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Glial-restricted precursors: Patterns of expression of opioid receptors and relationship to HIV-1 Tat and morphine susceptibility in vitro

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Abstract

Recent evidence suggests that HIV-induced pathogenesis is exacerbated by opioid abuse and that the synergistic toxicity may result from direct actions of opioids in immature glia or glial precursors. To assess whether opioids and HIV proteins are directly toxic to glial-restricted precursors (GRPs), we isolated neural stem cells from the incipient spinal cord of embryonic day 10.5 ICR mice. GRPs were characterized immunocytochemically and by RTPCR. At 1 day in vitro (DIV), GRPs failed to express µ (MOR or MOP) or κ-opioid receptors (KOR or KOP); however, at 5 DIV, most GRPs expressed MOR and KOR. The effects of morphine (500 nM) and/or Tat (100 nM) on GRP viability were assessed in GRPs at 5 DIV by examining the apoptotic effector caspase-3 and cell viability (ethidium monoazide exclusion) at 96 h following continuous exposure. Tat or morphine alone or in combination caused significant increases in GRP cell death at 96 h, but not at 24 h, following exposure. Although morphine or Tat caused increases in caspase-3 activity at 4 h, this was not accompanied with increased cleaved caspase-3 immunoreactive or ethidium monoazide-positive dying cells at 24 h. The results indicate that prolonged morphine or Tat exposure is intrinsically toxic to isolated GRPs and/or their progeny in vitro. Moreover, MOR and KOR are widely expressed by Sox2 and/or Nkx2.2-positive GRPs in vitro and the pattern of receptor expression appears to be developmentally regulated. The temporal requirement for prolonged morphine and HIV-1 Tat exposure to evoke toxicity in glia may coincide with the attainment of a particular stage of maturation and/or the development of particular apoptotic effector pathways and may be unique to spinal cord GRPs. Should similar patterns occur in vivo then we predict that immature astroglia and oligodendroglia may be preferentially vulnerable to HIV-1 infection or chronic opiate exposure.

Keywords

Human immunodeficiency virus (HIV); opiate drug abuse; glial progenitors; neural stem cells; gliogenesis; cell death

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INTRODUCTION

Injection drug abuse is a major cause of the spread of HIV/AIDS. Heroin, morphine and other opioid drugs of abuse not only promote HIV infection and the progression of AIDS (Nath et al., 2002; Steele et al., 2003), but also appear to intrinsically exacerbate the frequency and severity of HIV encephalitis (HIVE) in the CNS (Bell et al., 1998; Gurwell et al., 2001; Anthony et al., 2005). Despite the prevalence of HIV infection among drug abusers and importance of the opioid system in the pathogenesis of HIV, it is uncertain how opioid abuse augments the neuropathology of HIV (Nath et al., 2000; Nath, 2002).

Developing neural cells can express μ (MOR or MOP), δ (DOR or DOP), and/or κ (KOR or KOP) opioid receptors, which can be important in regulating cell proliferation, differentiation, and survival (Hauser and Mangoura, 1998). The particular effects seen for each receptor type are contextual and differ among cell types and at different stages of development. Immature neurons (Hauser et al., 2000; Eisch and Harburg, 2006; Narita et al., 2006a), astrocytes (Stiene-Martin and Hauser, 1990; Eriksson et al., 1990; Eriksson et al., 1991; Stiene-Martin and Hauser, 1991; Hauser et al., 1996; Stiene-Martin et al., 1998; Belcheva et al., 2005), oligodendrocytes (Knapp et al., 1998; Stiene-Martin et al., 2001), and their precursors (Persson et al., 2003a; Persson et al., 2003b; Persson et al., 2006; Kim et al., 2006) can respond uniquely to opioids. For example, MOR receptor activation can inhibit proliferation in immature astroglia, while activation of the same receptor type in immature oligodendroglia increases proliferation (Knapp et al., 1998).

We previously found that μ opioid receptor (MOR or MOP) activation exacerbated HIV-1 Tat cytotoxicity in oligodendrocyte-type 2 astrocyte (O-2A) progenitors (Khurdayan et al., 2004). The interactive toxicity was highly selective for O-2A progenitors and was not evident in more differentiated type 2 astrocytes and oligodendrocytes. To determine whether opiates and HIV-1 proteins are intrinsically toxic to glial precursors, we examined the effects of morphine, a prototypic opioid drug of abuse and preferential MOR agonist, on glial restricted precursors (GRPs) isolated from neural stem cells from embryonic mouse spinal cord.

