

# Molecular Regulation of Dendritic Spine Shape and Function

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## Key Words

Actin · Postsynaptic density · Glutamate receptors · Synaptic plasticity · Calcium homeostasis

## Abstract

Dendritic spines are discrete membrane protrusions from dendritic shafts where the large majority of excitatory synapses are located. Their highly heterogeneous morphology is thought to be the morphological basis for synaptic plasticity. Electron microscopy and time-lapse imaging studies have suggested that the shape and number of spines can change after long-term potentiation (LTP), although there is no evidence that morphological changes are necessary for LTP induction and maintenance. An increasing number of proteins have been found to be morphogens for dendritic spines and provide new insights into the molecular mechanisms regulating spine formation and morphology.

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## Introduction

Dendritic spines are morphological specializations of neuronal synapses. They are formed by small protruding pieces of membrane with a total volume ranging from less than 0.01 to 0.8  $\mu\text{m}^3$  [1, 2] and contain 90% of the excitatory synapses in mature brain. Spines were first observed by the Spanish neuroscientist Ramon y Cajal [3] using the Camillo Golgi 'reazione nera' protocol and, since then, many scientists and neuroscientists have tried to understand their cerebral function. Most of the principal glutamate- or GABA-releasing neurons (such as the pyramidal and Purkinje neurons), but not many other neuron classes (such as GABA-releasing interneurons), have synapses on dendritic spines. Dendritic spines are present at the squid giant synapses [4] but are rarely found in other lower organisms (such as *Drosophila melanogaster* or *Caenorhabditis elegans*), suggesting that they may have developed early in evolution in order to implement more complex nervous system functions. This review focuses on the cellular and molecular mechanisms that regulate spine shape and function. Recent knowledge on molecules that regulate spine morphology is also improving our understanding of their roles in brain plasticity and function.

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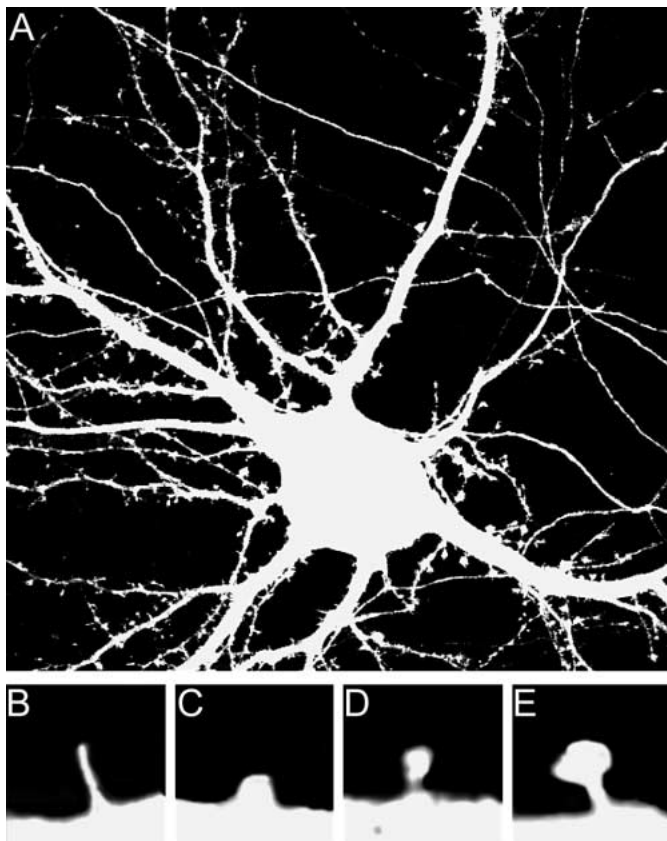


Fig. 1. Dendritic spine morphology can be observed in a hippocampal neuron transfected with GFP. A Low magnification of a GFP-transfected hippocampal neuron in culture. B-E Examples of different dendritic spine shapes viewed at high magnification: filopodium (B); stubby spine (C); thin spine (D), and mushroom-shaped spine (E).

