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Synaptic organizing complexes

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Abstract

A number of synaptogenic factors induce presynaptic or postsynaptic differentiation when presented to axons or dendrites. Many such factors participate in bidirectional trans-synaptic adhesion complexes. Axonal neurexins interacting in an isoform-specific code with multiple dendritic partners (neuroligins, LRRRTMs, or Cbln-GluR δ), and axonal protein tyrosine phosphatase receptors interacting with dendritic NGL-3, nucleate local networks of high affinity protein-protein interactions leading to aligned presynaptic and postsynaptic differentiation. Additional secreted target-derived factors such as fibroblast growth factors and glial-derived factors such as thrombospondin bind specific axonal or dendritic receptors stimulating signal transduction mechanisms to promote selective aspects of synapse development. Together with classical adhesion molecules and controlled by transcriptional cascades, these synaptogenic adhesion complexes and secreted factors organize the molecular composition and thus functional properties of central synapses.

Introduction

Synapses are the basic units of communication in the brain. Synaptic transmission relies on the coordinated development of highly specialized structures spanning both participating cell surface membranes and cytoplasm. Synaptic specializations on both sides of the cleft involve membranous organelles, cytoskeleton, and vast protein networks. Minimally, synaptic function requires that postsynaptic neurotransmitter receptors with associated scaffolding and signaling molecules be precisely aligned on the dendrite opposite chemically matched presynaptic vesicles with regulated release and recycling machinery in the axon.

We discuss in this review 'synaptogenic' proteins for vertebrate central neuron synapses, defined here as proteins that induce presynaptic or postsynaptic differentiation when presented to axons or dendrites, respectively. Clearly there are also other molecules that contribute in essential ways to synaptogenesis. For example, cadherin and immunoglobulin superfamily proteins are key mediators of synaptic adhesion [1], and transcription factors such as MEF2 and Npas4 control synaptogenesis by regulating expression of many genes including some discussed here [2]. We focus here on recent advances related to synaptogenic cell surface and cleft proteins that induce synaptic differentiation, also commonly known as synaptic organizing proteins.

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Classes of synaptic organizing proteins

Synaptic organizing proteins exist in two main classes: (i) synaptic adhesion complexes, and (ii) secreted factors. An inventory of synaptogenic proteins is presented in Figure 1.

The synaptogenic adhesion complexes are composed of transmembrane presynaptic and postsynaptic partners that bind *in trans* across the cleft, a classic example being presynaptic neurexin and postsynaptic neuroligin [3–5]. Such cleft-spanning synaptic organizing complexes often have bidirectional activity, inducing presynaptic and postsynaptic differentiation, and by their nature mediate cell adhesion and alignment of the pre- and postsynaptic specializations. At least initially, synaptogenic activity mediated by synaptic adhesion complexes does not involve enzymatic activity but rather recruitment via high affinity protein-protein interactions (Figure 2A). Three particularly interesting findings and principles have emerged recently.

First, neurexins bind multiple, structurally diverse partners across the cleft (Table 1). The 4–5 mammalian neuroligins were the first characterized neurexin binding partners [3–5]. Neuroligin-1 with an insert at its B splice site is the major glutamatergic neuroligin and binds only β -neurexins. Neuroligin-2 functions specifically at GABAergic synapses and appears to bind all neurexins. Leucine-rich repeat transmembrane neuronal proteins LRRTMs, initially discovered to be synaptogenic molecules in an unbiased expression screen [6••], were recently identified in three independent studies to be trans-synaptic partners for neurexins [7••,8••,-9••]. LRRTM1 and LRRTM2 are glutamatergic postsynaptic proteins and bind α and β neurexins specifically lacking an insert at splice site 4 (-S4) [6••, 9••]. LRRTM2 and neuroligin-1 compete with similar affinity for an overlapping face of β -neurexin(-S4) [9••]. Given their broadly overlapping expression patterns [10,11], neuroligin-1, LRRTM1, and LRRTM2 are likely to coexist at many glutamatergic postsynaptic sites..