Moreover, because morphine can have subtle actions via κ opioid receptors (KOR or KOP), and because MOR and KOR are expressed by developing neural cells (Hauser and Mangoura, 1998), and can independently affect cellular response/responsiveness to HIV-1 (Peterson et al., 1993; Chao et al., 1996; Peterson et al., 1999; Peterson et al., 2001; Chao et al., 2001), the patterns of expression of MOR and KOR were assessed in detail. GRPs were characterized by nestin, Sox2, A2B5 and Nkx2.2 immunoreactivity. Our findings indicate that opioid receptors are widely expressed by Sox2-positive and Nkx2.2-positive GRPs. In addition, both Tat and morphine alone are intrinsically lethal to GRPs, while in combination they display no additional interactive toxicity. This differs from the situation with A2B5-positive, bipotential O-2A progenitors, in which toxicity was seen only with combined Tat and morphine exposure (Khurdayan et al., 2004), suggesting a developmentally or regionally specific response of glial progenitors to HIV and/or opioid toxicity.

EXPERIMENTAL PROCEDURES

Cell culture

Pregnant ICR mice were obtained from Charles River Laboratories, Inc. (Wilmington, MA, USA). All experimental protocols conformed to local Institutional Animal Care and Use Committee (IACUC) and national (PHS) guidelines on the care and ethical use of animals. Experiments were designed and conducted to minimize the number of mice used and their discomfort. On gestational day 10.5 ± 0.5 (E10.5), mice were euthanized with ether and the embryos removed aseptically and kept on ice in cell culture media. Spinal cords were isolated from each embryo and pooled together such that the neural cells derived from the spinal cords of an entire litter (8-12 embryos) constituted a single statistical observation. As reported previously (Kalyani et al., 1997), cells were dissociated by incubating spinal cords in 0.05% Trypsin-EDTA solution (15 min) on ice, followed by incubation in cell culture medium containing 1% bovine serum albumin (BSA; Roche Diagnostics, Indianapolis, IN, USA) for 15 min and then by triturating 18-times through a fire-polished, glass Pasteur pipette. GRPs were isolated by immunopanning in dishes precoated with A2B5 antibodies prepared from an A2B5-expressing hybridoma cell line (Clone 105[HB-29]; ATCC, Manassas, VA, USA) for 1 h at room temperature at pH 7.0 (Mayer-Proschel et al., 1997; Mujtaba et al., 1999; Cao et al., 2005). Non-adherent cells were removed by rinsing with Dulbecco's phosphate buffered saline (PBS) and discarded. A2B5-adherent cells were gently removed, using a cell scraper, before plating onto poly-L-ornithine (15 µl/ml; Sigma, St. Louis, MO, USA) and laminin (20 µl/ml; GIBCO-Invitrogen Corporation, Gaithersburg, MD, USA) coated cell culture plates (24 or 48 wells). Approximately 95% of cells isolated by panning were A2B5 immunoreactive. Cultures containing less than 95% A2B5 positive cells were discarded. GRPs were cultured for 1 or 5 DIV at 35.5 °C in 5% CO₂/95% air and high humidity in Dulbecco's modified Eagle's medium/F12 (DMEM/F-12) containing N-2 supplement, recombinant murine fibroblast growth factor-2 (FGF-2, also known as basic FGF) (20 ng/ml; PeproTech Inc., Rocky Hill, NJ, USA) and platelet-derived growth factor-AA (PDGF-AA) (10 ng/ml; PeproTech Inc.) antibiotics (penicillin and streptomycin).

HIV-1 Tat protein

Recombinant Tat_{1-72} from HIV_{BRU} was prepared as described previously (Ma and Nath, 1997) with slight modifications (Gurwell et al., 2001). A deletion mutant variant of Tat_{1-72} (Tat ₃₁₋₆₁) lacking the core and basic neurotoxic domains (Nath et al., 1996) or immunoneutralized Tat_{1-72} were used as controls as previously described (Gurwell et al., 2001; Khurdayan et al., 2004).

