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# **Phosphoinositide 3-kinase enhancer (PIKE) in the brain: is it simply a phosphoinositide 3-kinase/Akt enhancer?**

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### **Abstract**

Since its discovery in 2000, phosphoinositide 3-kinase enhancer (PIKE) has been recognized as a class of GTPase that controls the enzymatic activities of phosphoinositide 3-kinase (PI3K) and Akt in the central nervous system (CNS). However, recent studies suggest that PIKEs are not only enhancers to PI3K/Akt but also modulators to other kinases including insulin receptor tyrosine kinase and focal adhesion kinases. Moreover, they regulate transcription factors such as signal transducer and activator of transcription and nuclear factor κB. Indeed, PIKE proteins participate in multiple cellular processes including control of cell survival, brain development, memory formation, gene transcription, and metabolism. In this review, we have summarized the functions of PIKE proteins in CNS and discussed their potential implications in various neurological disorders.

#### **Keywords**

Akt; GTPase; neuron; PI3K; PIKE

# **Introduction**

Phosphoinositide 3-kinase enhancer (PIKE) is a group of GTP-binding proteins that belong to the α 1 subgroup of centaurin GTPase family (Jackson et al., 2000). Through alternative splicing, three PIKE isoforms, designated as PIKE-S, PIKE-L, and PIKE-A, are generated from CENTG1 (Chan and Ye, 2007). They are multifunctional proteins that control diverse cellular activities including apoptosis, cell migration, transformation, receptor and endosome traffickings, gene transcription, and metabolism (Rong et al., 2003; Ahn et al., 2004a; Liu et al., 2007a; Tang et al., 2008; Zhu et al., 2009; Chan et al., 2010a,b, 2011b; Shiba et al., 2010). The core structure of PIKE comprises of a central GTPase domain, followed by a pleckstrin homology (PH) domain, a domain of GTPase-activating protein of Arf, and several ankyrin repeats (Chan and Ye, 2007). As a member of the GTPase superfamily, PIKE proteins share several common characteristics with other GTP-binding proteins. First, they possess intrinsic GTP hydrolysis activity (Ye et al., 2000). Nonetheless, PIKE GTPase domain demonstrates a nondifferential and highly eficient hydrolysis on GTP, ATP, UTP, and CTP (Soundararajan et al., 2007). Second, their GTPase activities can be augmented by interacting with specific guanine exchange factor. We have found that the PIKE GTPase activity was increased in the presence of phospholipase C  $\gamma$ 1 (PLC $\gamma$  1) (Ye et al., 2002). Furthermore, the cellular localization and the enzymatic activity of PIKE could be regulated by phosphoinositides (PIs). Several centaurin family members such as centaurin α binds to

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PI with high affinity to initiate its cell membrane tethering (Hammonds-Odie et al., 1996). We found that the PH domain of PIKE-L robustly bound to phosphatidylinositol trisphosphate (PIP<sub>3</sub>) (Hu et al., 2005). Yan et al. (2008) also reported that both PIKE-A and -L bound to the head groups of di- and triphosphoinositides with similar affinities. In addition, the PH domain which is responsible for lipid binding also functions to localize PIKE-L to the plasma membrane (Yan et al., 2008).

In the last decade, our research group endeavored to delineate the physiological roles of PIKE proteins, especially their functions in neurons. Using a combination of *in vitro* and *in* vivo models, we have discovered the antiapoptotic function of PIKE proteins by modulating the PI3K activity. With the availability of the whole body  $PIKE$  knockout ( $PIKE-/-$ ) mice, we surprisingly find that PIKE proteins are implicated in multiple signaling pathways in addition to the phosphoinositide 3-kinase (PI3K)/Akt cascade. In this review, we will discuss the functions of PIKE in the central nervous system (CNS) and how their functions are engaged in various neurological disorders.

