



Methamphetamine-Induced Neuronal Damage: Neurotoxicity and Neuroinflammation

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Abstract

Methamphetamine (METH) is a highly addictive psychostimulant and one of the most widely abused drugs worldwide. The continuous use of METH eventually leads to drug addiction and causes serious health complications, including attention deficit, memory loss and cognitive decline. These neurological complications are strongly associated with METH-induced neurotoxicity and neuroinflammation, which leads to neuronal cell death. The current review investigates the molecular mechanisms underlying METH-mediated neuronal damages. Our analysis demonstrates that the process of neuronal impairment by METH is closely related to oxidative stress, transcription factor activation, DNA damage, excitatory toxicity and various apoptosis pathways. Thus, we reach the conclusion here that METH-induced neuronal damages are attributed to the neurotoxic and neuroinflammatory effect of the drug. This review provides an insight into the mechanisms of METH addiction and contributes to the discovery of therapeutic targets on neurological impairment by METH abuse.

Key Words: Methamphetamine, Neurotoxicity, Neuroinflammation, Excitotoxicity, Apoptosis

INTRODUCTION

Methamphetamine (METH) is a well-known psychostimulant that can cause neurotoxicity and is one of the most widely abused drugs worldwide (Elkashaf *et al.*, 2008). The continuous use of METH promotes neurodegeneration and cognitive decline (Rusyniak, 2011; Dean *et al.*, 2013). In addition, chronic METH abuse is reported to cause selective patterns of brain deterioration leading to memory impairment (Meredith *et al.*, 2005).

The abuse of METH is closely related to the release of neurotransmitters such as dopamine (DA) (Saha *et al.*, 2014; Lin *et al.*, 2016). During the development of drug addiction, drug-seeking behavior proceeds from seeking the reward effect of drugs to being triggered by drug-associated cues (Robbins *et al.*, 2008). Therefore, greater decrease in dorsal striatal DA in METH abusers might promote habitual drug use (Wang *et al.*, 2012). METH increases DA neurotransmission via regulation of dopamine transporters (DATs) activity (Lin *et al.*, 2016; Sambo *et al.*, 2017). A recent study shows that a decrease of DATs in METH abusers increases the risk of developing Parkinson's disease (Chen *et al.*, 2013; Granado *et al.*, 2013). Parkinson's disease (PD) is caused by degeneration of DA

neurons in the midbrain. Biochemical and neuroimaging studies of human METH users revealed that the levels of DA and DATs were decreased, and microglia activation in striatum and other areas of the brain was also detected, which appears to be similar to that observed in PD patients (Granado *et al.*, 2013). METH is also known to cause neuronal inflammation, which eventually leads to neural degeneration (Cadet and Krasnova, 2009). Directly or indirectly, METH-induced neuroinflammation makes the brain more susceptible to neuropathology (Cadet and Krasnova, 2009).

Neuronal cells are highly susceptible to pro-inflammatory cytokine-induced damage, and exposure to pro-inflammatory cytokines has been shown to cause neuronal cell apoptosis (Castino *et al.*, 2007). Moreover, neuroinflammation can increase the oxidative stress by excessive release of harmful reactive oxygen species (ROS), which further promote neuronal damage and subsequent inflammation resulting in a feed-forward loop of neurodegeneration (Fischer and Maier, 2015). There are a number of excellent reviews outlining the health and societal concerns stemming from METH abuse and overdose, yet there remains a paucity of information related to neuroinflammation and neurotoxicity in METH abusers (Matsumoto *et al.*, 2014). Therefore, to provide a guide for future

Open Access <https://doi.org/10.4062/biomolther.2020.044>

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Received Mar 24, 2020 Revised Jun 23, 2020 Accepted Jun 25, 2020

Published Online Jul 15, 2020

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research, we want to review neuronal cell apoptosis through neurotoxic and neuroinflammatory mechanisms caused by METH.

