice stained with IBA1 (green) and NPCs from saline mice stained with Map2a (red) at 7 DIV. **H**, Representative images showing NPC stained with Map2a (red) and counterstained with DAPI (blue) in coculture with either saline–microglia or MPTP–microglia, with or without L-Nil treatment. Staining with Map2a shows the neurogenic-promoting effect of saline–microglia as opposed to the marked inhibition of neuron formation in coculture with MPTP–microglia, an effect reversed by L-Nil. Note the increased Map2a process extension in L-Nil-treated microglial–NPC coculture. **I**, DEVD-like fluorescent signal measured in NPCs cocultured with MPTP–microglia indicates activation of Caspase3-induced cell death. **J**, **K**, Western blot analysis of β -catenin (**J**) and active GSK-3 β (**K**; i.e., phosphorylated at tyrosine 216 residue, pTyr216 GSK-3 β). Data from the experimental bands were normalized to β -actin or GSK-3 β , as indicated, and values are expressed as percentage of PBS-treated cultures exposed to saline–microglia. * $p < 0.05$ versus saline; ** $p < 0.05$ versus PBS within experimental groups.

specificity of microglial-derived mediators in the observed inhibitory effects.

Further evidence for an inhibitory effect of MPTP–microglia was revealed by the sharp increase in the DEVD-like fluorescent signal measured in NPCs cocultured with MPTP–microglia (Fig. 5I), but not a ct-insert (data not shown), indicating activation of Caspase3-induced cell death. Thus, in the presence of MPTP-induced microglia overactivation *in vivo*, the harmful oxidative and inflammatory microenvironment likely increased NPC vulnerability, impairing NPC survival and, consequently, neurogenic potential. Indeed, free radicals such as superoxide and NO released by reactive microglia are the major players responsible for enhanced toxicity of MPTP after DAergic neurons both *in vivo* and *in vitro* (Gao et al., 2003, 2008; Mander and Brown, 2005; Hu et al., 2008; L'Episcopo et al., 2010a,b,c).

A PHOX–RNS–GSK-3 β /β-catenin-regulated signaling cascade is involved in microglial-induced inhibition of NPC neurogenic potential

Given the inhibitory role of NO in adult SVZ neurogenesis (Packer et al., 2003; Matarredona et al., 2004, 2005; Moreno-López et al., 2004; Torroglosa et al., 2007), we thus studied the effects of pharmacologically manipulating microglial oxidative and nitrosative status, by using the PHOX inhibitor apocynin and the specific inhibitor of iNOS-derived NO L-Nil, both in direct and indirect GFP⁺ cocultures established with microglia isolated *ex vivo* from either saline- or MPTP-treated mice, or with ct-inserts, as described. As observed, exposure of MPTP–microglia to Apo or L-Nil, significantly reduced PHOX-derived ROS and iNOS-derived RNS production, respectively (Fig. 5B–D). Accordingly, when compared with cocultures exposed to PBS,

MPTP–microglial–NPC cocultures exposed to Apo or L-Nil significantly mitigated MPTP–microglia-induced decreased proliferation and differentiation of GFP⁺ cells to levels measured in saline–microglial–NPC cocultures (Fig. 5E,F,H). On the other hand, the treated ct-inserts did not produce changes in either BrdU or Map2a expression in NPCs (data not shown). Interestingly, L-Nil treatment of microglial–NPC cocultures was characterized by increased MAP2a⁺ process length (Fig. 5H). In addition, we observed a significant counteraction of MPTP–microglia-induced Caspase3 activity (Fig. 5I) after Apo or L-Nil, whereas no effects followed coculture with treated ct-inserts, thereby supporting PHOX-derived ROS and iNOS-derived RNS as critical microglial harmful mediators of NPC neurogenic impairment.

We next addressed the mechanistic link between microglial harmful phenotype and signaling cascades impinging in NPC biochemical machinery. Recent studies on adult neural stem cell proliferation, migration of precursor cells, and neuronal differentiation have suggested a role for the *Wnt*/ β -catenin signaling system (Lie et al., 2005; Adachi et al., 2007; Zhang et al., 2011). Three different *Wnt* pathways have been described downstream of Frizzled receptor: the *Wnt*/ β -catenin pathway (known as the canonical pathway) and the noncanonical pathways, *Wnt*/Ca²⁺ and *Wnt*/planar cell polarity (Logan and Nusse, 2004; Gordon and Nusse, 2006). The hallmark of the *Wnt*/ β -catenin pathway is the stabilization of cytosolic β -catenin. GSK-3 β is part of a destruction complex that targets β -catenin for ubiquitination and degradation by the proteasome, whereas *Wnt* signaling inhibits GSK-3 β activity, thus increasing the amount of β -catenin, which enters the nucleus, and associates with T-cell factor/lymphoid enhancer binding factor (TCF/LEF) transcription factors, leading to the transcription of *Wnt* target genes involved in cell survival, proliferation, and differentiation (Logan and Nusse, 2004). We thus investigated the impact of microglial-derived mediators on the expression of β -catenin and GSK-3 β . Western blot analyses indicated a significant downregulation of β -catenin protein in NPCs exposed to MPTP–microglia compared with saline–microglia or ct-insert (Fig. 5J). Conversely, upregulation of phospho-Tyr216 GSK-3 β (i.e., the active pGSK-3 β) was observed 4 h after addition of the inserts containing MPTP–microglia (Fig. 5K). Time course experiments indicated an inverse correlation between upregulation of active pTyr216 GSK-3 β and the temporal loss of β -catenin protein in NPCs (data not shown), implicating activation of GSK-3 β and subsequent degradation of β -catenin in MPTP–microglial induced inhibition of NPC proliferation and neuronal differentiation. The implication of microglial-derived ROS and RNS in pTyr216 GSK-3 β upregulation and inhibition of neurogenesis were next investigated. In both Apo- and L-Nil-treated cultures, a significant counteraction of active GSK-3 β was observed (Fig. 5K). Accordingly, treatment with the ROS or iNOS inhibitors efficiently reversed the profound downregulation of β -catenin (Fig. 5J). Together, these results suggested that exacerbated oxidative and inflammatory microglial status may antagonize β -catenin signaling in NPCs and proposed a ROS–RNS/GSK-3 β / β -catenin signaling cascade contributing to microglial-induced inhibition of NPC neurogenic potential.

MPTP-induced dysregulation of *Wnt*/ β -catenin signaling *ex vivo* and *in vitro*

Wnt genes encode secreted glycoproteins and transduce signals via autocrine or paracrine mechanisms that regulate diverse developmental and postnatal processes, including neural stem proliferation and differentiation (Kalani et al., 2008; Kuwabara et al.,

2009; Munji et al., 2011). In the adult rodent, *Wnt* ligands (*Wnt*1, *Wnt*5a, *Wnt*7a) and Frizzled receptors are expressed in the SVZ (Shimogori et al., 2004). Of specific interest, using the *Axin2-d2EGFP* reporter (Jho et al., 2002), considered an accurate reporter mouse for canonical *Wnt* activity *in vivo*, Adachi et al. (2007) demonstrated that *Wnt*/ β -catenin signaling is activated in type B and C cells of the adult SVZ. Moreover, by injecting retroviral vectors to activate or inhibit β -catenin signaling specifically in dividing cells of the adult mouse SVZ, these authors also documented that activation of *Wnt*/ β -catenin signaling is sufficient to increase the percentage of dividing C-cells that give rise to new neurons in the SVZ. Importantly, β -catenin increases in the SVZ after intrastriatal TGF- α infusion in the 6-OHDA-lesioned rat PD model, suggesting a role for *Wnt* signals in the proliferative regulation of the SVZ (Cooper and Isacson, 2004). Coupled to our previously documented MPTP modulation of *Wnt*/ β -catenin signaling components in Str and midbrain (L'Episcopo et al., 2010a), these data prompted us to verify the hypothesis of a dysfunctional *Wnt*/ β -catenin signaling as a contributor of MPTP-dependent SVZ impairment. We first addressed changes in proliferative and differentiation potential associated with β -catenin and active GSK-3 β expression in NPCs acutely isolated *ex vivo* from MPTP-treated mice. NPCs isolated from naive and MPTP mice at 7 dpt were expanded, and proliferation/differentiation was measured as described. At 3 DIV, the capacity of NPCs isolated from MPTP mice to incorporate BrdU was reduced by threefold compared with NPCs from saline mice (Fig. 6A,B). To evaluate the neuronal differentiation potential of NPCs from saline and MPTP mice, cells were allowed to mature for 7 DIV in differentiation medium and then were stained with Map2a. Hence, NPCs from MPTP mice, *ex vivo*, produced Map2a⁺ neurons (Fig. 6A,B), albeit with a significantly lower efficiency, as revealed by the almost 50–60% decrease compared with Map2a⁺ NPCs from saline mice. In addition, Map2a⁺ process length was sharply reduced in MPTP compared with saline NPCs (Fig. 6A).

