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Neurodevelopmental effects of insulin-like growth factor signaling

John O'Kusky^a and Ping Ye^{b,*}

^aDepartment of Pathology and Laboratory Medicine, University of British Columbia, Vancouver, British Columbia, Canada V5Z 1M9

^bDepartment of Pediatrics, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599, United States

Abstract

Insulin-like growth factor (IGF) signaling greatly impacts the development and growth of the central nervous system (CNS). IGF-I and IGF-II, two ligands of the IGF system, exert a wide variety of actions both during development and in adulthood, promoting the survival and proliferation of neural cells. The IGFs also influence the growth and maturation of neural cells, augmenting dendritic growth and spine formation, axon outgrowth, synaptogenesis, and myelination. Specific IGF actions, however, likely depend on cell type, developmental stage, and local microenvironmental milieu within the brain. Emerging research also indicates that alterations in IGF signaling likely contribute to the pathogenesis of some neurological disorders. This review summarizes experimental studies and shed light on the critical roles of IGF signaling, as well as its mechanisms, during CNS development.

Keywords

IGF-I; IGF-II; IGF1R; CNS; Development; Neurons; Glial cells

1. Introduction

Experimental evidence accumulated during the past two decades has convincingly established an essential role for insulin-like growth factor (IGF) signaling in the normal growth and development of the central nervous system (CNS). IGF-I and IGF-II, two members of the IGF system, share homology with each other and with proinsulin. In developing brain, IGF signaling exerts pleiotropic actions on all major neural cell types [including neural stem cells (NSCs), lineage restricted neural precursor cells (NPCs), post-mitotic neurons, oligodendrocytes and astrocytes]. IGFs act to promote the proliferation, maturation, survival, and/or growth of neural cells, predominantly, if not exclusively, by interacting with the type 1 IGF receptor (IGF1R). The biological nature of the actions, however, appears to depend on the specific cell types in place, the local microenvironment, and the particular stage of development. IGF signaling also appears to influence specific biological processes in concert with additional neural signaling, which may provide primary instructive signals to steer NSC toward a specific cell lineage during early development.

Our current knowledge of IGF signaling in the developing brain comes predominantly from a wide variety of *in vitro* and *in vivo* experimental studies, the latter primarily derived from studies of mutant mouse models. Nevertheless, individual patients with mutation(s) in the *igf-I* gene (Camacho-Hubner et al., 1999; Woods et al., 1997) or in the *igf1r* gene (Abuzzahab et al., 2003; Kruis et al., 2010; Okubo et al., 2003; Wallborn et al., 2010; Woods et al., 1997) are found to be associated with severe body growth failure, microcephaly, and mental retardation, strongly arguing for a similar role for IGF signaling during CNS development in humans. Recent literature also has established that many of growth-related phenomena in neural cells, such as neurogenesis (Gage, 2002; Kelsch et al., 2010; Ming and Song, 2011), axon remodeling and *de novo* synaptogenesis (Bruehl-Jungerman et al., 2007; Butefisch, 2006; Carmichael, 2003; Cayre et al., 2009; Gogolla et al., 2007), persist throughout adult life. In parallel, IGFs and their receptors are steadily expressed in the adult brain in a spatial-specific pattern, albeit at relatively lower levels, and are thought to have a significant role in the pathogenesis of several growth-related neurological disorders. In this article, we will review the actions of IGF signaling on brain neural cells with a focus on IGF actions during prenatal and early postnatal life.

2. Overview of the IGF system

The IGF system is traditionally comprised of IGF-I, IGF-II, the IGF1R, the type 2 IGF receptor (IGF2R), and IGF binding proteins (IGFBPs). The growth-promoting actions of IGF-I and IGF-II are pre-dominantly, if not exclusively, mediated by the IGF1R. The receptor binding and biological activities of IGFs are modulated by IGFBPs. At least 10 IGFBPs, including 6 high-affinity IGFBPs and 4 low-affinity IGFBPs, have been identified.

In mice, the actions of IGF–IGF1R signaling appear to be significantly influenced by genomic background. For example, 95% of mice carrying a null mutation (knockout, KO) in the *igf-I* gene (*igf-I*KO mice) that are on C75B/6 background die perinatally (Powell-Braxton et al., 1993), while more than 50% of *igf-I*KO mice on a mixed genomic background of 129/MF1 survive postnatally (Liu et al., 1993). Similar phenomena are also observed in mice with ablated IGF1R expression specifically in nestin-positive (+) neural precursors (Kappeler et al., 2008; Liu et al., 2009). Gene modification of IGF signaling may also play an important role in the development and growth of the CNS. The details, however, remain to be elucidated. Both IGF-I and IGF-II at high concentrations also can bind to the insulin receptor (InR), and InR is capable of mediating IGF-II actions (Louvi et al., 1997; Morriane et al., 1997).

2.1. IGF-I, IGF-II and derivative forms

IGF-I and IGF-II are anabolic peptides (70 and 67 amino acids, respectively), sharing homology with each other and with proinsulin (Daughaday and Rotwein, 1989; Rotwein, 1991). Each of these growth factors is produced by a single large gene (95 kb and 35 kb, respectively), with expression beginning early in embryonic development. IGF-I is produced by all types of major neural cells in the brain. In the brain, IGF-II is more abundantly expressed than IGF-I during prenatal development. During postnatal development somatic IGF-I is regulated by pituitary growth hormone (GH), mediating most of GH's growth promoting actions. Brain IGF-I expression is also likely regulated by GH to certain extent during development (Hojvat et al., 1982; Hynes et al., 1987; Ye et al., 1997). The mechanisms regulating the expression of the *igf-I* and *igf-II* genes are still largely unclear.

Variant forms of IGF-I exist in the brain (Ballard et al., 1987). IGF-I is believed to undergo post-translational N-terminal cleavage by a specific protease into des-N-(1–3) IGF-I, which appears to be the dominant form in the brain (Ballard et al., 1987; Sara et al., 1986; Yamamoto and Murphy, 1995). In organ culture of new born rat olfactory bulb, des-N-(1–3)

IGF-I is shown to be potent supporter of viability, cell survival and differentiated cell growth (Russo and Werther, 1994). Des-N-(1–3) IGF-I has been found to have an increased potency for neuroprotective (i.e. anti-apoptotic) effects on neurons. This is likely due to its lower affinity for IGFBPs caused by the absence of glutamate at position 3 (Guan et al., 1996), and thus, relatively higher local concentration of its free form. The N-terminal gly-pro-glu tripeptide fragment (GPE tripeptide), generated along with des-N-(1–3) IGF-I during proteolytic cleavage of IGF-I, also is capable of mediating neuroprotective effects both *in vivo* and *in vitro* (Guan et al., 2000; Saura et al., 1999; Shapira et al., 2009), although it remains to be determined whether GPE actions depend on IGF1R. GPE tripeptide has more localized sites of action in the adult brain, including hippocampal CA1-2, pyriform cortex, amygdala, cerebral cortex, choroid plexus and blood vessels (Ikeda et al., 1995; Saura et al., 1999).

2.2. IGF1R and IGF2R

The IGF1R is a heterotetrameric glycoprotein composed of paired, disulfide-linked α - and β -subunits. The α -subunits are extracellular and bind IGFs. The β -subunits contain a long intracytoplasmic domain which contains intrinsic tyrosine kinase activity and critical sites of tyrosine and serine phosphorylation. The IGF1R shares a 46% homology with the InR. The IGF1R and the InR can form hybrid receptors by using the α - and β -subunits of each to form heterodimers. IGF1R/InR hybrids retain capability to transduce both IGF and/or insulin signaling, although the exact functional significance of these hybrid receptors is unknown.

The IGF1R binds IGF-I with high affinity and both IGF-II and insulin with lower affinities (approximately 10-fold and 100-fold, respectively). The IGF1R is expressed in NSC and in all neural cells evaluated (Baron-Van Evercooren et al., 1991). Binding of IGFs to the α -subunit of the IGF1R induces a conformational change in the receptor that results in autophosphorylation of the β -subunit, setting signaling cascades into motion that involve phosphorylation of a series of intracellular substrate proteins (LeRoith et al., 1995), such as insulin receptor substrate (IRS)-1 and IRS-2 as described in detail in Section 6 below.

The IGF2R, a single chain transmembrane protein, is identical to the cation-independent mannose-6-phosphate receptor, and acts to translocate proteins containing mannose-6-phosphate moieties and IGF-II to lysosomes for degradation. Global ablation of IGF2R expression leads to overgrowth, resulting from an increased accumulation of IGF-II (Efstratiadis, 1998; Eggenschwiler et al., 1997). No intrinsic enzymatic activity has been observed for the intracellular domain of the IGF2R. There is also little evidence that IGF-I interact with the IGF2R, and that the IGF2R mediates IGF growth promoting activity in the brain.

2.3. IGFBP

IGF-I and IGF-II are found in circulation and in the extracellular space of most tissues almost completely bound to members of the family of IGFBPs. There are at least 10 IGFBPs, i.e., 6 high-affinity IGFBPs and 4 low-affinity IGFBPs. Six high affinity IGFBPs, designated IGFBP-1 through IGFBP-6 (Jones and Clemmons, 1995), share structural homology with each other, and bind specifically to IGF-I and IGF-II with negligible affinity for insulin. IGFBP-2, -3, -4 and -5 are the most abundant IGFBPs in brain. IGFBP-6 is also expressed, IGFBP-1, however, is not detected in brain during normal development (Ocran et al., 1990). Each IGFBP is expressed in the CNS in a specific temporal-spatial pattern. The exact functions of IGFBPs in the CNS remain to be fully elucidated. The IGFBPs have been proposed to act: (1) as transport proteins in plasma, (2) to prolong the half lives of the IGFs in circulation, (3) to determine the tissue-and cell-specific localization of IGF-I and IGF-II, and (4) to control the biological actions of IGF-I and IGF-II by modulating their interactions

with their receptors (Jones and Clemmons, 1995). In addition, some IGFBPs can also exert IGF independent actions. For example, IGFBP-1 activates integrin-mediated intracellular signaling in trophoblast (Gleeson et al., 2001), and enhances oligodendrocytes migration (Chesik et al., 2010).

Approximately 75% of IGF-I and IGF-II in circulation is carried by a complex of IGF-I or IGF-II, IGFBP-3 and a non-IGF-binding component termed acid labile subunit (ALS). Binding of IGF-I or IGF-II to IGFBP-3 in the presence of ALS produces an IGF-IGFBP-3-ALS ternary complex, which is stabilized by IGF binding (Baxter and Martin, 1989). The IGFs do not readily leave the vascular compartment, when associated with this complex, and their half lives are significantly prolonged by 70- to 90-fold (Guler et al., 1989; Hodgkinson et al., 1989). This circulating pool of bound IGFs is thought to serve as a local reservoir in times of stress. IGFBPs with lower molecular weights (e.g. IGFBP-1, IGFBP-2 and IGFBP-4) have also been implicated as IGF carrier proteins, although their mechanisms of action are less well understood. Estimates of the half lives for these IGFBPs suggest they are cleared more rapidly than the IGF-IGFBP-3-ALS ternary complex (Rechler, 1993).

3. Ontogeny of IGF-I, IGF-II, IGF1R and IGF2R in brain

In the developing CNS, IGFs and IGF receptors are widely expressed in a tempo-spatial specific manner. Table 1 summarizes the expression of IGF-I, as well as that of IGF-II, IGF1R and IGF2R, in the major brain areas during perinatal development and in adult (Ayer-le et al., 1991; Bartlett et al., 1992; Bondy and Chin, 1991; Bondy and Lee, 1993; Bondy, 1991; Cavallaro et al., 1993; Dugas et al., 2008; Folli et al., 1994; Hawkes and Kar, 2003; Kar et al., 1993; Lee et al., 1993; Logan et al., 1994; Quesada et al., 2007; Stylianopoulou et al., 1988; Walter et al., 1999; Zhang et al., 2007). While the mechanisms that regulate the expression of each of these genes still are not completely understood, IGFs and IGF1R are found to be often expressed within close proximity, strongly indicating that IGF acts locally in developing brain in an autocrine and/or paracrine fashion. Nonetheless, IGF-I can be transported across the blood brain barrier (BBB) Pulford and Ishii, 2001; Reinhardt and Bondy, 1994, and thus, it is likely that circulating IGF can influence neurogenesis and their development (Aberg et al., 2007; Anderson et al., 2002).

IGF-I and the IGF1R also are expressed in non-neural tissues in early embryonic development before neural tissues have established (Alarcon et al., 1998; Ayaso et al., 2002; Bondy et al., 1990; de et al., 1993; Morales et al., 1997; Perez-Villamil et al., 1994; Scavo et al., 1991a; Scavo et al., 1991b). In chicken embryo, IGF-I mRNA can be detected at embryonic day (E) 3, while insulin is detected earlier at E0 (Perez-Villamil et al., 1994). The InR and IGF1R are present in the blastoderm at E0 through late organogenesis at E9 (Scavo et al., 1991b). IGF-I is preferentially expressed in cephalic regions during late neurulation and throughout organogenesis, and has been shown to be compartmentalized to the epithelial cells of developing eyes (de et al., 1993). The InR and the IGF1R are localized at their highest levels in Hensen's node, neural folds, neural tube and developing eyes (Girbau et al., 1989). In rodents, IGF-I mRNA is particularly abundant in undifferentiated mesenchymal tissue and does not become significantly evident until E14. IGF-II mRNA was also conspicuous in areas of vascular interface with the brain, such as the choroid plexus and the organum vasculosum of the lamina terminalis. IGF1R mRNA is widely distributed in embryonic tissues, but the highest levels are seen in the ventral floorplate of the hindbrain, where specialized neuroepithelial cells act as guides for axonal targeting (Bondy et al., 1990).

