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Regulation of spine and synapse formation by activity-dependent intracellular signaling pathways

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Abstract

Formation of the human brain during embryonic and postnatal development is an extraordinarily complex process resulting at maturity in billions of neurons with trillions of specialized connections called synapses. These synapses, composed of a varicosity or bouton from a presynaptic neuron that communicates with a dendritic spine of the postsynaptic neuron, comprise the neural network that is essential for complex behavioral phenomena and cognition. Inappropriate synapse formation or structure is thought to underlie several developmental neuropathologies. Even in the mature CNS, alterations in synapse structure and function continues to be a very dynamic process that is foundational to learning and memory as well as other adaptive abilities of the brain. This synaptic plasticity in mature neurons, which is often triggered by certain patterns of neural activity, is again multifaceted and involves post-translational modifications (e.g. phosphorylation) and subcellular relocalization or trafficking (endocytosis/exocytosis) of existing synaptic proteins, initiation of protein synthesis from existing mRNAs localized in dendrites or spines, and triggering of new gene transcription in the nucleus. These various cellular processes support varying temporal components of synaptic plasticity that begin within 1–2 min but can persist for hours to days. This review will give a critical assessment of activity-dependent molecular modulations of synapses reported over the past couple years. Owing to space limitations, it will focus on mammalian excitatory (i.e. glutamatergic) synapses and will not consider several activity-independent signaling pathways (e.g. ephrinB receptor) that also modulate spine and synapse formation [1,2].

Regulation of spine and synapse formation by small GTPases (see Figure 1)

Mature mushroom-shaped spines are unique micro-compartments that autonomously regulate the electrical and biochemical responses to synaptic activity. It is widely accepted that spine morphology and synapse function, via anchoring of key PSD proteins, is modulated by the actin cytoskeleton that is regulated largely by small GTPases (reviewed in [3]). The family of small GTPases (RhoA, Rac1, and Cdc42) cycle between an active GTP-bound form, promoted by guanine nucleotide exchange factors (GEFs), and an inactive GDP-bound form generated by GTPase-activating proteins (GAPs) that hydrolyze the GTP. GEFs and GAPs are major convergence points of upstream signaling pathways triggered by neuronal activity and growth factors that are often mediated by protein kinases. Major downstream effectors of these small GTPases that regulate the actin cytoskeleton include the p21-activated kinases PAK1 and PAK 3.

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Rac-1

Developing neurons exhibit spontaneous calcium spikes triggered by neurotransmitter-evoked activity [4] or growth factors such as brain-derived neurotrophic factor (BDNF) [5]. In mature neurons, Ca^{2+} entry through NMDA-type glutamate receptors (NMDARs) is crucial for synaptic plasticity. Elevated Ca^{2+} levels activate calmodulin (CaM) sensitive molecules including CaM-dependent protein kinases (CaMKs) [6]. CaMKs have been implicated in spine structural and synaptic plasticity, but the mechanisms were largely unknown until recently. CaMKII phosphorylates the RacGEF Kalirin-7 at Thr95, and they form a signaling complex with PSD95 and AMPA-type glutamate receptors (AMPA-Rs) [7•], the major transducer of rapid excitatory synaptic transmission in the mammalian CNS. In cultured neurons the general CaMK inhibitor, KN-62, suppressed NMDA-stimulated GEF activity of Kalirin-7, trafficking of AMPARs and spine maturation in cortical neurons. Mutant mice lacking Kalirin-7 show decreased spine density in CA1 pyramidal neurons, deficits in long-term potentiation (LTP), and impaired contextual fear learning [8], supporting the significance of Kalirin-7 in synaptic function *in vivo*. The RacGEF Tiam1 that associates with the NMDAR is also regulated by CaMKII, and dominant-negatives or RNAi inhibition of Tiam1 causes loss of NMDAR-dependent spine formation [9].