We next looked at *Wnt* signaling components using Western blot analysis and found a decreased β -catenin signal in NPCs from MPTP mice, compared with the β -catenin signal in NPCs from saline mice, whereas active the GSK-3 β signal was sharply increased (Fig. 6C). In keeping with these findings, using specific primers and quantitative real-time PCR, we found that expression levels of *Axin2*, a direct *Wnt* target induced by *Wnt*/ β -catenin activation (Jho et al., 2002), were downregulated in NPCs from MPTP mice compared with *Axin2* expression levels measured in NPCs from saline-injected controls (2.68 ± 0.31 arbitrary units in controls vs 1.15 ± 0.20 in MPTP; $p < 0.05$), thus corroborating MPTP-induced inhibition of *Wnt*/ β -catenin signaling activity in SVZ cells. These results appeared of interest in light of the recent report showing disruption of β -catenin signaling in glial progenitor cells from Alzheimer's disease (AD) and AD transgenic mice brain (He and Shen, 2009). We thus took advantage of the siRNA strategy (He and Shen, 2009; L'Episcopo et al., 2011b) to further examine the relationship between the β -catenin signaling pathway in MPTP-induced neurogenic impairment and tested the role of β -catenin and GSK-3 β . To this end, NPCs from MPTP mice were transiently transfected with GSK-3 β siRNA (Fig. 6D); alternatively, NPCs from saline mice were transiently transfected with β -catenin siRNA (Fig. 6E), as reported (He and Shen, 2009; L'Episcopo et al., 2011b). After 48–72 h of transfection, Western blot analysis indicated that treatment with GSK-3 β siRNA decreased GSK-3 β levels and increased the β -catenin levels in NPCs from MPTP mice (Fig. 6D), whereas

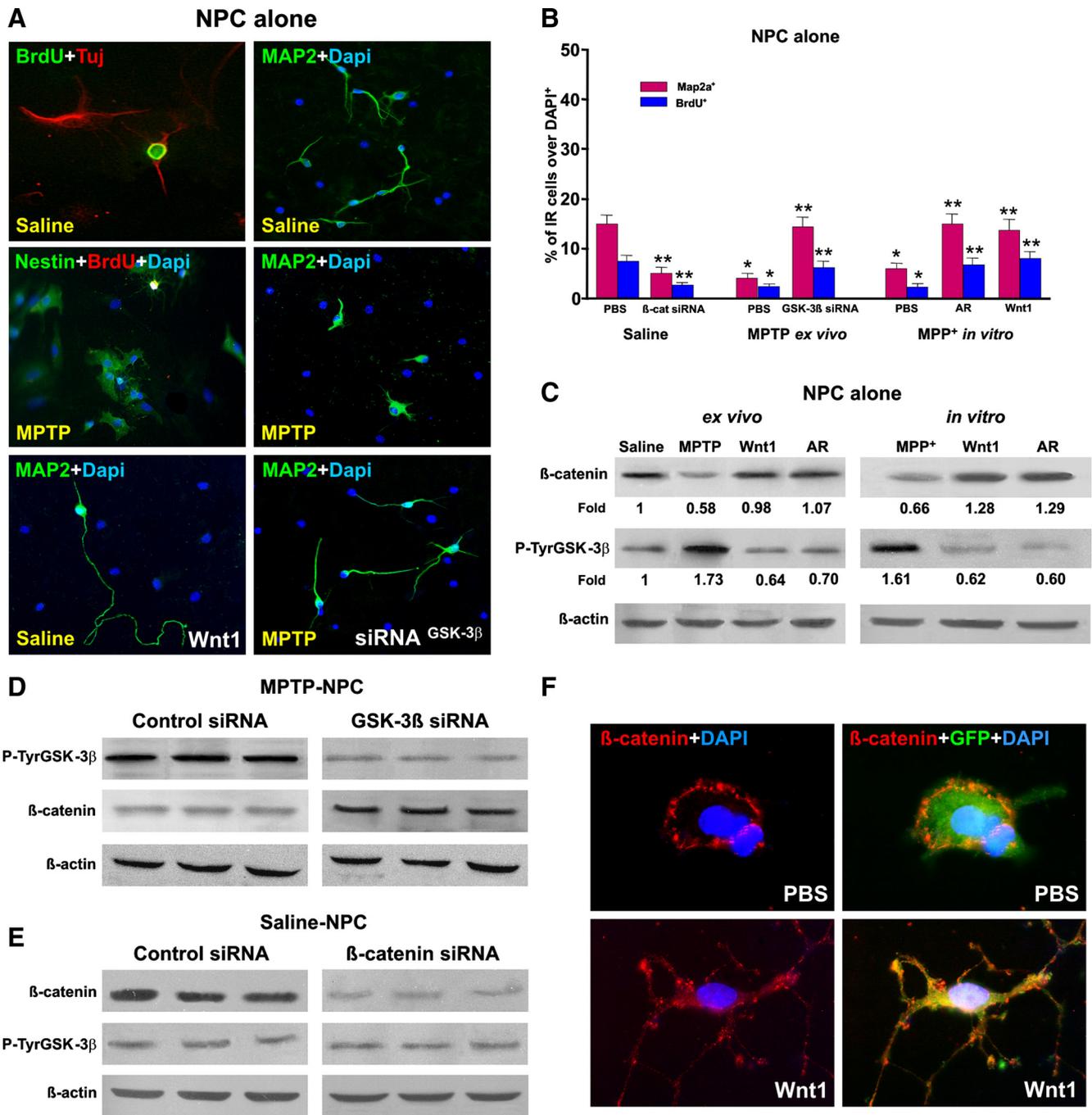


Figure 6. MPTP-induced impaired neurogenesis *ex vivo* and *in vitro* is associated to *Wnt*/ β -catenin signaling dysregulation. NPCs were acutely isolated *ex vivo* 7 d after MPTP or saline injection and expanded as described, and proliferation/differentiation was measured using BrdU, Tuj1, and Map2a. **A**, Representative immunocytochemical images of NPCs cultured alone and isolated from saline or MPTP, in the absence or presence of different treatments, as described. In NPCs from MPTP mice, a sharp loss of Map2a (green) and BrdU (red) was observed. **B**, Proliferation measured at 3 DIV and differentiation at 7 DIV in NPCs isolated *ex vivo* from saline and MPTP mice, and effect of MPP⁺ *in vitro*. NPCs of the indicated groups were exposed to different treatments, as described, and the percentage of BrdU⁺ or Map2a⁺ cells over the DAPI-stained nuclei was measured. **C**, Changes in β -catenin and P-TyrGSK-3 β studied in NPCs by Western blotting using the indirect coculture paradigm. Protein levels were studied in NPCs from saline and MPTP mice (*ex vivo*) and in NPCs from saline mice directly exposed to MPP⁺ (*in vitro*). Probing with an antibody to β -actin was used as an internal control. The values represent β -catenin and P-TyrGSK-3 β signal intensities compared with control. The effects of the *Wnt* agonist *Wnt1* (100 ng/ml) or the GSK-3 β antagonist (AR, 5 nM) are also illustrated. β -Catenin signal intensity is decreased in NPCs isolated *ex vivo* from MPTP mice, whereas active P-TyrGSK-3 β signal is increased. *Wnt1* and AR treatments increase β -catenin signal, whereas active GSK-3 β signal is decreased. Likewise, direct NPC exposure to MPP⁺ *in vitro* reduced β -catenin signal in the face of increased active GSK-3 β , whereas both *Wnt1* and AR treatments increased β -catenin signal intensity while reducing active GSK-3 β signal (**C**). Note the marked increase in Map2a⁺ (green) process length (**A**) and Map2a⁺ cell number (**B**) in NPCs exposed to the *Wnt* agonist *Wnt1* (100 ng/ml). **D**, **E**, NPCs isolated *ex vivo* from MPTP mice were transiently transfected with GSK-3 β siRNA (**D**), whereas NPCs from saline mice were transiently transfected with β -catenin siRNA (**E**) or control siRNA (see text for details). Western blotting documented that the transfection of NPCs from MPTP mice with GSK-3 β siRNA reduced GSK-3 β expression and promoted the expression of β -catenin (**D**), whereas β -catenin siRNA introduction in NPCs from saline mice decreased the expression of β -catenin without changing the expression of GSK-3 β signal (**E**). Note the increased Map2a (green) cell production (**A**) after introduction of GSK-3 β siRNA in NPCs from MPTP mice (**A**). **F**, GFP-expressing NPCs grown in differentiation medium were treated with PBS or *Wnt1* (100 ng/ml) for 72 h. Dual staining with β -catenin (red) and GFP (green) shows the distribution of the β -catenin-IF signal at the cell membrane of PBS-treated GFP, whereas *Wnt1* application sharply increased the β -catenin-IF signal at the membrane, cytoplasm, and growing GFP⁺ processes.