Experimental Treatments

GRPs were treated with morphine sulfate (500 nM), Tat₁₋₇₂, immunoneutralized Tat₁₋₇₂, or mutant Tat ₃₁₋₆₁ (100 nM). Some cultures were treated with the MOR antagonist β -funaltrexamine (β -FNA; Research Biochemical International, Natick, MA, USA) (1.5 μ M).

PCR

Opioid receptor mRNA was assessed by reverse transcriptase-polymerase chain reaction (RTPCR). GRPs were grown for 1 or 5 DIV and total RNA isolated from GRPs using GenEluteTM Mammalian total RNA kit (Sigma). The first strand of cDNA was synthesized using 0.2 µg total RNA and random decamers. The PCR cycles performed as 30 sec at 94 °C, 30 sec at 55 °C and 50 sec at 68 °C for 35 cycles using platinum Taq DNA polymerase (GIBCO-Invitrogen Corporation, Gaithersburg, MD, USA). Negative controls included all the reagents essential for amplification with the exception of template DNA. 15S ribosomal control primers (5'-TTCCGCAAGTTCACCTACC-3' and 5'-CGGGCCATGCTTTACG-3') (Ambion, Austin, TX, USA) were included as an internal standard. The sense primer used for MOR started from proximal initiation sites (P_{m1} :5'-CTCAGAGAGTGGCGCTTTGGGGGATGC-3') and the antisense primer was 5'-CGCTAAGGCGTCTGCCAGAGCAAG-3'. KOR primers consisted of 5'-ATCAGCGATCTGGAGCT-3' and 5'-GCAAGGAGCATTCAATGAC-3' (Wei et al., 2000). The final PCR products were analyzed on 2% agarose gels. The identities of the amplified PCR products were confirmed by sequencing (Interdisciplinary Center for Biotechnology Research, University of Florida, USA).

Immunocytochemistry

The presence of opioid receptors on GRPs was characterized immunocytochemically. After washing two times, GRPs grown for 1 or 5 DIV were fixed using 3% Zamboni's solution for 30 min, then permeabilized with 0.1% Triton X-100 in PBS for 15 min at room temperature. Immunodetection was performed by incubating with 1% donkey serum in PBS with rabbit anti-MOR-1 antisera (1:1,000 dilution; AB1580, Chemicon International, Temecula, CA, USA), rabbit anti-δ (DOR-1 or OPRD1; AB1560 N-terminus; Chemicon) or rabbit anti-κ (KOR-1 or OPRK1; H-70; Santa Cruz Biotechnology, Santa Cruz, CA) opioid receptor antisera (both at 1:500 dilutions) overnight at 4°C; general procedures as previously described (Hauser et al., 2000). The above anti MOR-1 opioid receptor antibodies were generated against a synthetic peptide (CTNHQLENLEAETAPLP) corresponding to Cterminus of the cloned µ receptor (Chen et al., 1996). This synthetic peptide contains the amino acid sequence (NHQLENLEAETAPLP) originally used to generate anti-MOR-1 antibodies (Arvidsson et al., 1995) (with an N-terminal cysteine for conjugation), which we have extensively employed to characterize MOR expression in developing glia (Hauser et al., 1996; Stiene-Martin et al., 2001). Anti-opioid receptor primary antibodies were detected using donkey anti rabbit secondary antibodies conjugated to Cy3 (Invitrogen-Molecular probes, Eugene, OR, USA) (1:250 dilutions) for 1 h at room temperature in the dark. Controls lacking primary antibodies were used to assess antibody specificity and MOR and KOR expression was independently confirmed by RT-PCR.

Cells were incubated overnight at 4°C in goat anti-Sox2 or Nkx2.2 polyclonal antibodies (both at 1:100 dilutions; Santa Cruz Biotechnology, Santa Cruz, CA, USA). Sox2 or Nkx2.2 primary antibodies were detected using secondary antibodies conjugated to Cy2 or Alexa Fluor 488 (Molecular Probes, Inc., Eugene, OR, USA) (1:250 dilution) by incubating for 1 h at room temperature in the dark. In some cases, mouse anti-Nkx2.2 IgG2b antibodies (1:200 dilution) obtained from the Developmental Studies Hybridoma Bank (DSHB, University of