Since KN-62 and KN-93 are general CaMK inhibitors, physiological roles previously ascribed to CaMKII based on inhibitory effects of these reagents have recently been re-evaluated. Indeed, other members of the CaMK family, CaMKI and its upstream activator CaMKK, have been demonstrated to play crucial roles in multiple aspects of neuronal development including axon formation [10] and elongation [11], and activity-dependent dendritic arborization [12] and spine/synapse formation [13•]. Enhanced dendritogenesis is mediated via CaMKI-mediated activation of MEK/Erk [14] with resultant CREB-dependent transcription of Wnt-2 [12], which enhances dendrite formation, and microRNA132 levels. MicroRNA132 suppresses translation of p250GAP, resulting in Rac1 activation [15•,16] (see below) and/or inhibition of Rho A [17]. CaMKK/CaMKI form a signaling complex with the GEF betaPIX and its scaffold protein GIT1 that also binds PAK1 [13•]. Activated CaMKI phosphorylates betaPIX at Ser516 and stimulates its Rac-GEF activity. Most of the above studies on CaMKI were performed in cultured hippocampal neurons but have also been replicated in cultured hippocampal slices. Different isoforms of CaMKI appear to specifically mediate these developmental effects as determined by RNAi knockdown.

RhoA

In contrast to Rac and Cdc42, RhoA inhibits spine formation and maturation [18] and should be inactivated during synaptogenesis. It has been reported that the RhoA pathway is regulated through the polarity complex PAR6/atypicalPKC [19]. Overexpression of PAR6, but not the PAR6 mutants lacking binding sites for PKC and PDZ, promotes spine density whereas RNAi suppression of PAR6 reduces spine density and increases RhoA activity. This spine phenotype is rescued by pharmacological inhibition of the RhoA effector kinase, ROCK. One of the RhoA GAPs, P190GAP, is a possible link between PAR6/aPKC and RhoA activity. Indeed, P190GAP co-immunoprecipitates with the PAR6/aPKC complex, and PAR6-induced RhoA inactivation is blocked by P190GAP knockdown. However, the molecular details of how PAR6/aPKC regulate P190GAP remain to be clarified.

Trafficking and activity of AMPARs, which are crucial for maintenance of spines and synaptic plasticity (reviewed in [20,21]), are dynamic processes. What effectors modulate AMPAR actions in spine morphology? The RhoA-specific GEF, GEF-H1, is an AMPAR-interacting protein [22]. A dominant-negative construct and RNAi-mediated knockdown of GEF-H1 increases spine density and length, perhaps due to the ability of RhoA to inhibit Rac1 [23]. Pharmacological inhibition of AMPARs activates RhoA, inactivates Rac1, and

decreases spine density—these changes are eliminated by GEF-H1 knockdown. These data suggest that GEF-H1 is a key molecule linking AMPAR signaling and spine morphology. Although GEF-H1 translocates into dendritic spines in response to glutamate in a Ca^{2+} -dependent manner [24], the regulations of GEF-H1-mediated RhoA activation/Rac1 inactivation in response to neuronal activity in spines warrants further investigation. The small GTPases Ras and Rap also appear to govern the synaptic insertion and removal of AMPARs, respectively. The trafficking of AMPARs via Ras is dependent upon its level of activation—low levels of Ras activity lead to incorporation of GluR2 via the MEK-ERK pathway whereas high levels of Ras activity induce GluR1 incorporation via PI3 kinase-PKB/AKT pathway [25]. It should be noted that active Ras is not limited to individual spines but can also spread to adjacent spines where it is thought to reduce the threshold of activation for LTP [26].

The Rho-GTPase signaling pathways have been linked to several forms of mental retardation [27]. A recent study provides a mechanism by which a genetic deficit in the Rho-GAP, oligophrenin-1 (OPHN1) causes impaired synaptic development and glutamatergic dysfunction [28•]. Spontaneous neuronal activity is required for OPHN1 localization in spines where OPHN1 couples to AMPARs, and in developing neurons knockdown of OPHN1 reduces mEPSC frequency. NMDAR-induced endocytosis of AMPAR is suppressed by overexpression of OPHN1 or inhibition of the Rho/ROCK pathway with a ROCK inhibitor, indicating that OPHN1 signaling controls synaptic functions by regulating AMPAR stabilization in the synapse through suppressing the RhoA pathway.

Cdc42

Although multiple functions for RhoA and Rac in regulation of synaptogenesis have been described, roles of Cdc42 are less characterized. For example, expression of constitutively active Cdc42 (V12 mutant) does not appear to affect spine morphology or density [27]. However, a recent study identified Cdc42 as a synaptic palmitoylated protein that is essential for synaptogenesis [29•]. A brain-specific splice variant of Cdc42 is palmitoylated, and glutamate stimulation of cultured neurons causes a rapid depalmitoylation of Cdc42 within 5 min, leading to loss of Cdc42 from dendritic spines. Mechanisms by which neuronal activity induces depalmitoylation of Cdc42 and the role of this pathway in synapse function need further exploration.