treatment with β -catenin siRNA decreased β -catenin protein levels without modifying GSK-3 β in NPCs from saline mice (Fig. 6E). When BrdU was added after siRNA introduction in NPCs from MPTP mice and cells were fixed after 24 h, we found that depleting GSK-3 β resulted in a significant increase in the percentage of cells expressing BrdU, in comparison with cells treated with siRNA control (Fig. 6B). Likewise, in NPCs from MPTP mice grown in differentiation medium and transfected with GSK-3 β siRNA, an increased percentage of MAP2a⁺ cells (Fig. 6A, B) compared with levels measured in NPCs pretreated with a control siRNA (data not shown) were observed. Reciprocally, β -catenin siRNA introduction in NPCs from saline mice significantly reduced the percentage of BrdU⁺ and MAP2a⁺ cells (Fig. 6B), compared with NPCs treated with a control siRNA (data not shown). These results showed that reduced β -catenin in the face of upregulated GSK-3 β may contribute to MPTP-induced SVZ impairment *ex vivo*; conversely, silencing GSK-3 β via upregulation of β -catenin can reverse the decreased neurogenesis of MPTP mice.

These results raised the possibility that MPTP/MPP⁺ might use this same signaling pathway to exert its direct action on SVZ cells. Based on the emerging evidence implicating active GSK-3 β upregulation in oxidative stress-induced neuronal cell death mechanisms (Bhat et al., 2000; Grimes and Jope, 2001; King et al., 2001; Kaytor and Orr, 2002), particularly DAergic neuronal cell death (G. Chen et al., 2004; Nair and Olanow, 2008; Duka et al., 2009; Petit-Paitel et al., 2009; L'Episcopo et al., 2011a,b), we next verified the effect of a direct exposure of NPCs to MPP⁺ *in vitro* and found decreased β -catenin signal associated with increased active GSK-3 β signal, compared with control (Fig. 6C). We next tested the ability of exogenous activation of *Wnt*/ β -catenin signaling using a selective GSK-3 β antagonist, AR-AO14418 (AR, 5 nM; Osakada et al., 2007; L'Episcopo et al., 2011a,b), or the *Wnt* ligand, *Wnt1* (100 ng/ml), in both NPCs from MPTP *ex vivo* and MPP⁺ *in vitro* (Fig. 6B, C). In the *in vitro* experimental paradigm, AR or *Wnt1* was applied 1 h before MPP⁺ exposure, and cells were fixed 24 h after MPP⁺. As observed, both AR and *Wnt1* efficiently reduced active GSK-3 β signal and reversed the decreased β -catenin signal in NPCs from MPTP mice and NPCs acutely exposed to MPP⁺ (Fig. 6C). These effects were associated with a significant reversal of a MPP⁺-induced decreased percentage of BrdU⁺- and Map2a⁺-expressing cells (Fig. 6B). In NPCs from saline mice and NPCs treated with PBS, *Wnt1* sharply increased the Map2a⁺ process length (Fig. 6A). Time course studies of *Wnt1* effect on GFP⁺ NPCs revealed an early increase in β -catenin immunofluorescence in the cytoplasm and nuclei of GFP⁺ cells within 6 h from *Wnt1* application, and by 24–48 h, *Wnt1*-induced β -catenin immunofluorescence was sharply upregulated and distributed in GFP⁺ cell membrane, cytoplasm, and particularly in the growing GFP⁺ processes (Fig. 6F), compared with the less differentiated PBS-treated GFP⁺ cells where β -catenin immunofluorescence appeared localized at the membrane (Fig. 6F).

Together, these *ex vivo* and *in vitro* results clearly established MPTP/MPP⁺-induced inhibition of *Wnt*/ β -catenin signaling activity in SVZ NPCs.

Astrocyte-derived factors and *Wnt*/ β -catenin signaling activation counteract MPTP-induced NPC impairment

The majority of currently identified neurogenic factors, including *Wnts*, are associated with a specific population of local astrocytes in neurogenic niches, implicating their specific role in instructing neurogenesis of NPCs in the niche (Alvarez-Buylla et

al., 2001; Castelo-Branco et al., 2004, 2006; Lie et al., 2005; Cahoy et al., 2008; Kazanis, 2009; Kuwabara et al., 2009; Munji et al., 2011). Given that astrocytes (1) protect NPCs from MPP⁺ exposure and efficiently reversed the decreased proliferation and neuronal differentiation (Fig. 4A–C) and (2) express *Wnt*/ β -catenin signaling components and promote neurogenesis *in vivo* and *in vitro* (Lie et al., 2005; Castelo-Branco et al., 2006; Cahoy et al., 2008; L'Episcopo et al., 2011a,b), we thus set to determine whether striatal astrocyte coculture might overcome the impaired NPC neurogenic capabilities of MPTP mice via β -catenin-mediated *Wnt* signaling activation. Hence, when astrocytes were layered on top of NPCs from MPTP mice (day 0), BrdU was next added (i.e., at 2 DIV), and cells were fixed after 24 h (i.e., at 3 DIV), a significant increase in cells expressing BrdU were observed (Fig. 7A, B). When the cells were allowed to mature for 7 DIV in differentiation medium, as above, the percentage of Map2a⁺ neurons was significantly increased compared with NPCs cultured alone (compare Fig. 6A, B with Fig. 7A, B), albeit the addition of the astrocyte monolayer to NPCs from saline mice more efficiently increased Map2a⁺ neuronal differentiation (Fig. 7A, B). These results supported our previous findings documenting the ability of midbrain astrocyte coculture to promote neurogenesis from neuroprogenitors isolated from adult midbrain and SVZ (L'Episcopo et al., 2011a). In addition, they further corroborated our present data showing that NPCs isolated from MPTP mice *ex vivo* are not intrinsically impaired. The ability of astrocytes to promote neurogenesis in MPTP-impaired NPCs could be the result of the effect of various growth/neurogenic factors recognized to promote neurogenesis. Although these unknown molecules might act either directly or indirectly and/or via paracrine/autocrine mechanisms, we used different approaches to address the involvement of β -catenin signaling modulation (L'Episcopo et al., 2011a,b). The first step in the activation of the *Wnt*/ β -catenin signaling pathway is the interaction between *Wnt* proteins and the receptor protein Fz and the coreceptor low-density LRP5/6. We thus tested the effect of *Wnt*/ β -catenin antagonism using the soluble inhibitor Dkk1, which prevents Fz/*Wnt*/LRP5/6 complex formation in response to *Wnts* (Semenov et al., 2001). This treatment was previously shown to counteract the neurogenesis-promoting effect of activated midbrain astrocytes (L'Episcopo et al., 2011a). We then found that Dkk-1 at a dose of 100 ng/ml significantly decreased astrocyte-induced increased Map2a⁺ neurons, as well as the percentage of BrdU⁺ cells (Fig. 7A, B) in NPCs from both saline and MPTP mice, suggesting that astrocyte activation of *Wnt* ligands by paracrine and/or autocrine mechanisms may contribute to the observed neurogenic effects. The implication of *Wnt* signaling was also indicated by Western blot analysis showing that β -catenin protein levels were not significantly decreased in NPC-Astro cocultures, whereas preventive application of the specific *Wnt*/ β -catenin antagonist Dkk1 efficiently reversed astrocyte-induced increased β -catenin protein (Fig. 7C), thus indicating *Wnts* as candidate activators. In support of this finding, the active GSK-3 β signal was upregulated in response to Dkk1 treatments of NPC-Astro, both in saline and MPTP groups (Fig. 7C). The potential role of β -catenin-mediated *Wnt* signaling in astrocyte–NPC cocultures was also supported by depleting β -catenin via the introduction of β -catenin siRNA in NPC cultures, as described, which resulted in sharp depletion of β -catenin in the face of increased active GSK-3 β expression (Fig. 6E). In β -catenin-depleted NPCs, the astrocyte insert failed to increase NPC proliferation and neuronal differentiation (Fig. 7A, B), supporting that

