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## **SMAD3-dependent Signaling Underlies the TGF-**β**1-mediated Enhancement in Astrocytic iNOS Expression**

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## **Abstract**

We previously demonstrated that transforming growth factor-β1 (TGF-β1), while having no effect alone, enhances nitric oxide (NO) production in primary, purified mouse astrocytes induced by lipopolysaccharide (LPS) plus interferon-γ (IFN-γ) by recruiting a latent population of astrocytes to respond, thereby enhancing the total number of cells that express *Nos2*. In this investigation, we evaluated the molecular signaling pathway by which this occurs. We found that purified murine primary astrocytes express mRNA for TGFβRII as well as the TGFβRI subunit ALK5, but not ALK1. Immunofluorescence microscopy confirmed the expression of TGFβRII and ALK5 protein in astrocytes. Consistent with ALK5 signaling, Smad3 accumulated in the nucleus of astrocytes as early as 30 min after TGF-β1 (3ng/ml) treatment and persisted up to 32 hr after TGF-β1 administration. Addition of ALK5 inhibitors prevented TGF-β1-mediated Smad3 nuclear accumulation and NO production when given prior to the *Nos2* induction stimuli but not after. Finally, astrocyte cultures derived from Smad3 null mutant mice did not exhibit a TGF-β1 mediated increase in iNOS expression. Overall, this data suggests that ALK5 signaling and Smad3 nuclear accumulation is required for optimal enhancement of LPS plus IFNγ-induced NO production in astrocytes by TGF-β1.

## **Keywords**

primary astrocytes; nitric oxide; LPS; IFNγ; ALK5; TGFβRI; heterogeneous

## **Introduction**

Neuroinflammation occurs during the pathogenesis of several neurological diseases/ disorders. One of the hallmarks of neuroinflammation is reactive gliosis, which is characterized by hypertrophy/hyperplasia of astrocytes and microglia (Hirsch et al. 2005; Malhotra et al. 1990; Ridet et al. 1997; Sofroniew 2009). Both microglia and astrocytes can modulate the inflammatory state by secreting either pro-inflammatory or anti-inflammatory mediators (Chung and Benveniste 1990; Lee et al. 1993; Meeuwsen et al. 2003; Romero et al. 1996). These mediators can act an autocrine and/or paracrine fashion to trigger the up- or down-regulation of other genes including the enzyme nitric oxide synthase-2 (NOS-2 or iNOS) (Hamby et al. 2008b; Romero et al. 1996).

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*Nos2* is the gene that encodes for the inducible isoform of NOS, the enzyme responsible for the catalytic conversion of L-arginine to the free radical nitric oxide (NO). Post-mortem brains of human patients who suffered from neurological diseases/disorders including Alzheimer's disease, Parkinson's disease, multiple sclerosis, cerebral ischemia and trauma (Fernandez-Vizarra et al. 2004; Forster et al. 1999; Katsuse et al. 2003; Liu et al. 2001; Luth et al. 2002; Sasaki et al. 2000; Wong et al. 2001) demonstrate iNOS immunoreactivity. Important to this study, immunoreactivity has been observed in astrocytes in post-mortem human brains from patients with neurodegenerative diseases (Katsuse et al. 2003; Liu et al. 2001; Luth et al. 2002; Wong et al. 2001), multiple sclerosis (Cross et al. 1998; Liu et al. 2001) and from those who suffered a acute neurological insults such as traumatic brain injury (Gahm et al. 2002; Luth et al. 2001). CNS tissue taken from mouse models of Alzheimer's diseases (Heneka et al. 2005), multiple sclerosis (Pozner et al. 2005; Tran et al. 1997) and traumatic brain injury (Luth et al. 2001; Wallace and Bisland 1994) also demonstrate marked astrocytic iNOS immunoreactivity. Notably, iNOS-derived NO products in these models have been demonstrated to be deleterious (Medeiros et al. 2007; Nathan et al. 2005; Wada et al. 1998)…

