

JNK/stress-activated protein kinase associated protein 1 is required for early development of telencephalic commissures in embryonic brains

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Abbreviations: AC, anterior commissure; CC, corpus callosum; FAK, focal adhesion kinase; JSAP1, JNK/stress-activated protein kinase-associated protein 1; MEF, mouse embryonic fibroblasts; TCAs, thalamocortical axons

Abstract

We previously reported that mice lacking JSAP1 (*jsap1*^{-/-}) were lethal and the brain of *jsap1*^{-/-} at E18.5 exhibited multiple types of developmental defects, which included impaired axon projection of the corpus callosum and anterior commissures. In the current study, we examined whether the early telencephalic commissures were formed abnormally from the beginning of initial development or whether they arose normally, but have been progressively lost their maintenance in the absence of JSAP1. The early corpus callosum in the brain of *jsap1*^{+/+} at E15.5-E16.5 was found to cross the midline with forming a distinct U-shaped tract, whereas the early axonal tract in *jsap1*^{-/-} appeared to cross the midline in a diffuse manner, but the lately arriving axons did not cross the midline. In the brain of *jsap1*^{-/-} at E17.5, the axon terminals of lately arriving collaterals remained within each hemisphere, forming an early Probst's bundle-like shape. The early anterior commissure in the brain of *jsap1*^{+/+} at E14.5-E15.5 crossed the

midline, whereas the anterior commissure in *jsap1*^{-/-} developed, but was deviated from their normal path before approaching the midline. The axon tracts of the corpus callosum and anterior commissure in the brain of *jsap1*^{-/-} at E16.5-E17.5 expressed phosphorylated forms of FAK and JNK, however, their expression levels in the axonal tracts were reduced compared to the respective controls in *jsap1*^{+/+}. Considering the known scaffolding function of JSAP1 for the FAK and JNK pathways, these results suggest that JSAP1 is required for the pathfinding of the developing telencephalic commissures in the early brains.

Keywords: axons; focal adhesion kinase 1; JNK mitogen-activated protein kinases; MAPK8IP3 protein, human

Introduction

JNK/stress-activated protein kinase-associated protein 1 (JSAP1), also called JIP3, functions as a scaffold protein for components of the c-Jun N-terminal kinase (JNK) pathway (Whitmarsh *et al.*, 1998; Yasuda *et al.*, 1999; Kelkar *et al.*, 2000). JSAP1 also interact with focal adhesion kinase (FAK), which is activated during cell adhesive interactions with the extracellular matrix (Takino *et al.*, 2002, 2005). Moreover, JSAP1 is distributed in growth cones and plasma membrane as well as in the soma of neurons during development (Kelkar *et al.*, 2000; Sato *et al.*, 2004, 2008, 2011; Abe *et al.*, 2009). Thus, these findings indicate that JSAP1 has a function to regulate axonal development. Indeed, mice lacking JSAP1 show pathfinding defects of the thalamocortical tracts in developing brain (Ha *et al.*, 2005).

FAK and JNKs are highly expressed in the developing brain, and are important in neurite outgrowth and cell migration (Burgaya *et al.*, 1995; Huang *et al.*, 2003; Zhan *et al.*, 2003; Rico *et al.*, 2004; Robles and Gomez, 2006). FAK-deficient cells form focal adhesion/complexes, but show a decreased rate of cell migration due to the delayed disassembly of focal adhesions (Ilić *et al.*, 1995; Reiske *et al.*, 1999; Sieg *et al.*, 1999; Jacamo *et al.*, 2007). Moreover, phosphorylated JNK (phospho-JNK) is colocalized with FAK in focal adhes-

ions of fibroblasts cultured on fibronectin (Almeida *et al.*, 2000; Takino *et al.*, 2005), suggesting that JNK is also involved in cell adhesion. And, *jnk1*^{-/-} mice exhibit disrupted tract formation of the anterior commissure (AC) in the brain resulting from the progressive loss of microtubules within axons and dendrites. It has been shown that MAP2 and MAP1B have a reduced ability to bind microtubules and to promote their assembly in the absence of JNK1 (Chang *et al.*, 2003). Consistent with the known roles of JNK in cell adhesion, the ectopic expression of the mixed lineage kinase, MUK/DLK/ZPK, in neural precursor cells impairs radial migration, whereas it allows these cells to leave the ventricular zone and differentiate into neural cells, suggesting that MUK/DLK/ZPK and JNK regulate radial cell migration *via* microtubule-based events (Hirai *et al.*, 2002; Eto *et al.*, 2010).

