

Signaling Pathways in Reactive Astrocytes, a Genetic Perspective

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Abstract Reactive astrocytes are associated with a vast array of central nervous system (CNS) pathologies. The activation of astrocytes is characterized by changes in their molecular and morphological features, and depending on the type of damage can also be accompanied by inflammatory responses, neuronal damage, and in severe cases, scar formation. Although reactive astrogliosis is the normal physiological response essential for containing damage, it can also have detrimental effects on neuronal survival and axon regeneration, particularly in neurodegenerative diseases. It is believed that progressive changes in astrocytes as they become reactive are finely regulated by complex intercellular and intracellular signaling mechanisms. However, these have yet to be sorted out. Much has been learned from gain-of-function approaches in vivo and culture paradigms, but in most cases, loss-of-function genetic studies, which are a critical complementary approach, have been lacking. Understanding which signaling pathways are required to control different aspects of astrogliosis will be necessary for designing therapeutic strategies to improve their beneficial effects and limit their detrimental ones in CNS pathologies. In this article, we review recent advances in the mechanisms underlying the regulation of aspects of astrogliosis, with the main focus on the signaling pathways that have been studied using loss-of-function genetic mouse models.

Keywords Astrogliosis · Gliosis · Reactive gliosis · Genetics

The CNS is susceptible to many types of pathological insults such as traumatic injury, ischemia, neurotoxic chemicals, tumors, infections, and neurodegenerative diseases. In all these cases, the response to damage includes at a minimum the activation of astrocytes, characterized by changes in their morphology and molecular expression profile. In addition to these changes, the response to damage often includes increased proliferation of several cell types, infiltration of leukocytes, scar formation, and neuronal death [1–4]. In this review, we limit our discussion primarily to what molecular mechanisms regulate astrocyte activation.

Many cell types including neurons, microglia, oligodendrocytes, endothelial cells, and leukocytes are likely to interact with astrocytes and influence the duration and amount of reactive astrogliosis [1–3, 5]. Although many extracellular signals are known to be up-regulated in different CNS pathologies, the cells that produce or respond to these signals in vivo in each case and how these signals affect astrocytes in particular is poorly understood. An informative approach will undoubtedly be to use mouse models of CNS damage combined with cell type-specific deletions of genes encoding intercellular and intracellular signaling components postulated to regulate astrogliosis. In this article, we discuss examples of what has been learned to date from such approaches.

In response to damage, the activation of astrocytes can be characterized by several features. First, the most commonly used marker of activated astrocytes is their upregulation of intermediate filaments (glial fibrillary acidic protein (GFAP), vimentin, and to some extent Nestin) coincident with cellular hypertrophy. Second, activated astrocytes are also likely to increase their expression of cytokines, chemokines, and extracellular matrix components [3]. However, in vivo data demonstrating this has remained sparse ([6]), since other cell types are also likely

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to secrete similar factors in response to damage making the cell source of these extracellular factors difficult to pinpoint. Hence, it remains to be determined just which factors are secreted by reactive astrocytes under what pathological conditions. Third, lineage tracing experiments in adult astrocytes in vivo using lentivirus and genetic approaches have shown that in some cases, such as after a stab wound injury, reactive astrocytes appear to become proliferative [7]. *Olig2*-expressing NG2 progenitors also proliferate in response to injury and may in some cases generate reactive astrocytes. NG2 cells have been reported to generate astrocytes in the cortex after stab wound or cryoinjuries [8, 9]. However, this is still a matter of debate as no new astrocytes appear to be generated from NG2 progenitors in other models of neurodegeneration, spinal cord injury, or cortical stab wound injury [10–12]. The differences may arise from the distinct combinations of fate mapping approaches and injury models. Further studies in which the progeny of defined glial cell types are marked are still needed to distinguish what cells can generate astrocytes under what injury conditions.