In a recent study, neurexins were identified as the trans-synaptic binding partners for another structurally distinct ligand pair: the complex of Cbln1-GluR δ 2 [12••]. Cbln1 was shown to bind β -neurexins containing but not lacking the insert at splice site 4 (+S4). The neurexin(+S4)-Cbln1-GluR δ 2 trans-synaptic triad, found to be essential for normal bidirectional parallel fiber-Purkinje cell synaptogenesis, illustrates the second principle: use of a secreted bridging protein (Cbln) to link transmembrane presynaptic and postsynaptic components [12••,13••]. Cbln1, a member of the C1q tumor necrosis factor superfamily secreted by granule cells, and GluR δ 2, a member of the ionotropic glutamate receptor family, were suspected to function in the same pathway since Cbln1 $^{-/-}$ and GluR δ 2 $^{-/-}$ mice had remarkably similar phenotypes [14,15]. Common defects include ataxia, deficient long-term depression, reduced numbers of parallel fiber-Purkinje synapses, increased numbers of free spines, and mismatches in the length of postsynaptic density relative to active zone at many remaining synapses. While Cbln1 and GluR δ 2 function selectively in parallel fiber synaptogenesis between cerebellar granule cells and Purkinje cells [12••,13••], related Cbln and GluR δ proteins are more widely expressed potentially resulting in considerable overlap with neuroligins and LRRTMs.

A third key finding was the discovery of an independent bidirectional synaptic organizing complex with high potency like neurexins and its partners: postsynaptic netrin G ligand NGL-3 and presynaptic LAR family protein tyrosine phosphatase receptors PTPRs [16••, 17]. NGLs were studied in the context of synaptogenesis based on their isolation as PSD-95 interacting proteins [18]. Like LRRTMs [6], the leucine-rich repeats of NGL-3 mediate presynaptic induction [16••]. However, a different family of presynaptic players are involved; NGL-3 binds the first two fibronectin domains of axonal LAR, PTP σ and PTP δ

[16••,17]. NGL-3 and the PTPR partners are broadly expressed in brain [17], suggesting substantial co-existence at synapses with neurexins and partners.

An unbiased cell-based screen to isolate the set of proteins able to trigger presynaptic differentiation identified LRRTMs, re-isolated neuroligins and NGLs and revealed the existence of cDNA pools containing novel potent synaptogenic factors [6••]. Thus, we have yet to identify the full complement of synapse organizing proteins. While this cell-based expression screen appears biased towards identifying transmembrane synaptic organizers, biochemical screens fractionating brain extracts [19] or glial conditioned media [20] have yielded several secreted synaptic organizing proteins.

Secreted synaptogenic factors can be derived from the target dendrite, the presynaptic cell, neighboring neurons, or neighboring astrocytes. They can selectively promote aspects of presynaptic differentiation or of postsynaptic differentiation, in part depending on the location of their receptor. Mechanisms of synaptic differentiation promoted by secreted factors can require activation of kinases, signaling pathways [21,22] and perhaps transcription (Figure 2B). It is not yet clear whether secreted synaptogenic factors are locally instructive, like the synaptic adhesion complexes, or perhaps more permissive in promoting synapses to form but not specifying their exact sites. Recent findings in this area include the differential roles of FGF22 and FGF7 in glutamatergic and GABAergic presynaptic differentiation [23•], and identification of dendritic surface receptors for the glial-derived synaptogenic factor thrombospondin [24•,25].

Presynaptic differentiation

Local instruction by synaptic adhesion complexes

The simplest and most likely mechanism whereby neuroligins, LRRTMs, and Cbln-GluR δ 2 induce presynaptic differentiation is by locally aggregating neurexins on the axon surface. Indeed, competition with neurexin1 β ectodomain reduced synaptogenic activity for each of these partners [7••,12••,26], mutation of LRRTM2 abolishing neurexin interaction abolished its synaptogenic activity [9], and RNAi-mediated knock-down of all neurexins reduced synaptogenic activity of GluR δ 2-Cbln1 [12••]. Moreover, artificially aggregating recombinant neurexin1 β on the axon surface is sufficient to cluster synaptic vesicles by a mechanism requiring the neurexin cytoplasmic domain [27]. A recent study suggested that binding to β -neurexins is not sufficient for neuroligin-1 to induce presynaptic differentiation and that binding to α -neurexins is necessary [28]. However, such a result contrasts with the findings from multiple groups that showed neuroligin-1(+B) binds only to β , not α , neurexins and robustly induces presynaptic differentiation [9••,26,29]. Most data suggest that aggregation of either α or β neurexins on the axon can mediate presynaptic differentiation. Considering that α -neurexins are required for presynaptic calcium channel function [30], induction of functional presynaptic differentiation by aggregation of β -neurexins may involve recruitment of α -neurexins.