Recent molecular genetic studies have reported that E18.5 brains with the JSAP1 null mutation show multiple developmental defects, such as impaired axon guidance of the corpus callosum (CC) and AC, pathfinding defects of thalamocortical tracts and migration defects of anti-calretinin positive cells in the developing cerebral cortex (Kelkar *et al.*, 2003; Ha *et al.*, 2005). The developmental defects identified in the E18.5 brain of *jsap1*^{-/-} should occur during developmental stages earlier than E18.5, probably as a consequence of the altered function of JSAP1-assisted signaling pathways (Ha *et al.*, 2005; Chae *et al.*, 2006). Moreover, because JSAP1 functions as a scaffold protein for JNK and FAK, the potential organization of JNK and FAK by JSAP1 might be important for the distribution or activation of these cellular factors in developing neurons. Thus, it would be intriguing to determine whether the JSAP1 dependent distribution and activations of JNK and FAK are involved in the impaired axon guidance of telencephalic commissures.

In the present study, we examined the appearance and progression of developmental defects in the early *jsap1*^{-/-} brain. In addition, we investigated whether the null mutation of JSAP1 affects the temporal distributions of phospho-JNK and phospho-FAK in the axon tracts of the telencephalic commissures.

Results

Temporal patterns of developmental defects of the telencephalic commissures in the brains of *jsap1*^{-/-}

In a previous study, we reported that the E18.5 *jsap1*^{-/-} brain showed impaired axon projections of the CC and AC (Ha *et al.*, 2005). In the current

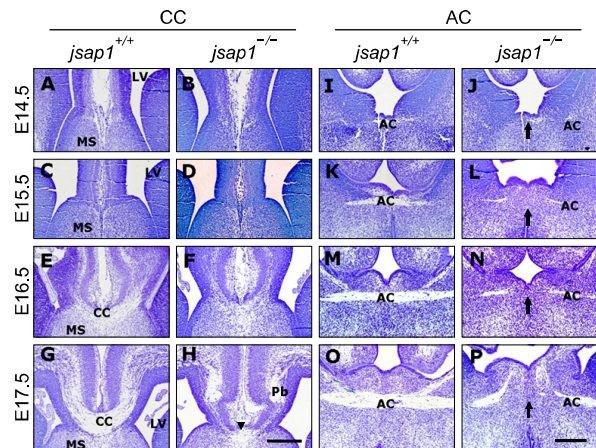


Figure 1. Developmental defects of the corpus callosum and anterior commissures in developing brains of *jsap1*^{-/-}. (A-N) Cresyl violet staining of coronal sections of paraffin-embedded brains at developmental stages; E14.5 (A, B, I, J), E15.5 (C, D, K, L), E16.5 (E, F, M, N) and E17.5 (G, H, O, P). In E14.5-E15.5 brains, the histologic configuration of the corpus callosum in *jsap1*^{-/-} was indistinguishable from that of *jsap1*^{+/+} (A-D). Note that the early axon tracts of the corpus callosum in E16.5 brains (E, F) were identifiable in both *jsap1*^{+/+} and *jsap1*^{-/-}. Probst's bundle-like structures (Pb) formed in the cingulate cortex at E17.5 in *jsap1*^{-/-} brain were indicated (H). Note that the axons of the anterior commissures were distracted from their normal path long before approaching the midline (I-P), and thus were disconnected during development (arrows). AC, anterior commissures; CC, corpus callosum; LV, lateral ventricle; MS, medial septum; Th, thalamus; arrows and arrow heads, defected area. Scale bars, 500 μ m.

study, we investigated the temporal patterns of the developmental defects during the early development of *jsap1*^{-/-} brains. Previous studies reported that the early axonal extension of the CC, which stems from the cingulate cortex, pioneers across the midline of the telencephalon during E15.5-E16.5, and subsequently the late arriving collaterals originating from other regions of the neocortex join the existing tract (Koester and O'Leary, 1994; Ozaki and Wahlsten, 1998; Rash and Richards, 2001). Cresyl violet-stained brain sections showed that the overall morphology of the telencephalon of *jsap1*^{-/-} at E14.5-E15.5 did not obviously differ from that of *jsap1*^{+/+} (Figures 1A-1D; see also Figures 2-5). Our cresyl violet-stained brain sections revealed that *jsap1*^{+/+} brains formed the CC during E15.5-E16.5 as reported previously (Koester and O'Leary, 1994; Ozaki and Wahlsten, 1998), while the early axons of the CC in the *jsap1*^{-/-} brain at E15.5-E16.5, at least in part, crossed the midline (Figures 1C-1F). In *jsap1*^{+/+} brains at E17.5, the axon bundle of the CC became thicker and formed a typical U-shaped tract, whereas in *jsap1*^{-/-} brains, late arriving neocortical-originated callosal axons failed to cross the midline. Instead, these axons projected only within each hemisphere, and formed Probst's bundle-like structures bilaterally in the

