# Multifaceted Role of Heat Shock Protein 70 in Neurons

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**Abstract** Heat shock protein 70 (Hsp70) plays important roles in neural protection from stress by assisting cellular protein folding. In this review we discuss the current understanding of inducible and constitutive Hsp70 in maintaining and protecting neuronal synaptic function under normal and stressed conditions.

**Keywords** Hsc70 · Hsp40 · HSPA · Synapse · Calcium · Neuronal protection

#### Introduction

Heat shock protein (Hsp) is a family of proteins that were originally found to respond to heat shock stimuli [1] but also associated with other cellular stressors such as alcohol [2–4], oxidants [4–7], heavy metals [8–10], amino acid analogues [11, 12], hypoxia [13, 14] and glucose deprivation [15, 16]. Hsp family proteins are in general molecular chaperones that serve cytoprotective roles by maintaining or assisting cellular protein folding. Although Hsp proteins come from diverse families and differ in their structures and cellular functions [17], they all contain a highly conserved heat shock element (HSE) in the promoter region of their corresponding genes [18]. Hsp proteins are subgrouped based on their molecular weights. The structural features and functional roles are highly conserved within the

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members of each subfamily but not observed among members of different subfamilies [19].

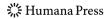
Hsp70, a subgroup of heat shock proteins with the molecular weight of 70 kDa, is the most ubiquitous Hsp subgroups, and structurally and functionally highly conserved across species. Under physiological conditions, Hsp70s serve as molecular chaperones. The major constitutive isoform of Hsp70, heat shock cognate 70 (Hsc70), has the ability to bind to exposed hydrophobic surfaces of various substrates and prevent nonproductive protein–protein interactions that would lead to aggregation, and it promotes protein refolding.

Under stressed conditions, upregulation of Hsps, such as the inducible Hsp70, confers protection at all levels from the whole organism [20], specific organs [21], cells [22] to specific organelles [23]. Increasing Hsp expression level by overexpression [24], drug induction [21], genetic manipulation [25] and preconditioning [20, 26–28] protects the cells/organisms from lethal conditions. The mechanisms underlying the cytoprotective effects of heat shock proteins, specifically the Hsp70s, have been well delineated. Hsp70 expression reduces protein aggregation, maintains mitochondria physiology, inhibits apoptosis/necrosis and suppresses inflammatory responses [29].

Of great importance is the role of neuronal Hsps at the synapse, a site that is sensitive to stress and prone to stress-induced damage. This minireview highlights the distinct functional roles of inducible and constitutive isoforms of the 70-kDa Hsp proteins at the neuronal synapse both under physiological conditions and in response to stress.

Structural Features of Hsp70s

Similar to other *Hsps* [30, 31], *Hsp*70s genes encode the HSE in their promoter region. The HSE consists of three



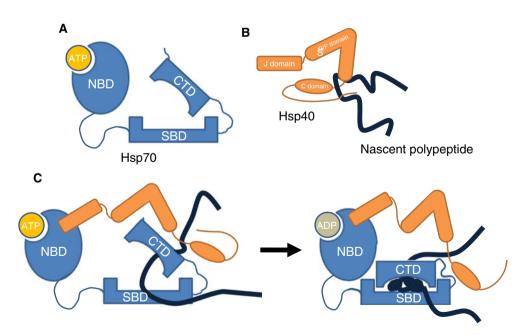
contiguous inverted repeats of the A/GGAAn sequence: each repeat interacts with one of the trimeric heat shock transcription factor (HSF) [32]. The number of pentameric repeats varies greatly among different heat shock proteins. For example, the promoter of *Hsp*40 has eight contiguous sequences [33], Hsp70 has five and Hsp90 has six such sequences [34]. Additional pentameric sequences results in greater affinity interactions between HSFs and HSE [35]. Hsp70s have three structural domains (Fig. 1a); a 44kDa N-terminal domain with ATPase activity (NBD), an 18-kDa substrate binding domain (SBD) and a 19-kDa Cterminal domain (CTD) that assists with substrate binding [36-38]. The concerted actions of all three domains are required for Hsp70 function. For example, the binding affinity of the SBD is dependent on the state of the NBD, such that there is increased substrate affinity when Hsp70 is adenosine diphosphate (ADP)-bound and decreased affinity when it is adenosine triphosphate (ATP)-bound [39].

