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Review

A Mechanistic Review on Toxicity Effects of Methamphetamine

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Abstract

Persistent methamphetamine use causes many toxic effects in various organs, including the brain, heart, liver, kidney and eyes. The extent of its toxicity depends on numerous pharmacological factors, including route of administration, dose, genetic polymorphism related to drug metabolism and polysubstance abuse. Several molecular pathways have been proposed to activate oxidative stress, inflammation and apoptosis: B-cell lymphoma protein 2 (Bcl-2)-associated X (Bax)/Bcl2/caspase-3, nuclear factor erythroid 2-related factor (Nrf2)/heme oxygenase-1 (HO-1), protein kinase B (Akt)/mammalian target of rapamycin (mTOR)/p70S6K, trace amine-associated receptor 1 (TAAR1)/cAMP/lysyl oxidase, Sigmar1/ cAMP response element-binding protein (CREB)/mitochondrial fission-1 protein (Fis1), NADPH-Oxidase-2 (NOX-2), renal autophagy pathway, vascular endothelial growth factor (VEGF)/phosphatidylinositol-3-kinase (PI3K)/ protein kinase B (Akt)/endothelial nitric oxide synthase (eNOS), Nupr1/Chop/P53/PUMA/Beclin1 and Toll-like receptor (TLR)4/MyD88/TRAF6 pathways. The activation promotes pathological changes, including the disruption of the blood-brain barrier, myocardial infarction, cardiomyopathy, acute liver failure, acute kidney injury, chronic kidney disease, keratitis, retinopathy and vision loss. This review revisits the pharmacological profiles of methamphetamine and its effects on the brain, heart, liver, eyes, kidneys and endothelium. Understanding the mechanisms of methamphetamine toxicity is essential in developing treatment strategies to reverse or attenuate the progress of methamphetamine-associated organ damage.

Keywords: cardiotoxicity, hepatotoxicity, renal, meth, neurotoxicity

Introduction

The World Drug Report 2023 reported 296 million drug users in 2021, with 39.5 million categorised as a clinical drug use disorder. Amphetamines (amphetamine and methamphetamine) were the third commonly used drug, contributing to 12.2% of the overall drug users, equivalent to 36 million people. In the USA, the estimated rates of adult lifetime and past-year users of methamphetamine from 2015 to 2018 were 14.7

million and 1.6 million, respectively, with half (52.9%) of the past-year users meeting clinical use disorder criteria [1]. The global geographical distribution of methamphetamine use varies, with extensive use reported in North America, Southern Africa, East and Southeast Asia, Australia and New Zealand [2].

Methamphetamine is a synthetic drug classified as an amphetamine-type stimulant together with amphetamine, methcathinone, and ecstasy-group

drugs including 3,4-methylenedioxymethamphetamine (MDMA). Its clinical use is limited to relieving nasal decongestion and treating ADHD (aged ≥ 6 years) and morbid obesity to a limited extent [3, 4]. The abuse potential of methamphetamine is attributable to its psychostimulant properties on the short-term general physical well-being, such as energy and sexual performance [5]. The reasons for methamphetamine use in patients on opioid replacement therapy are multifactorial. One possible factor is a short-term enhancement in sexual performance to alleviate sexual dysfunction, a common adverse effect in this population [5-7].

Persistent methamphetamine use, however, leads to many deleterious effects on many organs, including the brain, heart, liver, kidneys and eyes. There is a similarity of psychopathological symptoms between certain psychiatric disorders (schizophrenia spectrum and other psychotic disorders, bipolar and other related disorders, including anxiety disorders) and some patients with methamphetamine use disorders [8-10]. Moreover, methamphetamine may induce other diseases, including myocardial infarction, cardiomyopathy, acute liver failure, acute kidney injury, chronic kidney disease, hypertension, keratitis, retinopathy and vision loss. Various factors, including route of administration, dose, genetic polymorphism related to drug metabolism and polysubstance abuse, can modify the risk of methamphetamine toxicity [9]. Given the significant negative impacts of methamphetamine on public health, understanding the effects of methamphetamine is crucial. This review aims to revisit the pharmacological properties of methamphetamine along with its neurotoxic, cardiovascular, hepatotoxic and ophthalmologic effects in clinical populations and preclinical models.

Physicochemical properties

Methamphetamine exists in powder, crystal or tablet forms. Crystal meth is a physically solid, crystalline form of the drug (methamphetamine hydrochloride), appearing like shreds of glass or clear-white rocks. The crystal form can be vaporised by heating for inhalation by the users. Methamphetamine $(C_{10}H_{15}N)$ and amphetamine $(C_9H_{13}N)$ share a similar chemical structure: one aromatic ring with two carbon side chains. Both drugs contain a methyl group attached to a distal carbon side chain (alpha) of the ethylamine chain. A terminal secondary amine differentiates methamphetamine from amphetamine, which possesses a primary amine. A methyl group attached to the amine functional group compared to hydrogen in amphetamine is responsible for the enhanced lipid

solubility of methamphetamine. A functional amine group in methamphetamine contributes to its base property (PKa = 9.87) [11].

The first synthesis of methamphetamine from ephedrine took place in 1893 by Nagayoshi Nagai, followed by Akira Ogata, who first synthesised the crystallised form in 1919 [4, 11]. Methamphetamine has two isomeric forms: (R) (-) or l-isomer and (S) (+) or d-stereoisomer (Figure 1). The type of yield is usually determined by the methods used. Using phenyl-2-propanone (P2P) as a precursor, racemic methamphetamine can be produced by Leuckart or reductive amination methods. In the former technique, P2P reacts with formamides, producing intermediate amide, which can then react with hydrochloric acid to form racemic methamphetamine. The latter method involves an initial reaction of P2P with methylamine, aluminium and mercury chloride, forming an intermediate phenylacetone-N-methyl imine before further producing racemic methamphetamine via a reduction process. In addition, pure (S) (+) methamphetamine can be produced by reducing pseudoephedrine or ephedrine using chloride derivatives or hydroiodic acid and red phosphorus, respectively [11-13].

Pharmacokinetics

Route of administration and absorption

Numerous methods of methamphetamine administration have been recorded in the literature, including nasal (inhalation or insufflation), oral, intravenous, anal and vaginal routes [14]. The extent of methamphetamine absorption varies, depending on the routes of administration. The earliest methamphetamine serum levels detected after oral ingestion are 1.5±0.8 hours (0.3–2.0) and 1.1±0.5 hours (0.5–2.0) for low doses (10 mg daily four times a week) and high doses (20 mg daily four times in a week), respectively [15]. Meanwhile, the bioavailability for intranasal, inhalation and oral are 79%, 67%, and 67%, respectively [11, 16].

Distribution

Methamphetamine is distributed to all body parts, including the brain, heart, lungs, liver, stomach, kidneys, eyes, hair and breastmilk [16-19]. A positron emission tomography (PET) study utilising intravenous (S) (+) methamphetamine revealed that the lungs had the fastest uptake (55 seconds), one of the highest concentration peaks (22%), and the fastest clearance (7 minutes). Meanwhile, the brain had an intermediate uptake (nine minutes), one of the lowest concentration peaks (10%), and one of the slowest clearances (>75 minutes). The slow removal from the

brain explains the neurotoxicity effects due to prolonged methamphetamine exposure. The heart had the fastest uptake (60 seconds), one of the lowest concentration peaks (about 3%), and intermediate clearance (16 minutes). The liver displayed the slowest uptake rate, taking 30 minutes, comparable to the stomach, and reached one of the highest concentration peaks at 23%, with clearance occurring after 75 minutes. The primary excretory organ, the kidney, required about three minutes for the uptake and had a high concentration peak based on weight (7%) and intermediate clearance (22 minutes) [19].

