Original Article Cognitive impairment and changes of red blood cell components and serum levels of IL-6, IL-18, and L-tryptophan in methamphetamine abusers

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Abstract: The deficit in cognitive function is more concerning in methamphetamine (MA) users. The cognitive deficit was suspected to be the consequence of neuroinflammation-induced neurological dysregulation. In addition, activating the key enzyme in the tryptophan metabolic pathway by pro-inflammatory cytokines results in metabolite toxicity, further generating cognitive impairments. However, the evidence for the role of neuroinflammation and tryptophan metabolites involved in MA-induced cognitive deficit needs more conclusive study. Objectives: This retrospective study aimed to determine blood-inflammatory markers, tryptophan metabolite-related molecules, and cognitive function in MA abusers compared to healthy control (HC) participants. Methods: The cognitive functions were evaluated using Stroop, Go/No-Go, One Back Task (OBT), and Wisconsin Card Sorting Test-64 (WCST-64). Blood samples were analyzed for complete blood count (CBC) analysis, serum inflammatory cytokines interleukin (IL)-6 and IL-18 and tryptophan metabolites. Results: MA group exhibited poor cognitive performance in selective attention, inhibition, working memory, cognitive flexibility, concept formation and processing speed compared to HC. Reduction in red blood cell (RBC) components but induction in white blood cells (WBCs) and IL-6 were observed in MA abusers, which might indicate anemia of (systemic chronic low-grade) inflammation. In addition, the depletion of precursor in the tryptophan metabolic pathway, L-tryptophan was also observed in MA users, which might represent induction in tryptophan metabolites. Conclusion: These findings emphasize that blood biomarkers might be a surrogate marker to predict the role of neuroinflammation and abnormal tryptophan metabolite in MA-induced cognitive impairments.

Keywords: Methamphetamine, cognitive impairments, inflammation, tryptophan metabolites, anemia of inflammation, biomarkers

Introduction

Recent evidence demonstrated that methamphetamine (MA) could induce a deficit in neurological and neuropsychological functions [1, 2]. MA strongly affects the nervous system by enhancing the release of neurotransmitters such as dopamine (DA), serotonin, and norepinephrine [3]. MA administration demonstrates significant effects on the central nervous system (CNS), such as excessive sympathetic nervous system stimulation, neurologic complications, neuropsychiatric complications, and cognitive dysfunctions [4, 5]. Abusers can experience increased cognitive performance in acute low-dose exposure to MA [6]; however, chronic MA users come across with wide range of neuropsychological deficits [7] and cognitive impairments [4, 8-10]. MA-induced cognitive impairments mostly exhibit in the non-equally global aspect of cognition, including attention, inhibition, memory, and executive functions [1118]. Theoretically, MA promotes neurological and neuropsychological dysfunction via its effects of neurotoxicity and neuroinflammation.

MA can induce physiological and structural changes and altered inflammatory reactions associated with poor cognition. A study in MA abusers revealed cognitive deficits in the Stroop task accompanied by right inferior temporal gyrus hyperactivation, white matter hypertrophy, and high plasma levels of inflammatory-associated molecules releasing, including tumor necrosis factor-alpha (TNF-α) and S100 calcium-binding protein β (S100β) [19]. MA could be the critical causality of releasing molecular modulators from the CNS to the peripheral system. Additionally, the more defect in inflammatory function among MA users associates, the more exacerbate the deficit in cognition [20]. Yang and his colleagues [21] have also reported that elevated serum levels of IL-6, IL-8, and IL-10 are associated with cognitive impairments observed in MA-associated psychosis. Significant increases in serum inflammatory cytokines (TNF- α , IL-6, and IL-18) in chronic MA abusers are still observed even after abstinent from MA for almost six weeks [22].

Among ample pro-inflammatory cytokines produced by MA, IL-6 and IL-18 exclusively reported their association with cognitive impairments in various human conditions. For example, a high level of IL-6 with metabolic syndrome increases the risk of cognitive impairment [23]. Another study in post-ischemic stroke reported that patients with high IL-6 levels exhibited a high possibility of cognitive decline [24]. Notably, IL-6 negatively correlated with cognitive ability determined by Mini-Mental State Examination (MMSE) in the multiethnic cohort [25]. Additionally, the higher IL-18 is associated with more significant cognitive impairments in schizophrenia [26, 27]. The correlation between IL-18 and cognitive impairment is also found in Alzheimer's disease (AD) [28]. It seems that cytokines affect cognitive impairment in neuropsychological and neurodegenerative disorders.