Neurexins participate in both glutamatergic and GABAergic synaptogenesis. It will be important to determine exactly which neurexin isoforms, including splice variants, are expressed by specific GABAergic and glutamatergic neurons. We can make some educated guesses about the roles of neurexin variants in GABAergic versus glutamatergic synaptogenesis based on the synaptogenic activities and binding specificities of neurexin partners. Neuroligin-1(+B), LRRTM1, and LRRTM2 selectively induce glutamatergic presynaptic differentiation and collectively bind all neurexins except the α -neurexin(+S4) variants [6••,9••,29,31]. Neuroligin-2 selectively induces GABAergic presynaptic differentiation [31] and also binds α -neurexin(+S4) variants [29] thus suggesting that α -neurexin(+S4) variants may be selectively expressed by GABAergic neurons. A specific

role of α -neurexins(+S4) in GABAergic synaptogenesis is also supported by coculture studies [32] and by the finding that *in vivo* knockout of all α -neurexins, leaving β -neurexin expression intact, did not alter the number of morphologically defined glutamatergic synapses in cortex but reduced GABA synapse numbers by nearly half [30]. Interaction of α but not β neurexins with dystroglycan [33], which is present along with neuroligin-2 at a subset of mature GABAergic postsynaptic sites [34,35], may also contribute to long term stabilization of GABAergic synapses. Different glutamatergic synapses also express selective neurexin variants. Cerebellar neurons express mainly neurexin(+S4) variants, matching the cerebellar-specific expression of the (+S4)-selective Cbln1-GluR δ 2 partners, whereas (+S4) and (-S4) neurexins are highly expressed in hippocampus and cortex [12••]. The neurexin isoform partner code (Table 1), (+S4) with Cbln-GluR δ , (-S4) with LRRTMs, β with neuroligin-1(+B), and all neurexins with other neuroligins is intriguing, although the full physiological significance is not yet understood.

Aggregation of axonal neurexins likely induces presynaptic differentiation by nucleating a network of high affinity protein-protein interactions to recruit synaptic vesicles and active zone machinery. It is controversial whether neuroligin-1 dimerization is required for presynaptic inducing activity [27,28], but one can imagine how dimerization would facilitate nucleation by aggregated neurexins. The neurexin class II PDZ domain interaction site is presumed to be essential for mediating synaptogenic activity via binding to scaffolding proteins such as CASK [36], Mints [37], and/or syntenins [38]. Other PDZ-independent interactions of the neurexin intracellular domain may be important, including binding to protein 4.1N to promote local assembly of actin filaments [39] and binding to synaptotagmin to promote vesicle recruitment [40]. It is remarkable that aggregating neurexins by presenting neuroligins or LRRTMs appears sufficient to induce extensive differentiation of presynaptic sites capable of spontaneous and evoked transmitter release [6••,41,42]. Although the initial trigger is purely protein-protein interactions, it seems likely that catalytic processes must be invoked at some point to generate such complete presynaptic specializations. Whether phosphorylation of neurexins by CASK [43] or other kinases contributes to synaptic differentiation has not yet been explored. Artificial presynaptic differentiation induced by binding of poly-D-lysine coated beads to isolated axons, which presumably hijacks some of the normal cellular machinery, occurred as rapidly as one hour and required F-actin reorganization [44].

The most potent known synaptogenic complex independent of neurexins, postsynaptic NGL-3 acting on presynaptic LAR family PTPRs [16••], induces presynaptic specializations by a slightly different mechanism. These PTPRs do not terminate in PDZ domain binding sites. Intracellularly, they are composed of an active phosphatase domain and a phosphatase-like domain that binds α -liprins [45]. Given the role of liprins in presynaptic differentiation in *Drosophila* and *C. elegans*, and their ability to directly bind CASK, RIMs, and ERC/ELKS/CAST [46,47], recruitment of liprins is likely essential for presynaptic differentiation mediated by LAR family PTPRs. Whether the phosphatase activity plays a role has not yet been determined. NGL-1 and NGL-2 do not bind LAR. Their less potent presynaptic inducing activities may be mediated in part by binding to the selective ligands netrin-G1 and netrin-G2, but must involve additional factors since direct aggregation of the GPI-anchored netrin-Gs on the axon surface did not induce presynaptic differentiation [18].