MicroRNAs modulate spine formation and morphology

It is well established that localized protein synthesis, often initiated by activity-dependent regulation of translation factors, from selected mRNAs that are transported into dendrites and spines are important in modulating synaptic plasticity [30]. Recently, another mode of activity-modulated translational regulation in neurons via microRNAs (miRs) has been identified. MiRs are non-coding transcripts of approximately 19–24 nucleotides that regulate protein synthesis, either by destabilizing specific mRNAs or suppressing their translation [31]. MiRs have recently been implicated in several neuronal functions including apoptosis, neural patterning, and development of axons and dendrites [32]. Several miRs, together with their processing enzymes, such as Dicer, have been localized in dendritic spines, suggesting they may also play a role in spine/synapse formation and function. Dicer is present in the PSD in an inactive form where its RNAase III activity can be triggered in a Ca^{2+} -dependent manner via calpain cleavage, thereby converting pre-miRs into mature miRs [33]. Transgenic mice lacking forebrain Dicer in neurons lack several miRs, have a 50% decrease in cortical mass due to enhanced apoptosis in early development, and exhibit abnormal hippocampal patterning, decreased dendritic arborization and increased spine length in apical dendrites [34].

Studies in cultured neurons have identified some specific miRs and elucidated their targets that modulate dendrite and spine morphology (Figure 1). A miR that enhances dendritic development in cortical and hippocampal neurons is miR132 [15•,16]. Enhanced neuronal activity via the NMDA receptor and Ca²⁺-dependent activation of MEK/Erk by CaMKK/CaMKI stimulates CREB-dependent transcription of miR132 to suppress translation of the Rho family GTPase-activating protein p250GAP. Since p250GAP inhibits Rac1, activity-dependent decreases in p250GAP promote dendritic arborization, presumably by enhancing actin polymerization. However, it has been reported that RNAi suppression of p250GAP also activates RhoA and enhances spine size—this phenotype is rescued by dominant-negative RhoA [17]. Since RhoA is normally inhibitory to actin polymerization, the mechanism responsible for spine enlargement needs further clarification. Another miR, miR134, inhibits translation of LIM-kinase 1 that regulates the actin cytoskeleton [35]. Thus, basal miR134 tonically suppresses the size of spines, but this is reversed upon synaptic stimulation through release of BDNF that stimulates synthesis of LIM-kinase 1, perhaps through the mTOR pathway. Mechanisms by which BDNF inactivate the miR134 complex remain to be elucidated. A miR that regulates spine morphology without effect on spine density or dendritic arborization is miR138 [36]. MiR138 decreases spine volume through local suppression of acyl protein thioesterase 1 (APT1). APT1 catalyzes depalmitoylation of proteins, thereby modulating their membrane association. One of the relevant targets of APT1 is the small G protein subunit Gα13 whose membrane association is involved in Rho-dependent signaling.

It is clear that neurons contain numerous miRs, and an increasing number of targets are being identified [37]. Regulation of neuronal development as well as plasticity in mature neurons by miRs and their roles in neurodegenerative diseases is an emerging topic that promises to yield rich dividends.

LTP induces spine expansion and AMPAR trafficking

Several forms of synaptic plasticity result in morphological alterations of synapses, both pre- and postsynaptically. Mechanisms regulating trafficking of AMPARs during homeostatic synaptic scaling have recently been reviewed [38] and won't be dealt with here. It is known that LTP-inducing stimuli result in an initial robust and transient expansion of dendritic spines followed by a smaller but sustained increase in spine volume [39]. To date, few studies have examined the molecular mechanisms that underlie LTP-associated spine growth. However, several groups have now shown that pharmacological inhibition of CaMKs blocks the persistent but not the initial spine expansion associated with LTP [40,41••]. Additionally, the persistent increase in spine volume following LTP appears to require the kinase activity of CaMKIIα [42]. A likely target of CaMKIIα, Ser73 of PSD-95, a major synaptic protein, may regulate the termination of activity-dependent spine expansion [41••]. In hippocampal slice cultures, phosphorylation of PSD95 at Ser73 by CaMKII triggers the displacement of PSD95 and SHANK2 proteins from previously activated spines. SHANK proteins are thought to act as PSD scaffold proteins, linking many PSD proteins such as PSD95, GKAP, and Homer to both ionotropic and metabotropic glutamate receptors [43]. Overexpression of SHANK leads to enlarged spines [44], so translocation of this scaffold out of the spine may terminate spine growth. Indeed, expression of a phosphomimetic mutant of PSD95^(S73D) inhibits the increases in both synaptic strength and spine volume associated with LTP, suggesting that phosphorylation of PSD95^{S73} may contribute to activity-dependent changes in spine growth and synaptic strength [41••].

