Signalling through the type 1 insulin-like growth factor receptor (IGF1R) interacts with canonical Wnt signalling to promote neural proliferation in developing brain

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ABSTRACT

Signalling through the IGF1R [type 1 IGF (insulin-like growth factor) receptor] and canonical Wnt signalling are two signalling pathways that play critical roles in regulating neural cell generation and growth. To determine whether the signalling through the IGF1R can interact with the canonical Wnt signalling pathway in neural cells in vivo, we studied mutant mice with altered IGF signalling. We found that in mice with blunted IGF1R expression specifically in nestin-expressing neural cells (IGF1RNestin−/−KO mice) the abundance of neural β-catenin was significantly reduced. Blunting IGF1R expression also markedly decreased: (i) the activity of a LacZ (β-galactosidase) reporter transgene that responds to Wnt nuclear signalling (LacZTCF reporter transgene) and (ii) the number of proliferating neural precursors. In contrast, overexpressing IGF-I (insulin-like growth factor I) in brain markedly increased the activity of the LacZTCF reporter transgene. Consistently, IGF-I treatment also markedly increased the activity of the LacZTCF reporter transgene in embryonic neuron cultures that are derived from LacZTCF Tg (transgenic) mice. Importantly, increasing the abundance of β-catenin in IGF1RNestin−/−KO embryonic brains by suppressing the activity of GSK3β (glycogen synthase kinase-3β) significantly alleviated the phenotypic changes induced by IGF1R deficiency. These phenotypic changes includes: (i) retarded brain growth, (ii) reduced precursor proliferation and (iii) decreased neuronal number. Our current data, consistent with our previous study of cultured oligodendrocytes, strongly support the concept that IGF signalling interacts with canonical Wnt signalling in the developing brain to promote neural proliferation. The interaction of IGF and canonical Wnt signalling plays an important role in normal brain development by promoting neural precursor proliferation.

Key words: β-catenin, central nervous system (CNS), insulin-like growth factor (IGF), type 1 IGF receptor (IGF1R), signalling, Wnt.

INTRODUCTION

The growth and development of both neural stem cells and lineage-restricted neural progenitors in the CNS (central nervous system) is controlled and specified by multiple neural signals and their interactions. During the past two decades, accumulating experimental data have convincingly established an essential role for IGF (insulin-like growth factor) signalling in the normal development and growth of neural cells. IGF-I and IGF-II, two ligands of the IGF system that act predominately, if not exclusively, by interacting with the IGF1R (type 1 IGF receptor) (Liu et al., 1993; Efstratiadis,

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Abbreviations: CA, cornu ammonis; CNS, central nervous system; DAPI, 4',6-diamidino-2-phenylindole; DG, dentate gyrus; DMEM, Dulbecco’s modified Eagle’s medium; E, embryonic day; Erk, extracellular-signal-regulated kinase; GO, gestational day; GSK3β, glycogen synthase kinase-3β; HIP, hippocampus; IGF, insulin-like growth factor; IGF1R, type 1 IGF receptor; KO, knockout; LacZ, β-galactosidase; P, postnatal day; pAkt, phosphorylated Akt; PCL, pyramidal cell layer; PFA, paraformaldehyde; PI3K, phosphoinositide 3-kinase; pH3Ser10, phosphorylated histone H3 at Ser10; qRT-PCR, quantitative real-time-PCR; TCF, T-cell factor; Tg, transgenic; VZ, ventricular zone.

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studies of oligodendroglial cultures suggest that during development remain largely unclear. Recently, our pathways, as well as the functions of their interactions, extensively, the interactions of the two important signalling (Junghans et al., 2005).

2007) and an increase in apoptosis of neural precursors in neural proliferation (Machon et al., 2003; Adachi et al., 2003; Schuller and Rowitch, 2007), resulting from a decrease leads to a marked retardation of brain growth (Machon et al., 2003; Ye et al., 2002). In contrast, blunting the expression of IGF-I (Beck et al., 1995; Ye et al., 2002) or the IGF1R in specific neural cells (Zeger et al., 2007; Kappeler et al., 2008; Liu et al., 2009b; Lehtinen et al., 2011), or reducing the availability of IGF-I in brain (Ye et al., 1995; Ni et al., 1997) significantly retards brain growth by decreasing the proliferation of neural precursors and the survival of neurons and oligodendrocytes. These data convincingly show that signalling through the IGF1R is essential to the normal development of the CNS. The precise intracellular signalling pathway(s) that mediates each of these IGF actions and IGF interactions with other neural signalling, however, remain to be defined.

Canonical Wnt signalling, mediated through β-catenin (Aberle et al., 1997; Salic et al., 2000; Amit et al., 2002; Liu et al., 2002; Schwarz-Romond et al., 2002), also plays a critical role in neural development and growth. For example, Tg mice overexpressing β-catenin exhibit brain overgrowth with a greater proportion of neural precursors re-entering the cell cycle (Chenn and Walsh, 2002, 2003). Similarly, when β-catenin expression is increased in the SVZ (subventricular zone) of adult mice, the proliferation of neural precursors is increased (Adachi et al., 2007). In contrast, blunting the expression of β-catenin or Wnt3, an extracellular Wnt ligand, leads to a marked retardation of brain growth (Machon et al., 2003; Schuller and Rowitch, 2007), resulting from a decrease in neural proliferation (Machon et al., 2003; Adachi et al., 2007) and an increase in apoptosis of neural precursors (Junghans et al., 2005).

