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Cell Adhesion Molecules in Synapse Formation

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Neuronal transmission relies on signals transmitted through a vast array of excitatory and inhibitory neuronal synaptic connections. How do axons communicate with dendrites to build synapses, and what molecules regulate this interaction? There is a wealth of evidence suggesting that cell adhesion molecules (CAMs) provide much of the information required for synapse formation. This review highlights the molecular mechanisms used by CAMs to regulate presynaptic and postsynaptic differentiation.

Key words: adhesion; axon terminal; dendrite; neuron; synapse; synaptogenesis

In the nervous system, information is exchanged between neurons at sites of contact known as synapses. Before becoming a chemical synapse, a physical contact is typically formed between an axon and a dendrite. Contact and synapse formation is commonly thought to occur between an axonal filopodium (often from a growth cone) and a dendritic shaft (Ahmari et al., 2000; Jontes et al., 2000). However, dendritic filopodia have also been seen to play an active role in synapse formation (Ziv and Smith, 1996; Jontes et al., 2000). Initial contact establishment is followed by spatially and temporally controlled changes in morphology and molecular content to form a mature synapse characterized by the specific accumulation of synaptic vesicles and active zone components within the axon, in close apposition to a dendritic membrane studded with receptors (Fig. 1), which are held in place by a submembranous scaffold (Sheng and Kim, 2002).

Recent advances in subcellular fluorescence microscopy have revealed that the transformation of a contact site to a synapse involves the rapid recruitment and stabilization of both presynaptic and postsynaptic elements. These studies have shown that major components of the synaptic vesicle and active zone machinery travel in clusters together with other presynaptic proteins, such as calcium channels, and are rapidly recruited to new sites of contact (Ahmari et al., 2000; Zhai et al., 2001; Washbourne et al., 2002). On the postsynaptic side, receptor subunits and components of the scaffold or PSD (postsynaptic density) are recruited separately and with distinct time courses within minutes to hours after initial contact (Friedman et al., 2000; Bresler et al., 2001, 2004; Washbourne et al., 2002).

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These studies, however, have not shed light on key factors that control this process. Elucidation of the molecular mechanisms by which a transient contact between axonal growth cone and dendrite is transformed into a synapse is crucial for understanding the synaptic deficits that ultimately underlie disorders ranging from autism to schizophrenia (Zoghbi, 2003). In early studies using myoballs and motoneurons, it was apparent that adhesion played a strong role in setting up a communicative link between two cells (Chow and Poo, 1985). Although adhesion molecules had long been surmised to play a role in at least holding the contact together, it was not until very recently that membranebound CAMs were discovered to be major players in triggering synapse formation (Scheiffele et al., 2000; Biederer et al., 2002; Sytnyk et al., 2002; Fu et al., 2003) and thus acting as signal transducers. Furthermore, studies have shown that the action of adhesion molecules is not limited to initial contact formation but is also involved in specific target recognition (Yamagata et al., 2002; Shen and Bargmann, 2003; Shen et al., 2004) and regulation of synaptic size and strength (for review, see Scheiffele, 2003; Yamagata et al., 2003).

Because of the asymmetric nature of the synapse, the formation of the initial point of contact requires different responses in the presynaptic and postsynaptic structures (Fig. 1). Cell contact mediated by the heterophilic interaction between neuroligin and β -neurexin is the best understood mechanism for setting up the asymmetry at the synapse. However, the action of several other CAMs, including protocadherins and synaptic cell adhesion molecule 1 (SynCAM 1), are known to be or likely to be mediated by homophilic interactions (Obata et al., 1995; Biederer et al., 2002). It remains unclear whether these separate adhesive interactions may differentially modulate synaptic specificity and whether all act in concert to produce the huge variety of possible synaptic connections and strengths.

Here, we summarize recent findings on the roles of the following cell adhesion molecules in establishing synapses: neural cell

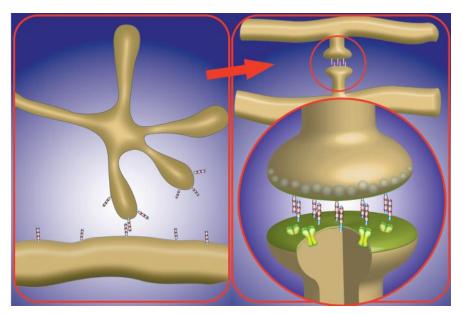


Figure 1. Diagram depicting the morphological transformation of an axodendritic contact to a mature synapse. Contact between an axonal growth cone filopodia and a dendrite via the homophilic adhesion molecule SynCAM 1 (blue—pink) develops to form a stereotypical excitatory synapse. SynCAM, an example of one of several synaptically localized adhesion molecules, remains at the synapse to hold the presynaptic and postsynaptic terminals together. The presynaptic terminal is filled with synaptic vesicles (white spheres), and neurotransmitter receptors (green) are recruited to the postsynaptic membrane.

adhesion molecule (NCAM), neuroligin/ β -neurexin, SynCAM 1, and γ -protocadherins. We also discuss the recent discovery that thrombospondins (TSPs), adhesive extracellular matrix (ECM) molecules, are novel glia-released factors involved in this process.