Numerous studies have been taken on animals to elucidate the mechanism underlying MAinduced neuroinflammation. It has demonstrat-

ed that acute MA administration increases IL-6 and TNF-a mRNA expression in the mouse hippocampus, striatum, frontal cortex [29], and IL-1 β in hippocampal tissue [30]. In addition, a study in the MA-treated rat brain demonstrates a significant increase in IL-1β mRNA expression in the hypothalamus [31]. These cytokines are related to neurotoxicity, which affirms by neurotoxic attenuation after blocking their production [30, 32, 33]. The molecular mechanism of MA elicits neuroinflammation suggested to exert the prominent role of glial cells. MA induces microglia and astrocytes activation and is related to the Toll-like receptor 4 (TLR4), which is involved in the immune surveillance of pathogens and exogenous small molecules. Consequently, TRL4 activation promotes nuclear factor-kappa B (NF-kB) to regulate the transcription of pro-inflammatory cytokines such as IL-1 β , IL-6, TNF- α and IL-18 [34, 35]. These insults activate the innate and acquired immune responses and initiate inflammatory reactions in the extracellular space and contribute to the pathophysiology of neuroinflammation [36].

Under the neuroinflammatory process, proinflammatory cytokines can alter the tryptophan metabolite by enhancing the expression of indoleamine 2,3-dioxygenase (IDO), the key enzyme in the tryptophan metabolic pathway [37]. Then IDO converts tryptophan into kynurenic acid (KYNA), which is an N-methyl-Daspartate receptor (NMDAR) antagonist in the human CNS [38, 39]. Elevated production of KYNA inhibits the NR1 subunit of NMDAR and alpha 7 nicotinic acetylcholine receptor (a7nAchR), leading to decreased NMDAR function and reduced a7nAchR-mediated glutamate release [40-42]. Thus, reduced levels of glutamate and NMDAR function were observed in the prefrontal cortex [43, 44]. Furthermore, IDO enzyme activation by pro-inflammatory cytokines can also increase another tryptophan metabolite, quinolinic acid (QUIN), an NMDAR agonist. A high level of QUIN results in excitotoxicity and neural cell death [45, 46]. Thus, these tryptophan metabolites might be key elements of extensive processes under inflammatory cytokines-induced neurological and neuropsychological dysfunctions. However, the effects of MA on neuroinflammation related to tryptophan metabolites and the extent of induced neural degeneration are not well understood.

The direct identification of target biomarkers in the human brain is limited to many ethical concerns. The interaction of neuroinflammation and tryptophan metabolites in neurodegenerative processes is mainly taken in animal studies [47, 48]. Hence, investigating biomarkers from the circulating blood system is one of the appropriately non-invasive approaches for exploring the neuropathological in the brain by taking advantage of blood-brain interaction [49]. The blood-brain barrier (BBB) could be disrupted following immune aggravation [50]; thus, biological changes discovered in the blood could trace the changes in the brain. Therefore, in this study, we aimed to determine the markers of inflammatory reactions and tryptophan metabolites in blood samples of MA abusers associated with their cognitive functions.

Methods

Participants

In this study, 15 MA abusers and 15 HC participants of the Thai population were recruited. The MA abusers were patients in Drug Abuse Treatment unit, Thanyarak hospital, Thailand. The study was approved by the Mahidol University Central Institutional Review Board (MU-CIRB 2016/043.3103). To perform experiments, all participants must previously voluntarily provide their permission in the informed consent form. The overall inclusion criteria include people 25-55 years old, normal eyesight and hearing (or had recovered to normal), and a Thai Mental State Examination (TMSE) score not less than 23. The MA addiction was diagnosed by a psychiatrist. Meanwhile, apart from cigarettes and alcohol, the HC must not represent any drug use. In addition, participants who experienced brain injury were excluded.

Questionnaires

The questionnaires contain information about demographic factors, diseases, drug administration, and drug addiction, including MA.

Screening tools

Participants were examined for psychotic symptoms by the Brief Psychiatric Rating Scale (BPRS) [51] and screening for mild cognitive impairment (MCI) by the TMSE.