While the role of IGF and canonical Wnt signalling pathways in neurogenesis has been independently studied extensively, the interactions of the two important signalling pathways, as well as the functions of their interactions, during development remain largely unclear. Recently, our studies of oligodendroglial cultures suggest that β-catenin acts as a signalling molecule downstream of the IGF-I-PI3K (phosphoinositide 3-kinase)-Akt pathway, partially mediating IGF proliferative and survival signalling (Ye et al., 2010). To further determine whether this new IGF signalling pathway also exists in neural cells during normal in vivo development, and if so, what role it plays in neurogenesis, we studied mutant mice with ablated IGF1R specifically in nestin-expressing neural cells (IGF1RNestin-ko mice). As almost all IGF1RNestin-ko mice die within 48 h after birth (Liu et al., 2009b), our study focused on prenatal development. In this report, we provide evidence that IGF signalling interacts with canonical Wnt signalling, at least at the level of β-catenin, to promote neural cell proliferation during normal brain development. Our data also strongly suggest that a portion of IGF-I and Wnt signalling converges by regulating the phosphorylation and activity of GSK3β (glycogen synthase kinase-3β) and the abundance of β-catenin.

**MATERIALS AND METHODS**

**Mutant mice**

Generation and characterization of IGF1RNestin-ko mice have been described previously (Liu et al., 2009b). Mutant mice carrying a LacZ (β-galactosidase) reporter transgene that is under the control of a promoter containing multiple copies of consensus TCF (T-cell factor)-binding motifs (LacZTg Tg mice, Maretto et al., 2003) were obtained from the Jackson Laboratory.

To suppress GSK3β activity in embryonic brains, pregnant dams bearing E11.5 IGF1RNestin-ko embryos were treated with LiCl at a concentration of 0.24% for 5–7 days as a supplement in drinking water (Wang et al., 2001; Willing et al., 2002). This concentration has been shown to be well below the level of toxicity for fetuses (S zabo, 1970). Sucrose (5%) was added to improve taste, and NaCl (0.9%) was added to maintain sodium balance and correct diuresis that may be caused by lithium (Leeman et al., 2007; Grunfeld and Rossier, 2009). IGF1RNestin-ko and control mice, given the same amount of sucrose and NaCl, served as treatment controls. No obvious abnormalities in gross brain morphology were observed in mice receiving LiCl. Some pregnant dams bearing E11.5 IGF1RNestin-ko embryos were treated with the GSK3β inhibitor AR-A014418 (EMD Millipore). AR-A014418 was administered by oral gavage at 30 μmol/kg, twice daily for 6 days, as previously reported (Noble et al., 2005). AR-A014418 was reconstituted in a modified solution (Noble et al., 2005) that contains 40% polyethylene glycol 400 and PBS, and dimethylamine was omitted from the original recipe because of its high toxicity in vivo. All procedures used were consistent with the guidelines of National Institutes of Health and approved by the institutional review committees of the University of North Carolina at Chapel Hill.

**LacZ histochemistry**

Brains were fresh-frozen in liquid N2, and coronally sectioned (18–20 μm in thickness) on a cryostat. Serial sections, comprising every sixth section, were obtained, and subjected to LacZ histochemical staining, as we previously described (Zeger et al., 2007). Briefly, sections were fixed with cold 2% PFA (paraformaldehyde) and 0.02% glutaraldehyde in PBS for 10 min on ice. After extensive washes with cold PBS, sections were incubated with a reaction buffer (Mercer et al., 1991) that contained 1 mg/ml X-Gal ([5-bromo-4-chloroindol-3-yl β-d-galactopyranoside]) and 2 mM MgCl2. For LacZ and

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immunohistochemical double staining, LacZ-stained sections were immunostained with Ki67 antibody (1:500, Vector Laboratories).

**Morphometric analysis of HIP (hippocampus)**

Brains from embryos were fixed by immersion in 4% PFA, paraffin-embedded, and sectioned in the coronal plane. Two to three sets of serial sections (8 μm in thickness), comprising every seventh section, were obtained. One set of the series sections was stained with Cresyl Violet. To determine the volume of HIP and CA (cornu ammonis) PCL (pyramidal cell layer), the area of the brain regions on each sections [corresponding to plates between number GD18 Cor. 12 and 14 in Schambra et al. (1992)] was measured under a microscope, assisted with Stereo Investigator software (Microbrightfield). The volume \( V \) was then estimated using the equation \( V = Σ A \times T \times I \), where \( Σ A \) is the sum of area measured on each section, \( T \) is the section thickness and \( I \) is the section intervals.

The numerical density of CA pyramidal neurons (neurons per mm\(^2\)) was determined using previously published methods (Dentremont et al., 1999). Briefly, stained sections corresponding to plate GD18 Cor. 14 in Schambra et al. (1992) from each brain was selected, and cell nuclei within delineated areas of interest were counted, assisted with Stereo Investigator software. The total number of CA neurons was calculated from the estimate of tissue volume and the numerical density of neurons.

