



PTEN in Autism and Neurodevelopmental Disorders

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Phosphatase and tensin homolog (PTEN) is a classical tumor suppressor that antagonizes phosphatidylinositol 3-phosphate kinase (PI3K)/AKT signaling. Although there is a strong association of *PTEN* germline mutations with cancer syndromes, they have also been described in a subset of patients with autism spectrum disorders with macrocephaly characterized by impairments in social interactions and communication, repetitive behavior and, occasionally, epilepsy. To investigate PTEN's role during neurodevelopment and its implication for autism, several conditional *Pten* knockout mouse models have been generated. These models are valuable tools to understand PTEN's spatiotemporal roles during neurodevelopment. In this review, we will highlight the anatomical and phenotypic results from animal studies and link them to cellular and molecular findings.

Autism spectrum disorders (ASDs) comprise a heterologous group of syndromes characterized by neuroanatomical network changes, impairment in social interactions, and verbal and nonverbal communication as well as restrictive and repetitive behavior or interests manifesting by 2–4 years of life. In about 10% of the cases, early brain overgrowth (macrocephaly) and intellectual impairment are observed in ASD patients (Courchesne et al. 2003, 2007; Kennedy and Courchesne 2008; Donovan and Basson 2017). Classically, ASDs include autism, Asperger syndrome, childhood disintegrative disorder, and pervasive developmental disorder-not otherwise specified (PDD-NOS). The definition can also include other monogenetic neurodevelopmental diseases with autistic behavior such as Rett syndrome or Fragile-X syndrome. Most ASD cases occur by unknown

etiology, but there are several risk genes increasing the likelihood to develop ASD (Ramaswami and Geschwind 2018).

Though heterozygous germline mutations in the gene encoding phosphatase and tensin homolog (*PTEN*) are mostly linked to cancer syndromes, including Cowden, Bannayan-Riley-Ruvalcaba, and Proteus syndromes—commonly termed PTEN hamartoma tumor syndromes (PHTS)—they have also been described in patients with ASD (Fig. 1A; Zori et al. 1998; Goffin et al. 2001; Butler et al. 2005; Spinelli et al. 2015; Leslie and Longy 2016). PTEN, in its classical function as tumor suppressor, is the main antagonist of the phosphatidylinositol 3-phosphate kinase (PI3K)/protein kinase B (PKB/AKT) pathway by hydrolyzing phosphatidylinositol 3,4,5-triphosphate (PIP₃) to phosphatidylinositol 4,5-bisphosphate (PIP₂) to regulate cellular pro-

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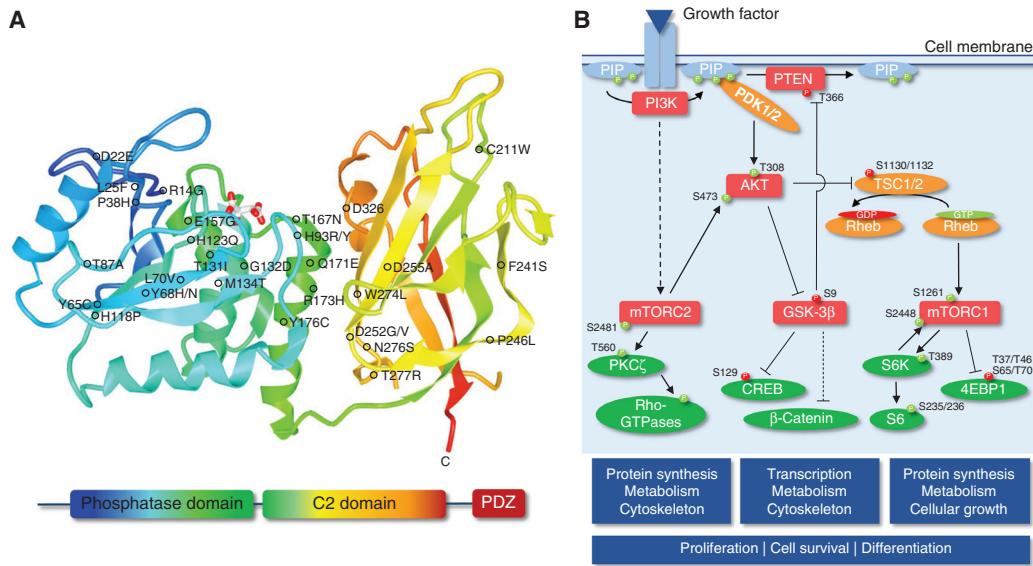


Figure 1. Phosphatase and tensin homolog (PTEN) structure and canonical phosphatidylinositol 3-phosphate kinase (PI3K)/PTEN signaling cascade. (A) Schematic representation of PTEN protein domains, together with the localization of germline missense mutations associated with ASD, developmental delay, and mental retardation (for a review, see Spinelli et al. 2015). Additional missense mutations not shown in the scheme: M1I, K6I/E. (B) Canonical PI3K/PTEN signaling cascade. Activating phosphorylation is depicted in green, and inhibitory phosphorylation is depicted in red. Dashed lines denote multiple steps.

liferation, differentiation, and cell death (Fig. 1B). Apart from its canonical lipid phosphatase function, PTEN is important for other cellular processes using its protein phosphatase activity in a compartment-specific fashion (Hopkins et al. 2014; Kreis et al. 2014). ASD-related *PTEN* mutations function mostly in a dominant-negative manner resulting in an unstable but catalytically active gene product, but can also lead to altered subcellular localization. For example, whereas most of the PHTS-linked *PTEN* mutations are loss-of-function mutations in line with loss of PTEN's canonical tumor-suppressive role, other ASD-associated mutations lead to impaired nucleocytoplasmic shuttling (Rodríguez-Escudero et al. 2011; Tilot et al. 2014; Spinelli et al. 2015; Fricano-Kugler et al. 2018; Mingo et al. 2018).

