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Regulation of Glutamate Receptors by Their Auxiliary Subunits

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Abstract

Glutamate receptors are major excitatory receptors in the brain. Recent findings have established auxiliary subunits of glutamate receptors as critical modulators of synaptic transmission, synaptic plasticity and neurological disorder. The elucidation of the molecular rules governing glutamate receptors and subunits will improve our understanding of synapses and of neural-circuit regulation in the brain.

1. Introduction

The brain is the primary coordinator of animal behavior. Neural circuits, which are composed of billions of neurons, are the functional units of the central nervous system (CNS). Neurons communicate with each other at synapses. Neurotransmitters released from the presynaptic terminal of one neuron act on neurotransmitter receptors at the postsynaptic membranes of another neuron to induce changes in membrane potential or to activate signaling cascades. This newly generated information at postsynaptic sites travels through dendrites and axons to presynaptic terminals and to adjacent neurons via synaptic transmission. This network of connections organizes neural circuits of the CNS. Therefore, the elucidation of the rules of synaptic transmission and of the changes in neuronal membrane potentials will allow us to generate blueprints of functional neural circuits to enhance our understanding of the brain.

Glutamate receptors

There are two types of synapse in the brain: excitatory and inhibitory synapses. Excitatory synapses, where neurotransmitters induce depolarization of postsynaptic membranes, utilize glutamate as a major neurotransmitter in the vertebrate brain. In contrast, inhibitory synapses utilize GABA and glycine as major inhibitory neurotransmitters in the vertebrate brain. At excitatory synapses, glutamate released from presynaptic terminals binds to glutamate receptors, which are classified as ionotropic or metabotropic glutamate receptors. Ionotropic glutamate receptors are further classified pharmacologically as AMPA-, NMDA-, and kainate-sensitive glutamate receptors. Postsynaptic membranes contain all three ionotropic glutamate receptors and each receptor plays distinct roles in the brain. NMDA- and kainate-type receptors play roles in synaptic plasticity or slower transmission (10–100 ms), whereas AMPA receptors (AMPA receptors) play dominant roles in fast synaptic transmission (faster than 10 ms) to induce membrane depolarization after glutamate binding. Therefore, fast synaptic transmission is determined by channel activity and the number of AMPARs at synapses. In this review, we will discuss recent progress in the research of the role of ionotropic glutamate receptors and their auxiliary subunits in the control of synaptic transmission.

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2. AMPA-type glutamate receptors and the transmembrane AMPAR regulatory protein (TARP) auxiliary subunit

AMPA receptors play major roles in fast synaptic transmission. Four subunits of AMPARs (GluA1–4) assemble as a tetramer. AMPAR tetramers can function as glutamate-gated ion channels. However, native AMPAR complexes comprise transmembrane AMPAR regulatory proteins (TARPs) as AMPAR auxiliary subunits to modulate channel activity and the trafficking of AMPARs.

TARP genes and proteins

The prototypical TARP stargazin/ γ -2 was identified as the causative gene in the spontaneous mutant mouse *stargazer*, which shows ataxia and absence epilepsy, and is considered to be a calcium channel γ subunit because of its 23% sequence homology with the γ -1 auxiliary subunit of the 1,4-dihydropyridine (DHP)-sensitive calcium channel from skeletal muscle (FIGURE 1A) (40). Eight γ -1 homologous proteins (γ -1–8) were identified from a genomic database (6, 7, 14, 35). Among the γ -1 homologous proteins, six proteins modulate AMPAR activity and are termed transmembrane AMPAR regulatory proteins (TARPs) (FIGURE 1A) (32, 33, 79). TARPs are tetramembrane-spanning proteins (FIGURE 1B) and each of the TARP isoforms is expressed distinctly in the brain (25, 35, 79). TARPs are classified into two classes according to their distinct functions in AMPAR modulation (discussed later). Class I TARPs include stargazin/ γ -2, γ -3, γ -4, and γ -8, which all contain a typical PDZ domain-binding motif (–TTPV) at their C terminus. In contrast, class II TARPs include γ -5 and γ -7, which both contain an atypical PDZ-binding motif (–S/TTPC) at their C terminus (FIGURE 1B). TARPs are well conserved among species, including vertebrates and invertebrates. Mammalian and *C. elegans* TARPs (STG-1 and 2) share low homology; however, both are tetramembrane-spanning proteins and modulate AMPAR functions (91–93). In addition, TARPs share homology with claudin, which plays roles in the formation of tight junctions, presumably as an adhesion molecule (86) (FIGURE 1A). Therefore, TARPs may function as claudin-like cell adhesion molecules (64, 81). However, synapses lacking TARPs show normal synaptic morphology (8); thus, TARPs may require specific circumstances to function as adhesion molecules.