**Culture of embryonic frontal cortex**

Frontal cortex from E14.5 LacZ\(^{Cr}\) Tg embryos were minced into small pieces and digested with papain (1 unit/ml) at 37°C for 3 min. Single cells were obtained by titration and passing through a screen, were seeded on to poly-L-lysine-coated coverslips individually placed in six-well plates, and cultured with a growth medium [DMEM/F12 medium supplemented with insulin-containing F27 medium (Invitrogen)]. After 2 days in culture, the growth medium was replaced by a low-insulin medium [DMEM/F12 medium supplemented with an alternative F27 medium containing no insulin (Invitrogen) and with 5 ng/ml of insulin]. At the concentration of 5 ng/ml, insulin does not activate IGF1R. Human IGF-I (100 ng/ml, Genentech), mouse Wnt3a (10 ng/ml, R&D Systems), or a combination of both IGF-I (100 ng/ml) and Wnt3a (10 ng/ml) were then added to cultures for 24 h.

To quantify LacZ-positive (+) cells, cells on each of the coverslips were immunostained with an antibody against LacZ (1:400, Abcam). Antibody–antigen complexes were detected by an Alexa Fluor\(^{®}\) 488-conjugated secondary antibody (Invitrogen), and cell nuclei were counterstained with DAPI. Staining without primary antibodies served as negative controls, and no significant non-specific background staining was observed. Fluorescent images were digitally captured and analysed using a fluorescent microscope and a Spot Jr. digital camera. To quantify the number of pH3\(^{Ser10}\)+ proliferating neural precursors in VZ (ventricular zone), the areas of VZ in immunostained sections, corresponding to plate GD18 Cor. 10 in Schambra et al. (1992) from each brain, were digitally captured. On captured images, pH3\(^{Ser10}\)-cells in VZ, each with a clearly visible nucleus, were scored, and their number within a defined linear length (100 μm) was calculated.

**Immunohistochemistry**

Brains were fresh-frozen and serially sectioned as described above. After being fixed with 4% PFA and washing with PBS, sections were subjected to immunostaining. Primary antibodies against the following proteins were used: β-catenin (1:500, BD Biosciences), LacZ (1:400, Abcam), activated caspase 3 (1:300, Cell Signaling), pH3\(^{Ser10}\) (histone H3 phosphorylated at Ser\(^10\); 1:2000, Cell Signaling), or Ki67 (1:500, Vector Laboratories). Antibody–antigen complexes were detected by an Alexa Fluor\(^{®}\) 488-conjugated or Alexa Fluor\(^{®}\) 594-conjugated secondary antibody (Invitrogen). Cell nuclei were counterstained with DAPI. Staining without primary antibodies served as negative controls, and no significant non-specific background staining was observed. Fluorescent images were digitally captured and analysed using a fluorescent microscope and a Spot Jr. digital camera.

To quantify the number of pH3\(^{Ser10}\)+ proliferating neural precursors in VZ (ventricular zone), the areas of VZ in immunostained sections, corresponding to plate GD18 Cor. 10 in Schambra et al. (1992) from each brain, were digitally captured. On captured images, pH3\(^{Ser10}\)+-cells in VZ, each with a clearly visible nucleus, were scored, and their number within a defined linear length (100 μm) was calculated.

**Protein Western immunoblot analysis**

Tissues were pulverized on solid CO\(_2\), and protein was extracted as previously described (Richards et al., 2001). Aliquots of protein (30–40 μg) were separated on polyacrylamide gels and transferred on to PVDF membranes (Amersham). Proteins of interest were detected using specific antibodies and visualized using ECL (enhanced chemiluminescence; Amersham). Primary antibodies against the following proteins were used: Akt (1:2000); pAkt\(^{Thr308}\) (1:2000); pAkt\(^{Ser473}\) (1:2000); pGSK3\(^{b}\) (1:2000); pGSK3\(^{b}\) (1:2000); pGSK3\(^{b}\) (1:2000); cell nuclei were DAPI stained. Nuclei were counterstained with DAPI. Staining without primary antibodies served as negative controls, and no significant non-specific background staining was observed. Fluorescent images were digitally captured and analysed using a fluorescent microscope and a Spot Jr. digital camera. To quantify the number of pH3\(^{Ser10}\)+ proliferating neural precursors in VZ (ventricular zone), the areas of VZ in immunostained sections, corresponding to plate GD18 Cor. 10 in Schambra et al. (1992) from each brain, were digitally captured. On captured images, pH3\(^{Ser10}\)+-cells in VZ, each with a clearly visible nucleus, were scored, and their number within a defined linear length (100 μm) was calculated.

**qRT-PCR (quantitative real-time PCR)**

RNA isolation and cDNA reverse-transcription qRT-PCR were performed as previously described (Liu et al., 2009a; Ye et al., 2010). The resultant mRNA-derived cDNA was quantified by
Statistics and data analysis

Either the Student’s t test or one-way ANOVA followed by the Newman–Keuls–Student test, assisted with the software SigmaStat for Windows (SPSS Inc.), was used to determine the statistical significance of differences between and among means.

RESULTS

Blunting IGF–IGF1R signalling significantly reduced the expression of β-catenin in embryonic brains, a finding that is consistent with our previous study of cultured oligodendrocytes (Ye et al., 2010). When judged by Western immunoblot analysis, IGF1RNestin−/− KO mice at E (embryonic day) 17.5 exhibited 60–65% reduction in the abundance of brain β-catenin protein and mRNA (Figures 1A and 1B). In parallel, the abundance of pAkt and pGSK3β, a signalling molecule that is known to regulate β-catenin stability, was also reduced by 40–50% in the brain of IGF1RNestin−/− KO mice (Figure 1C). In agreement with Western immunoblot analyses, immunofluorescence staining showed that in E17.5 IGF1RNestin−/− KO mice both nuclear β-catenin and membrane-bound β-catenin were markedly reduced in multiple brain regions, including HIP, VZ and SVZ (Figure 2), brain regions where neural precursors actively proliferate. Similarly, the abundance of β-catenin protein was also significantly decreased in the DG (dentate gyrus) and cerebellum of mice with an IGF-I null mutation [IGF-I KO (knockout) mice (Liu et al., 1993; Ye et al., 2002), see Supplementary Figure S1 at http://www.asnneuro.org/an/004/an004e092add.htm].