Although *Nos2* is subject to regulation via several means, a potent regulator of its expression is the pleiotropic cytokine transforming growth factor- $\beta$ 1 (TGF- $\beta$ 1) (Nelson et al. 1991; Perrella et al. 1996; Perrella et al. 1994; Vodovotz and Bogdan 1994; Vodovotz et al. 1993). TGF-β1 belongs to the TGFβ superfamily. It signals by binding to TGFβRII which then heterodimerizes and transphosphorylates the TGFβ signaling receptor activin-like kinase (ALK) 5 or 1 – the expression of which is cell-type specific – initiating an intracellular serine/threonine kinase signaling cascade (de Caestecker 2004; Konig et al. 2005; Lux et al. 2006; Miyazawa et al. 2002) (Moustakas et al. 2001) (Attisano and Wrana 2002) Whereas ALK1 phosphorylates Smad1/5/8, ALK5 phosphorylates Smad2/3, each resulting in nuclear translocation of distinct signaling complexes producing disparate changes in gene expression (Miyazawa et al. 2002). Like iNOS, TGF-β1 is upregulated under neuropathological conditions (Finch et al. 1993; Flanders et al. 1998; Grammas and Ovase 2002; Huang et al. 1997; Krupinski et al. 1996; Krupinski et al. 1998; Lehrmann et al. 1998; Lehrmann et al. 1995; Logan et al. 1994; Morganti-Kossman et al. 1997; Morganti-Kossmann et al. 1999; Peress and Perillo 1995; Tanuma et al. 1997; Wang et al. 1995; Zetterberg et al. 2004). While it is traditionally thought of as having anti-inflammatory and neuroprotective functions (Buisson et al. 2003; Dhandapani and Brann 2003; Flanders et al. 1998; Kim et al. 2004), several recent studies reveal a pro-inflammatory role for TGF-β1 in the brain (Burton et al. 2002; Grammas and Ovase 2002; Lesne et al. 2003; Wyss-Coray et al. 1997a; Wyss-Coray et al. 1995; Wyss-Coray et al. 1997b). Recently, we've demonstrated that in a pure population of astrocytes – i.e., cultures devoid of microglia – that TGF- $\beta$ 1 potentiates NO production and iNOS expression induced by various pro-inflammatory stimuli (Hamby et al. 2006a; Hamby et al. 2008b). Interestingly, this enhancement occurs specifically by increasing the population of astrocytes that expressed the protein (Hamby et al. 2008a; Hamby et al. 2006a; Hamby et al. 2008b). Herein, we demonstrate that this TGFβ1-mediated enhancement in the pool of astrocytes expressing iNOS requires signaling of ALK5 and nuclear translocation of Smad3. Portions of this work have been published in abstract form (Hamby et al., 2007).

## **Materials and Methods**

#### **Primary Astrocyte Culture**

Primary astrocytes were cultured from pooled cerebral cortices of CD1 pups (Charles River) or from single pups derived from Smad3 heterozygous  $[(+/-) \times (+/-)]$  breeding pairs maintained congenic on a C57Bl/6 background (Ashcroft et al. 1999; Yang et al. 1999). Tail snips from individual pups were used to assess the genotype via PCR as described (Yang et al., 1999). An aseptic dissection of the cerebral cortices of postnatal day 1-3 mice was performed and cells mechanically and enzymatically dissociated prior to plating (Hamby et al. 2006a). Plating medium consisted of a media stock (MS) containing 10% fetal bovine serum (FBS; Hyclone), 10% iron-supplemented calf serum (CS; Hyclone), 10ng/ml epidermal growth factor (Invitrogen), 2mM L-glutamine (Mediatech), 50IU/ml penicillin and 50μg/ml streptomycin (Mediatech). MS was comprised of modified Eagle's medium (Earle's salt; Mediatech) supplemented with glucose and sodium bicarbonate to a final concentration of 25.7mM and 28.2mM, respectively. In most protocols, cells were plated at a density of 1-1.5 hemispheres/24-well plate or 1.2-1.6 hemispheres/6-well plate (both from Falcon Primaria, BD Biosciences). In experiments involving assessment of SMAD3 nuclear translocation, cells were first grown in T25 flasks, removed and then replated onto 8-well glass chamberslides (LabTek). Upon reaching confluence, astrocyte monolayers were treated with 8μM cytosine β-D-arabinofuranoside (Ara-C; Sigma-Aldrich) once for 5-6 days to eliminate the growth of any rapidly dividing cells such as microglia. Cultures were subsequently maintained in growth medium consisting of MS containing 10% CS, 2mM Lglutamine, 50IU/ml penicillin and 50μg/ml streptomycin. One day prior to experimentation, astrocyte cultures were treated with 75mM L-leucine methyl ester (LME; 1hr) to remove any residual microglia (Hamby et al. 2006a; Hamby et al. 2006b). Cells were grown, maintained, and stimulated at  $37^{\circ}$ C in a humidified atmosphere containing 6% CO<sub>2</sub>. All studies were performed on purified monolayers between 14-31 days *in vitro* (DIV).