3.1. IGF-I

IGF-I is expressed in all regions of the CNS. In rodents, significant IGF-I expression is detected as early as E11 in most regions of rodent brain (Ayer-le et al., 1991). Brain IGF-I expression in rodents peaks in the second week of postnatal life, gradually decreases, but continues throughout life (Bach et al., 1991; Bartlett et al., 1991; Ye et al., 1997). The peak expression of local IGF-I is often spatially correlated with the active proliferation, development and growth of neural cells. During normal development, IGF-I production is predominately located in neurons, and to a lesser extent in glial cells (Bondy and Lee, 1993; Bondy, 1991; Bondy and Lee, 1993; Bondy et al., 1990; Shinar and McMorris, 1995). IGF-I mRNA also is detected in the postnatal subventricular zone (SVZ) (Bartlett et al., 1992; Perez-Martin et al., 2003 and in the Ki67+ proliferating precursors in the dentate gyrus (DG) (Zhang et al., 2007, strongly suggesting that IGF-I is likely to be produced by proliferating neural precursors.

3.2. IGF-II

IGF-II is highly expressed in mesenchymal tissues. In brain, unlike IGF-I which peaks during postnatal development, the highest levels of brain IGF-II expression are observed during embryonic development (Ayer-le et al., 1991; Bondy et al., 1990). With increasing age, parenchymal IGF-II mRNA expression is gradually reduced, and its expression becomes restricted to meninges and choroid plexus in adult (Hynes et al., 1988; Logan et al., 1994; Zhang et al., 2007). Consistent with studies using RNA *in situ* hybridization, analysis using laser capture microdissection followed by polymerase chain reaction (PCR) demonstrated that there is little IGF-II mRNA expression, if any, in the postnatal hippocampal neurons (Zhang et al., 2007). Meninges and choroid plexus are major sources of cerebro-spinal fluid (CSF) IGF-II, which appears to be a critically important stimulator for NSC proliferation in the ventricular zone (VZ) during embryonic development (Lehtinen et al., 2011). Nonetheless, IGF-II protein immunoreactivity is also observed in other brain regions in adult (Logan et al., 1994), suggesting that IGF-II is likely produced at meninges and choroid plexus and transferred to these locations.

3.3. IGF1R

The IGF1R also is ubiquitously expressed in all neural cell types, including NSC and NPC (Popken et al., 2005), and its abundance is positively correlated with cell proliferation and growth. In a detailed study of mouse hippocampus, we show that DG proliferating progenitors have the most abundant amount of IGF1R mRNA, although all hippocampal neurons express it (Zhang et al., 2007). An appropriate expression pattern of the IGF1R is likely to pertinently contribute to its proper functionality. During prenatal development when neuronal processes actively outgrow, IGF1R is significantly enriched in the growth cone (Quiroga et al., 1995). In the cortical VZ of murine embryonic brains, IGF1R expression is restricted to the apical domain of neural precursors, likely by protein associated with Lin 7 (Pals1) and phosphatase and tensin homolog (Pten) (Lehtinen et al., 2011). Disruption of Pals1 or Pten expression in embryos alters the expression pattern of IGF1R and the rate of cell proliferation and brain growth (Lehtinen et al., 2011). Focal adhesion kinase (FAK) also stabilizes IGF-1 receptor in cultured mouse embryonic fibroblasts (Andersson et al., 2009), whether FAK acts same in neural cells remains to be determined.

4. IGF-I actions in the CNS

The development of the mammalian brain occurs along specific stages, including neurulation, neurogenesis, differentiation into neurons and glia, neuronal migration, dendritic and axon out-growth, naturally occurring cell death, synaptogenesis, and

myelination. The time course for these stages differs among species. In general, there is a caudal-to-rostral gradient in the time course of these developmental stages for individual regions in a given brain. The growth promoting actions of IGF-I-IGF1R signaling in the developing brain have been documented at virtually every stage of CNS development.

4.1. Early embryonic development

IGF-I has been shown to increase the survival of mammalian embryos by increasing the proportion of preimplantation embryos that eventually become blastocysts. This occurs during normal development (Lima et al., 2006; Sirisathien et al., 2003) and during abnormal development in response to various stressors such as heat shock, oxidative stress, tumor necrosis factor- α and toxicity (Byrne et al., 2002; Fabian et al., 2004; Jousan and Hansen, 2007; Jousan et al., 2008; Moss et al., 2009). These protective actions of IGF-I appear to be developmentally regulated. Bovine embryos before implantation (16 cells) treated with IGF-I at gestational day 5 exhibit altered gene expression, including upregulation of 5 anti-apoptotic genes (*il6st*, *dyrk3*, *nfatc3*, *anp32*, and *EIF3A*), and downregulation of 5 pro-apoptotic genes (*dpysl4*, *mst1*, *tnfsf11a*, and *arhgef10l*), and upregulation of 2 genes involved in protection from reactive oxygen species (*gstm2* and *coo9*) Bonilla et al., 2011. Interestingly, these same embryos also exhibited downregulation of genes involved in neural development and differentiation. In animal models of maternal diabetes, hyperglycemic developmental conditions *in utero* have been shown to result in downregulation of IGF1R expression, a 40% drop in the number of preimplantation embryos surviving to become blastocysts, delayed onset and progression of gastrulation, and increased numbers of apoptotic cells in the embryonic disk (Ramin et al., 2010).

The default fate of embryonic neuroectoderm is to become neural tissue, and this process is inhibited in early stages of development (De Robertis et al., 2000) by bone morphogenic protein (BMP-4). In *Xenopus* embryos injection of IGF-I, IGF-II or IGFBP-5 mRNA promotes neural induction and head induction, likely by increasing the expression of the anterior neural transcription factors Six-3, Rx2a, Pax-6, Otx-2, while injection of a secreted dominant-negative IGF1R mRNA has the opposite effect (Pera et al., 2001). Injection of IGF mRNA also causes induction of ectopic eyes and ectopic head-like structures containing brain tissue. Conversely, blockage of IGF1R signaling in zebrafish by a dominant-negative IGF1R or specific IGF1R inhibitors not only delays the emergence of GnRH2 and GnRH3 neurons, but also results in an abnormal appearance of GnRH3 neurons (Onuma et al., 2011). This IGF action is developmental stage-dependent because IGF signaling blockade in advanced embryos has no such effect (Onuma et al., 2011). These data support a critical role for IGF signaling in anterior neural induction in non-mammalian animals. Whether IGF signaling is also capable of inducing neural tissues in mammalian remains to be determined. Current data, however, indicate otherwise. In mutant mice with ablated IGF-I-IGF1R signaling, brain cytoarchitecture in general appears to be normal, although its growth is severely retarded (Beck et al., 1995; Liu et al., 1993). However, it is possible that signaling through InR or other receptor(s) can compensate for the loss of IGF1R signaling during the early developmental stages.

During neural induction in *Xenopus* embryos, cortical rotation triggered by fertilization leads to an increase in β -catenin on the dorsal side, inhibiting BMP-4 transcription (Baker et al., 1999a), and predisposing cells in the ectoderm toward neural fates. The inhibition of Wnt signaling at the gastrula stage mediates the process of head induction (Glinka et al., 1998; Glinka et al., 1997). It has been shown that overexpression of IGF-I produces an anterior expansion of head neural tissues, while blunting IGF-I expression reduces head structures, by antagonizing the activity of the Wnt signaling pathway in the embryo at the level of β -catenin (Richard-Parpaillon et al., 2002). Furthermore, the IGF may act in concert with other potent inducers of neural induction, including the fibroblast growth factors and

Chordin, through the inhibitory phosphorylation of SMAD1 (Pera et al., 2003), culminating in a downstream inhibition of BMP-4.

4.2. Brain weight, regional brain volume and CNS growth

Scientific interest in IGF-I as a neuronal growth factor stemmed from *in vitro* studies demonstrating IGF-I's ability to promote proliferation in neural cells and to inhibit apoptosis and cell death in various culture systems (see Section 4.3 below). The true magnitude of IGF-I's role in promoting growth and development of the brain became apparent following *in vivo* studies using mutant mouse models (Fig. 1). Multiple lines of mice with genomic alterations in IGF system proteins have been generated and studied. Table 2 lists those mouse lines and changes in brain growth and cellular phenotypes. Briefly, these include transgenic (Tg) mice that overexpress IGF-I or IGF-II, mutant mice with a null mutation for IGF-I, IGF-II or IGF1R, and Tg mice with altered IGFBP expression.

Specific IGF actions likely depend on cell type, developmental stage, and local microenvironmental milieu within the brain, and the magnitude of altered brain regional growth may also in part reflect regional differences in IGF signaling in each specific mutant mouse line. In several lines of Tg mice that overexpress a metallothionein- I (MT-I) promoter-driven IGF-I transgene (IGF-I^{MT-I} Tg mice), which exhibit increased expression of IGF-I during early postnatal development, brain weight is increased from 22% to 91% with no significant change in body weight in adult (Gutierrez-Ospina et al., 1996). Consistently, morphometric studies of the primary somatosensory cortex in one IGF-I^{MT-I} Tg mouse line report an 81% increase in the volume of the cerebral cortex (Gutierrez-Ospina et al., 1996).

In Tg mice expressing an IGF-I transgene driven by IGF-II 5' regulatory sequences (IGF-I^{IGF-II} Tg mice), transgene expression and increased levels of IGF-I can only be detected in the brain, beginning at E18 and gradually increasing to plateau levels by postnatal days (P) 20 (Ye et al., 1996). Although transgene expression occurs throughout the brain in these Tg mice, the highest level of transgene expression is found in the cerebellum (Ye et al., 1996). At P30, total brain weight was increased by 28%, with regional increases observed in the cerebellum (43%), hippocampus (34%), diencephalon (28%), brainstem (28%) and cerebral cortex (9%) (Ye et al., 1996). Cerebellum weight was further increased by 90% in adult IGF-I^{IGF-II} Tg mice (Ye et al., 1996). The total volume of the medulla was found to be significantly increased in Tg mice by 27% at P35 (Dentremont et al., 1999), while individual medullary nuclei exhibited differential increases in volume, ranging from 29% in the facial nucleus to 84% in the dorsal motor nucleus of the vagus. In the hippocampal DG, the volumes of both the granule cell layer and the molecular layer were significantly greater in Tg mice after P7, exceeding control volumes by 55–66% at P130 (O'Kusky et al., 2000).

In IGF-I^{Nestin} Tg mice, IGF-I transgene expression is only observed in the brain, beginning in embryonic development as early as E13 and continuing throughout postnatal life (Popken et al., 2004). Brain weights of IGF-I^{Nestin} Tg mice were significantly greater than their non-transgenic littermate controls at E18 (6.5%) and P45 (23%), with no significant differences in body weight. Morphometric analysis of IGF-I^{Nestin} Tg embryos at E16 revealed a 25% increase in the volume of the dorsolateral telencephalic wall, corresponding to the primordial cerebral cortex. However, differential increases in tissue volume were observed in the cortical plate (52%), the intermediate zone (12%) and the combined volumes of the VZ and SVZ (26%) in IGF-I^{Nestin} embryos. Interestingly, the volume of the caudate-putamen anlage did not differ significantly between IGF-I^{Nestin} Tg and control embryos. At P12 regional tissue volumes in IGF-I^{Nestin} Tg mice were found to be significantly greater than controls in the forebrain (26%), cerebral cortex (29%), subcortical white matter (52%), caudate-putamen (37%), hippocampus (49%), DG (71%), and habenular complex (48%)

Popken et al., 2004. The relatively greater increases in volume, observed in the hippocampus, DG and habenular complex, were consistent with a significantly increased expression of the transgene in these regions as detected by *in situ* hybridization. The neocortical overgrowth in IGF-I^{Nestin} Tg mice at P12 was not uniform, differing as a function of cytoarchitectonic area. For example, significantly greater increases in cortical volume were found for the motor cortex (42%) compared to the somatosensory cortex (35%) Hodge et al., 2005.

The brain overgrowth observed in IGF-I overexpressing Tg mice also likely requires a continuous presence of IGF-I transgene expression. In a line of conditional IGF-I Tg mice, in which the expression of an IGF-I transgene in glial fibrillary acidic protein (GFAP)+ astrocytes (IGF-I^{GFAP} Tg mice) can be regulated by doxycycline (Ye et al., 2004), brains continuously overgrow from P5 to P110 when IGF-I transgene is allowed to express throughout the experimental period. Brain overgrowth, however, ceases after transgene expression is suppressed with doxycycline treatment (Ye et al., 2004).

In contrast, total IGF-I deficiency resulting from targeted disruption of the *igf-I* gene produces severe brain growth retardation. At 2 months of age mice homozygous for the null mutation (*igf-I* KO mice) exhibit a 38% decrease in brain weight and a 74% decrease in body weight relative to wild type controls (Beck et al., 1995). Decreased tissue volumes were reported in the pyramidal cell layer of the hippocampus (38%), the granule cell layer of the DG (59%), and the striatum (28%). Disproportionately greater decreases were detected in white matter regions, with a 69% decrease in the area of the anterior commissure and a 70% decrease in the thickness of the corpus callosum, due to decreased numbers of axons and oligodendrocytes (Beck et al., 1995). This preferential involvement of white matter in reduced brain volumes has been further confirmed in subsequent studies (Cheng et al., 1998).