To directly determine whether IGF signalling can influence the nuclear Wnt−/β-catenin signalling, we bred LacZTCF reporter Tg mice with IGF1RNestin−/− KO mice. LacZTCF reporter mice carry a LacZ transgene that is under the control of a promoter containing multiple copies of TCF-binding motifs and respond to β-catenin/TCF signalling (Maretto et al., 2003). Thus, this mouse model allows us to readily assess canonical Wnt−/β-catenin signalling in vivo by monitoring the expression pattern of LacZ. At E16.5–E18.5, LacZ+ cells were readily detected in many brain regions, including VZ, SVZ and DG, a finding that is consistent with previous reports (Maretto et al., 2003; Lie et al., 2005). In DG, LacZ expression was predominantly observed in granule cells and subgranule layer cells. When compared with control mice (i.e. LacZTCF Tg mice without altered IGF signalling), blunting IGF1R expression in LacZTCF/IGF1RNestin−/− KO double mutant mice significantly decreased LacZ staining. Figure 3(A) shows representative images of LacZ staining in the brain of an E18.5 LacZTCF/IGF1RNestin−/− KO double mutant mouse and that of a LacZTCF Tg control mouse. Immunostaining with a LacZ antibody exhibited a pattern that is identical with LacZ staining (Figure 3B).

In contrast, when LacZTCF transgene expression was assessed in the brain of IGF-I overexpressing Tg mice [IGF1R+/+/ Tg mice, (Ye et al., 1995)] at P (postnatal day) 12, stronger LacZ staining and more LacZ+ cells were observed (Figure 3C). Consistent with these in vivo results, IGF-I treatment of cultured embryonic cells derived from E14.5 LacZTCF frontal cortex also showed significant increases in the intensity of LacZ staining and in the number of LacZ+ cells (Figure 4). When compared with controls, the number of LacZ+ cells in IGF-I treated cultures was almost doubled (Figure 4B). Similarly, IGF-I treatment also increased the activity of a TOP–dGFP (TCF optimal promoter–destabilized green fluorescent protein) reporter transgene in IGF1RNestin−/− KO and control IGF1RNestin−/− KO mice [IGF1RNestin−/− KO mice (KO) and their littermate control (Cont)]. Values represent means ± S.E.M. from 4–5 samples. *P<0.05, compared with controls.

![Figure 1](image-url)  
**Figure 1**  
β-Catenin expression in IGF1RNestin−/− KO mice during embryonic development  
(A) Representative Western immunoblot analysis of brain β-catenin, as well as pAkt and pGSK3β, in an E17.5 IGF1RNestin−/− KO mouse (KO) and its littermate control (C). (B) Quantification of the abundance of brain β-catenin protein and its mRNA in E17.5 IGF1RNestin−/− KO mice (KO) and their littermate controls (Cont). (C) Quantification of the abundance of brain pAkt and pGSK3β protein in E17.5 IGF1RNestin−/− KO mice (KO) and their littermate controls (Cont). Values represent means ± S.E.M. from 4–5 samples. *P<0.05, compared with controls.
Figure 2 Representative microphotographs of β-catenin immunostaining in the VZ/SVZ of an IGF1R<sup>Nestin-KO</sup> mouse (KO) and a control mouse (Contl) at E17.5. Arrows indicate membrane-bound β-catenin, and arrowheads show nuclear β-catenin. LV, lateral ventricle.

Figure 3 IGF signalling alteration of LacZ TCF reporter transgene expression in embryonic brain
(A) Representative microphotographs of LacZ staining in brain sections from an E18.5 LacZ<sup>TCF</sup>/IGF1R<sup>Nestin-KO</sup> double mutant mouse (IGF1R<sup>Nestin-KO</sup>, right panels) or a LacZ<sup>TCF</sup> control Tg mouse (Control, left panels). Arrows indicate DG region. C 1, cortex layer 1; MH, medial habenular nucleus; LV, lateral ventricle; RS, retrosplenial cortex; V3, third ventricle. (B) Representative microphotographs of LacZ immunostaining in the VZ and DG of an IGF1R<sup>Nestin-KO</sup> mouse (KO) and a control mouse (Contl) at age of E17.5. (C) Representative microphotographs of LacZ histochemical staining in the HIP of LacZ<sup>TCF</sup>/IGF-I double Tg mice. Brains of a LacZ<sup>TCF</sup>/IGF-I double Tg mouse, a LacZ<sup>TCF</sup>/IGF-I double Tg mouse and its wild-type control were obtained at P12, a time when IGF-I<sup>MMT</sup>-transgene begins to be highly expressed in HIP and other brain regions (Ye et al., 1995).
green fluorescent protein) reporter gene, which also contains multiple copies of TCF-binding motifs linked to its promoter, in transfected B104 neuroblastoma cells (results not shown). Treatment with Wnt3a appeared to further increase the number of LacZ+ cells, as compared with IGF-I treatment, but the increase did not meet statistical significance (Figure 4B).