#### **Cytokine, Endotoxin and Drug Exposure**

Cultures were treated with recombinant human TGF-β1 (R&D Systems; 3ng/ml) or its vehicle in DMEM supplemented with 5% CS, 2mM L-glutamine, 50IU/ml penicillin and 50μg/ml streptomycin. To induce *Nos2*, cells were stimulated with lipopolysaccharide (LPS; 0127:B8; 2μg/ml) plus recombinant mouse interferon-γ (IFNγ; R&D Systems; 3ng/ml). These concentrations were chosen as they provide a saturating response with respect to astrocytic NO production (Hamby et al., 2006a). In cultures derived from CD-1 mice, this combination of LPS plus IFN $\gamma$  induces iNOS expression in roughly 5-10% of mouse pure primary astrocytes, which increases to 30-35% with TGF-β1 exposure (Hamby et al. 2008a; Hamby et al. 2006a). Similar results are found when IL-1β or TNF-α are used in lieu of LPS (Hamby et al., 2008a). In experiments assessing the effect of ALK5 activation, stock solutions of the ALK5 kinase inhibitors SB431542 or SB525334 (both from Tocris) were prepared in DMSO and diluted in incubation medium to their final concentration. All experimental conditions contained identical concentrations of DMSO, which never exceeded 0.15%.

#### **NO Production**

Production of nitric oxide (NO) was assessed indirectly by measurement of nitrite, an oxidative breakdown product of NO (Green et al. 1982; Hamby et al., 2006a,b). Nitrite accumulation was measured spectrophotometrically at 550nm in a microtiter plate reader (Thermolabs) and in most cases NO production was expressed as mean nitrite accumulation  $\pm$  SEM. In the experiments involving single pup dissections, this normalization followed correction to the mean mg protein for each individual culture. This correction controlled for the variability in plating density between culture wells that unavoidably occurs with single pup dissections.

#### **TGF**β **Receptor mRNA Expression**

mRNA expression was assessed via RT-PCR as previously described (Hamby et al. 2006b). cDNA samples (1μl) were amplified for 28 (ALK1, ALK5, TGFβRII) or 23 (β-actin) cycles in a Biorad iCycler using Taq DNA polymerase, PCR reagents (Invitrogen) and primers specific for either β-actin, ALK1, ALK5, or TGFβRII in a total reaction volume of 25μL. PCR amplimer pairs for analysis were as follows: ALK1, 5′- CTATGACATGGTACCCATGACC-3′ (sense) and 5′-ACACACTTTAGGCAGAG GAAGC-3′ (antisense); ALK5, 5′-ATCTTGTACCTTCTGATCCATCG-3′ (sense) and 5′- AGGAGCAGATATGAAGAGAGCAG-3′ (antisense); TGFβRII, 5′ACTTCACTT CCGGGTCATCATC-3′ (sense) and 5′-CATGAATATGGCCGAAGTGTTC-3′ (antisense); β-actin, 5′-GTGGGCCGCTCTAGGCACCAA-3′ (sense) and 5′- CTCTTTGATGTCACGCACGATTTC-3′ (antisense). PCR products were separated in a 2% agarose gel containing ethidium bromide (0.5μg/ml) and visualized with a UV transilluminator (UVP, Kodak). Ethidium bromide fluorescence was imaged using the Kodak Electrophoresis Documentation and Analysis System 120 and images processed using Adobe Photoshop.