Mice with a null mutation of the gene encoding the IGF1R (*igf1r* KO mice) invariably die at birth presumably due to respiratory failure and exhibit a 55% decrease in body weight compared to wild type controls (Liu et al., 1993). At E14 to E18, *igf1r* KO mutants exhibit corresponding growth retardation of the brain, as well as reduced volume and increased cell density in the mantle zone of the brainstem and spinal cord (Liu et al., 1993). These results suggest that brain growth retardation in these mice results predominantly from a reduction in the volume of neuropil in the CNS regions examined.

In mutant mice in which IGF1R expression is conditionally ablated in nestin-expressing neural precursors (*igf1r*^{Nestin-KO} mice), the whole-brain abundance of IGF1R protein is only ~52% and <2% of control values in heterozygous and homozygous *igf1r*^{Nestin-KO} mice, respectively (Liu et al., 2009). Homozygotes usually die within 48 h after birth. Surviving heterozygotes exhibit significant reductions in brain weight at P0 (56%), P5 (56%), P20 (59%) and P90 (60%), without significant changes in body weight (Liu et al., 2009). Regional tissue volumes for the hippocampus (CA 1–3), DG and granule cell layer of the DG were significantly reduced in heterozygotes by 44–54% at P0 and by 65–69% at P90.

Our unpublished morphometric and stereological analyses demonstrate that the volume of the telencephalic wall, corresponding to the primordial cerebral cortex, measured from the ventricular surface to the pial surface, is significantly reduced in both homozygotes (64%) and heterozygotes (31%) compared to controls by E16. In homozygous *igf1r*^{Nestin-KO} embryos, gross malformations of the telencephalon were observed at E16, but not at E12 or E14 (Fig. 2). In the worst cases, the malformation appears to result from an almost complete elimination of the developing telencephalic wall (Fig. 3). This cortical loss in homozygotes was preceded by a transient spike of apoptosis at E14. In heterozygotes at E14, a transient

wave of apoptosis was also observed, although the density of apoptotic cells was reduced by 67%, when compared to homozygotes. Compared to controls, the volume of hippocampus in homozygous *igf1r*^{Nestin-KO} mice also was reduced by ~69%.

IGFBP-1 inhibits the interactions of IGF-I and IGF-II with their cell surface receptors when present in molar excess. In Tg mice with ectopic brain IGFBP-1 expression driven by a MT-I promoter (IGFBP-1^{MT-I} Tg mice), brain weight was significantly decreased by 8–16% at P14 (D'Ercole et al., 1994). This brain growth retardation is most obvious in the cerebral cortex (18%), hippocampus (20%) and diencephalons (12%), respectively (Ye et al., 1995a). Decreased density of myelinated axons is prominent in the cerebral cortex, anterior commissure, corpus callosum and diencephalon, although the cerebellum and brainstem are relatively spared (Ye et al., 1995a). Morphometric analysis of the primary somatosensory cortex in these Tg mice revealed a 24% decrease in the volume of somatosensory barrels in cortical layer IV. In another line of IGFBP-1 Tg mice (IGFBP-1^{PGK} Tg mice), in which the IGFBP-1 transgene expression is driven by a mouse phosphoglycerate kinase (PGK) promoter, the authors reported more significant decreases in body weight (12%) and brain weight (40%), likely due to a more abundant and earlier expression of the transgene. Brain DNA content (16%) and total protein (23%) also are reduced in adult mice (Ni et al., 1997), consistent with a reduction in cell number. On representative histological sections, cross-sectional areas of the hippocampus and DG were significantly reduced in IGFBP-1^{PGK} Tg mice by 55% and 72%, respectively, with a decrease of 33% for the brain as a whole. White matter and fiber tracts were less well developed, and the thickness of the corpus callosum was decreased by 62% (Ni et al., 1997). Interestingly, in a line of Tg mice with hepatic IGFBP-1 overexpression and increased circulating levels of IGFBP-1, there was a significant reduction in brain weight by 2 months of age (Doublier et al., 2000), in line with the view that circulating IGF can pass across the BBB and regulate the brain growth. In IGFBP-2 Tg mice, as well as Tg mice with an increased expression of IGFBP-3 or IGFBP-5, modest reductions of brain weight were also reported (see Table 2).

4.3. Neurogenesis and apoptosis

The capacity of IGF-I to promote neuron progenitor proliferation and differentiation has been well-documented *in vitro* (Arsenijevic and Weiss, 1998; Arsenijevic et al., 2001; Cicco-Bloom and Black, 1988; Drago et al., 1991; Torres-Aleman et al., 1990b; Werther et al., 1993; Zackenfels et al., 1995). Additional *in vitro* studies have shown that IGF-I promotes cell survival through its anti-apoptotic actions (Russell et al., 1998; Takadera et al., 1999; Torres-Aleman et al., 1990a; Torres-Aleman et al., 1990b; Yamada et al., 2001). Morphometric and stereological analyses of the developing brain in IGF-I overexpressing Tg mice have reported substantial increases in the total number of neurons in the cerebral cortex (Gutierrez-Ospina et al., 1996; Hodge et al., 2005), cerebellar cortex (Ye et al., 1996), DG of the hippocampus (O'Kusky et al., 2000; Ye et al., 2004), and selected brainstem nuclei (Dentremont et al., 1999). By contrast, in *igf-1*KO mutants and IGFBP-1 overexpressing Tg mice, significant decreases in neuron number have been reported in the cerebral cortex, hippocampus, DG striatum, and cochlear nucleus (Beck et al., 1995; Camarero et al., 2001; Gutierrez-Ospina et al., 1996; Ni et al., 1997). Both *in vitro* and *in vivo* studies show that des-N-(1–3) IGF-I can block neuronal apoptosis in response to hypoxia/ischemia or excitotoxicity (Guan et al., 1996; Saura et al., 1999).

Morphometric analysis of IGF-I^{MT-I} mice revealed a 24% increase in the total number of neurons in somatosensory barrels in cortical layer IV by P90 (Gutierrez-Ospina et al., 1996). There is a 39% decrease in the numerical density of neurons (N_V , number per unit volume), indicating an increase in the volume of neuropil separating individual neuronal cell bodies, and a 33% increase in mean neuronal profile area. Neurons in layer IV of the cerebral cortex in mouse are generated during prenatal development (Hicks and D'Amato, 1968), while

apoptotic neuron death occurs predominantly from birth to P10 (Spreafico et al., 1995; Verney et al., 2000). Given that the transgene in IGF-I^{MT-I} mice is significantly expressed after birth (Ye et al., 1995a), it would appear that increased neuron number in these adult Tg mice results from decreased apoptosis during the regressive phase of neurogenesis. A similar morphometric analysis of the somatosensory cortex in IGFBP-1^{MT-I} Tg mice revealed a 15% decrease in the total number of neurons in somatosensory barrels and a 39% increase in the N_V of neurons (Gutierrez-Ospina et al., 1996). Decreased cortical volume in these Tg mice results from a decrease in both the number of neurons and the volume of neuropil separating individual cell bodies. Since the transgene expression in these mice also is driven by the same promoter and is observed first after birth (Ye et al., 1995a), decreased neuron number likely results from increased apoptosis during the regressive phase of neurogenesis.

In the cerebellar cortex of IGF-I^{IGF-II} Tg mice at P50, the total number of Purkinje cells and granule cells is increased by 20% and 82%, respectively (Ye et al., 1996). When bromo-2-deoxyuridine (BrdU) was employed to label proliferation of granule cell progenitors in the external granule cell layer, the total number of labeled cells was increased by 38% in IGF-I^{IGF-II} Tg mice at P15. Purkinje cells of the cerebellar cortex originate from mitotic neuroepithelial cells from E11 to E13, while granule cells originate from E17 to P15 (Mares and Lodin, 1970; Miale and Sidman, 1961). Given this protracted period of proliferation for granule cells and the increase in BrdU-labeled progenitors, it appears most likely that elevated levels of IGF-I during early postnatal development act to increase the rate of mitosis in the external granular layer. In a subsequent study, the anti-apoptotic effects of elevated IGF-I were investigated in the cerebellum of IGF-I^{IGF-II} Tg mice (Chrysis et al., 2001). Morphometric analysis of apoptotic cells in the cerebellum, detected by terminal deoxynucleotidyl transferase-mediated UTP nick end labeling (TUNEL), revealed a 47% decrease in Tg mice when compared to controls. The abundance of procaspase-3 and caspase-3 is also decreased in IGF-I^{IGF-II} Tg mice, accompanied by increased expression of the anti-apoptotic Bcl genes, *bcl-x_L* and *bcl-2*. Consistent with these studies, BCL-2 was found to be increased in immunohistochemical studies of cerebellum in these Tg mice (Baker et al., 1999b). These results provide direct evidence that elevated IGF-I acts to inhibit apoptosis during early postnatal development in a developmentally specific manner.

Increased growth of the hippocampal DG has been studied in IGF-I^{IGF-II} Tg mice throughout postnatal development (O'Kusky et al., 2000). In control mice the total number of neurons in the granule cell layer increased by 113% from P7 to P35, with no additional increase in neuron number by P130. In IGF-I^{IGF-II} Tg mice there was a 172% increase in neuron number from P7 to P35, with an additional and significant increase of 17% between P35 and P130, suggesting a protracted period of accelerated neurogenesis. The total number of neurons in the granule cell layer was significantly greater in IGF-I^{IGF-II} Tg mice by 56% at P35 and by 61% at P130. In IGFBP-1^{GKP} Tg mice with inhibited IGF-I actions, BrdU labeling of proliferating cells in the granule cell layer of the DG revealed a 41% decrease in the number of labeled cells in Tg mice at P3, as compared to controls (Ni et al., 1997). In the VZ and SVZ of the lateral ventricle, BrdU-labeled cells were decreased by 19%. The number of TUNEL-labeled apoptotic cells throughout the hippocampus was found to be increased by 55% in IGFBP-1^{GKP} Tg mice (Ni et al., 1997). These studies indicate that IGF-I acts to increase neuron proliferation while inhibiting apoptosis in a region of the brain where progressive and regressive phases of neurogenesis are known to occur throughout the life of the organism.

In the brainstem of IGF-I^{IGF-II} Tg mice at P35, the total number of neurons is significantly increased in the nucleus of the solitary tract (50%) and in the dorsal motor nucleus of the vagus (53%), but not in the hypoglossal nucleus or the facial nucleus (Dentremont et al., 1999). Neuron proliferation in mice occurs from E9 to E12 for the nucleus of the solitary

tract and from E9 to E10 for the dorsal motor nucleus of the vagus, the hypoglossal nucleus and the facial nucleus (Pierce, 1973). Given that transgene expression in IGF-I^{IGF-II} Tg mice is very low prior to birth, increased rates of neuronal proliferation are unlikely to account for increased neuron numbers. An inhibition of naturally occurring neuron death (apoptosis) appears to be more likely. Apoptotic death of motor neurons in rodent hypoglossal nucleus occurs exclusively from E16 to E21 (Friedland et al., 1995), as it does for motor neurons in the spinal cord (Oppenheim, 1986). Motor neurons in the facial nucleus are likely to undergo similarly early programmed cell death. Thus, the lack of effect of elevated IGF-I on neuron number in the hypoglossal and facial nuclei in these mice appears to stem from the age at which the transgene is expressed. Interestingly, although the total number of neurons did not differ significantly between Tg and control mice in the hypoglossal and facial nuclei, the N_V of neurons was significantly decreased in both regions while the mean neuronal profile area is significantly increased. Changes in these variables indicate an increased volume of neuropil and possibly more complex arborizations of the dendritic trees on individual neurons within these regions.

IGF also significantly increases neural precursor proliferation *in vivo*. In IGF-I^{Nestin} Tg embryos a 54% increase in the total number of cells in the dorsolateral wall of the telencephalon was observed within the cortical plate at E16 (Popken et al., 2004). Cumulative S phase labeling with BrdU revealed a significant decrease in total cell cycle length (T_C) in Tg embryos at E14 (Hodge et al., 2004). This decrease in T_C was found to result entirely from a reduction in the length of the G₁ phase of the cell cycle from 10.66 h to 8.81 h, with no significant changes in the lengths of the S, G₂ and M phases. Additionally, the proportion of daughter cells reentering the cell cycle was significantly increased by 15% in IGF-I^{Nestin} Tg embryos, compared with littermate controls (Hodge et al., 2004). These results demonstrate that IGF-I accelerates progenitor cell division in the VZ by reducing G₁ phase length and decreasing T_C but increases cell cycle re-entry.

In IGF-I^{Nestin} Tg mice at P12, significant increases in the total number of neurons were determined in the cerebral cortex (27%), caudate-putamen (27%), DG of the hippocampus (69%), the medial habenular nucleus (61%), and the lateral habenular nucleus (36%), when compared to littermate controls (Hodge et al., 2005; Popken et al., 2004). In the cerebral cortex of IGF-I^{Nestin} Tg and control mice, apoptotic cells were observed in all cortical layers from P0 to P7 by virtue of their immunoreactivity to antibodies against cleaved caspase-3 (Hodge et al., 2007; Popken et al., 2004). The vast majority of these cells (>80%) appeared to be neurons because of their atrophic dendritic arborizations. *In vivo* studies using these IGF-I^{Nestin} Tg mice clearly demonstrate that elevated levels of IGF-I in the developing brain, beginning as early as E13, can simultaneously accelerate mitosis in neural progenitors and inhibit apoptosis in post-mitotic neurons.

In *igf-I* KO mutant mice, immunohistochemical studies (Beck et al., 1995) have reported significant decreases in the number of parvalbumin-immunoreactive neurons in the striatum (52%), hippocampus (32%) and DG (59%). Interestingly, the numbers of cholinergic neurons in both the striatum and basal forebrain and dopaminergic neurons in the mesencephalon did not change, suggesting a differential susceptibility of neurotransmitter-specific neuron populations to the effects of IGF-I (Beck et al., 1995).