### Structure and Development of Dendritic Spines

Although spines may have different shapes, they are essentially formed by a neck and head attached to the dendritic membrane. Extensive electron microscopy studies of brain tissue have shown that they may be thin, stubby, mushroom-shaped or cup-shaped (fig. 1) [1, 5–8] and that these different shapes can be found at the same time with the same dendrites [2, 9]. This crude classification underlines the multiple forms and dimensions of the spinal head and neck: the classic mushroom-shaped spines have a large head and narrow neck, whereas thin spines have a smaller head and narrow neck and stubby spines show no obvious constriction between the head and its attachment to the dendritic shaft. However, this static view does not reflect the real *in vivo* situation because at least in devel-

oping neurons, about 50% of the spines change their shape over periods of minutes or hours and the other 50% retain their morphological classifications [10]. Spine motility is developmentally regulated and, in mature neurons, there are fewer transitions between categories [10, 11]. The fixed structures observed through an electron microscope are therefore probably a representative picture of spines at that particular moment during a possible morphological transition. In the brain, spines and presynaptic boutons are surrounded by glial cells in such a way as to form an intercommunicating tripartite complex and at least half of the circumference of about 57% of the synapses is covered by astrocyte processes [12]. Typical mature spines have a single excitatory synapse located at the head, but the same spines can also have an inhibitory input [13, 14]. However, spines essentially represent the main unitary postsynaptic compartment for excitatory input.

Remarkable differences can also be seen in the intracellular composition of each spine which consists of the postsynaptic density (PSD) facing the presynaptic bouton and a cytoskeletal structure mainly formed by F-actin. About 50% of the spines on hippocampal CA1 cells and virtually all Purkinje cell spines also have a smooth endoplasmic reticulum (SER) [15], some pyramidal cell spines contain the spine apparatus, an organelle formed by two or more disks of SER separated by electron-dense material [16]. Large spines usually have a proportionally large synapse and contain different organelles. Both the SER and spine apparatus are usually associated with larger spines and are formally absent in small spines [15]. As SER is known to play a role in  $Ca^{2+}$  handling [17, 18] differently sized spines may have different ways of controlling calcium homeostasis (see below). The final components are ribosomes in which proteins can be specifically synthesized in close relationship with each spine. Polyribosomes are frequently found in the spines of different neuron subtypes [19, 20].

PSD is an electron-dense thickening of the membrane usually found at the head of the spine, where the synaptic junction is located. It usually occupies about 10% of the surface area exactly opposite the presynaptic active zone, and is the site for the  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazole propionic acid (AMPA) and N-methyl-D-aspartate (NMDA) glutamate receptors. The PSD is probably the most complex spine organelle in which hundreds of components (including receptors, cytoskeletal and adapter proteins, and associated signaling molecules) are involved in a number of signaling pathways controlling synaptic plasticity [21, 22]. Proteins are associated with

each other in a complex based on a series of protein–protein interaction domains of which the PDZ domain is one of the most important [23].

Three immunogold-labeling studies of the hippocampus [24–26] have shown that the number of AMPA and NMDA receptors is proportional to the PSD area and spine volume because their density is constant within the PSD. The PSD and spine volume are also proportional to the area of the active zone, which is itself proportional to the number of docked vesicles [27], which in turn correlates with the amount of neurotransmitter release per action potential [28]. All of these data suggest that large spines represent stronger synapses for both presynaptic and postsynaptic properties, and that the growth of the spine head during development probably correlates with a strengthening of synaptic transmission.

Three-dimensional reconstructions of the PSD show that it may have a number of shapes: as macular (i.e., plaque-like and uninterrupted) or perforated (i.e., fenestrated appearance), horseshoe-shaped or segmented. Interestingly, perforated PSDs have been consistently found to be associated with more AMPA receptor immunoreactivity than non-perforated PSDs [24]. The mushroom-shaped and largest spines are more likely to contain perforated PSDs [29], whereas virtually all thin spines have macular PSDs [8]. The shape of the PSD is not fixed and may change with modifications in the strength of synaptic activity, such as during long-term potentiation (LTP) [30–32]. Perforated PSDs may reflect enhanced AMPA receptor insertion into the postsynaptic membrane (which occurs during synaptic potentiation or growth) as an early phase of synapse duplication and spine division, or may be the morphological correlate of the enhanced receptor turnover at the PSD that might occur during LTP [33–39].