Synapse formation mediated by NCAM

NCAM was one of the first membrane proteins implicated in adhesion between neurons. NCAM belongs to the Ig superfamily of CAMs and is involved in homophilic and heterophilic interactions with recognition molecules. Many biological functions of NCAM are mediated by the negatively charged carbohydrate polysialic acid, which appears to be exclusively carried by this molecule and is synthesized in a developmentally, cell type-, and activity-dependent manner (Muller et al., 1996; Eckhardt et al., 2000; Kleene and Schachner, 2004). Previous studies suggested functions for NCAM in fasciculation of axons to form bundles, such as the retinotectal and mossy fiber projections (Rutishauser, 1985; Cremer et al., 1997). More recent data suggest that NCAM is involved in both early synaptogenesis and subsequent synaptic maturation. Studies on developing hippocampal neurons maintained for 2-3 d in culture have shown that clusters of NCAM at the cell surface are linked via spectrin to trans-Golgi-derived organelles (Sytnyk et al., 2002). These NCAMorganelle complexes translocate along neurites to sites of contacts within minutes after initial contact formation. Experiments in heterogenotypic cocultures of NCAM-deficient and wild-type neurons provide evidence that both presynaptic and postsynaptic NCAM is required to anchor trans-Golgi-derived organelles at nascent synapses (Sytnyk et al., 2002). At later stages (4–7 d in vitro), the relative levels of postsynaptic but not presynaptic NCAM expression regulates both the number of synapses and strength of excitatory synaptic connections in an NMDA receptor-dependent manner (Dityatev et al., 2000). This mechanism involves the interaction between the polysialylated form

of NCAM and heparan sulfate proteoglycans and requires fibroblast growth factor receptor (FGFR)-mediated signaling (Dityatev et al., 2004). Strikingly, a peptide corresponding to the binding site of NCAM to FGFR promotes synapse formation and memory (Cambon et al., 2004), indicating that CAM-derived compounds have potential therapeutic value. The importance of mammalian NCAM for synapse dynamics is further supported by the following: (1) activity-dependent expression of NCAM (Schuster et al., 1998); (2) impaired long-term potentiation and depression in mice deficient in NCAM or associated polysialic acid (Muller et al., 1996; Eckhardt et al., 2000; Bukalo et al., 2004); (3) alterations in transmission, vesicle dynamics, and transmitter release machinery at NCAM-deficient neuromuscular junctions (Polo-Parada et al., 2004); and (4) pivotal roles of NCAM homologs, fasciclin II and Aplysia cell adhesion molecule (apCAM), in formation of synapses in Drosophila and Aplysia (for review, see Packard et al., 2003).

Role for neuroligin and β -neurexin in excitatory synapse formation

The heterotypic cell adhesion "couple" β -neurexin and neuroligin was proposed as a potential trigger of synaptogenesis when it was discovered that neuroligin, a transmembrane protein identified as an interacting protein of β -neurexin [a splice variant of the latrotoxin receptor α -neurexin (Ichtchenko et al., 1995)], was able to induce the formation of presynaptic terminals onto nonneuronal cells (Scheiffele et al., 2000; Fu et al., 2003). Further characterization of the presynaptic side of this heterotypic system established that it is indeed the clustering of β -neurexin that brings about the recruitment of synaptic vesicles (Dean et al., 2003).