Preliminary results from ongoing immunohistochemical studies of adult heterozygous *igf1r*^{Nestin-KO} mice in our laboratories indicate a disproportionate loss of GABAergic neurons in the ventral prefrontal cortex, when compared to controls (Fig. 4). In the cerebral cortex of *igf1r*^{Nestin-KO} mice, the N_V of all neurons did not differ significantly from controls, although the total number of neurons was reduced by 26% due to decreases in cortical volume. In the ventral prefrontal cortex, the N_V of GABAergic neurons in heterozygous

igf1r^{Nestin-KO} mice ($15,042 \pm 536$, mean \pm standard error) was significantly reduced (25%, $P < 0.001$), when compared to controls ($20,093 \pm 857$). In the dorsal prefrontal cortex, the N_V of GABAergic neurons did not differ significantly between heterozygous *igf1r*^{Nestin-KO} mice ($16,294 \pm 452$) and controls ($15,772 \pm 697$). Interestingly, this disproportionate loss of GABAergic neurons was due entirely to a loss of calbindin-immunoreactive GABAergic neurons (N_V in heterozygous *igf1r*^{Nestin-KO} mice: $13,484 \pm 745$; compared to N_V in controls: $17,508 \pm 432$, $P < 0.001$), with no significant change in the proportion of calretinin-immunoreactive GABAergic neurons (N_V in heterozygous *igf1r*^{Nestin-KO} mice and controls was 7309 ± 389 and 7765 ± 520 , respectively), as illustrated in Fig. 4.

Delayed maturation of the inner ear and neuron loss have been reported in homozygous *igf-I* KO mice during early postnatal development (Camarero et al., 2001). The volumes of the cochlea and cochlear ganglion were reduced by 34% at P20, accompanied by a 19% loss of cochlear neurons and a 31% decrease in the volume of the surviving neuron body. The number of apoptotic neurons, demonstrated by TUNEL labeling and caspase-3, also was increased in *igf-I* KO mutant mice (Camarero et al., 2001).

Clearly, IGF-I is a potent factor in the control of neurogenesis and cell survival through its pro-mitotic and anti-apoptotic actions. IGF-I continues to be a modulator of neurogenesis in the adult hippocampus (Aberg et al., 2000; Trejo et al., 2008). IGF-I signaling is also a potent mediator of axonal and dendritic circuit remodeling during postnatal development.

4.4. Neurite outgrowth and synaptogenesis

In vitro studies of fetal hypothalamic cells in culture (Torres-Aleman et al., 1990b) have reported that in the presence of IGF-I neurons have a more differentiated morphology and express significantly higher levels of protein kinase C, an enzyme that increases during neurite outgrowth and synaptogenesis. These data suggest that IGF-I can regulate neurite and synaptogenesis.

In vivo studies of mice support the findings derived from culture studies. Injection of an IGF-I antisense nucleotide in the inferior olive of adult rats results in a significant decrease of IGF-I levels in the contralateral cerebellum and a marked reduction in the size of dendritic spines and in the numerical density of dendritic spines on Purkinje cells (Nieto-Bona et al., 1997). Morphometric analyses of the developing fronto-parietal cortex of *igf-I* KO mice (Cheng et al., 2003) and the hypothalamus of mice with a null mutation specifically in gonadotropin-releasing hormone (GnRH) neurons (*igf1r*^{GnRH-KO} mice) (DiVall et al., 2010) demonstrate that blunting IGF-I–IGF1R signaling decreases dendritic length, branching complexity, and dendritic spine formation. Interestingly, both male and female *igf1r*^{GnRH-KO} mice exhibit delayed onset of puberty, although reduced IGF1R signaling during development produces no change in the distribution and total number of GnRH neurons (DiVall et al., 2010). Mice with a mutation in the *InR* gene specifically in GnRH neurons, however, do not exhibit delayed puberty and GnRH neuronal pathology (DiVall et al., 2010). Therefore, these data suggest that IGF1R signaling, but not InR signaling, in GnRH neurons is necessary for normal pubertal onset, likely involving dendritic tree growth and the prepubertal maturation of afferent input to these neurons.

In *igf-I* KO mice, there is also a 16% reduction in the density of dendritic spines, correlating with a 30% reduction in synaptotagmin levels. Consistently, P20 *igf-I* KO mice exhibit an abnormal distribution of synaptophysin in the organ of Corti (Camarero et al., 2001), as the pattern of its immunoreactivity in the cell bodies of cochlear ganglion neurons and sensory hair cells more closely resemble control mice at P5. These data indicate the persistence of an immature pattern of synapse distribution in the absence of IGF-I, and suggest that IGF-I deficiency is associated with a reduction of synaptogenesis in the developing brain.

More direct evidence that IGF-I promotes synaptogenesis comes from studies of IGF-I Tg mice and IGFBP-1^{MT-I} Tg mice (Gutierrez-Ospina et al., 2004; O'Kusky et al., 2003). The density and total number of synapses in individual barrels of the somatosensory cortex were determined using histochemical methods for the oxidative enzyme succinic dehydrogenase. The density of synapses did not differ significantly among Tg and control mice. However, the total number of synapses was found to be 38% greater in IGF-I^{MT-I} Tg mice and 43% less in IGFBP-1^{MT-I} Tg mice (Gutierrez-Ospina et al., 2004). These changes resulted from the increased or decreased volumes of individual barrels in IGF-I^{MT-I} Tg and IGFBP-1^{MT-I} Tg mice, respectively, rather than from changes in the packing density of individual synaptic contacts (Gutierrez-Ospina et al., 2004).

The effects of elevated IGF-I on both the progressive and regressive phases of synaptogenesis in the hypoglossal nucleus were also investigated in IGF-I^{IGF-II} mice (O'Kusky et al., 2000; O'Kusky et al., 2003). In control mice the total number of synapses in the hypoglossal nucleus was increased by 354% from P7 to peak values at P21, followed by a significant decrease of 33% by P130 (O'Kusky et al., 2003). IGF-I^{IGF-II} Tg mice exhibited a similar trend of changes. When compared to control mice, however, IGF-I^{IGF-II} Tg mice had 42% and 52% more synapses at P21 and P130, respectively (O'Kusky et al., 2003). Given that the total number of hypoglossal neurons does not differ significantly between Tg and control mice at the ages examined, the synapse-to-neuron ratio is significantly greater in IGF Tg mice after P14. Thus, the increased *in vivo* expression of IGF-I during postnatal development augments the progressive phase of synaptogenesis, although it does not prevent synapse elimination during the regressive phase.

Similarly, in the hippocampal DG of IGF-I^{IGF-II} Tg mice, significant increases in the total number of synapses in the molecular layer were observed at P14 (by 61%), P21 (42%), P28 (105%), P35 (96%), and P130 (54%) (O'Kusky et al., 2000). Interestingly, the density of synapses is significantly greater only at P28 (35%) and P35 (21%) in IGF-I^{IGF-II} Tg mice, and accordingly, a greater synapse-to-neuron ratio, which returns to normal values by P130. The increase in synapse number in the DG tends to reflect the increased number of neurons in the granule cell layer in IGF-I^{IGF-II} Tg mice. It has been shown that reduced neurogenesis and synaptogenesis in the hippocampus of animal models of gestational–neonatal iron deficiency is secondary to reduced levels of IGF-I during development (Tran et al., 2012), suggesting that IGFI can mediate the effects of iron on synaptogenesis.

Taken together, these data strongly indicate that IGF signaling plays an important role in neurite outgrowth and synaptogenesis throughout postnatal development and in adult. In addition, IGF signaling may act by regulating the expression of synaptic proteins, such as synaptophysin, because synaptophysin, by binding synaptobrevin, is believed to play an important role in regulating SNARE assembly, vesicle fusion (Edelmann et al., 1995; Valtorta et al., 2004), and synapse formation (Tarsa and Goda, 2002).

4.5. Oligodendrocytes and myelination

IGF actions on oligodendrocyte lineage cells have been well-documented. In cultures both IGF-I and IGF-II are capable of promoting the proliferation, survival and development of rodent and human oligodendrocyte lineage cells (Armstrong et al., 1992; Barres et al., 1992; Cui et al., 2012; Masters et al., 1991a; McMorris et al., 1986; Mozell and McMorris, 1991; Ye and D'Ercole, 1999). Many of these results have been confirmed in studies of mutant mice.

Overexpression of IGF-I in the brain of Tg mice results in dramatic increases in brain growth, myelin content, the number of oligodendrocytes and their precursors (Carson et al., 1993; Ye et al., 1995a; Ye et al., 1995b; Ye et al., 2007; Ye et al., 2004). Compared to

control mice, adult IGF-I^{MT-I} Tg mice have 130% more myelin content (Carson et al., 1993) and 2–3-fold more abundant mRNAs for myelin-basic protein (MBP) and proteolipid protein (PLP) (Ye et al., 1995a), the two most abundant myelin-specific proteins. The increased myelin is the result of increases in the thickness of myelin sheath and in the number of myelinated axons, which does not proportionally depend on axon size (Ye et al., 1995a; Ye et al., 1995b). Taken together with the fact that the number of PLP+ oligodendrocytes is only modestly increased (by 18% in corpus callosum and 36% in cerebral cortex) (Ye et al., 1995a), these data suggest that IGF-I promotes the production of myelin by each oligodendrocyte. In contrast, blunting IGF-I expression in *igf-I* KO mice results in a dramatic reduction in oligodendrocytes and their precursors, and much less abundant MBP and PLP during early postnatal development (Beck et al., 1995; Ye et al., 2002b). The abundance of the myelin-specific proteins, however, gradually increases with increasing age, and becomes normal in adult (Ye et al., 2002b). Because the abundance of IGF-II is significantly increased during the same developmental period (Ye et al., 2002b), it is likely that the recovery of reduced myelin-specific protein expression induced by IGF-I deficiency is, at least in part, due to the compensatory effects of IGF-II. This speculation is supported by the findings that myelination and the expression of MBP and PLP remain suppressed in the adult mice with an IGF1R null mutation specifically in the cells of oligodendrocyte lineage (see below), or in the adult mice that ectopically express IGFBP-1 (Ni et al., 1997; Ye et al., 1995a), an IGF binding protein that inhibits IGF-I and IGF-II actions.

Direct interactions of IGF with the IGF1R on the cell surface of oligodendrocyte lineage cells are essential to the normal oligodendrocyte development and myelination, although indirect IGF effects via neurons and other neural cell types also are likely to be involved. When IGF1R expression is specifically blunted in the Olig1+ oligodendrocyte precursors and their progeny, the volume of corpus callosum and anterior commissure, two regions that are rich in oligodendrocyte lineage cells, and the number of oligodendrocytes and their precursors are markedly decreased (Zeger et al., 2007). The reduction in these cells apparently is a result of decreased proliferation and increased apoptosis (Zeger et al., 2007).

While IGF signaling has been convincingly shown to promote the development of oligodendrocyte precursors, the details about its role in each of the developmental stages remain to be delineated. In cultured oligodendrocyte precursors from adult corpus callosum, IGF-I, in the presence of insulin at a concentration that also stimulates IGF1R, promotes the development of O4+ oligodendrocyte precursors to O1+ early oligodendrocytes, but has no effects on the progression of A2B5+ early oligodendrocyte precursors to O4+ precursors (Mason and Goldman, 2002). These data suggest that IGF signaling preferentially promotes the differentiation of adult oligodendrocyte precursors at one or more stages of their development. The actions of IGF signaling on oligodendrocyte specification also have been reported, but conclusion remains discordant. In cultured adult rat hippocampal NSC IGF-I appears to preferentially direct NSC towards oligodendrocyte lineage (Hsieh et al., 2004). In contrast, a more recent study of cultured NSC from neonatal mouse forebrain shows that overexpression of an IGF-I transgene in the cells markedly increases neuronal number without an appreciable change in the number of oligodendrocytes (Kouroupi et al., 2010). The exact reason(s) for the discrepancy between these two studies is not known. Possible explanations likely include different NSC sources and culture conditions employed. It is worth noting that in both studies GFAP was used as a marker for astrocytes. Because a subpopulation of NSC express GFAP, thus, the reduction in GFAP reported in the both studies may also reflect a reduction in the NSC population due to their development to mature neural cells.

Currently, there is no available data from mutant mice that support an essential role for IGF in oligodendrocyte specification *in vivo*. In *igf1r^{Olig1-KO}* mice that carry an *igf1r* null mutation in the oligodendrocyte precursors positive for Olig1, a transcription factor that is specifically expressed in early oligodendrocyte precursors and motor neurons (Lu et al., 2002; Woodruff et al., 2001), both mature oligodendrocytes and their precursors continue to exist, although markedly reduced in number (Zeger et al., 2007). In addition, the total number of astrocytes and neurons is not inversely increased, rather modestly decreased. In contrast, in the developing brains of each of the 4 IGF-I overexpressing Tg mouse lines that we have studied (see Table 2), including IGF-I^{Nestin} Tg mice which overexpress an IGF-I transgene from E13, the number of neurons, astrocytes and oligodendrocytes are all increased, although the magnitude of the increase in each neural cell type varies depending on specific lines (Dentremont et al., 1999; Gutierrez-Ospina et al., 1996; Popken et al., 2004; Ye et al., 1995a; Ye et al., 2000; Ye et al., 2004; Ye et al., 1996). In addition, mice that conditionally overexpress IGF-I in GFAP+ cells exhibit a proportional increase in the number of astrocytes, oligodendrocytes and neurons in hippocampus (Ye et al., 2004).