To uncover the involvement of PTEN loss during early neuronal development of ASD, researchers have generated several promoter-driven cre/lox animal models allowing conditional *Pten* knockout (KO) in a spatiotemporal manner. These animals mimic several phenotypes observed in PTEN-ASD patients ranging from

severe neuroanatomical alterations to the development of autistic behavior. Most importantly, these phenotypes strongly correlate with the time point of PTEN depletion. Although homozygous deletion is embryonically lethal (Di Cristofano et al. 1998), KO prior to neuronal differentiation (*Pten*^{ff}; *Gfap-cre* mice) results in hypertrophy, aberrant lamination of hippocampus and cerebellum, as well as seizures starting between 4 and 8 weeks and early death (Backman et al. 2001; Kwon et al. 2001). Deletion shortly after neuronal differentiation (*Pten*^{ff}; *Nse-cre*) leads to delayed hypertrophy, increased arborization, and spine density together with seizures obvious after 8 months, although first autistic behavior is already present after 6 weeks (Kwon et al. 2006). Deletion in fully mature neurons (*Pten*^{ff}; *CamKIIα-cre*), however, does not lead to morphological changes of the hippocampus, but induces severe defects in neuronal transmission (Sperow et al. 2012). In contrast, focal adeno-associated virus (AAV)-mediated *Pten* KO in the motor cortex leads to neuronal hypertrophy, laminar disruption, and increased neuronal

arborization after 12–18 months without, however, the emergence of severe motor defects or seizures (Gutilla et al. 2016; Gallent and Steward 2018). These data show that PTEN is not only important during whole organismal development and tumor suppression, but also during neurogenesis and neuronal maintenance.

But what are the molecular mechanisms leading to these phenotypes? During embryonal and early postnatal neurodevelopment, neuronal stem cells (NSCs) and neuronal precursor cells (NPCs) proliferate, migrate to their correct position, and finally differentiate into specialized neurons or glia cells in a timely defined manner. Differentiation additionally includes neurite outgrowth, dendritic arborization, and the formation of synapses to establish neuronal circuits, which, in turn, control motor function, memory formation, speech, anxiety, or social interactions (Fig. 2; de Graaf-Peters and Hadders-Algra 2006; Reemst et al. 2016; Thion and Garel 2017). In the following sections, we will review the most important anatomical, cellular, and molecular findings from animal studies that increased the understanding into the pathogenesis of PTEN-ASD.

Proliferation and Differentiation

Early brain overgrowth and lamination defects are a commonality in multiple patients suffering from ASD. One may speculate that this pheno-

type is caused by hyperproliferation of NSCs and NPCs and/or neuronal hypertrophy. Indeed, numerous studies indicate that PTEN-depleted premature NSC/NPCs show shorter cell-cycle duration concomitant with higher proliferation rates, less stress response with less (naturally occurring) apoptosis, accelerated growth, migration defects, aberrant outgrowth of axons and dendrites, and altered connectivity (Gao et al. 2000; Backman et al. 2001; Groszer et al. 2001, 2006; Chenn and Walsh 2002; Li et al. 2002; Jo et al. 2012; Chen et al. 2015; Lyu et al. 2015). In the absence of PTEN, NPCs are able to differentiate both into neurons as well as into glia cells, but show a higher tendency to differentiate into neurons with enhanced neuritogenesis (Groszer et al. 2001; Jo et al. 2012; Chen et al. 2015; Lyu et al. 2015). In contrast, at postnatal stages, *Pten*-deficient NPCs preferentially differentiate into glia cells rather than into neurons (Chen et al. 2015). Interestingly, these outcomes seem to be restricted to NSCs but not to mesenchymal stem cells in which PTEN depletion leads to senescence rather than hyperproliferation (Duan et al. 2015).

In NSCs, PTEN predominantly localizes to the nucleus (Duan et al. 2015), which is in line with PTEN-regulating G₀-G₁ cell-cycle entry, genomic integrity, and preventing NSCs/NPCs from excessive self-renewal, and reaching the premature cell division limit (Ginn-Pease and Eng 2003; Groszer et al. 2006; Jo et al. 2012;

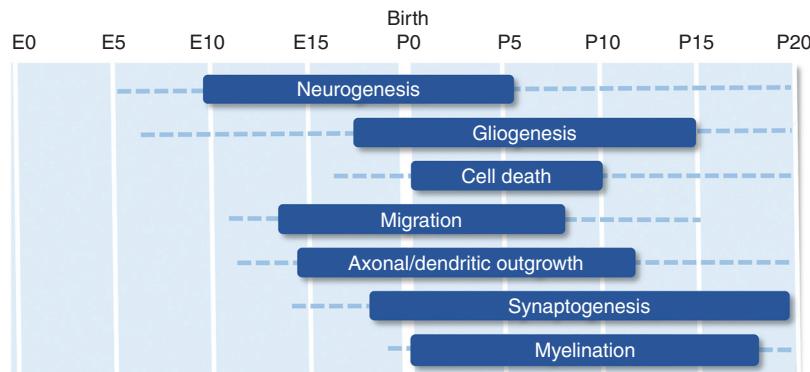


Figure 2. Stages of murine neurodevelopment. Boxes indicate peaks in the processes, dashed lines indicate that the process is active to a distinct extent. (Figures created from data in Reemst et al. 2016 and Thion and Garel 2017.)



Feng et al. 2015; Liu et al. 2017). In postmitotic neurons, such as retinal ganglion cells, PTEN's cytosolic role is important to antagonize PI3K signaling as enriched PI3K signaling facilitates cell-cycle exit and differentiation (Jo et al. 2012). Importantly, as well as Ras/ERK signaling, the PI3K/PTEN/AKT axis is one central node in the fate of NSCs/NPCs, for example in the control of proliferation, differentiation, size regulation, arborization, or survival. This involvement is required in the tight and spatiotemporal integration of different signaling cues or cross talking to other signaling pathways such as Wnt/β-catenin, Notch, brain-derived neurotrophic factor (BDNF), fibroblast growth factor 2 (FGF2), insulin-like growth factor (IGF), or epidermal growth factor (EGF) signaling.