Aspects of reactive astrogliosis can exert both beneficial and harmful effects. For example, in transgenic mice in which reactive astrocytes are ablated or disabled by loss of GFAP and vimentin, traumatic injury leads to the lack of normal scar formation, prolonged and more widespread inflammation, and a failure to reconstruct the blood–brain barrier (BBB) and to maintain tissue integrity [13, 14]. However, ablation or impairment of reactive astrocytes also

leads to increased nerve fiber growth in the immediate vicinity of the injury site [13]. Despite this example, the positive roles and negative consequences of astrogliosis in most CNS pathologies remain largely unclear. For example, whether the reactive astrogliosis that occurs in neurodegenerative diseases is beneficial or detrimental is unknown. Moreover, the genetic pathways, of which there are likely many (examples are illustrated in Fig. 1), that regulate the different aspects and extent of astrogliosis in vivo remain largely unaddressed. An understanding of these pathways will be necessary to target specific aspects of reactive astrogliosis to maximize its beneficial effects and minimize its unwanted effects for each type of pathology.

NF- κ B and Inflammation Nuclear transcription factor- κ B (NF- κ B), which can be activated by stimuli associated with damage and in turn can promote expression of pro-inflammatory cytokines, is thought to play a central role in most inflammatory responses. When the CNS is damaged, the resident cell types that activate NF- κ B to mount a response have not been fully defined, but are likely to include astrocytes. Mice deficient for a major subunit of NF- κ B, p50, exhibit a smaller infarct size after middle cerebral artery occlusion (a model of stroke), which presumably reflects less neuronal death [15]. However, in this study, activated NF- κ B was only detected in neurons, not astrocytes, of control animals. Likewise, activated NF- κ B was only detected in neurons and microglia, but not astrocytes, after spinal cord injury in rats [16]. On the other

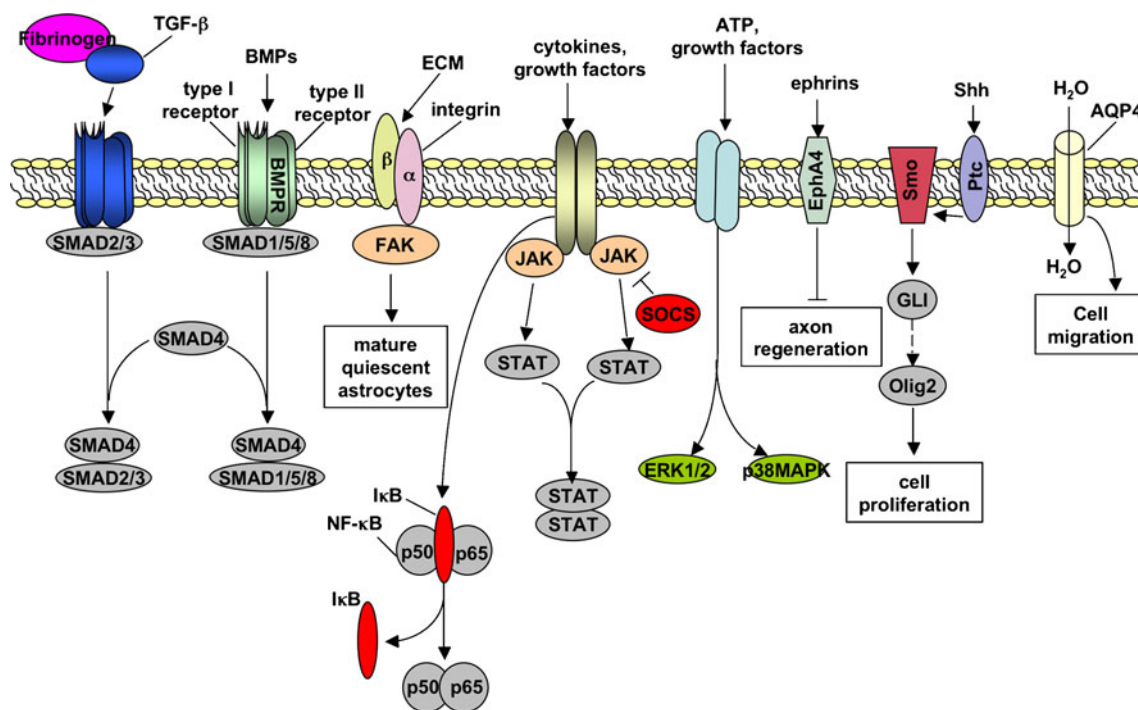


Fig. 1 Molecules and signaling pathways involved in activating different aspects of reactive astrogliosis. See text for details