Previously, it had been established that neuroligin contained a PDZ (postsynaptic density 95/Discs large/zona occludens 1) binding domain that could bind PSD-95 (Irie et al., 1997) and could therefore establish a link between the presynaptic cell and the postsynaptic density. It was further determined that neuroligin-mediated adhesion promotes the functional recruitment of postsynaptic NMDA receptors probably via the interaction with PSD-95 when transiently expressed in non-neuronal cells (Fu et al., 2003). Increasing neuroligin-mediated adhesion by overexpression stimulates recruitment of postsynaptic scaffolding molecules such as PSD-95 and Homer and synaptic recruitment of NMDA receptors (B. Chih and P. Scheiffele, unpublished observations). This indicates that β -neurexin– neuroligin adhesion complexes promote synapse assembly in a bidirectional manner and provide a platform for postsynaptic neurotransmitter receptor recruitment. Suppression of neuroligin-1, -2, and -3 expression by RNA interference leads to a loss of synapses and dendritic spines in hippocampal neurons (Chih and Scheiffele, unpublished observations). Interestingly, also a loss of inhibitory terminals formed on cells with reduced neuroligin-1, -2, and -3 expression was observed. Therefore, the function of neuroligins in synaptogenesis might not be restricted to excitatory synapses, as anticipated previously. This is consistent with the observation that endogenous neurexins are concentrated at excitatory and inhibitory presynaptic terminals (Ullrich et al., 1995). These in vitro studies suggest that the β -neurexin– neuroligin adhesion system is essential for the appropriate assembly of synaptic circuits. Such an essential role is further supported by the recent realization that mutations in neuroligin-3 and -4 genes are associated with mental retardation and autism (Philibert et al., 2000; Jamain et al., 2003; Laumonnier et al., 2004), two neurodevelopmental disorders that are characterized by aberrant spine morphogenesis and defects in dendritic arbor growth. All presently identified neuroligin-3 and -4 mutations result in a loss of function, either attributable to truncation or a point mutation of the cholinesterase domain, which result in intracellular retention (Chih et al., 2004; Comoletti et al., 2004). Furthermore, it appears that point mutations and chromosomal rearrangements in regions of the genome containing the genes for neuroligin-2 and PSD-95 are associated with autism (Mariner et al., 1986; Risch et al., 1999).

New work from Prange et al. (2004) demonstrates that concerted actions of cell adhesion molecules and scaffolding proteins regulate the morphology, number, and type of synapses formed. For example, the presence of PSD-95 dictates what kinds of synapses are induced by neuroligin-1: excitatory or inhibitory. PSD-95 restricts neuroligin-1 to excitatory synapses, and manipulations that alter endogenous levels of these proteins result in an overall change in the ratio of excitatory to inhibitory synaptic contacts. Therefore, improper expression and/or targeting of molecules that control synaptic specificity may result in formation of aberrant synapses and a change in the balance of neuronal excitation—inhibition that underlie complex psychiatric disorders (Rubenstein and Merzenich, 2003).

SynCAM

SynCAM 1 and neuroligins are the only two known CAMs that are sufficient to drive formation of presynaptic terminals (Scheiffele et al., 2000; Biederer et al., 2002; Fu et al., 2003) (Y. Sara, T. Biederer, D. Atasoy, M. G. Mozhayeva, T. C. Sudhof, and E. T. Kavalali, unpublished observation). SynCAM 1 is a vertebratespecific Ig superfamily member with homophilic binding properties. It is localized to both sides of the synaptic cleft (Fig. 1) and shares similarities with the invertebrate molecules fasciclin II and apCAM (Biederer et al., 2002). Like neuroligin, when SynCAM 1 is expressed in non-neuronal cells and cocultured with neurons, the formation of presynaptic terminals is induced. In such a coculture system, the activities of SynCAM 1 and neuroligin-1 are highly comparable with respect to the properties of synaptic vesicle recycling in the induced presynaptic terminals. Furthermore, recent results show that either SynCAM 1 or neuroligin-1 can be used to reconstitute both evoked release and spontaneous miniature events (minis) (Biederer et al., 2002; Fu et al., 2003) (Sara, Biederer, Atasoy, Mozhayeva, Sudhof, and Kavalali, unpublished observation). This is probably attributable to the fact that both adhesion molecules may share the same presynaptic signaling pathways. The known protein interaction motifs in the cytosolic tails of SynCAM 1 and β -neurexins are highly conserved. They therefore presumably bind the same adaptor proteins at the presynaptic terminal. However, binding proteins, such as CASK (calcium/calmodulin-dependent serine kinase) have so far only been characterized in vitro (Biederer and Südhof, 2001; Biederer et al., 2002), and the downstream partners of SynCAM 1 and β -neurexin in vivo remain to be identified. Studies with dominant-negative constructs strongly indicate that PDZ domain containing adaptor proteins are involved in this process

(Biederer et al., 2002). Via this presynaptic signaling, both Syn-CAM 1 and neuroligins are capable of inducing presynaptic terminals with a complete physiological complement.