#### **TGF**β **Receptors, Smad3 and iNOS Protein Analyses**

Cultures were fixed with 4% paraformaldehyde (15 min) and then permeabilized using 0.25% Triton X-100 (PBS) for 7 min. Non-specific binding sites were then blocked via incubation with 10% normal goat serum (NGS) in PBS (25°C, 1 hr). TGF-β1 Receptors: A rabbit polyclonal TGFβRI (ALK5) antibody (8μg/ml, Santa Cruz Biotechnology) or TGFβRII antibody (2μg/ml, Santa Cruz Biotechnology) was added in PBS containing 5% NGS (4°C overnight) and the binding visualized with a Cy3-conjugated secondary antibody directed against rabbit IgG (7.5μg/ml; Jackson ImmunoResearch). DAPI (2μg/ml) was used to visualize nuclei. Images  $(40 \times$  magnification) were captured using a CRX digital camera (Digital Video Camera Co) mounted on an Olympus IX50 inverted microscope outfitted with epifluorescence and processed identically using Adobe Photoshop software. Smad3 and iNOS: For Smad3 single labeling, a rabbit polyclonal Smad3 antibody (2μg/ml, Santa Cruz Biotechnology) was added in PBS containing 5% NGS (4°C overnight) and its binding was visualized using either a Cy3-conjugated (7.5μg/ml; Jackson ImmunoResearch) or Alexa488-conjugated (10μg/ml; Molecular Probes) secondary antibody directed against rabbit IgG. For Smad3 and iNOS co-labeling experiments, a mouse monoclonal iNOS antibody (2.5μg/ml, BD Transduction Labs) and the aforementioned Smad3 antibody were

added simultaneously in PBS containing 5% NGS (4°C overnight) and their binding visualized using Cy3-conjugated (7.5μg/ml; Jackson ImmunoResearch) and an Alexa488 conjugated (10μg/ml; Molecular Probes) secondary antibodies directed against mouse and rabbit IgG, respectively. Control experiments demonstrated that secondary antibodies showed no non-antigen cross-reactivity. DAPI (2μg/ml) was used to visualize nuclei. Images were captured using an AxioCam MR digital camera (Zeiss) mounted on an Axiovert 200 inverted microscope (Zeiss) and processed identically using Adobe Photoshop software.

#### **Quantification of Smad3 nuclear accumulation**

Images from five microscopic fields  $(40\times$  magnification) were acquired. For each image, the total number of DAPI positive nuclei was automatically calculated using Scion NIH Image software while the number of cells exhibiting Smad3 nuclear accumulation was manually counted. The percentage of cells exhibiting Smad3 nuclear accumulation per image was calculated by dividing the number of cells exhibiting Smad3 nuclear accumulation by the total number of cells in each field (i.e., DAPI-labeled nuclei) followed by averaging the percentages from 5 fields/well. Data are expressed as the mean % cells with Smad3 nuclear accumulation + SEM. For statistical analyses, data were transformed prior to post-hoc analyses (Steel and Torrie 1980).

#### **Statistical Analyses**

All statistical analyses were performed using GraphPad Prism (Version 4.03, GraphPad Software, Inc.) as described in each figure legend. In all experiments, significance was assessed at  $p < 0.05$ .

## **Results**

Astrocytes express mRNA for TGFβRII and the TGFβRI subunit ALK5 but not ALK1 as indicated via RT-PCR analyses (Fig. 1). Immunocytochemical analyses of TGFβRII and ALK5 expression reveals that astrocytes also homogeneously express both TGFβRII and ALK5 protein (∼100% of cells) staining (Fig. 2). No qualitative change in TGFβRII and ALK5 mRNA or protein expression levels was observed following a 24hr exposure to TGFβ1 (Fig. 1, data not shown, respectively). Consistent with its expression, the TGF-β1 mediated enhancement in LPS plus IFNγ-induced NO production required ALK5 signaling as evidenced by the concentration-dependent diminution in NO production that occurred in the presence of the ALK5 kinase inhibitors SB431542 and SB525334 (Figure 3).

