

Retinoic acid signaling in axonal regeneration

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Radhika Puttagunta and Simone Di Giovanni, Laboratory for Neuroregeneration and Repair, Hertie Institute for Clinical Brain Research, University of Tuebingen, Otfried-Mueller Strasse 27, D-72076 Tuebingen, Germany. e-mail: radhika.puttagunta@ medizin.uni-tuebingen.de; simone.digiovanni@medizin. uni-tuebingen.de Following an acute central nervous system (CNS) injury, axonal regeneration and functional recovery are extremely limited. This is due to an extrinsic inhibitory growth environment and the lack of intrinsic growth competence. Retinoic acid (RA) signaling, essential in developmental dorsoventral patterning and specification of spinal motor neurons, has been shown through its receptor, the transcription factor RA receptor β2 (RARβ2), to induce axonal regeneration following spinal cord injury (SCI). Recently, it has been shown that in dorsal root ganglion neurons (DRGs), cAMP levels were greatly increased by lentiviral RARβ2 expression and contributed to neurite outgrowth. Moreover, RARβ agonists, in cerebellar granule neurons (CGN) and in the brain *in vivo*, induced phosphoinositide 3-kinase dependent phosphorylation of AKT that was involved in RARβ-dependent neurite outgrowth. More recently, RA-RARβ pathways were shown to directly transcriptionally repress a member of the inhibitory Nogo receptor (NgR) complex, Lingo-1, under an axonal growth inhibitory environment *in vitro* as well as following spinal injury *in vivo*. This perspective focuses on these newly discovered molecular mechanisms and future directions in the field.

Keywords: RA, RAR_β, axonal regeneration

INTRODUCTION

The limited regenerative capacity observed following an acute central nervous system (CNS) lesion, such as stroke or brain/spinal cord traumatic injuries, is due to both the presence of an extrinsic inhibitory growth environment and the lack of intrinsic growth factors. Interestingly, the peripheral nervous system (PNS) does regenerate to a certain extent following a lesion, which has led to further research to determine the distinction between the CNS and PNS. The inhibitory environment of the CNS is vastly different than that of the PNS (Richardson et al., 1980). Following PNS injury, an inflammatory response is activated, Schwann cells infiltrate the injured area, debris is removed, neurotrophic factors are released, regeneration is initiated, and sheathing of these growing axons occurs (George and Griffin, 1994). Conversely, the glia cells in the CNS, oligodendrocytes, and astrocytes, provide an inhibitory environment for growth (McKeon et al., 1991; Qiu et al., 2000; Domeniconi et al., 2002; Niederost et al., 2002; Yiu and He, 2006).

Specifically, the neuronal insulating layer, myelin, is fragmented following a CNS lesion, releasing the extrinsic inhibitory molecules, myelin associated glycoprotein (MAG), Nogo, and Oligodendrocyte myelin glycoprotein (OMgp) (DeBellard et al., 1996; Huber and Schwab, 2000; Wang et al., 2002; He et al., 2003) that inhibit axonal outgrowth and functional recovery following injury. These myelin proteins signal through the neuronal membrane bound Nogo Receptor (NgR) complex, which includes NgR1 (Chen et al., 2000; GrandPre et al., 2000), Lingo-1 (Mi et al., 2004), and p75^{NTR} (Domeniconi et al., 2002; Wong et al., 2002) or TROY (Park et al., 2005). Myelin protein engagement of the NgR complex activates RhoA, which induces ROCK-dependent phosphorylation of cofilin, thus actin depolymerization and growth cone collapse (He and Koprivica, 2004). When the RhoA pathway has been blocked following a CNS lesion, regeneration has been observed (Lehmann et al., 1999; Dergham et al., 2002; Fournier et al., 2003). The Paired immunoglobulin-like receptor B (PirB) is another receptor with high affinity for myelin inhibitory molecules that mediates outgrowth inhibition through dephosphorylation of tropomyosin receptor kinase (Trk) neurotrophin receptors (Atwal et al., 2008; Fujita et al., 2011).