#### **PIKE as a PI3K enhancer**

PI3K is a group of enzymes that phosphorylate the inositol ring of phosphatidylinositol to generate a number of structurally different PIs (Wymann and Pirola, 1998). PIs are important secondary messengers that participate in a great variety of functions in the CNS including cell survival, apoptosis, neuronal migration, neurotransmission, and morphological development of neurons (Katso et al., 2001; Deane and Fruman, 2004; Cosker and Eickholt, 2007; Waite and Eickholt, 2010; Yamazaki et al., 2010). While most attentions on PI signaling have been paid on their activities at the plasma membrane, it is now clear that nucleus has a distinct set of PI signaling machinery and effectors with different regulatory mechanisms (Martelli et al., 2006, 2007). In 2000, our research group isolated PIKE-S (the first isoform of PIKE) from PC12 cells, which is a well-established model for studying neuronal differentiation (Ye et al., 2000). It is a neuronal specific protein that associates with class I PI3K in the nucleus. Indeed, PIKE-S is responsible for initiating the PIP<sub>3</sub> production in the nucleus after nerve growth factor (NGF) stimulation (Ye et al., 2000). Once NGF receptor TrkA is activated, it induces the translocation of PLC  $\gamma$ 1 from the cytoplasm to the nucleus, where it interacts with PIKE-S and enhances its GTPase activity (Ye et al., 2002). PIKE-S then augments the nuclear PI3K activity in a GTPasedependent manner by binding to the regulatory subunit (p85) of PI3K, presumably stabilizes the effect of p85 on p110. Nuclear Akt is consequently activated, which in turns phosphorylates and interacts with other proteins (e.g., nucleoplasmin) to execute the antiapoptotic function of NGF (Ahn et al., 2004c, 2005). In addition, intact GTPase activity of PIKE-S is necessary for cyclin-D1 expression in PC12 cells as overexpressing GTPasemutated PIKE-S abolishes NGF-induced cyclin D1 expression (Ye et al., 2000). Thus, PIKE-S is a novel nuclear effector of NGF to protect against apoptosis, induce cell cycle arrest, and initiate differentiation.

# **Association of PIKE with glutamate receptors in neurons**

Using cDNA library screening, we have cloned the second PIKE isoform (PIKE-L) from the rat brain cDNA library. In contrast to the nuclear residency of PIKE-S, PIKE-L is enriched in the postsynaptic densities (PSD) (Rong et al., 2003). As PSD is a region at the membrane of the postsynaptic neuron, where it concentrates and organizes neurotransmitter receptors for efficient transmission of neuronal signals, it is reasonable to find that PIKE-L functionally couples with various glutamate receptors such as α-amino-3-hydroxy-5 methyl-4-isoxazolepropionate receptor (AMPAR) and metabotropic glutamate receptor I  $(mGlu<sub>1</sub>)$ . Glutamate is the neurotransmitter for fast excitatory transmission (Greenamyre,

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1986). However, while excessive glutamate is a major inducer for excitotoxicity-induced cell death (Lau and Tymianski, 2010), it can also rescue neuronal injury by activating mGlu<sub>1</sub> and AMPAR in a PI3K-dependent fashion (Copani et al., 1995; Wu et al., 2004; Nishimoto et al., 2008; Lau and Tymianski, 2010). PIKE-L couples to both mGlu<sub>1</sub> and AMPAR by distinct mechanisms. Upon mGlu<sub>1</sub> activation, PIKE-L associates with mGlu<sub>1</sub> through Homer Ic (Rong et al., 2003). Homer is a group of scaffold proteins for protein-protein interaction, which link mGlu<sub>1</sub> to inositol 1,4,5-triphosphate receptor for the increase of intracellular calcium (Tu et al., 1998). By forming a complex with Homer Ic and PIKE-L, the activated mGlu<sub>1</sub> can transmit the signal to cytosolic PI3K and subsequently protects the neurons from staurosporine-induced cell death (Rong et al., 2003). In contrast to linking the mGlu<sub>1</sub> via an adaptor protein, PIKE-L interacts with the GluA2 subunit of AMPAR directly (Chan et al., 2011b). This receptor/PIKE association is not constitutive; instead, it depends on the activation of another ionotropic glutamate receptor N-methyl D-aspartate receptor (NMDAR). When NMDAR is activated by its co-agonist glycine, the AMPAR-associated PI3K will be induced, possibly by the influx of  $Ca^{2+}$  and the subsequent activation of calmodulin kinase II (Man et al., 2003). The activated PI3K on the AMPAR complex then provides a signal to recruit PIKE-L to GluA2, which further potentiates the activity of the receptor-linked PI3K, thus leading to the membrane delivery of AMPAR and surface retention (Chan et al., 2011b). In addition, PIKE-L functions as a molecular linkage between GluA2 and glutamate receptor interacting protein 1 (GRIP1) (Chan et al., 2011b). As GRIP1 is a critical protein for AMPAR surface expression (Dong et al., 1997; Hoogenraad et al., 2005), the increased association between PIKE-L and GluA2 in response to glycine stimulation thus provides an additional docking site for GRIP1 to the AMPAR complex to further facilitate its cell surface anchorage. It is interesting to note that 4.1N, an interaction partner of PIKE-S (Ye et al., 2000), associates with AMPAR subunit GluA1 and anchors the AMPAR on the plasma membrane by coupling the receptor to the cytoskeleton (Shen et al., 2000; Lin et al., 2009). It is unknown if PIKE-L/4.1N interaction has any physiological significance on the AMPAR functions; nevertheless, it is clear that PIKE-L is an important effector in the glutamate receptors complex.