Interaction of TARPs with AMPARs

Immunoprecipitation of the TARP complex from the brain identified all AMPAR subunits (GluA1–4) as major interactors (23,80). Purification of the native AMPAR complex identified TARPs as major binding proteins (23,57). Furthermore, Blue-Native PAGE analysis of the TARP and AMPAR complexes revealed that all stargazin/ γ -2 interact with AMPARs and that most AMPARs interact with TARPs in the cerebellum (89). These results established TARPs as major components of the AMPAR complex in the brain.

Where do TARPs interact with AMPARs in neurons?

The total levels of AMPAR are decreased in the cerebellum of stargazin/ γ -2 disrupted mice and in the hippocampus of γ -8 knockout mice, where each TARP isoform is expressed as a major TARP (24,67,79). Furthermore, the ratio of EndoH-sensitive immature to EndoH-resistant mature AMPAR is increased in both mouse models (67,79). Interestingly, the expression of ER chaperones, BiP/Grp78, is increased in stargazin/ γ -2 disrupted mice, as part of the AMPAR unfolded protein response. Therefore, TARPs are likely to interact with AMPARs at the ER (88).

Interaction domains between TARPs and AMPARs

Interaction domains remain unclear, probably because of the difficulty in handling two transmembrane proteins. Single-particle analysis, which revealed the structure of the native AMPA receptor complex with and without TARPs at a 40 Å resolution, suggests that the transmembrane domains could act as interaction domains (57,58). The determination of the atomic structure of the complex is necessary to determine the precise mechanism of this interaction. As described later, TARPs modulate the pharmacology of AMPARs. The difference between the pharmacology of AMPARs alone (TARPlless AMPARs) and AMPARs with TARP (TARPin AMPARs) suggests that TARPs could assume variable stoichiometry (0/2/4) on AMPARs in neurons (71).

3. AMPAR trafficking and synaptic localization

Excitatory synapses in the vertebrate brain show two characteristic features: an electron-dense area beneath postsynaptic sites, the so-called postsynaptic density (PSD), and the use of glutamate as a major excitatory neurotransmitter. The molecular link between the PSD and glutamate receptors has been studied extensively. In a series of studies, a PDZ domain-containing protein, PSD-95, was identified as a major component of the PSD (12). In addition, overexpression of PSD-95 in neurons increases AMPAR activity at synapses, as shown by the increase in the levels of AMPAR and in excitatory postsynaptic currents (EPSCs) (4,20,21,59,68). However, PSD-95 cannot interact directly with AMPARs, which suggests that an additional molecule links PSD-95 to AMPARs. TARP, which binds to both AMPARs and PSD-95, was identified as such a molecule.

TARP-mediated AMPAR trafficking

Cerebellar granule cells express stargazin/ γ -2 as the sole TARP. The *stargazer* mouse does not exhibit AMPAR activity at cerebellar mossy fiber/granule cell synapses (8,28). Interestingly, overexpression of full-length stargazin/ γ -2 in primary cerebellar granule cell cultures from *stargazer* mice restores both synaptic and surface AMPAR activity, whereas overexpression of stargazin/ γ -2 lacking the C-terminal PDZ domain-binding motif (four amino acids, -TTPV) restored surface, but not synaptic, AMPAR activity (8). This result indicates that stargazin/ γ -2 modulates AMPAR activity via two distinct mechanisms, i.e., TARPs regulates the surface expression of AMPARs and the C-terminal PDZ-binding motif of TARPs (-TTPV) controls the synaptic localization of AMPARs (FIGURE 2B, 2A).