The control of the precise transition of NSC/NPC proliferation and differentiation involves two important pathways, the β-catenin and Notch axes, both of which regulate cAMP response element-binding protein (CREB)-mediated transcription parallel or downstream of PI3K/AKT, respectively. β-Catenin promotes symmetric division of radial glia (RG) in early development, but promotes differentiation later during neurogenesis (Wrobel et al. 2007). The cross talk between PI3K and β-catenin signaling is important in this decision as AKT directly phosphorylates β-catenin and inhibits GSK-3β-induced β-catenin destruction (Polakis 2000; Fang et al. 2007). Thus, PI3K/AKT signaling leads to nuclear β-catenin accumulation and transcriptional activation via coactivators, including CREB-binding protein (CBP) or p300 (Fig. 3A). Here, either CBP or p300 may decide whether to divide or to differentiate by distinctly regulating *survivin*, *c-myc*, and/or *cyclinD1* expression (Ma et al. 2005; Teo et al. 2005). Additionally, β-catenin promotes RG differentiation by up-regulating *n-myc* and inhibiting Notch signaling (Kuwahara et al. 2010; Zinin et al. 2014). The Notch axis is another important regulator of NPC fate, which depends on PTEN activity. Upon activation, Notch intracellular domain (NICD) forms a complex with Mastermind (MAM) and recombining binding protein suppressor of hairless (RBPF) to recruit CBP/CREB and to induce expression of the transcrip-

tional repressor *Hes1*. HES1, in turn, down-regulates proneural genes and prevents neuronal differentiation in an oscillatory manner (Shimojo et al. 2008), but also inhibits PTEN (Jo et al. 2012). Thus, AKT activity temporally increases and prevents NICD:MAM:RBPF complex formation to terminate the Notch signaling cascade and *Hes1* expression. At a later stage, PTEN activity again increases to down-regulate AKT signaling and maintain NSCs in a proliferative state (Fig. 3B; Palomero et al. 2007; Jo et al. 2012).

Similar to cancer syndromes, in which the PTEN dosage influences tumor progression (Trotman et al. 2003), relative PTEN protein abundance has also been observed as an important regulatory characteristic in the nervous system of *Pten* mouse models. For instance, heterozygous *Pten* mutant mice show elevated AKT/β-catenin signaling that coincides with macrocephaly because of excessive neuron numbers. In homozygous *Pten* mutant mice, however, increased AKT/mechanistic target of rapamycin (mTOR) signaling regulates cell size (Chen et al. 2015). These results are in line with studies showing that β-catenin overexpressing mice develop macrocephaly (Chenn and Walsh 2002), whereas β-catenin mutations result in microcephaly, ataxia, and intellectual disabilities (Dubruc et al. 2014). In contrast, PTEN depletion and accelerated AKT activity in the retina leads to decreased NICD translocation, altered CREB-mediated transcription, and premature loss of RPCs (Jo et al. 2012). Together, these results show that CREB is a central hub in neuronal development (for a review, see Sakamoto et al. 2011) and dysregulation may result in aberrant NSC/NPC proliferation and differentiation. Of importance, mutations in other CREB-related genes lead to various syndromic autism (e.g., *MeCP2* or *CREBBP/EP300*), which are themselves causative for Rett syndrome and Rubinstein-Taybi syndrome, respectively (Spina et al. 2015; Lyu et al. 2016; Negri et al. 2016; Bu et al. 2017). However, additional mechanisms may be involved in the cross talk of β-catenin/Notch/CREB axis as PTEN directly dephosphorylates CREB independently of PI3K/AKT and PTEN loss leads to up-regulation of several

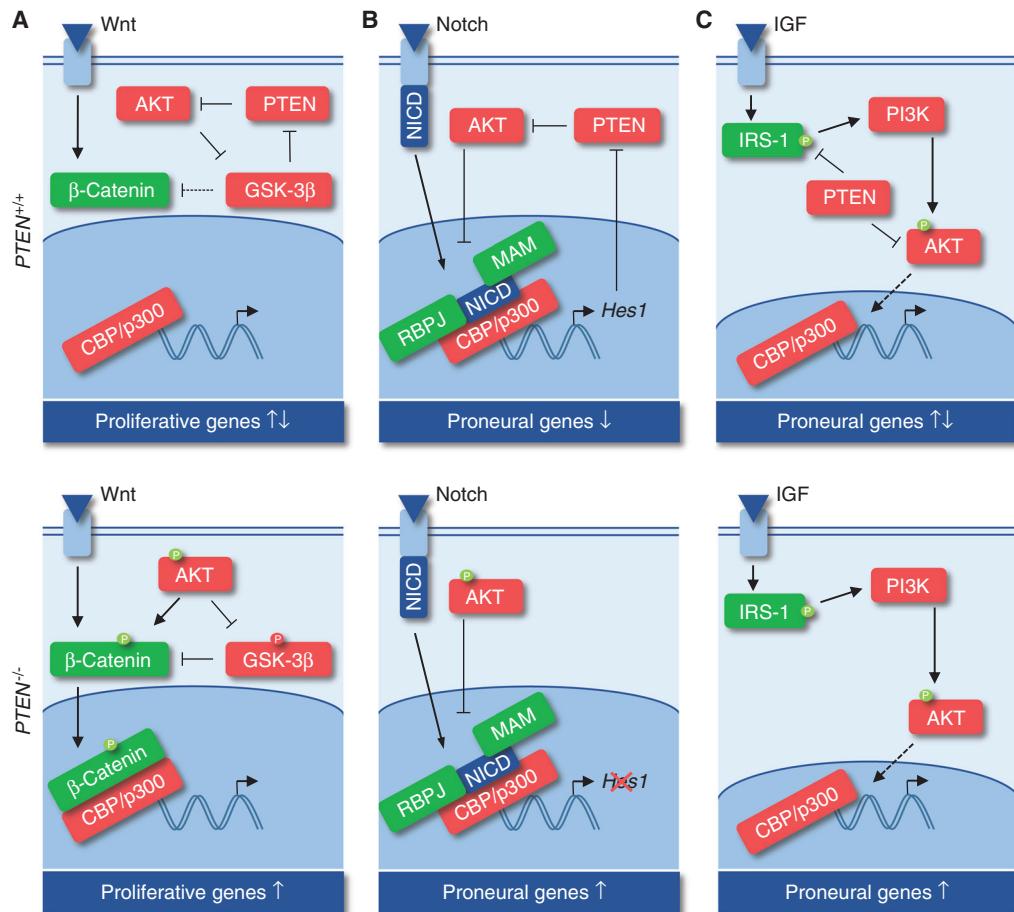


Figure 3. Cross talk between phosphatidylinositol 3-phosphate kinase (PI3K)/AKT/phosphatase and tensin homolog (PTEN) cascade with other signaling axes during proliferation and differentiation. (A) Cross talk with Wnt/β-catenin signaling. (B) Cross talk with Notch signaling. (C) Cross talk with insulin-like growth factor (IGF) signaling.