TMSE is the Thai cognitive screening test developed from MMSE [52]. This Thai version screening test is purposed to screen dementia for Thai elderly and patients. The cut-off testing to distinguish dementia is 23 from 30 marks. TMSE is divided into six dimensions concerning 6 points of orientation, 3 points of registration, 5 points of attention, 3 points of registration, 10 points of language, and 3 points of recall. Therefore, the cut-off point to determine cognitive impairment is 23 out of 30.

The BPRS is a rating scale that a clinician or researcher may use to measure 18 psychiatric symptoms, including somatic concern, anxiety, guilt feeling, grandiosity, depressive mood, hostility, suspiciousness, hallucination, unusual thought content, disorientation, conceptual disorganization, emotional withdrawal, tension, mannerism and posturing, motor retardation, uncooperativeness, blunted affect and excitement. The subjects specified each area of the symptoms in the rating range from 1-7; 1 mark represents no psychiatric sign, while 7 marks represent the most severe symptom. The total score is from 18-126 (lowest to highest). A score equal to or lesser than 36 is interpreted as a low level of psychiatric symptoms; however, a score of more than 36 demonstrates a high level of psychiatric symptoms.

Determination of cognitive functions

A computerized Stroop test: A computerized Stroop test is a tool used to assess inhibition, selective attention, and cognitive flexibility [53, 54]. The task comprises a set of word-naming colors, e.g., red, green, blue, and yellow. Those words are represented in a random colors of red, green, blue, and yellow. Thus, the task consists of 2 different modes; congruent and incongruent words. The congruence represents in the same way as color and word meaning. The incongruence illustrates different ways between color and word meaning. Participants must click the right mouse button when a congruent word appears on the screen. Conversely, participants must click the left mouse button when an incongruent word appears. This task contains 200 trials to perform within 5 minutes. The measurement parameters are the percentage of correct congruence, miss, incongruence, and reaction time (millisecond: msec.). From these parameters, the percentage of congruence and incongruence represent selective attention and inhibition, respectively. Cognitive flexibility is the ability to switch responses between congruence and incongruence.

Go/No-Go task: The task comprises Go and No-Go conditions for assessment attention and inhibition [55, 56]. This task consisted of a set of random numbers between 0 to 9 representing on the screen by which 0 is a target clue, and 1 is a target probe. Participants must respond Go-condition when 0 is followed by 1 by immediate clicking on the left mouse button. On the contrary, they do nothing for the No-Go condition when 0 follows with a non-target probe (2-9). The task comprises 60 events of the Go-condition, 30 events of the No-Go condition, and 20 other interfering pictures. The overall measuring time is about 5 minutes. The psychometric measurement parameters include the response percentage of Go, No-Go, miss, commission error, and reaction time (msec.). The percentage of Go correct responses represents attention.

In contrast, the miss percentage should reversely correlate with this score. The percentage of No-Go and commission error reveal response inhibition. Since commission error makes a Go response on the No-Go trial, lesser commission error represents better response inhibition.

One-back test: This test mainly uses for measuring working memory [57]. The computer screen randomly displayed one Thai letter at a time. Participants must respond to a target that represents 2 continuing orders of letters by clicking on the left mouse button. The test contains 30 events of the target (15%) from a total of 200 events. It took about 6 minutes to accomplish. The psychometric scores generated from the test are composed of the percent of correct, incorrect, and reaction time (msec.).

Wisconsin card sorting test-64 (WCST-64): The WCST-64 is designed for prefrontal brain ability determination, which mainly investigates cognitive flexibility in the face of changing schedules of reinforcement. Cognitive functions such as

working memory, attention, and visual processing are involved. Besides, in case of frontal lobe damage leads to poor performance determined by this test [58-61]. The WCST-64 test contains 64 cards with different geometric designs for the participants to decide which matches the best with the stimulus cards. The criteria for matching are composed of colors (red, blue, yellow, or green), forms (triangle, star, cross, or circle), and numbers (1, 2, 3, or 4). The matching process is according to the unknown rule. In this test, the matching rule will change when the correct scores continuously reach 10 cards without warning. The test generates various psychometric scores consisting of total correct (%), total error (%), perseverative response (%), perseverative error (%), non-perseverative error (%), number of trials to complete the first category, and reaction time (msec.) [62].

