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Synapse to Nucleus Signaling

Toh Hean Ch'ng1 and **Kelsey C. Martin**1,2,3

Toh Hean Ch'ng: tchng@mednet.ucla.edu

¹Department of Biological Chemistry, University of California, Los Angeles, California 90095-1737, United States of America.

²Department of Psychiatry and Biobehavioral Sciences, University of California, Los Angeles, California 90095-1737, United States of America.

³Semel Institute of Neuroscience and Human Behavior, David Geffen School of Medicine, University of California, Los Angeles, California 90095-1737, United States of America.

Abstract

Signals generated in distal subcellular compartments of neurons must often travel long distances to the nucleus to trigger changes in gene expression. This retrograde signaling is critical to the development, function and survival of neural circuits, and neurons have evolved multiple mechanisms to transmit signals over long distances. In this review, we briefly summarize the range of mechanisms whereby distally-generated signals are transported to neuronal nuclei. We then focus on the transport of soluble signals from the synapse to the nucleus during neuronal plasticity.

Introduction

Activity-dependent changes in gene expression allow eukaryotic cells to alter their structure and function in response to a range of environmental stimuli. The signal transduction cascades within neurons that couple extracellular stimulation with transcription face a unique set of challenges, since signals received in distal subcellular compartments often travel long distances to reach the nucleus. The transport of distal signals into the nucleus is critical to many processes in neurons. For example, signals generated at distal growth cones trigger transcriptional changes that are essential for neuronal development [1,2]; synaptically generated signals elicit changes in transcription that are required for persistent forms of learning-related synaptic plasticity [3,4]; and signals transported from sites of axonal injury to the nucleus are critical to axonal regeneration [5•,6]. In this review, we briefly summarize the various mechanisms whereby signals are transported to the neuronal cell body and nucleus, and then focus on the active nucleocytoplasmic trafficking of soluble, synaptically generated signals during neuronal plasticity. We conclude by addressing future challenges in the field.

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Corresponding Author Kelsey C. Martin, University of California, Los Angeles, Department of Biological Chemistry, 615 Charles E. Young Drive South BSRB 390-B, Los Angeles, CA 90095, Phone - (310)-794-9502, Fax - (310)- 206-1929, kcmartin@mednet.ucla.edu (KCM).

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Rapid electrochemical signaling

Neurons utilize multiple mechanisms for signaling between distal subcellular compartments and the nucleus. Electrochemical signaling allows for extremely rapid communication to the cell body and nucleus (Fig. 1-A). Local activation of ion channels in distal compartments triggers action potentials that reach the soma within milliseconds, leading to the opening of somatic voltage-gated calcium channels. The influx of calcium at the soma activates calcium-sensitive signaling cascades that in turn activate transcription factors, thereby coupling neuronal activity to changes in gene expression within minutes [7–9].

Rapid signaling through regenerative calcium waves in the ER

Another method for rapid, calcium-dependent nuclear signaling in neurons involves regenerative calcium waves propagated along the endoplasmic reticulum (ER; Figure 1-B) [10]). In addition to its functions in the secretory pathway, the ER serves as an internal calcium store that is continuous with the nuclear membrane and extends into distal neuronal axons and dendrites [11,12]. The ER contains two calcium channels on its surface: the inositol triphosphate receptor $(InsP_3R)$ and the ryanodine receptor (RYR) , both of which are calcium-sensitive calcium channels [13]. Hence, calcium influx via cell surface voltagegated calcium channels also triggers the release of internal calcium stores from the ER, with the local release of calcium activating neighboring $InsP₃R$ and RYR, creating a regenerative calcium wave that propagates toward the nucleus.

Physical translocation of signaling molecules from distal compartments to the nucleus

Yet another mechanism for synapse to nucleus signaling involves the physical transport of signaling molecules from their site of initiation to the nucleus. This type of signaling includes the transport of signaling endosomes (Fig. 1-C) and the diffusion (Fig. 1-D) or active transport (Fig. 1-E) of soluble signaling molecules. The physical translocation of signals from distal sites to the nucleus is slower than electrochemical signaling or regenerative calcium waves in the ER and can also persist over greater lengths of time.