Also less potent than NGL-3 or the neurexin partners, SynCAMs/Necl5 [48], EphB2 [49], and ephrinB3 [50,51] are reported to induce presynaptic differentiation in co-culture. Like neurexin, the corresponding presynaptic partners, SynCAMs, ephrinB1/2, and EphB2, respectively, all terminate in class II PDZ domain binding sites and bind syntenins and/or CASK. However, initial analyses of knockout mice suggest that SynCAMs may be as or more important for axon guidance and myelination [52,53]. A surprising new player

reported to induce presynaptic differentiation is amyloid precursor protein (APP), by a mechanism requiring APP or APLP2 in the contacting axons [54•]. Although APP does not terminate in a PDZ domain binding site, its direct interaction with Mint1 was suggested to mediate presynaptic differentiation. SALM3/Lrln4 and SALM5/Lrln5 also induce presynaptic differentiation by mechanisms not yet identified [55], perhaps involving transcellular association of SALM5 homophilically or with SALM4/Lrln3 [56]. The immunoglobulin domain protein SIRP α induces presynaptic differentiation by a mechanism at least partially involving axonal CD47 [22]. However, SIRP α may be more like the secreted factors in a couple of ways, in that it is active even in cleaved ectodomain form, and its mode of action involves G protein signaling [22].

In the co-culture or bead-induced hemi-synapse formation assay, where a single protein such as a neuroligin or LRRTM or NGL-3, or just the corresponding ectodomain, induces presynaptic differentiation in contacting axons, the triggering protein is presented at high local concentration. How are such high local concentrations achieved at *bona fide* synapses, or are they? Two additional mechanisms are likely important. First, other proteins such as cadherin and immunoglobulin superfamily proteins contribute to adhesion of the axon-dendrite contact site. Such adhesion mechanisms may be necessary to allow the axonal and dendritic synaptogenic partners to come in proximity. In support of such cooperative function, the ability of neuroligin-1 overexpression to increase mEPSC frequency in neurons differentiated from ES cells was lost by targeted deletion of N-cadherin [57]. Second, evidence is accumulating that none of these synaptogenic proteins acts alone at synapses. At a single glutamatergic synapse, we envision dendritic neuroligins, LRRTMs, and perhaps GluR δ -Cbln binding to overlapping sets of neurexin variants, dendritic NGL-3 binding to axonal LAR family PTPRs, and additional cell adhesion and soluble synaptogenic factors acting in concert to achieve presynaptic differentiation (Figure 2). The recently discovered overlapping interactions and the multiple sets of synaptogenic factors may be necessary to drive synaptogenesis by a dynamic network of local high affinity protein-protein interactions. For example, the interactions of neurexins with CASK, LAR with liprin, and CASK with liprin may reinforce each other. Multiple individual synaptogenic adhesion proteins may serve as nucleating factors, and thus loss of any single synaptogenic protein may result in only partial defects in synaptogenesis. Indeed, relatively subtle defects have been found in individual neuroligin-1, neuroligin-2 and LRRTM1 knockout mice [6•, 10,58,59••,60,61].

Promotion by secreted factors

Growth factors and neurotrophic factors comprise one class of secreted synaptogenic proteins. BDNF and NT-3, not discussed in detail here, have long been appreciated to promote aspects of synaptogenesis through activating their tyrosine kinase receptors TrkB and TrkC, respectively [62]. GDNF in a complex with GFR α 1 was reported to locally induce presynaptic differentiation by a mechanism mediated in part by GFR α 1 in the axons [63]. Since GFR α 1 is GPI-anchored, presumably a coreceptor must be involved to achieve presynaptic differentiation. GDNF could also mediate adhesion between cells expressing GFR α and was proposed to be a bridging molecule [63], like Cbln1. FGFs have been a focus of recent research. FGF22 was isolated from an unbiased biochemical screen for proteins that induce vesicle clustering in cultured motoneurons, and multiple FGFs were found to be active [19]. FGF2 was also previously found to increase clustering of presynaptic and postsynaptic markers and their apposition [21]. Analyses of knockout mice revealed selective roles for target-derived FGF22 in glutamatergic synaptogenesis and FGF7 in GABAergic synaptogenesis onto hippocampal CA3 neurons [23•].