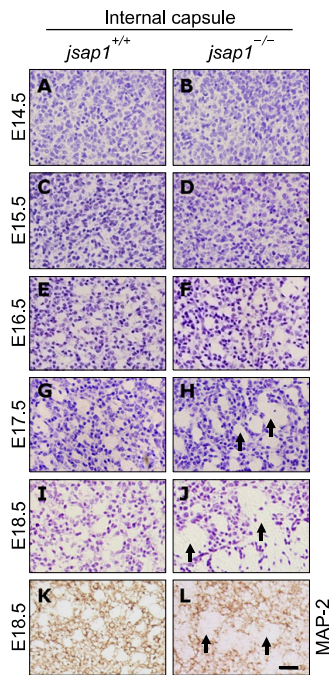


Figure 2. Developmental defects of the internal capsule in developing brains of *jsap1*^{-/-}. (A-L) Cresyl violet (A-J) or anti-MAP-2 (K-L) stained coronal sections of E14.5-E18.5 brains. The fiber tracts of the internal capsule were not clearly detected until E15.5 in both genotypes. The thickness of the fiber tracts in *jsap1*^{-/-} at E17.5-E18.5 was larger than that of *jsap1*^{+/+}. Arrows indicate the fiber tracts of the internal capsule. Scale bar, 100 μ m.

cingulate cortex (Figures 1G and 1H). The Probst's bundles are known to be comprised of axonal tips that have failed to develop properly to cross the midline after their initial extension to the medial hemispheric walls (des Neves *et al.*, 1999; Shen *et al.*, 2002).

The second telencephalic commissure, AC, comprises of two major axon pathways; the anterior limb that connects the anterior olfactory nucleus and anterior piriform cortex, and the posterior limb that connects the piriform and temporal cortices (Abbie, 1940). The early extension of the AC in *jsap1*^{+/+} was clearly detected and appeared to cross the midline at E14.5. The axonal projection of the AC in *jsap1*^{-/-} appeared at E14.5, but did not cross the midline (Figures 1I and 1J). The developing axons of the AC during E15.5-E17.5 appeared to be thinner than *jsap1*^{+/+}. Moreover, developing axons of the AC were deviated from their normal path before approaching the midline and the AC axons in *jsap1*^{-/-} headed to the thalamus (Figures 1K-1P). The average diameter of axonal tracts of the AC in *jsap1*^{-/-} was relatively thinner than *jsap1*^{+/+} (Figures 1K-1P).

In contrast to the telencephalic commissures described above, the average diameter of axonal

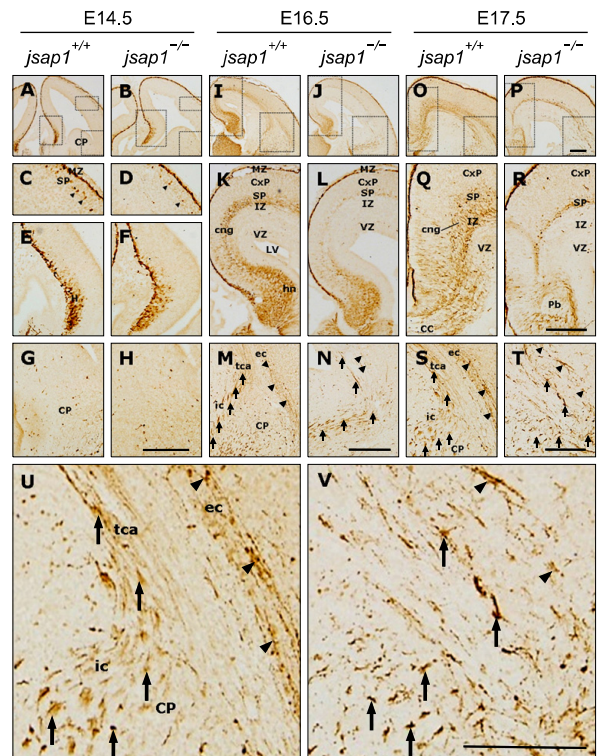


Figure 3. Anti-calretinin staining showed altered expression levels and developmental defects in thalamocortical fibers in the brains of *jsap1*^{-/-}. (A-V) Anti-calretinin stained coronal sections of the cortices of developing brains at E14.5 (A-H), E16.5 (I-N) and E17.5 (O-V). Anti-calretinin immunoreactivity in *jsap1*^{-/-} brains at E16.5-17.5 was reduced in many brain regions compared to *jsap1*^{+/+} brains (I-V). Anti-calretinin labeled thalamocortical fibers (arrows) in *jsap1*^{-/-} brains at E16.5-17.5 appeared to have a pathfinding defect (M, N, S-V). High magnification of the panels S and T (U, V). cng, cingulum bundle; CP, caudo-putamen; CxP, cortical plate; hn, hippocampal neuroepithelium; ic, internal capsule; IZ, intermediate zone; ec, external capsule; LV, lateral ventricle; MZ, molecular zone; SP, subplate; tca, thalamocortical axons; VZ, ventricular zone. Scale bars, 500 μ m.

tracts (internal capsule) that pass through the striatum in *jsap1*^{-/-} was thicker than that in *jsap1*^{+/+} at E16.5-E18.5 (Figures 2A-2L). In E16.5, the fiber tracts of the internal capsule were clearly present and their thickness was similar in both genotypes (Figures 2E and 2F). After E17.5, however, the thickness of the fiber tracts in *jsap1*^{-/-} was larger than that of *jsap1*^{+/+} (Figures 2G-2L). It remained unknown whether these axonal tracts all reach to their target neurons in the telencephalic cortex properly.