Neurotrophins and signaling endosomes

Compelling evidence supports a role for motor-driven retrograde transport of ligand-bound, activated neurotrophin receptors in axons via signaling endosomes (for reviews, see [14]. Many studies have focused on the binding of trophic factors such as nerve growth factor (NGF) to TrkA tyrosine receptors at axon terminals, a process that regulates neuronal survival and axonal outgrowth [15]. The NGF-bound activated TrkA receptor is endocytosed into a signaling endosome and then trafficked to the cell body via microtubules [14,16–19]. Once the endosome arrives in the soma, the active complex interacts with a range of kinases or second messenger molecules including Ras-MAPK, PI3K, PLCγ and ERK5, which in turn activate transcription factors such as CREB to trigger gene expression in the nucleus [18,20,21]. A recent study reported that signaling endosomes travel not only from distal axons to the cell body, but further into dendrites where they regulate development of the postsynaptic compartment [22••]. It is important to note that while there is a wealth of data to support the Signaling Endosome hypothesis in neurotrophic signaling, there are several alternative models such as the "wave" or the "signaling effector" models to describe the transport of signals from axon terminals [14,20,23].

Transport of soluble proteins from synapse to nucleus

One of the biggest challenges for synapse-to-nucleus trafficking is the distance proteins traverse from distal synapses to the nucleus. It is unlikely that diffusion-mediated signaling

is efficient enough to allow distally generated signals to reach the nucleus in adequate concentrations to alter transcription (Fig 1-D). Indeed, using large scale Monte Carlo simulations, Howe argues that fast local signaling via diffusion can occur effectively only at distances of less than 200 nm [24]. Hence, long distance movement of signaling proteins likely occurs via "information packets" that are transported by molecular motors [24]. Kholodenko and colleagues have also demonstrated that simple diffusion alone is insufficient for signal transfer in the MAP kinase cascade [25,26]. In contrast, Wiegert and colleagues recently reported that stimulus-induced ERK1/2 nuclear translocation is largely mediated by facilitated diffusion [27].

Analysis of motor-driven nuclear import requires consideration of neuronal polarity. Vertebrate central nervous system neurons are polarized into axons and dendrites, while many invertebrate neurons elaborate unpolarized processes, or neurites, which contain both pre- and post-synaptic elements. Microtubules in axons are uniformly aligned with minusends directed toward the cell body, and thus, the majority of signals that originate from the axons, growth cones or axon terminals engage the motor protein dynein for the retrograde movement of the signaling endosomes toward the nucleus [17,28–30]. In contrast, dendritic microtubules are of mixed polarity, although recent studies have suggested that dendritic microtubules may be polarized in an age-dependent and cell type specific manner [31,32]. It is not known whether synaptic cargoes engage only one class of motor proteins for dendritic transport, constantly "hopping" between microtubules of the same polarity, or whether they engage both plus-end (kinesin) and minus-end (dynein) directed motor proteins with regulation of these motor proteins yielding a net movement toward the cell body [33]. There is also evidence of crosstalk between microtubules and actin-driven anterograde transport [34•,35•,36], although this type of molecular motor crosstalk has not been well-examined during retrograde trafficking.

Importin-mediated nuclear transport from synapse to nucleus

In the early 1990s, Ambron and colleagues microinjected rhodamine-labeled human serum albumin coupled to the nuclear localization signal (NLS) of SV40 large T antigen into the distal growth cones of cultured *Aplysia californica* sensory neurons and found that the NLS was necessary and sufficient for translocation of the microinjected protein into the nucleus [37]. This study implicated a role for the classical nuclear import pathway (via binding of the importin α/β 1 heterodimeric complex to NLS-bearing cargo, [38]) in long-distance retrograde transport from distal sites to the soma.