Iowa, Iowa City, USA) were used to co-localize Nkx2.2 with Sox2. Nestin was detected using anti-nestin monoclonal IgG1 (1:1 dilution; DSHB). Mouse anti-A2B5 IgM (1:200, Chemicon) was used to detect oligodendrocyte-type 2 astrocyte bipotent (O-2A) progenitors (Raff et al., 1983; Raff et al., 1984; Gard and Pfeiffer, 1990). In some instances, Hoechst 33342 (15 μ g/ml in 0.1% BSA in PBS for 15 min at room temperature) was used to counterstain cell nuclei. To assess the proportion of cells possessing opioid receptor and/or developmental marker immunoreactivity, cell cultures at 1 to 6 DIV were double-labeled for Sox2 or Nkx2.2 and MOR, DOR, or KOR immunoreactivity. Nestin and A2B5 immunoreactivity was also assessed in the cell cultures as previously reported (Khurdayan et al., 2004). For each combination of opioid receptors and developmental markers, approximately 100 cells were counted in separate cell cultures within each experiment. The proportion of immature glia that were immunoreactive for an opioid receptor type or developmental marker are given as the mean \pm SEM from 3-4 experiments.

The effects of opioids and HIV-1 Tat on individual GRPs were assessed through immunocytochemical detection of cleaved (activated) caspase-3. To detect cleaved caspase-3, cells were incubated in anti-active caspase-3 antibodies (1:4,000 dilution; R & D Systems, Inc., Minneapolis, MN, USA) overnight at 4°C followed by incubation in secondary antibodies tagged with Alexa 488 (1:250 dilution) for 1 h at room temperature in the dark. Coverslips containing GRPs were mounted on microscope slides using ProLong[®] Anti-fade mounting medium (Molecular Probes).

Cell Viability assay

Cells were incubated in calcein AM, 4 μ M; ethidium homodimer-1, 3.5 μ M for 40 min (Live/Dead assay kit, Molecular Probes). Gentle rinsing with PBS stopped the calcein AM and ethidium homodimer reactions. Cells were stored at 4°C in DMEM-F12 and counted within 24 h of the reaction. In some instances, the photoaffinity label, ethidium monoazide (EMA), was substituted for ethidium homodimer as previously described (Singh et al., 2004).

Caspase-3 activity

Caspase-3 enzyme activity was measured as previously described (Singh et al., 2004). Briefly, cell lysates were incubated at 37°C in a buffer (25 mM HEPES, pH 7.5; 10% sucrose; 0.1% CHAPS and 1 mM dithiothreitol supplemented with 50 µM Ac-DEVD-7amino-4-methylcoumarin [AMC]) in 96 well Costar plates. Fluorescent activity due to cleavage of the fluorogenic AMC moiety was measured using a Victor³ microplate reader (Perkin-Elmer, Boston, MA) at 360 nm excitation and 460 nm emission wavelengths. Caspase-3 activity was expressed as fluorescence units (cleaved AMC) per milligram of total cytosolic protein.

Statistics

For measures where opioid receptors were co-localized with Sox2 and Nkx2.2, approximately 100 cells were counted per condition in each experiment and experiments were repeated at last 3 times. For assays of apoptosis, 400 cells were sampled per treatment group in each experiment. Experiments were repeated 4 to 11 times. Data were expressed as

the mean \pm the standard error of the mean (SEM). Differences among experimental groups were assessed statistically using ANOVA followed by Duncan's multiple comparisons test (Statistica version 6.1, StatSoft, Tulsa, OK).

RESULTS

Expression of glial markers

A majority of cells expressed Sox2 and Nkx2.2 (Table 1), as well as nestin (not shown) and A2B5, immunoreactivity at 1 DIV after immunopanning and generally retained this pattern of expression at 5 DIV, although there was a tendency for the proportion of Nkx2.2+ cells to decline with progressive maturation *in vitro*. The pattern of expression of these markers suggests that a majority of GRPs retain Sox2 expression, although subsets of cells are likely to be differentiating toward astroglial and/or oligodendroglial fates (Qi et al., 2001; Wei et al., 2005). In addition, cells appeared to retain their A2B5 antigenicity, since $94.3 \pm 1.1\%$ of the cells in our GRP cultures retained A2B5 immunoreactivity at 5 DIV. To assess whether opioid receptors are expressed in glial precursors, MOR, DOR, or KOR antigenicity were co-localized with Sox2 and/or Nkx2.2 immunoreactivity in the same cell.