In a previous study, we demonstrated that calretinin (calcium-binding protein) labels a subset of neurons and axonal tracts in the brain of E18.5 embryo, permitting to detect distinct developmental deficits in *jsap1*^{-/-} brain (Ha *et al.*, 2005). In the present study, we continued to explore the distribution of calretinin-positive cells and axonal

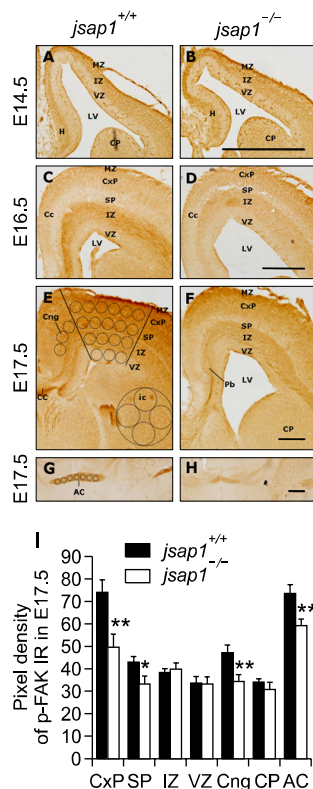


Figure 4. Reduced expression of phospho-FAK levels in developing brains of *jsap1*^{-/-}. (A-H) Anti-phospho-FAK stained coronal sections of cryo-cut brains at E14.5 (A, B), E16.5 (C, D), and E17.5 (E-H). Phospho-FAK was distributed in the soma and neuritic processes of neuronal cells in broad regions in the developing *jsap1*^{+/+} and *jsap1*^{-/-} brains. The expression level of phospho-FAK in the U-shaped axon tract of the corpus collasum (C-F) and anterior commissure (G, H) in *jsap1*^{-/-} brains was lower than that in *jsap1*^{+/+} brains. (I) Quantification of the expression levels of phospho-FAK in indicated brain regions in both genotypes. The pixel density of immunoreactivity was measured, as described in Materials and Methods. * $P < 0.05$ and **, $P < 0.01$, differences between *jsap1*^{+/+} and *jsap1*^{-/-} brains (Student's *t*-test). AC, anterior commissure; CC, corpus callosum; CxP, cortical plate; IZ, intermediate zone; LV, lateral ventricle; MZ, molecular zone; Pb, Probst's bundle; VZ, ventricular zone. A, C, and E; *jsap1*^{+/+}. B, D, and F; *jsap1*^{-/-}. Scale bars, 500 μ m.

tracts in earlier embryonic brains. In the E14.5 brain, anti-calretinin labeled the marginal zone, subplate, hippocampus and caudoputamen, and their expressions were similar between the two genotypes (Figures 3A-3H). In E16.5-E17.5 *jsap1*^{+/+} brains, anti-calretinin labeled cell bodies and axonal processes of neurons in the marginal zone, cortical plate, subplate, intermediate zone, hippocampal neuroepithelium, and caudoputamen. While in *jsap1*^{-/-}, anti-calretinin similarly labeled the cell bodies and axonal processes of neurons in these regions, but staining levels were lower than those of *jsap1*^{+/+} (Figures 3I-3T).

Developing thalamocortical axons (TCAs) were also stained by anti-calretinin. Anti-calretinin barely

labeled TCAs at E14.5 (Figures 3G and 3H). In E16.5-E17.5 *jsap1*^{+/+} brains, anti-calretinin labeled TCAs running along the intermediate zone of the cortical cortex. Whereas in E16.5-E17.5 *jsap1*^{-/-}, anti-calretinin staining showed the pathfinding defects of TCAs at the junction between the caudoputamen and the cortical wall, and their axon collaterals ran diffusely in the intermediate zone (Figures 3N and 3T). Stained axons at high magnification revealed that in *jsap1*^{+/+}, calretinin was distributed in a patched manner along the thalamocortical and/or corticofugal projections, whereas in *jsap1*^{-/-}, it was detected in the axonal tracts, but with an infrequent abnormal heavy accumulation in certain axons (Figures 3S and 3T).