#### **Binding of PIKE to netrin receptors in neurons**

We have also found that PIKE-L is a component of the netrin 1 signaling to protect the neurons from apoptosis (Tang et al., 2008). Netrin 1 is a chemotropic cue for cells and axon migration, which also controls the axon arborization and synapse formation during neural development by acting on its cognate receptors deleted in colorectal cancer (DCC) or UNC-5 homologs (UNC5) (Lai et al., 2011). In addition, it is a survival factor to neurons in the ventricular zone of brainstem, which express UNC5 and DCC (Llambi et al., 2001). In cultured hippocampal neurons, netrin-1 stimulation protects the cells from glutamateprovoked apoptosis (Tang et al., 2008). However, the protection is abolished if PIKE-L is depleted, suggesting that PIKE-L is an essential downstream effector of netrin-1 to mediate its protective functions. Further studies on the functional interactions between PIKE-L and UNC5B indicate that a molecular complex of these two proteins is formed after netrin-1 stimulation, which induces the activation of PI3K subsequently (Tang et al., 2008). The PIKE-L/ UNC5B association is controlled by Fyn, a protein kinase that phosphorylates the tyrosine residues of numerous proteins, and Fyn phosphorylations on the receptor and PIKE-L are necessary for their interaction (Tang et al., 2008; Saito et al., 2010). As Fyn is constitutively associated with DCC but not UNC5, presumably, PIKE-L may not solely interact with UNC5 but also tethers to the UNC5/DCC heteroreceptor (Mille et al., 2009). Indeed, PIKE-L and DCC could be co-immunoprecipitated from the rat brain lysates, which further provides confidences on this hypothesis (Tang et al., 2008).

### **PIKE as an** *in vivo* **neuroprotectant**

The above in vitro studies indicate that PIKE-L has a neuroprotective role against neurotoxic insults in cultured neurons via PI3K activation, but the role of PIKE in vivo has not been well explored. In intact animals, the concentrations of extra-neuronal glutamate are elevated during acute damages such as status epilepticus, mechanical trauma, or ischemia as a result of cellular leakage or depolarization-induced exocytosis (Lau and Tymianski, 2010). The raise of glutamate concentration leads to intensive activations of ionotropic glutamate receptors, causing excessive calcium influxes to the neurons and triggering apoptosis (Szydlowska and Tymianski, 2010). It is reported that the process involves multiple pathways including activation of nitric oxide synthase, calcium-sensitive proteases, caspases, and mitochondrial damage (Wang and Qin, 2010). As PI3K/Akt is the central pathway against apoptosis, the low PI3K/Akt activity in the cortex of PIKE−/− mice may make the animals more susceptible to neurotoxic damages (Chan et al., 2011d). As anticipated, the knockout animals developed larger infarct volume with higher number of cells perusing apoptosis after middle cerebral artery occlusion (Chan et al., 2011d). A similar observation was found in mice injected with kainic acid (KA), where higher caspase 3 activity and poly (ADP-ribose) polymerase cleavage were detected in the PIKE−/− hippocampus (Chan et al., 2011a). Interestingly, KA stimulation induces more  $Ca^{2+}$  infl ux into PIKE−/− neurons, which results in a lower threshold toward KA-induced seizure (Chan et al., 2011a). In vitro culture of PIKE−/− neurons also reveals a high vulnerability under glutamate stimulation, suggesting that PIKE-L is necessary for cell protection during stroke and glutamate challenges both *in vitro* and *in vivo* (Chan et al., 2011a).