In adult hippocampal CA1 pyramidal and granule cells, dendritic spine density ranges from two to four spines per micrometer of dendrite [8, 29, 40, 41], whereas it is more than ten spines per micrometer in Purkinje cells [42, 43]. The density is lower in dissociated cultures of hippocampal neurons usually being 3–4 spines for each 10  $\mu\text{m}$  of dendrites [44] (fig. 1A). The two-dimensional nature of cultured neurons provides a viable and convenient means of studying the factors regulating the development and functions of dendritic spines. In the brain, the density of spines is not homogeneous throughout the dendritic tree but increases at each layer, thus suggesting that the afferent system independently regulates different parts of the dendritic tree [45, 46]. Spine density also varies across cortical areas. Apparently, in macaque monkeys

and humans the density on basal dendrites in the cortical areas of the frontal pole and orbitofrontal cortex is generally higher than in neurons of the primary visual and somatosensory cortices [47, 48], and it has been hypothesized that there may be a link between spine density and overall number and level of cortical processing in these regions. Although not totally proved, it is believed that these ‘higher’ order areas are involved in a greater degree of convergent processing, which may create a need for more synapses and therefore more spines.

How are mature spines formed? Early spines are often very long and have frequent filopodia-like shape (fig. 1B) but, later during development, their mean length decreases and the number of filopodia is greatly reduced. Three major changes can be observed during the maturation process: an increase in spine density; a decrease in overall length, and a decrease in the number of dendritic filopodia with a simultaneous decrease in spine motility [11, 49]. Recent studies have shown that filopodia rapidly protrude and retract from dendrites, especially during the early stages of synaptogenesis [50–52], and it is widely believed that dendritic filopodia are the precursors of dendritic spines, various hypotheses as to how the transition from filopodia to spines takes place have been put forward.

In the first proposed model, filopodia actively seek out synaptic partners in the developing neuropil and when a filopodium makes contact with an axon, it becomes shorter and draws the axonal terminal closer to the dendrite shaft. Subsequently, a fully mature synapse is formed on the spine head, spine motility gradually decreases and the structure is stabilized [50–52]. However this model does not explain why the density of asymmetric synapses is much higher on dendritic shafts than on filopodia during early development [50, 53]. Another model has proposed that, after their contacts with axons, the filopodia retract completely, thus leading to the formation of an asymmetric shaft synapse from which a spine emerges with a mature synapse at its head [2, 50]. In a recent study using two-photon time-lapse microscopy, Parnass et al. [10] demonstrated that stubby and other types of spines can originate from filopodia in developing hippocampal neurons and, depending on the state of the afferent input, spines turning into filopodia were also observed. In this model a filopodium is like a spine in a state of morphological instability and is not a necessary intermediate for spine formation. In other words, the transition from filopodia to spines is less likely to be predestined than a reversible process regulated by local factors, such as synaptic activity.

In two recent time-lapse studies spine morphogenesis and PSD-95 fused to green fluorescent protein (GFP), a PSD marker, were simultaneously imaged. In both cases the authors suggested that synapses initially form on dynamic filopodia-like spines that soon convert directly into stable spines at the same time as the formation of postsynaptic specialization [54, 55]. However, only Marrs et al. [54] showed the emergence of stable spines from shaft synapses.

Overall, these data suggest that it is probably still incorrect to assume that all spines go through the same stages (beginning as filopodia, proceeding to thin or stubby spines, and ending as mushroom spines), and that spine maturity does not necessarily correlate with spine morphology.

### Spine Function and Morphogenic Spine Molecules

What is the function of dendritic spines in synaptic transmission? The currently prevailing opinion is that the spine may function as a microcompartment for segregating postsynaptic chemical responses [56, 57]. There is no clear evidence that spines work as electrical compartments [57], at least in the case of CA1 synapses [58]. Interestingly, the shape of the spine neck might control postsynaptic calcium responses: in spines with long necks  $[Ca^{2+}]$  rises faster and decays slower than in those with short necks [59, 60], and spine motility induces modifications in neck length that correlate with altered calcium kinetics within the spine [61, 62, but see also 63].