Following the intranasal administration of the powder type, the peak plasma concentration (t_{max}) was achieved at 2.7 hours post-exposure, reaching about 113±23.1 μg/L. The smoking route demonstrated a similar t_{max} of 2.5 hours, but the peak concentration (C_{max}) achieved was about half $(50.9\pm24.7 \text{ µg/L})$ of that of the intranasal route [20]. The distribution of methamphetamine for the oral route depends on the dose administered but typically has a lower t_{max} and C_{max} than the other two routes. For repeated low doses (10 mg daily four times a week), the drug serum levels reached a peak at 5.4±2.5 hours (2.0–8.0) with maximum concentrations of 20.2±6.4 μg/L (14.5–33.8). The high doses (20 mg daily four times a week) peaked at 7.5+3.4 hours (2.0–11.5) with highest concentrations of 32.4 ± 7.7 μ g/L (26.2– 44.3). The volume of distribution after low and high doses of oral methamphetamine were 1.7±0.9 and 2.6 ± 2.1 L/kg, respectively [15].

Metabolism

The metabolism of methamphetamine occurs primarily in the liver. The primary cytochrome P450 (CYP) isoenzyme involved in phase I is CYP2D6, which metabolises methamphetamine via aromatic hydroxylation and N-demethylation (Figure 1). The former reaction produces p-hydroxymethamphetamine, which can be further conjugated with either glucuronide or sulphate in a phase II reaction before the urinary excretion. The N-demethylation process yields amphetamine, which can be further metabolised by CYP2D6 to form p-hydroxyamphetamine or converted into norephedrine via beta-hydroxylation [11, 21, 22].

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Figure 1. Chemical structures of methamphetamine and its products of metabolism. A. (S) (+) methamphetamine. B. ® (-) methamphetamine. C. Metabolic pathway of methamphetamine Adapted from Abbruscato and Trippier [11].

Genetic polymorphisms of metabolising enzymes contribute to differing metabolism extent and toxicity of methamphetamine. For CYP2D6, the number and functionality of alleles determine the phenotypes for methamphetamine metabolism. Compared to CYP2D6**10*, CYP2D6**1* metabolises (+) and (-) enantiomers of methamphetamine more efficiently, particularly via N-demethylation than aromatic hydroxylation [23]. In a study on the Japanese population, methamphetamine abusers with intermediate metaboliser (IM*-*4/*10, *5/*10, *5/*14B, *10/*10, *10/*18, *10/*36, *10/*10xn*) phenotype carrying two or one reduced (**10, *14B, *10xn*) and one non-functional allele (**4, *5, *14A, *18, *36*) predicted lower metabolism rate than extensive metaboliser (EM) phenotype carrying one or two functional alleles (**1, *1xn, *2, *2xn*) based on urine and bone marrow autopsy samples [24].

Excretion

The main excretion route of methamphetamine is urine, eliminating approximately 37–54% of unchanged methamphetamine [18, 25]. Active tubular transport plays a pivotal role in the renal excretion of methamphetamine, other than glomerular filtration clearance. The organic cation transporter (OCT)-2 and multidrug and toxin extrusion (MATE) transporters 1/2 transport methamphetamine and its metabolite amphetamine, while OCT1–3 and MATE-1 are responsible for transporting another metabolite p-hydroxymethamphetamine. The urine pH is an essential determinant for urinary excretion, with acidic pH promoting the elimination of methamphetamine and amphetamine. The reason for increased excretion is attributable to the weak base properties of methamphetamine and amphetamine, which become more ionised in the acidic environment, thus decreasing partition-mediated reabsorption. The pH-dependent MATE transporters further enhance its excretion in acidic urine pH [25].

Drug half-life and clearance are influenced by administration routes and dosage levels. Intranasal and smoking routes have the same half-life of 10.7 hours, while a 15-minute intravenous infusion of 10 mg methamphetamine shows a slightly longer half-life of 11.4 hours [20]. The half-life of low oral doses (10 mg daily four times a week) is 9.3+3.7 hours (2.1–14.0), and 11.1+7.2 hours (2.2+21.2) for the high dose group (20 mg daily four times a week). The clearance of low and high doses of ingested methamphetamine are 13.2+6.5 (3.4–19.6) and 12.9+7.5 (4.5–24.7) L/h, respectively [15].

Interestingly, methamphetamine also diffuses into breast milk, which exposes danger to infants if the milk containing methamphetamine is consumed. The methamphetamine half-lives in the breast milk of two patients who smoked methamphetamine were 11.3 and 30.3 hours, which became undetectable after 100 hours of methamphetamine use. For injected methamphetamine, the predicted half-lives for two patients were 13.6 and 43.0 hours [18]. Given the fact that urinary methamphetamine has an additional 30– 75-hour detection window, a negative urine test for at least 24 hours can be utilised as a safety indicator for reinitiating breastfeeding [17].

Hair is another area that accumulates methamphetamine, which is eliminated from the body once the hair sheds from the scalp. The knowledge of the elimination of methamphetamine in hair is useful in detecting and monitoring drug intake in the long term. In the study by Wang and colleagues, the half-life of (S) $(+)$ - and (R) $(-)$ methamphetamine ranged from 0.5–1.0 (0.6±0.1) and 0.4–0.9 (0.6±0.2) months, respectively. The former isomer predominated in hair samples, potentially due to higher content than the latter in abused methamphetamine [16]. Since a fraction of methamphetamine is still detected at four months, they proposed a 6-month cut-off period to indicate abstinence from the drug after six months.

Pharmacodynamic

One essential reason for methamphetamine abuse is the enhancement of subjective feelings of well-being, alertness and energy. Harris and colleagues [20] in their study reported increments in numerous subjective feelings, including good drug effects, drug liking, feeling high and intoxication for intranasal and smoked amphetamine with concurrent intravenous methamphetamine infusion. Notably, the intranasal route effects peaked earlier in good drug effect, feeling high and intoxication at 15 minutes. Smoked amphetamine reached the peak later at various time points: 15, 30, and 45 minutes for good drug effects, feeling high and intoxication, respectively. The earlier drug peak and earlier methamphetamine effects (good drug effects, feeling high, and intoxication) in pharmacodynamic assessment, along with higher t_{max} and C_{max} in pharmacokinetic evaluations for intranasally than smoking might partly explain the higher addictive potential of the former route [20, 26]. Bad drug effects were reported in both groups, about less than 20% at peak [20]. Another study discovered that acute administration of oral methamphetamine increased spatial and numeric working memory, vigilance, reaction time, and stimulant effects [27].

The enhancement of the subjective well-being of methamphetamine is attributable to its effects on monoamine neurotransmission, particularly dopamine. Other than subjective feelings, acute administration of methamphetamine produces numerous physiologic effects, such as increased blood pressure and heart rate. Harris *et al.* (2003) reported that systolic blood pressure peaked at 141 ± 13 mm Hg from 121 ± 10 mm Hg in the intranasal methamphetamine group, while the smoked methamphetamine group had a peak of 137 + 11 mm Hg with a similar baseline. For diastolic blood pressure, the pressure peaked at 86 + 7 mm Hg from 76 ± 7 mm Hg and 83 ± 8 mm Hg from 73 ± 10 mm Hg for intranasal and smoked methamphetamine groups, respectively. The heart rate was also increased in both routes of administration. The intranasal group demonstrated an increase from 73 ± 9 beats/minute to 94 ± 13 beats/minute, while the smoked amphetamine group reported an increase from 76 ± 10 beats/minute to 106 ± 19 beats/minute [20]. Oral methamphetamine also has similar stimulating effects on heart rate and systolic and diastolic blood pressure [27].