Opioid receptor expression

At exactly1 DIV after cells were isolated, GRPs did not express detectable levels of MOR or KOR mRNA (Fig. 1). By contrast, at 5 DIV, MOR and KOR transcripts were expressed by isolated GRPs, suggesting that the expression of these receptor types is developmentally regulated and increases with maturation (Fig. 1).

The proportion of glial precursors that expressed MOR or KOR immunoreactivity increased from 2-3 DIV compared to 5 DIV (Fig. 2). MOR, DOR, and KOR immunoreactivity was evident in some cells at 2-3 DIV, suggesting the onset of opioid receptor maturation occurs within a few days of plating. By contrast, by 5 DIV a vast majority of Sox2 or Nkx2.2 positive precursors possessed opioid receptor immunoreactivity (Table 2). In fact, approximately 90% of Sox2 positive cells possess MOR, DOR or KOR, while 69-88% of Nkx2.2 positive cells possess MOR, DOR or KOR immunoreactivity, suggesting that many cells co-expressing Sox2 and Nkx2.2 also possessed opioid receptors. Moreover, expression was not limited to a particular opioid receptor type; rather, MOR, DOR, and KOR immunoreactivity could be readily co-localized in Sox2 or Nkx2.2 positive cells.

Caspase-3 activation

At 1 DIV, exposure to morphine or Tat_{1-72} for 4 h caused an approximate 1.6-fold increase in caspase-3 activity compared to control cultures (Fig. 3A), while combined exposure did not increase caspase-3 activity beyond levels seen with either substance alone. By contrast, caspase-3 activity was unaffected by morphine and/or Tat following 24 h of continuous exposure, suggesting that the response of immature glia is dependent on the duration of exposure (Fig. 3B).

Morphine and HIV-1 Tat-induced GRP cell death

Previous studies indicated that Tat and/or morphine, or gp120-induced increases in caspase-3 activity were accompanied by increased neuronal or O-2A progenitor death; however, 24-96 h were required before actual cell losses became evident (Singh et al., 2004; Khurdayan et al., 2004). For this reason, cleaved caspase-3 product was assessed at 24 h and viability was examined at 24 and 96 h in individual GRPs. Despite increases in caspase-3 activity at 4 h, the proportion of caspase-3 immunoreactive cells were not increased above numbers seen in untreated controls at 24 h (Fig. 3B).

The effects of morphine, HIV-1 Tat₁₋₇₂, inactive mutant Tat ₃₁₋₆₁ and/or the MOR-specific antagonist β -funaltrexamine (β -FNA) on GRP cell survival were assessed by EMA exclusion. At 24 h, there were no significant effects of morphine and/or Tat on GRP viability (Fig. 4A). Unlike the results at 24 h, continuous exposure for 96 h to morphine or HIV-1 Tat₁₋₇₂ increased the number of EMA-labeled GRP cells (Fig. 4B). In combination, Tat and morphine displayed no additional cytotoxicity compared to either substance alone. Importantly, Tat toxicity was highly specific since GRP viability was unaffected by continuous exposure for 96 h to mutant Tat ₃₁₋₆₁ or immunoneutralized Tat₁₋₇₂ (Fig. 3B). Lastly, when morphine was co-administered with the MOR selective antagonist β -FNA, morphine failed to cause significant increases in cell death. Approximately 12.3 ± 0.1% of control GRPs were EMA+ at 5 DIV.

DISCUSSION

Our findings suggest that opioid receptor expression by GRPs isolated from the incipient spinal cord is developmentally regulated *in vitro* and suggest that opioid receptors are necessary, but not sufficient, in defining critical periods of vulnerability of GRPs and their progeny to opiates. At 1 DIV, GRPs were characteristically nestin, Sox2, and A2B5 immunoreactive (Liu and Rao, 2004), but failed to express MOR and KOR mRNA. However, at 1 DIV, cells exposed to morphine for 4 h showed significant increases in caspase-3 activity, suggesting that opioid receptor mRNA levels were below detection by PCR or that the onset of opioid receptor expression occurs very rapidly after 1 DIV. After 5 DIV, large proportions of glial precursors expressed MOR and KOR and this coincided with increases in the percentage of Nkx2.2 immunoreactive cells. The lack MOR and KOR expression by GRPs at 1 DIV was unexpected. Prior findings indicate that neural and glial precursors can express opioid receptors (Reznikov et al., 1999; Persson et al., 2003a; Persson et al., 2003b; Khurdayan et al., 2004; Kim et al., 2006; Hahn and Knapp, unpublished). The stress of isolating GRPs in culture and/or the loss of key factors available within the extracellular milieu in vivo may temporarily shut down opioid receptor expression. Alternatively, opioid receptor expression in developing glia may normally be regulated by multiple maturational factors and is extremely dynamic (Hauser and Mangoura, 1998).