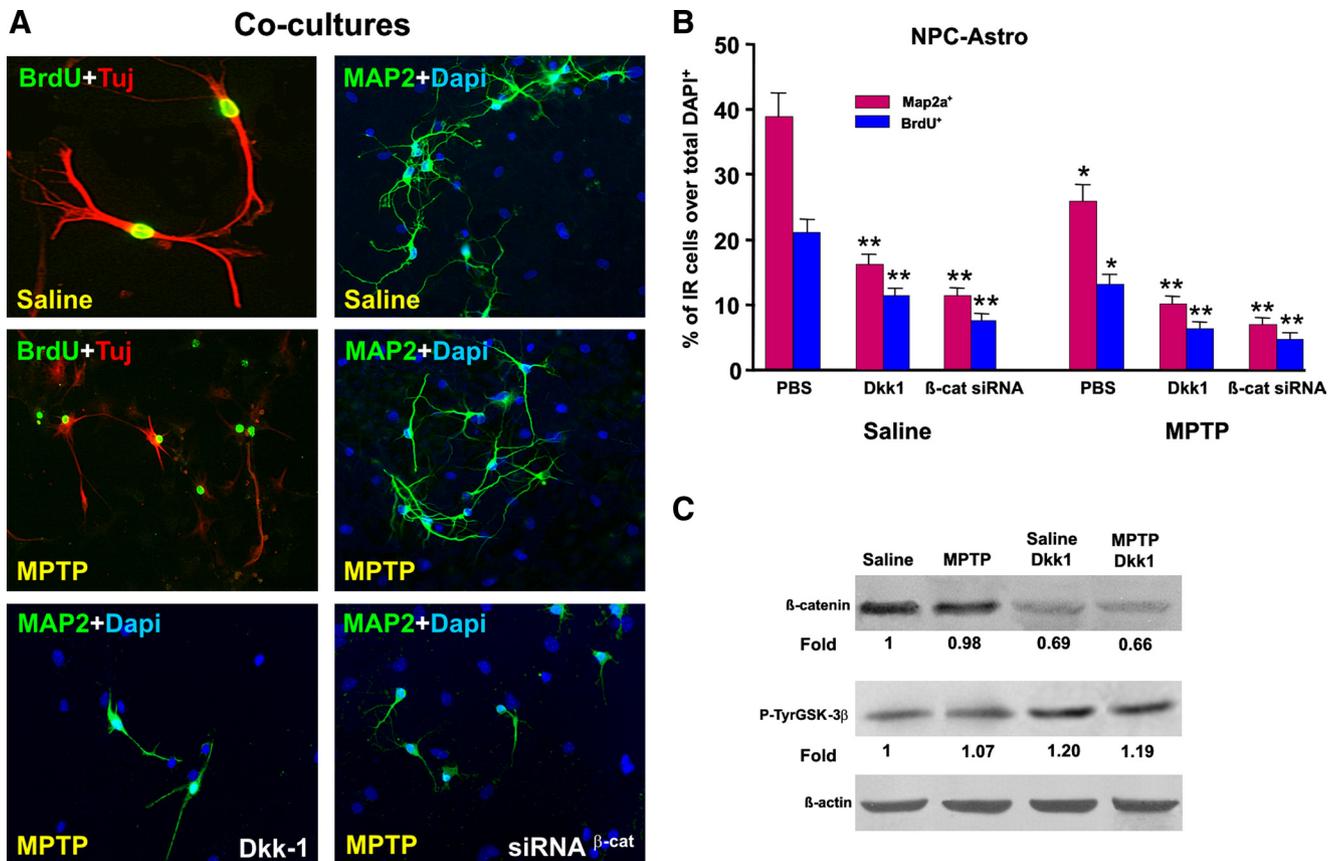


Figure 7. Astrocyte coculture and *Wnt*/ β -catenin signaling activation counteract MPTP-induced impaired neurogenesis *ex vivo*. NPCs were acutely isolated *ex vivo* 7 d after MPTP or saline injection, expanded as described, and cocultured with striatal astrocytes for proliferation and differentiation studies using BrdU, Tuj1, and Map2a. **A**, Representative images of NPCs isolated from saline or MPTP mice and cocultured with astrocytes stained with BrdU (green) and Tuj1 (red) at 3 DIV, or stained with Map2a (green) and DAPI (blue) at 7 DIV. Note the marked increase in Map2a⁺ (green) neuron production and process length by coculture with astrocytes both in saline and MPTP-NPCs. *Wnt*/ β -catenin antagonism with Dkk1 or β -catenin silencing markedly decreases Map2a⁺ (green) neuron production and process extension. **B**, Proliferation measured at 3 DIV and differentiation at 7 DIV in NPCs isolated *ex vivo* from saline and MPTP mice. NPCs of the indicated groups were exposed to different treatments, and the percentage of BrdU⁺ or Map2a⁺ cells over the DAPI-stained nuclei were measured. Astrocyte coculture significantly decreases the MPTP-induced impaired proliferation and differentiation (compare with Fig. 6, *A* and *B*). Note that *Wnt*/ β -catenin antagonism with Dkk1 (100 ng/ml) significantly reduced astrocyte-induced reversal of NPC impairment of MPTP mice (*A*, *B*). In NPCs from saline mice, transient transfection with β -catenin siRNA (see legend for Fig. 6) and coculture with astrocytes sharply decreased BrdU and Map2a⁺ cells, compared with NPCs transfected with control siRNA (see text for details). In NPCs from MPTP mice, both Dkk1 and transient transfection with β -catenin siRNA significantly reduced the neurogenic promoting effect of astrocyte coculture (*A*, *B*). **p* < 0.05 versus PBS within saline and MPTP groups, respectively; ***p* < 0.05 versus saline in NPC alone and NPC-Astro, respectively. **C**, Western blotting was performed in NPCs of saline and MPTP mice and cultured with astrocytes with the indirect coculture paradigm, as described. Probing with an antibody to β -actin was used as an internal control. The values represent β -catenin and P-TyrGSK-3 β signal intensities compared with control. Comparable β -catenin signal is observed in saline- and MPTP-NPCs cocultured with astrocytes, whereas Dkk1 treatment decreased β -catenin signal in both saline- and MPTP-NPC, while increasing the intensity of the active GSK-3 β signal.

besides others, β -catenin-mediated *Wnt* signaling may contribute to astrocyte neurogenic effects.

All together, the present findings indicated β -catenin-activated signaling as a point of convergence of MPP⁺, microglial-, and astrocyte-derived factors and prompted us to begin elucidating the contribution of *Wnt*/ β -catenin signaling *in vivo*.