Complete blood count (CBC) analysis

Whole venous blood of about 3 mL was collected in an ethylenediamine tetra-acetic acid (EDTA) tube and determined as a part of a routine laboratory test in the hospital using a Beckman Coulter AU 2700 analyzer (Brea, CA).

Blood samples measurement of inflammatory cytokines

Inflammatory markers in human serum were detected by guantitative sandwich Enzyme-Linked Immunosorbent Assay (ELISA) using paired antibodies, including IL-6 (Biolegend Inc., USA) and IL-18 (R&D Systems, USA). First, venous blood was collected and left to clot at room temperature for about 30 minutes. Then, clotted blood was centrifuged at 3,000 rpm for 15 minutes at room temperature to separate serum fractions. Next, the serum samples were kept in new tubes at -80°C until examined. Inflammatory cytokines were measured by following guide protocols. The absorbance of color reaction was read at 450 nm with a microplate reader from BioTek® Instruments (Vermont, USA) for IL-6, while IL-18 was detected via EZ Read 2000 microplate reader (Biochrom, USA). Finally, the concentration of inflammatory cytokines was calculated by comparing them with their standard curves.

Measurement of tryptophan metabolites

The serum samples were also determined of tryptophan metabolites by liquid chromatogra-

participants and MA users		
Demographic data	HC (N = 15)	MA (N = 15)
Age (years)	32.20±6.81	34.13±6.40
Gender (males/female)	8/7	6/9
Years of education	13.07±3.83	11.13±2.48
Duration of methamphetamine use (years)	-	11.80±6.11
Methamphetamine use (gram/day)	-	0.30±0.57
Abstinence duration (days)	-	12.13±1.85
TMSE	28.87±1.06	26.47±2.53**
BPRS	20.00±2.27	22.87±3.48*

 Table 1. Demographic data, TMSE and BPRS scores of HC participants and MA users

Values are mean \pm SD. *P < 0.05 and **P < 0.01 compared to HC.

phy with tandem mass spectrometry (LC-MS/ MS). The LC-MS/MS system consists of a liquid chromatography part (Dionex Ultimate 3000) in combination with an ESI/mass spectrometer (Model Q-ToF Compact II, Bruker, Germany). All samples were kept in low-protein binding microcentrifuge tubes. In the first step, serum was overnight mixed with methanol and allowed protein aggregation before collecting the supernatant. The solvent was dried using speed vacuum centrifugation and kept at -80°C until use. Then the pellets were dissolved in H₂O containing 0.1% formic acid and centrifuged at 12,000 rpm, 4°C for 5 min. At this step, the supernatants were ready to measure. Notably, a pool tube for each group was necessary to control all discovered metabolites. In the stage of measurement, the samples run in machine DIONEX Ultimate[™] 3000 Ultra High-Performance Liquid Chromatography (UHPLC) system (Thermo Fisher Scientific, USA) through Acclaim[™] Polar-Advantage II C18 (PA2) column (3 µm, 120 Å 2.1 × 100 mm) Dionex Bond Silica Products (Thermo Fisher Scientific, USA) with mobile phase A (0.1% formic acid in H_2 O) and B (0.1% formic acid in acetonitrile) about 20 minutes. The compounds containing either positive or negative charge were determined by mass spectrometry (Model Q-ToF Compact II, Bruker, Germany).

Statistical analysis

The normality and lognormality were applied to determine the normal distribution in every pair of comparisons for appropriate parametric and nonparametric statistical tests. In addition, the Mann Whitney and Unpaired T-test were used for the nonparametric and parametric of the two classes differentiation, respectively. These were performed using GraphPad Prism 8.0.0, GraphPad Software, San Diego, California USA, www. graphpad.com. The statistical significance is determined at *p*-values less than 0.05.

Results

Demographic data of participants

There were no differences in demographic data such as substance use information, age and years of education between the two groups of participants. However, MA abusers showed a significant decrease in TMSE scores (P = 0.0021) along with significantly higher BPRS scores (P = 0.0124) when compared to HC (**Table 1**).

Cognitive performance in healthy control (HC) participants and methamphetamine abusers (MA)

Selective attention assessment by a computerized Stroop test: The results showed a significantly lower percentage of correct congruence (P = 0.0026) in MA compared to HC. Consequently, the percentage of the miss was significantly higher (P = 0.0028) in the MA group. Moreover, MA also spent more reaction time and was significantly different compared to HC (P = 0.0236) (Table 2).