Recent data indicate that recycling endosomes bring AMPARs as well as the additional membrane necessary for activity-dependent spine growth and remodeling into spines during LTP [45,46]. It should be noted that a mobile pool of AMPARs has been shown to reside at

sites adjacent to the PSD and probably play a contributing role in supplying AMPARs to the synapse via means of lateral diffusion [47,48]. Two groups examining the exocytic delivery of AMPARs to the membrane have independently implicated Rab 11 in this process [49,50]. Rab proteins are crucial regulators of the endosomal membrane system and control membrane trafficking within the exocytic, endocytic and recycling pathways [51]. Using independent techniques, both groups demonstrate that trafficking of GluR1 into spines during LTP requires association with Rab11 containing vesicles. Work by Esteban's group has also indicated a role for Rab8 in the synaptic delivery of GluR1-containing AMPARs during LTP [52]. Not only are numbers of AMPARs modulated by synaptic activity, but their subunit composition and therefore biophysical properties may also be regulated. The AMPARs at the CA3/CA1 synapse in hippocampus is normally comprised of GluR1/GluR2 subunits, but certain LTP paradigms may promote synaptic incorporation of GluR2-lacking AMPARs ([53,54] but see [55,56]). This possibility is of particular interest as GluR2-lacking AMPARs have higher unitary conductances, are permeable to Ca^{2+} and have been implicated in several neuropathologies (reviewed in [57,58]). Synaptic insertion of GluR1, in addition to increasing synaptic strength, may provide a stable platform to promote spine expansion [59].

Although there is a consensus for the role of Rab proteins in endosome-mediated trafficking of AMPARs during LTP, it is not clear which actin-based myosin motors contribute to the trafficking of these vesicles. Candidate motors include myosin Va [49], Vb [50] and myosin VI [60]. In the first report, GST-pull downs demonstrated that Rab11 and GluR1, but not a truncated form of GluR1 lacking the last 30 amino acids, binds to the globular tail of Myosin Va. Furthermore, using dominant-negative constructs they show that inhibition of Myosin Va, but not Vb or VI significantly reduces AMPAR-mediated evoked responses, and Myosin Va RNAi impairs pairing-induced LTP. In an independent report, vesicular trafficking of metalloproteinase 9, which has been shown to contribute to persistent spine expansion and synaptic potentiation following LTP [61], into spines was also found to be associated with Myosin Va [62]. It should be noted, however, that synaptic transmission and LTP are not altered in mice that express a functional null mutation in the myosin Va gene [63].

By contrast, Wang *et al.* [50] show that endogenous myosin Vb may be the relevant motor protein induced by LTP. Using live-cell imaging, these authors demonstrate that a population of myosin Vb colocalizes with recycling endosomes following NMDAR activation. Binding of MyoVb to Rab11-FIP2 was also shown to be Ca^{2+} dependent, providing a mechanistic link between NMDAR activation and association of myosin Vb to Rab11-FIP2. Furthermore, acute inhibition of MyoVb by a nonhydrolyzable ADP analog that suppresses the mobility of a MyoVb mutant along F actin, impaired LTP.

If indeed myosin V motors are involved in mediating the enduring synaptic and structural changes associated with LTP, it will be important to learn the mechanism(s) by which these motors are activated. Both groups assume that the trigger leading to myosin activity is increased intracellular Ca^{2+} . While this assumption is in line with the necessity for calcium in LTP, work in the myosin field has shown that Ca^{2+} binding to the calmodulin light chains actually impairs myosin motility [64]. Therefore, how could calcium influx, necessary for LTP, also be triggering myosin-dependent vesicular movement? According to Sellers *et al.*, myosin motility is most probably being regulated by cargo receptor proteins, where cargo binding to the globular tail domain increases its enzymatic as well as mechanical activity through steric or allosteric regulation [65]. Which cargo-receptor proteins and whether disruption of these proteins interferes with LTP will be of future interest.

LTD and morphological plasticity of spines?