hand, in a number of brain pathologies, NF- κ B levels are up-regulated in cell types that include astrocytes based on cell morphology [17–20]. Consistent with a role in astrocytes, mice in which expression of a constitutively active form of the inhibitor of NF- κ B, I κ B α is driven by the hGFAP promoter exhibit reduced lesion volumes, reduced glial scar formation, increased preservation of white matter, and a significant improvement in their ability to recover mobility after spinal cord injury [21]. Similarly, using the same transgenic mice, expression of pro-inflammatory genes, neuronal death, and overall disease severity were reduced following experimental autoimmune encephalomyelitis [22]. Importantly, however, the cell-type specificity of expression of the hGFAP-I κ B α transgene in these mice was not examined for adult tissues in vivo, leaving open the possibility that NF- κ B function was directly inhibited in cells other than astrocytes and that astrocytes were subsequently activated. Nevertheless, these studies together suggest that inhibition of NF- κ B signaling results in protective effects after injury and is a potential target for pharmacological intervention in CNS pathologies.

MAPK Signaling Reports describing whether the mitogen-activated protein kinase (MAPK) pathways are upregulated in astrocytes in vivo are mixed. For example, an increase in the level of phosphorylated extracellular signal-regulated kinase (ERK), a component of MAPK signaling, is detected specifically in astrocytes after traumatic spinal cord injury [23]. On the other hand, activation of p38 MAPK is detected solely in microglia, and not in astrocytes, in a mouse model of Alzheimer's disease [24]. Moreover, pharmacological inhibition of p38 MAPK or ERK appeared protective in some models of ischemia or brain injury [25–28], whereas inhibition of p38 MAPK appeared to worsen brain damage after ischemia in another study [29]. Further experiments are clearly needed to delineate which components of MAPK signaling are activated under what pathological conditions and in which cell types.

In addition, the ligands and receptors that activate components of MAPK signaling in astrocytes remain to be identified and will likely require deleting the genes encoding candidate receptors specifically in these cells upon CNS damage. Such candidates include the P2 purinergic receptor P2X7, FGF receptors, and EGF receptors, all of which are upregulated in astrocytes upon CNS damage [30–36]. The functionally relevant downstream genes that are regulated by MAPK signaling in vivo also remain to be identified.

SHH and Olig2 in Astrocyte Activation and Proliferation The factors that regulate cell proliferation in response to CNS damage are not fully known, but are likely to include the mitogen SHH and the basic helix–loop–helix transcription factor Olig2, which can be induced by SHH. Olig2

expression is upregulated in most GFAP-positive reactive astrocytes after injury, ischemia, and a model for multiple sclerosis [37–40]. Conditionally knocking out *Olig2* in neurons or oligodendrocytes does not affect reactive astrogliosis after cortical injury whereas conditionally knocking this gene out in all cortical cells results in reduced activation and proliferation of reactive astrocytes after injury [37]. This strongly suggests that *Olig2* is required specifically in astrocytes for their activation. SHH expression and activity (measured using a *Gli1*-luciferase reporter transgene) are induced after cortical injury and are required to promote the proliferation of *Olig2*-expressing cells [41]. Although likely, whether SHH-induced proliferation is dependent on Olig2 activity remains to be confirmed.

Endothelins in Astrocyte Activation and Proliferation Endothelins are a family of potent vasoactive peptides. Following brain injury, expression of the endothelin ET-1 and endothelin receptor ET-B is strongly upregulated in reactive astrocytes [42, 43]. Moreover, infusion of exogenous ET-1 or ET receptor agonists causes astrocytes to become hypertrophic and proliferate [44], whereas receptor antagonists attenuate astrocyte activation and proliferation after stab wound injury [45]. In these studies, although the effect of endothelins is likely to be directly on astrocytes, indirect effects through other cell types cannot be ruled out. Targeting of endothelin receptors directly in astrocytes and other cell types, for example, will clarify the role of endothelins in astrogliosis.

Fibrinogen, TGF β and Scar Formation Reactive astrocytes build a glial scar to contain damage and protect surrounding tissue. However, the glial scar is likely to form a biochemical and mechanical impediment to axon regeneration. Given the therapeutic potentials of manipulating scar formation, understanding how this process is regulated is of great interest.