METH-INDUCED NEUROTOXICITY

Dopaminergic pathway

Methamphetamine is a psychostimulant that primarily induces the release of dopamine, serotonin and norepinephrine (Rothman *et al.*, 2001). These neurotransmitters are involved in neuronal cell inflammation and necrosis in the mesolimbic region of the brain (Panenka *et al.*, 2013). The process of intoxication of METH is closely related to the induction of DA release. Chronic METH intake regulates dopamine release by acting primarily on vesicle monoamine transporter-2 (VMAT-2) and plasma membrane DATs, two major molecules of the dopaminergic neuronal terminal (Fig. 1) (Kahlig and Galli, 2003). DATs are responsible for dopamine reuptake into the presyn-

aptic dopaminergic neurons from the extracellular area, which is extremely important for regulating and maintaining dopamine homeostasis (Fleckenstein *et al.*, 2007). Under normal circumstances, neuronal activation promotes the release of DA into the synapse (Nickell *et al.*, 2014). The DATs removes DA from the synapse, and the VMAT-2 transports cytoplasmic DA into vesicles for storage, release, and protection from oxidation and reactive consequences (Riddle *et al.*, 2006). However, METH causes abnormal trafficking of DATs, which means that METH increases extracellular dopamine levels by inhibiting dopamine reuptake, stimulating dopamine efflux, and internalizing DATs from the plasma membrane (Riddle *et al.*, 2006). Moreover, METH increases the excitability of dopaminergic neurons in a DATs-dependent manner. The DAT is a member of Na⁺/Cl⁻ dependent co-transporters (Sonders *et al.*, 1997), and bidirectional transport of dopamine through DATs is achieved by the movements of Na⁺/Cl⁻ ions. METH enhances DATs-mediated inward current and promotes the excitability of dopamine neurons (Chu *et al.*, 2008; Schmitt and Reith, 2010; Saha *et al.*, 2014).

VMAT-2 is an integral membrane protein that transports monoamines from the intracellular cytosol into synaptic vesicles (Fleckenstein *et al.*, 2009). However, METH causes synaptic vesicles to leak monoamines into the cytosol by disrupting the hydrogen pump-mediated proton gradient (Fleckenstein *et al.*, 2007). Moreover, METH binds to VMAT-2 and competitively inhibits the uptake of monoamines leading to high concentrations of monoamines in the cytoplasm (Sulzer *et al.*, 1992, 1993). Moreover, dysfunction of VMAT-2 due to METH interferes with physiological storage of DA, resulting in a significant increase in DA levels in endogenous cells (Lazzeri *et al.*, 2007; German *et al.*, 2012). Thus, high concentrations of DA, which can freely diffuse in cells, can easily cause large amounts of oxidative damage, which is associated with the neurotoxic effects of large amounts of METH (Hogan *et al.*, 2000; Volkow *et al.*, 2001; Eyerman and Yamamoto, 2007).

METH-induced neurotoxicity

Upon METH stimulation, large amounts of DA from cytosol and synaptic clefts are oxidized to quinone or semi-quinone. And, increasing of DA oxidation further leads to significant production of reactive oxygen species (ROS) such as hydroxyl radicals (OH⁻), hydrogen peroxide (H₂O₂) and superoxide anions (O²⁻) (Yang *et al.*, 2018). These ROS can inhibit mitochondrial adenosine triphosphate (ATP) production, which in turn results in a depolarized mitochondrial membrane potential and mitochondrial dysfunction (Stokes *et al.*, 1999; Zhu *et al.*, 2006; Dawson and Dawson, 2017). Dysfunction of mitochondrial metabolism has been reported to play a very important role in METH-induced neurotoxicity, because it inhibits the Krebs cycle and electron transport chain (ETC) and potentiates oxidative stress (Ares-Santos *et al.*, 2013). Therefore, defects in mitochondrial respiration can cause neuronal cell death and neurodegenerative diseases.