Although following a severe CNS lesion a glial scar forms to repair the site of blood brain barrier disruption and limit inflammation (Rolls et al., 2009) it also represents a physical barrier to axonal growth (McKeon et al., 1991). The glial scar consists mainly of reactive astrocytes proteoglycans, and collagen, of which collagen IV provides the basement membrane scaffold for chondroitin sulphate proteoglycans (CSPGs) to bind (McKeon et al., 1991). It has been shown that CSPGs, of the glial scar, are inhibitory to axonal outgrowth in culture via binding to the recently discovered receptor PTPsigma (McKeon et al., 1991; Snow et al., 1996; Shen et al., 2009). Furthermore, removing CSPG glycosaminoglycan chains with chondroitinase ABC (ChABC) promotes functional recovery after spinal cord injury (SCI) (Bradbury et al., 2002).

More recently, the axonal regeneration field has partially shifted to elucidating the intrinsic growth capacity of CNS neurons. Not surprisingly, many well-defined embryonic developmental pathways have become validated in adult stem cell proliferation, regeneration, and differentiation (Tsokos et al., 1987; Terenghi, 1999; Esposito et al., 2005; Harel and Strittmatter, 2006; Reimer et al., 2009). This supports the theory that regeneration can be accomplished by a revival of developmental signals. It is believed that CNS neurons possess the ability, albeit limited, to regenerate. For example, the growth of CNS neurons into a PNS environment is possible (David and Aguayo, 1981; Benfey and Aguayo, 1982). It has been shown that the elongation of axons is linked to de novo transcription, and that blocking transcription immediately after injury inhibits regeneration from occurring (Smith and Skene, 1997). Furthermore, regeneration associated genes (RAGs) have been found following PNS injury that are absent following CNS injury in adults (Makwana and Raivich, 2005). These include GAP-43 and CAP-23 (growth cone proteins) (Chong et al., 1994; Mason et al., 2002), Sprr1a and SCG-10 (cytoskeletal proteins) (Mason et al., 2002; Starkey et al., 2009), Galanin (neuropeptide) (Holmes et al., 2000), brain derived neurotrophic factor (BDNF) (Tonra et al., 1998), as well as Chl1 and Lgals1 (cell adhesion molecules) (Zhang et al., 2000), and galectin-1 (McGraw et al., 2004). Furthermore, following a conditioning lesion, upregulation of cAMP activates protein kinase A (PKA), which in turn phosphorylates the transcription factor cAMP response element-binding protein (CREB), inducing transcription thus leading to neurite outgrowth (Neumann et al., 2002; Qiu et al., 2002; Gao et al., 2004). Interestingly, it has been shown that when neurons mature they lose their ability to overcome an inhibitory environment and this transition coincides with a decrease in cAMP levels (Cai et al., 2002). Another transcription factor thought to be involved in the retrograde signal following a conditioning lesion is signal transducer and activator of transcription 3 (STAT3) (Bareyre et al., 2011), which is activated via phosphorylation by tyrosine kinase Janus kinase (JAKs) associated with the receptor subunit glycoprotein 130 (gp130) initiated by the growth factor ciliary neurotrophic factor (CNTF) (Schwaiger et al., 2000). When either CREB or STAT3 pathways are blocked, regeneration in the CNS following a conditioning lesion is abolished (Gao et al., 2004; Qiu et al., 2005). Furthermore, it has also been shown in vitro that the conditioning lesion effect is completely abolished in a gp130 conditional knockout (Hyatt Sachs et al., 2010). However, Il-6 has been shown to be sufficient but not necessary for conditioning lesion-dependent axonal regeneration (Cao et al., 2006).

A number of injury responsive transcription factors have also been shown to affect neurite outgrowth and axonal regeneration, such as p53, c-Jun, activating transcription factor 3 (ATF3), nuclear factor of activated *T* cells (NFATs), nuclear factor kappalight-chain-enhancer of activated B cells (NF κ B), SRY-box containing gene 11 (Sox11), SnoN, and the Krüppel-like factor (KLF) family (reviewed in (Moore and Goldberg, 2011).