After vascular damage with disruption of the BBB, the soluble blood coagulation protein fibrinogen leaks into the CNS and is converted to insoluble fibrin by the action of thrombin. Depletion of fibrin pharmacologically or genetically results in decreased inflammation and delayed onset of demyelination in mouse models of multiple sclerosis, which leads to increased lifespan and delayed symptoms [46]. Both in vivo and in vitro studies show that fibrinogen can activate astrocytes, leading to deposition of scar components after CNS injury. In fibrinogen knockout mice or mice treated with the fibrinogen-depleting agent anicrod, GFAP expression and deposition of neurocan, an inhibitory chondroitin sulfate proteoglycan (CSPG) secreted by astrocytes, are significantly reduced after stab wound injury [47]. However, the effect of fibrinogen on astrocytes is likely to be indirect through the transforming growth factor

beta (TGF β) pathway. Fibrinogen can act as carrier of latent TGF β to sites of injury. In both fibrinogen-deficient mutants and anicrod-treated mice, active TGF β levels after injury are dramatically reduced. Moreover, injection of fibrinogen into the cortex is sufficient to induce astrogliosis, while inhibition of the TGF β pathway abolishes the fibrinogen-induced effects on glia scar formation [47].

Multiple Roles for BMPs Interestingly, the bone morphogenetic proteins (BMPs), members of the TGF β superfamily of ligands, are also involved in astrocyte reactivity. Both BMP ligands and receptors are increased following CNS injury in neurons, astrocytes, oligodendrocytes, and microglia [48–51]. Using a GFAP-Cre transgene to conditionally delete the type 1 BMP receptor gene, *Bmpr1a*, specifically in astrocytes has no obvious effect under normal conditions. However, after spinal cord injury, *Bmpr1a* mutant astrocytes exhibit less hypertrophy and lower levels of GFAP than control astrocytes, leading to increased infiltration of inflammatory cells and greater loss of tissue [52]. Interestingly, mice that are null for a different type 1 receptor gene, *Bmpr1b*, exhibit an opposite phenotype after spinal cord injury including higher levels of GFAP, a more compact inflammation area, and smaller lesion volume [52]. These data suggest that BMPRI A and 1B signaling exert opposite effects on astrocytes following injury. However, because BMPRI B is a null allele that is lost in all cell types, the effects observed on astrogliosis in the *Bmpr1b* mutant may not be direct. Alternatively, the function of the two receptors within astrocytes may differ. In any case, the difference in phenotypes underscores the complexity of astrogliosis regulation.

STAT3 and Astrocyte Migration The Signal transducer and activator of transcription 3 (STAT3) is a member of the Jak-STAT signaling family. It may transduce signals for several cytokines and growth factors implicated in the injury response. The activation of STAT3 by phosphorylation increases markedly in astrocytes, microglia, endothelial cells, and neurons shortly after CNS insults [53–58]. GFAP-Cre and Nestin-Cre were used to conditionally delete *Stat3* in astrocytes from developmental stages onward [58, 59]. Although conditional deletion of *Stat3* specifically in adult astrocytes are still needed to exclude potential developmental and indirect effects, the results obtained from these studies nevertheless suggest a key role for this gene in the healing process after spinal cord injury. In the mutants, astrocyte hypertrophy and scar formation fail after injury and there was an increased spread of inflammation and lesion volume with impaired motor recovery [58, 59]. Consistent with these findings, the conditional deletion of *Socs3* in astrocytes, which negatively feedback on STAT3, exhibits the opposite

phenotypes [59]. Interestingly, STAT3 appears to function by promoting the migration of reactive astrocytes to the injury site [58, 59].

Aquaporin-4 and Astrocyte Migration Aquaporins (AQPs) are water channels that regulate water homeostasis in physiological and pathological conditions. Analysis of mutant mice lacking AQP4 indicates that this protein facilitates water flux into and out of the brain parenchyma and may be involved in brain edema in multiple pathological conditions [60–65]. In mutant mice lacking AQP4, glia scar formation after cortical stab injury is significantly impaired, perhaps due to a failure of astrocytes to migrate, as suggested by the behavior of AQP4-deficient astrocytes in culture [66, 67].