In contrast, the γ -8 knockout mouse exhibits a 90% reduction in surface AMPAR activity, but only a 30% reduction in AMPAR-mediated EPSCs in hippocampal pyramidal cells (67). The milder deficit in AMPAR-mediated synaptic transmission in γ -8 knockout mice compared with *stargazer* mice could be due to redundancy by other TARPs in the hippocampus, because all class I TARPs are expressed in hippocampal pyramidal cells (25,79). In support of this, γ -8 and stargazin/ γ -2 double-knockout mice exhibit a more severe reduction (50%) in synaptic transmission (67). Other possibilities to explain the difference in the extent of reduction in synaptic transmission between *stargazer* and γ -8 knockout mice could be the differences in the expression of AMPAR in distinct brain regions or in the expression of TARP subunits. For instance, TARP-dependent AMPAR trafficking is dominant in the cerebellum, but not in the hippocampus, and TARPlless AMPARs may be localized at synapses in the hippocampus. Because mice carrying disruption of three class I TARPs (stargazin/ γ -2, γ -3, and γ -8) exhibit lethality at postnatal day 0 (49), it is difficult to study adult mice carrying disruption of all six TARP isoforms. Conditional targeting disruption is required to examine this possibility. Alternatively, stargazin/ γ -2 and γ -8 are preferentially targeted to synapses and extrasynapses, respectively. In support of this assumption, biochemical fractionation showed that stargazin/ γ -2 is more

abundant in the synaptic fraction, whereas γ -8 is more abundant in the extrasynaptic fraction (30).

TARP interactors

TARPs interact with PSD-95-like MAGUKs (8,18). Compensatory mutations in both the PDZ domain 1 of PSD-95 and in the C-terminal PDZ domain-binding motif of stargazin/ γ -2 increase AMPAR-mediated EPSCs, whereas mutation in only one of these proteins does not (69). Furthermore, TARP interaction with PSD-95 slows AMPAR diffusion at the cell surface (1). These results suggest that TARPs interact directly with PSD-95 to control synaptic AMPARs. Other TARP interactors have also been reported. PDZ domain-containing proteins (OMP25, MUPP1, PIST, and MAGI2) and non-PDZ-containing proteins (light chain 2 of the microtubule associate protein (LC2)) were identified, in addition to PSD-95-like MAGUKs (17-19,31). It would be important to examine the distinct roles of each of these interactors in the regulation of the TARP/AMPAR complex.

4. TARPs modulate the channel properties and pharmacology of AMPARs

Synaptic strength is determined by the number and channel properties of AMPARs at synapses. TARPs modulate not only the trafficking, but also the channel properties of AMPARs. *Xenopus laevis* oocytes are widely used as a system to evaluate receptor activity. Glutamate-evoked currents and AMPAR surface expression in oocytes coinjected with GluA1 and Stargazin/ γ -2 cRNAs are significantly larger than those evoked by GluA1 alone (9,80,94). Furthermore, TARPs increase glutamate-evoked currents about four times more than they increase the surface expression of AMPARs, which suggests that TARPs increase both the trafficking and the individual channel activity of AMPARs (77).

TARPs slow the decay kinetics of AMPARs

AMPARs open their channel pore after glutamate binding, which is followed by closing of the channel pore after glutamate removal (deactivation) or with glutamate binding (desensitization). During synaptic transmission, the decay of AMPAR-mediated EPSCs is determined by deactivation and desensitization. TARPs slow both the deactivation and the desensitization processes in heterologous cells (65,77,87). Furthermore, γ -4 and γ -8 slow the decay kinetics of AMPARs to a greater extent than do γ -2 and γ -3 in heterologous cells and at synapses (11,37,54,76). Single-channel analysis revealed that TARPs increase AMPAR open channel probability by increasing burst length without changing open-dwell time, with no effect on conductance (77). This result indicates that TARPs accelerate the gating of AMPARs (77). Importantly, the decay of AMPAR-mediated EPSCs is controlled by TARPs, as overexpression of a dominant-negative form of TARP in neurons accelerated the decay of AMPAR-mediated EPSCs (77). In addition, TARPs render AMPARs more inwardly rectifying channels (11,72,73). Mutations in the channel pore of AMPARs changed the magnitude of TARP modulation of AMPAR activity (36). These observations suggest that TARPs may change the molecular environment surrounding the AMPAR-channel pore.