In addition to gene regulation it has been recently discovered that signaling pathways involving protein translation can induce axonal regeneration both in the PNS and CNS (Park et al., 2008; Christie et al., 2010; Liu et al., 2010). Classically known as a tumor suppressor, phosphatase and tensin homolog (PTEN), has been shown to counteract phosphoinositide 3-kinases' (PI3K) conversion of the lipid second messenger phosphatidylinositol (4,5) bisphosphate (PIP2) into phosphatidylinositol (3,4,5) triphosphate (PIP3) (Song et al., 2005). When PTEN is inhibited or deleted, there is an increase in activation of AKT by phosphorylation (Song et al., 2005). This in turn has led to an increase in axonal regeneration, which has been determined to be through the downstream activation of the mTOR leading to protein synthesis and cell growth (Guertin and Sabatini, 2007; Park et al., 2008; Christie et al., 2010; Liu et al., 2010).

RETINOIC ACID SIGNALING PATHWAYS IN AXONAL REGENERATION

Retinoic acid (RA), a degradation product of retinol after food ingestion, is a lipophilic vitamin A derivative that readily transverses the blood-brain barrier (Le Doze et al., 2000), and is a ligand for several nuclear receptors such as RA receptors (RAR, α , β , or γ , each of which have several isoforms due to alternative splicing) that play a role as transcription factors (Chambon, 1996). RA signaling is dependent upon its nuclear availability, controlled among others by retinol-binding protein-1 (RBP1), which is the carrier protein involved in the transport of retinol from the liver storage site to peripheral tissue and by cellular retinol binding protein (CRBP), which is the intracellular carrier involved in intracellular movement of retinol (Le Doze et al., 2000). Interestingly, the administration of RA not only activates the transcription factor RARB by direct binding that releases corepressors from promoters and recruits co-activators, but also increases its gene expression (Sucov et al., 1990; Leid et al., 1992). Importantly, a cAMP response element was found on RAR^β2 promoter, which binds CREB in response to cAMP and induces RARβ2 expression (Kruyt et al., 1992). In the presence of ligand, RA bound RAR β typically forms a heterodimer with retinoid X receptor (RXR, α , β , or γ) at RA response elements (RAREs) in gene promoters, recruits co-activators (CBP/p300, the CBP/p300 associated factor PCAF, SRC1, p160 pCIP, CoA, SWI/SNF, and ACTR) and activates transcription (Chambon, 1996; Glass and Rosenfeld, 2000). However, in the absence of ligand, RAR^β binds DNA in concert with co-repressors (nCo-R, SMRT, HDAC, and mSin3) and inhibits transcription (Glass and Rosenfeld, 2000). There have been several documented cases where RA bound RARB was found to occupy promoters independently from RXR and to repress transcription (Glass et al., 1989; Lipkin et al., 1992; Schoorlemmer et al., 1994). RA signaling typically involves direct transcriptional regulation, even though there are some less-defined cases involving non-transcriptional dependent RA signaling (Lopez-Carballo et al., 2002; Masia et al., 2007; Ohashi et al., 2009).

RA SIGNALING IN NEURITE OUTGROWTH AND AXONAL REGENERATION

Classically involved in development, neuronal differentiation, ventral neural patterning, and motor neuron specification (Maden et al., 1996; Diez del Corral et al., 2003; Novitch et al., 2003; Sockanathan et al., 2003), more recently RA also became a prime candidate to induce neurite outgrowth in neurons. The addition of exogenous RA and/or RAR β 2 has been shown to induce neurite outgrowth *in vitro* in dorsal root ganglion neurons (DRGs), adult cortical neurons, and postnatal cerebellar granule neurons (CGN) (Corcoran and Maden, 1999; Corcoran et al., 2000; Wong et al., 2006; Yip et al., 2006; Puttagunta et al., 2011). In embryonic and adult DRGs as well as CGN, RA upregulated the expression RAR β 2 in correlation with induced neurite outgrowth (Corcoran and Maden, 1999;