Pharmacological activation of *Wnt*/ β -catenin signaling *in vivo* reverses MPTP- or Dkk1-induced impaired proliferation

We first used immunohistochemistry to localize β -catenin in the SVZ of saline- and MPTP-treated mice. In accordance with the study by Adachi et al. (2007), in the SVZ of control mice, the β -catenin-IF signal was detected at membrane, cytoplasm, or perinuclear regions of the DAPI-counterstained cell nuclei (Fig. 8*A*). Dual localization of β -catenin and BrdU revealed that a proportion of β -catenin⁺ cells coexpressed BrdU (Fig. 8*A*). In contrast, MPTP treatment sharply downregulated the β -catenin-IF signal and the proportion of β -catenin⁺ cells coexpressing BrdU (Fig. 8*A*). Because β -catenin is expressed in type C cells and since MPTP reduced the proliferation of C-cells, if the

Wnt/ β -catenin pathway is relevant for MPTP-induced neurogenic impairment of SVZ, then activation of β -catenin signaling should counteract MPTP-induced NPC impairment. We thus used pharmacological inhibition of GSK-3 β , resulting in the activation of β -catenin signaling to verify the functional importance of this pathway in cells of the SVZ. To test this hypothesis, we selected the GSK-3 β antagonist AR, used *in vitro*, and performed systemic injections or local SVZ administration by intracerebroventricular infusion, as reported (Adachi et al., 2007). Pilot experiments were performed to verify doses and timing capable to decrease active GSK-3 β in the SVZ of both intact and MPTP mice. To test the systemic effect, AR was injected intraperitoneally at a dose regimen (10 mg/kg twice per day) (L'Episcopo et al., 2011*a,b*), starting 3 h after MPTP and continuing for 3 d. For intracerebroventricular infusion, AR or the vehicle alone was infused into the left lateral ventricle of the brain. Groups of saline-injected mice received AR via intraperitoneal injections or intracerebroventricularly and served as controls. In agreement with previous findings (Adachi et al., 2007), direct GSK-3 β antagonism by AR intracerebroventricular infusion re-

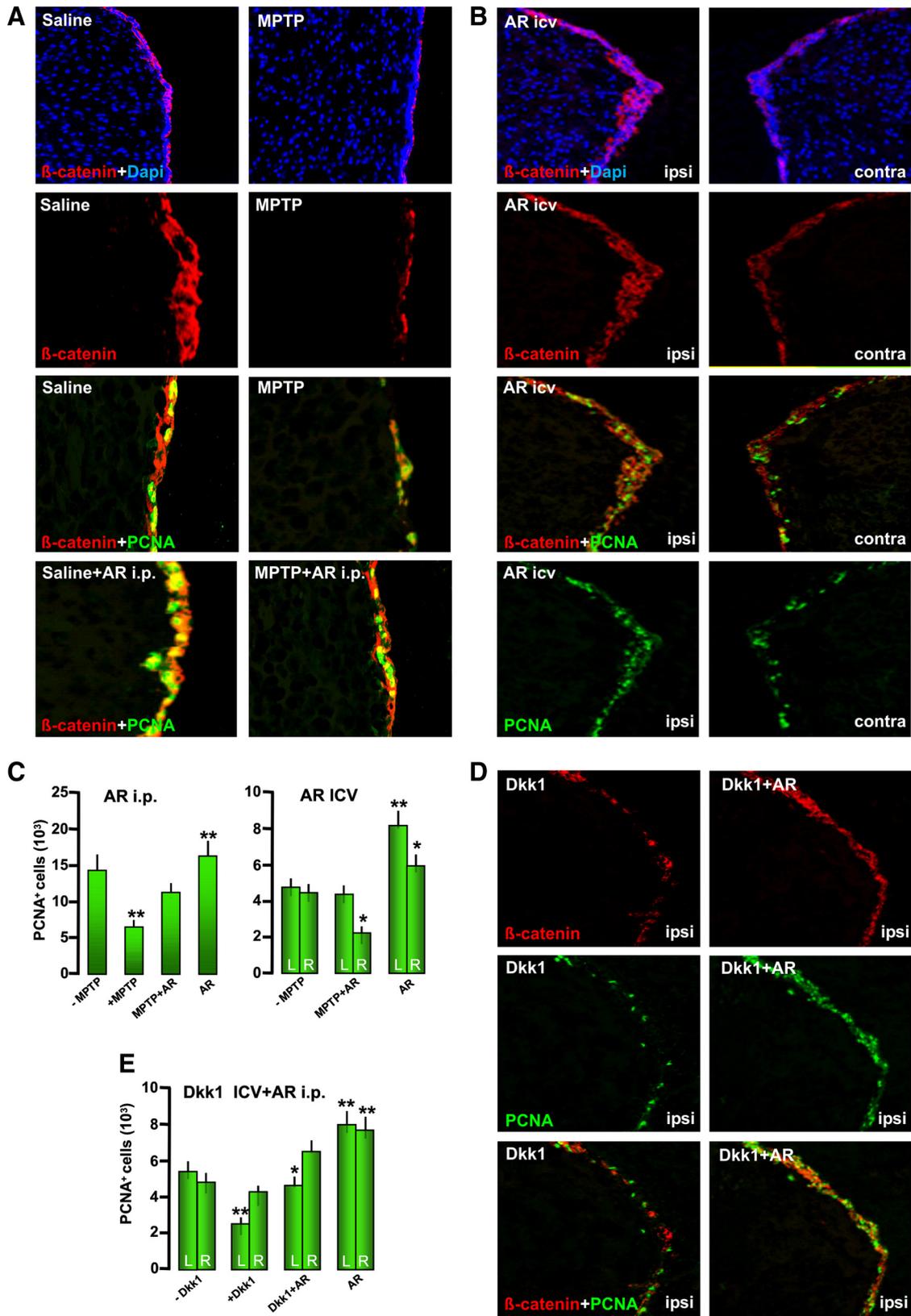


Figure 8. Activation of *Wnt*/ β -catenin signaling by GSK-3 β antagonism *in vivo* counteracts MPTP-induced impaired neurogenesis. The effect of pharmacological activation of *Wnt*/ β -catenin signaling using the specific GSK-3 β inhibitor AR was studied by either systemic or SVZ administration, starting 3 h after MPTP treatment. The effect of *Wnt*/ β -catenin signaling antagonism with Dkk1 intracerebroventricular infusion was studied in the absence or presence of AR systemic administration (for details see text). Mice were killed 3 d after MPTP, corresponding to maximal neurogenic impairment and exacerbated microglial reaction. **A**, Representative images of coronal sections of striatal SVZ stained with β -catenin (red) counterstained with the nuclear marker DAPI (blue) showing β -catenin expression in the SVZ of saline and MPTP mice. Dual localization of β -catenin and BrdU revealed that a proportion of β -catenin⁺ cells coexpressed BrdU and certain β -catenin⁺ cells exhibited nuclear colocalization. MPTP reduces the β -catenin-IF signal and β -catenin⁺ cells coexpressing BrdU. **B**, **C**, Representative images showing the effect of unilateral infusion of GSK-3 β antagonist (AR) in β -catenin expression ipsilateral (ipsi) compared with the noninfused contralateral (contra) side in saline-injected mice (**B**). (Figure legend continues.)

sulted in a sharp increase in β -catenin expression and cell proliferation in the SVZ ipsilateral to the infusion (Fig. 8B), compared with contralateral noninfused SVZ (Fig. 8B). Likewise, systemic AR treatment of saline mice resulted in a significant increase in β -catenin expression and cell proliferation in the SVZ (Fig. 8A,C). When mice exposed to MPTP received, after 3 h, AR by either intracerebroventricular infusion or systemic injection, both treatments resulted in a significant counteraction of the MPTP-induced decreased PCNA⁺ cells and β -catenin expression in the SVZ, 3 d after MPTP (Fig. 8A–C), indicating that the acute exogenous activation of β -catenin signaling during the temporal window of maximal neurogenic impairment and at the peak of microglial exacerbation (Fig. 3F) may overcome the disrupted neurogenesis observed in the SVZ of MPTP mice *in vivo*. Conversely, *Wnt/β-catenin* antagonism using the soluble inhibitor *Dkk1*, infused intracerebroventricularly unilaterally (Zhang et al., 2009), mimicked MPTP systemic injection (Fig. 8D,E), as reflected by a significant decrease in the proportion of PCNA⁺ and β -catenin⁺ cells ipsilateral compared with contralateral noninfused SVZ (Fig. 8D,E). Importantly, this inhibitory effect was efficiently counteracted by concomitant activation of the downstream transcriptional effector, β -catenin, with AR systemic injections (Fig. 8D,E).

Together, the *ex vivo*, *in vitro*, and *in vivo* findings suggested a disruption of *Wnt/β-catenin* signaling in SVZ associated with decreased proliferation of MPTP-injured SVZ. Although further time-dependent in-depth analyses will elucidate the effects *Wnt/β-catenin* manipulation on a longer-term basis, these results showed the ability of pharmacological activation of β -catenin-mediated signaling to counteract the impaired neurogenic potential of MPTP- or *Dkk1*-infused mice.