TARPs modulate the pharmacology of AMPARs

Native AMPARs respond to kainate more than to glutamate, whereas AMPARs expressed in heterologous cells respond to glutamate more than to kainate. Interestingly, AMPARs coexpressed with TARP respond to kainate more than to glutamate, which suggests that native AMPARs contain TARPs (11,39,54,71,76-78,87). TARPs also modulate the efficacy of AMPAR potentiators, e.g., cyclothiazide or PEPA, which slow the desensitization and deactivation of AMPARs (82,95). Importantly, cyclothiazide and TARPs shows additive effects on AMPAR activity, indicating that cyclothiazide and TARPs modulate AMPAR activity via distinct mechanisms, which is consistent with the fact that TARPs accelerate

gating and cyclothiazide slows entry into desensitization. TARPs also modulate the sensitivity to AMPAR antagonists. Surprisingly, TARPs convert the AMPAR competitive antagonists CNQX and DNQX into partial agonists (38,52). Furthermore, TARPs change the potency of the non-competitive AMPAR inhibitor GYKI53655 (15). These results suggest that inclusion of TARP in AMPARs is necessary for future drug screening.

Structural and functional analyses of TARPs and AMPARs

Extensive structure/function studies revealed that the first extracellular loop of TARPs is necessary and sufficient for the modulation of the channel properties of AMPARs (FIGURE 2A) (77, 87). Furthermore, mutations in the ligand-binding domain of AMPAR prohibit TARP interaction with AMPARs (83), which indicates that the first extracellular loop of TARPs may interact directly with the ligand-binding domain of AMPAR to modulate its channel properties. In addition to the extracellular domain, the cytoplasmic domain of TARPs was recently suggested to be involved in the modulation of channel properties (2, 53); however, its mechanism remains unclear. In contrast, chimeric and deletion studies showed that the cytoplasmic domain of TARPs is necessary and sufficient for the surface expression of AMPARs (3, 77). The C-terminal PDZ-binding motif is necessary for the synaptic localization of the AMPAR/TARP complex, as described above (1, 8, 69). Interestingly, AMPAR and TARPs accumulate in the axons of mice carrying a disruption of the β subunit of AP-4, which is an adaptor protein for protein sorting, via the interaction between AP-4 and the cytoplasmic domain of TARPs (48).

5. Regulation of AMPAR activity by TARPs

One of the intrinsic features of AMPAR is that neuronal activity modulates synaptic AMPAR activity. Two mechanisms have been proposed for the TARP-mediated dynamic regulation of AMPAR activity.

TARP phosphorylation regulates AMPAR activity at synapses

Neuronal activity increases calcium influx through NMDA-type glutamate receptor (NMDAR) to activate calcium-dependent kinases, which is followed by the increase in AMPARs at synapses (FIGURE 3) (16,43,45-47). However, the substrate of calcium-dependent kinases is unknown. TARPs are highly phosphorylated at the PSD (84). TARP phosphorylation is bidirectionally regulated by PKC and CaMKII under NMDAR activity in neurons (30,84). Furthermore, TARP carrying replacement mutations of its phosphorylated serine residues to aspartic acid (constitutive phospho-mimic) were generated and overexpression of phospho-mimic stargazin/ γ -2 increases AMPAR-mediated EPSCs specifically in neurons (34,77). These results indicate that TARP phosphorylation is a regulator of synaptic AMPAR activity and may be a substrate for NMDAR-mediated synaptic plasticity (FIGURE 3). In support of this, neuronal activity phosphorylated TARPs and PSD-95 better than other proteins in glutamate receptor complex (85).