Corcoran et al., 2000; Puttagunta et al., 2011). Neurotrophins, such as nerve growth factor (NGF), essential in the development of the PNS (Piirsoo et al., 2010), have been shown to upregulate retinaldehyde dehydrogenase-2 (RALDH-2) and RAR^β2 expression to induce neurite outgrowth of adult mouse DRGs (Corcoran and Maden, 1999). Moreover, a RAR β agonist, but not a RAR α or RARy agonist, induced neurite outgrowth from embryonic DRGs (Corcoran et al., 2000). Additionally, both cortex and DRG explants from adult rats treated for 14 days with a RARB agonist induced neurite outgrowth in contrast to vehicle treated animals (Agudo et al., 2010). As a proof of principle that RA mediates its effect on neurite outgrowth exclusively through RARB, it was shown in postnatal CGN that RARB null CGN fail to extend neurites in response to RA (Puttagunta et al., 2011). Whereas, embryonic spinal cord explants respond to RA by inducing neurite outgrowth, adult spinal cord explants do not, and the limited RAR^{β2} expression in the adult spinal cord in comparison to the embryonic spinal cord is believed to be the reason for this difference (McCaffery and Drager, 1994; Zetterstrom et al., 1999; Corcoran et al., 2002; So et al., 2006; Yip et al., 2006). Furthermore, when RAR\u00c62, yet not RAR\u00f64, was virally introduced into the adult spinal cord in organotypic preparations, it induced neurite outgrowth (Corcoran et al., 2002). Therefore, it can be concluded that RA acts specifically through RARβ2 in inducing neurite outgrowth in both embryonic and adult neurons.

Having established RAR β 2 as the receptor responsible for RA induction of neurite outgrowth, the next logical step was to show that overexpression of RARB2 in vivo would induce axonal regeneration. RAR^{β2} lentiviral infection of adult rat DRGs induced axonal growth and functional recovery of injured sensory neurons into the dorsal root entry zone (DREZ) following dorsal root lesion (Wong et al., 2006). In support of these findings, it was also shown that RARB2 null mice have less axonal regeneration following a peripheral nerve crush versus wildtype mice (So et al., 2006). Additionally, RAR^{β2} lentiviral infection of the sensorimotor cortex three weeks prior to spinal cord lesion induced axonal and functional regeneration of the cortical spinal tracts (CST) in adult rats (Yip et al., 2006). Finally, RARB specific agonist, CD2019, brain treated rats following a dorsal column lesion had increased growth of CST axons and functional recovery of the forelimb versus those administered with vehicle (Agudo et al., 2010). These in vivo studies confirmed previous in vitro studies, showing that indeed RA-RARB signaling is involved in neurite outgrowth and axonal regeneration.

THE MOLECULAR MECHANISMS OF RA SIGNALING IN AXONAL REGENERATION

While it is known that RA signaling does induce neurite outgrowth and axonal regeneration only limited research has been done on the precise molecular mechanisms involved. It was shown that lentiviral RAR β 2 expression induces neurite outgrowth in DRGs and increases cAMP levels (Wong et al., 2006). The effect of RAR β 2 on neurite outgrowth was significantly decreased when an adenylate cyclase inhibitor, 2',5'dideoxyadenosine (DDA) or a cell–permanent inhibitor of cAMPdependent PKA was used (Wong et al., 2006). While this suggests that cAMP is involved in RA signaling, it is not sufficient, as use of dibutyryl cAMP in adult DRGs in the DREZ model did not promote functional recovery as robustly as RAR β 2 alone (Wong et al., 2006). In fact, it was shown in postnatal CGN that the positive affect of a RAR β agonist (CD2019) on neurite outgrowth was fully attenuated by a PI3K inhibitor but not a PKA inhibitor. Furthermore, it was shown that RAR β induced phosphorylation of AKT in CGN and *in vivo*, exclusively in injured neurons (Agudo et al., 2010). Interestingly, it has been previously shown that RA-RAR β 2/RXR directly activates AKT during the differentiation of human neuroblastoma cells (Lopez-Carballo et al., 2002; Ohashi et al., 2009).