The gene sequences of Hsp70s are highly conserved from bacteria to human. The human hsp70-gene family is evolutionarily diverse [40]. There are at least 14 isoforms of Hsp70 in humans encoded by 17 different genes [40, 41] (Table 1). There are another 30 sequences coded as hsp70 pseuodogenes, structurally characterized with variations including frame shifts and in-frame stop codons [40]. The hsp70 genes are located on 18 chromosomes except 15–17, 19 and 22 [40, 41]. The classification and localization of

the identified Hsp70s are summarized in Table 1. The mitochondrial Hsp70 (Hsp70-9; GRP75) is encoded by a gene at chromosome segment 5q31.1 [42, 43]. The gene for ER homologue of Hsp70 (GRP78) [44] is localized to human chromosome 9q34 [45]. The major constitutively active form of Hsp70, Hsc70, is encoded by a gene located on chromosome segment 11q24 [46]. Hsc70 is abundantly expressed under physiological conditions and is not significantly heat inducible. Its major functions include protein folding and translocation, clathrin uncoating and peptide binding to prevent aggregation [47]. A number of Hsp70s are stress-inducible (Table 1). The major inducible Hsp70s (or Hsp72) are encoded by HSP1A and HSP1B [48, 49], which are located on chromosome segment 6p21.3 [50, 51]. The expression level of these Hsp70s is minimal under normal conditions but is greatly upregulated after exposure to stress such as hypoxia, ischemia, hyperthermia and toxins [52]. Most genes encoding Hsp70 do not contain introns; an exception is found in the case of Hsc70 and Grp78, offering the possibility of alternative splicing [41].

### Cofactors of Hsp70s

Two main cofactors have been identified for Hsp70s, namely, Hsp40s [33] and nucleotide exchange factors (NEFs) [53]. Hsp40s are homologues of the bacterial DnaJ [33], which generally contains three distinct domains



**Fig. 1** Schematic illustration of Hsp70 and Hsp40 complex. **a** Hsp70 has three major functional domains. The NBD has ATPase activity and drives the conformational change that allows its interaction with the two other domains. The SBD contains the substrate affinity site and CTD forms the lid on top of the SBD. **b** Three structural domains of Hsp40: the J domain consisting of histidine, proline and aspartic acid motif, the glycine/phenylalanine-rich G/F domain, and cysteine-

rich region of the C domain. Hsp40 interacts with nascent peptide via a peptide-binding site located on the C-terminal. c The peptide-bound Hsp40 is anchored to Hsp70 through interaction between the G/F domain of Hsp40 with a C-terminus sequence of the CTD of Hsp70. The J domain of Hsp40 interacts with the NBD domain of Hsp70 to increase ATPase activity

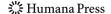


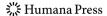
Table 1 Classification of human heat shock proteins