CREB downstream targets, including the survival-promoting transcripts *c-myc* and *Bcl-2* (Gu et al. 2011) or *Pax7* (Duan et al. 2015). Thus, further experiments deciphering the timely and specific role of CREB targets in *Pten* KO mice are needed to fully understand the precise function of PTEN in controlling proliferation and differentiation.

Migration and Lamination

During cortical development, excitatory neurons are generated from three types of NPCs: (1) RG in the ventricular zone with long ascend-

ing fibers and short apical processes, which can self-renew by asymmetric division to generate one neuron and one RG; (2) apical interprogenitor cells in the ventricular zone undergoing symmetric division to generate two neurons; and (3) intermediate progenitor cells in the subventricular zone with symmetric division that form two neurons (for a review, see Paridaen and Huttner 2014; Hirota and Nakajima 2017). Several mouse models, in which *Pten* was conditionally deleted under the *Gfap* promoter, develop lamination defects especially in the cortex and hippocampus, but also in the amygdala and cerebellum (Fig. 4). These lami-

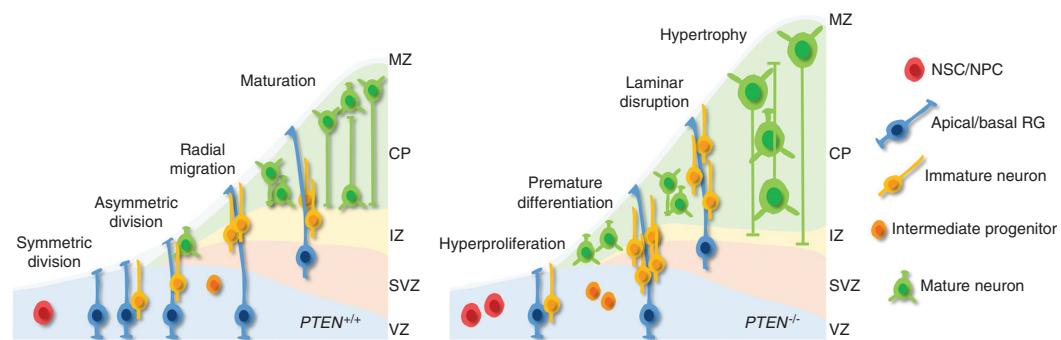


Figure 4. Cortical development in *phosphatase and tensin homolog* (*PTEN*)-positive and -negative brains. NSC, Neuronal stem cell; NPC, neuronal precursor cell; RG, radial glia; MZ, marginal zone; CP, cortical plate; IZ, intermediate zone; SVZ, subventricular zone; VZ, ventricular zone.

nation defects are commonly associated with seizures and ataxia (Backman et al. 2001; Kwon et al. 2001; Wen et al. 2013). Deletion of *Pten* in Bergmann glia results in premature differentiation of granule cells concomitant with aberrant cerebellar lamination (Yue et al. 2005). Additionally, a reduction of Müller glia with increase of other neuronal cells is observed upon *Pten* deletion in the retina (Jo et al. 2012). In another study, the migration of NSCs from the subventricular zone to the olfactory bulb was analyzed in heterozygous *Pten* mice (Li et al. 2002). In these mice, NPCs show a more rapid radial migration into the olfactory bulb. However, consistent with other studies, this increased migration behavior does not lead to bulb overgrowth (Chen et al. 2015). Although studies commonly show lamination defects upon *Pten* deletion, it is not clear whether such defects result from aberrant migration or from premature differentiation (Yue et al. 2005). As stated in the section Proliferation and Differentiation, NPCs from *Pten* KO models show a higher tendency to differentiate into neurons rather than glia. An important regulator of asymmetric division of RGs is Notch, which exhibits an asymmetrical distribution of its intracellular signaling molecules during cell division. Here, Notch signaling maintains one daughter in a proliferative RG state, whereas the cell receiving less Notch signaling differentiates into an intermediate progenitor or neuron (Bultje et al. 2009; Dong et al. 2012). Given that *Pten* KO

neurons exhibit aberrant Notch signaling, this could be one possible explanation for increased neurogenesis, premature loss of RGs, and lamination defects. However, further studies are needed to investigate the cell-autonomous and non-cell-autonomous effects of PTEN functions in RGs.

Hypertrophy

One of the dominant phenotypes in ASD patients with *PTEN* germline mutations is the development of macrocephaly. Thus, NSC/NPC hyperproliferation, together with aberrant migration, may only play a minor role in brain overgrowth. There is evidence that PTEN loss-induced neuronal hypertrophy of soma, axon, and dendrites may be responsible for this phenotype. For instance, cerebellar overgrowth in *Pten*^{ff}; *Gfap-cre* animals is caused by postmitotic granule cells with no indication of hyperproliferation (Kwon et al. 2001). Similarly, *Pten*^{ff}; *Nse-cre* *Pten* KO mice reveal hypertrophic mossy fibers, broader innervation of the CA3 area and misplaced cells, together with ectopic positioning of synapses (Kwon et al. 2006). This hypertrophic phenotype is a cell-autonomous effect as a result of accelerated AKT/mTOR/S6K signaling and increased messenger RNA (mRNA) translation, rather than a response to the cellular microenvironment (Backman et al. 2001; Kwon et al. 2001, 2003; Arafa et al. 2019). In support of this, injection of the mTOR inhib-



itor CCI-779 prevents neuronal hypertrophy in *Pten^{ff};Gfap-cre* mice and also reduces premature death and seizure frequency (Kwon et al. 2003). In addition, nuclear PTEN has a significant impact on the regulation of soma size, although the studies are contradictory. This may be because of the differences in the models used (i.e., expression of nuclear-excluded *PTEN* patient mutations) (Fricano-Kugler et al. 2018), CRISPR/Cas-based introduction of the K13R mutation to prevent nuclear shuttling (Igarashi et al. 2018), or the use of the *Pten^{m3m4}* mouse model showing a drop of total and nuclear PTEN (Tilot et al. 2014). Seemingly, the amount of residual PTEN load, the time point of induced PTEN-loss, and the area of modulation may have a pivotal role in the severity and specificity of phenotypes.