The PI3K-enhancing activity by PIKE is not only critical for protecting neurons from pathological damage; it is also necessary for normal brain development. During embryonic growth, markedly enhanced apoptosis is observed in the nestin-positive progenitor cells in the ventricular zone of PIKE−/− neocortex (Chan et al., 2011d). However, a comparable density of proliferating cells is detected between wild-type and PIKE−/− brain at the same developmental stage, indicating that neuronal survival but not proliferation is affected by PIKE depletion. The apoptosis in cortical neurons is sustained beyond the developmental stages as we observed positive caspase-3 staining in the somatosensory cortex of adult PIKE −/− mice but in not wild-type controls. As a result, the number of neurons in the neocortex is reduced in PIKE-null animals, leading to a thinner cortical layers and reduced brain mass (Chan et al., 2011d).

# **PIKE as a kinase regulator**

Akt is a downstream kinase of PI3K. It is a serine/threonine kinase that expresses ubiquitously in numerous tissues. In 2004, we isolated the third isoform of PIKE (PIKE-A) when we characterized glioblastoma multiforme (GBM). PIKE-A displays an elevated expression in glioblastomas and astrocytomas (Ahn et al., 2004b). In addition, it is highly expressed in a variety of cancers which originate from breast, ovary, kidney, prostate, skin, uterus, and colon, suggesting that PIKE-A is an ubiquitous proto-oncogene (Liu et al., 2007a). Structural analysis of PIKE-A shows that it does not contain the N-terminal domain in PIKE-S and PIKE-L for PI3K association, indicating that it might not act as a PI3K enhancer. As Akt is highly active in many cancer cells, we thus hypothesized that PIKE-A may be a novel Akt partner to modulate its kinase activity. As expected, overexpression of PIKE-A in cells enhanced the growth factor-induced Akt activation in a GTPase-dependent manner and depletion of PIKE-A abolished EGF-initiated Akt activity (Liu et al., 2007a). We also observed a diminished Akt activity in PIKE−/− brain (Chan et al., 2011b). However, it is difficult to distinguish if such Akt reduction is a direct consequence of PIKE-

A depletion, or a secondary effect of PIKE-L knockout as the upstream PI3K activity is impaired in the absence of PIKE in intact brain (Chan et al., 2011d).

To date, the majority of researches on PIKE-A focus on its role in neoplasia and little is known about its role in neurons. Given its high expression levels in the brain, it is believed that PIKE-A has a significant contribution to the neuronal function (Nagase et al., 1996; Xia et al., 2003). Nie et al. (2005) reported that PIKE-A interacts with the clathrin adaptor protein AP1 and controls its cellular localization. Moreover, PIKE-A overexpression redistributes AP1 from the perinuclear region to punctuate structure throughout the cells and decreases the transferring recycling (Nie et al., 2005). As AP1 is responsible for proper synaptic endocytosis in neurons, it is tempting to speculate that PIKE-A also functions as a part of the coated vesicles during the endosomes- trans Golgi network transportation for neurotransmitters (Glyvuk et al., 2010; Shiba et al., 2010). Indirect evidence also suggests that PIKE-A might be involved in neurotransmitter release/uptake. McFarland et al. (2008) has reported that PIKE-A is a protein found in the syn-aptosome. The authors also demonstrated that PIKE-A is one of the association partners to the C-terminus of α synuclein, a neuronal protein that localizes in the presynaptic terminals to inhibit synaptic vesicle exocytosis and neurotransmitter release (Larsen et al., 2006; McFarland et al., 2008). However, their studies provide neither mechanistic insights on the PIKE-A/ $\alpha$ -synuclein interaction nor any functional consequences of the association. Thus, it remains to be determined if PIKE-A plays any role in neurotransmitter exocytosis.