Distribution of the phosphorylated forms of FAK and JNK in developing neurons in the early brains of *jsap1*^{-/-}

In E14.5 *jsap1*^{+/+} brain, phospho-FAK was distributed in broad brain areas with enhanced expression in the intermediate and molecular zones of the cortical plate. This distribution pattern did not obviously differ from that in the respective brain of *jsap1*^{-/-} (Figures 4A and 4B). In E16.5 brain, anti-phospho-FAK clearly labeled U-shaped early callosal axons in both genotypes, though staining level in the axon tract in *jsap1*^{-/-} was lower than that in *jsap1*^{+/+} (Figures 4C and 4D). In E17.5 *jsap1*^{+/+} brain, phospho-FAK distributed in most brain regions, including the corpus callosum, anterior commissure intermediate zone, and cortical plate, wherein phospho-FAK was distributed mainly in neuritic processes and soma of neurons, and also labeled radial glial fibers in the cortical plate (Figure 4E). While in the E17.5 brain of *jsap1*^{-/-}, the immunoreactivity of p-FAK was decreased and particularly in the axon tract of the corpus callosum (Figures 4E, 4F and 4I) and the anterior commissure (Figures 4G-4I). In E14.5-E17.5 brains of both genotypes, FAK was expressed in most brain regions, and this expression pattern was similar to that of phospho-FAK (data not shown).

Similarly, anti-phospho-JNK also broadly labeled developing brains. In E14.5 brains, enhanced phospho-JNK immunoreactivities were detected in the marginal zone and intermediate zone of the cortical plate, caudoputamen, ventricular zone, and hippocampus in *jsap1*^{+/+}. The anti-phospho-JNK levels in the corresponding regions of *jsap1*^{-/-} were slightly, though not dramatically, lower than *jsap1*^{+/+} (Figures 5A and 5B). In E16.5-E17.5 brains, anti-phospho-JNK labeled developing neurons in the marginal zone, subplate, intermediate zone, subventricular zone, caudoputamen, and cingulum

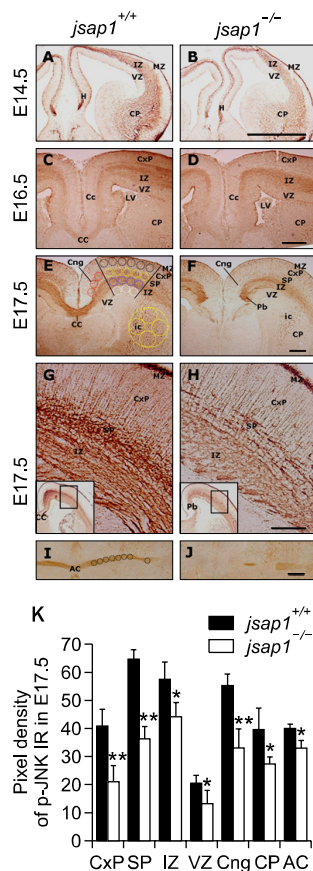


Figure 5. Reduced expression of phospho-JNK levels in developing brains of *jsap1*^{-/-}. (A-J) Anti-phospho-JNK stained coronal sections of cryo-sectioned brains at E14.5 (A, B), E16.5 (C, D), and E17.5 (E-J). Phospho-JNK was distributed in the soma and neuritic processes of neuronal cells in broad regions of developing *jsap1*^{+/+} and *jsap1*^{-/-} brains. Anti-phospho-JNK labeled neuronal cells in the marginal zone, subplate, intermediate zone, subventricular zone, caudoputamen, and cingulum bundles, and also labeled radial glial fibers in the cortical plate (G, H). G and H are magnified images of the rectangles in the insets. The expression level of phospho-JNK in the U-shaped axon tract of the corpus callosum (C-F), and anterior commissure (I, J) in *jsap1*^{-/-} brains was lower than that in *jsap1*^{+/+} brains. (K) Quantification of the expression levels of phospho-JNK in indicated brain regions in both genotypes. * $P < 0.05$ and ** $P < 0.01$, differences between *jsap1*^{+/+} and *jsap1*^{-/-} brains (Student's *t*-test). AC, anterior commissure; CC, corpus callosum; cng, cingulum bundle; CP, caudo-putamen; CxP, cortical plate; ic, internal capsule; IZ, intermediate zone; LV, lateral ventricle; MZ, molecular zone; Pb, Probst's bundle; SP, subplate; tca, thalamocortical axons; VZ, ventricular zone. A, C, E, and G; *jsap1*^{+/+}. B, D, F, and H; *jsap1*^{-/-}. Scale bars: A-F, 250 μ m; G and H, 100 μ m.