MOR, KOR, and DOR were widely expressed by Sox2 and Nkx2.2 immunoreactive GRPs. Sox2 is a stem cell marker expressed by self-renewing, pluripotent neural cells within the ventricular zone of the developing CNS (Ellis et al., 2004; Wegner and Stolt, 2005), while Nkx2.2 is a glial specific transcription factor suggestive of an oligodendroglial fate.

Although Nkx2.2 and Olig2 are required for oligodendrocyte maturation (Zhou et al., 2001), the expression of Nkx2.2 alone is not sufficient to indicate that a cell is an incipient oligodendrocyte (Qi et al., 2001). The patterns of Sox2 and Nkx2.2 expression suggest opioid receptors are extensively expressed by progenitors very early during gliogenesis. This is interesting because only a subpopulation of their mature astroglial or oligodendroglial progeny will subsequently express MOR, DOR, or KOR (Stiene-Martin et al., 1998; Knapp et al., 1998), although a very large percentage of young, O4⁺ oligodendrocytes are MOR-positive (Knapp et al., 1998; Tryoen-Toth et al., 2000; Stiene-Martin et al., 2001). Patterns of Sox2 and Nkx2.2 expression also suggest that some opioid-receptor expressing GRPs in the present study are likely to be differentiating into more mature O-2A progenitors, since the proportion of Sox2 and Nkx2.2 cells decline from 1 to 5 DIV. Thus, while opioid receptor expression is dynamic and highly coordinated with glial maturation, the presence of a particular receptor type does not uniquely delineate a particular stage of glial maturation.

Besides temporal differences, regional disparities in the pattern of opioid receptor expression by GRPs may also exist. Regional differences in MOR, DOR, and KOR are reported for astrocytes (Stiene-Martin et al., 1998). Few studies have examined opioid receptor expression in spinal cord glia (Cheng et al., 1997; Xu et al., 2004; Narita et al., 2006b). Prior studies report that MOR and KOR expression occurs prenatally in the developing rodent spinal cord, while the onset of DOR sites occurs approximately at birth (Attali et al., 1990; Leslie et al., 1998). In contrast, other investigators find MOR and KOR mRNA in embryonic day 12.5 (E12.5) mouse spinal cord and DOR transcripts by E15.5 (Zhu et al., 1998). Moreover, DOR selective radioligand binding is present in the gray matter of 14 week human fetal spinal cord (Sales et al., 1989), while the apparent absence of DOR transcripts in human adults suggests that DOR expression maybe transient (Peckys and Landwehrmeyer, 1999). The reported variability and late onset in DOR expression during maturation prompted us to focus on MOR and KOR. Importantly, the patterns of opioid receptor expression and susceptibility to opiates and HIV-1 Tat may be unique to GRPs from the spinal cord glia and may differ from GRPs from other brain regions.

Exposure to either morphine or HIV-1 Tat alone significantly increased GRP death. GRP death was preceded by increases in caspase-3 enzyme activity at 4 h. As noted, 1 day old GRPs showed elevations in caspase-3 activity, but no increases in cell death as measured by EMA exclusion, while after 96 h cells showed marked increases in cell death with longer exposure durations. The lack of significant cell losses when morphine was combined with β -FNA, suggests that morphine is acting through MOR; however, β -FNA only partially attenuated the cytotoxicity inferring that morphine might additionally modulate cell death via a second opioid receptor type. Whether KOR or potentially DOR might be involved awaits further study, but it is feasible that morphine's complex actions in glial precursors are mediated by multiple opioid receptor types. Cytotoxicity coincided with the onset of MOR and KOR expression and progressive glial differentiation *in vitro*. The present results in GRPs were interesting because they differed markedly from the response of more mature O-2A progenitors assessed in previous studies (Khurdayan et al., 2004). We previously found that MOR activation exacerbates HIV-1 Tat cytotoxicity in O-2A progenitors, while

morphine or Tat alone were not cytotoxic and morphine actually reduced the proportion of O-2A progenitors displaying cleaved caspase-3 immunoreactivity (Khurdayan et al., 2004).