The research on FGFs illustrates apparent differences in the mode of action of secreted synaptogenic proteins compared with synaptogenic adhesion complexes. Whereas the

adhesion complexes trigger relatively complete synaptic differentiation, FGF22 and FGF7 selectively mediated synaptic vesicle clustering and not active zone formation or postsynaptic differentiation [23•]. Thus, secreted factors may be utilized to promote selective aspects of synaptogenesis. FGFRs are tyrosine kinases, and the effects of FGF2 and FGF22 on synaptic vesicle accumulation were blocked by kinase inhibitors [21,22]. Thus, whereas the synaptogenic adhesion complexes act primarily through nucleating networks of high affinity protein-protein interactions, secreted synaptogenic factors may act primarily by stimulating catalytic signaling pathways.

An unresolved issue is how these secreted factors act in soluble form to promote development of highly localized structures. Several mechanisms can be envisioned. One, the secreted factors may bridge presynaptic and postsynaptic components, like Cbln1. Two, they may bind to presynaptically localized receptors, implying that other mechanisms have already generated a partial presynaptic site with the localized receptor. Three, they may be secreted in a highly localized manner, with the secretion site controlling the zone of action. Alternatively, secreted factors may promote synaptogenesis in a more permissive way, for example by increasing the aggregation of synaptic vesicles along the axon but not instructing the precise site of synapse formation. FGF22 and FGF7 increase axon branching as well as vesicle clustering in cultured neurons [19], suggesting non-localized actions, and yet they can localize specifically to excitatory or inhibitory synapses [23•], whether by localized secretion or binding to specific receptors is not yet known.

Wnts are another class of secreted factors that promote synaptogenesis. Wnt7a is involved transiently in cerebellar glomerular development *in vivo* [64] by a mechanism involving Dishevelled [65]. In hippocampal cultures, Wnt7a increases synaptic vesicle clustering and mEPSC frequency without altering postsynaptic properties [66] by a mechanism involving Frizzled5 [67]. Additional Wnt family members contribute specific synaptogenic activities through multiple signaling cascades [68,69]. Importantly, as shown in *C. elegans*, Wnts can also act as anti-synaptogenic factors, creating regions of a neuron refractory for synaptogenesis [70]. Such anti-synaptogenic factors are likely to play important roles in directing synaptogenesis but have not yet been well studied in mammals.

Postsynaptic differentiation

Local instruction by synaptic adhesion complexes

Just as aggregation of neuroligins on the axon surface mediates presynaptic differentiation, aggregation of the neuroligin postsynaptic partners, or NGL-3, on the dendrite surface mediates postsynaptic differentiation. Direct aggregation of recombinant neuroligins [71], LRRTM2 [6•], GluR52 [13•], or NGL-3 [16•] with antibody-coated beads was sufficient to recruit multiple postsynaptic proteins including PSD-95/93, Shank, GKAP, Homer, SynGAP, and NMDA receptor NR1. As for the presynaptic side, the mechanism of postsynaptic differentiation induced by these synaptogenic adhesion complexes occurs initially, and perhaps completely, by nucleating a dynamic network of high affinity protein-protein interactions. Neuroligins, LRRTMs, GluR δ , and NGLs all terminate in PDZ domain binding sites which can bind PSD-95, PSD-93, SAP102, and/or SAP97 and other glutamatergic scaffolding proteins [6•,18,72–74]. All bear consensus class I PDZ domain binding sites except for LRRTMs for which the atypical - ECEV mediates the interaction. It seems likely that non-PDZ domain interaction sites in the intracellular domains of these postsynaptic adhesion proteins are also important in postsynaptic differentiation; for example, GluR δ 2 directly binds and recruits Shank2 [75], and neuroligin-1 and LRRTM2 can each target to glutamatergic postsynaptic sites via intracellular sequences distinct from the PDZ domain binding sites [6•,76].

Whether AMPA receptors are recruited by these synaptogenic adhesion complexes is not entirely clear, and may vary among complexes. AMPA receptor recruitment has not been demonstrated by simple coculture with LAR family PTPRs or neurexins, but was reported for neurexin coculture with neurons overexpressing PSD-95 upon treatment with glutamate [77]. Direct aggregation of NGL-3 but not NGL-2 on dendrites recruited surface GluR2 [16••,78]. Whether such direct aggregation of neuroligin-1 or LRRTM2 recruits AMPA receptors has not been tested, although recombinant LRRTM2 was reported to co-immunoprecipitate individually with GluR1, GluR2, and NR1 [8••]. SALM 1–3 also terminate in class I PDZ domain binding sites, SALM 2 and 3 but not 5 can recruit postsynaptic proteins [55,78], and SALM 1 coimmunoprecipitates with NR1 [79]. EphB2 interacts with AMPA receptors indirectly through a class II PDZ domain intermediate such as GRIP or PICK1 [49], and the extracellular domain of EphB2 directly binds the extracellular domain of NR1 [80].