Gene	Approved names	Alternative/synonyms	Localization	Stress inducible	References
HSPA1A	Heat shock 70 kDa protein 1A	Hsp70-1	Nucleus, cytoplasm, liposome	Y	Wu et al [49] Mol Cell Biol 5:330-341; Milner and Campbell [48] Immunogenetics 32:242-251.
HSPA1B	Heat shock 70 kDa protein 1B	Hsp70-2	Nucleus, cytoplasm, lysosome	Y	Wu et al [49] Mol Cell Biol 5:330-341; Milner and Campbell [48] Immunogenetics 32:242-251.
HSPA1L	Heat shock 70 kDa protein 1-like	Hsp70-HOM, Hum70t	Nucleus, cytoplasm	N	Goate et al. [50] Hum Genet 75:123-128.
HSPA2	Heat shock 70 kDa protein 2		Nucleus, cytoplasm	N	Bonnycastle et al. (1994) Genomics 23: 85-93.
HSPA4	Heat shock 70 kDa protein 4	Hsp70, RY	Cytoplasm	Y	Kang CM et al (2002) Radiat Res 157: 650-655.
HSPA4L	Heat shock 70 kDa protein 4 L	APG-1, Osp94	Cytoplasm	Y	Held et al. (2006). Mol Cell Biol 26: 8099-8108.
HSPA5	Heat shock 70 kDa protein 5	BiP, GRP78	ER	N	Munro and Pelham [44] Cell 46: 291-300.
HSPA6	Heat shock 70 kDa protein 6	HSP70B'	Nucleus, cytoplasm	Y	Leung et al. [5] Biochem J 267: 125-132.
HSPA7	Heat shock 70 kDa protein 7	HSP70B	Nucleus cytoplasm	Y	Leung et al. [5] Biochem J 267: 125-132; Siddiqui et al. (2008) Cancer Invest 26: 553-561.
HSPA8	Heat shock 70 kDa protein 8	Hsc71, Hsc70, Hsc73	Nucleus, cytoplasm	N	Dworniczak and Mirault (1987) Nucleic Acids Res 15: 5181-5197.
HSPA9	Heat shock 70 kDa protein 9	GRP75, PBP74, mot-2, mtHSP75	Mitochondria	N	Domanico et al. [43] Mol Cell Biol 13: 3598-3610; Bhattacharyya et al. [42] J Biol Chem 270:1705-1710.
HSPA12A	Heat shock 70 kDa protein 12A	FLJ13874, KIAA0417	Nucleus, cytoplasm	_	Pongrac et al (2004) Biol Psych 56:943-950
HSPA13	Heat shock 70 kDa family, member 13	STCH	ER	N	Yamagata et al (2008) Biochem Biophys Res Commun 376:499-503.
HSPA14	Heat shock 70 kDa protein 14	Hsp70-4, Hsp70L1	Cytoplasm	Y	Wan et al (2004) Blood 103:1747-1754; Otto et al (2005) Proc Natl Acad Sci U S A 102:10064-10069.

(Fig. 1b). (1) A highly conserved 70-amino-acid J domain near the N-terminus. The J domain consists of a highly conserved sequence of histidine, proline and aspartic acid (HPD motif); it mediates the interaction between Hsp40 and Hsp70 and stimulates the ATPase activity of Hsp70. (2) A glycine/phenylalanine (G/F) rich region acting as a flexible linker. (3) A cysteine-rich region near the C-domain containing four CXXCXGXG motifs. All DnaJ homologues contain the N-terminal J domain and either one or both the other two domains [54]. Based on this variability, DnaJ homologues are classified into three groups. Type I homologues contain all three domains. Type II homologues have the J and G/F domain but lack the C domain. Type III Hsp40s contain only the J domain. The human Hsp40 is a type II homologue [54].

There are 41 DnaJ homologues identified in mammals, which greatly outnumbers the amount of Hsp70s [55]. Thus, it is suggested that each Hsp70 chaperone complex consists of multiple Hsp40s [33]. Functionally, Hsp40s

have no chaperone activity, but their intracellular localization may determine the functional specificity of each Hsp70/Hsp40 chaperone complex [55]. The substrate polypeptides first bind to Hsp40, and this complex then interacts with Hsp70/ATP [53]. The substrate is transferred to the open peptide-binding cleft of Hsp70 and locked in by subsequent Hsp40 stimulated hydrolysis of ATP (Fig. 1c). For example, the simian virus 40 virus large T antigen protein contains a J domain that recruits Hsp70 to drive mitosis [56]. Similarly, the Hsp40 homologue in yeast Sec63 recruits Hsp70 to the ER through its J domain to assist in protein unfolding prior to membrane transport [57]. NEFs [53] regulate the rate ADP/ATP exchange and serve as an important regulatory point for Hsp70 function [52]. Depletion of NEFs Fes1 or Sse 1 in Saccharomyces cerevisiae impairs prion propagation. The prolonged binding between Hsp70 and its substrate in the absence of NEFs prevents the protein from reverting back to its prion state [52].



#### Hsc70 at the Synapse

Under physiological conditions, the Hsc70/Hsp40 complex plays a prominent role in maintaining and regulating various components of the synapse. Presynaptically, Hsc70 regulates the N-type calcium channels, the rate of synaptic vesicle recycling and the rate of neurotransmitter synthesis and packaging. The potential mechanisms of Hsc70 involvement at the synapse are proposed in Fig. 2 (endocytotic mechanisms) and Fig. 3 (exocytotic mechanisms).