TARPs are phosphorylated at nine serine residues in their cytoplasmic domain. It is not known how many TARP phosphorylation sites are required for the regulation of synaptic AMPAR activity. Interestingly, these phosphorylated residues are located within a short consecutive region and the total negative charge of this short stretch changes in a gradient manner. If the total negative charge is the mediator of synaptic AMPAR activity, the short stretch containing the nine phosphorylated serine residues could serve as a molecular rheostat for synaptic AMPAR activity. In addition to the nine phosphorylated serine residues within this short stretch, TARPs are phosphorylated at the threonine residue in the C-terminal PDZ-binding motif. Moreover, the TARP mutant that is phosphorylated at this threonine residue, which can be phosphorylated by PKA, does not interact with PSD-95. In

addition to regulation by calcium-dependent kinases, the localization of the AMPAR/TARP complex may be regulated by PKA (10,13).

Dynamic interaction between TARP and AMPAR

TARPs interact with AMPARs and modulate the trafficking and channel properties of AMPARs. Glutamate-induced AMPAR desensitization could induce partial or complete dissociation of AMPARs from TARPs (55,80).

In some neurons, AMPARs show a bell-shaped dose-response curve, where the amplitude of the steady-state current declines at glutamate concentrations above 100 μ M. The mechanism underlying this observation is not completely understood, as AMPAR expression in heterologous cells does not exhibit a bell-shaped dose response (27,66,90). However, AMPARs coexpressed with TARPs show such a response. Interestingly, an AMPAR–TARP covalently linked tandem protein (TARPed AMPAR) shows similar channel properties to AMPARs coexpressed with TARPs; however, TARPed AMPAR does not show a bell-shaped dose response (55). In addition, cyclothiazide blocks the reduction of AMPAR currents at higher glutamate concentrations (55). These results indicate that glutamate induces desensitization of the AMPARs that interact with TARPs and that, subsequently, TARPs dissociate from AMPARs to reduce AMPAR activity via loss of TARP modulation. Interestingly, AMPAR desensitization regulates synaptic AMPAR distribution (29). This mechanism could be due to TARP–AMPA dissociation, although it remains unclear whether AMPARs dissociate from TARPs completely or partially after AMPAR desensitization in a short time scale (10–50 ms). In contrast, relatively long exposure of AMPA (over a few minutes) induces complete dissociation of AMPAR from TARPs, which subsequently induces the internalization of AMPAR, but not of TARPs, within a few minutes (80). The regulation of the TARP–AMPA interaction allows diverse responses of AMPARs in a glutamate concentration- and exposure-time-dependent manner.

6. Neurological aspects

Ataxia and absence epilepsy

TARPs play multiple roles in AMPAR modulation in normal conditions, but also in disease conditions. The *stargazer* mouse is a spontaneous mutant mouse that exhibits ataxia and absence epilepsy (61). Although the ataxic phenotypes could be explained by loss of AMPAR activity in the cerebellum (28), the absence epilepsy phenotype is the opposite of what one would expect from the loss of AMPAR activity, because epilepsy is in general caused by synchronized and enhanced neural activity. One possible explanation for the absence epilepsy observed in the *stargazer* mouse is the disinhibition of interneurons. Stargazin/ γ -2 is also expressed in interneurons (79) and strong reduction of AMPAR activity is observed in interneurons of TARP knockout mice (50). Thus, loss of stargazin/ γ -2 could cause loss of AMPAR in some unidentified interneurons, which would lead to loss of inhibition of inhibitory neurons, i.e., hyperexcitation of neural activity and induction of absence epilepsy. Mice in which both stargazin/ γ -2 and γ -4 have been disrupted shows progression of absence epilepsy compared to mice disrupting stargazin/ γ -2 alone (41). Because each TARP functions in a redundant fashion in mice (51), the detailed analysis of TARP-isoform expression could lead to the identification of the neurons that cause these phenotypes.