In two papers [63, 64] it has recently been proposed that  $[Ca^{2+}]$  in spines may regulate the activation of LTP or LTD induced by NMDA receptor stimulation. The different kinetics of spine calcium concentration obtained using different stimulation protocols might activate CAMKII and LTP or calcineurin and LTD [63]. Differences in  $[Ca^{2+}]$  kinetics and synaptic plasticity also depend on the location of the spines along the dendritic tree [64],  $[Ca^{2+}]$  decay is faster on distal spines whose synapses are less susceptible to depression. Under physiological conditions, the spine neck acts as a barrier to diffusion and its head is isolated from the dendrite during an action potential. Spines can act as semi-autonomous chemical compartments separated from the dendritic shaft by a thin neck of up to a few micrometers in length. In brief, they can compartmentalize calcium and other second messengers (such as  $IP_3$ , cAMP and cGMP), and this function is regulated by their morphology. However, their physiological signifi-

cance for brain function will probably remain a major unresolved question until all of the molecular mechanisms regulating spine formation and morphology have been clarified.

Considerable efforts have recently been made to characterize the molecular mechanisms controlling spine morphology, formation and plasticity (fig. 2). Most of these studies were carried out using cultured neurons in which putative molecules were overexpressed and the modifications in spine morphology were observed using GFP overexpressed protein. Few molecules have been analyzed in genetic experiments. One of these, the fragile X mental retardation protein (FMRP) has a direct connection with a human pathology, in which its deletion is responsible for severe mental retardation. An abnormal density of dendritic spines has been observed in human patients affected by the fragile X syndrome [65, 66]. In KO mice with the *Fmr1* gene, the number and length of dendritic spines are greater than in normal animals [67], especially during the first 2 weeks after birth when the spines develop [49]. FMRP is an mRNA-binding protein that may regulate the synthesis of the molecules involved in the developmental stabilization, elimination or pruning of dendritic spines or synapses.

Defects in spine morphology during the development process have been observed in two other KO animals, one for the NR3A and the other for spinophilin. The NR3A KO mice show enhanced NMDA responses associated with increased spine density during early postnatal development [68]. Similarly, a marked increase in spine density during development in vivo and altered filopodial formation in cultured neurons have been observed in spinophilin KO mice [69].

As filamentous actin (F-actin) is the main constituent of the cytoskeleton of dendritic spines, actin-binding proteins are likely to be involved in spine formation and shape [70]. A number of actin-binding proteins are highly localized to spines:  $\alpha$ -actinin [71], drebrin [72], spinophilin/neurabin II [73], SPAR [74], adducin [75], synaptopodin [76] and cortactin [77]. There is evidence that some of them are involved in regulating spine morphology. We have already mentioned spinophilin KO mice [69]. Hayaishi and Shirao [72] have shown that the overexpression of drebrin in cultured cortical neurons (whose localization to spine depends on the actin-binding domain) induces a statistically significant elongation of dendritic spines. However, how drebrin or spinophilin regulates actin dynamics is still not totally clear.

Small GTPases of the RHO/RAC/CDC42 family are regulators of the actin cytoskeleton in many cell types.