***Wnt/β-catenin* signaling disruption of MPTP mice is reversed by pharmacological modulation of inflammation: correlation with neuroprotection**

Finally, to support the plasticity of adult NPCs and test the potential to modulate this endogenous system with anti-inflammatory drug therapy *in vivo*, we used the CINOD HCT1026, endowed with a safe profile and previously shown to mitigate microglial activation by reducing PHOX and iNOS-derived RNS and to efficiently protect nigrostriatal DAergic neurons in PD rodent models (L'Episcopo et al., 2010a,b, 2011c). In accordance with our *in vitro* and *in vivo* findings, using Western blot analysis (Fig. 9A,B) and immunohistochemistry (Fig. 9C), MPTP induced downregulation of β -catenin protein levels in the SVZ. Specifically, dual labeling with IBA1 and β -catenin and with GFAP and β -catenin (Fig. 9C) supported β -catenin downregulation during the phase of maximal microglial exacerbation, i.e., at 3 dpt. On the other hand, active GSK-3 β (pTyr216 GSK-3 β) was

markedly upregulated (Fig. 9B). In contrast, HCT1026, although inactive in saline-treated mice, efficiently increased β -catenin protein back to controls after MPTP (Fig. 9A) and sharply decreased active GSK-3 β (Fig. 9B) compared with MPTP mice fed with a control diet. Double labeling with β -catenin and glial markers further revealed normalization β -catenin-IF staining in SVZ of HCT-MPTP mice and reduced GFAP⁺ and IBA1⁺ cell numbers and reactivity (Fig. 9C).

These effects of HCT1026 were associated with a significant increase in the number of BrdU⁺- and DCX⁺-expressing cells in the striatal SVZ (Fig. 9D,E,H). In addition, in dual localization experiments with GFAP and DCX and with GFAP and BrdU (Fig. 9H), GFAP⁺ astrocytes adjacent to the SVZ ensheathed the young DCX⁺ neuroblasts and displayed a reduced reactive phenotype after MPTP treatment in HCT1026, compared with mice fed with a control diet, showing disrupted astrocyte–NPC interactions at 3 dpt. Stage 4 microglial cell density was sharply decreased in the Str and SVZ of HCT1026 mice, as opposed to MPTP mice fed with a control diet (Fig. 9F–H). As observed at the ventral mid-brain level (L'Episcopo et al., 2010b), these morphological effects were associated with a substantial downregulation of microglial pro-oxidant and inflammatory mediators, such as Mac1, PHOX, and iNOS in HCT1026 as opposed to mice fed with a control diet (data not shown).

As far as striatal DAergic end points are concerned, in agreement with our previous studies (L'Episcopo et al., 2010b), by 14 d after MPTP, DAT-IF fiber density and DA uptake in striatal synaptosomes were markedly reduced by MPTP in mice fed with a control diet, compared with HCT1026-fed mice, showing a significant degree of DAergic reinnervation (Fig. 9I–K), in line with a significant protection of midbrain cell bodies (data not shown), confirming our previous report (L'Episcopo et al., 2010b).

Discussion

Impaired neurogenesis in the SVZ of PD patients and PD experimental models has previously been reported. Loss of the neurotransmitter dopamine, from midbrain DAergic cell bodies innervating type C cells in the SVZ, was causally related to decreased neurogenic potential (see Introduction). In addition, certain dopamine agonist therapies can rescue NPC proliferation in PD (van Kampen et al., 2004; Winner et al., 2009). The present work focused on glia and shows that in addition to dopamine, the PD neurotoxin MPTP/MPP⁺ either directly and/or in conjunction with astrocyte- and microglial-derived mediators may contribute to regulate SVZ plasticity in this PD mouse model. Hence, the unfavorable conditions of the SVZ niche during the early degenerative phase, resulting from decreased DAergic innervation and MPTP/MPP⁺-dependent striatal oxidative and nitrosative status, likely inhibit the survival and/or the expansion/differentiation and/or migration of endogenous NPCs, at least in part via disruption of β -catenin-mediated *Wnt* signaling in the SVZ. With time, a shift toward a less reactive microglial “harmful” phenotype likely permits the mitigation of the niche microenvironment and the return of astrocyte “beneficial” expression of growth factors and neurogenic signals, including Wnts, together with a progressive striatal reinnervation, likely contributing to NPC recovery (Fig. 10). In keeping with these findings, exogenous activation of β -catenin signaling or pharmacological mitigation of microglia overactivation upregulated β -catenin in the SVZ and successfully rescued NPC proliferation and neuroblast formation. Additional studies are clearly required to decipher whether activation of *Wnt/β-catenin* signaling, either directly or indirectly via inflammation-dependent SVZ modula-

←

(Figure legend continued.) β -Catenin (red) and cell proliferation measured with PCNA (green) are significantly increased in the ipsilateral AR-infused left SVZ (B, C). When mice exposed to MPTP received after 3 h AR by either intracerebroventricular infusion or systemic injection, both treatments resulted in a significant counteraction of the MPTP-induced decreased PCNA⁺ cells and β -catenin expression in the SVZ, 3 d after MPTP (A–C). Systemic treatment with AR alone also increased proliferation in the SVZ (A–C). D, E, *Dkk1*, infused intracerebroventricularly unilaterally, mimicked MPTP systemic injection, resulting in decreased PCNA⁺ and β -catenin⁺ cells ipsilateral compared with contralateral noninfused SVZ. The inhibitory effect of *Dkk1* was efficiently counteracted by concomitant AR systemic injections. * $p < 0.05$ between ipsilateral and contralateral, within the saline and treated groups, respectively; ** $p < 0.05$ versus MPTP/*Dkk1* within the different groups. icv, Intracerebroventricular; i.p., intraperitoneal.