Postsynaptic differentiation that is artificially induced by any of the above synaptogenic factors is incomplete in another major way. These synaptogenic adhesion complexes are not sufficient to induce dendritic spine morphogenesis. Yet many of these proteins contribute to spine morphogenesis, since overexpression, knock-down, or knockout alters spine numbers or spine morphology [7••,18,81]. Signaling through EphB-ephrinB complexes in particular may be more important for spine morphogenesis [82•,83] than for the other aspects of synaptic differentiation that are the focus of this review. Perhaps the coculture or bead induction assays preclude spine morphogenesis, or perhaps no individual molecule presented to dendrites will be sufficient to trigger the full signaling cascade necessary for spine morphogenesis; it may prove fruitful to test co-presentation of multiple factors.

Another incompletely resolved issue is the role of neuroligins in selective GABAergic versus glutamatergic postsynaptic differentiation. In fact, the roles of neuroligins may be even broader, given the evidence for a role of neuroligin-1 in ciliary ganglion cholinergic synaptogenesis [84]. In the central nervous system, neuroligin-2 localizes specifically to GABAergic synapses [71,85], neuroligin-1 localizes selectively to glutamatergic synapses [86], and individual knockout mice exhibit selective deficits in GABAergic/glycinergic or glutamatergic transmission, respectively [58,59••]. Yet the phenotype of neuroligin-1,2,3 triple knockout mice is more severe than any of the single knockouts [10], indicating some redundancy among neuroligins. All neuroligins terminate in class I PDZ domain binding sites and can bind PSD-95 [72], and a recent study showed that all neuroligins can bind the GABAergic scaffolding protein gephyrin via a conserved cytoplasmic motif [59••]. Importantly, only neuroligin-2 also bound collybistin/ArhGEF9 and could recruit gephyrin bound to collybistin [59••]. The importance of this interaction is supported by defects in GABAergic synaptogenesis in collybistin knockout mice [87]. All neurexins were recently reported to interact directly with GABA_A receptors, although neurexin knock-down did not alter GABAergic transmission [88]; further studies will be required to determine whether this low affinity interaction has physiological significance. The potential interaction of all neuroligins with common glutamatergic and GABAergic scaffolding proteins may be an important design feature. Overexpressing PSD-95 not only enhances excitatory transmission but also reduces inhibitory transmission via translocating neuroligin-2 from GABAergic to glutamatergic synapses; such mechanisms may be utilized for controlling the balance of excitatory and inhibitory input onto neurons [89]. It is becoming clear that chemical matching of presynaptic and postsynaptic components cannot be achieved by neurexins and known partners; there must be additional GABA-specific and glutamate-specific adhesion complexes.

Promotion by secreted factors

One of the earliest identified synaptogenic factors for glutamatergic synapses was the secreted neuronal pentraxin NP2/NARP [90]. NP2 and the related NP1 interact via their pentraxin domain with the extracellular domain of all AMPA receptor subunits, and NP1/2 expressing cells recruit AMPA receptors to contact sites [90–92]. This mechanism of direct aggregation is rather like that of the synaptogenic adhesion complexes. Recent analyses of NP1/2 knockout mice have revealed a developmentally restricted role for NP1/2 in AMPA receptor clustering *in vivo*. During early postnatal stages, NP1/2 knockouts exhibited reduced clustering of GluR4 in hippocampus [92] and reduced AMPA-mediated but normal NMDA-mediated retinogeniculate transmission [93]. AMPA-mediated transmission in the knockouts subsequently surpassed that in wild type, possibly due to later-developing compensatory mechanisms. Perhaps in part due to the altered transmission, deficits were also found in segregation of eye-specific retinogeniculate projections [93,94].