The accepted model to date has been that bidirectional alterations of synaptic strength occur in parallel with corresponding changes in spine geometry. This concept is supported by studies on LTP (see above), and previous studies indicated that LTD is accompanied by a shrinkage in dendritic spines [66,67]. Whether changes in structural plasticity are necessary to adjust synaptic weights or vice versa, however, still remains an open question. Recently, two independent groups challenged this model by testing whether physiological and morphological changes associated with LTD are linked to either a common signaling mechanism or physical process. The first report [68•], using combined live-cell electrophysiology and two-photon microscopy, determined that LTD at identified parallel fiber–Purkinje cell synapses was not associated with structural changes in dendritic spines. These data were not simply the result of an inability to detect changes in spine volume since these same authors were able to detect reductions in spine size associated with depolarization-induced spine retraction. It should be noted, however, that the later manipulation failed to induce changes in evoked synaptic responses, further illustrating a clear disconnect between LTD and spine morphological plasticity.

A separate study [69] using acute hippocampal slices found that when neurons were internally loaded with a phospho-cofilin peptide, to prevent actin depolymerization, LTD was no longer associated with a reduction in spine size, while synaptic depression was still observed. Additionally, application of insulin, which has been shown to result in synaptic rundown and rapid clathrin-mediated internalization of synaptic AMPARs failed to elicit changes in spine volume despite reducing synaptic responses. Alternatively, inhibiting endocytosis with D15 peptide, to disrupt interaction of dynamin with amphiphysin, enhanced synaptic responses but without effects on spine size. Therefore, the lack of association found between LTD and spine morphology as seen in these studies suggests that reductions in synaptic strength may not necessarily correlate with changes in spine morphology as previously thought.

There is now growing evidence that morphological plasticity associated with LTD may be more prominent in presynaptic boutons. Using combined two-photon time lapse microscopy and electrophysiology to monitor labeled presynaptic boutons in hippocampal slice cultures, LTD was found to induce a significant reduction in the size of presynaptic boutons [70•,71]. Therefore, LTD induction can elicit structural remodifications on both sides of the synapse leading to a separation between post and presynaptic contacts. In addition, Becker *et al.* [71] observed that a major contribution to the loss of synaptic contacts was due to the loss of the partnered presynaptic bouton. Furthermore, they found that while the loss of dendritic spines led to reductions in the size of paired boutons, spine volumes did not change following the gain or loss of partnered boutons. What is the relationship between the loss of synaptic contacts and LTD? Bastrikova *et al.* [70•] found that the greater the magnitude of synaptic depression they observed following LTD, the greater the reduction in synaptic contacts. These findings suggest that under certain conditions activity-dependent structural changes associated with LTD may predominate in the presynaptic bouton. Whether LTD expression is predominantly due to post-synaptic or presynaptic mechanisms appears to depend on several factors such as brain area, mode of induction, and genetic background (e.g. state of FMR1 gene; reviewed in [72,73]). Future mechanistic studies regarding the signaling involved in determining how an existing synaptic connection is eliminated will be needed. Furthermore, it will be important to understand the contributions or circumstances leading to presynaptic terminal withdrawal, reduced spine size, and AMPAR properties (i.e. internalization, phosphorylation) as a result of LTD stimuli.

Future directions

It is clear that signaling pathways that regulate the actin cytoskeleton via the small GTPases are major players in dictating spine morphology. In fact, several neuropathologies are associated with mutations in these proteins that lead to abnormal spine/synapse maturation. As described in this review, these signaling pathways act on multiple GEFs and GAPs to fine-tune the balance between opposing roles of Rac1 and RhoA. Furthermore, miRs have recently been found to be important regulators of these GTPases. As more cellular targets of various miRs become identified, their roles in spine/synapse maturation will expand. Although numerous regulators of RhoA and Rac1 have emerged, the molecular details of how these regulatory proteins act in concert to promote normal spine/synapse maturation is still lacking. Application of new technologies in live-cell imaging of signaling molecules should further define these intricate cross-talk mechanisms.

Activity-dependent changes in spine/synaptic structure continue to be an area of intense research given that perturbations in signaling molecules associated with activity-dependent structural plasticity lead to cognitive deficits. While the molecular mechanisms contributing to activity-dependent spine expansion/retraction are now starting to emerge, they have also lead to more questions. For example, recent data have implicated a role for myosin-based motors in the establishment of LTP; however, the mechanisms linking LTP and myosin mobility are unclear. Since Ca^{2+} does not appear to directly enhance myosin mobility, it will be interesting to determine whether any of the cargo-receptor proteins, which most probably effect myosin mobility, are regulated by shifts in intracellular Ca^{2+} . Finally, the association between activity-dependent synaptic plasticity and spine morphology appears to be more complex than previously thought. While LTP remains associated spine enlargement and AMPAR recruitment, structural modification of spines as a result of LTD-inducing stimuli is more complex. Current research has now shown that some forms of synaptic depression are not associated with alterations in spine size, questioning whether postsynaptic morphological changes are necessary for LTD. Future studies examining modulation of presynaptic structures will probably reveal novel mechanisms associated with activity-dependent synaptic pruning.