bundles in the cortical plate in both genotypes (Figures 5C-5J). Radial glial fibers, which function in neuronal migration and laminar patterning of developing cortex (Parnavelas, 2000; Ever and Gaiano, 2005), were also labeled by anti-phospho-JNK in similar levels in both genotypes (Figures 5E-5H). Whereas phospho-JNK levels in the subplate and the cingulum bundle in E16.5- E17.5

jsap1^{-/-} brains were lower than those in *jsap1*^{+/+}. Anti-phospho-JNK also labeled the callosal axons crossing the midline in the E16.5-E17.5 brains in both genotypes, though the staining levels in *jsap1*^{-/-} were lower than those in *jsap1*^{+/+} (Figures 5C-5F and 5K). The expression level of anti-phospho-JNK in the anterior commissure was also decreased in E17.5 *jsap1*^{-/-} (Figures 5I-5K). At E17.5, anti-phospho-JNK stained callosal axon tracts were disrupted and formed Probst's bundle-like structures bilaterally in the cingulate cortex (Figure 5F). Quantification of immunoreactive images revealed that the immunoreactivity of phospho-JNK in *jsap1*^{-/-} brains was reduced in all brain regions examined compared with those in *jsap1*^{+/+} (Figure 5K). In E14.5-E17.5 brains of both genotypes, JNK1 was expressed in most brain regions, and the distribution of JNK1 was overlapped with that of phospho-JNK (data not shown).

Because JSAP1 knockout mice showed reduction in the cortical thickness of the brain (Ha *et al.*, 2005) and overexpression of JSAP1 increased JNK activity and apoptosis following ischemia-reperfusion (Xu *et al.*, 2010), we investigated whether the JSAP1 deficiency affect cell death during development. TUNEL staining showed that TUNEL-positive cells in each brain section were distributed in many brain regions, including the frontal cortex, dorsal striatum, and the areas around the lateral ventricles (data not shown). TUNEL-positive cells distributed around the corpus callosum at the midline area of the brain of *jsap1*^{-/-} were tended to be reduced in numbers compared to *jsap1*^{+/+} (Supplemental Data Figure S1), although this study did not examine the possibility that such reduction was due to the delayed progression of development in *jsap1*^{-/-} compared to *jsap1*^{+/+}.

Discussion

We recently reported that the JSAP1 deficient E18.5 brain exhibits various types of developmental defects in the brain, including impaired axon guidance of the CC and AC (Ha *et al.*, 2005). The CC is the largest axon bundle that connects many areas between the two hemispheres, including the inferior frontal lobes, deeper limbic structures, supplementary motor cortices and cingulate gyri (Abbie, 1940). Early callosal axons originating from the cingulate cortex (cingulate cortex-originated axons) pioneer and cross the midline of the telencephalon during E15.5-E16.5, and late arriving collaterals of the CC that originate from other regions of the neocortex (neocortex-

originated axons) follow the pre-existing axon tract, and join to form the typical U-shaped axonal tract (Koester and O'Leary, 1994; Ozaki and Wahlsten, 1998; Rash and Richards, 2001). As described, the early axonal extension of the CC in *jsap1^{-/-}* was detected during E15.5-E16.5 (Figure 1). A careful examination of the callosal axons in the E16.5 brain revealed that some collaterals of these early axons lost U-shaped axonal integrity, though their tract trace was identified in all cases, and that the remaining axon tract, as examined at E16.5, was lightly stained by anti-phospho-JNK or anti-phospho-FAK (Figures 4 and 5). In contrast, the axon tract of the CC in E17.5 *jsap1^{-/-}* brain was interrupted at the midline, and instead formed Probst's bundle-like structures in the cingulate cortex bilaterally (Figures 1G and 1H; Figures 5E and 5F). The disrupted tract formation of the CC is likely to be due to the degeneration of early developed axons, and the pathfinding defect of late developing axons. Thus our results suggest that the JSAP1-regulated signal pathway may play a role in the development of the telencephalic commissure, particularly for axon pathfinding and probably the maintenance of the early tract.

The finding that the callosal axons in *jsap1^{+/+}* were strongly labeled by antibodies for the active forms of FAK and JNK (that is, phospho-JNK and phospho-FAK, respectively) during E14.5-E17.5 (Figures 4 and 5). Moreover, the expression levels of phospho-JNK and phospho-FAK in developing callosal axons in *jsap1^{-/-}* were lower than those in *jsap1^{+/+}*, and this reduction was correlated with the disruption of axon tract formation. In consistent with these findings, previous studies have shown that FAK and JNK pathways are involved in axonal or dendritic development. For examples, targeted neuron-specific FAK deletion alters dendritic morphology in the developing brain (Beggs *et al.*, 2003; Shi *et al.*, 2009), and JNK deletion induces disrupted tract formation of the AC and the progressive loss of microtubules within axons and dendrites in neonate brains, although the CC was unaffected (Chang *et al.*, 2003). Regarding the JSAP1-dependent mechanism of axon development, it is of interest that the JSAP1 deficiency resulted in reduced levels of phospho-FAK and phospho-JNK in developing callosal axons, but it did not completely abolish their expression. These results suggest that FAK and JNK signaling pathways have a role in the development and/or maintenance of this axon tract, though it remains to be determined whether the JSAP1-dependent down-regulation of the FAK or JNK pathways is the primary cause of defective telencephalic commissure development. We recently demonstrated that