Neurite Outgrowth and Arborization and Spine Formation

PTEN is involved in neuronal polarization, axonal and dendritic outgrowth, and arborization, but each of these processes underlies different downstream mechanisms. Experiments using an RNAi approach show that PTEN depletion by more than 50% in hippocampal neurons results in the formation of multiple axons at the cost of dendrite formation by regulating GSK-3β activity (Jiang et al. 2005). This is in line with the *in vivo* findings from *Pten^{ff};Nse-cre* mice, which develop longer axons but also elongated and thickened dendrites in the dentate gyrus (Kwon et al. 2006). In contrast, *PTEN* overexpression prevents polarization of hippocampal neurons (Shi et al. 2003). These studies suggest that the PTEN/GSK-3β axis may be pivotal in polarization as well as axodendritic outgrowth and identity and that *Pten*-deficient neurons lack proper termination of axonal and dendritic growth.

Once an axon has formed, it starts to extend toward its target area. Axonal pathfinding is selectively regulated by the cellular microenvironment. Chemorepulsive or chemoattractive cues are secreted by surrounding cells that bind to specific receptors on axonal growth cones (e.g., semaphorin/plexin, draxin/DCC, or neurotro-

phins) (for a review, see Russell and Bashaw 2018). A common downstream event of these repulsive ligand–receptor interactions is the reorganization of the actin/microtubule cytoskeleton and β-integrin removal to induce growth cone collapse and prevent outgrowth (Henle et al. 2013). There is growing evidence that chemorepulsion depends on PTEN by downregulating PIP₃ levels and antagonizing AKT signaling, whereas chemoattraction is PTEN independent (Henle et al. 2013). PTEN distributes in the growth cone and translocates to the membrane upon repulsive stimulation with Sema3A, which results in decreases in PIP₃/AKT signaling and activation of GSK-3β (Chadborn et al. 2006). It has been shown that GSK-3β is, at least in part, crucial to induce Sema3A-mediated growth cone collapse by the modulation of F-actin dynamics (Fig. 5A; Eickholt et al. 2002). Apart from F-actin remodeling during chemorepulsion, the PTEN/GSK-3β axis organizes microtubule dynamics in a number of different ways. First, PTEN regulates microtubule detyrosination and PTEN depletion results in hyperstabilization of microtubules together with increased axonal outgrowth in hippocampal neurons (Kath et al. 2018). Second, GSK-3β phosphorylates several microtubule-associated proteins (e.g., MOB1, Tau, MAP1B) to regulate microtubule assembly and to prevent neurite outgrowth (Wagner et al. 1996; Meli et al. 2015; Song et al. 2018). For instance, the chemo-repellent draxin uses GSK-3β to phosphorylate MAP1B and to prevent axonal growth concomitant with *Draxin* KO in mice leading to severe morphological changes of different brain structures including the corpus callosum (Ahmed et al. 2011; Meli et al. 2015). There are numerous examples clearly indicating that PI3K/PTEN signaling is essential for accurate guidance of axons; it is furthermore relevant for mediating cellular adhesion to the extracellular matrix as PTEN depletion results in less removal of focal adhesions in zebrafish spinal cord (Henle et al. 2013).

Cytoskeletal remodeling further requires local translation of proteins to regulate neurogenesis and arborization. Mechanistically, a well-regulated interplay between the mTORC1