PIKE-A has also been reported as a focal adhesion kinase (FAK) enhancer. Zhu et al. (2009) show that PIKE-A is a novel interaction partner of FAK, which enhances its kinase activity in response to growth factor stimulation. The authors also speculate that PIKE-A regulates focal adhesion dynamics by at least two mechanisms: (1) it activates FAK to disassemble focal adhesion and (2) controls the trafficking of regulators to or from the focal adhesion when the focal adhesion is being remodeled. As a major component of focal adhesion and involved in actin organization, it is not surprising to know that FAK is implicated in dendrite formation, axon outgrowth, and neuronal migration, which require proper cytoskeleton remodeling (Kawauchi and Hoshino, 2008). Although the functional interaction between PIKE-A and FAK has not been tested in neurons, several lines of indirect evidence might provide hints on these protein interplays in modulating neuronal migration. It has been reported that phosphorylation of FAK by cyclin-de-pendent kinase 5 (Cdk5) is critical for FAK-mediated microtubule fork formation and neuronal migration in the developing cortex (Xie et al., 2003). However, Cdk5 phosphorylation does not affect the catalytic activity of FAK. As activating the kinase activity of FAK to disrupt the focal adhesion is a critical step for cell migration, it is suggested that phosphorylation of FAK by Cdk5 affects FAK ' s ability to assemble a particular protein complex in mediating cell migration (Ilic et al., 1997; Nikolic, 2004). Given that PIKE-A is also a physiological substrate of Cdk5, which enhances the binding affinity of PIKE-A toward other protein partners, conceivably, PIKE-A might regulate FAK activity after Cdk5 activation (Liu et al., 2008).

Recently, we have discovered that PIKE-A is a novel regulator of insulin receptor tyrosine kinase (IRTK) activity. The association of PIKE-A with an insulin receptor (IR) is important for insulin to fully initiate the hepatic IRTK (Chan et al., 2011c). We also found that the brain IR could not be fully activated in PIKE−/− mice after systemic insulin injection, suggesting the IR/PIKE-A interaction is also valid in neuronal tissues (data not shown). Insulin has profound effects in the CNS to regulate food intake, energy homeostasis, reproduction, and neuronal survival (Plum et al., 2005). Our studies in PIKE−/− mice suggest that PIKE-A is not involved in all of insulin functions in the CNS. For example, the food intake in PIKE−/− mice is comparable to their wild-type littermates (Chan et al., 2010b). Moreover, PIKE−/− mice are fertile with no significant differences in the number of

pups born from knockout dams, although the mammary gland defect in the PIKE−/− mothers causes a low survival rate of newborns (Chan et al., 2010a). Nevertheless, *PIKE*−/− mice are vulnerable to KA-induced neuronal death and seizure, which is in good consensus with the neuroprotective role of insulin, as insulin can protect neurons from apoptosis caused by AMPAR overactivation and KA-induced tonic-clonic convulsions (Uysal et al., 1996; Ryu et al., 1999; Kim and Han, 2005; Chan et al., 2011a). Presumably, PIKE-A is selectively implicated in a portion of IR-expressed neurons in the CNS to mediate insulin ' s neuronal survival actions but not energy homeostasis or reproduction.