mouse embryonic fibroblasts (MEF cells) prepared from *jsap1^{-/-}* embryos showed a whole sequence of cell spreading and cell adhesion on culture plates like *jsap1^{+/+}* MEF cells, but they look significantly longer to go through the sequence of cell spreading and adhesion-induced FAK activation was slightly reduced. These *jsap1^{-/-}* MEFs showed the reduced cell spreading and cell adhesion properties in *in vitro* cell adhesion assay (Chae *et al.*, 2006). Moreover, in a previous study, we reported that axon guidance defects of the CC and AC, and the reduced levels of phospho-FAK and phospho-JNK in the JSAP1-deficient brain were partially rescued by the transgenic expression of JIP1/SKIP (Ha *et al.*, 2005), although JSAP1 and JIP1/SKIP are distinct proteins. Thus, it is likely that overlapping components of the JSAP1- and JIP1-dependent functional networks play a role in axon guidance, which suggest that JSAP1-like proteins play a compensatory role in unaffected neurons of *jsap1^{-/-}*, and that they might participate to an extent even in affected neurons, and mitigate JSAP1-deficient phenotypes. The developmental phenotypes displayed by the *jsap1^{-/-}* brains are highly complex, yet available information gathered from molecular biologic and knockout mouse studies is too limited to explain the observed phenotypes. Although JSAP1-dependent regulation of FAK and JNK occurs and their proper regulation might be important for the tract formation of the CC, further work is needed to reveal the mechanisms by which the FAK and JNK pathways regulate axon tract maintenance and axon pathfinding of the CC.

Because JSAP1 is known to function as a scaffold protein for FAK and JNK (Whitmarsh *et al.*, 1998; Yasuda *et al.*, 1999; Takino *et al.*, 2002), we were interested in determining whether JSAP1 regulates the subcellular localization of FAK and JNK in neuronal cells and their subcellular distributions are correlated with developmental defects. We found no evidence that the JSAP1 null mutation restricts the distribution of FAK and JNK in axons. Consequently, though the *jsap1^{-/-}* brain showed reduced phospho-JNK and phospho-FAK expression in the callosal axons, it is not yet known whether JSAP1 is required for the activation of FAK and JNK, or for the sustained maintenance of activated FAK or JNK.

The impaired axon guidance of the late arriving CC collaterals in the *jsap1^{-/-}* brain is particularly interesting because these axons showed pathfinding defects. Previous studies have reported that mice lacking growth cone-related genes, such as GAP-43 and MAP1B, display impaired axon guidance of the CC and the AC (Richard, 2002). JSAP1 forms a complex with FAK (Takino *et al.*,

2002), which is also a component in growth cones (Burgaya *et al.*, 1995), and JSAP1 is strongly expressed in growth cones and affects neurite outgrowth (Xu *et al.*, 2003). Therefore, it will be interesting to determine whether JSAP1 interacts any of these growth cone components or whether the intracellular signals essential for growth cone function are commonly regulated by JSAP1 and these growth cone components. Phospho-JNK positive TCAs running through the internal capsule also showed pathfinding defects, and projected with an aberrant lateroventral, rather than dorso-lateral, trajectory (Figure 5F). Subplate neurons and their fibers are known to play a role in the guidance of cortical afferents, such as thalamo-cortical afferents (Zhou *et al.*, 1999; Soria and Fairén, 2000; Deng and Elberger, 2001; Shinozaki *et al.*, 2002). Though it is not known whether the pathfinding defects of callosal axons and TCAs are due to cell autonomous or not, organized regulation of cell adhesion-related factors by JSAP1 might play a role.

The present study and our previous study (Ha *et al.*, 2005) together show that almost all neuro-anatomical abnormalities, such as impaired axon guidance of the CC and AC, the pathfinding defects of thalamocortical tracts and migration defects of anti-calretinin positive cells in the developing cerebral cortex, with the exception of calretinin transport along axons (Ha *et al.*, 2005), are primarily related to various forms of defective cell adhesion. Thus, our results strongly support the notion that JSAP1 regulates signal factors essential for cell adhesion *in vivo*. Given the known functions of JNK and FAK pathway in neurite outgrowth and cell migration, proper distribution and formation of functional modules of JNK and FAK pathway components might be essential to afford specificity and efficiency of these signaling pathways, and JSAP1 may participate in these processes.

Methods

JSAP1 deficient embryos

JSAP1 knockout mice were described in a previous study (Ha *et al.*, 2005). JSAP1 deficient embryos were obtained by intercrossing JSAP1 knockout heterozygotes, and the genotypes of individual embryos were determined by PCR using tail tip DNA, as previously described (Ha *et al.*, 2005).