4.6. Astrocytes and microglia

There are several lines of evidence indicating that IGF also promotes the development and growth of astrocytes. In astrocyte cultures, IGF-I significantly increases their proliferation (Aberg et al., 2003b; Ballotti et al., 1987; Chernauek, 1993; Han et al., 1987; Han et al., 1992). Consistently, injection of IGF-I into developing sheep brains following ischemic injury (Cao et al., 2003) or overexpressing IGF-I in the brain of IGF-I^{GFAP} Tg mice (Ye et al., 2004) increases the number of astrocytes. Intriguingly, unlike IGF-I^{MT-I} Tg mice that show a predominate expression of IGF-I transgene in neurons, adult IGF-I^{GFAP} Tg mice have 50–270% more GFAP protein (depending on the brain regions) (Ye et al., 2004), suggesting that astrocyte-derived IGF-I can differentially influence astrocyte functions. In line with this speculation, IGF-I treatment of cultured astrocytes also increases the expression of the gap junction protein connexin43 (Aberg et al., 2003b), alpha 1 isoform of Na-ATPase (Matsuda et al., 1993), glucose transporter (Masters et al., 1991b), and glial glutamate transporter GLAST (Suzuki et al., 2001), as well as their activity (Masters et al., 1991b; Matsuda et al., 1993). These data also suggest that IGF-I is capable of promoting the cross-talk between astrocytes and other neural cells. Taken together, these data suggest that IGF signaling is likely to promote the development of both oligodendrocyte, neuron and astrocyte lineages in the developing brain, and that the site of IGF-I expression influences the nature and magnitude of its actions.

Following various brain injuries in adult, IGF expression is significantly increased in astrocytes (Garcia-Estrada et al., 1992; Gudi et al., 2011; Komoly et al., 1992; Li et al., 1998; Liu et al., 1994), parallel to an increased astroglial activation and proliferation. Decrease in IGF-I availability by ectopic expression of IGFBP-1 in the brain of IGFBP-1^{PGK} Tg mice significantly reduced astrocyte responses to injury (Ni et al., 1997). These data indicate that IGF signaling is critically important to astrocyte proliferation and functions following injury.

Microglia are derived from hemangioblastic mesoderm (Streit, 2001), and populate the developing neuroectoderm as early as E8 in rodents. As with the case in astrocytes, there are few reports about IGF actions in microglia generation and development during prenatal and postnatal development. Like astrocytes, IGF production is markedly increased in activated microglial cells following various brain injuries (Beilharz et al., 1998; Breese et al., 1996; Choi et al., 2008; Gudi et al., 2011; O'Donnell et al., 2002). In cultures, IGF-I increases microglia proliferation (O'Donnell et al., 2002), and microglia-derived IGF-II protects oligodendrocytes from tumor necrosis factor- α induced cell death (Nicholas et al., 2002).

These data suggest that microglial IGF-I acts on neural cells in a paracrine and/or autocrine pattern.

5. IGF-II actions

In cultures, IGF-II exerts actions on the development and growth of neural cells in a manner that is similar to that of IGF-I. Genetic studies also clearly demonstrate that IGF-II has an important role in growth during early development. Global ablation of IGF-II gene expression in *igf-II* KO mice results in a marked retardation in body growth (Baker et al., 1993; DeChiara et al., 1990; Liu et al., 1993) and in brain growth, leading to a 24% reduction in brain weight at postnatal day 8 (Lehtinen et al., 2011). The decreased brain growth, at least in part, is a result of reduced cell proliferation of NSC/NPC during early development (Lehtinen et al., 2011). In contrast, adding exogenous IGF-II to culture medium significantly increases the number of neurospheres derived from NSC/NPC and the number of proliferating cells in the VZ of embryonic explants (Lehtinen et al., 2011). Similarly, during late embryonic development in mice, IGF-II, derived from choroid plexus and signaling through the IGF1R, appears to be a dominant growth factor that stimulates the proliferation of VZ NSC (Lehtinen et al., 2011). Nonetheless, postnatal overexpression of IGF-II in mutant mice does not alter brain growth (Van Buul-Offers et al., 1995). It is possible that during postnatal development IGF1R signaling derived from abundantly-expressed IGF-I, which has 10–100 fold higher affinity for IGF1R compared to IGF-II, has already operated at its maximal level. Thus, increasing IGF-II postnatally may not be able to further increase signaling through the IGF1R and to elicit additional growth actions.

However, during late brain development or in the adult brain, IGF-II is likely to have an important role in modulation of non-growth neuronal functions. For example, in adult rodents injection of IGF-II into hippocampus improves memory (Chen et al., 2011). The IGF-II action appears to be specifically mediated by IGF2R, because co-injection of an IGF2R inhibitor, but not IGF1R inhibitors, abolishes the memory improvement induced by IGF-II (Chen et al., 2011). In addition, Leu²⁷-IGF-II, an IGF-II analog that preferentially binds to the IGF2R, attenuates depolarization-evoked GABA release in cultured adult hippocampal and cortex (Amritraj et al., 2010), further supporting that IGF-II, via the IGF2R, can affect neuronal functions in adult.

6. IGF signaling

6.1. Signaling through IGF1R

Most IGF growth actions, if not all, are mediated by IGF1R. Direct interaction of IGF with the IGF1R in CNS neural cells is essential for the normal neural development and the proper brain cytoarchitecture, as summarized in the sections above. While our understanding of IGF intracellular signaling has been significantly advanced in recent years, the intracellular signaling pathways that mediate each of IGF actions in the CNS remain to be precisely elucidated. Fig. 5 depicts largely simplified IGF signaling pathways in the CNS.

It is known that binding IGF-I or IGF-II to the IGF1R leads to activation of the tyrosine kinase in the β -subunits, which in turn autophosphorylates its tyrosine residues and recruits docking proteins (LeRoith et al., 1995). Docking proteins that are involved in IGF signaling include five isoforms of IRS [IRS-1 to IRS-4 and Grb2-associated binder-1 (Gab-1)], Ras, and Src homology domain-containing proteins. Phosphorylated docking proteins then subsequently recruit downstream signaling molecules and transduce IGF signaling. The Ras–Raf–MAP and PI3–Akt kinase pathways are two extensively studied signaling pathways that play key roles in IGF neural actions. There is also evidence that G protein mediates some of IGF neural actions (Keummerle and Murthy, 2001). More recently, it has

been reported that in cultured non-neural cells IGF1R can be modified by small ubiquitin-like modifier protein-1 (SUMO), and transported into the nucleus, where the SUMO-modified IGF1R acts as a transcription factor to directly regulate its target gene expression (Sehat et al., 2010). It is highly possible that the IGF1R can also act as a transcription factor in neural cells, because neural cells are capable of internalizing the IGF1R (Romanelli et al., 2007). It is worth noting that most, if not all, IGF signaling molecules and pathways described below also are involved in many other signaling pathways. For example, IGF and insulin share many signaling molecules and pathways, although how they exert distinct functions remains to be understood.

6.1.1. IRS—IRSs, molecules that are also involved in insulin signaling, are widely expressed in the CNS. IRS-1, -2, -4 and Gab-1 are expressed in a tempo-spatial specific pattern (Fantin et al., 1999; Folli et al., 1994; Holgado-Madruga et al., 1996; Numan and Russell, 1999; Sciacchitano and Taylor, 1997; Ye et al., 2002a), and little IRS-3 is detected in adult brain (Sciacchitano and Taylor, 1997). It appears that while it mediates some IGF-I neural actions, IRS-1 is not essential in IGF neural signaling. Brain growth (Pete et al., 1999; Schubert et al., 2003; Ye et al., 2002a) and myelination (Engleka et al., 1996) are modestly reduced in *irs-1* KO mice. Unlike skeletal muscle in which IGF-I dependent growth is impaired by IRS-1 ablation (Pete et al., 1999), blunting IRS-1 expression does not block IGF-I-stimulated brain growth and myelin-specific protein expression (Ye et al., 2002a). The abundance of IRS-2 and IRS-4 is significantly increased in the brain of *irs-1* KO mice, suggesting that increased IRS-2 and/or IGF-4 may compensate for the loss of IRS-1.

On the other hand, blunting IRS-2 expression globally (Schubert et al., 2003) or specifically in nestin+ neural cells (Taguchi et al., 2007) results in a much more significant reduction in brain weight (by 30–38% in adult mice), compared to that in *irs-1* KO mice. These data suggest that IRS-2 appears to be more important in mediating IGF/insulin signaling in the CNS. It is worthwhile to note that in *irs-2* KO mice blunting one *igf1r* allele further reduces brain weight by 15%, while brain growth retardation is similar in *igf1r* KO and *irs-2/igf1r* double KO embryos (Schubert et al., 2003). These data strongly indicate that IGF–IGF1R signaling can act through molecule(s) other than IRS-2. At E14–E16, *irs-2* KO mice have 37% less proliferating neuronal cells without apparent changes in cell apoptosis (Schubert et al., 2003). This suggests that at this developmental stage IRS-2 primarily transduces proliferation signaling, but is not essential to pro-survival signaling. IRS-2, however, is capable of mediating pro-survival signaling, at least during postnatal development. For example, the number of apoptotic photoreceptor cells is increased by 50% in the retina of 2 week old *irs-2* KO mice (Yi et al., 2005).

IRS-2 also appears to play an important role in the growth of oligodendrocyte lineage cells. Blunting IRS-2 expression transiently decreases the abundance of multiple myelin/oligodendrocyte-specific proteins during the first 2 weeks of postnatal life (Freude et al., 2008). The expression of these proteins then gradually recovers with increasing age, likely due to the compensatory actions of increased IRS-1 and/or Akt (Freude et al., 2008). The details of IRS-2 actions in the generation and differentiation of oligodendrocyte lineage cells, as well as other neural cells remain to be further defined.

Taken together, IRS-2 appears to mediate a significant portion of growth signaling from IGF, as well as insulin, in a cell- and development- dependent manner, and one or more IRS molecules are capable of compensating for the loss of others.

6.1.2. Phosphoinositide-3 (PI3) kinase–Akt kinase pathway—IGF treatment rapidly increases Akt phosphorylation and activation in neural cultures. Unlike the transient

activation of Raf–Erk (see below), however, IGF-I-stimulated PI3–Akt activation is sustained for at least 24 h, at least in cultured oligodendrocytes (Ness and Wood, 2002; Romanelli et al., 2007; Ye et al., 2010). In line with these *in vitro* studies, IGF-I overexpression markedly increases the abundance of phosphorylated Akt in developing brains, while reducing IGF availability decreases its abundance during the same developmental period (Sun and D'Ercole, 2006).

It has been well-documented that the PI3–Akt kinase pathway plays a key role in the survival of neural cells, including NSC/NPC, neurons (Johnson-Farley et al., 2007; Johnson-Farley et al., 2006) and oligodendrocytes (Dudek et al., 1997; Ebner et al., 2000; Sinor and Lillien, 2004; Vemuri and McMorris, 1996; Yao and Cooper, 1995). Multiple downstream signaling mechanisms are involved in these survival-promoting actions, including: (1) inhibition of the activity of the pro-apoptotic protein Bad (Kulik and Weber, 1998) and the forkhead transcription factor FKHRL1 (Brunet et al., 1999), which regulates the transcription of proapoptotic protein genes; (2) suppression of Bax translocation (Ness et al., 2004); (3) enhancement of the expression of the pro-survival proteins Bcl2 (Baker et al., 1999b; Chrysis et al., 2001; Minshall et al., 1999) and inhibitor of apoptosis proteins (IAPs) Liu et al., 2011; (4) reduction of the expression of the apoptotic enzymes caspase-3 and caspase-9 (Cui et al., 2005; Ness et al., 2004); and (5) increase in the activity of mammalian target of rapamycin (mTOR) Sinor and Lillien, 2004.

Accumulating data also show that the PI3–Akt pathway is required for neural proliferation and IGF-stimulated proliferation. In culture cells inhibition of PI3 kinase significantly reduces the incorporation of ³H-thymidine into oligodendrocyte precursors (Ebner et al., 2000), and markedly suppresses the IGF-I-stimulated proliferation in cortical NSC/NPC (Mairet-Coello et al., 2009) and cerebellar granule cell precursors (Cui et al., 1998). Conversely, overexpression of Akt in cortical precursor cells increases PCNA and BrdU labeling (Sinor and Lillien, 2004), markers of cell proliferation.

Glycogen synthase kinase 3 β (GSK3 β) and β -catenin appear to be two critical downstream effectors in the PI3–Akt pathway that mediates IGF proliferation actions. Phosphorylation and inactivation of GSK3 β by PI3–Akt kinases leads to stabilization of cyclin D1 protein (Ryves and Harwood, 2003; Takahashi-Yanaga and Sasaguri, 2008), a molecule key to the cell cycle progression through the G1/S phases. IGF can also signal through β -catenin, a key member of Wnt signaling pathway. In studies of cultured oligodendrocytes we (Ye et al., 2010) have show that IGF-I promotes Akt kinase and GSK3 β phosphorylation. As a result, the abundance of β -catenin and the mRNA expression of its target gene cyclin D1 are increased. Inhibiting PI3–Akt kinase activity significantly suppresses the IGF-I-stimulated expression of β -catenin mRNA and its protein. Importantly, knocking down β -catenin expression markedly inhibits the IGF-I actions on cyclin D1 mRNA expression, cell proliferation and survival. These data suggest that β -catenin is a common effector mediating a portion of IGF-I and Wnt signaling to promote neural cell proliferation and survival.