In both VZ/SVZ and DG regions, LacZ was expressed in a pattern similar to that of Ki67 (Figure 5A), and often co-located in Ki67+ proliferating neural precursors (Figure 5B). These data are consistent with the previous results that Wnt–

β-catenin signalling is active in proliferating neural precursors (Lie et al., 2005) and is required for normal cell proliferation during postnatal life (Solberg et al., 2008). IGF1R deficiency significantly decreased the number of LacZ+ cells and Ki67+ cells. In control mice, ~47% and ~30% of total cells were LacZ+ cells in DG and VZ/SVZ respectively (Figure 5A and Table 1). Compared with controls, the number of LacZ+ cells in IGF1R

Nestin–KO mice was decreased by ~53% in DG and by ~24% in VZ/SVZ. Similarly, the number of Ki67+ cells was also decreased in IGF1R

Nestin–KO mice by ~40% and ~26% in DG and VZ/SVZ respectively (Table 1). Intriguingly, there is only a fraction of Ki67+ proliferating cells that also exhibit active expression of the LacZ reporter transgene. The reason(s) for this is not clear. One of the possible explanations is that β-catenin can activate nuclear Wnt signalling in a TCF-independent manner (Filali et al., 2002). Regardless, however, the number of cells positive for both LacZ and Ki67 in the DG and VZ/SVZ of IGF1R

Nestin–KO

KO mice.

Figure 4  IGF-I increase of LacZTCF reporter transgene expression in cultures derived from E14.5 frontal cortex (A) Representative microphotographs of LacZ immunostained cortical cultures treated without (C) or with 100 ng/ml IGF-I (I), 10 ng/ml Wnt3a (W), or a combination of IGF-I and Wnt3a (I+W) for 24 h. Arrows indicate LacZ+ cells. (B) Quantification of LacZ+ cells. Values represent means ± S.E.M. from 3–4 samples. *P<0.05; **P<0.01, compared with untreated controls.

Figure 5  LacZ+ cells and Ki67+ cells in E17.5 LacZTCF/IGF1R

Nestin–KO mice (A) Representative microphotographs of LacZ and Ki67 double immunostaining in the HIP and DG regions of a LacZTCF/IGF1R

Nestin–KO mouse (KO) or of a LacZ

control mouse (Contl). DG, dentate gyrus; LV, lateral ventricle. Arrows indicate CA regions and arrowheads point to lateral ventricles. (B) Representative microphotographs of double-immunostained DG at a high magnification. Arrows point to cells that are positive for both LacZ and Ki67, and arrowheads indicate LacZ+ and Ki67—cells.
mice was altered in a pattern similar to LacZ+ cells or Ki67+ cells, being ~30% and ~63% of that in controls respectively (Table 1).

It has been shown that, in *Xenopus* embryo, IGF-I is capable of regulating the expression of several Wnt mRNAs (Carron et al., 2005). To determine whether IGF signalling also can regulate the expression of other canonical Wnt signalling molecules in mouse brain, we quantified the abundance of mRNA for Wnt ligands, frizzled receptors and TCF-4 transcription factor, which are highly expressed in the CNS in a spatial- and temporal-specific pattern (Roelink and Nusse, 1991; Parr et al., 1993; Hollyday et al., 1995; Cho and Dressler, 1998). As shown in Figure 6, the mRNA abundance of the Wnt ligands and Frizzled receptors examined, except for Wnt 3a and Frizzled 3, was similar in the brain of E17.5 IGF1R* Nestin* KO mice and control mice. The abundance of TCF-4 mRNA in IGF1R* Nestin* KO mice also did not differ from that in control mice (Figure 6). In contrast, the mRNA abundance of both Wnt 3a and Frizzled 3 in the brain of IGF1R* Nestin* KO mice was significantly increased, being ~400% and 130% of that in control mice respectively (Figure 6).

Blunting IGF–IGF1R signalling in mutant mice significantly reduces neural cell proliferation, survival and development (Zeger et al., 2007; Liu et al., 2009b; Lehtinen et al., 2011; Liu et al., 2011). If β-catenin is a downstream signalling molecule in the IGF–IGF1R–Akt pathway, increasing β-catenin abundance could, at least in part, rescue the phenotypic changes induced by IGF1R deficiency. To test this possibility, we increased the abundance of brain β-catenin by treating pregnant dams bearing E11.5 IGF1R* Nestin* KO embryos with lithium. Lithium, a widely used clinical medicine to control mood (Emilien et al., 1995; Cookson, 2001), has been shown to suppress GSK3β activity (by increasing GSK3β phosphorylation), resulting in an increase in β-catenin stability, and, thus, its abundance. In addition, lithium has no obvious effects on Akt activity in cultured oligodendrocyte precursors (Ye et al., 2010) or in brain (De et al., 2002), or the activity of PKA (protein kinase A), Erk1 (extracellular-signal-regulated kinase 1) and CKII (casein kinase II) in vitro (Klein and Melton, 1996).