Preparation of brain sections

Fetuses were removed from dams by caesarian section at

embryonic ages E12.5 to E18.5. To prepare paraffin-embedded sections, brains were quickly removed, fixed in fresh 4% paraformaldehyde in 0.1 M phosphate buffer (pH 7.4) at 4°C overnight, dehydrated through a graded alcohol series, cleared with xylene, and embedded in Paraplast (Oxford, UK). Serial 4 µm coronal sections were then prepared, and subjected to cresyl violet staining to examine cellularity. For immunohistochemistry, brains were fixed in 4% paraformaldehyde at 4°C overnight and equilibrated in 10%, 20%, and 30% sucrose solutions and then frozen in OCT compound (Tissue Tek, Elkhardt). Brains were sectioned coronally into 20 µm sections using a cryostat (CM1850, Leica Ins., Germany). Sections were immunologically stained after mounting onto gelatin-coated subbed slides or following a floating method. The numbers of *jsap1*^{+/+} or *jsap1*^{-/-} embryos examined in detail at each age were as follows: E14.5, *n* = 4 and 5; E15.5, *n* = 2 and 2; E16.5, *n* = 10 and 10; E17.5, *n* = 10 and 10; E18.5, *n* = 13 and 15.

Immunohistochemistry and quantification

For immunohistochemistry, sections were incubated for 30 min with 3% H₂O₂ in 0.1 M PBS (pH 7.4) to remove endogenous peroxidase activity, and washed in PBS. Sections were then blocked with a solution containing 5% normal goat/or horse serum, 2% BSA, 2% FBS and 0.1% triton X-100 for 2 h at room temperature (RT). Primary antibody in blocking buffer was then added to sections, and sections were incubated overnight at 4°C and washed in PBS. They were then incubated with biotinylated secondary antibody (Vector Laboratories, Burlingame, CA) at a dilution of 1:200 for 1 h at RT, avidin and biotinylated HRP complex (Vector Laboratories) at 1:200 for 1 h at RT, and visualized with 0.05% 3,3'-diaminobenzidine and 0.001% H₂O₂ in 0.1 M Tris, pH 7.4. After counterstaining, the brain sections were dehydrated and cover-slipped using a standard procedure.

Primary antibodies used were as follows: rabbit anti-phospho-JNK (Biosource, CA), rabbit anti-phospho-FAK (Y397) (Biosource, CA), mouse anti-FAK (Santa Cruz, CA), mouse anti-JNK1 (Pharmingen, San Diego, CA) and rabbit anti-calretinin (Chemicon International Inc. Temecula, CA), mouse anti-neuronal nuclei (NeuN) (Chemicon), and anti-MAP-2 (Upstate, Lake placid, NY).

For quantification, immunoreactive images were uploaded on the Adobe Photoshop software (Adobe Systems, San Jose, CA). The pixel density of immunoreactivity was measured using Image J software (<http://rsbweb.nih.gov/ij/index.html>) in a blinder manner, with a modification of the quantification method described previously (Kopniczky *et al.*, 2005). Briefly, open circular cursors with a diameter of 50, 100, or 200 µm were placed on the corpus callosum (6 circles), cingulum bundle (3 circles), ventricular zone (5 circles), intermediate zone (5 circles), subplate (5 circles), cortical plate (5 circles), caudo-putamen (4 circles) and anterior commissure (5-10 circles), as illustrated in Figures 4 and 5. The average of 10 background determinations performed near the brain areas being counted was subtracted from the average pixel densities measured for the brain regions to study.

In situ detection of fragmented DNA (terminal deoxynucleotidyl transferase-mediated UTP nick end labelling, TUNEL)

The fragmentation of DNA was examined using an ApopTag[®] Peroxidase In situ Apoptosis Detection Kit (S7100) (Millipore, MA) according to the manufacturer's instructions. Briefly, after deparaffinization and hydration using xylene and graded alcohol, brain sections were placed to enzymatic digestion with 20 µg/ml of proteinase K for 5 min, treated with 5% hydrogen peroxide for 20 min to exhaust endogenous peroxidase activity, and washed with phosphate-buffered saline (PBS, 0.1 M, pH 7.4). They were then immersed in an ApopTag[®] Equilibration Buffer to label the 3'-OH ends of fragmented DNA for 10 min and incubated with terminal deoxynucleotidyl transferase enzyme at 37°C for 1 h. After washing with PBS, sections were incubated with anti-digoxigenin conjugated peroxidase and the peroxidase substrate (diaminobenzidine) to detect signs of apoptotic cell death.

Statistical analysis

Differences between two groups were assessed using a Student's *t*-test, at a minimum confidence level of $P < 0.05$.

Supplemental data

Supplemental data include a figure and can be found with this article online at http://e-emm.or.kr/article/article_files/SP-43-8-04.pdf.

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