Toxicity effects

Neurotoxicity

Enhancement of monoamine release into the synapse

The high lipophilic property of methamphetamine allows a simple diffusion of drug across the blood-brain barrier. In the brain, methamphetamine can diffuse into the nerve

terminals via monoaminergic transporters owing to its shared chemical structure with monoamines (dopamine, serotonin, and norepinephrine). Given its weak base characteristics, methamphetamine accumulates and alkalinises intracellular environment (Figure 2). The intracellular alkalinisation affects the uptake and accumulation of neurotransmitters in storage vesicles, which depends on the pH gradient between the internal and external sides of the vesicles. Methamphetamine can also accumulate in the vesicles. These intracellular effects prevent the uptake of monoamines into the vesicles and cause the release of monoamines into the cytosol (via vesicular monoamine transporter 2 (VMAT2) and synapse (via monoamine transporters) by a reverse transport mechanism. Interestingly, its effect on VMAT2 is not attributable to the inhibition of ATPase but rather the reversal of the transporter. Excessive accumulation of neurotransmitters in the synapse is also attributable to methamphetamine effects on neurotransmitter uptake and metabolism [4, 28, 29]. The presence of high levels of neurotransmitters stimulates various monoamine receptors located in the postsynaptic membrane. Increased dopaminergic neurotransmission is the mechanism for reward, learning and memory. Serotonin regulates impulsivity, stress, mood, learning and memory, whereas norepinephrine promotes arousal and attention, as well as regulates stress, mood, learning, and memory [4].

Figure 2. The neurotoxic effects of methamphetamine (denim blue). Methamphetamine causes a reverse transport mechanism of dopamine transporter and VMAT2, increasing the levels of dopamine (and other monoamine neurotransmitters) in the synapse. Subsequent dopamine deficit results from reduced VMAT2 and dopamine uptake into the

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vesicle, dopamine transporters and dopamine synthesis secondary to reduced tyrosine hydroxylase production and activity. Also, methamphetamine activates astrocytes and microglia, further contributing to neuroinflammation. Differential inhibitory and stimulatory effects of methamphetamine on various dopamine receptor subtypes might explain its effects on locomotion, learning, memory and thermoregulation. The positive symbols represent the stimulatory effects of methamphetamine, while the negative symbols represent the inhibitory effects. Abbreviations: DM: dorsomedial; DL: dorsolateral; DR: dopamine receptor; MDA: malondialdehyde (orange); ROS: reactive oxygen species (red); TH: tyrosine hydroxylase; VMAT2: vesicular monoamine transporter 2. Additional colour coding: dopamine (mint green), L-dopa (fern green), quinone (pine green), pro-inflammatory cytokines (mulberry pink), tyrosine (asparagus green).

Effects on dopaminergic neurotransmission in numerous brain regions

Dose and duration are two determining factors contributing to methamphetamine effects on dopaminergic neurotransmission. At a low concentration, acute administration of methamphetamine induces dopaminergic neuronal firing via dopamine transporter-mediated excitation and synaptic transmission. In contrast, methamphetamine reduces cell firing at high concentrations via dopamine D2 auto-receptor activation, evidenced by the reduction of dopamine inhibitory postsynaptic current amplitudes [30]. Regarding the duration, acute binge-like dosing produces an increase in dopamine levels, while prolonged administration produces the opposite effects. A study utilising a rodent model reported a significant elevation in dopamine levels within the frontal cortex (at 2- and 24-hours post-treatment) and amygdala (at 24 hours post-treatment) following one-day binge-like dosing of methamphetamine. Interestingly, a decrease in 3,4-dihydroxyphenylacetic acid (DOPAC) levels at 24- and 48-hour post-treatment was observed in the striatum, indicating a reduction in dopamine turnover [31].

The dopaminergic deficit in the striatum is attributable to reduced VMAT2 and dopamine uptake into the vesicle, dopamine transporters and dopamine synthesis secondary to reduced tyrosine hydroxylase production and activity (Figure 2) [32]. Methamphetamine enhances dopamine release when administered acutely. Upon secretion, dopamine binds to its receptors, initiating a cascade of diverse cellular, molecular and behavioural changes. All dopamine receptors (D1R–D5R) play various roles in drug addiction and relapse, including methamphetamine-induced behavioural sensitisation. Interestingly, the dopamine D3 receptor (D3R) exerts a positive regulatory effect on methamphetamineinduced locomotor function but not the basal spontaneous motor activity which is regulated by other dopamine receptors [33].

Further, binge methamphetamine use decreases the striatal dopamine transporter (DAT) (Figure 2) uptake in dorsolateral and dorsomedial areas, which coincides with the increase in body temperature [34, 35]. Repeated injections of a specific dopamine D3R antagonist, PG01037, mitigate the effects of methamphetamine on DAT and body temperature.

These findings proposed the role of D3R in methamphetamine-induced hyperthermia, well-documented phenomenon contributing to neuronal damage [34, 35]. Interestingly, similar effects of PG01037 on the striatal DAT uptake are not replicable in warmer and more ambient conditions, suggesting a complex interaction between intrinsic and environmental factors [34]. Another methamphetamine-induced neurotoxic effect in the striatum is impaired long-term potentiation in the dorsomedial region mediated by D1R. The attenuation of long-term potentiation might explain the neurotoxic effects on memory and learning [34].

Nucleus accumbens (NAc) is another area affected by methamphetamine use, evidenced by reduced expressions of D1R and D2R [33]. D1R- and D2R-medium spiny neurons (MSNs) are the primary neurons found in the NAc. The activation of the former neurons facilitates drug-seeking behaviour, while the latter causes the opposite effects. D2R-MSNs project to the ventral pallidum, which is connected to the ventral mesencephalon. The activation of the D2R-MSNs-ventral pallidum-ventral mesencephalon pathway inhibits the thalamus and, thus, drug-seeking behaviour (indirect pathway). The updated concept informed that D2R-MSNs can also directly cause inhibition through the D2R-MSNs-ventral pallidum-thalamus pathway [36]. The NAc-specific D2R-knockdown mice have been reported to exhibit reduced acute methamphetamine-mediated hyperlocomotion and attenuated the extent of locomotor sensitisation upon repeated methamphetamine dose. These findings further support the role of D2R in regulating motivated and addictive behaviours [37].

Oxidative stress and inflammatory response

Methamphetamine induces neurotoxicity in various brain areas attributable to impaired dopaminergic and serotonergic circuits, neuronal apoptosis and neuroinflammation induced by astrocytes and microglia activations (Figure 2). Several studies reported that activation of neuronal nitric oxide synthase (NOS) promoted overproduction of nitric oxide radicals (such as peroxynitrite), contributing to oxidative stress [38]. Other than dopamine, methamphetamine also induces glutamate release. The binding of glutamate to NMDA receptors causes calcium influx and activation of neuronal NOS. Activated neuronal NOS catalyses the generation of nitric oxide which can react with superoxide, forming peroxynitrite [38]. Several inhibitors of nitric oxide synthase, such as 7-nitroindazole (7-NI), *N*G-nitro-L-arginine and *N*G-nitro-L-arginine methyl ester have been reported to attenuate methamphetamine-induced dopaminergic neurotoxicity, suggesting the essential role of nitric oxide in the pathophysiology of methamphetamine neurotoxicity [38, 39]. Interestingly, the results from clinical studies on the central effects of methamphetamine on glutathione, a primary antioxidant, are equivocal. One study reported that methamphetamine use disorder patients had a significantly higher level of glutathione (GSH) in the dorsolateral prefrontal cortex compared to healthy controls [40]. In contrast, another study found negligible differences between the two groups despite higher peripheral immunoinflammatory markers in patients than in healthy controls [41].