As we reported above, the abundance of inactive pGSK3βSer9 and β-catenin was significantly reduced in the brain of IGF1R* Nestin* KO mice treated with NaCl, being ~40% Table 1 The number of LacZ+ cells, Ki67+ cells and LacZ+/Ki67+ cells in the VZ/SVZ and DG of IGF1R* Nestin* KO mice and controls

| Brain Region | Control mice | IGF1R* Nestin* KO mice |
|-------------|--------------|------------------------|
| VZ/SVZ      |              |                        |
| LacZ+ cells | 29.96 ± 0.90 | 22.87 ± 1.44 *         |
| Ki67+ cells | 52.74 ± 0.52 | 39.18 ± 0.17 **        |
| LacZ+/Ki67+ | 19.29 ± 0.94 | 12.16 ± 0.43 *         |
| DG          |              |                        |
| LacZ+ cells | 46.69 ± 3.79 | 22.10 ± 1.08 *         |
| Ki67+ cells | 50.10 ± 6.79 | 30.15 ± 1.08 *         |
| LacZ+/Ki67+ | 23.82 ± 0.23 | 6.92 ± 0.74 **         |

Figure 6 mRNA expression of Wnt ligands, Frizzled receptors, and TCF-4 transcription factor (T) in the brains of E17.5 IGF1R* Nestin* KO mutant (KO) and control (Contl) mice

Values represent means ± S.E.M. from 4–5 samples. *P<0.05; **P<0.01, compared with control mice.
of that in NaCl-treated littermate controls (Figure 7). Compared with LiCl-treated littermate controls, the abundance of pGSK3βSer9 and β-catenin also was reduced in the IGF1RNestin−K0 mice treated with LiCl for 6 days. The magnitude of the reduction, however, was much smaller, and the abundance of pGSK3βSer9 and β-catenin proteins in LiCl-treated IGF1RNestin−K0 mice was twice as much as that in NaCl-treated IGF1RNestin−K0 mice (Figure 7). Similar results were also observed for non-phosphorylated and active β-catenin proteins (results not shown).

Next, we examined the morphology of the HIP, a brain region with a distinct cytoarchitecture that facilitates our...
LiCl treatment markedly mitigated the detrimental effects of IGF1R ablation on the growth of brain and HIP (Figure 8). When compared with NaCl-treated IGF1R Nestin KO mice, the brain weight of E17.5 IGF1R Nestin KO mice that were previously treated with LiCl for 6 days was ~40% greater (Figure 8A). At other developmental ages, i.e., E16.5 and E18.5, similar results were also observed in IGF1R Nestin KO mice that were respectively treated with LiCl for 5 or 7 days (Supplementary Figure S2 available online at http://www.asneuro.org/an/004/an004e092add.htm), further validating LiCl effects on brain growth in IGF1R Nestin KO mice. Consistently, the size of HIP in LiCl-treated E17.5 IGF1R Nestin KO mice was ~48% greater than that in NaCl-treated IGF1R Nestin KO mice (Figures 8B and 8C). The cortex plate in LiCl-treated IGF1R Nestin KO mice also was ~20% thicker (results not shown).

As with HIP, the volume of CA PCL was significantly reduced in NaCl-treated mice and LiCl-treated IGF1R Nestin KO mice. The density of PCL neurons was similar in all groups of mice (Table 2), and thus, the reduction in the PCL neuron number in IGF1R Nestin KO mice was largely due to the decrease in PCL volume (Figure 8D). The magnitude of the reduction in CA PCL volume and the number of PCL neurons, however, was much smaller in LiCl-treated IGF1R Nestin KO mice. Compared with NaCl-treated IGF1R Nestin KO mice, the PCL volume and PCL neurons in LiCl-treated IGF1R Nestin KO mice was 63–66% greater (Figure 8D). Because DG and hilus could not be accurately delineated in IGF1R Nestin KO mice due to dramatic growth retardation (Figures 3 and 7), the number of neurons in the DG region was not analysed.

Next, we quantified the effects of LiCl on neural proliferation by determining the number of the VZ neural precursors positive for pH3 Ser10, a nuclear marker for the proliferating cells that are in the M phase of the cell cycle. Consistent with our previous report that IGF increases the number of pH3 Ser10+ neural cells in embryonic VZ explants (Lehtinen et al., 2011), blunting IGF1R expression significantly reduced the number of pH3 Ser10+ cells in the VZ of IGF1R Nestin KO mice. Compared with NaCl-treated controls, NaCl-treated IGF1R Nestin KO mice exhibited ~64% reduction in proliferating pH3 Ser10+ precursors in VZ (Figure 9). LiCl-treated IGF1R Nestin KO mice also exhibited a decreased number of pH3 Ser10+ cells in the VZ (Figure 9).

Table 2

| NaCl-treated | LiCl-treated |
|--------------|--------------|
| Control      | IGF1R Nestin KO | Control | IGF1R Nestin KO |
| CA 1–2       | 11379.13 ± 583.90 | 11074.60 ± 173.08 | 10731.84 ± 353.62 | 11412.36 ± 172.06 |
| CA3          | 10228.13 ± 493.22 | 11710.94 ± 367.40 | 10459.41 ± 498.51 | 10889.19 ± 213.92 |
| CTX          | 11623.20 ± 926.54 | 12685.73 ± 511.19 | 10556.29 ± 605.07 | 11894.49 ± 910.18 |

Figure 9 Proliferating neural precursors in the VZ/SVZ of E17.5 IGF1R Nestin KO mutant mice (KO) and their control littermates (Conti) that were previously treated with LiCl or NaCl for 6 days.