Mammalian target of rapamycin (mTOR) is yet another important downstream effector of the IGF–PI3–Akt pathway. In recent years, details about the role of mTOR in IGF pleiotropic neural actions begin to emerge. It appears that mTOR is mainly involved in signals promoting cell maturation and function, because blocking mTOR signaling suppresses the development of cultured immature oligodendrocytes (Guardiola-Diaz et al., 2012) and myelination in mice (Narayanan et al., 2009). mTOR activation, as well as that of PI3 and Akt kinases, is also required for the IGF-I stimulated protein synthesis in cultured oligodendrocyte precursors (Bibollet-Bahena and Almazan, 2009). Similarly, insulin at a concentration that also activates the IGF1R promotes the differentiation of cultured neuronal precursors via mTOR (Han et al., 2008). PI3K–Akt–mTOR signaling appears also to be

involved in IGF-I enhanced GLUT3 expression in P12 neuronal cells (Yu et al., 2012), long-term potentiation in the medial prefrontal cortex (Sui et al., 2008), and both cell growth and dendritic elaboration in hippocampus neurons (Jaworski et al., 2005). These data suggest that IGF-mTOR signaling likely promotes cellular maturation and functions by increasing protein synthesis. Intriguingly, it is reported that in cultured oligodendrocyte precursors mTOR is directly downstream of PI3 kinase rather than Akt kinase (Bibollet-Bahena and Almazan, 2009). Whether this signaling pathway also occurs *in vivo* or in other neural cell types, however, is not known.

6.1.3. Ras–Raf MAP pathway—IGF signaling through mitogen-activated protein (MAP) kinases is another pathway that has been abundantly studied. MAP kinases, a family of serine/threonine protein kinases, are conserved and widely distributed throughout the body. MAP kinase signaling cascades are organized in three-tiers. MAP signaling is initiated by the activation of MAP kinase kinase kinases, which in turn activates MAP kinase kinases and subsequently MAP kinases. MAP kinases include Erk1/2, p38, SAPK/JNK1/2/3, and Erk5/BMK1. While there are few reports on the role(s) of SAPK/JNK1/2/3 or Erk5/BMK1 in IGF neural actions, evidence has indicated that IGF can act in part by signaling through the Erk1/2 and p38 MAP pathways.

In cultured oligodendroglial cells and neurons, IGF-I transiently induces a rapid phosphorylation and activation of Erk1/2 and p38 MAP kinases within minutes (Cui and Almazan, 2007; Feldman et al., 1997; Hallak et al., 2001; Ye et al., 2010). Intriguingly, IGF-I treatment also leads to a delayed repression of Erk1/2 activity in cerebellar granule cells (Subramaniam et al., 2005). The biological significance of this late Erk suppression induced by IGF-I is not fully understood, although Erk suppression or ablation in adult neuronal cells has been associated with an enhanced cell survival (Subramaniam et al., 2005; Subramaniam et al., 2004) and synaptic plasticity, learning and memory abilities in mice (Mazzucchelli et al., 2002).

Studies of cultured neural cells indicate that Erk and p38 MAP signaling play an important role in IGF actions on cell maturation and differentiation. In oligodendroglial cell cultures, suppressing Erk activity blocks IGF-I stimulated protein synthesis (Bibollet-Bahena and Almazan, 2009), and halts the progression of early oligodendrocyte precursors to late oligodendrocyte precursors (Guardiola-Diaz et al., 2012). Similarly, suppression of p38 MAP pathway arrests the development of oligodendrocytes and myelination (Fragoso et al., 2007; Haines et al., 2008). The role of the Raf–Erk pathway in IGF promotive actions on oligodendrocyte maturation has been confirmed in mutant mice. Blunting the expression of brain B-Raf (the dominant activating isoform of the 3 Rafs in brain) decreases the abundance of phosphorylated Erk1 and Erk2 by more than 60% (Galabova-Kovacs et al., 2008; Zhong et al., 2007), and markedly reduces the number of mature oligodendrocytes and myelination in mutant mice (Galabova-Kovacs et al., 2008).

Based on culture studies using pharmacological inhibitors, activation of the Raf–Erk pathway has been believed to be key to IGF-stimulated cell proliferation (D'Ercole et al., 1996). In line with this concept, in cultured NSC inhibiting Erk MAP activity also ablates IGF-I-stimulated proliferation (Aberg et al., 2003a). Furthermore, in cultured oligodendrocytes IGF-I can synergize FGF stimulatory effects on Erk MAP kinase activation and on the expression of cyclin D1 (Frederick et al., 2007; Frederick and Wood, 2004), a factor that is involved in cell proliferation and the cell cycle progression. These data suggest that in neural cells the Raf–Erk pathway plays an important role in mediating IGF proliferative signaling. In these culture studies, however, Erk inhibitors and IGFs remain in medium for the entire culture period, rather than transiently during the initial IGF-induced Erk activation. Thus, it remains unclear about the nature of IGF-I-induced initial Erk

activation. In addition, because the inhibitors employed have multiple actions, inhibition of IGF-stimulated activation of other signaling pathways cannot be excluded in these studies.

The view that the Raf–Erk pathway plays a dominant role in the IGF-dependent proliferation has been further challenged by recent genetic studies, at least in oligodendrocyte precursors. IGF-I overexpressing Tg mice, which exhibit an increased neural proliferation, have a brain abundance of phosphorylated Erk similar to that in control mice (Sun and D'Ercole, 2006). Similarly the abundance of phosphorylated Erk does not change in the brain of E17 *igf1r*^{Nestin-KO} embryos (our unpublished data). In contrast, embryos with a global Erk2 KO exhibit normal progenitor proliferation (Yao et al., 2003). Because further ablation of Erk1 (i.e. the other known Erk), in cultured Erk2-null stem cells does not affect this outcome, the normal proliferation in Erk2 null mutants is unlikely due to Erk1 compensation. Studies of mice with an Erk deletion specifically in neural precursors have proven inconclusive (Heffron et al., 2009; Samuels et al., 2008; Satoh et al., 2007; Satoh et al., 2011). These data suggest that Erk is not essential in cell proliferation mediated by IGF signaling and/or other growth signaling.

In addition, data from mice with a Raf deletion specifically in neural precursors argue strongly against the idea that the Raf–Erk pathway mediates the proliferative actions of IGF and/or other signaling. As indicated above, blunting the expression of brain B-Raf significantly decreases phosphorylated Erk1 and Erk2 (Galabova-Kovacs et al., 2008; Zhong et al., 2007). Despite B-Raf ablation and haplo-insufficiency in phosphorylated Erk1/2, however, the brain of these mice exhibits minimal undergrowth (by 5–10%) and an increased number of oligodendrocyte precursors. Because IGF-I increases oligodendrocyte precursor proliferation, these findings, albeit indirectly, do not support an essential role for the Raf–Erk pathway in IGF proliferation signaling. Further evidence supporting this conclusion comes from a recent report showing that inhibition of Erk does not suppress IGF-I-stimulated proliferation of NSC/NPC in cultured cortex (Mairet-Coello et al., 2009). However, these data do not preclude the possibility that IGF signals through Erk to stimulate neural proliferation under pathological conditions.

Taking together, the MAP kinase pathway in neural IGF signaling, although not fully comprehended, appears to be primarily involved in cellular maturation and survival. In addition, the actions of the IGF-MAP kinase pathway in brain are likely cell-dependent. More detailed studies are needed.

6.2. Signaling through IGF2R

There is little evidence that the IGF2R mediates the growth actions of both IGF-I and IGF-II. Several studies, however, have strongly suggested that the IGF2R is capable of mediating some of IGF biological actions. For example, specific stimulation of the IGF2R potentiates acetylcholine release from cultured neurons (Hawkes et al., 2006), while suppressing IGF2R signaling by an IGF2R specific antibody blocks IGF-II-enhanced choline acetyltransferase activity in cultured neurons (Konishi et al., 1994) and memory in adult rodents (Chen et al., 2011).

As indicated in Section 2.2 above, there is no enzymatic activity in the intracellular domain of the IGF2R, and at present the IGF2R intracellular signaling pathway(s) remains largely unclear. Multiple studies have identified G protein as a key signaling molecule to IGF-II–IGF2R actions in neuronal cells (Hawkes et al., 2006). Involvement of G protein activity in IGF2R signaling also has been observed in non-neuronal cells (McKinnon et al., 2001). Protein kinase C (Hawkes et al., 2006), MAP kinase (McKinnon et al., 2001), and GSK3 (Chen et al., 2011) are also likely to participate IGF2R signaling.

7. IGF actions in the human CNS

IGF-I, IGF-II, IGF1R and IGF2R are widely expressed in human brain and CSF (Carlsson-Skwirut et al., 1986; Chesik et al., 2006; Connor et al., 1997; De Keyser et al., 1994; Mashayekhi et al., 2010; Wilczak et al., 2000; Wilczak and De, 1997). There is, however, little information available about the actions of IGF signaling in human neural development. Several individuals with a mutation in the *igf-I* gene or the *igf1r* gene have been reported. The available findings from these patients are consistent with the CNS abnormalities seen in mutant rodents, as summarized above in Section 4, and clearly point to an important role for IGF signaling in the human CNS.

Three homozygous *igf-I* gene mutations have been identified in three individuals, respectively. These mutations include a deletion of exons 4 and 5 of the *igf-I* gene (Camacho-Hubner et al., 1999; Woods et al., 1997), and a single nucleotide substitution (G → A) mutation each at a distinct location in the *igf-I* gene, resulting in a change of a valine residue at position 44 of the mature IGF-I to a methionine (Denley et al., 2005; Walenkamp et al., 2005) or an arginine residue at position 36 to an glutamine (Netchine et al., 2009), respectively. The mutations lead to a severe deficit in IGF1 expression (Camacho-Hubner et al., 1999; Woods et al., 1997), or to a marked reduction in its affinity for the IGF1R (Denley et al., 2005; Netchine et al., 2009; Walenkamp et al., 2005). Another patient with a homozygous T → A nucleotide substitution within exon 6 of the *igf-I* gene also has been reported (Bonapace et al., 2003), but a later study indicate that the change also is observed as polymorphism in normal persons and is not associated with growth retardation observed (Coutinho et al., 2007).

Each of these three patients exhibited intrauterine growth retardation, with a birth weight from 1400–2350 g [−4 to −2.5 standard derivation scores (SDSs)] and birth length 37.8–44 cm (−4.9 to −3.7 SDS). They had microcephaly with a head circumference 27–32 cm (−4.9 to −2.5 SDS). During postnatal development, the children persistently exhibited growth failure, and had severe deafness and mental retardation (Camacho-Hubner et al., 1999; Denley et al., 2005; Walenkamp et al., 2005; Woods et al., 1997). Despite a smaller head and likely smaller brain size, magnetic resonance imaging revealed that myelination and hippocampal architecture seemed to be normal (Camacho-Hubner et al., 1999; Netchine et al., 2009; Woods et al., 1997). While serum IGF-II was increased in the patients with *igf-I* gene mutations, it is not clear whether brain IGF-II expression also was increased, nor whether IGF-II could compensate for the IGF-I deficiency in the children, as in the apparent case in mice (Ye et al., 2002b).

More recently, a heterozygous duplication mutation in the *igf-I* gene has been reported in two patients from the same family (van Duyvenvoorde et al., 2010). The duplication mutation of four nucleotides at the position 35 of the mature IGF-I results in a frame shift and a premature termination of protein synthesis. The two patients exhibited postnatal growth retardation, but their nature, however, remains to be investigated.

There have been at least 8 individuals with a heterozygous mutation in the *igf1r* gene reported. However, there are no reports of humans born with mutations leading to absence of IGF1R expression. Heterozygous *igf1r* mutation (Abuzzahab et al., 2003; Inagaki et al., 2007; Kawashima et al., 2005; Kruis et al., 2010; Okubo et al., 2003; Raile et al., 2006; Walenkamp et al., 2006; Wallborn et al., 2010) or monoallelic IGF1R expression due to a deletion of the terminal 15q segment (Hammer et al., 2004) results in reduced IGF1R abundance and/or aberrant IGF1R functions (Abuzzahab et al., 2003; Inagaki et al., 2007; Kawashima et al., 2005; Kruis et al., 2010; Okubo et al., 2003; Raile et al., 2006; Wallborn et al., 2010), associated with an increased serum IGF-I (Inagaki et al., 2007; Raile et al.,

2006; Wallborn et al., 2010). As with the individuals carrying an *igf-I* gene mutation, patients with an *igf1r* gene mutation exhibited prenatal and postnatal growth retardation, microcephaly and mild mental retardation, learning disorders, and/or altered behavioral characteristics (Abuzzahab et al., 2003; Inagaki et al., 2007; Kawashima et al., 2005; Kruis et al., 2010; Okubo et al., 2003; Raile et al., 2006; Walenkamp et al., 2006; Wallborn et al., 2010).

Taken together, these data strongly indicate that IGF–IGF1R signaling plays a key role in the developing CNS in human. Further support for this conclusion comes from limited studies of cultured human neural cells, showing that IGF exerts pro-growth actions similar to those observed in rodents (Armstrong et al., 1992; Cui et al., 2012; Satoh and Kim, 1994; Wilson et al., 2003). Nonetheless, the details of IGF actions in the development and growth of each of the human neural cell lineages require further study. More precise brain imaging and functional studies performed during the course of development are needed to better understand the actions of IGF signaling in human brain during development.

8. Injuries and neurological disorders

Emerging data have shown that the expression of IGF system proteins is significantly altered in a variety of brain injuries and neurological disorders (Popken et al., 2005), which are characterized by a marked loss of neural cells due to either increased cell death and/or reduced proliferation. Given that IGF signaling has a critical role in proliferation, survival and differentiation of neural cells, IGFs (primarily IGF-I) have been implicated in the pathogenesis of many neurological disorders during development.