The hypothesis about the involvement of glutamate (Glu) in METH toxicity is supported by the discovery that METH causes Glu release in the brain (Baldwin *et al.*, 1993; Abekawa *et al.*, 1994). Glu is a major excitatory neurotransmitter in the brain and has been reported to play an important role in the excitotoxicity induced by METH (Moratalla *et al.*, 2017). Specifically, large amounts of Glu by METH activate the N-methyl-D-aspartate receptor (NMDAR) and metabolic glutamate

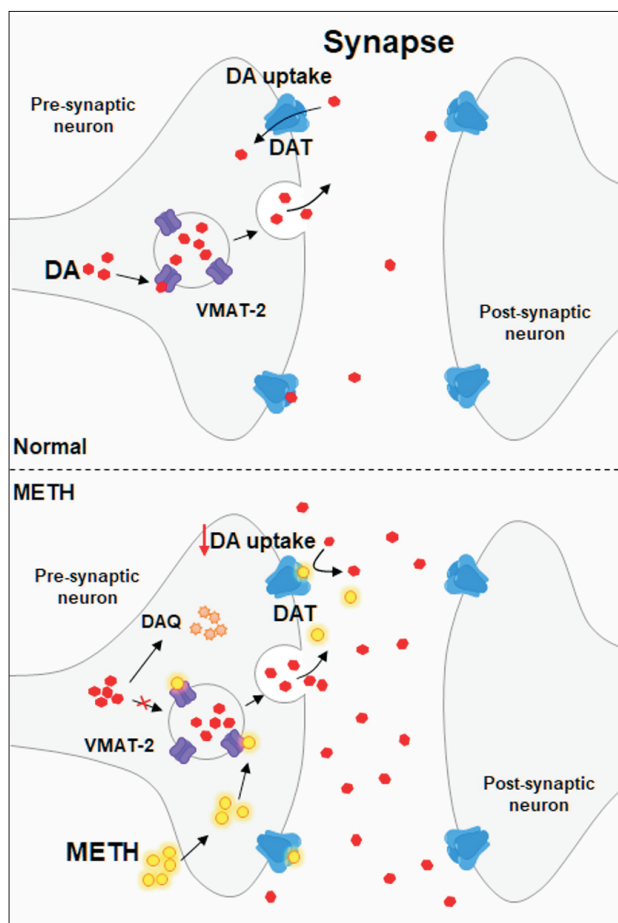


Fig. 1. METH regulates dopamine release by acting on DAT and VMAT-2. Vesicles containing DA are transferred to the extracellular space (synapse) and DA is released. In normal conditions, DAT mediates the DA reuptake, however, METH causes DA accumulation in the synapse by blocking DA uptake via interaction with DAT. METH also causes synaptic vesicles to leak monoamines into the cytosol and promotes the generation of dopamine-quinone (DAQ), which results in neurotoxicity.

receptor (mGluR) (Ohno *et al.*, 1994; Battaglia *et al.*, 2002; Tseng *et al.*, 2010). Glu accumulation overstimulates various downstream signal transduction pathways associated with Ca^{2+} influx, which leads to increased intracellular Ca^{2+} concentrations (Chamorro *et al.*, 2016). The excessive production of Ca^{2+} in cells activates protein kinases, phosphatase, and nitric oxide synthase (NOS) and promotes NO production (Moratalla *et al.*, 2017). Excessive NO production leads to endoplasmic reticulum (ER) stress, activation of the apoptotic pathway, and eventually causes neurotoxicity by METH (Moratalla *et al.*, 2017). Previous report supported that glutamate-mediated NO formation may also be involved in METH toxicity because knockout mice lacking neuronal nitric oxide synthase (nNOS or iNOS) are protected from METH-induced damage from monoaminergic axons (Itzhak *et al.*, 1998). In addition, in many studies, various nNOS inhibitors are also known to protect against the depletion of monoaminergic axons caused by METH administration (Itzhak *et al.*, 2000; Sanchez *et al.*, 2003). These evidences indicate a glutamate/NO pathway plays a major role in METH-induced neurotoxicity (Fig. 2).

METH-induced neuroinflammation

METH is also known to contribute to neuronal inflammation through excessive release of DA and Glu (Kohno *et al.*, 2019). The released DA is oxidized to form toxic quinones, leading to presynaptic membrane damage via oxidative stress, mitochondrial dysfunction and the subsequent production of peroxide radicals and hydrogen peroxide (Kohno *et al.*, 2019). The impairment of mitochondrial energy metabolism as well as the release of inflammatory cytokines increases the response to synapses and neuroinflammation (Li *et al.*, 2008; Tocharus *et al.*, 2010; Panenka *et al.*, 2013; Loftis and Janowsky, 2014). It has been reported that these METH-induced neuroinflammation is caused by targeting microglia, the innate immune cells of the central nervous system (Sekine *et al.*, 2008).