Hsc70 is not a native protein at the synapse but is recruited there by several DnaJ homologues (Hsp40s) that are natural residents of the presynaptic terminal. One such protein is the cysteine string protein (CSP) [58]. CSP is mainly expressed in the presynaptic terminal of neurons and other secretory cells, where it is involved in exocytosis of neurotransmitters, hormones and enzyme precursors. CSP knockout (KO) flies show impaired presynaptic neuromuscular transmission with loss of synaptic vesicles [59]. CSP is a type III Hsp40 protein and is composed of three domains [58]; it has an N-terminus containing a J domain (residues 1–82) that allows binding to Hsc70, a 30residue (83–112) linker region followed by the C-terminus (residues 113-198) region. The C-terminus is the least conserved region among all three CSP isoforms and is believed to be responsible for its flexibility and diversity in cellular processes. The Hsc70/CSP complex forms a trimer with a third protein, the small glutamine-rich tetratricopeptide repeat protein (SGT) that binds to the C-terminus of

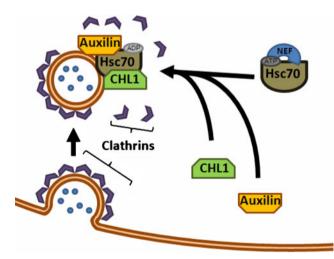


Fig. 2 Schematic of Hsc70 associated proteins in endocytosis. Cytosolic Hsc70 is targeted to the synapse by CHL1 and auxilin. Upon binding, the CHL1/Hsc70 or the auxilin/Hsc70 complex stimulates the clathrin decoating activity of Hsc70, greatly enhancing synaptic recycling and thus increasing the number of available synaptic vesicles for subsequent exocytotic events. *ADP* adenosine diphosphate; *ATP* adenosine triphosphate; *CHL1* close homolog of L1; *Hsc70* heat shock protein cognate 70; *NEF* nucleotide exchange factor

CSP [60]. SGT contains three tandem tetratricopeptide repeat domains (TPRs) with a degenerate 34-amino-acid sequence [58]. The complex is further stabilized by the interactions between Hsc70 and SGT, mediated by the binding between the TPR domain of SGT with the C-terminus of Hsc70 [60]. Binding of the protein complex is dependent on the presence of ADP, while ATP promotes the disassembly of the complex. This complex formation stimulates the ATPase activity of Hsc70 by 19 folds and thus greatly enhances its protein refolding abilities on the synaptic vesicle surface [58].

# Inhibition of N-Type Calcium Channel

G-proteins are heterotrimeric GTP-binding proteins composed of  $\alpha$ ,  $\beta$  and  $\gamma$  subunits [61]. Activation by exogenous ligand results in exchange of GDP for GTP on  $G_{\alpha}$  subunit and dissociation of  $G_{\beta\gamma}$  subunit from the trimeric G-protein complex. This step is regulated by the Hsc70/CSP/SGT trimer ([61]; see also [62]). Full-length CSP protein interacts with  $G_{\alpha}$  and increases the steadystate GTP hydrolysis, and the rate of hydrolysis is further increased in the presence of the Hsc70/SGT. The Nterminal (residues of CSP1-112) of CSP alone can bind to the GDP-bound inactive form of  $G_{\alpha}$  and stimulate GDP/ GTP exchange in a manner that is unaffected by the presence or absence of Hsc70/SGT. Thus, it was proposed that the C-terminal of CSP (residues 112-192) is a regulatory domain that inhibits the activity of CSP and this inhibition is abolished by the binding of Hsc70/SGT [61].