An inhibition assessment by Go/No-Go task: The percent of correct Go and No-Go responses were determined for assessment attention and inhibition, respectively. The results demonstrated a significantly lower percentage of No-Go correct response ($P \le 0.0001$) and higher commission error ($P \le 0.0001$) performed by MA users compared to HC (**Table 2**).

Working memory assessment by One-back test: The percentage of a correct response on target was determined to assess working memory. MA performed a significantly lower percentage of correct responses (P = 0.0029) than HC. The result reasonably represents along with a

Cognitive test	HC (N = 15)	MA (N = 15)
Stroop test		
Congruence (%)	95.30±5.42	87.78±6.29**
Miss (%)	4.69±5.42	12.22±6.29**
Congruence Reaction Time (msec.)	587.60±35.53	643.10±82.32*
Go/No-Go		
Go (%)	97±4.64	93.22±7.25
Miss (%)	3.00±4.64	6.78±7.25
No-Go (%)	98.67±3.03	91.56±7.44****
Commission error (%)	1.33±3.03	8.44±7.44****
Go Reaction Time (msec.)	425.10±103.9	434.80±142.50
One-back test		
Correct (%)	78.21±15.22	57.10±19.95**
Miss (%)	21.61±15.01	42.67±19.32**
Hit Reaction Time (msec.)	612.60±142.70	647.70±191.10
WCST-64		
Total correct (%)	85.36±4.38	74.48±7.03****
Total error (%)	14.64±4.38	25.96±6.80****
Perseverative response (%)	13.47±3.36	15.17±6.14
Perseverative error (%)	9.00±1.73	11.73±4.60*
Non-perseverative error (%)	5.27±3.41	14.23±5.28****
Trials to complete the first category	11.53±1.96	18.80±9.17**
Reaction time (msec.)	2253±630.2	3461± 491.9****

Table 2. Cognitive test in HC participants and MA users

Values are mean \pm SD. **P* < 0.05, ***P* < 0.01, and *****P* < 0.0001 compared to HC.

CBC parameters	HC (N = 15)	MA (N = 15)	Normal range [102]
RBCs (× 1,000,000 µl)	4.96±0.60	4.45±0.79	4.00-6.10
WBCs (× 1000 µl)	6.77±1.55	8.09±1.75*	4.40-11.30
Platelet count (× 1000 µl)	283.70±61.50	316.70±62.71	179.00-356.00
Hb (g/dL)	13.71±1.77	11.38±1.37***	12.00-16.90
Hct (%)	40.90±4.73	35.99±4.52**	37.00-51.90
Neutrophil (%)	56.60±7.88	57.98±10.05	40.00-70.30
Eosinophil (%)	3.33±2.32	5.73±5.02	0.40-9.20
Basophil (%)	0.27±0.46	0.73±0.59*	0.20-1.40
Lymphocyte (%)	35.13±6.88	33.40±5.37	20.30-48.30
Monocyte (%)	4.53±1.46	4.80±1.15	3.40-12.30
MCV (fL)	82.64±5.26	81.85±8.00	80.40-98.80
MCH (pg)	27.83±1.98	25.95±3.08	25.00-31.20
MCHC (g/dL)	33.67±0.97	31.67±0.83****	30.20-34.60

Values are mean ± SD. **P* < 0.05, ***P* < 0.01, ****P* < 0.001, and *****P* < 0.0001 compared to HC. MCV: Mean Corpuscular Volume; MCH: Mean Corpuscular Hemoglobin; MCHC: Mean Corpuscular Hemoglobin Concentration; g: gram; dL: deciliter; fL: femtoliter; pg: picogram.

significantly higher percentage of miss (P = 0.0024) (Table 2).

Cognitive flexibility, concept formation, and processing speed assessment by WCST-64: WCST-64 was employed to determine executive functions, especially in cognitive flexibility. MA abusers exhibited cognitive deficits compared to HC. MA group performed a significantly lower total correct score than HC $(P \leq 0.0001)$. Contradictory, MA showed a significantly higher total error (%) ($P \leq$ 0.0001), perseverative error (P = 0.0403), non-perseverative error ($P \leq 0.0001$) and trials to complete the first category (P = 0.0025) than HC, respectively. In addition, a significantly longer reaction time was observed in MA compared to HC ($P \leq$ 0.0001) (Table 2).