Indeed, METH-mediated activation of microglia is associated with Toll-like receptor 4 (TLR4), which is involved in immune surveillance of pathogens and exogenous small molecules (Bachtell *et al.*, 2015; Du *et al.*, 2017). TLR4 is a receptor that can activate both the Myd88-dependent and Myd88-independent pathways (Billod *et al.*, 2016). In the Myd88-dependent pathway, Myd88 activates tumor necrosis factor receptor-related factor 6 (TRAF6), interleukin-1 receptor related kinase (IRAK) to induce nuclear factor- κ B (NF- κ B) activation (Shen *et al.*, 2016). Consequently, the activation of TLR4 due to METH increases inflammatory mediators such as interleukin (IL)-1 α , 1 β , tumor necrosis factor (TNF)- α and IL-6 (Wan *et al.*, 2017). In contrast, the MyD88-independent pathway leads to the induction of IFN- γ through the activation of TRIF-related adapter molecule (TRAM) and interferon regulatory factor 3 (IRF3) (Brempeles *et al.*, 2017). The MyD88-independent pathway also induces NF- κ B activation, but it occurs later than activation through the MyD88-dependent pathway (Liu *et al.*, 2012). NF- κ B is a well-known transcription factor involved in neurodegenerative progression, and it is considered to be a key target for prevention and treatment of neurodegenerative diseases (Majdi *et al.*, 2019).

Sig-1R is an ER chaperone protein that is widely expressed throughout the brain and has a high affinity for METH (Hayashi *et al.*, 2010). Sig-1R is closely related to toxicity and inflammation caused by METH (Hedges *et al.*, 2018) via regulation of various mechanisms such as calcium homeostasis, gluta-

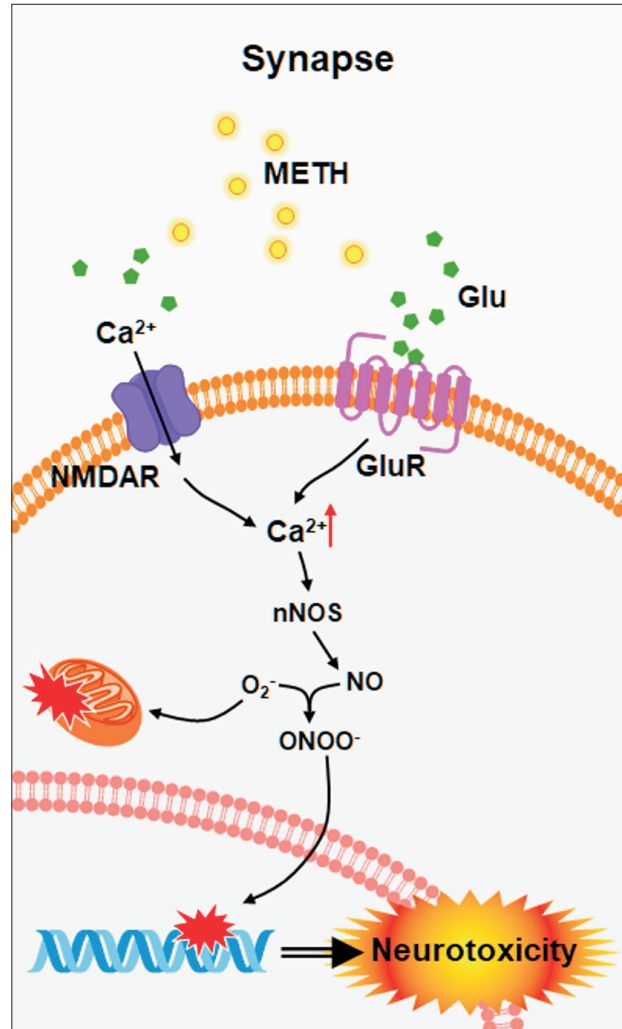


Fig. 2. METH induces Glu-mediated neurotoxicity. METH causes Glu accumulation in the synapse, and high concentration of Glu stimulates downstream pathways associated with Ca^{2+} influx. The excessive Ca^{2+} mediated neurotoxicity by activation of various enzymes related to DNA damage and ER stress.