The two activated subcomplexes,  $G_{\alpha}$  and  $G_{\beta\gamma}$ , have distinct cellular functions [63]. The  $G_{\alpha}$  subunit binds to the N-terminal J domain of CSP in an ATP-dependent manner, whereas the  $G_{\beta\gamma}$  subunits associate with the C-terminus of CSP independently of ATP. The complex of CSP and  $G_{\beta\gamma}$ subunits binds to the  $\alpha_{1B}$  subunit of N-type calcium channels, resulting in tonic inhibition of calcium influx. The association of either  $G_{\alpha}$  or  $G_{\beta\gamma}$  to CSP can cause Ntype calcium channel inhibition via different mechanisms. Inhibition of calcium channel by CSP requires direct physical binding of  $G_{\beta\gamma}$  subunit to CSP C-terminus. Interaction between  $G_{\alpha}$  subunit and CSP J domain, on the other hand, does not require this direct interaction. This association is thought to inhibit the assembly of the trimeric G-protein, freeing the  $G_{\beta\gamma}$  subunits to interact with CSP Cterminus, thus leading to inhibition of N-type calcium channel activity [63]. Hsc70, in conjunction with CSP and SGT, promotes the activity of G proteins with the end effect of tonic inhibition of N-type channels. In the presence of Hsc70 and SGT, CSP is activated and promotes the GDP/ GTP exchange on  $G_{\alpha}$  subunit, releasing the  $G_{\beta\gamma}$  subunit to inhibit N-type calcium channels.



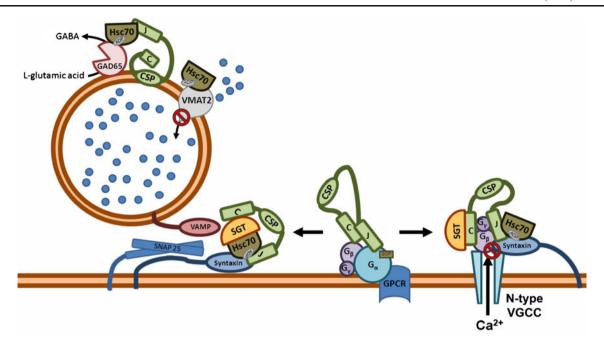


Fig. 3 Schematic of Hsc70-associated proteins in exocytosis. Hsc70/CSP/SGT trimer catalyzes the steady-state GTP hydrolysis on  $G_{\alpha}$ , thereby promoting the dissociation of the trimeric G-protein complex into the  $G_{\alpha}$  and  $G_{\beta\gamma}$  subunits. Subsequent association of  $G_{\beta\gamma}$  with CSP tonically inhibits calcium influx through N-type calcium channels. Hsc70/CSP complex also anchors GAD65 to the synaptic vesicle and stimulates GAD65 activity, thereby promoting synthesis of GABA. Lastly, Hsc70 interacts with and inhibits the activity VMAT2.

ADP adenosine diphosphate; ATP adenosine triphosphate; C Cterminus on CSP; CSP cysteine string protein; GDP guanosine diphosphate; GPCR G-protein coupled receptor; GTP guanosine triphosphate; Hsc70 heat shock protein cognate; J J domain on CSP; N-type VGCC N-type voltage gated calcium channel; SGT small glutamine-rich tetratricopeptide; VAMP vesicle-associated membrane protein (synaptobrevin); VMAT2 vesicle monoamine transporter 2; GAD65 L-glutamic acid decarboxylase 65

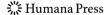
Syntaxin 1A, a tSNARE protein, is critical for neurotransmitter release. Syntaxin 1A incorporation into the SNARE complex is required for vesicle fusion and exocytosis. Syntaxin 1A associates with Hsc70, CSP and N-type calcium channels [64] and binds to the  $\alpha_{1B}$  subunit of N-type calcium channels, in the same region that CSP binds. Its specific function in this complex has yet been determined. The association between syntaxin and Hsc70 has been shown to promote syntaxin incorporation into the SNARE complex [64]. Thus, it appears that the Hsc70/ syntaxin complex performs antagonistic roles at the synapse. Syntaxin and Hsc70, as part of the multimeric complex with CSP, SGT and G proteins, inhibit calcium influx through N-type calcium channels [64] and thus may prevent exocytosis. This may serve as a negative feedback mechanism to control synaptic transmission.