Importin-mediated nuclear transport from sites of axonal injury to the nucleus in mammalian neurons has also been shown to occur during injury-induced regeneration. Hanz et al. [29] localized importin α and β1 to the axoplasm of sciatic nerve and cultured dorsal root ganglion neurons. They found that axonal injury triggered the local translation of importin $β1$ mRNA in axons, and that the newly synthesized importin $β1$ formed a transport complex with importin α and NLS-containing cargo proteins in axons to mediate nuclear import of injury-induced signals. Later studies by these authors identified phosphorylated ERK1/2 as an injury-induced, retrogradely transported cargo of importin β 1 [39]. The nuclear import of ERK1/2 by importins is thought to initiate a transcriptional program that underlies injuryinduced axonal regeneration. More recently, the Fainzilber lab used a combination of phosphoproteomics and microarray analysis to identify the axonal signaling networks and transcriptional factors that are recruited by nerve crush injury [5•].

Our laboratory defined a role for importin-mediated active nuclear import in mediating synapse-to-nucleus trafficking during long-term plasticity. We demonstrated that importins localize to distal synapses and accumulate in the nucleus in response to plasticity-inducing stimuli in both rodent hippocampal neurons and cultured *Aplysia* sensory-motor neurons

[40]. In *Aplysia*, we further demonstrated that the active nuclear import pathway is required for long-term, but not short-term forms of plasticity. More recently, we proposed a mechanism whereby activity regulates the synaptic localization of importins in hippocampal neurons. We showed that importin $α$ binds to an NLS encoded by exon 21 in the alternatively spliced NMDA receptor subunit NR1-1a. This interaction was regulated by activity, such that stimuli that induced transcription dependent long-term potentiation of hippocampal synapses triggered protein kinase C phosphorylation of serine residues within and flanking the NLS, disrupting the binding of importin α . The released importin α was then free to bind soluble cargoes and transport them to the nucleus [41•].

Studies in *Drosophila melanogaster* have revealed a role for importin-mediated nuclear import during development of the nervous system. Importin α 3 has been shown to be required for transport of dSmad2 from the growth cone to the nucleus, with null mutations in importin α 3 showing defects in the axonal tiling of photoreceptors [42]. More recently, Schwarz and colleagues reported that importin α 2, together with importin β 11, relays Wnt signals from synapse to nucleus [43••]. Specifically, they demonstrated that binding of the ligand wingless to the postsynaptic *Drosophila* Frizzled-2 (dFz2) receptor leads to proteolytic cleavage of Frz2, with the carboxy-terminal cleavage product binding to importin α 2 and importin β 11, followed by translocation to the nucleus. This synapse-tonucleus transport also requires the PDZ protein dGRIP [44]. The subsequent transcriptional activation by Frz2 promotes the elaboration of the postsynaptic plasma membrane [43••]. Additional genetic studies from the Schwarz lab revealed roles for importin α 2 in determining active zone density and axonal projections [43••,45].

Together, these studies describe roles for importin-mediated retrograde nuclear transport in axonal injury and regeneration, synaptic plasticity, axon guidance and synapse formation.

Synaptically localized signaling proteins that undergo activity-dependent nuclear import

In this section, we review specific synaptically localized proteins that undergo retrograde transport into the nucleus following distinct types of stimuli. Many of these proteins are transcription factors. As such, their nuclear import directly couples synaptic activity with changes in transcription.

cAMP Responsive Element Binding protein -2 (CREB2/ATF4)

We recently showed that the transcriptional repressor CREB2 (also known as ATF4) translocates from distal neuronal processes to the nucleus during long-term depression (LTD) of cultured rodent hippocampal neurons and FMRF-amide induced LTD of *Aplysia californica* sensory motor synapses [46]. Biochemical studies revealed that ATF4/CREB2 binds specifically to importins α 1 and α 6. Consistent with importin-mediated transport, retrograde trafficking was blocked by saturating concentrations of NLS peptides.