mate activity, ROS formation, ER and mitochondrial function (Nguyen *et al.*, 2015; Ruscher and Wieloch, 2015). Another study reported that activation of microglia due to METH stimulation can be mediated by Sig-1Rs via ROS generation and activation of mitogen-activated protein kinase (MAPK) and phosphoinositide 3-kinase (PI3K)/Akt pathways (Chao *et al.*, 2017). The MAPK signaling pathway is also closely related to the NF- κ B signaling pathway (Zanassi *et al.*, 2001; Lee *et al.*, 2006; Chen *et al.*, 2009), with both playing key roles in the induction of inflammatory cytokines by METH (Liu *et al.*, 2012).

ERK is a representative kinase that plays an important role in regulating neuronal and behavioral processes mediated by DA and Glu (Shiflett and Balleine, 2011). ERK is activated by neurotrophin or growth factor (Sun *et al.*, 2016), and phosphorylated ERK is translocated to the nucleus and subsequently phosphorylates Elk-1 (Besnard *et al.*, 2011). Activated Elk-1 promotes immediate early gene (IEG) transcription associated with neural adaptation (Davis *et al.*, 2000). Another study

demonstrated that the ERK signaling pathway is linked to the regulation of dopamine D1 receptor involved in rewarding effects induced by METH (Mizoguchi *et al.*, 2004). It has also been reported that METH can increase the activation of ERK phosphorylation in certain brain regions (Son *et al.*, 2015). Once activated, ERK causes cAMP response element binding protein (CREB) phosphorylation and enhances the expression of c-Fos (Valjent *et al.*, 2005). CREB is a transcription factor that is phosphorylated by different kinases, including protein kinase A (PKA) and protein kinase C (PKC) (Johannessen and Moens, 2007; Shin *et al.*, 2012). CREB phosphorylation sequentially promotes the recruitment of co-activators such as CREB-binding protein (CBP)/p300 to the basal transcriptional machinery, which is followed by increased expression of target genes such as Arc, c-Fos, Egr1, Fos-b, and brain-derived neurotrophic factor (BDNF) (Barco *et al.* 2005; Beaumont *et al.*, 2012). A previous study supported that METH self-administration was accompanied by increased recruitment of phosphorylated CREB on the promoter of c-Fos (Krasnova *et al.*, 2016). These are important processes that promote neurological inflammation by releasing various pro-inflammatory factors such as IL-6, IL-1 β , TNF- α , monocyte chemoattractant protein 1 (MCP-1), and cell adhesion molecule (ICAM-1) (Fig. 3) (Snider *et al.*, 2013; Yang *et al.*, 2018).

APOPTOSIS DUE TO METH-INDUCED NEUROTOXICITY AND INFLAMMATION

Mitochondria-mediated death pathway

As mentioned above, cytotoxicity and inflammation caused by METH leads to neuronal cell death. Besides ROS and NO, B-cell lymphoma 2 (Bcl-2) family proteins are also involved in METH-induced neurotoxicity and inflammation (Jayanthi *et*