The reasons for the discrepancy between GRPs (present study) and more mature O-2A progenitors (Khurdayan et al., 2004) are uncertain, but may relate to maturational differences. Indeed, maturational differences were noticed between GRPs at 1 and 5 DIV in the present study. The protracted delay between caspase-3 activation (seen at 4 h) and onset of cell death (evident at 96 h) has been reported following viral protein \pm morphine exposure in other cell types (Singh et al., 2004; Khurdayan et al., 2004). One explanation is there may be a critical threshold for caspase-3 activation before apoptosis is initiated, and the levels and/or duration of caspase-3 activation seen at 4 h were insufficient to trigger apoptosis. Others have shown large temporal discrepancies between caspase activation and cell losses (McEwen and Springer, 2005). Additionally, caspase-3 activation may not necessarily result in cell death (Bellizzi et al., 2006). Another explanation may be that components of Tat- or morphine-mediated apoptotic signaling cascades are not fully developed in GRPs at 1 DIV, although this seems unlikely since caspase-3 activation is one of the final steps in apoptosis (Yuan et al., 2003). Acute opioid agonist exposure can cause oxidative stress in astroglia; however, astroglia may be able to adapt (perhaps temporarily) to sustained exposure (Hauser et al., 1998). A speculative notion is that caspase-3 activation alone is insufficient and additional factors may be needed before cell death can proceed; perhaps the immature glia lack this factor. We previously found that Tat activates caspase-3 and causes endonuclease G release from mitochondria in striatal neurons (Singh et al., 2004). Interestingly, caspase inhibition does not prevent Tat-induced death of striatal neurons suggesting that an alternative, caspase-3-independent mechanism is operative (Singh et al., 2004).

Another consideration is the contribution of astrocytes. Unlike the present studies where cultured cells consisted of GRPs largely uncontaminated by other cell types, our past studies assayed O-2A cells in mixed cultures with astrocytes exposed to morphine and/or Tat. Morphine and Tat cause synergistic increases in the release of cytokines, including IL-6 by astrocytes (see El-Hage et al., 2005), as well as massive increases in oxyradical production (N El-Hage, unpublished), which may contribute to the cytotoxicity seen in O-2A progenitors (Khurdayan et al., 2004). IL-6 is toxic to neural stem cells in the subgranular layer of the dentate gyrus (Monje et al., 2003), and could mediate O-2A cytotoxicity in astrocyte co-cultures.

Morphine's effects during development are cell-type-specific and highly contextual (Hauser and Mangoura, 1998). For example, epidermal growth factor (EGF) ligands negate the antiproliferative effects of morphine in neuronal precursors (Opanashuk and Hauser, 1998). Coscia and coworkers have shown that FGF-2 and EGF signals can converge with MOR signals in astrocytes (Belcheva et al., 2002; Belcheva et al., 2003; Belcheva et al., 2005), and FGF-2 and EGF signals profoundly affect the survival of neural precursors. Importantly, opiates can modulate the response of cells to apoptotic signals (Singhal et al., 1998; Polakiewicz et al., 1998; Suzuki et al., 2003; Bhat et al., 2004) and have been paradoxically described as both neurotoxic and neuroprotective (Hauser and Mangoura, 1998; Tegeder and Geisslinger, 2004; Samways and Henderson, 2006). Thus, morphine's effects on survival are

likely to be unique for GRPs and modulated by extracellular factors such as the FGF-2 supplement to the cell culture medium, which is necessary for GRP maintenance.

In summary, we find that GRPs isolated *in vitro* are intrinsically vulnerable to opiate drug and HIV-1 Tat exposure. Together with previous findings that glial precursors can be infected with HIV-1 (Lawrence et al., 2004), the present results indicate that glial precursors are important cellular targets of drug abuse and HIV-1 encephalitis. Assuming similar patterns of susceptibility occur *in vivo*, then our findings would suggest that the production and maintenance of astroglial and oligodendroglial populations is preferentially vulnerable to chronic opiate exposure and HIV-1 infection. Importantly, the vulnerability of glial progenitors appears to depend, in part, on stage-specific ontogenetic events and factors that remain to be defined.