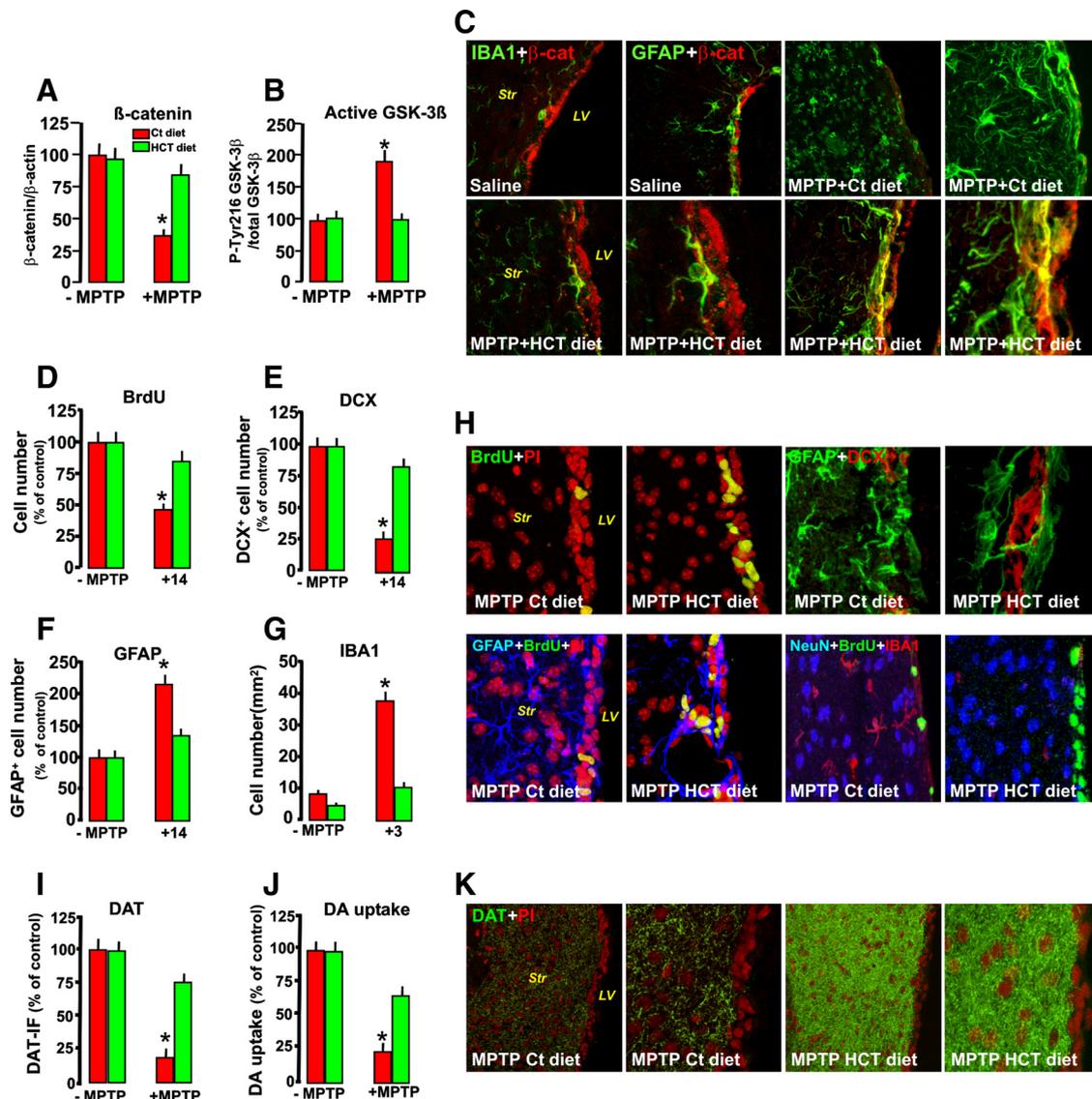


Figure 9. Pharmacological modulation of inflammation *in vivo* prevents β -catenin downregulation and reverses MPTP-induced SVZ impairment. Mice were fed with a control (Ct) or HCT1026 (HT) diet, starting 7 d before saline or MPTP injection and continuing until they were killed. **A, B**, Western blot analysis of β -catenin (**A**) and active GSK-3 β (i.e., phosphorylated at tyrosine 216 residue, pTyr216 GSK-3 β , **B**) in protein extracts from the SVZ isolated from control and treated mice. * $p < 0.05$ versus MPTP, within experimental groups. **C**, Representative confocal images of dual localization of β -catenin (red) and IBA1⁺ microglia (green) and of β -catenin (red) and GFAP (green) in mice fed with a control or HCT1026 diet, with or without MPTP treatment. Note the marked loss of the β -catenin-IF signal after MPTP in mice fed with a control diet and HCT-1026-induced reversal of the β -catenin-IF signal loss in SVZ. **D**, BrdU⁺ cell number in the SVZ of mice fed with a control or HCT1026 diet in the absence or presence of MPTP treatment. **E**, Percentage of NPCs expressing DCX in mice fed with a Ct or HCT1026 diet. **F, G**, GFAP⁺ (**F**) and stage 4 microglial (**G**) cell numbers in the Str of mice fed with a Ct or HCT1026 diet. Differences were analyzed by ANOVA, followed by the Newman–Kuls test, and were considered significant when $p < 0.05$. *Versus saline. **H**, Confocal images of BrdU⁺ (green) counterstained with the nuclear marker PI (in red), showing loss of BrdU⁺ cells in MPTP mice fed with a Ct diet and the remarkable reversal in HCT1026-fed mice. Note that dual localization of DCX⁺ neuroblasts (red) and GFAP⁺ astrocytes (green) in SVZ of MPTP mice fed with HCT1026 mice reverses DCX impairment. Dual staining of GFAP⁺ astrocytes (blue) and BrdU (green) counterstained with PI (red) in MPTP mice fed with a control diet similarly show the SVZ rescue effect of the HCT1026 diet. Triple staining with the neuronal cell markers NeuN (blue), BrdU (green), and IBA1 (red) in MPTP mice fed with HCT1026 show a significant reversal of MPTP-induced microglialosis and reduced proliferation of mice fed with a Ct diet. **I, J**, Striatal DAergic innervation as determined by measurements of FI of DAT-IR fibers in Str (**I**) and DA uptake in striatal synaptosomes (**J**). Data are expressed as percentage values of saline-injected controls. **K**, Representative images showing DAT immunofluorescence (green) in MPTP mice fed with a Ct diet, compared with HCT1026-fed mice, at 14 dpt. Note the remarkable DAergic neuroprotection of MPTP-HCT1026 compared with MPTP mice fed with a Ct diet.

tion, has potential implications for DAergic neuroprotection/self-repair (Wang et al., 2007; Maiese et al., 2008; Toledo et al., 2008; Chong et al., 2010; Inestrosa and Arenas, 2010; Kim et al., 2010; L'Episcopo et al., 2010b, 2011a,b,c; Shrueter et al., 2011).

Neuroinflammatory astrocyte- and microglial-derived signals act in concert within the niche microenvironment to shape the SVZ response to MPTP

Previous and more recent studies indicate that within the neurogenic microenvironment, factors arising from glial cells may play

important roles, especially after acute or chronic brain damage, where proliferation, and/or fate choice, migration, and maturation, may be differentially affected according to the severity of the lesion, the region affected, and the type of insult (Ekdahl et al., 2003, 2009; Monje et al., 2003; Butowski et al., 2006; Jakubs et al., 2008; Pluchino et al., 2008; Schwartz et al., 2008; Thored et al., 2009). In the MPTP-lesioned mouse model, which recapitulates many of the pathogenetic processes operative in PD (Jackson-Lewis and Przedborski, 2007), glial inflammatory mechanisms are known to contribute to nigrostriatal degenera-

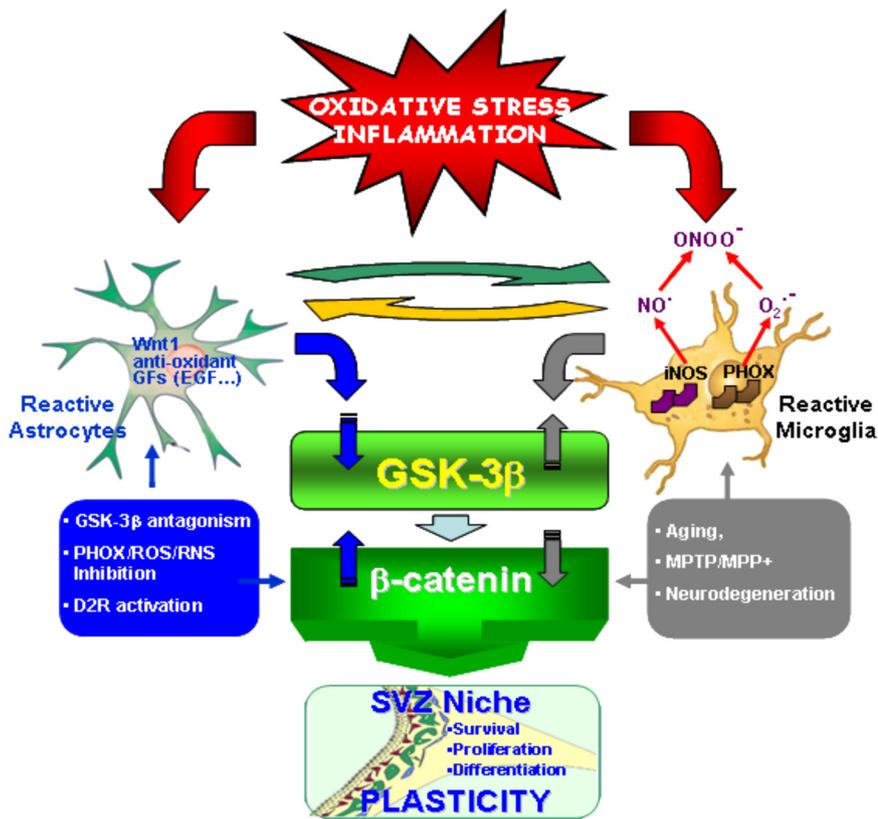


Figure 10. Cross talk between inflammatory and *Wnt/β-catenin* signaling pathways in MPTP-induced SVZ plasticity. A simplified scheme summarizing MPTP-induced neuroinflammation and SVZ plasticity via modulation of *Wnt/β-catenin* signaling is shown. During the early degeneration phase of MPTP toxicity, hyperactivated microglia contributes to the impairment of SVZ neurogenesis at different levels. By increasing oxidative and nitrosative stress and in synergy with MPTP/MPP⁺ direct toxicity, microglial-derived mediators (PHOX-derived ROS, iNOS-derived NO, and peroxynitrite) may act as molecular switch for cell signaling pathways critically involved in the physiological control of NPC homeostasis, with harmful consequences for astrocyte and NPC physiology, at least in part through GSK-3β activation, followed by phosphorylation and consequent degradation of β-catenin. In contrast, pharmacological mitigation of inflammation and oxidative stress with Apo, L-Nil, or HCT1026 upregulates β-catenin and successfully rescues NPC proliferation and neuroblast formation, a process associated with striatal DAergic neuroprotection, with further positive modulation of SVZ proliferation via D₂ receptor (D2R) activated mechanisms. The mutual role of astrocyte–microglial interactions in the plasticity of SVZ response to MPTP is exemplified by the astrocyte’s ability to overcome microglial inhibitory effects, also via cross talk with *Wnt/β-catenin* signaling.