Excitotoxicity

Kainate is a natural toxin from a type of seaweed and induces seizures and neurotoxicity in humans and other animals. High-affinity kainate receptors are believed to be involved in kainate-induced phenomena. Kainate receptor (GluK2) knockout mice show an altered

threshold for kainate-induced seizures and some gliosis (56). Interestingly, kainate-induced cell loss in the hippocampus is reduced in γ -8 knockout mice, whereas kainate-induced seizure was not altered (78). The difference in kainate-induced neurotoxicity is probably due to a loss of kainate sensitivity by AMPARs via the absence of TARPs in the AMPAR complex. This could mean that TARPs may be involved in excitotoxicity in neurons.

TARPs as potential drug targets

TARPs may be potential drug targets for the potentiation or suppression of AMPAR activity at synapses. AMPAR potentiators (AMPAkines) enhance cognitive function and are currently being investigated as a potential treatment for a variety of neurological disorders, including schizophrenia, Alzheimer's disease, and Parkinson's disease (42,44,62,74). Each AMPAkinase has a specific preference for the flip or flop AMPAR splicing isoform (22,63), although in the presence of stargazin/ γ -2, these potentiators can act on either isoform (82). One drawback of AMPAkinase treatment is the fact that AMPARs are ubiquitously expressed, which makes it difficult to target AMPARs in specific regions of the brain. Drugs that alter the stability of the AMPAR/TARP complex could, in principle, up- or downregulate AMPAR activity via TARP modulation of channel activity, and TARP isoform-specific drugs would allow the efficient targeting of specific brain regions (25,79). Therefore, targeting TARPs may be an effective method for regulating AMPAR activity and synaptic strength.

7. Other transmembrane auxiliary subunits of glutamate receptors

TARPs are auxiliary subunits of AMPARs; however, extensive studies have been performed to identify other subunits of ionotropic glutamate receptors, which resulted in the identification of several proteins as candidate subunits (not yet fully confirmed) (FIGURE 4).

AMPARs

A high-throughput proteomics approach identified cornichon (CNIH2 and 3) as a novel auxiliary subunit of AMPARs (FIGURE 4) (70). Cornichon is a trimembrane-spanning protein that modulates the channel properties and surface expression of AMPAR in heterologous cells (70). The precise localization of the AMPAR/CNIH complex remains unclear. Because *Drosophila* cornichon acts as a cargo receptor for ER export (5), cornichon may promote the exit of AMPAR from the ER. Further studies are required to elucidate this mechanism.

C. elegans AMPAR (GLR-1) contains another auxiliary subunit (SOL-1), which was identified in a genetic screening as a *suppressor of lurcher*, which AMPAR carrying Lurcher mutation is a constitutive active form of AMPAR (98). SOL-1 is a single-transmembrane protein that contains four CUB domains in its extracellular domain. In heterologous cells and *C. elegans*, SOL-1 slows the decay kinetics of GLR-1 AMPAR (97). Interestingly, *C. elegans* AMPAR (GLR-1), TARP (STG-1 and 2), and SOL-1 form a tripartite complex that exhibits distinct channel properties (91,92). This result suggests the existence of an unidentified mammalian SOL-1 homolog (FIGURE 4).

NMDA receptors

The neuronal protein Neto1 was identified as a retina-specific protein; subsequently, other splicing isoforms of Neto1 expressed in the brain were identified (75). Neto1 is a CUB domain-containing protein. Two other CUB domain-containing proteins, SOL-1 and LEV-10, were identified as subunits of the worm glutamate receptor (GLR-1) and of the worm acetylcholine receptor, respectively (26,98). Neto1, a protein with unknown function, is

a vertebrate protein containing two CUB domains that shared highest homology with CUB domains in SOL-1 and LEV-10. Neto1 was examined as a subunit of ion channels, NMDARs. Neto1 knockout mice exhibit NMDAR-mediated synaptic plasticity and impairment in learning tasks (60). These results propose Neto1 as a novel subunit of NMDARs (FIGURE 4).