8.1. Hypoxia/ischemia (H/I) brain injury

Associations of IGF and H/I brain injury are relatively well-documented. H/I in the prenatal brain is a major cause of neural cell loss that leads to cerebral palsy, epilepsy, cognitive deficits and growth retardation in affected children (Vannucci, 1990). Following H/I brain injury, the expression of IGF-I and its downstream signaling molecules (Akt and β -catenin) sharply declines in injured and surrounding areas within 24 h following H/I (Beilharz et al., 1993; Brywe et al., 2005; Costain et al., 2008; Lee et al., 1996; Lee et al., 1992; Mastroiacovo et al., 2009), concurrent with significant neuronal apoptosis during this time (Clawson et al., 1999). IGF-I expression, followed by IGF-II expression, then progressively increases in astrocytes (Lee et al., 1996) and/or microglia (O'Donnell et al., 2002), in association with tissue recovery.

Direct evidence that IGF signaling plays an important role in protecting neural cells from H/I injury comes from studies of rodents administered with IGF-I or IGF-II. In rodents IGF-I administered directly to the H/I brain prevents the loss of neurons and oligodendrocytes (Cao et al., 2003; Guan et al., 2001; Lin et al., 2005), increases the number of newly born cells in the DG (Dempsey et al., 2003), and improves neurological signs (Zhong et al., 2009). Increasing the abundance of β -catenin, a molecular in both IGF and Wnt signaling pathways, by lithium treatment or by blunting the expression of the Wnt antagonist DICKKOPF-1 markedly ameliorates ischemic cortical damage (Mastroiacovo et al., 2009).

More recently, it has been reported that peripherally administered IGF-I through subcutaneous injection (Zhong et al., 2009) or nasal insufflation (Lin et al., 2009) mitigates H/I brain injury significantly, in part by promoting the survival of neuronal cells and proliferation of neural precursor cells. Notably, IGF-I remains effective when given by subcutaneous injection 24 and 48 h after H/I injury (Zhong et al., 2009), a finding that supports the clinical utility of IGF-I in treating ischemic neural injury. Consistent with these findings, an early clinical trial shows that IGF-I, given intravenously, appears to improve

clinical outcome in patients with moderate-to-severe head injury (Hatton et al., 1997), whose BBB was likely compromised, facilitating translocation of blood IGF-I into the brain.

The capacity of IGF-I to promote neuronal cell survival during H/I injury is, at least in part, mediated by IGF1R. Blunting the IGF1R expression in neuronal cells exacerbates H/I-induced neuronal apoptosis (Liu et al., 2011). Mice with ablated neural IGF1R expression also exhibit a reduction in the expression of both cellular IAP-1 (cIAP-1) and X-linked IAP (XIAP) Liu et al., 2011, two members of IAP family proteins that are capable of suppressing the activity of caspases by either enhancing caspase degradation (cIAP) or directly inhibiting their activity (XIAP) Eckelman and Salvesen, 2006; Eckelman et al., 2006. These data suggest that IGF-stimulated IAP expression partially contributes to IGF-IGF1R protection against neuronal apoptosis following H/I injury.

8.2. Schizophrenia

Substantial evidence suggests that schizophrenia is a developmental disorder of the brain, reflecting a genetically conferred vulnerability combined with early insults from the environment (Boksa, 2004; Fatemi and Folsom, 2009; Tsuang, 2000). Epidemiological studies have provided evidence that exposure to certain obstetric complications is associated with a significantly increased risk for the later development of schizophrenia (Boksa, 2004; Fatemi and Folsom, 2009). Obstetric complications with the greatest effect size for schizophrenia include maternal diabetes, infections during pregnancy, preeclampsia, intrauterine growth retardation, and fetal/neonatal hypoxia.

Plasma levels of IGF-I are significantly decreased in schizophrenic patients (Kyle and Pichard, 2006; Venkatasubramanian et al., 2007; Wu et al., 2008). Microarray studies of the cerebral cortex and hippocampus from cases of schizophrenia and of the DG in a mouse model of human 22q11 deletion have revealed decreased gene expression and proteins for mitochondrial oxidative energy metabolism and specific cytoskeletal proteins implicated in synaptic functions (Altar et al., 2005; Jurata et al., 2006; Mirnics et al., 2000; Prabakaran et al., 2004). IGF-I and insulin have been shown to modulate these genes in a dose-dependent fashion, increasing the expression of those genes that are decreased in schizophrenia (Altar et al., 2008).

Brain imaging and postmortem histological studies of schizophrenia patients show gross pathological abnormalities, including marked reductions in the volumes of cortical gray matter, the hippocampal formation, and subcortical myelin (Andreone et al., 2007; Bartzokis et al., 2003; Di et al., 2009; Fornito et al., 2009; Friedman et al., 2008; Goldstein et al., 1999; Harrison, 1999; van Haren et al., 2008; Voineskos et al., 2010). Perhaps the most robust lesion found in human schizophrenia is a preferential loss of a particular subset of GABAergic interneurons in the dorsal prefrontal cortex (DPFC). By immunohistochemical analysis, this neuron loss involves those GABAergic interneurons which are GABA+/parvalbumin+ and GABA+/calbindin+ interneurons, but not those that are GABA+/calretinin+ (Beasley and Reynolds, 1997; Beasley et al., 2002; Benes et al., 1986; Benes et al., 1991; Reynolds and Beasley, 2001).

In heterozygous *igf1r*^{Nestin-KO} mice, whose IGF1R expression is conditionally ablated in nestin+ neural cells, our ongoing morphometric and stereological studies have revealed a loss of GABAergic neurons that are positive for calbindin, but not for calretinin, in the ventral prefrontal cortex (VPFC) (see Section 4.3. and Fig. 4). The VPFC in rodents corresponds to human DPFC (Heidbreder and Groenewegen, 2003), which is known to be associated with schizophrenia pathogenesis. Our unpublished behavioral studies further show that adult heterozygous *igf1r*^{Nestin-KO} mice exhibit behavioral abnormalities. These abnormalities include impaired learning and memory and abnormal acoustic startle/pre-pulse

inhibition (AS/PPI), which are consistent with hallmark signs of schizophrenia in mouse models (Arguello and Gogos, 2006). These data suggest that reduced IGF-I signaling during prenatal and early postnatal development may play a role in the pathogenesis of schizophrenia during development.

9. Conclusion

Over the past 20 years or more, literature illustrates that IGF signaling plays critical roles in the growth and development of the CNS at virtually every stage of development. IGF-I increases the survival of pre-implantation embryos reaching the blastocyst stage through its anti-apoptotic actions and through its ability to protect against reactive oxygen species. Both IGF-I and IGF-II promote NSC/NPC proliferation during development, and IGF-I accelerates neurogenesis by reducing G₁ phase length of the mitotic cell cycle while increasing cell cycle re-entry. IGF-I promotes neural cell survival during normal development and in response to pathological insults. In postmitotic neurons, IGF-I acts to promote elongation and branching of dendrites and the relative density of dendritic spines. IGF-I also is capable of augmenting the progressive phase of synaptogenesis and the myelination of maturing axons. Clearly, the IGFs and their related IGF family proteins as therapeutic targets have tremendous potential in the treatment of injuries and developmental disorders of the brain and in various neurological diseases in the adult brain.

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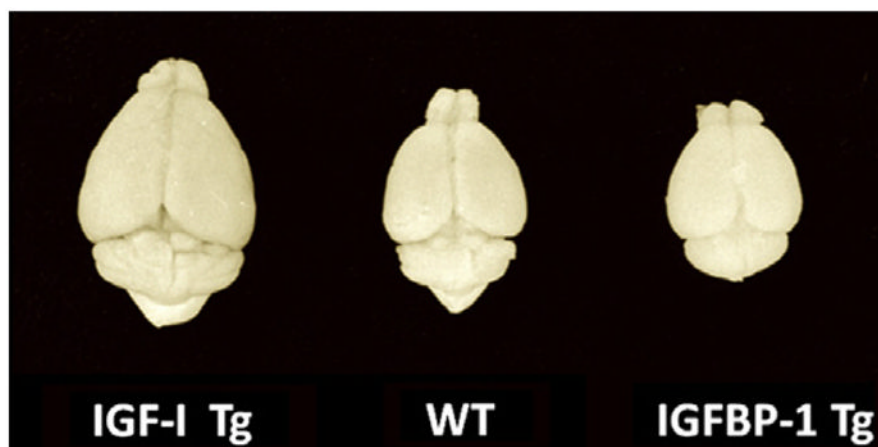


Fig. 1. Brains from an IGF-I^{MT-I} overexpressing Tg mouse (IGF-I Tg), a wild type control mouse (WT), and an IGFBP-1 Tg mouse that ectopically express brain IGFBP-1, which reduces IGF availability.

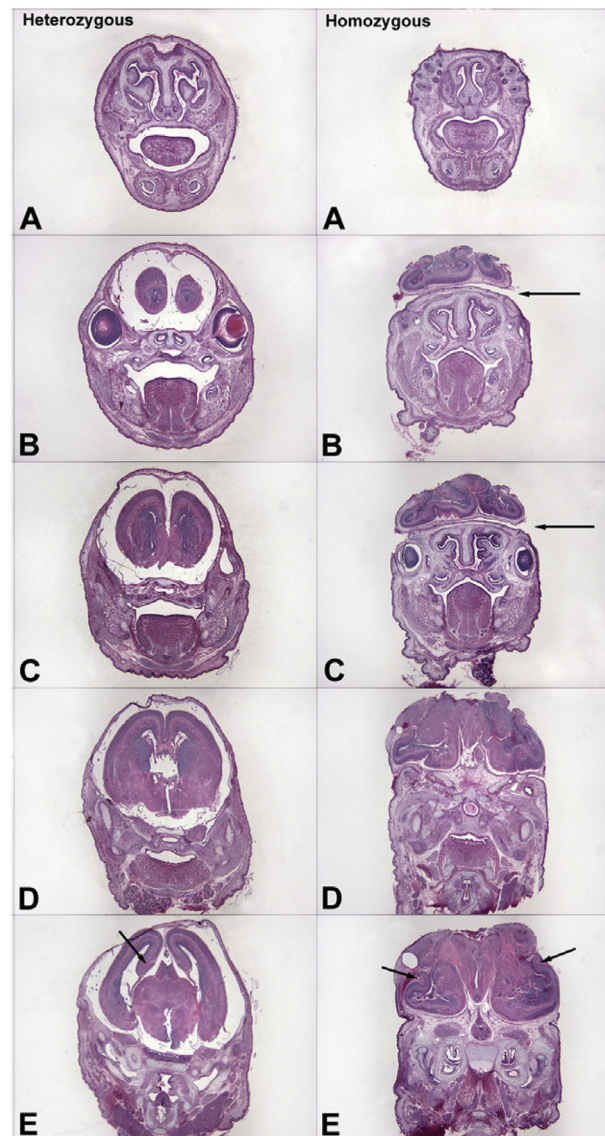


Fig. 2.

Representative sections through the head and brain of an E16 homozygous *igf1r*^{Nestin-KO} embryo (right panels) and its heterozygous littermate (left panels), stained with hematoxylin and eosin. In heterozygous embryos the brain exhibits relatively normal morphology at all levels, although smaller in size compared to littermate controls (not illustrated). In homozygous embryos the brain exhibits gross malformations in some mice. There is a failure to close the longitudinal suture of the skull in homozygous embryos, accompanied by the extrusion of gray matter through the skull and sloping forward (Panels B and C, arrows). In heterozygous embryos the developing hippocampus and DG are located along the medial edge of the telencephalic wall as in controls (E, arrow). In homozygous embryos the hippocampus and DG rotated to the lateral edge of the telencephalic wall (E, arrows) following extensive apoptosis in the dorsolateral wall at E14 and protrusion of the underlying telencephalic structures.

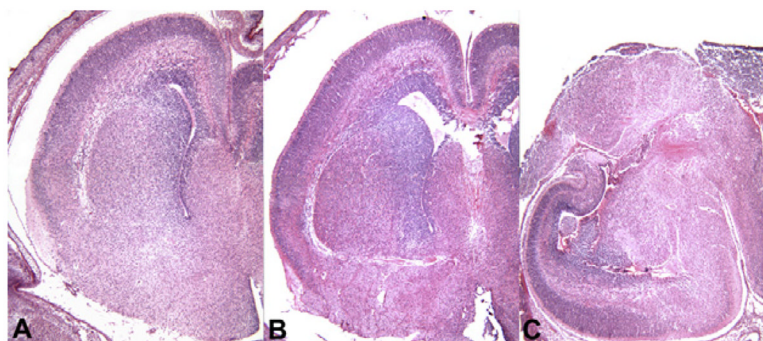


Fig. 3. Representative sections of the right cerebral hemisphere in a control embryo (A), a heterozygous *igflr*^{Nestin-KO} embryo (B), and a homozygous *igflr*^{Nestin-KO} embryo at E16.