Signaling via ALK5 often, but not always, involves downstream Smads (Smad2/3) (Dai et al. 2003; Engel et al. 1999; Yu et al. 2002). To examine whether Smad3 was activated in primary astrocytes following TGF-β1 treatment, its nuclear accumulation was assessed immunocytochemically. Smad3 was excluded from astrocyte nuclei (i.e., was predominantly cytosolic) in vehicle-treated cultures (Figure 4A,B). However, 30min after TGF-β1 administration – the earliest time point evaluated – Smad3 accumulation was observed in approximately 50% of astrocyte nuclei (Fig. 4 A,B). Smad3 was still present in ∼20% of astrocytic nuclei 24 hr post-TGF-β1 addition (Fig. 4 A,B).

Given that nuclear Smad3 was evident at the time of iNOS induction, we next tested whether Smad3 nuclear accumulation correlated with astrocyte iNOS expression following TGF-β1 treatment. As expected, iNOS immunoreactivity resulting from exposure to LPS plus IFNγ alone occurred independently of Smad3 nuclear accumulation (Fig. 5; left panel). However, the majority of cells expressing iNOS also exhibited nuclear localization of Smad3 in cultures treated with LPS plus IFN $\gamma$  in the presence of TGF- $\beta$ 1 (Fig. 5, right panel, arrows vs. arrowheads).

To determine the time-frame in which ALK5 signaling was required to facilitate *Nos2*  induction, SB431542 (20μM) was added at various times after TGF-β1 administration and Smad3 translocation and NO production were assessed. Addition of SB431542 (20μM) rapidly and completely reversed the TGF-β1-mediated nuclear translocation of Smad3 (Fig. 6), demonstrating its effectiveness in terminating ALK5 signaling. When ALK5 – hence Smad3 activation – was inhibited with SB431542 ( $20\mu$ M) at any time prior to LPS plus IFNγ addition (arrowhead), the TGF-β1-mediated enhancement in NO production was blocked (Fig. 7). However, the TGF-β1-mediated enhancement in astrocytic NO production persisted when the ALK5 inhibitor was added after (4-20hr) LPS plus IFNγ addition (Fig. 7). Thus, ALK5 signaling is required at the time of iNOS induction.

To specifically examine whether Smad3 was required for the TGF-β1-mediated enhancement in *Nos2* induction, iNOS expression and NO production were examined and compared in astrocyte cultures derived from wild-type  $(+/+)$  and Smad3 null  $(-/-)$  mice (Fig. 8). While addition of LPS plus IFN $\gamma$  led to comparable levels of iNOS expression and NO production in astrocyte cultures derived from both wildtype and Smad3 null mice, the TGFβ1-mediated enhancement was largely ablated in cultures lacking Smad3 (Fig. 8).

## **Discussion**

The intracellular effectors and biological outcome of TGF-β1 signaling vary depending on the identity of TGFβRI, which can be either ALK5 or ALK1 (Miyazawa et al. 2002). When activated, ALK1 phosphorylates Smad1/5/8 whereas ALK5 phosphorylates Smad2/3. Herein, we find that the ability of TGF-β1 to enhance astrocytic *Nos2* expression and NO production (Hamby et. al, 2006, 2008a,b) requires Smad3, the activation of which follows ALK5 signaling. Further, results support the notion that ALK5 signaling must occur prior to and at least concomitant with the induction stimuli in order for the TGF-β1-mediated enhancement of iNOS induction to occur.

Demonstration of expression of TGFβRII and ALK5, but not ALK1, mRNA in purified primary mouse astrocyte cultures (Fig. 1) suggests that the TGF-β1-mediated effect occurs via ALK5 signaling. This is in agreement with a previous report assessing TGFβRI and TGFβRII expression in astrocytes cultures derived from rat (Konig et al. 2005). The punctate staining pattern of TGFβRII protein found herein (Fig. 2) is also similar to that described in mouse astrocytes (Sousa et al. 2009). Likewise, the staining pattern for ALK5 protein expression in astrocytes (Fig. 2) agrees with the immunocytochemical assessments of ALK5 in other cell types (Riser et al. 1999). Finally, ALK5 inhibition, afforded by the addition of the pharmacological inhibitors SB431542 or SB525334 to astrocyte cultures

(Grygielko et al. 2005; Hjelmeland et al. 2004; Laping et al. 2002), completely suppresses the TGF-β1-mediated enhancement in NO production (Fig. 3), confirming that TGF-β1 enhances iNOS expression through ALK5.