tion and self-repair (Morale et al. 2004, 2006; Marchetti and Abbraccio, 2005; Marchetti et al., 2005a,b; L’Episcopo et al., 2010a). Here, we linked SVZ impairment with the early and marked upregulation of microglial PHOX and iNOS, generating the highly toxic peroxynitrite fingerprint 3-NT, possibly contributing to NPC mitochondrial dysfunction, increasing NPC vulnerability to cell death, and/or rendering the SVZ microenvironment unfavorable for neuroblast proliferation and/or differentiation. In particular, the studies of Estrada and coworkers (Estrada and Murillo-Carretero, 2005; Torroglosa et al., 2007) have clearly implicated NO as a key factor that regulates stem cell fate in the adult SVZ. The critical role of the microenvironment was also suggested by *ex vivo* analyses in NPC isolated from MPTP mice, showing a significant early inhibition in their growth rate during the temporal window of SVZ impairment. In analogy with studies performed in a chronic model of brain inflammation (Pluchino et al., 2008), the intrinsic mitogenic properties of NPC from MPTP mice were recovered after 12 d *in vitro* in the presence of FGF-II and EGF. In keeping with this concept, the abatement of proinflammatory and oxidative mediators herein observed with time *in vivo* likely represents a contributing factor in NPC recovery.

Astrocyte–microglia cross talk can switch the balance toward inhibition/recovery of NPC neurogenesis upon MPTP/MPP⁺ injury

Astrocytes are key players in brain homeostasis (Magistretti, 2006; Allaman et al., 2011) and in the context of adult neurogenesis (Alvarez-Buylla et al., 2001; Song et al., 2002; Jiao and Chen, 2008). In addition, microglia, when moderately active, are beneficial and support proliferation, survival, and differentiation of adult NPCs (Ekdahl et al., 2009). Importantly, the chief role of astrocyte–neuron cross talk in neurodegeneration, neuroprotection, and neurorepair is long recognized (Marchetti et al., 2005a,b; Magistretti, 2006; L’Episcopo et al., 2010a; Allaman et al., 2011), but the direct action of MPTP/MPP⁺ in NPCs, the contributory role of glial cells, and the signaling mechanisms involved, were not yet addressed. Here, by examining *in vitro* NPC vulnerability to MPP⁺, we found that doses ≤5 μM did not affect NPC survival, whereas doses at ≥10 μM, MPP⁺ induced Caspase-3 activation in a dose-dependent fashion. In addition, the ability of adult NPCs to proliferate and differentiate into Map2a⁺ neurons was dose-dependently reduced by MPP⁺, indicating the direct ability of the PD neurotoxin to impair SVZ neurogenic potential. The fact that reactive astrocytes but not microglia efficiently reversed NPC impairment is in line with findings indicating that inflammatory factors can be involved in astroglial modulation of adult neurogenesis, whereas microglia activation with LPS or with certain cytokines can impair neurogenesis (Monje et al., 2003; Ekdahl et al., 2003, 2009; Barkho et al., 2006; Butowski et al., 2006; Pluchino et al., 2008). The fact that increasing astrocyte-to-microglia ratio significantly reversed MPP⁺-induced NPC impairment also suggests that besides the degree of microglia activation, astrocyte–microglial cross talk may also play a critical role in MPTP modulation of NPC neurogenic potential.

Cross talk between inflammatory and *Wnt/β-catenin* signaling in MPTP/MPP⁺-induced neurogenic impairment: β-catenin as a common final pathway

Most of the mechanisms affecting adult neurogenesis in brain diseases are primarily unknown. Oxidative, and especially nitrosative, stress is recognized to influence fundamental cellular processes linked to aging and the development of age-related diseases, especially PD. *Wnt/β-catenin* signaling is a vital pathway regulating self-renewal and differentiation of neural stem cells, and β-catenin represents a chief transcriptional regulator (Logan and Nusse, 2004). Hence, stabilized β-catenin can enter the nucleus and associate with T-cell factor/lymphoid enhancer binding factor (TCF/LEF) transcription factors, leading to the transcription of *Wnt* target genes involved in cell survival, proliferation, and differentiation. Emerging evidence (Lie et al., 2005; Adachi et

al., 2007; Wexler et al., 2008; Kuwabara et al., 2009; Munji et al., 2011; Zhang et al., 2011) indicates a role for this pathway in adult neurogenesis. Because activation of *GSK-3β* causes *β-catenin* depletion (Aberle et al., 1997), *GSK-3β/β-catenin* disruption appeared a potential candidate mediator of MPTP–microglia-induced NPC impairment. The fact that pharmacological inhibition of microglial ROS and RNS resulted in reversal of NPC impairment associated with *β-catenin* upregulation coupled to normalization of *GSK3β* activity also suggested cross talk between inflammatory and *Wnt/β-catenin* signaling components. The observed upregulation of active *pGSK-3β* in the face of *β-catenin* and *Axin2* downregulation in SVZ of MPTP mice further implicated disruption of *Wnt/β-catenin* signaling. Because *Wnt/β-catenin* signaling controls the expression of a variety of target genes, misregulation of this signaling cascade is involved in various diseases, particularly neurodegenerative disorders associated with impaired neurogenesis such as AD and PD (Toledo et al., 2008; He and Shen, 2009; Inestrosa and Arenas, 2010; Kim et al., 2010; L'Episcopo et al., 2011a,b; Shrueter et al., 2011). Consistently, the signaling mechanisms involved in MPTP/MPP⁺-induced neuronal and NPC impairment seem to target the *Wnt/β-catenin* signaling pathway (Duka et al., 2009; Petit-Paitel et al., 2009; L'Episcopo et al., 2011a,b). Accordingly, exogenous activation of *Wnt/β-catenin* signaling by *GSK-3β* inhibition/silencing, *Wnt1* exposure, or astrocyte coculture overrode MPTP-induced impaired neurogenesis. Likewise, exogenous manipulation of *Wnt/β-catenin* signaling in primary mesencephalic neurons exerts potent neuroprotective effects against oxidative stress, 6-OHDA, and MPP⁺-induced DAergic cell death *in vitro* and *in vivo* (L'Episcopo et al., 2011a,b), and additional studies are clearly needed to address the significance and implications of *Wnt/β-catenin* signaling disruption in conditions associated with exacerbated inflammation, neurodegeneration, and impaired neurogenesis, such as PD. Because ≥100 *β-catenin* target genes (<http://www.stanford.edu/~rnuusse/pathways/targets.html>) have been identified, different upstream and downstream signaling cascades may converge in finely tuning *β-catenin* transcriptional activity (Jin et al., 2008). For example, NO can inhibit EGF-R and the phosphoinositide 3-kinase (PI3K)/AKT survival pathway (Estrada et al., 2005; Torroglosa et al., 2007), and *GSK-3β* is a downstream target of Akt (Maiese et al., 2008). In addition, peroxynitrite-induced nitration regulates the p85 subunit of PI3 kinase (Zhang et al., 2005; Trowbridge et al., 2006; Phukan et al., 2010). Moreover, a variety of molecules, including growth factors and dopamine via D₂ receptor, can signal through (PI3K)/AKT/*GSK-3β* and *Wnt* signaling activation (Spiegel et al., 2007; Beaulieu et al., 2011; Oieda et al., 2011); thus, “cross talk” among *Wnt/β-catenin* and prominent intracellular pathways may be envisaged in fine tuning SVZ neurogenic potential/plasticity observed herein (Fig. 10).

In conclusion, within the complex cell–cell interactions and signaling cascades at play in the SVZ, we herein uncovered antagonism of *Wnt/β-catenin* signaling by MPTP and microglial-derived oxidative and nitrosative mediators via upregulation of *GSK-3β* and *β-catenin* degradation as candidate contributors of neurogenic impairment in PD. Coupled to the critical involvement of astrocyte-derived Wnts and *Wnt/β-catenin* signaling in rescuing the impaired neurogenic potential observed herein, an in-depth knowledge of the mechanisms involved in the protection, activation, and migration of these endogenous neural progenitors to the site of injury would represent a strategy to promote endogenous restoration of the diseased/injured DAergic neurons in these models (Borta and Hoglinger, 2007; Hermann

and Storch, 2008; Ouredinik et al., 2009; Fernando et al., 2010; Deleidi et al., 2011).

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