## Synaptic Recycling—CHL1 and Auxilin

Hsc70 at the synapse is involved in clathrin uncoating during endocytosis [65]. Synaptic vesicles, upon releasing their content at the synaptic cleft, is internalized via formation of clathrin-coated pits on the synaptic plasma membrane, which leads to formation of clathrin-coated vesicles (CCVs). Removal of clathrin from vesicles is an

essential process for repackaging synaptic vesicles and for subsequent exocytotic events [66]. Hsc70 binds to clathrin molecules and catalyze uncoating of clathrin from CCVs, but this proceeds at a low rate. CHL1 is a cell adhesion molecule of the immunoglobulin superfamily. It normally accumulates in presynaptic membranes and contains an HPD tripeptide sequence (J domain) in its intracellular Cterminal, which allows interaction with Hsc70 [67]. It is proposed that upon synaptic activation (depolarization), CHL1 recruits Hsc70 to the synaptic membrane from the cytosol and forms a dimeric complex in an ADP-dependent manner. Binding with CHL1 enhances the uncoating ability of Hsc70 [67] and thus aids in synaptic recycling. Neurons from CHL1 KO mice show decreased level of Hsc70 at the synapse, an increase in CCVs and decreased number of available synaptic vesicles in subsequent exocytotic events [67]. CHL1 KO mice show reduced synaptic plasticity and mental retardation [68]. In humans, this gene has been linked to schizophrenia, which is characterized by abnormal synaptic neurocircuits [69]. In addition to CHL1, CSP has been proposed as an alternate for Hsc70 targeting to the synapse, and may play similar roles as CHL1 in endocytosis, as it has been found to localize on CCVs [67, 70].

Synaptic recycling also involves the association between auxilin and Hsc70. Auxilin is a type III member of DnaJ



chaperone family and a native resident of CCVs [71]. It contains a J domain that allows it to bind to Hsc70s [71] and to recruit Hsc70 to the synaptic vesicle membrane. ATP-dependent binding of auxilin stimulates the ATPase activity of Hsc70, which in turn catalyzes clathrin decoating. Blocking of binding between auxilin and Hsc70 inhibits synaptic transmission in the giant squid axon [72].

There are several similarities between auxilin and CHL1. They both reside mainly in the presynaptic terminal and are critical for successful synaptic vesicle recycling. They recruit Hsc70 to the synapse with their J domain, in a manner similar to that of Hsp40. Thus, it is likely that Hsc70 interacts with both auxilin and CHL1 in a functional synapse at the synapse, although the exact relationship between CHL1 and auxilin has yet been determined.

## Neurotransmitter Packaging

Hsc70 and VMAT2 Vesicle monoamine transporter (VMAT) is responsible for storing monoamines such as dopamine, norepinephrine and serotonin into synaptic vesicles [73, 74]. Hsc70 binds to the N-terminus of VMAT2 with its own SBD and CTD domains. This interaction results in a decrease in VMAT2 activity [75] and thus may result in regulation of neurotransmitter packaging into synaptic vesicles.

Hsc70 and GABA synthesis γ-Aminobutyric acid, GABA, is one of the major inhibitory neurotransmitters in the central nervous system [76]. It is synthesized from Lglutamate via the enzyme L-glutamic acid decarboxylase (GAD) [77]. There are two forms of GAD, GAD65 and GAD67, so named because of their molecular weights. The membrane-associated GAD (MGAD) is concentrated on synaptic vesicle membranes and is composed mainly of GAD65. The soluble GAD pool is concentrated in the cytosol and is composed mainly of GAD67. MGAD is the major contributor for GABA synthesis, and the activity of MGAD is dependent on its association with the synaptic membrane. GAD65 proteins lack the integral membrane stretch that is characteristic of most membrane proteins [78-80]. Binding to Hsc70 and CSP, GAD65 can become associated with the synaptic vesicle membrane, thus increasing its ability to synthesize GABA [76]. After exocytosis, these synaptic vesicles are recycled through clathrin-mediated endocytosis, whose activity is regulated by the Hsc70/CSP/SGT trimer previously described.

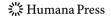
## Hsp70 in the Stressed Synapse

Stressed cells display several distinct characteristics, including an increase in resting calcium concentration, decrease in pH, generation of reactive oxygen species and alteration of membrane potential. At the synapses, there is impairment in both the presynaptic and postsynaptic function. Induced expression of Hsp70 is vital for counteracting the effects of stress-induced abnormalities at the synapse.