Kainate receptors

The channel properties of native and recombinant kainate receptors are significantly different. A proteomics approach identified Neto2 as a kainate receptor-binding protein (96). Neto2 modulates the channel properties of kainate receptors in heterologous cells (96). In contrast, kainate receptors modulate the surface expression of Neto2 in heterologous cells and in neurons (96). Furthermore, overexpression of Neto2 increases the frequency of kainate receptor (KAR)-mediated events and slows the decay kinetics of KAR-mediated EPSCs (96). These results propose Neto2 as a subunit of kainate receptors (FIGURE 4). Interestingly, Neto2 shares very high homology with Neto1 identified as an auxiliary subunit of NMDARs. It is important to examine possible dual roles of Neto1 and 2 as auxiliary subunits of both KAR and NMDAR, simultaneously.

8. Concluding remarks

Recent extensive studies established TARPs as auxiliary subunits of AMPARs in the brain. TARPs interact with AMPARs and modulate AMPAR channel gating, channel pharmacology, and trafficking to the cell surface and to synapses. Several questions remain unanswered in this field, including “do all AMPAR complexes in the brain contain TARPs?”, “how does neuronal activity modulate AMPAR activity?”, and “what is the role of TARPs in neurological disorders and how should these disorders be treated?” These mechanisms should be revealed in the future. Furthermore, growing knowledge of the auxiliary and accessory subunits of ionotropic glutamate receptors will shed light on the fundamental rules that govern excitatory synaptic transmission.

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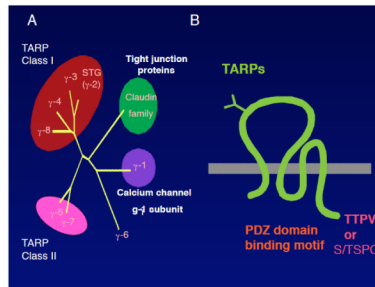


FIGURE 1. TARP structure and phylogenetic tree of TARP-related proteins

A. γ -1 is a calcium channel gamma subunit (CACNG-1). Subsequently, eight homologous genes were identified that were termed γ -1–8. Among the eight γ -1 homologous proteins, six proteins modulate AMPAR activity and were classified as class I and class II TARPs, functionally. Class I TARPs comprise stargazin/ γ -2, γ -3, γ -4, and γ -8 and Class II TARPs include γ -5 and γ -7. The roles of γ -6 remain unclear. B. TARPs are tetramembrane-spanning proteins that contain typical (–TTPV) and atypical (–S/TTPC) binding motifs for the PDZ domain in their C terminus.

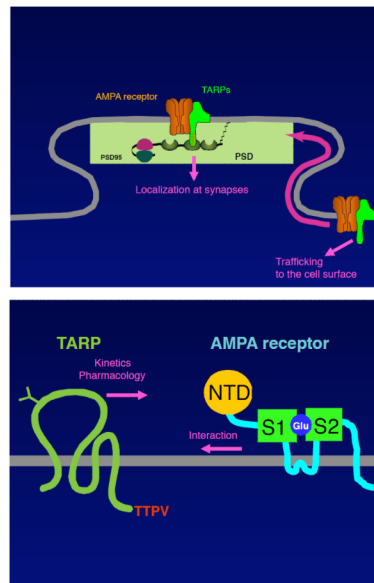


FIGURE 2. Functional domains of TARPs and AMPARs

A. The cytoplasmic domain of TARPs is necessary and sufficient for the surface expression of AMPARs. The C-terminal PDZ-binding motif is necessary for the synaptic localization of the AMPAR/TARP complex. The interaction domains of TARPs with AMPARs remain unclear. B. Extensive structure/function studies revealed that the first extracellular loop of TARPs is necessary and sufficient for the modulation of the channel properties of AMPARs. Furthermore, mutations in the ligand-binding domain of AMPARs prohibit TARP interaction with AMPARs.