# **PIKE and neurological disorders**

Loss of neurons during pathological conditions such as stroke or traumatic injuries could be devastating, as neurons have limited regenerative capacity. Depending on the regions affected, neuronal loss in brain causes impaired cognitions, memory, locomotion, and homeostasis. Neuronal death has also been attributed as a major feature implicated in various neurodegenerative diseases such as Alzheimer ' s disease (AD), Parkinson ' s disease, and Huntington ' s disease (Dong et al., 2009). Thus, understanding the mechanisms to cope with toxic insults in neurons is extremely important for clinically preserving the integrity of CNS. Observations over the past decades have identified that PI3K cascade is one of the central pathways to promote neuronal survival during development and after neurotoxic insults (Yuan and Yankner, 2000; Waite and Eickholt, 2010), but the mechanisms controlling the temporal and spatial activation of the protective PI3K/Akt pathway are not well understood. Studies have suggested that neurotro-phins such as brainderived neurotrophic factor (BDNF) are the key molecules to initiate the neuroprotective mechanism during these catastrophic damages. It has been reported that BDNF protects against glutamate-induced apoptotic cell death via PI3K and extracellular signal-regulated kinase pathways in vitro (Almeida et al., 2005). Moreover, in pathological conditions such as forebrain ischemia, infusion of BDNF shortly after stroke can effectively reduce total infarct volume (Yamashita et al., 1997). Furthermore, increased BDNF expression after ischemia and seizure induction is suggested as a protective mechanism against excessive neuronal death (Tsukahara et al., 1994; Kokaia et al., 1995, 1998). Insufficient BDNF production or response has also been proposed as a significant contributor to chronic neurodegenerative diseases. For example, the amount of BDNF is reduced in the cortex and hippocampus of AD patients (Narisawa-Saito et al., 1996). Studies in TrkB knockout mice demonstrate that impairing the BDNF signaling provokes a negative outcome in learning and memory – the two features which are markedly affected in AD (Minichiello et al., 1999). Most importantly, BDNF has neuroprotective effects against Aβ peptide toxicity *in vitro* and *in vivo* – the key molecules leading to the neural damage in AD models (Liu et al., 2007b; Arancibia et al., 2008). We have demonstrated that the presence of PIKE is necessary for BDNF to fully execute its neuroprotection activity as *PIKE*−/− neurons are ' partially ' resistant to BDNF stimulation (Chan et al., 2011d). Moreover, PIKE−/− mice shared striking phenotypic similarities with cortex-specifi c BDNF/TrkB knockout mice – both of them have compressed cortical thickness in layer II/III, diminished soma size of cortical neuron, decreased cortical neuron number, reduced dendritic complexity, enhanced locomotor activity, and impaired memory, implying that PIKE is the downstream effector of BDNF (Xu et al., 2000; Gorski et al., 2003; Chan et al., 2011d). Although there is no report on the genetic correlation between CENTG1 polymorphism and various neurodegenerative diseases in human, the animal studies have provided strong evidence that PIKEs are intrinsic protective molecules against various neurotoxic challenges.

#### **PIKE in memory formation**

The integrity of PIKE is also important for memory formation. In Morris water maze test, *PIKE−/*−animals showed poor performance to locate the platform, although the animals possess comparable learning behavior as their wild-type control (Chan et al., 2011d). Similar results were observed in the Y-maze test in which PIKE−/− mice had difficulties to identify the route which has already been explored (Chan et al., 2011d). The molecular mechanism of this memory defect in the absence of PIKE is not clear but our data suggest two possible explanations. First, we observed that PIKE−/− neurons had defective long-term potentiation (LTP), which is a result of lower GluA2-containing AMPAR density on cell surface (Chan et al., 2011b). AMPAR and NMDAR are critical ion channels in modulating the synaptic plasticity and memory formation in the hippocampus (Kullmann et al., 2000; Rao and Finkbeiner, 2007). In particular, the activity and distribution of AMPAR in the synapse is crucial for LTP expression (Isaac et al., 2007). The insufficient GluA2-containing AMPAR on the cell surface of PIKE−/− neurons thus hinders the LTP expression and memory formation. Second, PIKE−/− neurons have less dendritic arbors (Chan et al., 2011d). Development of highly branched dendrites is essential for establishing functional connections between neurons. Several researches have shown that dendritic patterning of pyramidal cells in cerebral cortex is escalated after learning (Turner et al., 2003; Kolb et al., 2008; Gelfo et al., 2009). When PI3K/Akt pathway is augmented in PIKE−/− neurons by over-expressing constitutively active Akt or p110 subunit of PI3K, the arborization pattern is rescued (Chan et al., 2011d). We extended the studies by generating the PIKE−/−PTEN−/− double mutant animals to mend the defective Akt signaling in vivo. PTEN is a phosphatidylinositol-3,4,5-trisphosphate 3-phos-phatase that antagonizes the PI3K/Akt pathway (Myers et al., 1998). In PTEN knockout mice, the brain mass and neuronal size are augmented (Kwon et al., 2001, 2006). Therefore, PTEN ablation in PIKE−/− neurons should restore the PI3K/Akt function, thus rescuing the defective arborization. As expected, the dendritic patterns as well as the total dendritic length were increased in the double mutant neurons both *in vitro* and *in vivo* (Chan et al., 2011d). These observations are also in good agreement with the reports that the integrity of PI3K/ Akt is necessary for proper dendritic arborization and memory formation (Mizuno et al., 2003; Chen et al., 2005; Jaworski et al., 2005; Kumar et al., 2005). Presumably, the brain circuitry in PIKE−/− rain may be improperly or insufficiently wired, which provides another possibility for the memory defect.