Study by Karunanithi et al. [27] first showed that in *Drosophila melanogaster*, induction of Hsp70 prior to heat shock protects individual synapses from subsequent hyperthermic insult. Specifically, heat shock preconditioning prevents the reduction of the quantal content (number of vesicles liberating neurotransmitter per stimulus) and stabilizes presynaptic functions. It also maintains the postsynaptic currents at the neuromuscular junction at a constant amplitude in heat-shocked flies as temperature increased [27]. These synaptic changes correlate well with the time course of elevation of Hsp70 after heat shock. Thus, synaptic protective effects of thermal preconditioning are partly attributable to the upregulation of Hsp70 [81, 82].

Not all organisms show upregulation of Hsp70 in response to stress. For example, populations of the Rana temporaria tadpoles from southern climates show less induction of Hsp70 expression following heat shock than those from a northern population [83]. Also, the thermotolerant species of hydra, Hydra attennata, is able to synthesize large amounts of Hsp60 after heat shock, whereas the thermosensitive Hydra oligactis cannot [84]. Furthermore, the desert species of fruit flies Drosophila arizonae show increased thermotolerance compared to the standard D. melanogaster fruit flies despite lower amount of upregulation of Hsp70 [85]. It was suggested that as D. arizonae flies had higher basal expression of Hsp70, this was able to afford heat shock protection [85]. Recent data suggest that a different sets of heat shock proteins are activated in thermosensitive species of D. melanogaster (hsf<sup>4</sup> mutant) [86]. Upon heat shock, there is significant upregulation of Hsp83 and Dna-J 1 protein without significant changes in expression of Hsp70. The function of Hsp83 is vague. The homologue of Dna-J1, Hsp40, is involved in repairing damaged proteins in a complex with Hsc70/Hsp70 [86]. Furthermore, in response to heat shock, Hsp40 forms a complex with CSP at the synapse, disrupting CSP dimerization and promoting CSP-mediated GTP hydrolysis on  $G_{\alpha}$  subunit, in a manner that is similar to that of Hsc70 [87]. Given the expression level and functions of Hsp83 and Dna-J/Hsp40, these heat shock proteins can provide significant thermoprotection in the absence of inducible Hsp70.

Presynaptically, elevation of Hsp70 may stabilize presynaptic protein expression. Work from our laboratory show that early induction of Hsp70 stabilizes presynaptic protein expression and prevents hypoxia-induced motor and sensory depression in *Lynmaea stagnalis*, fresh pond water snails.



The snails exposed to chronic hypoxia show slowed sensory reaction to light stimuli and reduced motor movements [13, 14, 88]. The neural suppression of the snail coincides with an increase in Hsp70 expression and decreased expression of several proteins critical for exocytosis, including syntaxin I, synaptic vesicle protein 2 (SV2) and synaptotagmin [13, 14, 88]. Time-course analysis shows an early induction of Hsp70 within 6 hours, downregulation of syntaxin in the first 24 hours, a delayed reduction of synaptotagmin after 4 days and a biphasic response in SV2. Prevention of upregulation of Hsp70 with RNAi leads to further suppression of these presynaptic proteins and exacerbation of neurobehavioural dysfunction. Coimmunoprecipitation assay shows that Hsp70 directly interacts with syntaxin [13]. Consistent to the previous findings in neuromuscular junction in D. melanogaster [27], our results indicate that early induction of Hsp70 during hypoxic stress is critical in maintaining the expression level of presynaptic proteins, thus preventing aggravation hypoxic-induced neurobehavioural dysfunction [13, 14, 88].

Hsp70 may also confer protection through association with calcium channels. In addition to the N-type calcium channel, Hsp70s interact with the cytosolic loop of Ca<sub>V</sub>2.3 R-type voltage-gated calcium channel [89]. The interaction may be important in decreasing the intracellular calcium overload that accompanies hyperthermic insults [25]. Prior to heat shock or induced elevation, Hsp70 at the *D. melanogaster* neuromuscular junction stabilizes the intracellular calcium concentration by maintaining near-resting calcium level and promoting calcium clearance in subsequent hyperthermic episodes. Furthermore, stabilization of the intracellular calcium level is correlated with better synaptic transmission and locomotory behaviour [25].