Cardiotoxicity

Many studies have reported the cardiotoxicity effects of methamphetamine, leading to numerous diseases, such as coronary heart disease, arrhythmias, myocardial infarction, cardiomyopathy and heart failure [10]. Further, methamphetamine use is not only associated with various heart-related illnesses but also accelerates cardiovascular diseases (CVD) a few years earlier than the typical CVD onset [42-44]. Compared to CVD patients without a history of methamphetamine use, those using methamphetamine had worse dysfunction in various parameters, including left ventricular (LV) ejection fraction (EF), LV mass index, end-diastolic volume (EDV) index and myocardial perfusion [43-45]. The risk factors for methamphetamine-associated CVD development include hypertension, chronic kidney disease, diabetes and smoking [10]. Mechanisms of methamphetamine-induced cardiotoxicity are described in detail below.

Oxidative stress

Evidence from animal studies found that methamphetamine can induce oxidative stress in the cardiovascular system as evidenced by the elevation of a reactive oxygen species (ROS) superoxide and a lipid peroxidation product, malondialdehyde (MDA) [46, 47]. NADPH oxidases (NOX) are primary ROS generators, and methamphetamine has been shown to stimulate catalytic subunits (p47phox and gp91phox) of NOX2 (Figure 3) [47]. An inhibitory effect of methamphetamine on cardiomyocyte mitochondrial electron transport chain via suppression of the Sigmar1/cyclic adenosine 3,5-monophosphate (cAMP) response element-binding protein

(CREB)/mitochondrial fission-1 protein (Fis1) pathway is also partly attributable to oxidative stress (Figure 3) [48].

Oxidative stress leads to the depletion of various endogenous antioxidants, including superoxide dismutase and hydrogen sulphide (H2S). The reduction of H2S is partly attributable to reduced production and excessive use to fight against oxidative stress. The former cause is potentially due to reduced cystathionine gamma-lyase (CSE) expression, one of the primary H₂S-producing enzymes [47]. H₂S can neutralise various ROS, including hydrogen peroxide, superoxide, and peroxynitrite. Further, H2S upregulates nuclear factor-erythroid factor 2-related factor 2 (Nrf2)-signalling pathway by interacting with the Kelch-like ECH-associating protein 1 (Keap-1), which is attached to Nrf2 in the cytoplasm. The release of Nrf2 allows its translocation into the nucleus to activate various promoter regions responsible for antioxidant system activation (Figure 3) [49].

Interestingly, methamphetamine exposure promotes the expression of Nrf2, which in turn induces the heme-oxygenase-1 (HO-1) promoter region. The activation of the Nrf2/HO-1 pathway exerts cytoprotective effects, attenuating oxidative stress by promoting antioxidant effects [46]. Besides oxidative stress, methamphetamine also promotes apoptosis via the Bax/Bcl2/caspase-3 pathway (Figure 3) [46].

Inflammatory response

Regarding inflammation, cardiac transcriptional analysis in the animal models reported upregulations of various inflammation-related genes, including Bpifa/Plunc, RegIIIg and Scgb3a2 [50]. Together, oxidative stress, inflammation and apoptosis effects of methamphetamine lead to structural and functional abnormalities.

Fibrotic, vascular, structural and functional effects

One of the most common structural changes induced by long-term methamphetamine administration is cardiac fibrosis, which is observed in various animal models (zebrafish and mice) and humans [46-48, 51, 52]. Fibrotic histopathological changes observed in preclinical and clinical samples include excessive collagen deposition and elevated fibrotic markers of periostin and smooth muscle actin [48, 50]. Methamphetamine-induced fibrotic properties are partly attributable to its ability to stimulate G-protein-coupled receptor (GPCR) trace amine-associated receptor 1 (TAAR1), which utilises a cAMP second messenger system in cardiomyocytes. Elevated cAMP levels then promote lysyl oxidase (LO) production, an essential enzyme that forms crosslinking between collagens in the extracellular matrix (Figure 3) [51]. Other histopathological changes include nucleolysis, nuclear fission and hyperfused and extended mitochondrial networks [46, 48]. Clinical findings reported in the literature include intracardiac thrombus and elevated LV mass [52]. Histopathological findings of autopsy cases of methamphetamine revealed various cardiac pathologies, such as myocardial fibre hypertrophy, myocardial infarction, interstitial and perivascular fibrosis, atherosclerosis (mild to severe), congestion and focal degeneration/necrosis [53].

Molecular and structural changes observed are parallel to elevated biochemical markers of cardiac injury, including creatine kinase MB (CKMB), lactate dehydrogenase (LDH) and cardiac troponin I (cTnI) [46]. These pathological changes can cause disturbances in cardiac function. Cardiac dysfunction associated with methamphetamine use varies in pre-clinical models, ranging from reduced LVEF,

reduced fractional shortening (FS) and increased LV internal diameter (LVID) systole and diastole [47]. In clinical studies, similar cardiac dysfunctions were reported, including elevated end-systolic/diastolic volumes, reductions in ventricular EF and reduced coronary microcirculation [43, 45, 52].

Sex-specific effects

Methamphetamine causes sensitisation effects as early as nine days in a rodent model characterised by elevated heart rate and EF but reduced stroke volume [54]. Although methamphetamine-cardiotoxicity effects are sex-dependent, various pre-clinical models reported inconsistent results. Marcinko *et al.* [50] reported that only male mice developed dilated cardiomyopathy at five months of methamphetamine exposure, characterised by reduced EF and FS but elevated LVID at systole and diastole. Interestingly, the mortality rate for male mice was more than 50% compared to none in female mice.

Figure 3. Molecular-signalling pathways activated by methamphetamine contribute to cardiotoxicity. Methamphetamine induces ROS formation by stimulating NADPH oxidases 2 to generate ROS. ROS then causes lipid peroxidation and activates S1R/CREB/Fis1 and Nrf2/HO1 pathways. Also, methamphetamine promotes cell apoptosis by stimulating the Bax/Bcl2/caspase-3 pathways. Fibrotic-inducing properties of methamphetamine are mediated through the TAAR1/cAMP/LO pathway. The positive symbols represent the stimulatory effects of methamphetamine (denim blue), whereas the negative symbols represent the inhibitory effects. Abbreviations and colour coding: cAMP: cyclic adenosine 3,5-monophosphate (Uranian blue); CSE: cystathionine gamma-lyase; CREB: cAMP response element-binding protein; CytC: cytochrome C (baby blue); Fis1: mitochondrial fission-1 protein; HO-1: heme-oxygenase-1; H2S: hydrogen sulfide (pastel purple); keap1: Kelch-like ECH-associating protein 1; LO: lysyl oxidase; MDA: malondialdehyde (orange); NOX: NADPH oxidases; Nrf2: nuclear factor-erythroid factor 2-related factor 2; ROS: reactive oxygen species (red); S1R: Sigmar 1, TAAR1: trace amine-associated receptor 1. Additional colour coding: Caspase-9 (light orange), caspase-3 (salmon pink).

Contradictory, a 10-day methamphetamine exposure caused prominent changes in circadian clock gene expression in only female rats, upregulating *Per2* and *Per3* while downregulating *BMal1*, *Npas2 and Clock* genes. Circadian clock genes play a pivotal role in the diurnal regulation of blood pressure, heart rate, and heart metabolism. Altered expression of these genes can increase the risk of cardiovascular disease. Interestingly, one-month abstinence from methamphetamine reversed the altered circadian clock genes [55]. Another study utilising the same treatment protocol in addition to an ischemic-reperfusion period also displayed a higher susceptibility of female rats to methamphetamine toxicity effects. More extensive post-ischemic infarct size and contractility dysfunction at systolic and diastolic were reported in only female rats after a 10-day treatment as well as a 10-day-and-one-month abstinence [56]. Different treatment protocols in dosing (constant vs increasing), duration (ten days vs five months), frequency (daily vs week and administration route (subcutaneous vs intraperitoneal) might partly explain the differing outcomes between studies.