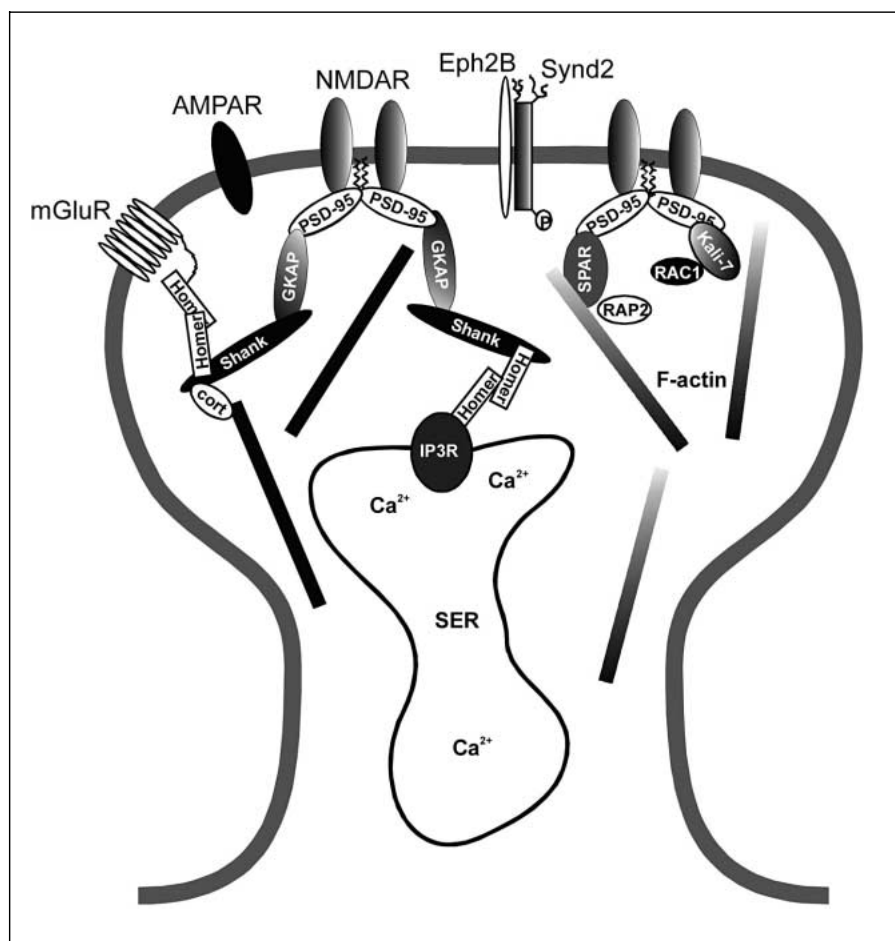


Fig. 2. Schematic representation of a mushroom-shaped spine and some of the proteins within the PSD and postsynaptic membrane that regulate spine morphology. NMDAR = N-Methyl-D-aspartate receptor; AMPAR = AMPA receptor; F-actin = filamentous actin; GKAP = guanylate-kinase-associated protein; Kali-7 = Kalirin-7; mGluR = metabotropic glutamate receptor; SPAR = spine-associated RapGAP; IP3R = IP3 receptor; Cort = Cortactin; Synd2 = Syndecan-2.

Transgenic mice overexpressing active Rac1 have an increased number of smaller dendritic spines in developing and mature cerebellar Purkinje cells, which form supernumerary synapses without modifying the dendritic trees [78]. A similar effect has been observed in pyramidal neurons transfected with constitutively active Rac1, which showed an increase in the number of filopodia-like processes and lamellipodia-like ruffles [79], whereas transfection with dominant-negative Rac1 reduced the number [80]. The involvement of Rac1 in spine formation has been further supported by Penzes et al. [81] who overexpressed Kalirin-7 (a guanine nucleotide exchange factor (GEF) for Rac1) in cultured cortical neurons and found that Kalirin-7 increased the number and size of spine-like protrusions. This effect is dependent on GEF activity because the overexpression of a Kalirin-7 mutant lacking GEF activity reduced the number of spines [81]. Interestingly, by binding to PSD-95 (and other PDZ domain-containing proteins) Kalirin-7 remains concen-

trated to the postsynapse compartment where its activity is required, whereas a mutant that is unable to interact with PDZ proteins remains in the cell soma and induces the local formation of aberrant filopodial neuritis [81]. The effect of RhoA on dendrite spines is less clear. Tashiro et al. [80] showed that the overexpression of constitutively active RhoA can reduce the number of spines only in a subset of neurons whereas the inhibition of Rho activity may lead to the formation of supernumerary spines or the elongation of spine necks. It has been proposed that Rac and Rho signaling might antagonize the molecular mechanisms regulating spine formation and/or growth [80].