Hsp70 confers protection to the presynaptic terminal by maintaining the normal expression level of presynaptic proteins and by regulating calcium fluxes through interaction with presynaptic calcium channels. The protective role of Hsp70 at the postsynaptic terminal is unclear. Recent research shows that other heat shock proteins (e.g., Hsp90) associate with postsynaptic potassium [90, 91] and chloride channels [22] to produce hyperpolarization that may counteract the stress-induced depolarization. Thus, in response to stress, the actions of heat shock proteins strive to return the physiological environment of the neuronal synapse to a resting condition, thereby stabilizing both presynaptic and postsynaptic functions.

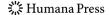
## Hsp70 in Neuroprotection

The protective role of Hsp70 extends in neurons beyond the confines of the synaptic terminal. The detailed evidence on

the neuroprotective role of Hsp70 has been thoroughly reviewed preciously [92, 93]. Hsp70 exerts antiinflammatory actions [94], upregulates anti-apoptotic molecules and downpregulates proapoptotic molecules [95] in ischemic cells. It also prevents protein aggregation and facilitates protein trafficking, and thus leads to its potential applications in "protein misfolding disorders" such as Parkinson's and Alzheimer's diseases [95-97]. The neuroprotective effects of Hsp70 are not limited to its role in neurons. In astrocytes, overexpression of Hsp70 reduces apoptosis and necrosis induced by glucose deprivation and oxygen glucose deprivation [95, 98]. In mice cocultures of microglia and astrocytes, Hsp70 suppresses activation of the inflammatory transcription factor nuclear factor kB (NFkB) via direct binding, and reduces the expression level of NFkB regulated genes, resulting in higher survival of neighbouring astrocytes [94]. This interplay between different cell populations of the CNS is also observed between astrocytes and neurons [92]. Neurons internalize the extracellular Hsp70s secreted by neighbouring glial cells during cell stress. The released Hsp70s during stress may function as a physiological alarm signal for cell trauma; however, whether the endogenous Hsp70s play a similar role remains unclear [92]. Thus, in the stressed CNS, Hsp70 may transcend its classic role as a molecular chaperone and emerge as a multifaceted protector for the specialized synapse, the damage-prone neuronal protoplasm and the supporting astrocytes.

## **Perspectives**

The current understanding is that Hsp70 acts as the major protective molecule against stress. However, Hsc70 is mostly expressed in the cytosol, and thus must be targeted to the specific sites such as synapse, by local proteins in order to perform specific functions. That is, the functional role of Hsc70 is largely determined by its essential cofactors, such as the Hsp40s [33]. At the synapse, Hsc70 participates in both exocytosis and endocytosis through association with different sets of proteins, although the overall mechanisms share great similarity. In all cases, Hsc70 is recruited to the synaptic site by native synaptic proteins containing a J domain. Binding of Hsc70 to these proteins is ATP/ADP dependent. Therefore, manipulation of the levels of native synaptic proteins, their Hsc70 binding abilities or the levels of ATP/ADP can alter the levels of Hsc70 at the synapse, and consequently promote or hinder Hsc70-mediated processes (e.g., clathrin uncoating and synaptic protein refolding). Thus, identification the targets and/or binding partners of Hsp70 is critical for our understanding of the role of Hsp70 in neuronal functions.



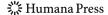
There are several differences between Hsc70 and Hsp70s. First, Hsc70 contains several introns, whereas Hsp70 is intronless [41]. The structural simplicity of Hsp70 may enable its quick induction in response to stress. However, the functional and perhaps structural consequence of this genetic difference is unclear. Next, with regard to the synapse, Hsc70 mainly participates in the maintenance of normal synaptic functions such as exocytosis and endocytosis. Hsp70, however, functions to counteract the effects of stress so that the processes involving Hsc70 can proceed, thus illustrating a functional distinction between Hsc70 and Hsp70. This distinction is further exemplified by the fact that, currently, there is no common target observed (at the synapse). Also, study has shown direct antagonistic effects of Hsc70 and Hsp70, such as in the case of the epithelial sodium channels [99]. Currently, it is premature to conclude whether Hsc and Hsp70 share more differences or similarities. Further research thus is warranted to elucidate this matter.

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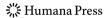
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