Electrophysiological effects

A study using a zebrafish model exposed to methamphetamine demonstrated unexpected outcomes in electrophysiological aspects [51]. Heart rate was reported to reduce [51] in contrast to other preclinical and clinical findings [45, 54, 57, 58]. The possible explanation is that elevated blood pressure caused by methamphetamine stimulates a baroreceptor reflex, leading to reduced heart rate. An increase in heart rate variation (HRV) during the first week of methamphetamine exposure might also be attributed to a similar mechanism. Interestingly, Zhang and colleagues [51] found a reduction in the HRV of zebrafish exposed to methamphetamine at week two, possibly due to persistent inflammation and cardiac injury. Another interesting finding is a reduced QTc interval, a contradictory phenomenon observed in clinical cases. The possible reason for this condition is attributable to early changes caused by methamphetamine before overt cardiomyopathy is established [51].

Cardiovascular changes following abstinence

Regarding cardiovascular effects following abstinence, a preclinical model of rhesus macaques on long-term methamphetamine (3.6–8.6 years) provided some insight into the dynamic changes of the heart. In the study, researchers found tolerance effects of methamphetamine on diastolic pressure (day 1) and systolic pressure (day 1 and week 1) post-abstinence

and sensitisation effects on diastolic pressure (week 12) and heart rate (week 26). Other functional effects included reductions in EF and cardiac output (CO). Interestingly, the abnormal physiological measures returned to control values in one year, indicating the reversibility of the methamphetamine-induced cardiac effects [58]. In a clinical population, a complete cessation of methamphetamine improved EF and heart failure-related hospital admission in heart failure patients with reduced EF [44].

Effects of prenatal exposure

Another essential area is the prenatal exposure effect of methamphetamine on the heart in adults. In the pre-clinical model, methamphetamine exposure during the first half, second half, or whole pregnancy period increased the infarct size following an episode of ischemic-reperfusion injury in adult female rats. No significant differences were reported between treatment exposures, indicating that full abstinence is required to avoid the cardiac sensitisation effect of methamphetamine. In contrast, the authors reported no significant effects of prenatal methamphetamine exposure on functional parameters, including heart rate, developed pressure, +dP/dT and -dP/dT [57].

Other effects of prenatal methamphetamine exposure in an adult animal model are the long-term epigenetic changes. Females are more susceptible to cardiac epigenetic changes. Dague and colleagues [59] in their study reported that female adult rats exposed to methamphetamine exposure during a prenatal period exhibited two times higher non-similar gene expressions. Two primary changes were observed in dimethylarginine dimethylaminohydrolase-2 (DDAH2) and 3-hydroxybutyrate dehydrogenase 1 (BDH1). Interestingly, the protein expression of BDH1 was also reduced in adult rats of both sexes. BDH1 is particularly useful as an energy production using ketones in pathological conditions, including cardiac hypertrophy and heart failure. Reduced expression is associated with cardiac dysfunctions [59]. In contrast, although reduced gene expression of DDAH2 was observed in both sexes, only female rats reported a decrease in protein expression, suggesting a possible compensatory mechanism in male rats. DDAH2 degrades asymmetric dimethylarginine which is a nitric oxide synthase inhibitor. The effect of methamphetamine on DDAH2 expression can potentially cause asymmetric dimethylarginine accumulation and vascular dysfunction [59].

Hepatotoxicity

The liver is among the most susceptible organs to methamphetamine toxicity. The extent of methamphetamine-induced hepatotoxicity ranges from mild liver injury to severe fulminant hepatic impairment. Pathological changes of hepatic complications include fibrosis, necrosis, ballooning degeneration in centrilobular zones, hepatomegaly and hepatitis. Further, methamphetamine's effect on the liver is associated with other health issues, such as cognitive deficits and psychiatric symptoms [60].

The liver hosts several transaminases essential for synthesising and degrading amino acids and converting energy-storing compounds. The levels of transaminases are usually low but increase in liver injury, secondary to increased cell membrane permeability, allowing intracellular enzymes to escape into the blood. Numerous preclinical studies reported that methamphetamine exposure led to increased blood levels of alanine transaminase (ALT), aspartate transaminase (AST), alkaline phosphatase (ALP) and ammonia compared to the control group [61-66]. Also, patients with methamphetamine use disorder had significantly higher ALT and AST levels compared to healthy controls [67]. The biochemical changes indicative of liver injury are parallel to histopathological findings observed in methamphetamine users (Eskandari *et al.* 2014; Wang *et al.* 2017). These changes include nuclear abnormalities (shrinkage, karyopyknosis and nuclear migration), cytoplasm damage (loosening, vacuolar degeneration, microvesicular lipid and increased glycogen), cellular changes (hydropic change and hypertrophy), mitochondrial aggregation, and inflammatory cell infiltration in the portal and lobular areas [61-64, 66, 68, 69]. In contrast, Azizi *et al.* (2023) [70] reported vascular degeneration and congestion not only in the exposed group but also in the control group. The inconsistent findings might be attributable to different study designs, with various rodent types and dose and duration of methamphetamine exposure.

The precise mechanisms underlying methamphetamine-induced liver damage remain unclear. Potential factors contributing to methamphetamine toxicity are hyperthermia, disruption of the CYP1A2 metabolic pathway, oxidation of biogenic amines, hyperammonaemia, mitochondrial impairment, apoptosis, inhibition of cell division, elevated neurotransmitter efflux and the impairment of bile acid homeostasis by gut microbes [64, 66, 71, 72]. The following subsections further discuss on the mechanism of methamphetamine-induced hepatotoxicity.

Oxidative stress

ROS formation induced by methamphetamine elicits pathological downstream events, such as lipid peroxidation and antioxidant depletion (glutathione peroxidase (GPx), superoxide dismutase (SOD)), promoting oxidative stress [63].

Inflammatory response

Oxidative stress activates a toll-like receptor (TLR)4/MyD88/TRAF6 pathway, promoting inflammation [61, 64, 68]. TLRs play a key role in regulating innate and adaptive immunity, pathogen detection and inflammatory signalling (Lai *et al.*, 2015; Xie *et al.*, 2018). Activated adaptor proteins MyD88 and TRAF6 by TLR4 trigger downstream inflammation (Xie *et al.*, 2018), promoting the expression of various inflammatory markers, including TNF-α, IL-6, IL-1β, and IL-18 [61, 64, 66, 68]. Methamphetamine-induced cell cycle arrest and apoptosis in hepatocytes may result from the disruption of essential cellular bioprocesses [65].

In a clinical population, a study examining the association of methamphetamine use with liver pathology in autopsy samples of 527 (413 cases and 114 controls) reported similar occurrences of fatty liver and cirrhosis in both groups. Hepatitis and infiltration of inflammatory cells (lymphocytes and plasma cells) in the portal triads (areas of the liver that contain bile ducts, veins and arteries) were significantly higher in cases but no significant differences were observed between methamphetamine and other intravenous drug users [73]. The exact reason for negligible differences between groups is unknown, but potentially due to other factors such as shared pathological changes in intravenous drug users regardless of the type of drugs and inclusion criteria for the control group (drug-free (based on the post-mortem toxicological exam) trauma victim).

Vascular effects

A case report from another autopsy finding of a case of a 35-year-old man due to methamphetamine overdose revealed multi-organ ischemic insults, secondary to methamphetamine-induced vasoconstriction in the liver, heart (left ventricle) and brain (cerebellum). Histopathological examination revealed necrotic hamartoma in the liver and pancreas and extensive fibrosis and thrombi in the liver. Also, fat necrosis with infiltration of acute inflammatory cells, oedema and fibrinous exudate were observed in the pancreas [74].