Another PSD-95 and actin-binding protein, SPAR, has been found to be involved in regulating spine shape [74]. SPAR is a GTPase-activating protein (GAP) for Rap, binds to the GK domain of PSD-95 and is enriched in spines. In COS-7 cells SPAR dramatically reorganizes F-actin into large aggregates, dispersed clusters or smaller

well-defined star-like clusters. PSD-95, which binds to SPAR with the GK domain, is recruited to these clusters. SPAR seems to bind directly to actin through two distinct actin-interacting domains (Act1 and Act2), which are separated by the GAP domain. The overexpression of SPAR in neurons causes the enlargement and elaboration of spine heads, thus making the spines more irregular, thorny and multilobed. SPAR-enlarged spines are also frequently associated with multiple synaptic contacts and, as many of them appear to be branched, they might be dividing. Three SPAR domains are required to induce an increase in the size of spine heads: the GAP domain, the second actin-binding domain (Act2), and the domain that binds to PSD-95 (GK-binding domain). The overexpression of a dominant-negative SPAR mutant lacking Rap-GAP activity causes an elongation and thinning of spines, some of which resemble filopodial-like structures [74]. These results indicate that the SPAR modulation of Rap GTPases plays a regulatory role on the actin cytoskeleton and changes spine shape.

Not only Rap2 but also Ras is a component of the postsynaptic NMDA receptor complex. Indeed filopodia can be induced in cultured neurons by means of multiple depolarizing stimuli and this effect depends on activation of the Ras/MAPK pathway [82]. In conclusion Rac1, Rap2 and Ras together with their regulatory enzymes, are components of the postsynaptic NMDA receptor complex [83] linking receptor-mediated signals to actin dynamics. What could still remain to be discovered are the possible specific effectors of spine actin.

It is known that some of the receptors and scaffold proteins localized to dendritic spines are involved in spine morphogenesis. One of these is a cell-surface heparin-sulfate proteoglycan, called syndecan-2. The overexpression of syndecan-2 in hippocampal neurons accelerates the maturation of dendritic spines [84]. Interestingly, EphB2 (Eph family receptor) phosphorylates syndecan-2 on two cytoplasmic tyrosines, and this phosphorylation is required for the localization of the two proteins to developing spines and their maturation [85]. The idea is that EphB2, activated by ephrinB binding, phosphorylates the cytoplasmic tail of syndecan-2 and, in the process, the two proteins physically associate in clusters that lead to spine formation [85].

Among the PSD proteins, PDZ domain-containing scaffold proteins are believed to represent a molecular interface between glutamate receptors in synaptic membrane and spine cytoskeleton [21, 23]. It is consequently logical to think that some of these proteins may also link glutamate receptor activation to spine actin dynamics.

The overexpression of PSD95, which binds directly to the NR2 subunits of the NMDA receptors, increases the number and size of spines in hippocampal cultured neurons [86]. The overexpression of Shank, which links the NMDA receptor and the metabotropic glutamate receptor (mGluR) complexes through multiple protein interactions [77, 87], promotes the maturation of mushroom-shaped spines in developing hippocampal neurons, and increases the size of spine heads in mature neurons without affecting spine number [87]. The enlargement of spines by Shank depends on and cooperates with Homer, a protein that also binds to mGluRs and inositol-1,4,5-trisphosphate receptors (InsP<sub>3</sub>R). Indeed, Shank and Homer seem to mediate the recruitment of InsP<sub>3</sub>R (and presumably SER) to dendritic spines. Dominant-negative Shank mutants reduce spine density, possibly by decreasing the stability of affected spines or by inhibiting spine formation. Interestingly, postsynaptic overexpression of PSD-95 or Shank/Homer significantly enhances presynaptic function in addition to spine enlargement [86, 87], this emphasizes the close functional relationship between the two sides of the synapse.

#### Dendritic Spine Motility and Morphological Modification

Spine motility was originally proposed by Blomberg et al. [88] and Crick [89] but was first demonstrated by Fischer et al. [90]. Using time-lapse imaging and fluorescent proteins, considerable spine motility over a time scale of seconds to minutes has been documented in dissociated cultures [90–92], brain slices [11] and in vivo [93]. Motility is developmentally regulated, and is more pronounced during the critical period [11, 93–95]. Spine motility is actin-dependent, involves the remodelling of the actin cytoskeleton in the spine [11, 90], and is inhibited by volatile anesthetics [96]. At least in cultured neurons, the activity-dependent suppression of spine movement, has been found to inversely correlate with developmental age and contact with active presynaptic terminals, and is stimulated by inhibiting the basal activity with tetrodotoxin [95]. In brain slices, others have failed to detect any change in spine motility after the blockade or stimulation of neuronal activity, or in correlation with presynaptic contact [11, 97]. This difference can be explained by the different experimental preparations: dispersed hippocampal cultures [90, 95, 96] versus hippocampal slices [11, 51, 97]. Interestingly, a recent study using GFP-transfected pyramidal neurons in the somato-