Integrins and Astrocyte Maturation Integrins act as receptors for extracellular matrix proteins and as such are likely to play multiple roles in response to CNS damage. These roles have yet to be clearly outlined, although β 1-integrin is likely to be required specifically in astrocytes for their normal maturation in the uninjured brain. Conditional deletion of the β 1-integrin gene in astrocytes and neurons during embryogenesis leads to many features of astrogliosis in postnatal and adult cortex and spinal cord including astrocyte hypertrophy, upregulation of GFAP, and microglia activation [68]. Deletion of this gene specifically in neurons, however, has no phenotype, suggesting that β 1-integrin is required specifically in astrocytes. A conditional knockout of *Fak* in embryos, a gene encoding a kinase that can be activated by integrins, also exhibits numerous GFAP-positive astrocytes in postnatal cortex [69]. However, it remains unclear to what extent the observed astrogliosis in the adult brain is due to developmental defects or due to abnormal astrocytes activation in the absence of obvious injury. In addition, whether the response of astrocytes to exogenously induced injury would be exacerbated in these mutants is not known. In any case, it is reasonable to postulate that the down-regulation of integrin after injury may be required for astrocyte activation.

Eph Signaling in Axon Regeneration Although factors such as myelin-associated glycoprotein, Nogo-A, oligodendrocyte myelin glycoprotein, CSPGs, and KSPGs have previously been identified as potential inhibitors of axon regeneration after injury, experiments aimed at perturbing these factors in vivo to promote axon regrowth have met with limited success [70–79]. The Eph receptor tyrosine kinase family and its ligands, the ephrins, regulate axon guidance through contact repulsion during CNS development. Normally, continued low level expression of Ephs and ephrins occurs in the adult CNS. After injury or in CNS diseases, many Ephs and ephrins are upregulated in different cell types including astrocytes, neurons, and

oligodendrocytes [80]. In vitro evidence has suggested that the increased Eph/ephrin signaling in CNS pathological conditions is inhibitory to axon regeneration of spinal cord, cortical neurons, and retinal ganglion neurons [81–84]. Analysis of *EphA4*^{−/−} mutant mice has clearly demonstrated the inhibitory effects of Eph/ephrin signaling on axon regeneration following spinal cord hemisection [85]. In *EphA4*^{−/−} mice, large numbers of newly formed axons grow through the lesion site. Functionally, several parameters of movement recovery are greatly improved in *EphA4*^{−/−} mice compared with the controls presumably due to improved axon regeneration. In addition to inhibiting axon regrowth, upregulation of Ephs and ephrins in astrocytes in CNS pathologies appears to play a role in mediating astrogliosis and scar formation. In *EphA4*^{−/−} mice, astrogliosis, assessed by GFAP expression, and scar formation are greatly reduced [85]. Together, these data indicate that Eph/ephrins are produced during astrogliosis and are inhibitory molecules for axon regeneration.

Proteoglycans A hallmark of reactive astrocytes is their secretion of proteoglycans, extracellular matrix molecules comprised of a protein core and variable glycosaminoglycan side chains. One major class of proteoglycans, CSPGs, are upregulated and secreted rapidly after a wide range of CNS injuries and can persist for months [86–88]. Although it is likely that these proteoglycans can interact with various signaling pathways [89], these have yet to be identified in vivo in the context of CNS damage. One exception is EGFR signaling, which can mediate inhibition of axon regeneration by myelin and CSPGs in a Ca²⁺-dependent manner [90]. Consistent with this finding, blocking EGFR function by an inhibitor can promote axonal regrowth and functional recovery following optic nerve and spinal cord injuries [90, 91]. Despite this example, further in vivo studies are clearly required to define the effects of CSPGs on reactive astrocytosis.

Conclusion and Prospects We currently only have a superficial grasp of the mechanisms regulating the initiation and extent of astrocyte activation under different pathological conditions. These mechanisms likely involve interactions with several cell types including infiltrating leukocytes, endothelial cells, microglia, and neurons. Our understanding of how the inflammatory response to CNS damage is eventually resolved, including the de-activation of astrocytes, is even more rudimentary. As with most biological processes, the state of astrocytes even under non-pathological conditions may be regulated by a balance between pro- and anti-reactive signals.

Nevertheless, with the advent of new genetic tools and experimental paradigms our knowledge of how astrocytes

and other cells respond to CNS damage is growing quickly. Of particular interest would be to develop a therapeutic ability to modulate the negative effects of certain aspects of inflammation to maximize tissue viability and in some cases promote neuronal and axonal regeneration.

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