Thrombospondins (TSPs), glial derived synaptogenic factors, act in a complementary manner to NP1/2, promoting formation of ultrastructurally normal synapses that are functional presynaptically but lack AMPA receptors [20]. We consider TSPs in the postsynaptic section because they act via binding receptors on dendrites. It was recently found that the 3 EGF repeats of TSPs mediate the synaptogenic activity by binding the $\alpha 2\delta$ -1 auxiliary calcium channel subunit, a target of GABA-pentin [24•]. The mechanism of TSP action appears to be independent of calcium channel function, but the events downstream of binding to the GPI-anchored $\alpha 2\delta$ -1 and potential coreceptors for intracellular signal transduction have not yet been delineated. Interestingly, $\alpha 2\delta$ -3 was found to be necessary presynaptically in *Drosophila* for bouton morphogenesis, also by a mechanism independent of calcium channel activity [95]. Another report suggested that TSPs accelerate synaptogenesis by binding to and aggregating neuroligin-1 on dendrites [25], although whether neuroligin-1 with bound TSP could simultaneously bind neurexins to induce apposing presynaptic differentiation is not clear. Based on the several day period required to see the effects of TSPs, along with the broad distribution and developmental window of action, it was suggested that TSPs may induce neurons to alter the expression of genes involved in synaptogenesis, such as the locally acting factors discussed here, rather than necessarily acting locally themselves [20].

Beyond basic synaptic differentiation

Synaptic development is a protracted process involving axon and dendrite contact, presynaptic and postsynaptic differentiation, morphogenesis, maturation, maintenance, and plasticity. In addition to their roles in differentiation, the synaptogenic proteins discussed here may function in other aspects of synaptogenesis, including synapse specificity. Thorough analyses have not been performed for most of the synaptogenic adhesion proteins, except the cerebellar-specific Cbln1-GluR δ 2 as discussed above, but the initial analyses suggest that there may be considerable synaptic selectivity. A role of neurexins in synaptic partner choice was first suggested upon the discovery of the possibility of thousands of neurexin splice variants. Although neuroligin-2 is localized to nearly all GABAergic synapses [85], its function appears to be more selective. Neuroligin-2 knockout mice show selective deficits at perisomatic synapses in hippocampus [59••] and at pyramidal cell synapses made by fast-spiking interneurons but not by somatostatin-positive interneurons in cortex [60]. LRRRTMs, NGLs, and LAR family PTPRs show interesting laminar-specific distributions [6••,11,17]. NGL-1 and NGL-2 concentration in specific dendritic layers was dependent on their axonal netrin-G partners [96]. Conditional knockout of GluR δ 2 in adult cerebellum revealed an essential function in synapse maintenance [97], but the role of most other proteins in synapse maintenance is not known. The contribution of each of the individual synaptogenic adhesion complex proteins to cellular partner choice and synapse

maintenance are major questions that will require careful *in vivo* analyses of localization and effects of targeted deletion to resolve.

Synaptic organizing proteins are well-suited to mediate plasticity because they integrate presynaptic and postsynaptic domains and influence the size, stability and morphology of synapses. A role of ephrins and EphRs in synaptic plasticity has been well documented [83], TSPs were implicated in experience dependent plasticity in mouse barrel cortex [24•], and Wnts in experience dependent plasticity at hippocampal mossy fiber synapses [98]. Clear evidence for neurexins and partners as mediators of synaptic plasticity has not yet emerged, but activity regulation of neurexin splicing [32] hints at such roles, and altering postsynaptic levels of neuroligin-1 can retrogradely modulate release probability [99]. Clever strategies may be required to test roles in plasticity separately from roles in synaptic differentiation for these potent synaptogenic adhesion complexes. Cerebellar long-term depression, which is abolished by loss of either Cbln1 or GluR δ 2, surprisingly was rescued by viral expression of GluR δ 2 lacking the N-terminal Cbln1-binding domain, even without rescue of the structural synaptic defects associated with loss of GluR δ 2 [100]. Perhaps high level expression in the absence of Cbln1 binding allowed sufficient postsynaptic accumulation of the GluR δ 2 C-terminal domain to rescue plasticity.