sensory cortex found that spine motility is sensitive to input deprivation, but only during the period of peak neocortical synaptogenesis, thus suggesting the presence of a link between experience-driven synaptogenesis and spine motility [93].

There is evidence that actin-based motility is controlled by synaptic activity [91, 92]. The activation of either AMPA or NMDA receptors greatly inhibits spine actin dynamics and actin-based protrusive activity from the spine head, and the spine becomes more rounded and regular [91]. In this case, the inhibition of spine motility by AMPA receptors was dependent on postsynaptic membrane depolarization and calcium influx through voltage-activated channels [91]. Using a clever and technically advanced system to stimulate individual neurons, Colicos et al. [92] were able to show coordinated presynaptic and postsynaptic actin motility after tetanus stimulation. On the postsynaptic side, actin expands laterally outward from the central core, and new filopodia-like protrusions emerge that are eventually contacted by new presynaptic puncta of actin that become functional in a period of hours [92]. These morphological modifications are reminiscent of what can be observed during developmental synaptogenesis [50]. Finally, another fast and rapid movement of spines has been described by Korkotian and Segal [98], who found that they produce a tiny and rapid 'twitch' under back-propagating action potential and coincide with a transient increase in the intraspine calcium concentration. This fast movement is apparently independent on the age of the neurons and is present on spines contacted by an active presynaptic terminal [98].

However, although all of these findings indicate that spines are constantly moving and change morphology under physiological conditions, the functional significance of spine motility occurring over seconds or minutes is still completely obscure.

As mentioned above glutamate receptor stimulation plays a role in controlling spine motility but also regulates spine numbers. NMDA application to hippocampal cultures causes an almost complete collapse of dendritic spines and removes actin from the spine head [99]. Calcium and calcineurin seem to be involved in regulating spine stability: the NMDA-induced loss of spine actin and spine collapse is reduced in the presence of calcineurin, a calcium/calmodulin-dependent phosphatase [99]. On the contrary a low level of AMPA receptor activation with an intensity similar to that of spontaneous neurotransmitter release is required to maintain spine numbers in hippocampus organotypic cultures [100]. As described above AMPA receptor stimulation can also stop rapid spine

movements by stimulating calcium influx [91], but it has also been reported that the release of intracellular calcium by caffeine stimulates spine elongation [101]. In order to reconcile these data, we can imagine a bimodal relationship between calcium concentration and spine growth that is similar to what has been invoked to explain the different calcium requirements of LTD and LTP. A moderate transient increase in spine calcium provided by SER release, or through voltage-gated calcium channels mediated by AMPA receptor stimulation is necessary to promote spine stability and growth, whereas the higher concentrations of spine calcium induced by a prolonged activation of NMDA receptors induce spine shrinkage or collapse [102–104].

But the main question concerning spine plasticity is whether LTP or LTD is the expression of morphological modification in spine structure [105] and a number of electron microscopy studies have been carried out to test this hypothesis. The first report on the effect of LTP on spine morphology was published in 1975 by Fifkova and Van Harrefeld [106] who used a similar experimental protocol to that used a few years before by Bliss and Lomo [107] when they discovered LTP in the dentate gyrus. In this and a subsequent study [106], the authors describe a significant increase in spine area and volume from 2 min to 23 h after LTP induction with a peak at 10–60 min. Together with the observation that the spine neck becomes wider and shorter, these data suggest that tetanization is able to increase synapse size by increasing spine volume and recruiting an actin cytoskeleton, and that these changes last for hours [108, 109]. In a controlled series of more electrophysiological studies, Desmond and Levy [110] showed that LTP increases the density of large spines that have spinules or U-shaped profiles, and decreases the number of simple and ellipsoid spine profiles. The PSD area was also modified, with the total PSD surface area per unit volume of concave spines increasing significantly, whereas that of the non-concave spines decreased and an increase in the mean PSD length was observed for at least 60 min [30, 31, 110, 111]. Together, the studies of Fifkova et al. [106, 108, 109] and Desmond and Levy [110, 111] suggest that LTP induces the morphological modification of existing spines and synapses without the formation of new entities.