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Fig. 1.

Detection of MOR and KOR opioid receptor transcripts by RT-PCR. MOR (A) or KOR (B) mRNA was not detected at 24 h, but was expressed at 5 DIV; GRPs (+) and blank lanes (-) lacking MOR or KOR mRNA, respectively.

Buch et al.



Fig. 2.

Co-localization of μ (MOR), δ (DOR), and κ (KOR) opioid receptors with Sox2+ and Nkx2.2+ glial precursors at 2-3 DIV (A-I) and 5 DIV (J-P). Opioid receptors are widely expressed by subpopulations of Sox2+/Nkx2.2+ glial precursors (A-P). Note, MOR, Nkx2.2 and Sox2 immunoreactivity can be localized to the same cell (arrow in M-P) and overlapping Sox2 and Nkx2.2 immunoreactivity was evident in some DOR or KOR expressing cells (data not shown); scale bar = 20 μ m (A,D,G); scale bar = 25 μ m (B-C,E-F,H-I); scale bar = 10 μ m (J-P).

Buch et al.



Fig. 3.

Effects of morphine and/or Tat₁₋₇₂ on caspase 3 activity at 4 h (A) and on the proportion of cleaved caspase-3 immunoreactive GRPs at 24 h (B). Although morphine or Tat₁₋₇₂ increased caspase-3 activity at 4 h, the proportion of cleaved caspase-3 immunoreactive GRPs was unchanged at 24 h. Data represent (n=4) independent experiments performed on cells derived from separate litters cultured for 1 DIV. Values are mean fold-change from control \pm SEM (**P* < 0.05 vs. vehicle-treated control cultures).

Buch et al.



Fig. 4.

Effects of opiates and/or Tat₁₋₇₂ (Tat), Tat ₃₁₋₆₁ [(mut)Tat], or immunoneutralized Tat [(neutr)Tat] on the viability of glial precursors after 24 h (A) or 96 h (B) of continuous exposure. Both morphine (Morph) and Tat₁₋₇₂ alone caused significant increases in the proportion of dying, ethidium monoxide-positive (EMA⁺) glial precursors (*P < 0.05 vs. vehicle-treated controls), while (neutr)Tat and (mut)Tat did not increase the percentage of EMA⁺ cells (#P < 0.05 vs. treatment with Tat₁₋₇₂). In combination, morphine and Tat₁₋₇₂ did not increase the proportion of EMA⁺ cells beyond the rate of death seen with either substance alone. In the presence of the MOR selective antagonist β-funaltrexamine (FNA), morphine did not cause significant increases in cell death. Values are the mean ± SEM fold-change in EMA⁺ cells compared to vehicle-treated controls from 4-11 experiments initiated at 1 DIV.

Table 1

Characterization of Sox2 and Nkx2.2 immunoreactivity in glial restricted precursors at 1 and 5 DIV.

Days in vitro (DIV)	Percentage (%) of cells expressing Sox2+	Percentage (%) of cells expressing Nkx2.2+
1 DIV	75.5 ± 16.5	67.3 ± 4.5
5 DIV	64.1 ± 7.5	41.9 ± 4.5

* The proportion of glial precursors that are Sox2 or Nkx2.2 immunoreactive. Mean ± SEM from 3-4 experiments and approximately 100 cells per group were sampled in each experiment.

Table 2

Characterization of opioid receptor immunoreactivity in glial restricted precursors at 5 DIV.

Developmental marker	Opioid receptor	Percentage (%) co-localized with opioid receptors $*$
Sox2	MOR	87.9 ± 4.4
	DOR	90.5 ± 1.4
	KOR	93.9 ± 3.4
Nkx2.2	MOR	88.2 ± 4.0
	DOR	77.8 ± 10.7
	KOR	69.0 ± 13.1

* The proportion of Sox2 or Nkx2.2 immunoreactive glial precursors that expressed μ (MOR), δ (DOR), or k (KOR) opioid receptors at 5 DIV. Mean \pm SEM from 3-4 experiments and approximately 100 cells per group were sampled in each experiment.