Nuclear factor kappa of light chain enhancer of activated B-cells (NF–κB)

The NF–κB complex consists of a transcriptionally active dimer that is anchored in the cytoplasm through its interaction with an inhibitory subunit [47]. In neurons, the complex most commonly contains a p65 and p50 transcriptionally active dimer and the $I\kappa Ba$ inhibitory subunit [48], and is found at synapses [49–51]. Several labs have demonstrated that transcriptionally active p65 subunits fused to eGFP translocate into the nucleus in an NLS-dependent manner upon glutamatergic stimulation [50,52]. This nuclear translocation of active p65 requires association with microtubules via the dynein/dynactin molecular motor protein complex [53,54]. An intriguing report by Marcora et al. suggested that NF–κB

translocate from activated synapse to the nucleus by associating with wildtype (but not mutant) huntington protein via association with importin α 2 [51].

ApCAM Associated Protein (CAMAP)

Long-term synaptic strengthening of *Aplysia* sensory-motor synapses is accompanied by new synaptic growth, which has been shown to involve the internalization and degradation of the Aplysia cell adhesion molecule ApCAM [55]. A yeast two-hybrid screen for ApCAM-interacting proteins identified a novel transcriptional coactivator of CRE-driven gene expression, CAMAP [56]. Local synaptic stimulation triggered CAMAP phosphorylation by PKA, dissociation from the cytoplasmic tail of ApCAM and translocation of CAMAP into the sensory neuron nucleus. In the nucleus, CAMAP bound CREB1a and promoted transcription of specific CRE-driven genes [56].

Amyloid Precursor Protein Intracellular Domain Associated Protein-1 (AIDA-1)

Mass spectrometric analysis of PSD fractions from rat brain led identified AIDA-1 as a synaptically localized protein [57]. Subsequent analysis of AIDA-1 revealed that it binds PSD95 and that NMDA receptor activation drives proteolysis and subsequent nuclear translocation of AIDA-1 into the nucleus [58]. The proteolytically cleaved N-terminus of AIDA-1 contains a functional NLS that mediates its transport into the nucleus [58,59]. In the nucleus, AIDA1 accumulates in nucleoli, sites of ribosomal RNA transcription and assembly [58,60]. Functional studies suggest that the synapse-to-nucleus transport of AIDA-1 regulates nucleolar number and global protein synthesis during persistent synaptic stimulation [58].

Jacob

Jacob was identified as a binding partner of the neuronal calcium sensor caldendrin that undergoes regulated synapse-to-nucleus trafficking. NMDA receptor activation and calcium influx was found to trigger dissociation of Jacob from caldendrin, unmasking of an NLS, subsequent binding to importin α 1, and nuclear import [61•]. Unlike AIDA-1, the nuclear translocation of Jacob requires activation of extrasynaptic NR2B-containing NMDA receptors, activation of which are thought to trigger the cell death pathway by activating a CREB "shutoff" mechanism [9]. Subsequent studies by Kindler et al. revealed that NMDA receptor-induced and calpain-mediated proteolysis of the myristoylated N-terminal fragment of Jacob is required for nuclear translocation [62].

Abelson Interacting Protein-1 (Abi-1)

Abi-1 is a substrate for the non-receptor tyrosine kinase c-Abl and is a binding partner for the postsynaptic density protein ProSAP2/Shank3. In developing neurons, Abi-1 is responsible for regulating dendritic outgrowth and determining the morphology and number of synaptic contacts [63,64]. In mature neurons, Abi-1 is found in dendrites, and a brief application of NMDA in both cultured neurons and acute rodent brain slices resulted in the reversible, stimulus-dependent, translocation of Abi-1 from dendrites into the nucleus [63,64]. Importantly, this nuclear translocation is dependent on an active microtubule network and does not require new protein synthesis. Abi-1 was also shown to associate with the Myc/Max complex of transcription factors that enhance E-box regulated gene transcription [64].