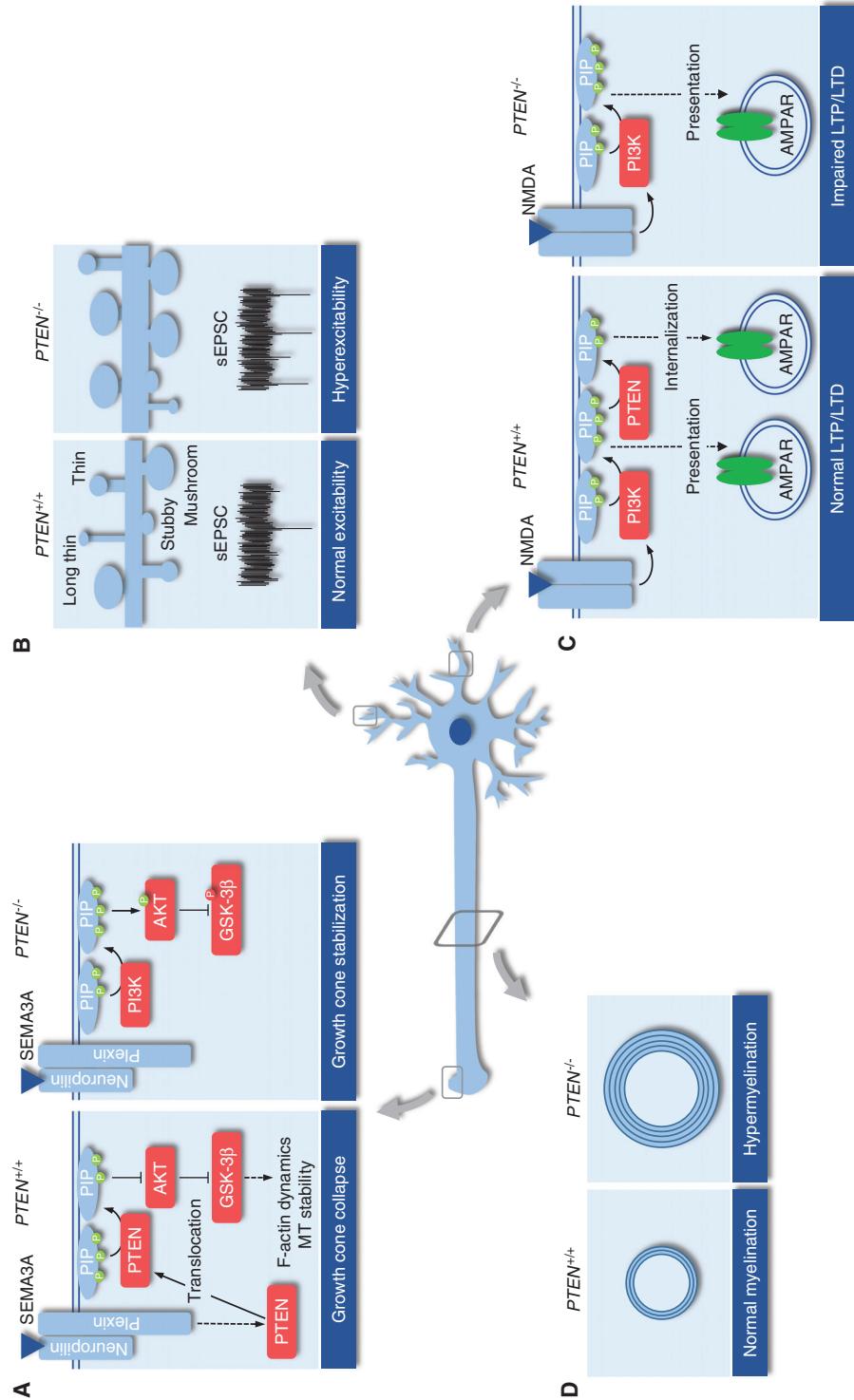


Figure 5. Processes in postmitotic neurons that are altered in the gene encoding phosphatase and tensin homolog (*PTEN*) loss. (A) Regulation of growth cone collapse. (B) Regulation of spine density, morphology, and excitability. (C) Regulation of synaptic plasticity. (D) Regulation of axonal myelination. sEPSC, Spontaneous excitatory postsynaptic current; PI3K, phosphatidylinositol 3-phosphate kinase; NMDA, *N*-methyl-D-aspartate; LTP, long-term potentiation; LTD, long-term depression.

and mTORC2 complexes drives these processes in hippocampal neurons (Urbanska et al. 2012), distinctly by mTORC1 promoting mRNA translation via p70S6K or by mTORC2 regulating actin dynamics via RhoGTPases (Jacinto et al. 2004; Sarbassov et al. 2004; Jaworski et al. 2005). To allow for increased local mTOR activity and dendrite growth or branching to occur, local PTEN depletion is essential. Indeed, it has been shown that NEDD-4, an E3-ligase that catalyses PTEN ubiquitination and proteasomal degradation (Wang et al. 2007), is a regulator of axonal branching (Drinjakovic et al. 2010; Christie et al. 2012; Hsia et al. 2014).

NEDD-4 furthermore regulates PTEN nuclear translocation by monoubiquitination (Trotman et al. 2007; Wang et al. 2007). PTEN's nuclear role during axonogenesis is not completely understood, but it has been shown that PTEN interacts with the APC-CDH1 complex (Song et al. 2011) and that this complex drives the expression of different cues important for axonal growth (Lasorella et al. 2006). PTEN localization has been systematically analyzed in the developing brain during neurito- and synaptogenesis. Interestingly, during neuritogenesis, PTEN is mainly localized in the nucleus and sparsely found in dendrites, but retranslocates into mature dendrites to antagonize mTOR signaling, pause arborization, and promote synaptogenesis (Perandones et al. 2004). These data overlap with the narrow time window, in which mTOR inhibitors are able to reverse aberrant outgrowth defects in a tuberous sclerosis complex (*Tsc1^{ff};Gfp-cre*) model of autism (Cox et al. 2018). Thus, a spatial regulation of PTEN localization is important for proper neuritogenesis, and constant PTEN (nuclear) depletion leads to increased axodendritogenesis with increased synaptogenesis possibly caused by hyperactive mTOR signaling. Given that *Pten* loss leads to diminished growth cone collapse, increased axonal and dendritic outgrowth, and more focal adhesions, it is plausible that *Pten*-deficient neurons cannot stop growing in a timely fashion and form ectopic synapses. Consequently, this effect will have a significant impact on neuronal connectivity, synaptic plasticity, and the development of autistic behavior.

PTEN in Connectivity and LTP/LTD

PTEN-ASD patients as well as *Pten* cKO mice develop behavioral abnormalities, including aberrant social interactions, problems in communication, impaired cognition and memory, repetitive behavior, and altered fear/anxiety (Kwon et al. 2006; Sperow et al. 2012; Lugo et al. 2014; Smith et al. 2016; Hodges et al. 2018). *Pten* cKO animals have been analyzed for synaptic connectivity and plasticity, showing that there is an imbalance between synaptic excitation and inhibition (E/I balance), which is now well accepted as a key pathological mechanism linked to developing ASD-related behavior (Lee et al. 2017). Systematic approaches involve studies in which *Cre*-expressing virus or shRNA against *Pten* has been injected into the dentate gyrus (Luikart et al. 2011; Williams et al. 2015; Skelton et al. 2019), amygdala (Haws et al. 2014), or auditory cortex (Xiong et al. 2012) of neonatal mice. Although differences in spine density upon *Pten* deletion have been observed, which may be because of different time points of virus injection and tissue specificity (P7 dentate gyrus vs. P49 basolateral amygdala vs. *Pten^{ff};Nse-cre*), the number of mature mushroom-like spines is increased compared to controls (Haws et al. 2014; Williams et al. 2015). Additionally, *Pten* KO neurons receive accelerated afferent input from multiple neurons (Skelton et al. 2019). As a consequence, neurons require a higher firing threshold to evoke an action potential but, at the same time, are more sensitive to afferent stimulation and fire more readily as shown by higher excitatory postsynaptic current (EPSC) frequency and amplitude (Fig. 5B; Luikart et al. 2011; Haws et al. 2014; Williams et al. 2015; Skelton et al. 2019). Inhibitory input on these neurons is unchanged (Williams et al. 2015), indicating that *Pten* deletion in excitatory neurons makes them hyperexcitable. On the other hand, compelling studies show an involvement of GABAergic inhibitory interneurons. *Pten^{ff};Nkx2.1-cre* mice with *Pten* deletion in the medial ganglionic eminence demonstrate a preferential loss of somatostatin- over parvalbumin-positive GABAergic interneurons, but, at the same time, show a two-fold increase of induced pluripotent stem cells





(iPSCs) on layer II/III excitatory neurons concomitant with autistic behavior (Vogt et al. 2015). In this study, we have also analyzed the effect of ASD-related *PTEN* mutations in a knockin model and observed similar effects on the ratio of somatostatin/parvalbumin-positive interneurons (Vogt et al. 2015). Together, these results indicate that changes in afferent input render neurons either hyper- or hypoexcitable, potentially providing the strongest causal link to the development of ASD-related behavior. Such a mechanistic relationship has also been observed in other ASD models including Rett syndrome (Wood et al. 2009).