Recently, we have discovered a direct transcriptional target of RA signaling in neurite outgrowth in an inhibitory environment. RA signaling increases neurite outgrowth, decreases RhoA activation and inhibits Lingo-1 gene and protein expression in a myelininhibitory environment (however, not on a CSPG-inhibitory substrate) in CGN, specifically through RARB. By in silico analysis, we discovered a RARE in the Lingo-1 promoter, which bound RARB but not RXR upon RA treatment in a myelin-inhibitory environment. Moreover, we showed this RARE to be functional by a luciferase assay that when mutated did not respond to RA signaling. Furthermore, addition of Lingo-1 in RA-treated CGN in a myelin-inhibitory environment abrogated RA-RARB induced neurite outgrowth. Finally, in vivo RA treatment decreased Lingo-1 protein expression following SCI in wildtype but not in RARβ null mice, providing physiological relevance to the in vitro findings (Puttagunta et al., 2011) (Figure 1A). Importantly, it has previously been shown that Lingo-1 antagonists promoted axonal sprouting, improved functional recovery, decreased RhoA activation, and increased oligodendrocyte and neuronal survival following rubrospinal or CST transection (Ji et al., 2006). It will be important to determine if this RA-dependent decrease in Lingo-1 is found exclusively in the neurons or also in the oligodendrocytes that make up the myelin sheathing and if RA signaling, such as seen with Lingo-1 inhibtion (Mi et al., 2005, 2007, 2009), leads to an increase in remyelination. In fact, for proper functionality following SCI, axons must re-grow, reinnervate their targets, and remyelinate their axons.

RA AND EPIGENETICS

Epigenetics changes can affect gene regulation by modifying the surrounding chromatin environment. They include DNA methylation and a number of histone post-translational modifications such as phosphorylation, acetylation, and methylation. For example, methylation of promoters or histones often inhibits gene expression, yet acetylation of histones or transcription factors induces gene expression by promoting transcription factor occupancy within a more accessible chromatin environment (reviewed in (Kouzarides, 2007; Mueller and von Deimling, 2009). These epigenetic changes can be transient or more long lasting thereby allowing extrinsic and intrinsic cellular signals to immediately and strongly influence gene expression (Guasconi and Puri, 2009). The acetyltransferases, CBP/p300, are essential for mammalian cell proliferation and development (Yao et al., 1998). Importantly, p300 appears to be a critical cofactor for RAR transcriptional activity, even leading to differentiation and somitic

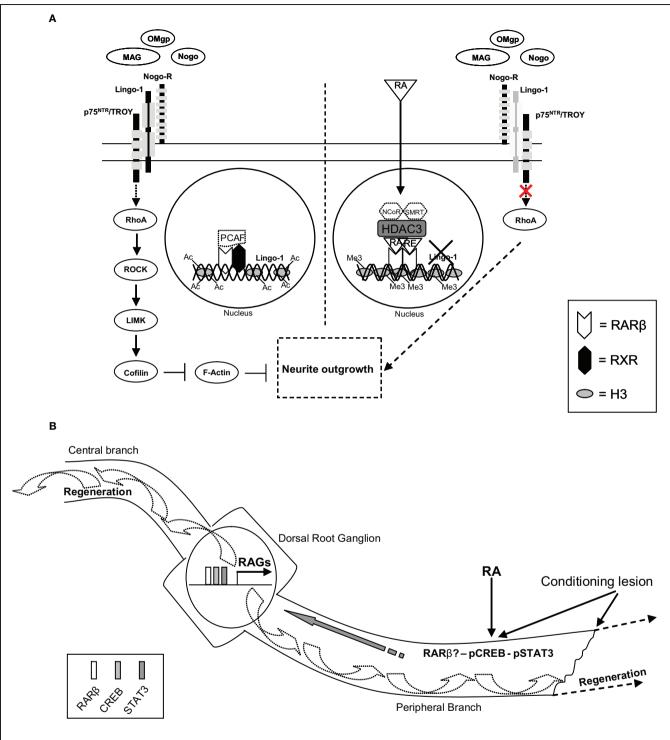
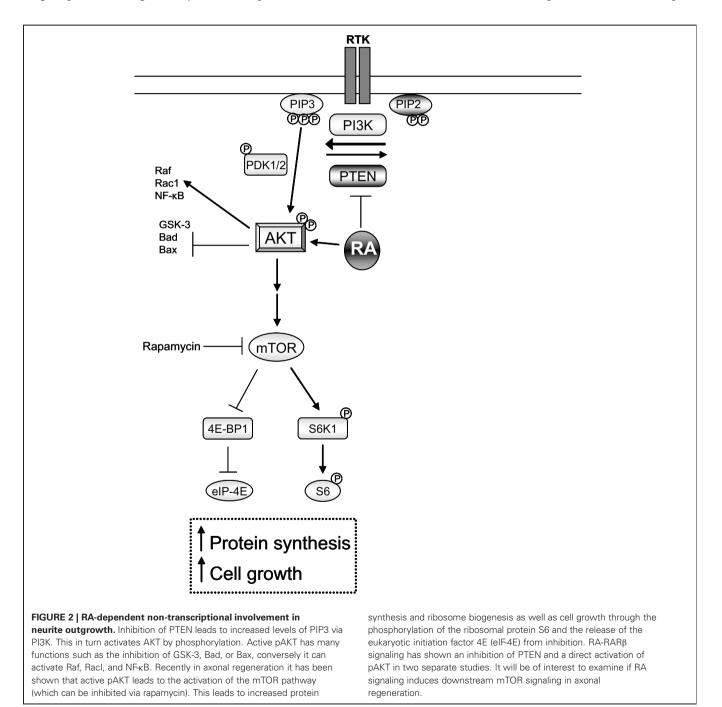