CBC analysis

The results of CBC analysis showed that WBC numbers significantly increased in MA groups compared to HC (P =0.0375). In addition, basophil, an allergic immune ce-II, significantly increased in MA compared to HC (P = 0.0227). Apart from WBC, RBC indices also showed significant differences in the MA group compared to HC. Significant decreases in hemoglobin (Hb) (P = 0.0004), hematocrit (Hct) (P = 0.007), and mean corpuscular Hb concentration (MCHC) ($P \leq$ 0.0001) were observed in MA abusers compared to HC (Table 3).

Serum levels of pro-inflammatory cytokines and L-tryptophan

The serum levels of IL-6 slightly increased (Figure 1A) while the serum levels of IL-18



Figure 1. Serum levels of (A) IL-16, (B) IL-18, and (C) L-Tryptophan in HC participants and MA abusers. Values are mean \pm SD. *****P* < 0.0001 compared to HC.

slightly decreased (**Figure 1B**) in the MA group compared to HC, respectively. In addition, the results showed a significant decrease in L-Tryptophan levels in the MA group compared to HC ($P \le 0.0001$) (**Figure 1C**).

Discussion

The results of the present study showed that demographic data matches HC participants and MA abusers. However, the mild cognitive assessment by TMSE [52] demonstrated low TMSE scores in MA abusers compared to the HC group. On the opposite, the evaluation of the psychotic symptoms by BPRS [51] revealed higher BPRS scores in MA abusers compared to the HC group. The neurocognitive assessment showed that MA abusers exhibited deficits in multiple cognitive functions, including selective attention, inhibition, working memory, cognitive flexibility, concept formation, and processing speed. Blood sample analysis demonstrated a decrease in RBC components, especially Hb, Hct and MCHC but an increase in WBCs and basophils in MA abusers compared to the HC group. Moreover, a slight increase in serum IL-6 levels, a slight decrease in IL-18, and a significant decrease in serum L-tryptophan levels were observed in MA abusers compared to the HC group. The key finding of this study might postulate the neuroinflammation-induced pathophysiological mechanisms of cognitive deficits in MA abusers. The inflammation in MA abusers was classified as chronic low-grade inflammation, represented by anemia of inflammation (or anemia of chronic disease) and an increase in inflammatory cytokines in blood analysis [63-66]. Taken together, the induction of inflammatory reaction was also alternatively demonstrated by the activation of tryptophan metabolites [67, 68].

Recent evidence has emphasized that oxidative damage and chronic systemic inflammation might be the initial factor to activate the anemia of inflammation. The study in phenylhydrazine (PHz)-intoxicated C57BL/6J mice model of anemia demonstrated that the antioxidant molecule, nano-complex of manganese and citrate could abolish PHz-induced reduction in Hb, RBC count, antioxidant enzyme (superoxide dismutase, glutathione peroxidase, and catalase) activities, and CD4+/CD8+ T-lymphocyte ratio, and induction in serum levels of lipid peroxidation, TNF- α , IFN-y and IL-6 [66]. Concomitant with oxidative damage and neuroinflammation, substantial evidence has revealed that MA can induce oxidative damage and inflammation in CNS. MA in blood circulation can directly pass BBB into the brain according to its high lipid solubility [69]. The action of MA in the brain involves the generation of oxidative stress and excitotoxicity [70-72] and also induces neuroinflammation through a striking correlation to microglial activation [34, 35, 70, 73-76]. In addition, MA possesses oxidative damage by induced DA overflow and auto-oxidized, resulting in reactive oxygen species (ROS) formation [77-79]. Taken together, MA can disturb the electron transport chain in the ATP production process of mitochondria which also gives rise to the leakage and accumulation of ROS [70, 80]. High production of ROS further induces oxidative stress and oxidative damage. It has been reported that MA-induced oxidative stress may cause the generation of damage-associated molecular patterns (DAMPs) [81]. The correlation between MA and inflammation has been investigated for the activation of Toll-like receptors (TLRs) by DAMPs [82]. The inflammatory cascades of TLRs activation in glial cells [34, 82] subsequently induce NF-κB signaling [83] leading to an increase of inflammatory cytokines production such as IL-1β, TNF- α , IL-18, and IL-6 [29, 32, 35, 81, 82, 84-90].