(A) Representative microphotographs of VZ pH3 Ser10 immunostaining. Arrows indicate pH3 Ser10+ positive proliferating cells. (B) Quantification of pH3 Ser10+ positive proliferating cells. Values represent means ± S.E.M. from 3–4 samples. *P<0.001, compared with their respective control mice; **P<0.01, compared with NaCl-treated IGF1R Nestin KO mice.
of pH3Ser10+ precursors, but the magnitude of the reduction was much smaller. When compared with NaCl-treated IGF1R
Nestin−/− KO mice, LiCl-treated IGF1R Nestin−/− KO mice had 
~72% more pH3Ser10+ cells in VZ (Figure 9). Quantification of proliferating neural precursors positive for Ki67, a marker for proliferating cells in the G1, G2 and S phases of the cell cycle, showed changes in a similar pattern (Table 3 and Supplementary Figure S3 available online at http://www.asnneuro.org/an/004/an004e092add.htm). Treatment of E11.5

| Control mice | IGF1R Nestin−/− KO mice |
|--------------|------------------------|
| VZ/SVZ       |                        |
| NaCl treated | 70.60 ± 2.71           |
| LiCl treated | 69.70 ± 1.95           |
| DG           |                        |
| NaCl treated | 43.79 ± 2.20           |
| LiCl treated | 40.93 ± 2.57           |

Table 3 The number of Ki67+ proliferating neural precursors in the VZ/SVZ and DG of IGF1R Nestin−/− KO mice and controls treated with LiCl or NaCl at age of E11.5 were treated with LiCl or NaCl for 6 days and killed at age of E17.5. Ki67+ cells were scored on the sections corresponding to plate GD18 Cor. 10 (VZ/SVZ) or plate Cor 14 (DG) respectively in Schambra et al. (1992). Values are expressed as a percentage of total number of cells, and represent means ± S.E.M. from 3–4 samples. *P < 0.05; **P < 0.01; ***P < 0.001, compared with their respective control mice treated with NaCl or LiCl; P<0.05; †P<0.01, compared with NaCl-treated IGF1R Nestin−/− KO mice.

In contrast, only a few apoptotic cells (as judged by immunolabelling for activated caspase 3) were detected in the VZ and other brain regions in E17.5 IGF1R Nestin−/− KO mice and control mice (Supplementary Figure S4 available at http://www.asnneuro.org/an/004/an004e092add.htm) regardless of treatment. The number of apoptotic cells in the VZ of

Figure 10 Hippocampal growth and neural precursor proliferation in the VZ/SVZ of E17.5 IGF1R Nestin−/− KO mutant mice (KO) and their control littermates (Contl) that were previously treated with AR-A014418 (AR) or PBS for 6 days (A) Representative microphotographs of HIP. Arrows in the bottom panels indicate CA region. (B) CA pyramidal neuron number. (C) Representative microphotographs of VZ pH3Ser10 immunostaining. Arrows indicate pH3Ser10+ proliferating cells. (D) Quantification of pH3Ser10+ proliferating cells. In (B, D), values represent means ± S.E.M. from 3 samples. *P<0.05; **P<0.01, compared with their respective control mice; †P<0.05, compared with PBS-treated IGF1R Nestin−/− KO mice.
IGF1R\textsubscript{Nestin–KO} mice did not differ from that in control mice (results not shown), a finding that is consistent with our previous report (Lehtinen et al., 2011).

**DISCUSSION**

The data from our current study strongly support the concept that, in developing neural cells, \(\beta\)-catenin is a signalling molecule downstream of IGF–I–PI3K–Akt–GSK3\(\beta\), and that IGF signalling can interact with canonical Wnt–\(\beta\)-catenin signalling to promote neural precursor proliferation in vivo. Specifically, we have shown that, in the brain of mutant mice, blunting the expression of neural IGF-I or IGF1R significantly reduces: (i) the abundance of pGSK3\(\beta\), an enzyme known to regulate \(\beta\)-catenin stability, (ii) the expression of \(\beta\)-catenin, a key member of the canonical Wnt signalling pathway, at both mRNA and protein levels and (iii) the activity of the LacZ\textsuperscript{TCF} reporter transgene that responds to nuclear Wnt signalling stimulation. In contrast, overexpressing IGF-I in LacZ\textsuperscript{TCF} Tg brain and IGF-I treatment of cultured neuronal cells derived from LacZ\textsuperscript{TCF} brain markedly increases the expression of the LacZ\textsuperscript{TCF} reporter transgene. Furthermore, increasing \(\beta\)-catenin abundance by suppressing pGSK3\(\beta\) activity drastically reduces the detrimental effects of IGF1R deficiency on neural growth, as evidenced by increases in brain weight, the volume of HIP and CA, the number of PCL neurons, and the neural proliferation capacity in IGF1R\textsubscript{Nestin–KO} mice.

Consistent with our previous report showing a critical role for neural IGF signalling during postnatal life (Liu et al., 2009b), our current study also demonstrates that blunting IGF1R expression specifically in the nestin+ neural precursors and their progeny significantly reduces the weight of developing brain, the volume of HIP and the number of pyramidal neurons during embryonic development. These results support a critical role for IGF–IGF1R signalling in neurogenesis and brain growth during normal prenatal development. At E17.5, when neural precursors actively proliferated in both VZ/SVZ and HIP, deficiency in IGF–IGF1R signalling leads to a significant reduction in the number of proliferating neural precursors. In contrast, less than 0.05% of active caspase 3 positive apoptotic cells are detected in VZ and other brain regions, and no significant differences are observed between IGF1R\textsubscript{Nestin–KO} mice and control mice. These data are consistent with our previous study of Tg mice overexpressing IGF-I in nestin+ neural precursors (Popken et al., 2004), and suggest that, at this stage of brain development, neural apoptosis is unlikely a major factor in neurogenesis; rather, precursor proliferation plays a significant role in response to IGF stimulation.