FIGURE 1 | RA-RAR β dependent direct transcriptional repression of inhibitory myelin signaling and a hypothetical schematic of RA involvement in retrograde signaling, RAGs expression, and axonal regeneration. (A) Following RA treatment HDAC3 binding to the RARE of the Lingo-1 promoter decreases acetylation of H3 lysine 9 and possibly displaces PCAF binding. Meanwhile, an increase in methylation of H3 lysine 27 is observed, coupled with strong binding of a RAR β homodimer to the RARE of the Lingo-1 promoter repressing Lingo-1 expression. This is believed to displace the equilibrium of the NogoR complex and block the activation of RhoA, thus leading to neurite outgrowth. Other possible members of this transcriptional repressor complex may include NCoR and SMRT (indicated with dashed lines). **(B)** Following a conditioning peripheral lesion and RA treatment an increase in RAR β , pCREB, and pSTAT3 levels are observed. All three have been shown to be vital to PNS axonal regeneration. CREB and STAT3 have been shown to play a role in retrograde signaling involved in the induction of RAGs expression prior to axonal regeneration. It will be intriguing to examine if RA signaling activates these three transcription factors to work in concert regulating RAGs expression to induce both axonal CNS and PNS regeneration *in vivo*.

symmetry (Kawasaki et al., 1998; Yao et al., 1998; Vilhais-Neto et al., 2010). Furthermore, in spinal motor neuron development, CBP is recruited by RA activated RAR to promoters for acetylation of histones, increasing accessibility of the promoter to transcription factors and inducing the expression of spinal motor neuron genes (Lee et al., 2009). RA signaling initiates PCAF binding to RAR/RXR heterodimers displacing co-repressor complexes and inducing gene expression (Blanco et al., 1998). Preliminary findings show (personal observations) that acetylation of histone 3 at lysine 9 (AcH3K9) at the RARE of the Lingo-1 promoter is significantly decreased upon RA treatment

in a myelin-inhibitory environment. In accordance, we saw an increase in histone deacetylase 3 (HDAC3, known to form a repressor complex with NCoR/SMRT at RA regulated genes) (Hartman et al., 2005) on the RARE of the Lingo-1 promoter. In addition, we observed an increase in methylation on H3 at lysine 27 (H3K27Me3, also known to be previously linked to RA mediated differentiation) (Gillespie and Gudas, 2007). These observations further extend upon recent published work showing a decrease in Lingo-1 expression upon RA treatment in a myelin-inhibitory environment (Puttagunta et al., 2011) (**Figure 1A**), however, further examination is required to determine complete



activator and repressor complexes involved in RA-dependent Lingo-1 regulation.