Bile acid profile derangements

The effect of methamphetamine on the liver can disrupt bile acid production, leading to decreased bile acid production and deficiency in secondary bile acids. Ma and colleagues [67] conducted a study to investigate the association of various parameters concerning lipid profile, bile acids and psychiatric psychopathology in methamphetamine use disorder patients during the withdrawal period. The effects of methamphetamine were the most prominent at a 3-month withdrawal point when patients showed the greatest derangement in transaminases (AST and ALT), lipid parameter triglyceride and psychiatric symptomatology of depression and anxiety. Interestingly, the bile acid profiles, including total, primary (CA and CDCA) and secondary bile acids (HCA and UDCA), were significantly lower in patients than in HC. The findings suggest the role of bile acids as potential mediators of liver injury and psychiatric comorbidities in methamphetamine withdrawal, which could be linked to disease progression, potentially through the crosstalk between the liver and brain [67]. The association between liver injury and mental health problems or cognitive impairment has long been established, particularly in patients with non-alcoholic fatty liver disease (NAFLD). NAFLD patients possess a higher risk of cognitive impairments (memory, visual attention and coding ability) and depression [75, 76].

Ocular toxicity

A range of ophthalmologic complications in methamphetamine use disorder patients has been reported in the literature. The methamphetamine ophthalmologic-related sequela includes keratitis, endophthalmitis, vision loss, retinopathy, intra-retinal haemorrhages, retinal vasculitis, amaurosis fugax, optic neuropathy and acute closed-angle glaucoma [77-81].

Local or direct effects

Keratitis is the commonest case reported, partly attributable to chemical irritation, vasoconstriction (secondary to catecholamine release), decreased blink reflex, scratching and self-epilation exacerbations, increased infection susceptibility and elevated pain threshold. Direct contact with methamphetamine vapour can irritate the eyes (Figure 4). Given the close location between the eyes and nasopharynx, inhaled methamphetamine can also cause local toxic effects. Accidental introduction of methamphetamine and its diluting agents when the contaminated hand touch the eyes is another possible cause. Diluting agents like bicarbonate can cause chemical burns [82].

Figure 4. Ophthalmologic effects of methamphetamine. Methamphetamine can cause direct effects (chemical burn), reduced sweeping reflex (may increase the risk of infection), oxidative stress, inflammation (↑ TNF-α), vasoconstriction (may increase the risk of retinopathy), neovascularisation (secondary to ischemic-induced increased in HIF-1α/ VEGF), and dysregulation of extracellular matrix proteins (through MMPs). The positive symbols (and colour) represent the stimulatory effects of methamphetamine (denim blue), while the negative symbols represent the inhibitory effects. Abbreviations: HIF-1α: hypoxia-inducible factor-α (fern green); MMP: matrix metalloproteinases; NE: norepinephrine (baby blue); PECAM-1: platelet endothelial cell adhesion molecule-1; TNF-α: tumor necrosis factor-α (purple); VEGF: vascular endothelial growth factor (mint green). Additional colour coding: asparagus green (diluting agent), lace pink (MMP-2), mulberry pink (MMP-9), and orange (MMP-14).

Suppression of weeping and blinking reflexes

Besides, anaesthetic agents in diluting solution (including lidocaine) can suppress weeping and blinking reflexes, reducing corneal protective ability while increasing infection risks. Infective agents such as gram-positive (Staphylococcus aureus, Streptococcus spp. and Propionibacterium acnes) and negative (Pseudomonas and Capnocytophaga) bacteria and fungi (Candida albicans) have been isolated from the sensitivity tests of keratitis cases [77, 83, 84]. Although infections are commonly found, keratitis with no evidence of infection has also been previously reported [82]. The agent can also disturb the re-epithelialisation via its anti-mitotic effects and disruption in cellular respiration [83, 85].

Vascular effects

Vascular effects via sympathomimetic effects of methamphetamine can cause vasospasm and vasoconstriction, leading to ischemic retinopathy and ischemic optic neuropathy. Increased plasma levels of norepinephrine, a potent vasoconstrictor, have been reported in methamphetamine animal models (Figure 4) [86]. In the eyes, the short posterior ciliary arteries are the primary blood supply to the optic nerve head (optic disk). The ischemic episodes can cause ischemic optic neuropathy [79]. Ischemic retinopathy cases have been reported in methamphetamine users involving occlusions of the central retinal vein and bilateral branch retinal arteries [87, 88]. Ischemic insults favour the formation of hypoxia-inducible factor 1 α (HIF-1 α), which in turn upregulates the expression of vascular endothelial growth factor (VEGF). Retinal neovascularisation is the consequence of HIF-1α-induced VEGF elevation, as seen in clinical cases and preclinical studies [87, 89].

Extracellular matrix protein dysregulation

Apart from the vasoconstrictive effects of norepinephrine, its excessive levels increase metalloproteinase (MMP)-14, which promotes the activation of MMP-2 and MMP-9, dissolving numerous endothelial surface molecules such as syndecan-1, glypican-1 and platelet endothelial cell adhesion molecule-1 (PECAM-1) (Figure 4) [86].

Oxidative stress and inflammatory response

Excessive norepinephrine promotes a pro-inflammatory state, evidenced by the increase in tumour necrosis factor-alpha (TNF-α) and oxidative stress markers in the retina and blood [90, 91]. The TNF-α cell death-signalling pathway and oxidative stress might partly explain retinal neurodegeneration as shown by thinning retinal nerve fibre layer

thickness (RFNL) and a reduction in Bruch's membrane opening minimum rim width reported in clinical and animal studies [86, 90, 92]. Other pathologies include marked activation of astrocytes and microglia secondary to methamphetamine exposure [93].

Talc deposition

Talc deposition in the macula is another form of retinopathy known as crystalline retinopathy. Kumar, Kaiser [94] reported a case of crystalline retinopathy in a patient who had been snorting methamphetamine for years. Although talc deposition is more common in patients who administer crushed oral drug suspension intravenously, nose or lung are possible routes of delivering talc to retinal circulation [94].

Neural effects

Another pathological change induced by methamphetamine is the abnormality in the sensory receiving area and visual pathway, which might cause a delay in signal delivery due to altered processing in the primary visual area [78].

Renal toxicity

Methamphetamine can cause acute kidney injury (AKI) and chronic renal disease (CKD) [95-98]. Acute manifestations of methamphetamine-induced AKI may include acute tubular necrosis (the most prevalent), tubulointerstitial nephritis and thrombotic microangiopathy [96, 99-102]. Hyperthermia, rhabdomyolysis, tubular obstruction, vasoconstriction and hypovolemia are possible causes of AKI secondary to methamphetamine intoxication [101].

Thermoregulatory effects

The effect of methamphetamine on serotonin and dopamine levels can lead to hyperpyrexia, one of the contributing factors of rhabdomyolysis [34, 35, 100]. As previously stated, D3R stimulation by methamphetamine causes hyperthermia [34, 35]. Hyperthermia is one of the common signs reported in rhabdomyolysis cases, particularly in severe methamphetamine intoxication [96, 102]. Further, elevation of 70 kDa heat shock protein, a protective molecule against heat stress, in myoglobin-positive kidneys of methamphetamine users may support the role of this protein in the pathophysiology of hyperthermia and AKI secondary to methamphetamine intake [102].

Rhabdomyolysis

Rhabdomyolysis is a common manifestation in acute methamphetamine intoxication, accompanied by excessive release of creatine kinase from damaged skeletal muscle. A meta-analysis study involving acute methamphetamine intoxication cases reported that the weighted mean values of five studies were seven times higher than the normal upper limits of creatine kinase [103]. The creatine kinase levels with a cut-off point of 10,000 IU/l in acute poisoning cases may be a useful predictive tool for AKI (84% sensitivity, 69% specificity) [104].