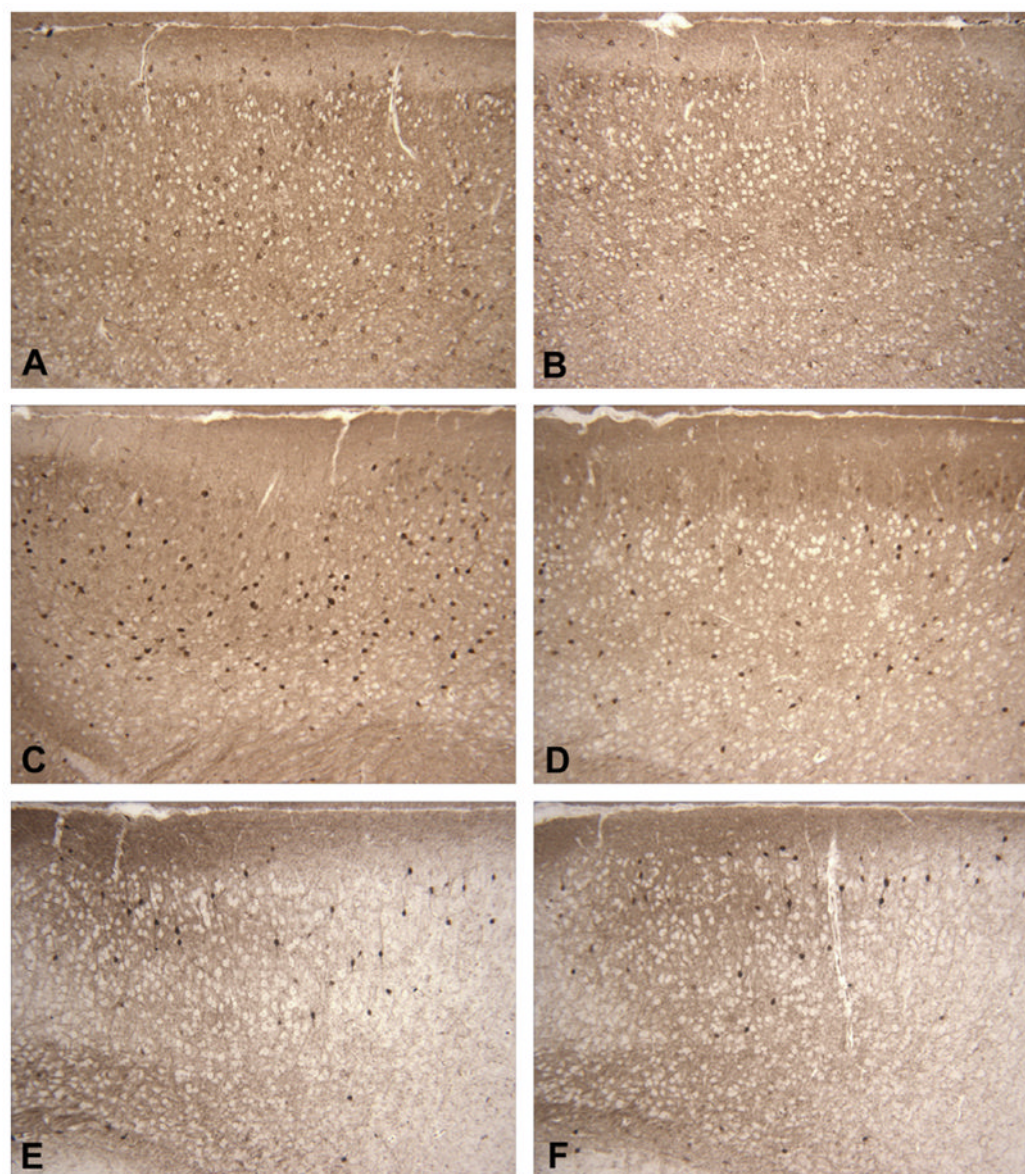


Fig. 4.

Immunostaining for GAD67, calbindin, and calretinin in the ventral prefrontal cortex of an adult heterozygous *igflr*^{Nestin-KO} mouse (B, D, and F) and its littermate control (A, C, and E). Coronal sections through the ventral prefrontal cortex have been stained with antibodies against GAD67 (A and B), calbindin (C and D) and calretinin (E and F). Note the decreased density of GAD67-immunoreactive GABAergic neurons in (B) and the decreased density of calbindin-immunoreactive GABAergic neurons in (D) with no change in the density of calretinin-immunoreactive GABAergic neurons in (F), when compared to controls.

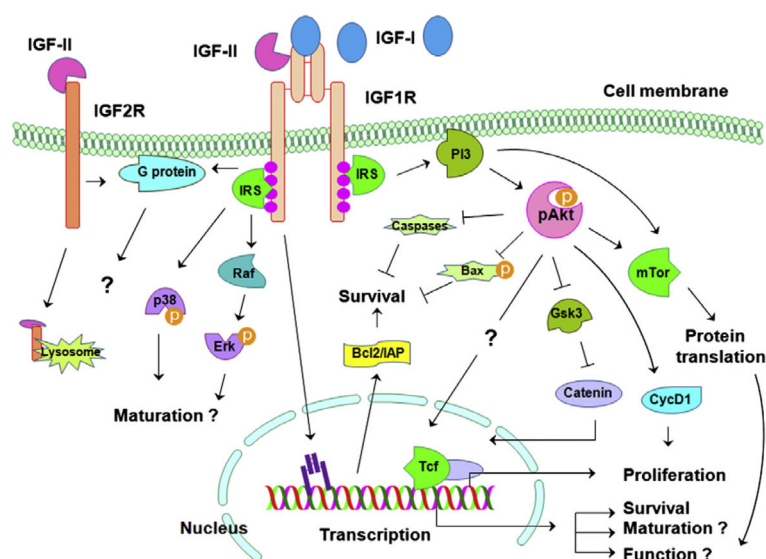


Fig. 5.

IGF signaling in the CNS. In this largely simplified diagram the IGF signaling pathways in the CNS are schematically depicted. Other signaling molecules and pathways described in non-neural cells systems are not included in the Figure, although it is possible that they are also involved. \perp = inhibitory modification, and \downarrow = stimulatory modification.

Table 1

mRNA expression of IGF system proteins in neural cells in developing and adult brain.

Brain regions	IGF-I	IGF-II	IGF1R	IGF2R
Cerebrum				
Cerebral cortex	+		+	+
VZ/SVZ	+		+	
Corpus callosum			+	+
Hippocampus				
CA	+		+	+
DG	+		+	+
Olfactory bulb				
Mitral cell layer	+		+	+
Tufted cell layer	+		+	
Hypothalamus	+		+	+
Periventricular nucleus			+	+
Supraoptic nucleus			+	+
Thalamus	+		+	+
Basal Ganglia			+	+
Midbrain	+		+	+
Brainstem	+		+	+
Cerebellum	+			
External granule cells			+	+
Internal granule cells	+		+	+
Purkinje cells	+		+	+
Deep cerebellar nuclei	+			+
Others				
Meninges	+	+	+	+
Choroid plexus	+	+	+	+
Blood vessels		+	+	

1. Data comprised in Table 1 are primarily derived from mRNA *in situ* hybridization studies of rodents at ages of E18 to adult, and adapted from references (Ayer-le et al., 1991; Bartlett et al., 1992; Bondy and Chin, 1991; Bondy and Lee, 1993; Bondy, 1991; Cavallaro et al., 1993; Lee et al., 1993; Walter et al., 1999; Zhang et al., 2007). Immunohistochemical staining and ¹²⁵I-IGF-II receptor binding studies also are used for IGF1R and IGF2R (Folli et al., 1994; Hawkes and Kar, 2003; Kar et al., 1993; Quesada et al., 2007).

2. IGF-II is expressed in vascular endothelial cells (Ayer-le et al., 1991; Dugas et al., 2008; Logan et al., 1994; Stylianopoulou et al., 1988).

Table 2

Phenotype changes in IGF system protein mutant mice.

Protein	Abbreviation	Neural outcomes	References
IGF-I			
	IGF-I ^{MT-1} Tg	↑ Brain growth, ↑ myelination	(Behringer et al., 1990; Carson et al., 1993; Mathews et al., 1988)
	IGF-I ^{MT-1} Tg ^a	↑ Brain growth, ↑ myelination, ↑ oligodendrocyte and oligodendrocyte precursor number, ↑ oligodendrocyte and oligodendrocytes precursor proliferation, ↑ neuron number, ↓ neuronal apoptosis, ↑ synaptogenesis, ↑ uptake of [³ H]2-deoxyglucose	Gutierrez-Ospina et al., 1996; Gutierrez-Ospina et al., 1997; Mason et al., 2000; Ye et al., 1995a; Ye et al., 1995b
	IGF-I ^{Nestin} Tg	↑ Brain growth, ↓ mitotic cell cycle duration, ↑ neuron number	Popken et al., 2004
	IGF-I ^{IGF-II} Tg	↑ Brain growth, ↑ neuronal proliferation, ↑ neuron number, ↓ neuronal apoptosis, ↑ synaptogenesis, ↓ levels of 14-3-3	Chrysis et al., 2001; Dentremon et al., 1999; O'Kusky et al., 2000; O'Kusky et al., 2003; Ye et al., 1996; Zhang et al., 2003
	IGF-I ^{GFAP} Tg	↑ Brain growth, ↑ neuron number, ↑ neuronal proliferation, ↑ astrocyte number, ↑ GFAP expression, ↑ myelination, ↑ histone H3 and H4 acetylation	Lehtinen et al., 2011; Sun and D'Ercole, 2006; Ye et al., 2004
	<i>igf-I</i> KO	↓ Brain growth, ↓ myelination, ↓ oligodendrocyte number, ↓ neuron number	Ye et al., 2002b
	<i>igf-I</i> KO	↓ Brain growth, ↓ myelination, ↓ oligodendrocyte number, ↓ neuron number ↓ dendritic length and branching	Beck et al., 1995; Cheng et al., 2003
	<i>igf-I</i> ^{m/m} KO ^b	Brain growth not reported	Lembo et al., 1996
	<i>igf-I</i> ^{Alb-KO}	No growth changes in body, liver, spleen, kidney and heart. Brain growth not reported	Sjogren et al., 1999; Yakar et al., 1999
IGF-II			
	<i>igf-II</i> KO	↓ Brain growth, ↓ neuron number	Lehtinen et al., 2011
	IGF-II ^{H-2Kb} Tg	No obvious brain phenotypic changes	Reijnders et al., 2004; Smink et al., 1999; Van Buul-Offers et al., 1995
IGF1R			
	<i>igf1r</i> KO		
	Heterozygous	No phenotypic changes	Liu et al., 1993
	Homozygous	↓ Brain growth	(Liu et al., 1993)
	<i>igf1r</i> ^{mini-KO} b	↓ Brain growth	(Holzenberger et al., 2000)
	<i>igf1r</i> ^{Meu} KO ^d	↓ Brain growth	Holzenberger et al., 2001
	<i>igf1r</i> ^{Nestin-KO}	↓ Brain growth, ↓ neuron number, ↓ neuronal proliferation, ↑ neuronal apoptosis	(Liu et al., 2011; Liu et al., 2009)
	<i>igf1r</i> ^{Nestin-KO}	↓ Brain growth, ↓ neuron number	(Kappeler et al., 2008)
	<i>igf1r</i> ^{Olig1-KO}	↓ Brain growth, ↓ myelination, ↓ oligodendrocytes and oligodendrocytes precursor number	(Zeger et al., 2007)
	<i>igf1r</i> ^{PLP-KO}	↓ Brain growth, ↓ oligodendrocytes number, ↓ myelination	(Zeger et al., 2007)
	<i>igf1r</i> ^{GnRH-KO}	↓ dendritic branching, ↓ spine formation	(DiVall et al., 2010)
IGFBPs			
	IGFBP-1 ^{MT} Tg	↓ Brain growth, ↓ myelination, ↓ oligodendrocyte and oligodendrocytes precursor number, ↓ neuron number	(Gutierrez-Ospina et al., 1996; Ye et al., 1995a; Ye et al., 1995b)
	IGFBP-1 ^{PGK} Tg	↓ Brain growth, ↓ myelination, ↓ neuron number	(Ni et al., 1997; Rajkumar et al., 1995)
	IGFBP-2 ^{CMV} Tg	↓ Brain growth	(Hoeflich et al., 2001; Hoeflich et al., 1999)

Protein	Abbreviation	Neural outcomes	References
	IGFBP-3 ^{GKG} Tg	↓ Brain growth	(Modric et al., 2001)
	IGFBP-3 ^{GKG mini} Tg ^e	↓ Brain growth, ↓ neural proliferation in the periventricular zone	(Silha et al., 2005)
	ALS ^{CMV} Tg	↔ Brain growth	(Modric et al., 2001)
	IGFBP-5 ^{Actin} Tg	↓ Body growth, ↑ relative brain growth	(Salih et al., 2004)
	IGFBP-5 ^{MT} Tg	↔ Brain growth	f
	IGFBP-6 ^{GFAP} Tg	↓ Brain growth, ↓ astrocyte number	Bienvenu et al., 2004

^aTransgenic mice were generated using the same transgene construct that was employed for the original line of mice described in Behringer et al. (1990), Carson et al. (1993), Mathews et al. (1988).

^bUnlike IGF-I KO and IGF1R KO mice in which IGF-I and IGF1R expression is completely ablated, *igf-I*^{m/m}-KO and *igf1r*^{mini}-KO mutant mice exhibit marked reduction in IGF-I or IGF1R expression, respectively, due to an insertion mutation.

^c*igf-I*^{Alb}-KO mice carry a *igf-I* null mutation specifically in liver, and exhibit a severe reduction (by 70%) in serum IGF-I.

^d*igf1r*^{Meu} KO mice carry a null mutation on one *igf1r* allele and a partial inducible null mutation on the other *igf1r* allele during late embryonic development.

^eA human Gly(56)/Gly(80)/Gly(81)-mutant IGFBP-3 cDNA was inserted downstream of a GKG promoter. The mutant has a markedly reduced affinity for the IGFs, but retains the IGF-independent effects. No phenotypic changes, except for brain, are observed.

^fOur unpublished data.