Conflicting results can be found in the literature concerning changes in spine number after LTP or exposure in an enriched or altered sensory environment. Trommald et al. [112] Andersen et al. [113] used three-dimensional reconstructions of serial EM micrographs, and observed an up to 50% increase in spine numbers, as well as

changes in the diameter of the spine neck and an increase in the so-called bifurcated spines, after inducing LTP in the dentate gyrus. They also found a small but significant increase in spine density in the CA1 region of the hippocampus of rats exposed to an enhanced sensory environment [114, 115], but others [6, 40, 116] found no changes in absolute spine number. However, some of these studies used the CA1 region instead of the dentate gyrus, or stimulated brain slices instead of *in vivo* stimulation. All of these studies were also based on statistical comparisons of two samples, whereas a better way would be to observe single spine in living tissue and show morphological changes in real time. Three seminal studies in this direction were published in 1999. Two of these used two-photon microscopy and brain slices. Maletic-Savatic et al. [117] visualized individual neurons infected by sindbis-virus-eGFP constructs, and found the outgrowth of dendritic protrusions similar to filopodia (often more than 4  $\mu\text{m}$  longer) after a strong tetanus-inducing LTP. The formation of these new protrusions that may turn into spine-like structures could be blocked by APV, the NMDA and LTP blocker [117]. In a more sophisticated way, in an attempt to localize where the potential morphological changes may occur, Engert and Bonhoeffer [118] showed a close correlation between successful functional synaptic enhancement and the generation of new spines.

Using another elegant approach, the same question was addressed by means of an EM analysis of spines from neurons in which an LTP had been previously induced [38]. This technique was able to select where morphological changes can be expected to occur in stimulated synapses by accumulating an EM-visible calcium precipitate in a postsynaptic spine that had just been subjected to a strong stimulus. Scrutinizing these spines for morphological changes, the authors found that, in many cases, there were pairs or triplets of labelled spines making contact with the same presynaptic terminal, or what were called 'same-dendrite, multiple synapse boutons' (sdMSBs). Similar ultrastructure modifications were observed by the same authors after short anoxic-hypoglycemic episodes and NMDA receptor activation in brain slices [119]. This may be the result of a rapid morphogenetic sequence of events after LTP-inducing stimulation that leads to PSD segmentation and spine splitting [37, 38]. However, using classical EM reconstruction, Fiala et al. [120] have more recently suggested that the sdMSBs formed after LTP are not due to spine splitting but probably the growth of new spines or the maturation of filopodia-like structures.

In conclusion although there seems to be a correlation between LTP and spine morphological modifications, there is no direct proof that these morphological changes contribute to synaptic strength as well as to the generation of potential new contact sites.

## Conclusions and Open Questions

The molecular exploration of spines is just at its beginning, but it is not surprising that a number of structural proteins and signaling pathways are involved. Spine structure is complex and dynamically regulated by different factors over short and long time scales. Many fundamental issues must still be addressed in order to understand spine function in brain physiology: for instance, although several actin-binding proteins are known to be involved in spine actin dynamics, none of them seems to be dedicated to dendritic spine regulation. The identification of specific actin modulators localized to spines might increase the possibility of analyzing spine function by means of a genetic approach. Alternatively it will be important to identify intracellular or extracellular factors that determine spine formation or absence in various neuron types. Another important question is to determine the relationship between spine shape and function in detail, as this will perhaps allow us to adjust spine morphology in order to make a neuron work or survive better. A number of cognitive disorders are associated with subtle spine malformations, such as changes in spine length, distribution, number or morphology, but still a detailed morphological, physiological and biochemical analysis has to be made in order to correlate clinical phenotypes and spine abnormalities. All of these key questions and others will be addressed in the near future using new and advanced technologies, this making an exciting and rapidly developing field.



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