SynCAM 1 overexpression in hippocampal neurons specifically promotes excitatory synaptic transmission and increases the frequency of minis (Biederer et al., 2002). This activity of SynCAM 1 depends on its cytosolic sequence, highlighting the importance of finding the downstream signaling partners to understand the function of synaptic adhesion molecules (Sara, Biederer, Atasoy, Mozhayeva, Sudhof, and Kavalali, unpublished observation). SynCAM 1 is expressed throughout the CNS, and its developmentally controlled expression and glycosylation pattern (see below) strongly suggests that it exerts its synaptogenic activity predominantly during early development (Biederer et al., 2002). However, a role for SynCAM 1 may still be determined at adult synapses.

Protocadherins

In addition to the requirement for homotypic and heterotypic CAM interactions for the general formation of synaptic contacts, CAM interactions have the potential to provide molecular mechanisms for determining synaptic partners. This could be considered a corollary to Sperry's classic "chemoaffinity hypothesis" (Sperry, 1963) in which molecular cues present on presynaptic and postsynaptic neurons determine synaptic specificity in a "lock-and-key" manner. Because of the vast complexity of synaptic connections, such a mechanism would require great combinatorial diversity; the discovery of the α -, β -, and γ -protocadherin genes (Kohmura et al., 1998; Wu and Maniatis, 1999) has provided one potential source of such diversity. Over 50 alternative exons, each encoding the extracellular, transmembrane, and proximal cytoplasmic domains of a particular protocadherin isoform, are arrayed in tandem on a single chromosome in the mouse and human genomes (Wu and Maniatis, 1999; Wu et al., 2001). Each such "variable" exon is, in the α and γ clusters, spliced to three "constant" exons that encode a common C-terminal domain. Multiple α - and γ -protocadherin genes are highly expressed by many neurons, and the proteins are concentrated at synapses (Kohmura et al., 1998; Wang et al., 2002a,b; Phillips et al., 2003). Thus, the protocadherins have the potential to channel a great multiplicity of presynaptic and postsynaptic specificities into a common signal transduction pathway.

Evidence that protocadherins are important molecules in synaptic development was obtained recently from studies by J. A. Weiner, X. Wang (Northwestern University, Evanston, IL), and J. R. Sanes (Harvard University, Cambridge, MA) on the analysis of mice mutant for the 22 γ -protocadherins. Mice lacking the entire locus ($Pcdh-\gamma^{del/del}$) die at birth, exhibiting little voluntary movement or reflex action (Wang et al., 2002b). This is attributable to massive late embryonic apoptosis of spinal interneurons and a concomitant neurodegeneration and loss of spinal synaptic density. To differentiate between defects associated with neurodegeneration and synapse loss, apoptosis was blocked by crossing the $Pcdh-\gamma^{del/del}$ mice with mice lacking the gene encoding Bax, a pro-cell death member of the Bcl-2 family (Deckwerth et al., 1996). As expected, the neurodegenerative phenotype is completely rescued in the resulting double mutants; however, the mice still die at birth with reduced motor activity and exhibit a reduction in spinal synaptic density similar to that of the *Pcdh*- $\gamma^{del/del}$ single mutants. Similar synaptic defects (and neonatal lethality) in the absence of neurodegeneration are also observed in a hypomorphic mouse mutant in which the common *Pcdh*-γ C terminus is truncated. This result confirms the phenotype in the

 $Pcdh-\gamma/Bax$ double mutants and suggests that synapse loss in the double mutants is unlikely to reflect simply aberrant neuronal phenotypes attributable to artificial blocking of the cell death program (J. A. Weiner, X. Wang, J. C. Tapia, and J. R. Sanes, in preparation).

Together, these results confirm that γ -protocadherins are required for normal synaptic development. Initial electrophysiological analyses of mutant spinal neurons *in vitro* have revealed reductions in the amplitude of both EPSCs and IPSCs, corresponding to reductions in size and number of immunostained synaptic puncta. However, it remains unclear whether these defects reflect primarily a disruption of synapse formation, specificity, or maturation. Examination of the dynamics of synapse assembly in *Pcdh-\gamma* mutant neurons can be addressed in spinal neuron cultures, but analysis of synaptic specificity will likely require further genetic manipulations *in vivo* of particular neuronal subtypes with defined projection patterns.

Thrombospondins

Very little is known on how synaptic adhesion and synapse formation is modulated. Surprisingly, recent evidence suggests that synapse formation may be regulated by key secreted factors that are produced by astrocytes. Consistent with an important role for glia in synapse formation, the bulk of CNS synapse formation takes place only after astrocytes develop. The discovery that CNS synapse number is profoundly enhanced by signals secreted by astrocytes (Ullian et al., 2001) prompted the search for soluble factors involved in this process. Recently, TSPs were discovered as soluble factors that are released from astrocytes. TSPs are necessary and sufficient for synapse formation and are capable of inducing ultrastructurally normal synapses that are presynaptically active but postsynaptically silent (K. S. Christopherson, E. M. Ullian, and B. A. Barres, unpublished observation). In vivo, TSPs are concentrated at astrocyte processes and synapses throughout the developing brain. Furthermore, mice deficient in two of the TSP isomers expressed in the CNS (TSP1 and TSP2) show a significant decrease in synapse number in cortex and superior colliculus (Christopherson, Ullian, and Barres, unpublished observation).