The above examples represent a subset of the synaptically localized proteins that are known to translocate into the nucleus following specific stimuli. Additional examples include transmembrane proteins that undergo Regulated Intramembrane Proteolysis (RIP) followed by nuclear import, such as Amyloid Precursor Protein (APP) and erbB4. APP is found in

both axons and dendrites and undergoes γ-secretase cleavage to generate a cytoplasmic fragment (AICD) that forms a potent transcription complex with Fe65 and Tip60 [65–67]. ErbB4 is a receptor tyrosine kinase that is cleaved upon neuregulin-induced stimulation to produce the NLS-bearing intracellular fragment E4ICD, which subsequently binds to the signaling protein TAB2 and corepressor N-CoR for nuclear translocation [68,69]. Another transmembrane protein that is proteolytically cleaved in the C-terminal is the L-type voltage-gated calcium channel $Ca_v1.2$. The cleaved fragment, CCAT (calcium channel associated transcription regulator) is regulated by intracellular calcium and translocates into the nucleus where it binds the nuclear protein p54(nrb)/NonO to activate transcription [70].

Future challenges

At present, the kinetics of soluble protein transport is not well-characterized. It is not known if soluble proteins are transported individually, in a signaling complex, or how they engage the molecular motor proteins for active transport. Some technical challenges, such as visualizing the long distance transport of cargo proteins from distal synapses to the nucleus, can be surmounted through the use of photoconvertible fluorescent fusion proteins in conjunction with time-lapse imaging [71]. Choosing a microscope and an image capture system with high spatial and temporal resolution is critical since nucleocytoplasmic shuttling is rapid. Newly developed superresolution microscopy techniques provide unprecedented resolution, and advances in instrumentation promise to decrease acquisition time [72]. Equally important is the design of software capable of identifying, tracking and quantifying the movement of soluble proteins, which often exists as a rapidly moving "wave."

Synaptic inputs received by neurons are discrete and localized, and thus it is important that stimuli be delivered to small regions such as dendritic branches, groups of, or even individual synapses rather than to the entire cell. Many parameters of long-range protein transport in dendrites, such as the kinetics and directionality of movement, may be altered if stimuli are not compartmentalized. Methods for local, synaptic stimulation include photolysis of caged compounds and local perfusion of stimuli. Recent developments in microfluidic technology provide a means of culturing neurons in compartmentalized chambers in which soma and distal processes can be stimulated with great spatial resolution [73•].

Fundamental questions about the function of synapse-to-nucleus signals remain unanswered: What types of stimuli trigger the translocation of synaptically localized signals to the nucleus, and what changes in gene expression and neuronal function do such signals induce? Is there a threshold number of stimulated synapses that is required for effective synapse-tonucleus signaling? What types of plasticity (e.g. Hebbian versus homeostatic) recruit synapse to nucleus signaling? And finally, can the cell keep track of individual synapses that have been stimulated to send a signal to the nucleus, such that any downstream transcriptional changes function to specifically alter the efficacy of those synapses? Future experiments taking advantage of technical advances in local stimulation and live-cell microscopy hold the promise of providing answers to these questions.

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neuronal processes. This allows users to manipulate specific dendritic and axonal segments separately from cell bodies.

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1. . The different mechanisms of synapse to nucleus signaling

Signals generated in distal compartments (axons and dendrites) in neurons are transmitted to the nucleus by multiple mechanisms. Electrochemical signaling (A) and regenerative calcium waves in the endoplasmic reticulum (B) allow for extremely rapid signaling. Signals received at growth cones and axon terminals can be internalized into signaling endosomes that are transported back to the nucleus by molecular motors (C). Soluble proteins can be physically transported from distal sites to the nucleus by passive or facilitated diffusion (D) or by active motor-driven transport (E). All black arrows indicate the net movement of proteins, vesicles or ions in neurons following stimulation. All solid black lines indicate cytoskeletal filaments.