#### **PIKE-A in human brain tumors**

The most prominent clinical evidence on PIKE ' s activity in disease onset is its role in neuro-oncology. High levels of PIKE-A are found in tumors that originate from the CNS including astrocytoma and glioblastoma, which result from gene amplification (Ahn et al., 2004b). Knobbe et al. (2005) further demonstrated that PIKE-A expression is also increased in 93 % of brain tumors without CENTG1 amplification. Our studies suggest that PIKE-A amplification leads to constitutive Akt activation, which prevents glioblastoma from apoptosis (Ahn et al., 2004b). This model is further supported by the observation that CENTG1 depletion in glioblastoma reduces Akt activity and increases apoptosis (Ahn et al., 2004a). Indeed, PIKE-A associates with the C-terminal regulatory and partial catalytic domains of Akt, which substantiates its kinase activity by increasing the phosphorylations on  $\text{Ser}^{473}$  and Thr<sup>308</sup>. As *CENTG1* is frequently co-amplified with cyclin-dependent kinase 4 (Cdk4) (a well-known proliferation activator by promoting E2F- and Cdk2-dependent cell cycle progression in human sarcoma and brain tumors; Reifenberger et al., 1994), it would be logical to imagine if PIKE-A amplification or overexpression coordinately acts with Cdk4 amplification to drive tumorigenesis. Using an integrated network analysis, Cerami et al. (2010) confirms that PIKE-A is a driver gene; the mutations of which contribute directly

to tumorigenesis in GBM (Cerami et al., 2010). A recent report further suggests that micro-RNA hsa-miR-26a, CDK4, and PIKE-A comprise a functional integrated oncomir/oncogene DNA cluster which promotes the aggressiveness in GBM (Kim et al., 2010). Interestingly, Cai et al. (2009) reported that PIKE-A promotes tumor progression by increasing the transcriptional activity of nuclear factor κB by interacting directly with its p50 subunit (Cai et al., 2009). However, it remains unknown whether the association also occurs in GBM.

Recently, we have found that PIKE-A also associated with UNC5B in glioblastoma cell lines (He et al., 2011). UNC5 are dependence receptors, which regulate apoptosis depending on its interaction with netrin-1 (Llambi et al., 2001). They have also been proposed as tumor suppressors which inhibit tumor extension by inducing apoptosis in areas without netrin-1 (Thiebault et al., 2003). Thus, association of PIKE-A with UNC5B may modulate the apoptotic function of the receptor. Indeed, PIKE-A/UNC5B interaction represses the receptor-induced cell death (He et al., 2011). This PIKE-A/UNC5B binding is tightly regulated by Akt in which Akt-mediated phosphorylation of PIKE-A provokes its interaction with UNC5B. Interestingly, PIKE-A also represses UNC5B transcription through downregulating p53 expression, which is in good alignment to the report that UNC5B is a direct target gene of p53 (Tanikawa et al., 2003; He et al., 2011). As such, netrin-1 might initiate Akt activation, which subsequently phosphorylates PIKE-A and escalates its interaction with UNC5B to prevent the receptor cleavage and consequent pro-apoptotic action. Moreover, Akt-phosphorylated PIKE-A feedbacks positively to further elevate the Akt kinase activity and leads to p53 degradation through the Akt-MDM2 pathway, resulting in down-regulation of UNC5B transcription (He et al., 2011).

#### **Conclusions**

The role of PIKE GTPases in maintaining neuronal survival is evident based on the studies performed in the past 10 years, yet we believe that the functional activities of PIKE in neurons have not been fully elucidated. Notably, PIKEs are implicated in regulating the activity of transcription factors such as signal transducer and activator of transcription 5A (STAT5A) after prolactin (PRL) stimulation (Chan et al., 2010a). As PRL/STAT5A cascade controls critical activities of neurons including neurogenesis and neurotransmitter release (Chen and Ramirez, 1982; Shingo et al., 2003), it remains to be explored if PIKE takes part in any of these processes under the control of PRL. Whether inactivation of PIKE is involved in chronic neurodegenerative diseases is another interesting question to be resolved. Genetic analysis of CENTG1 in patients of various neurodegenerative diseases would provide a definitive answer. Thus, further studies on the PIKE signaling in neurons will not only enhance our knowledge about its physiological activities but also have significant implications to disease prevention and treatments.

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### **Biographies**



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