Consistent with signaling through ALK5, TGF-β1 treatment of astrocytes leads to activation and nuclear translocation of Smad3, a process blocked by inhibition of ALK5 kinase activity (Fig. 6). In astrocytes, the accumulation of Smad3 in the nucleus following TGF-β1 stimulation not only occurs fairly rapidly but is sustained (Fig. 4, 5). Although it is possible for Smad3 to remain in the nucleus for a long period of time, prolonged Smad activation in endothelial cells has previously been attributed to the continuous shuttling of Smads between the nucleus and cytoplasm in response to sustained receptor activation (Inman et al. 2002). Our results suggest that a similar mechanism may be occurring in astrocytes as astrocyte Smad3 nuclear accumulation induced by TGF-β1 rapidly dissipates following treatment with SB431542 (Fig. 6).

Notably, not all astrocytes exhibit Smad3 activation (i.e., nuclear localization) when stimulated with TGF-β1 (Fig 4). Likewise, the ability of TGF-β1 to up-regulate astrocytic iNOS expression in purified murine primary astrocyte cultures treated with LPS plus IFN $\gamma$  is not uniform (Hamby et al. 2006a). Importantly, the percentage of cells that show nuclear translocation of Smad3 (Fig. 4) and iNOS expression (Hamby et al. 2006a) following stimulation with all three mediators is very similar, suggesting that differential Smad3 signaling could account for the heterogeneous enhancement of iNOS by TGF-β1. The correlation of Smad3 with iNOS in cells treated with TGF- $\beta$ 1, LPS and IFN<sub> $\gamma$ </sub> support this notion (Fig. 5). Importantly, when Smad3 is depleted from the nucleus just prior to *Nos2*  induction, the TGF-β1-mediated enhancement in NO production is ablated (Fig. 7), suggesting that Smad3 signaling is required. In support, the ability of TGF-β1 to facilitate iNOS induction in Smad3 null-derived astrocytes was largely ablated (Fig. 8). Altogether, data indicate that Smad3 is required to be in the nucleus at the time of *Nos2* induction in order for TGF-β1 to enhance the total number of cells that express iNOS in response to LPS plus IFNγ stimulation.

The Smad3 nuclear translocation found in TGF-β1-treated astrocytes herein is consistent with biochemical evidence showing that astrocytes derived from the Smad2/3 specific Smad binding element (SBE)-luciferase reporter mice display increased luciferase activity when treated with TGF-β1 (Lin et al. 2005). The present data extend these findings by demonstrating at the single cell level that only a fraction of astrocytes exhibit Smad3 nuclear accumulation following TGF-β1 treatment. The reason for the differential activation of Smad3 in the astrocyte population (Fig. 4) despite homogeneous expression of TGFβ receptors (Fig. 2) is presently not known. However, heterogeneous activation of Smad3 in cultured cells is not unprecedented and has been seen in TGF-β1 treated cultures of rat ovarian granulosa cells (Xu et al. 2002). Interestingly, heterogeneous I-Smad, Smad7 expression has also been shown to exist in other cell types including mouse endothelial and smooth muscle cells (Banas et al. 2007) and human fibroblasts (Ishida et al. 2006). Since, Smad7 competes with Smad2/3 for phosphorylation which can, through the recruitment of additional mediators, result in the inactivation of TGFβRI and even elicit TGFβ receptor degradation (Yan et al. 2009), it is possible that differential Smad7 expression/localization

might account for inability of TGF-β1 to facilitate Smad3 nuclear accumulation and *Nos2*  induction in a subset of astrocytes. Future studies to understand the mechanism underlying heterogeneous Smad3 activation in astrocytes would be worthwhile.