Since the initial linkage of mutations in neuroligins 3 and 4 to autism in 2003 [101], evidence has rapidly accumulated for contribution of neuroligin and neurexin variants to the neurodevelopmental disorders autism, schizophrenia, and mental retardation [5,102]. Although occurring in only a small fraction of patients, alterations include copy number variants, protein-truncating frameshifts, and function-altering missense variants, some *de novo*, thus constituting strong evidence of contribution to the disease. Mimicking these variants in mice can phenocopy some aspects of the disease; for example, neuroligin-4 knockout mice exhibit selective deficits in social behavior and ultrasonic communication [103]. Most of the other synaptogenic proteins have not been systematically studied in this context, although associations were also found for LRRTM1 and GluR δ 1 with schizophrenia [104,105] and SynCAM1 with autism [106]. Based on their function to recruit components to control synaptic properties in circuits important for cognitive processing, we suspect that many of these synaptogenic genes are at high risk for predisposing to neurodevelopmental disorders. Further understanding of the molecular pathways and circuit events down-stream of these synaptogenic molecules, and assessment of the validity of genetically-based mouse models of these disorders, may eventually contribute to effective targeted therapies.

Conclusions

An often-asked question is, "why are there so many synaptogenic proteins"? A recent variation of this question might be, "why are there multiple neurexin postsynaptic partners"? One can imagine several reasons. (1) Redundancy is one obvious reason, so that synaptogenesis will not be so susceptible to single-gene deleterious events. (2) The nature of the initial events in synaptic differentiation, nucleation by high affinity protein-protein interactions rather than catalytic events, may indicate another reason, a sort of mechanical stability. A network of such high affinity interactions where each protein connects with multiple other proteins may be necessary to drive local assembly. Such a network of multiply interacting proteins is thought to maintain both the presynaptic density and the glutamatergic postsynaptic density. Cross-interactions of neurexins with multiple postsynaptic partners as well as parallel systems may help stabilize interactions across the cleft by tying into the presynaptic and postsynaptic networks at multiple points. (3) Convergence onto signaling pathways and signal amplification may be another reason. Multiple signaling pathways and cytoskeletal networks must be invoked to achieve complete presynaptic and postsynaptic differentiation including morphogenesis in a physiologically

relevant time-frame. (4) These synaptogenic proteins may contribute to synaptic specificity, pairing specific presynaptic and postsynaptic cells, controlling exactly where and when synapses form. Specificity may apply not only to partner choice but may also direct synapses to subcellular domains. (5) Different synaptogenic proteins may confer differences in molecular composition, structure, and functional properties. Although the cell culture experiments with synaptogenic adhesion proteins revealed surprisingly little variability in such properties, loss of function *in vivo* may reveal more variability in the synaptic properties conferred by each molecule. (6) A primary function of some of these proteins may be in synaptic plasticity. The need for differential regulation by activity and signaling pathways would create a need for diversity in secreted and adhesion complex synaptic organizing proteins.

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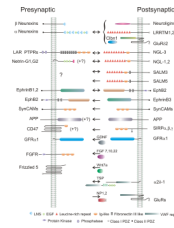


Figure 1.

An inventory of synaptogenic molecules, defined here as proteins that induce presynaptic (\leftarrow) or postsynaptic (\rightarrow) differentiation when presented to axons or dendrites, respectively. Many of the adhesion complexes have bidirectional synaptogenic activity (\leftrightarrow). The main receptors are also shown for the secreted synaptogenic factors. PDZ domain binding sites and common protein domains are indicated.

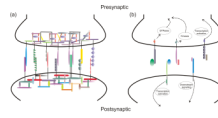


Figure 2.

Different initial mechanisms by which synaptogenic molecules promote synaptic differentiation. (a) Many synaptogenic adhesion complexes function primarily by nucleating a dynamic network of local high affinity protein-protein interactions in which each component interacts with multiple other components. (b) Many of the secreted factors directly activate signal transduction cascades involving kinases and GTPases, and perhaps regulate transcription. However, there are no strict boundaries, synaptogenic adhesion proteins can be kinases (e.g. Ephs) and secreted factors can act by local aggregation (e.g. NP1/2). These initial mechanisms are likely to converge on common downstream pathways mediating aspects of synaptic differentiation.

Table 1

Isoform-specific interactions between neurexins and postsynaptic partners

Neurexin isoform:	β (-S4)	β (+S4)	α (-S4)	α (+S4)
LRRTM 1,2	Yes	No	Yes	No
Neuroigin 1(+B)	Yes	Yes	No	No
Neuroigin 1(-B),2,3,4	Yes	Yes	Yes	Yes
Cbln1-GluR δ 2	No	Yes	?	?