How do TSPs induce synaptogenesis? TSPs are large oligomeric ECM molecules that act as multifunctional regulators of cell-cell and cell-ECM adhesive interactions (Adams and Tucker, 2000; Lawler, 2000; Adams, 2001; Bornstein, 2001). There are five TSP family members that form either homotrimers or homopentamers. TSP1 and TSP2, the closely related trimeric TSPs, share the same domain structure and are both able to promote synapse formation. Many of the known effects of TSPs in other systems are mediated by direct or indirect modulation of integrin function, although other TSP-interacting molecules have been identified, including CD36, LRP (lipoprotein receptor-related protein), calreticulin, heparan and chondroitin sulfate proteoglycans, and heparan-binding growth factors such as TGF-β1. Although little is known about how TSPs could be functioning in the CNS (Iruela-Arispe et al., 1993; Scott-Drew and ffrench-Constant, 1997; Adams and Tucker, 2000), it is plausible that they promote synapse formation by aligning or maintaining adhesion of presynaptic and postsynaptic specializations. This idea is supported by the observation that treatment of neurons with TSP-depleted astrocyte conditioned medium results in the induction of presynaptic and postsynaptic puncta that are no longer colocalized (Christopherson, Ullian, and Barres, unpublished observation). Alternatively or concurrently, TSPs could be mediating certain intracellular signaling pathways by binding to

one of their known cell surface receptors. To elucidate the cellular and molecular mechanism by which TSPs induce synaptogenesis, the B. A. Barres laboratory at Stanford University (Stanford, CA) is investigating the domain(s) of TSP responsible for promoting synaptogenesis. Together, these findings show that TSP1 and TSP2 promote CNS synaptogenesis and add to the growing evidence that astrocytes are active participants in CNS synaptogenesis *in vivo*.

Concluding remarks

In light of the remarkable heterogeneity of CNS synapses, it is not surprising that the factors that govern the diversity of synaptic contacts have been elusive. Recently, functional studies of individual adhesion molecules have begun to provide a wealth of information on their role in synapse assembly, spine morphogenesis, and synaptic plasticity. Although the disparate adhesion systems share the ability to mediate adhesive interactions, they individually control specific aspects of synapse formation. Because it appears that multiple systems cooperate at individual synapses, it is of great interest to determine whether separate CAM systems act in a parallel or in a hierarchical manner. Cadherins, neurexins, and several other recently identified synaptic adhesion proteins are coupled to common cytoplasmic scaffolds, and their functions might be integrated at that level, with molecules of one class triggering recruitment of adhesive factors from a different family. In contrast, the strong synaptogenic activity displayed by SynCAM and β -neurexin–neuroligin suggests that these may be positioned upstream of other CAMs. It could be envisioned that these "early" CAMs are required for the initial synaptogenic event and that "late" CAMs are then required to decide what type of synapse it becomes or whether the synapse is stabilized and matures. It is conceivable that certain CAMs may also regulate the size of specific neuronal populations. The requirement for γ-protocadherins in both synapse development and neuronal survival suggests that they may provide a molecular link between the processes of neuronal apoptosis and synaptogenesis during development.

Another outstanding question is how the activity of a particular CAM may be regulated either developmentally or in response to activity. For instance, palmitoylation and polysialylation drastically modify adhesive properties, binding partners, and signaling mediated by NCAM (Rutishauser and Landmesser, 1996; Niethammer et al., 2002). Interestingly, the SynCAM 1 glycoform primarily expressed during the peak of synaptogenesis is the one that displays the strongest homophilic binding (A. Fogel and Biederer, unpublished observation), and it is tempting to speculate that controlled N-glycosylation regulates both Syn-CAM 1 binding and synapse-inducing activity. Another possibility may be activity-dependent alternative splicing of CAM transcripts, which may be used to modulate function and/or expression levels, as has been described for NCAM and NMDA receptors (Doherty et al., 1992; Mu et al., 2003). Understanding the mechanisms that govern the synthesis and degradation of cell adhesion molecules and their subsynaptic distribution and signaling will be relevant for understanding how these events contribute to the regulation of synapse stability and number.

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