The Erk MAPK (mitogen-activated protein kinase) pathway has been thought to play a major role in cell proliferation. However, accumulating evidence indicates that in neural cells the PI3K-Akt pathway plays a critical role in cell proliferation and is likely required for a full mitogenic response to IGF-I. Overexpression of Akt enhances the proliferation of cortical neural precursors (Sinor and Lillien, 2004). Conversely, inhibiting PI3K-Akt activity markedly suppresses the IGF-I-stimulated proliferation in cultured cortical neural precursors (Mairet-Coello et al., 2009) and cerebellar granule cell precursors (Cui et al., 1998). In line with these reports, our previous studies of cultured oligodendrogial precursor cells also demonstrate that inhibition of PI3K-Akt activity hinders the IGF-I-stimulated expression of mRNA for cyclin D1 protein (Ye et al., 2010), a molecule key to the cell-cycle progression through the G\(_1\)/S phases. The IGF-I stimulatory effects on cyclin D1 expression are partially mediated by \(\beta\)-catenin (Ye et al., 2010), a key member of the canonical Wnt signalling pathway, indicating that IGF signalling can interact with Wnt canonical signalling, via the PI3K-Akt-\(\beta\)-catenin pathway, to promote their proliferations. While the predominant neuronal expression of the nestin-driven Cre transgene in our mutant mice (Liu et al., 2009b) precludes studies on oligodendrocyte lineage cells in vivo, our current study clearly supports the concept that \(\beta\)-catenin plays an important role in IGF proliferation signalling during neuronal development by showing that in neural precursor cells: (i) IGF signalling regulates the abundance of \(\beta\)-catenin and the activity of the LacZ\textsuperscript{TCF} reporter transgene, which responds to nuclear Wnt signalling stimulation, and (ii) increasing \(\beta\)-catenin abundance mitigates the detrimental effects of IGF1R deficiency on neural proliferation.

Our data also show that ablating neural IGF1R expression significantly reduces the abundance of \(\beta\)-catenin protein and its mRNA in the brain of mutant mice, suggesting that IGF signalling regulates the expression of \(\beta\)-catenin at both the protein and mRNA level. These findings are consistent with our previous study of cultured neural cells (Ye et al., 2010). In cultured oligodendrocytes, IGF-I rapidly increases the abundance of \(\beta\)-catenin protein as early as 1 h after treatment, likely by enhancing its stability. In contrast, the expression of \(\beta\)-catenin mRNA is not altered during first 4 h of treatment and becomes significantly up-regulated after 24 h of treatment (Ye et al., 2010). Despite the distinct patterns of \(\beta\)-catenin protein and its mRNA expression in response to IGF-I stimulation, inhibition of PI3K-Akt kinases significantly suppresses the IGF-I-stimulated increases in both \(\beta\)-catenin protein and mRNA (Ye et al., 2010), indicating a critical role for PI3K-Akt in IGF-I stimulatory actions on both \(\beta\)-catenin protein and mRNA in cultured oligodendrocytes. Whether IGF signalling also regulates the expression of \(\beta\)-catenin protein and its mRNA in a similar pattern in vivo remains to be determined.

While the brain expression of individual members in the Wnt signalling pathway has not been fully defined, multiple Wnt ligands, Frizzled receptors and TCF/LEF (lymphoid enhancer factor) family transcription factors are highly expressed in the CNS in a spatial- and temporal-specific pattern (Roelink and Nusse, 1991; Parr et al., 1993; Hollyday et al., 1995; Cui and Bulleit, 1998). Consistent with these reports, we also observed abundant expression of mRNA for multiple Wnt ligands and Frizzled receptors, as well as TCF-4 transcription factor, in developing forebrains. The
mRNA abundance of these proteins, however, is similar in E17.5 IGFR1\textsuperscript{Nestin\textsuperscript{−/−}KO} mice and their littermate controls, except for Wnt 3a mRNA and Frizzled 3 mRNA, both of which are increased in IGFR1\textsuperscript{Nestin\textsuperscript{−/−}KO} mice. As β-catenin, a molecule downstream of Wnt and Frizzled receptor, is significantly decreased in IGFR1\textsuperscript{Nestin\textsuperscript{−/−}KO} mice, we interpret the increased expression of Wnt 3a and Frizzled 3 mRNA as compensatory to reduced β-catenin signalling, and deem that IGF signalling and Wnt signalling are likely to interact mainly at the levels of GSK3β and β-catenin, although other mechanisms may also exist.

In summary, our new findings strongly point to a critically important role for the IGF and Wnt-β-catenin signalling interaction during normal in vivo neurogenesis, and suggest that β-catenin is a common effector mediating a portion of IGF and Wnt signalling to promote neural cell proliferation. Our conclusion is further supported by an earlier report that IGF-I signalling is capable of interacting with Wnt-JNK (c-Jun N-terminal kinase) signalling in Xenopus embryo (Carron et al., 2005), albeit their actions are likely different in early stages of embryonic development. More studies are needed to precisely define IGF signalling pathways and their interaction with Wnt signalling.

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IGF and Wnt signalling interacts in neuronal cells

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