PTEN-ASD subjects and mice often develop (or are prone to) seizures, a condition of synchronized pathological firing, and it is accepted that neuronal hyperexcitability is a driving force for epileptogenesis. Aberrant circuitry of the dentate gyrus because of ectopic migration, hypertrophy, and hyperexcitability has been shown to result in increased spontaneous EPSCs, fewer iPSCs, and multiple population spikes and is also a well-known factor for temporal lobe epilepsy (Nadler 2003; Pun et al. 2012; LaSarge et al. 2016; Santos et al. 2017). From a molecular point of view, an involvement of potassium channels in hippocampal hyperexcitability of 8-week-old *Pten^{ff};Gfap-cre* mice has been implicated (Lugo et al. 2014; Nguyen and Anderson 2018). Another interesting point is that induction of status epilepticus in *Pten^{ff};Gfap-cre* mice, in which granule cells are mainly affected by *Pten* deletion, results in more severe hyperactivity and impairment in social behavior compared to animals that have not undergone seizures (Smith et al. 2016).

Along with E/I imbalance and development of seizures, *Pten* deletion alters synaptic plasticity in an age- and brain region-dependent manner. For instance, *Pten* cKO (*Pten^{ff};Nse-cre*) enhances theta burst-induced long-term potentiation (LTP) in dentate granule neurons of young (8–12 weeks) animals, but completely impairs LTP in older (20–30 week) animals (Takeuchi et al. 2013). Comparably, *Pten* cKO decreases LTP at CA3-CA1 synapses of 8-week-old *Pten^{ff};CamKIIα-cre* mice (Sperow et al. 2012) or *Pten^{ff};Gfap-cre* mice (Fig. 5C; Fraser et al. 2008). Additionally, *N*-methyl-D-aspartate

(NMDA) receptor (NMDAR)- and metabotropic glutamate receptor (mGluR)-dependent long-term depression (LTD) are impaired in these mice at all time points analyzed (Sperow et al. 2012; Takeuchi et al. 2013), whereas acute pharmacological inhibition of PTEN lipid phosphatase activity prevents only NMDAR-dependent LTD but not mGluR-dependent LTD nor LTP (Jurado et al. 2010). These data suggest that PTEN depletion early during development leads to aberrant synaptic strength and connectivity along with morphological changes, whereas mature neurons respond with altered synaptic plasticity. Anatomical and morphological correlates, together with the frequency and severity of seizures upon early or late PTEN modulation, further strengthen this hypothesis. Importantly, the first signs of ASD-related behavior and seizures occur prior to severe morphological changes in *Pten^{ff};Nse-cre* mice (Kwon et al. 2006; Takeuchi et al. 2013).

But how does this happen? On the one hand, long-term *Pten* deletion may lead to further synaptic/cellular defects that remain to be determined. On the other hand, it may be the case that aberrant plasticity induced by *Pten* loss may involve additional PI3K-dependent as well as PI3K-independent actions of the phosphatase. For example, the F-actin-stabilizing protein drebrin is an important regulator of spine formation, maintenance, synaptic plasticity, context-dependent fear learning, and social behavior (Hayashi and Shirao 1999; Kojima et al. 2010; Koganezawa et al. 2017). In young adults, *drebrin* loss does not result in any discernible phenotypes nor alter synaptic plasticity under physiological conditions (Willmes et al. 2017), whereas its overexpression leads to abnormal filopodia formation in immature neurons, spine elongation in mature neurons, and E/I imbalance (Hayashi and Shirao 1999; Mizui et al. 2005; Ivanov et al. 2009). Moreover, gabazine treatment, which induces seizure-like network activity, results in drebrin hyperphosphorylation (Kreis et al. 2013), potentially involving hyperactivity-induced local production of reactive oxygen species (ROS) and activation of the ATM kinase to protect spines during aging (Kreis et al. 2019). Because PTEN dephosphor-

ylates drebrin independently of PI3K signaling leading to drebrin destabilization (Kreis et al. 2013; Kreis et al. 2019), it could be assumed that *Pten* loss does not counteract the increased drebrin phosphorylation during seizures. It may be the case that *Pten* deletion further promotes spine enlargement and synaptic strengthening by using the actin-stabilizing activity of drebrin. Such a PI3K-independent effect of PTEN could be one explanation of why co-KO of *Pdk1* (*Pten^{ff};Pdk1^{ff};CamKIIα-cre*), a positive regulator of mTOR, rescues LTP and LTD deficits, but is ineffective in rescuing impairments in spatial memory formation in *Pten^{ff};CamKIIα-cre* mice (Sperow et al. 2012). In addition to drebrin, PTEN interacts with PSD-95 via its PDZ-binding domain and is recruited to the PSD upon NMDA stimulation in a biphasic fashion to regulate PIP₃ signaling during LTP and LTD (Jurado et al. 2010; Arendt et al. 2014). Interestingly, the protein levels of PSD-95 as well as other postsynaptic proteins, including mGluR5 or Kv4.2, are down-regulated in *Pten^{ff};Gfap-cre* animals, whereas other ASD-related scaffolding proteins, including FMRP, Shank, or Sapap1, are up-regulated (Lugo et al. 2014), further suggesting molecular commonalities between neurodevelopmental disorders.