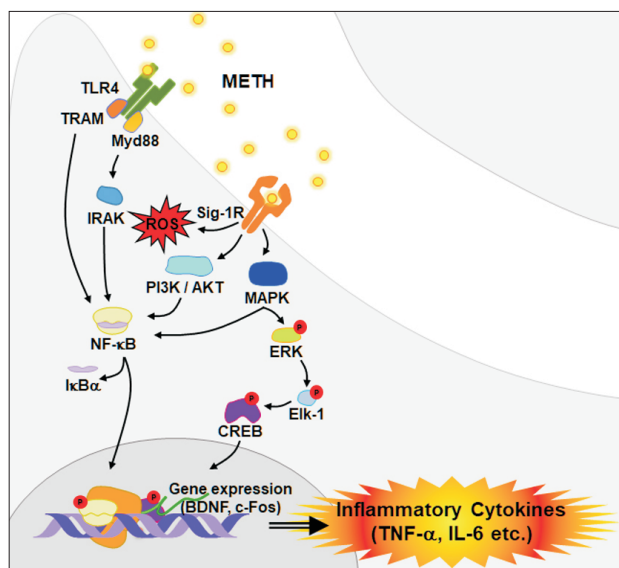


Fig. 3. METH contributes to neuroinflammation. METH activates TLR4 and Sig-1R, triggering downstream signal pathways including NF- κ B, MAPK and PI3K/Akt. Activation of CREB, c-Fos and BDNF promotes nerve inflammation through expression of various inflammatory cytokines.

al., 2001). Previous studies have reported that METH exposure increases the expression of pro-apoptotic proteins such as Bax, Bad, Bid and decreases the expression of anti-apoptotic proteins such as Bcl-2 and Bcl-xL (Jayanthi *et al.*, 2001, 2004; Beauvais *et al.*, 2011). The increase of pro-apoptotic proteins by METH is due to the release of mitochondrial intermembrane space (IMS) proteins, including apoptosis inducing factor (AIF) and cytochrome c (Galluzzi *et al.*, 2009). AIF and second mitochondria-derived activator of caspases/direct IAP-binding protein with low isoelectric point, PI (SMAC/DIABLO), which are released from mitochondria, activate the caspase-9 and -3 to induce neuronal cell death (Cadet *et al.*, 2005). The release of cytochrome c is another key step in the caspase-dependent mitochondrial apoptotic pathway (Shin *et al.*, 2018). Cytochrome c forms apoptosome, which is composed of Apaf-1, dATP and procaspase-9, and then induces sequential activation of the executioner caspases-3, -6 and -7 (Shin *et al.*, 2018). Many studies regarding METH-mediated apoptosis show increased cytochrome c release from mitochondria and subsequent caspase activation after METH exposure *in vitro* (Nam *et al.*, 2015; Park *et al.*, 2017) and *in vivo* (Deng *et al.*, 2002; Jayanthi *et al.*, 2004; Beauvais *et al.*, 2011; Dang *et al.*, 2016). Another study suggested that activation of caspase-3 and PARP in the brain was also associated with METH toxicity (Deng *et al.*, 2002). Therefore, these findings suggest that METH also affects neuronal cell death via regulation of mitochondrial pathway in the brain (Fig. 4).

ER-Dependent Death Pathway

In addition to the mitochondria-mediated apoptosis path-

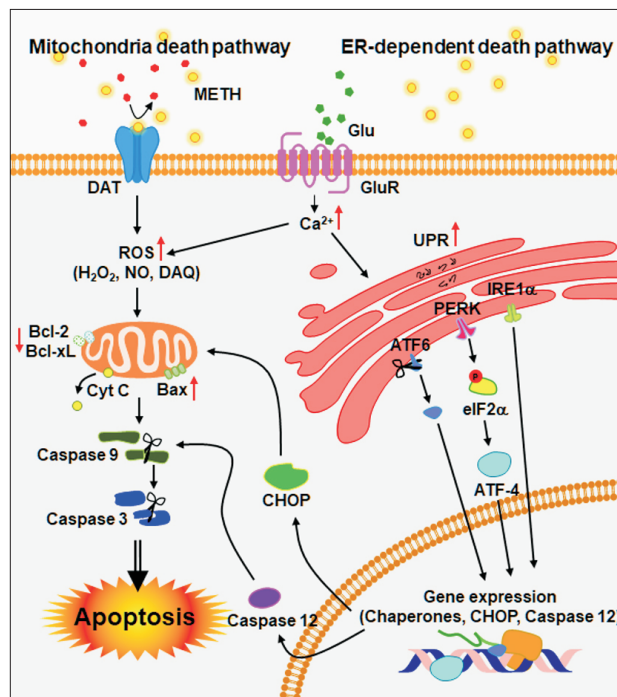


Fig. 4. METH-induced neurotoxicity and neuroinflammation cause neuronal cell apoptosis. Neurotoxicity and neuroinflammation pathways are involved in METH-induced apoptosis. Increasing of DA and Glu by METH produce ROS and Ca²⁺ that act as secondary messengers for mitochondria- and ER-mediated apoptosis.