The requirement for nuclear localization of Smad3 at the time of iNOS induction is consistent with a role for Smad3 in facilitating *Nos2* transcription. However, the murine iNOS promoter does not have a canonical Smad-binding element (SBE), though Smad3 has been shown to bind to non-canonical elements including AP-1 (Zhang et al. 1998), which is present on the iNOS promoter (Xie et al. 1993). Additionally, Smad3 has been demonstrated to facilitate gene transcription without binding to DNA directly but, rather, via interaction with other transcription factors and co-activators in a transcriptional activation complex (Heldin et al. 2009). Notably, Smad3 has been shown to facilitate transcription through complexes with AP-1 and NFκB, both which are known to be involved in facilitating *Nos2*  transcription (Dhandapani et al. 2003; Zhang et al. 1998; Zhu et al. 2004).

In sum, we find that TGF-β1 modulates the induction of iNOS by inflammatory mediators in an ALK5/Smad3-dependent manner. These data additionally provide one explanation that accounts for the heterogeneous enhancement in LPS plus IFNγ-induced *Nos2* expression that follows TGF-β1 treatment of astrocytes *in vitro*, namely, differential Smad3 activation. Additionally, results described herein may explain, in part, the reported heterogeneous expression of iNOS that occurs in astrocytes under neuroinflammatory conditions *in vivo*  (Luth et al. 2001; Oleszak et al. 1998). Future studies are necessary to determine the exact mechanism for both the heterogeneous activation of Smad3 and its facilitation of *Nos2*  expression.

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#### **Figure 1. Analysis of TGF**β **receptor mRNA expression in murine primary astrocytes**

Astroctye cultures were exposed to TGF-β1 (3ng/ml) or its vehicle for 24 hr prior to addition of medium alone or that containing  $LPS + IFN\gamma$  (final concentration=  $2\mu g/ml + 3ng/ml$ , respectively; 4hr). Thereafter, total RNA was isolated and TGFβRII, ALK5 and ALK1 mRNA expression was assessed via RT-PCR. Lanes 1,5, Basal; Lanes 2,6, TGF-β1; Lanes 3,7, LPS + IFNγ; Lanes 4,8, TGF-β1 + LPS +IFNγ. Positive control (+) used was mRNA isolated from brain microendothelial cultures (BMECs). Negative control (-) is H<sub>2</sub>O. β-actin mRNA expression was also assessed in astrocyte samples to demonstrate RNA integrity and approximate equal loading. Data from two separate experiments (lanes 1-4 and 5-8, respectively) from two separate dissections are shown.



**Figure 2. Immunocytochemical assessment of TGF**β **receptor expression in murine astrocytes** Naïve astrocyte cultures were fixed and immunolabeled for TGFβRII (red) or ALK5 (red) in the presence  $[ (+)$  BP; negative control] or absence  $[ (-)$  BP] of their respective blocking peptides followed by DAPI counterstaining (blue) to illustrate the number of nuclei per field. A representative photomicrograph  $(40 \times$  magnification) is shown per treatment condition. Scale bar= 40μm.



#### **Figure 3. Effect of ALK5 inhibition on astrocytic iNOS-derived NO production**

Primary astrocytes were treated with the indicated concentrations of either SB431542 (A) or SB525334 (B) for 1 hr prior to addition of medium containing vehicle or TGF- $\beta$ 1 (final = 3ng/ml). Twenty-four hr later, cultures were cultures were spiked with medium alone or that containing LPS plus IFN $\gamma$  (final = 2μg/ml and 3ng/ml, respectively). (A) Twenty-four hr or (B) 29-32hr later, cell culture supernatants were collected and NO production (mean μM nitrite accumulation + SEM) was assessed. (A),  $n = 6$  cultures from 2 separate dissections, (B) n=10-11 from 4 separate dissections. (\*) indicates a significant increase over LPS plus IFNγ alone, whereas (#) indicates a significant diminution from control as determined by one-way ANOVA followed by the Student Newman Keul's post-hoc test. Significance was assessed at p<0.05.