Other than creatine kinase, myoglobin released from injured skeletal muscle is another essential biomarker. Rhabdomyolysis induces the release of myoglobin into the systemic circulation, which is subsequently deposited in the kidney [105]. The iron component (heme protein) of myoglobin can undergo redox cycling between ferric and ferryl forms, initiating lipid peroxidation reactions. Potent renal vasoconstrictors, F_2 -isoprostanes, are one of the lipid peroxidation products responsible for renal vasoconstriction $[106]$. Furthermore, F₂-isoprostanes can stimulate endothelin-1 production (another potent vasoconstrictor) [107]. In the kidney, endothelin-1 is produced primarily by endothelial and tubular cells. Overactivity of the endothelin-1 type A over type B receptors may be responsible for developing renal disease [108]. Interestingly, urinary alkalinisation reduces urinary F2-isoprostane levels and improves renal function, potentially owing to a decreased ferryl myoglobin reactivity [106].

Oxidative stress

Myoglobin also increases other oxidative stress markers, including 8-hydroxy-2' -deoxyguanosine (an oxidative DNA damage marker), 4-hydroxy-2 -nonenal (a product of lipid peroxidation) and superoxide dismutase (a free radical scavenger) in a clinical population [102]. In a preclinical study, Zhang *et al.* reported increased oxidative stress in animals exposed to methamphetamine, evidenced by increased GSH levels and GPx activities, besides reduced levels of MDA and protein carbonyls and activities of catalase [98, 109]. Interestingly, pre-treatment with pharmacological agents with antioxidant properties, such as caffeic acid and *N*-acetylcysteine amide, confer protection against methamphetamine-induced oxidative stress [98, 109, 110]. However, the exact causes (either direct methamphetamine toxicity or other factors, such as myoglobin) cannot be concluded, as the study did not assess the presence of myoglobin casts in the kidney [109, 110]. Two potential pathways activated by methamphetamine use are the Nrf2/HO-1 pathway, which enhances antioxidant effects, and the renal autophagy pathway, which stabilises cellular homeostasis by degrading damaged macromolecules [111].

Oxidative stress and inflammation caused by methamphetamine can induce various pathological changes. Histopathological changes of renal biopsy in methamphetamine intoxication secondary to rhabdomyolysis (confirmed/suspected) include myoglobin-positive granular casts in the distal tubules, detached renal epithelium in the lumen and interstitial oedema of the renal medulla in clinical populations [96, 100, 102]. Interestingly, myoglobin casts are not always detected, with the detection rate ranging from 19–66% in the most prevalent renal toxicity findings (ATN), suggesting that other pathophysiologies might be essential for AKI among methamphetamine abusers [99, 102].

Tubular obstruction

Intratubular myoglobin in the kidney can react with Tamm-Horsfall protein in acidic urine, forming intratubular casts. These casts may block the renal tubular system, contributing to AKI [70, 112]. Hypovolemia, acidosis and ischemia are other contributing factors that can worsen AKI [112].

Volume depletion

Isoardi and colleagues [101] in their prospective observational study reported 90% presentations of AKI in the Emergency Department among
methamphetamine intoxication patients with methamphetamine intoxication patients with abnormal creatinine levels. The incidence of rhabdomyolysis was almost half (44%) among 50 presentations at the Emergency Department [101]. Most AKI presentations were mild in severity and successfully attenuated by crystalloid therapy. The possible reasons for volume depletion-induced AKI in this situation are attributable to concurrent psychomotor agitation and reduced fluid intake [101].

Repeated renal insults

Persistent renal insults secondary to chronic methamphetamine abuse and other associated factors (including malignant hypertension and uncontrolled diabetes) can increase the risk of CKD. Chronic renal pathologies associated with methamphetamine consumption include tubulointerstitial nephritis (active or inactive), diabetic glomerulosclerosis, thrombotic microangiopathy and focal segmental glomerulosclerosis (commonest variant – not otherwise specified) [98, 99].

Baradhi and colleagues [98] reported a case of methamphetamine-induced end-stage renal failure (ESRF) in a patient who had been taking methamphetamine for several years. It was then concluded that continuous methamphetamine use might induce persistent malignant hypertension in this patient, evidenced by high blood pressure (210/124 mm Hg) and multi-end-organ damages, such as left ventricular hypertrophy and ESRF. The renal biopsy of the patient identified multiple renal pathologies, including subacute thrombotic microangiopathic injury, advanced glomerulosclerosis, striking mucoid intimal hyperplasia, cellular crescent formation and tubular atrophy or interstitial fibrosis [98].

Malignant hypertension is a common manifestation in methamphetamine users. A retrospective study involving methamphetamine users referred to the renal unit reported that 96% of them had chronic kidney disease (55% ESRF) and 89% had hypertension (45% malignant hypertension) [97]. Among patients with renal biopsy results, half showed hypertensive changes, while a quarter demonstrated malignant changes. Also, 58% of patients with mesangiocapillary glomerulonephritis had positive IgM and C3 complement [97]. The vasoconstrictive effects of methamphetamine-mediated via α_1 and β_1 receptors may explain the reason for increased systemic vascular resistance and blood pressure [97, 98].

Endothelial toxicity

Endothelial cells are flattened cells that form the inner cellular lining of all blood vessels [113]. The cells are connected by tight junctions and anchored to a continuous basal membrane. The endothelium can be found in most arteries, veins and capillaries of the brain, skin, lungs, heart, kidneys and muscle. The cells form a natural barrier between blood and tissues, controlling the movement of substances and fluid across tissue [114]. Impaired endothelial cell functions can lead to serious health issues [115]. For instance, the blood-brain barrier, a highly selective membrane formed by endothelial cells using tight junctions to prevent large and potentially toxic molecules from entering the brain, can be compromised due to vascular dysfunction in certain conditions [116]. One potential cause of endothelial toxicity is methamphetamine mediated through oxidative stress, inflammation and vasoconstriction.

Oxidative stress and inflammatory response

Oxidative stress induced by methamphetamine in brain endothelial cells can cause impaired blood-brain barrier function. Methamphetamine increases the permeability of the blood-brain barrier *in vivo*. The exposure of brain microvascular endothelial cells (BMVECs) to methamphetamine reduces the expression of cell membrane-associated tight junction proteins, resulting in a reduction in the tightness of BMVEC monolayers. Other pathological changes include increased ROS production, enhanced monocyte migration across the endothelium, altered glucose transporter protein-1 expression, reduced tight junction proteins (occludin and zonula occludens-1) and activation of myosin light chain kinase (MLCK) in BMVECs [117-119]. Further, methamphetamine increases plasma levels of oxidative marker MDA, inflammatory marker C-reactive protein and endothelial injury marker endothelial-derived microparticle [120].

eNOS/NO-mediated transcytosis

Endothelial NOS (eNOS)/NO-mediated fluid-phase transcytosis is among the proposed mechanism for the increased permeability of methamphetamine and lymphocytes at the blood-brain barrier [121]. The BMVECs respond to low concentrations of methamphetamine by a rapid activation of endothelial NOS. Inhibiting this enzyme reduces the methamphetamine-induced blood-brain barrier permeability through caveolar transport without compromising the integrity of tight junctions in preclinical models of *in vitro*, *in vivo* and *ex vivo* [121, 122]. Remarkably, a similar effect is absent at higher methamphetamine concentrations [121, 122].

Endothelial-dependent and independent vasoconstriction

Another proposed mechanism of methamphetamine-induced vasoconstriction is the endothelin-dependent pathway. The synthesis of endothelin-1 (ET-1) by brain endothelial cells is enhanced following methamphetamine exposure. In a study using cultured mouse brain endothelial cell lines, Seo and co-workers reported that methamphetamine exposure caused vasoconstriction, which was prevented by concurrent administration of endothelin receptor antagonists [123]. As previously stated, myoglobin-induced formations of potent vasoconstrictors F2-isoprostanes and ET-1 are among the mechanisms of renal toxicity induced by the use of methamphetamine use [106, 107].