PTEN and Myelination

During neuritogenesis, oligodendrocytes and Schwann cells myelinate the newly formed axon in the central nervous system (CNS) or peripheral nervous system (PNS), respectively. Tremendous thickening of myelin sheaths with aberrant myelin compaction, abnormal formation of nodes of Ranvier, and reduced g-ratio (i.e., the ratio of the inner axonal diameter to the total axonal diameter) have been observed in *Pten^{ff};Gfap-Cre* (Fraser et al. 2008), *Pten^{ff};Olig2-cre* (Harrington et al. 2010; Maire et al. 2014), *Pten^{ff};Cnp1-cre* (Goebels et al. 2010), or *Pten^{ff};Pdgfra-cre* (González-Fernández et al. 2018) mouse models, especially in the corpus callosum and spinal cord (Fig. 5D). In contrast, reduced myelin sheath thickness with increased g-ratio in the corpus callosum caused by the inability of oligodendrocytes to properly target

and ensheathe axons has been described in homozygous *Pten^{m3m4}* mice (Lee et al. 2019). Although different models have been used to analyze axonal myelination, the mice collectively show increased cerebral white matter, which is a common phenotype in patients harboring *PTEN* mutations (Vanderver et al. 2014; Frazier et al. 2015). Additionally, these studies report on aberrant oligodendrocyte precursor cell proliferation, migration, and differentiation (Maire et al. 2014; González-Fernández et al. 2018; Lee et al. 2019), increased myelin synthesis (Harrington et al. 2010; González-Fernández et al. 2018; Lee et al. 2019), defective myelin storage (Lee et al. 2019), and decreased myelin compaction (Fraser et al. 2008; Lee et al. 2019). As such, the inability of *Pten^{m3m4}* oligodendrocytes to correctly wrap axons has been proposed to be caused by aberrant maturation and precocious myelin spreading making the myelin sheath dysfunctional (Lee et al. 2019).

Mechanistically, CNS myelination has been linked to mTOR signaling, as myelination defects in *Pten^{ff};Cpn1-cre* mice can be rescued by the application of rapamycin (Goebels et al. 2010). However, mTOR double cKO (*Pten^{ff};Mtor^{ff};Pdgfra-cre*) does not ameliorate the phenotypes seen in *Pten^{ff};Pdgfra-cre* animals, which has led to the hypothesis that PTEN's effect on myelin synthesis involves GSK-3β (González-Fernández et al. 2018). Based on this model, constant PTEN depletion in mature oligodendrocytes results in decreased GSK-3β and increased mTORC1 activity to facilitate myelination. However, increased GSK-3β phosphorylation is not observed in oligodendrocytes from mice expressing hyperactive AKT (Narayanan et al. 2009). A similar model has been proposed for Schwann cells, in which high mTORC1 activity suppresses Schwann cell differentiation, whereas physiological activity promotes myelin synthesis and axon wrapping (Figlia et al. 2017). As well as the cell-autonomous effects upon altered mTOR and GSK-3β activities, other signaling pathways or regulatory factors may influence myelination. For instance, transient PTEN inhibition potentiates IGF-1-mediated oligodendrocyte precursor differentiation and myelination in collaboration of PI3K/





AKT and MEK/ERK pathways (De Paula et al. 2014). Similarly, Wnt/β-catenin signaling is absolutely essential for normal axonal myelination (for a review, see Gaesser and Fyffe-Maricich 2016), which has not yet been studied in *Pten* cKO oligodendrocytes. PTEN interacts also with several scaffolding and polarization proteins, including Dlg1 or Par3, and interferes with β-integrin signaling. In Schwann cells, Dlg1 has been identified as a negative regulator of axon ensheathing by up-regulating PTEN (Cotter et al. 2010). However, in oligodendrocytes, myelination of CNS axons appears to be independent on Dlg1 (Brinkmann et al. 2008), indicating that further molecules may be important. Interestingly, it has been shown that β1 integrin is important during myelination by up-regulating AKT activity (Barros et al. 2009). Given that β1 integrin removal upon chemorepulsive stimulation is PTEN dependent in neuronal growth cones (Henle et al. 2013), one could speculate that PTEN depletion in oligodendrocytes has a similar effect on β1 integrin removal. Axon caliber is another determinant during myelination in the CNS and recent evidence comes from one study showing that secreted factors from *Pten^{ff}*; *Nex-cre* granule cell axons regulate their myelination by wild-type oligodendrocytes (Goebels et al. 2017).

In conclusion, there has been significant progress in understanding the function of PTEN during myelination. It will be relevant to analyze whether *Pten*-deficiency-induced myelination phenotypes impact on the progression of developmental disorders and epilepsy. For example, it will be interesting to analyze in detail whether the PTEN-depletion-induced myelination defect in the corpus callosum contributes to epilepsy and behavioral deficits, especially as corpus callosotomy is considered as a surgical procedure for some children suffering from interhemispheric seizures (Luat et al. 2017).

CONCLUDING REMARKS

cKO mouse models have emerged as valuable tools to understand specific spatiotemporal roles of PTEN during development and its impact

on human developmental disorders including ASD. Using these models, significant progress has been made in determining cell-autonomous and cell-intrinsic effects that result as a consequence of *Pten* deletion. However, cKO models do not reflect entirely the patients' situation, because the latter harbor heterozygous *PTEN* germline mutations but do not possess specific *PTEN* null neurons. The timing of the neurodevelopmental processes described in this review (proliferation, migration, differentiation, neuronal connection, synaptic plasticity, and axonal myelination) is intermingled and the understanding of how the actual pathomechanism of PTEN-ASD relies on a dysregulation of a specific event is unclear. Comprehensive analysis of the cellular, anatomical, electrophysiological, and behavioral observations of models, in which ASD-related mutations have been introduced in the *Pten* allele(s), will be valuable. More physiological models, which have been adopted to date in only a few studies (Vogt et al. 2015; Lee et al. 2019), will have the advantage of showing mutant *Pten* expression in all tissues, cells, and progenitors at the same time. Similarly, PTEN PI3K-independent but also phosphatase-independent processes need to be considered further to comprehend the multifaceted contributions of PTEN to neuronal development and function. Correlating these results with the data obtained from cKO models could be a valuable step in understanding ASD development and to optimize the time window of treatment strategies.

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