#### **Figure 4. Immunocytochemical assessment of Smad3 nuclear translocation following TGF-**β**1 treatment**

**(A)** Astrocyte cultures were treated with either vehicle (0 hr) or 3ng/ml TGF-β1. After the indicated treatment times, cultures were fixed and immunolabeled for Smad3 (red) followed by DAPI counterstaining (blue) to illustrate the number of nuclei per field. A representative photomicrograph (40 $\times$  magnification) is shown for each treatment condition. Scale bar = 40μm. **(B)** The percentage of cells exhibiting Smad3 nuclear accumulation per treatment condition was calculated (from a total of ∼100 cells present /field of view) and data plotted as mean % cells with Smad3 nuclear accumulation  $+$  SEM (n = 3-4 wells from 2 separate dissections). (\*) indicates a significant increase in % of cells with Smad3 nuclear accumulation compared to control (0; non-TGF-β1 treated cells) as determined by one-way ANOVA followed by the Student Newman Keul's post-hoc test following appropriate transformation of the percentage data. Significance was assessed at  $p<0.05$ .





**Figure 5. Immunocytochemical assessment of astrocytes exhibiting Smad3 nuclear accumulation and iNOS expression**

Cultures were treated with either vehicle [(-) TGF-β1] or TGF-β1 [(+) TGF-β1; 3ng/ml] for 24 hr prior to the addition of medium containing LPS plus IFN $\gamma$  (final = 2μg/ml and 3ng/ml, respectively). Eight hr later, cultures were fixed and immunolabeled for SMAD3 (green) and iNOS (red) followed by DAPI counterstaining (blue) to illustrate the number of nuclei per field. Representative photomicrographs  $(63 \times$  magnification) from at least three experiments are shown for each treatment. Scale  $bar = 20 \mu m$ .



#### **Figure 6. Time course of Smad3 nuclear translocation after SB431542 addition**

Astrocyte cultures were treated with either vehicle [(-) TGF-β1] or TGF-β1 [3ng/ml; (+) TGF-β1] for 60 min. Thereafter, cultures were either fixed **(A)** or exposed to vehicle [DMSO; (-) SB431542] or SB431542 [20μM; (+) SB431542] **(B)**. Thirty or 90 min later, cultures were fixed (**A,B**), immunolabeled for Smad3 and counterstained for DAPI. Images from the same field of view per treatment group are shown. A representative photomicrograph (63× magnification) from at least three experiments is shown for each condition. Scale bar= 40μm.



#### **Figure 7. Effect of SB431542 following TGF-**β**1 addition on NO production**

Cultures were exposed to either vehicle  $[(-)$  TGF- $\beta$ 1] or TGF- $\beta$ 1  $[(+)$  TGF- $\beta$ 1; 3ng/ml] followed by the addition of SB431542 (20μM) at the indicated times. At 24 hr (arrow), medium containing LPS plus IFN $\gamma$  (final = 2μg/ml and 3ng/ml, respectively) was added to cultures. Twenty-four hr later, culture supernatants were collected and NO production (mean μM nitrite accumulation + SEM) was assessed (n = 6 from 3 separate dissections). (\*) indicates a significant increase due to TGF-β1, whereas (#) indicates a significant within group SB431542-mediated diminution as compared to CON. Significance, assessed at p<0.05, was determined by two-way ANOVA followed by the Bonferroni's post-hoc test.



#### **Figure 8. Effect of TGF-**β**1 in Smad3 null versus wildtype-derived astrocytes**

Astrocyte cultures derived from either Smad3 null  $(-/-)$  or wildtype  $(+/-)$  littermates were treated with vehicle  $[(-)$  TGF- $\beta$ 1] or TGF- $\beta$ 1  $[(+)$  TGF- $\beta$ 1; 3ng/ml] for 24 hr prior to the addition of 2μg/ml LPS plus 3ng/ml IFNγ. **(A)** Twelve-14 hr later, culture supernatants were collected and NO production (mean  $\mu$ M nitrite accumulation/mg protein + SEM) was assessed ( $n = 15-18$  from 3-4 separate dissections). (\*) indicates a significant within group increase, whereas (#) indicates a significant between group diminution as determined by two-way ANOVA followed by the Bonferroni's post-hoc test. **(B)** Thereafter, cultures were fixed and immunolabeled for iNOS followed by DAPI counterstaining. The percentage of cells expressing iNOS per culture well was calculated and data plotted as mean % iNOS positive cells  $+$  SEM (n = 6 from 3 separate dissections). Following transformation of the percentage data, significance was assessed as in (A).