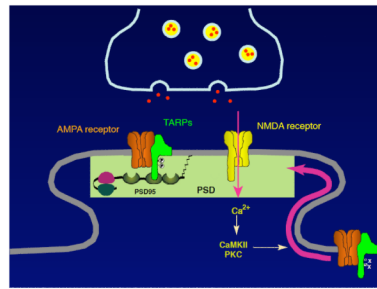


FIGURE 3. A model for the regulation of AMPAR localization

Neuronal activity increases calcium influx through NMDAR to activate calcium-dependent kinases, which is followed by an increase in AMPARs at synapses. TARPs are highly phosphorylated at the PSD. TARP is a substrate of PKC and CaMKII *in vitro*. These observations indicate that TARP phosphorylation is a regulator of synaptic AMPAR activity and may be a substrate for NMDAR-mediated synaptic plasticity.

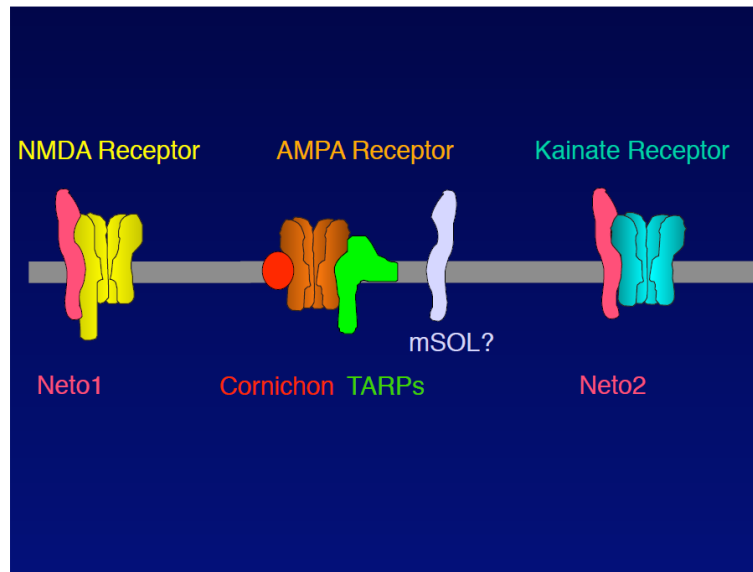


FIGURE 4. Auxiliary and accessory subunits of ionotropic glutamate receptors

Ionotropic glutamate receptors are pharmacologically classified as AMPA-, NMDA-, and kainate-sensitive glutamate receptors. Several transmembrane subunits have been proposed to date, but remain unconfirmed. The NMDAR complex comprises Neto1, which is a CUB domain-containing protein. The AMPAR complex comprises TARPs and cornichon. Notably, it remains unclear whether AMPAR/TARP/cornichon form a tripartite complex. The *C. elegans* AMPAR complex comprises SOL-1, which is a CUB domain-containing protein. Therefore, mammalian AMPAR complexes may comprise a mammalian homolog of SOL-1 (mSOL). The kainate receptor complex comprises Neto2, which is a CUB domain-containing protein. The high sequence homology between Neto1 and 2 raises the question of whether Neto1 and 2 act as auxiliary subunits for both NMDA and kainate receptors.

Table 1

TARPs modulate AMPAR functions

Function	TARP modulation	TARP domains involved in each modulation
AMPA Interaction	Interaction	No report Note, AMPAR desensitization induces TARP dissociation
Channel properties	EPSC Decay kinetics (Synaptic transmission)	Extracellular domain, TARP subfamily (γ -2/3 and γ -4/8)
	Decay kinetics (deactivation/desensitization)	Extracellular domain, Cytoplasmic domain, TARP subfamily (γ -2/3 and γ -4/8)
	AMPA open probability (Accelerating gating)	No report
	Channel rectification	No report
	KA efficacy	Extracellular domain, Cytoplasmic domain, TARP subfamily (γ -2/3 and γ -4/8)
	AMPA potentiator	No report (AMPA splicing isoform, flip/flop)
Trafficking	Convert AMPAR antagonists into partial agonists	No report
	Synaptic localization	C-terminal PDZ binding domain (-TTPV)
		TARP Phosphorylation
Surface expression	TARP cytoplasmic domain	