way, METH is related to the ER-dependent cell death pathway (Koumenis *et al.*, 2002; Shah and Kumar, 2016). Oxidative stress due to METH exposure can cause cellular damage by causing dysfunction of cellular organelles such as the ER (Choi *et al.*, 2010; Wang *et al.*, 2016). Moreover, METH-mediated oxidative stress increases the expression of ER-resident chaperones such as BiP/GRP-78, P58IPK, and heat shock protein (HSP), which are important regulators of abnormal protein folding. ER stress can initiate an unfolded protein response (UPR) to restore proteolysis or to induce apoptosis (Shen *et al.*, 2004). ER stress is also closely linked to three major signaling molecules: (1) activating transcription factor 6 (ATF6), (2) inositol requiring protein-1 (IRE-1), and (3) protein kinase RNA (PKR)-like ER kinase (PERK) (Shah and Kumar, 2016). The activity of these three molecules collectively constitutes an UPR (Tabas and Ron, 2011). ATF6 acts as a transcription factor for UPR induction, while phosphorylation of IRE-1 leads to the expression of ER-resident proteins such as BiP/GRP-78, GRP94 and C/EBP homologous proteins (CHOP)/growth arrest, and DNA damage-inducing gene 153 (Gadd153) (Tabas and Ron, 2011). In addition, PERK induces phosphorylation of eukaryotic initiation factor-2 α (eIF2 α), which results in the stimulation of activating transcription factor 4 (ATF-4), C/EBP homologous protein (CHOP), and caspase-12 (Gorlach *et al.*, 2006). Since the ER contains the majority of intracellular Ca²⁺, the released Ca²⁺ from the ER is absorbed by the mitochondria which then promotes ATP production (Gorlach *et al.*, 2006).

As such, previous studies have shown that METH induces the expression of several ER stress genes, including 78kDa glucose regulated protein (GRP-78), CHOP, and ATF4, which leads to neurotoxicity in rat striatum (Bahar *et al.*, 2016). Another study suggests that METH-induced apoptosis is mediated by ER-dependent mechanisms including CHOP, spliced X-box binding protein 1 (XBP1), caspase-12, and caspase-3 (Xiong *et al.*, 2017). In addition, a relatively high dose of METH promotes dopaminergic neuronal apoptosis via nuclear protein 1 (Nupr1)/CHOP pathway (Xu *et al.*, 2017). ER stress and dysregulation of calcium homeostasis appear to be involved in neuronal cell death because METH can induce the activation of calpain (Suwanjang *et al.*, 2010). The increased calpain in METH exposure is associated with the cytoskeleton protein spectra and microtubule tau activity in rat striatum and the hippocampus (Fig. 4) (Warren *et al.*, 2005; Staszewski and Yamamoto, 2006).

CONCLUSIONS

METH is an addictive psychostimulant that acts on the central nervous system through various physiological pathways. Chronic use of METH can lead to memory deficit, and the deterioration of attention and executive functioning, which can be attributed to the direct neurotoxic and inflammatory effects of the drug. Cumulative studies have revealed the neurological effects of METH intake, however, specific mechanisms underlying METH-mediated neuronal damages remain unclear.

In this review, we focused on the neurotoxicity and neuroinflammation caused by METH, which lead to neuronal cell death and impairment of brain function. We demonstrate that the process of neuronal damage by METH is closely related to oxidative stress, regulation of transcription factor, DNA dam-

age, and various apoptosis pathways.

We hope that this review will help understanding the molecular mechanisms related to METH-induced brain damage and studies targeting the discovery of METH addiction therapy.

ACKNOWLEDGMENTS

This research was supported by Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Ministry of Education (NRF-2016R1A6A1A03011325), and by the Keimyung University Research Grant of 2018 (BDP).

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