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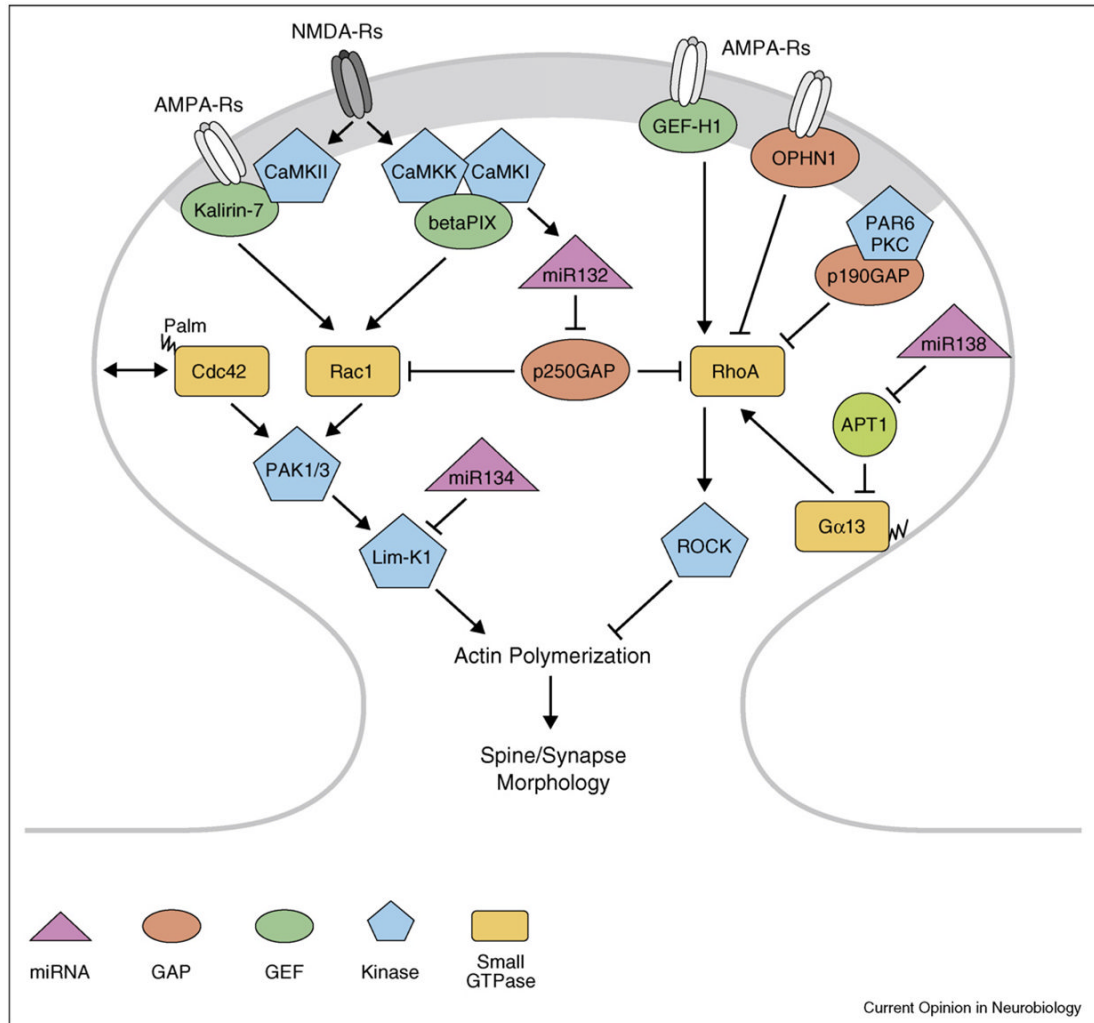


Figure 1. Schematic outline of small GTPase regulation of spine morphology via remodeling of the actin cytoskeleton. See text for details. Palm, palmitoylation; miRNA, microRNA.