Contradictory, a case-control study involving methamphetamine users and healthy controls reported no significant differences between groups in the flow-mediated dilatation (FMD), a measure of endothelium-dependent vasodilation [124]. However, the researchers found a marked difference between groups in endothelium-independent vasodilation, as measured using exogenous vasodilator nitroglycerin [124]. Smoker status (cigarette) in both groups might confound the effect of methamphetamine on FMD, as smoking can cause FMD impairment. Methamphetamine effects on vascular media smooth muscle may reduce the reactivity of smooth muscle cells towards nitroglycerin, causing reduced dilation in response to nitroglycerin [124]. The altered reactivity of vascular smooth muscle might predispose patients to vascular diseases like stroke. A recent case-control study investigating the risk of cerebral small vessel disease in patients with acute ischemic stroke reported that patients who used methamphetamine had a higher total burden of cerebral small vessel disease and more pathological changes in white matter hyperintensities and lacunes [125].

Activation of apoptosis-induced pathways

Methamphetamine induces apoptosis in endothelial cells through various signalling pathways, including blockade of kappa opioid receptor, inactivation of the protein kinase B (Akt)/mammalian target of rapamycin (mTOR)/p70S6K, upregulation of the extracellular signal-regulated kinase 1/2 (ERK

1/2) [126], as well as stimulation of NADPH-Oxidase-2 (NOX-2) [127] in primary human and rat brain microvascular endothelial cells [127]. Another apoptotic pathway affected by methamphetamine exposure in human umbilical veins and rat cardiac microvascular endothelial cells is Nupr1/Chop/P53/PUMA/Beclin1 [128]. Similarly, activation of the VEGF/PI3K/Akt/eNOS-signalling pathway by methamphetamine can also increase cardiac microvascular permeability [129]. Collectively, the pathways represent potential therapeutic targets for methamphetamine-induced endothelial cell apoptosis in blood-brain barrier impairment and cardiovascular toxicity. Table 1 summarises the studies on methamphetamineassociated toxicity effects.

Table 1. Summary of human and animal studies related to ophthalmologic complications of methamphetamine use.

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↑ elevated/higher ↓ reduced/lower + positive association – negative association

7-NI: 7-nitroindazole; AKI: acute kidney injury; Akt: protein kinase B;ALP: alkaline phosphatase; ALT: alanine transaminase; AST: aspartate transaminase; ATF4: activating transcription factor 4; ATN: acute tubular necrosis; BA: bile acid; Bax: B-cell lymphoma protein 2-associated X; BBB: blood-brain barrier; Bcl-2: B-cell lymphoma protein 2; BDNF: brain-derived neurotrophic factor; BP: blood pressure; CA: d4-cholic acid; cAMP: cyclic adenosine monophosphate; CDCA: chenodeoxycholic acid; CKD: chronic kidney disease; CK-MB: creatine kinase-MB; CMECs; cardiac microvascular endothelial cells; CO: cardiac output; Cr: creatine; CRA: central retinal artery; CSE: cystathionine γ-lyase; cSVD: cerebral small vessel disease; cTnI: cardiac troponin I; CYP: cytochrome P450; CVD: cardiovascular disease; DAB: diaminobenzidine; DA: dopamine; DAT: dopamine transporter; DCA: deoxycholic acid; DHE: dihydroethidium; DLPFC: dorsolateral prefrontal cortex; D-NAME: NG-nitro-D-arginine-methyl ester; DOPAC: 3,4-dihydroxyphenylacetic acid; dp/dt: pressure change; DR: dopamine receptor; EB-Alb: Evans Blue-albumin; ECG: electrocardiogram; EF: ejection fraction; EDV: end-diastolic volume; eNOS: endothelial nitric oxide synthase; ERK: extracellular signal-regulated kinase 1/2; ET-1: endothelin-1; Fis1: mitochondrial fission 1 protein; FS: fractional shortening; FSGS: Focal segmental glomerulosclerosis; GABA: γ-aminobutyric acid; GCL: ganglion cell layer; GFAP: glial fibrillary acidic protein; GLUT-1: glucose transporter protein-1; Glx: glutamine + glutamate; GN: glomerulonephritis; GPx: glutathione peroxidase; GSH: glutathione; GPC: glycerophosphorylcholine; GPx: glutathione peroxidase; HAM-A: Hamilton anxiety rating scale; HAM-D: Hamilton depression rating scale; HBECs: human brain endothelial cells; HBMECs: human brain microvascular endothelial cells; HC: healthy control; HCA: hyocholic acid; H&E: hematoxylin and eosin; HDL: high-density lipoprotein; HIF-1α: hypoxia inducible factor 1-alpha; HO-1: heme oxygenase-1; HR: heart rate; HRa: hazard ratio; H2S: hydrogen sulfide; HUVECs: Human umbilical vein endothelial cells; HVA: homovanillic acid; ICAM: intercellular cell adhesion molecule; IgAN: Immunoglobulin A nephritis; IL: interleukin; INF: interferon; INL: inner nuclear layer; Ins: inositol; IP: intraperitoneal; IPL: inner plexiform layer; IPSC: inhibitory postsynaptic currents; ISON: isosorbide dinitrate; IV: intravenous; LCA: lithocholic acid; LDH: lactate dehydrogenase; LDL: low-density lipoprotein; LGE: late gadolinium enhancement; L-NAME: NG -nitro-L-arginine methyl ester; LV: left ventricle; LVID: left ventricular internal diameter; MCP: monocyte chemoattractant protein; MDA: malondialdehyde; MDC: macrophage-derived chemokine; MIP: macrophage inflammatory protein; MMP: matrix metalloproteinases; MRS: magnetic resonance spectroscopy; MP: microparticles; MPO: myeloperoxidase; MRW: Bruch's membrane opening minimum rim width; mTOR: mammalian target of rapamycin; NAA: *N*-acetyl aspartate; NAAG: *N*-acetylaspartylglutamate; NAc: nucleus accumbens; NACA: *N*-acetylcysteine amide; NF-κB: nuclear factor kappa B; NO: nitrogen oxide; NOARG: N^G-nitro-L-arginine; NOS: nitric oxide synthase; NOX: NADPH oxidase; Nrf2: nuclear factor erythroid 2-related factor; NS: not significant; ONL: outer nuclear layer; PCh: phosphorylcholine; PCr: phosphocreatine; PECAM-1: platelet endothelial cell adhesion molecule-1; PK: penetrating keratoplasty; PI3K: phosphatidylinositol-3-kinase; RFNL: retinal nerve fibre layer thickness; ROS: reactive oxygen species; RPE: retinal pigmen epithelium; RV: right ventricle; SFN: sulforaphane; SIRT: sirtuin; SNP: sodium nitroprusside; SOD: superoxide dismutase; SPF: specific-pathogen-free; SV: stroke volume; TARC: thymus- and activation-regulated chemokine; TG: triglycerides; TLCA: taurolithocholic acid: TLR: Toll-like receptor; TNF-α: tumour necrosis factor alpha; TR: tricuspid regurgitation; TRP: transient receptor potential; TUNEL: terminal deoxynucleotidyl transferase deoxyuridine triphosphate nick end labelling; UDCA: ursodeoxycholic acid; VCAM: vascular cell adhesion molecule; VE: vascular endothelial; VEGF: vascular endothelial growth factor; WT: wild-type; ZO-1: zonula occludens-1.

Conclusion

Methamphetamine is a toxic psychostimulant with various detrimental effects on the brain, heart, liver, kidneys and eyes. Various pharmacokinetic and pharmacodynamic factors may alter the toxicity effect of methamphetamine. Understanding the mechanisms of methamphetamine toxicity is essential to develop treatment strategies to reverse or attenuate the progress of methamphetamine-associated organ damage.

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Competing Interests

The authors have declared that no competing interest exists.

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