FUTURE PERSPECTIVES

In order to achieve functional axonal regeneration and recovery following SCI several obstacles need to be overcome: (a) an inhibitory glial environment, (b) excessive inflammation, (c) lack of neuronal intrinsic capacity for axonal outgrowth, and (d) axonal demyelination. In fact, RA signaling seems to have the potential to tackle several of these limitations.

RA has long been known for its anti-inflammatory benefits (Orfanos and Bauer, 1983; Nozaki et al., 2006). Interestingly, RA has been shown to inhibit the interferon-gamma induced inflammatory response in primary rat brain cultured astrocytes by inducing suppressors of cytokine signaling (SOCS3) and inhibiting JAK and STAT3 activation (Choi et al., 2005). Thus, it will be of interest to elucidate weather RA treatment *in vivo* following SCI reduces the formation of a glial scar by inhibiting the inflammatory response.

As the RA-RAR β pathway seems to overcome the myelin inhibitory environment, but not necessarily proteoglycan inhibitory signaling, combining ChABC with enhancement of RA-RAR β should strongly reduce the inhibitory signaling following SCI and further promote axonal regeneration than either does alone.

Given that neuronal overexpression of RAR β promotes axonal sprouting and regeneration of the CST and dorsal columns after SCI, it will be interesting to explore whether RA-RAR β is able to enhance the intrinsic ability of CNS neurons to regenerate by driving the expression of specific RAGs. Besides a possible direct transcriptional role for RAGs, RA-RAR β signaling may be involved in the cAMP-dependent retrograde signal following a PNS lesion that induces RAGs expression (Qiu et al., 2002). In fact, RAR β null mice have significantly reduced neurite outgrowth following peripheral nerve crush compared to wildtype mice (So et al., 2006), and, as mentioned previously, cAMP is induced by RAR β overexpression and involved in its ability to induce neurite outgrowth (Wong et al., 2006). Moreover, STAT3 is thought to be a retrograde signal following a PNS lesion and its inhibition

limits axonal regeneration (Qiu et al., 2002, 2005; Schweizer et al., 2002; Bareyre et al., 2011). While STAT3 is inhibited by RA signaling in astrocytes, conversely it is induced in a CNTF-dependent manner in differentiated neuroblatoma cells, retinal, and ciliary chick ganglion neurons (Malek and Halvorsen, 1997; Wang and Halvorsen, 1998). It will be of interest to inspect if RA-RARβ, CREB, and STAT3 transcriptional signaling work in concert to induce expression of RAGs and axonal regeneration (**Figure 1B**).

RA may also intersect with PTEN-mTOR signaling after SCI. The inhibition of PTEN results in the activation of AKT (pAKT) inducing axonal regeneration through mTOR by new protein synthesis and growth (Park et al., 2008). Accordingly, pAKT, which decreases after axonal injury (Liu et al., 2010), was shown to be only upregulated in injured CST axons following treatment with a RAR β agonist (Agudo et al., 2010). Therefore, it will be interesting to determine if AKT activation by RA signaling is through the inhibition of PTEN or the direct phosphorylation of AKT following axonal lesions and if this results in mTOR activation (**Figure 2**). In fact, dissection of this pro-regenerative molecular pathway may offer novel molecular targets for the effective enhancement of axonal regeneration.

Moreover, as RA signaling may take place via either paracrine or autocrine pathways and as Schwann cells, astrocytes, meningeal fibroblasts, and macrophages differentially express RA, RAR, or RALDH2 after PNS and CNS injury (Mey, 2006), further investigation of the relative weight of the two modes of action in axonal regeneration is warranted.

While RA pathways seem to positively affect several pathogenic aspects of SCI, further investigation is needed to clarify more direct transcriptional targets of RA-RAR β signaling following SCI and the interaction of RA pathways with other pro-regenerative signals. Only then, may we have the possibility to design RA signaling-dependent molecular therapies that may specifically enhance spinal axonal regeneration and recovery, with limited off target side effects.

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