Whether several lines of evidence suggest that MA can induce inflammation in CNS, leading to disturbing neurological functions; however, most studies have been done in animals [30, 91-93]. Moreover, the brain structure system is mainly closed, and manipulation of the human brain components is invasive. Thus, investigating the association between neuroinflammatory reactions and neurological function in MA abusers needs a more conclusive study. Interestingly, recent data by Tipton and colleagues suggest that MA elicits human inflammation via increased peripheral cytokine production from monocyte or macrophage [90]. Therefore, the study of blood components might be an alternative biomarker to determine the role of MA-induced neuroinflammation, which affects cognitive deficits in MA abusers.

Furthermore, our results showed a slight increase in serum IL-6 levels. Still, a slight decrease in serum IL-18 level in MA abusers compared to the HC group might explain the restorative effect of MA withdrawal [94, 95]. Thus, drug abstinence is a limitation in this study by which the more prolonged abstinence, the more recovery of brain pathology. Therefore, further study of blood inflammatory markers during MA administration are remarkably required for better elucidation.

Moreover, a new finding was observed in the present study: a significant perturbation of RBC indices such as Hb, Hct, and MCHC in MA users. These decreasing levels might be occurred by which anemia of inflammation. The evidence

possibly occurs due to prolonged immune activation by several inflammatory cytokines, such as IL-1β, IL-6, IL-10, and IFN-y which control iron homeostasis by prohibiting Hb production via promoting iron uptake into macrophage and directly damage erythroid progenitors via apoptotic activation [64]. In addition, the percentage of serum basophils up-regulated. Basophil is derived from mast cells and is responsible for allergic reactions via releasing histamine. Previous studies in rodent brains demonstrated a significant increase of histamine in the hypothalamus after a moderate dose of MA injection [96]. The same evidence might also be occurred in blood circulation as in this study. The involvement of IL-6 in the pathophysiology of anemia of inflammation has been investigated in various chronic/inflammatory diseases such as end-stage renal disease, chronic kidney disease, and rheumatoid arthritis. IL-6 mediates anemia through the induction of hypoferremia and ends with iron-restricted erythropoiesis [97, 98].

Further study on the role of neuroinflammation on neurological function has emphasized the association between low levels of tryptophan and cognitive decline [99]. The evidence of a high ratio of tryptophan metabolite, kynurenine and tryptophan was also found to conversely correlate with white matter integrity and low glutamate levels in schizophrenia [100]. Recent evidence has demonstrated that pro-inflammatory cytokines can induce the activity of IDO, the key enzyme of tryptophan metabolites, leading to decreased tryptophan levels and increased the concentration of tryptophan metabolites such as QUIN and KYNA [38, 39]. KYNA acts as an NMDAR antagonist, while QUIN acts as the NMDAR agonist in the human CNS [38, 39]. Excessive production of the tryptophan metabolic pathway results in either low glutamate levels-induced reduction in NMDAR function [40-44] or cvtotoxicity-induced neural degeneration [39, 101]. This evidence might support our findings on the correlation between the low level of L-Tryptophan and cognitive deficits in MA abusers (Figure 2).

Taken together, the results of this study might emphasize the association between perturbations in tryptophan metabolites and low-grade systemic inflammatory conditions in MA abus-



Cognitive deficit and neuroinflammation in methamphetamine abusers

Figure 2. The postulated mechanism of blood-brain interaction in MA-induced anemia of inflammation, tryptophan metabolites and neurological dysfunction. MA affects the immune system by stimulating NF-κB cascades in macrophages in peripheral blood or microglia and astrocytes in the brain. In the brain, pro-inflammatory cytokines are produced from activated microglia and astrocytes. A similar mechanism occurs in the peripheral system by which MA stimulates pro-inflammatory cytokine production and release from macrophages. The pro-inflammatory cytokines disturb iron homeostasis and Hb production in red blood cells leading to anemia of inflammation. In addition, pro-inflammatory cytokines by activating IDO1 result in the production of KYNA or QUIN. Finally, KYNA and QUIN provoke cognitive deficits by dysregulating NMDAR function, inducing cytotoxicity and neural degeneration. TRY: Tryptophan; KYNA: Kynurenic Acid; QUIN: Quinolinic Acid; 3-HK: 3-Hydroxykynurenine; KATs: Kynurenine Aminotransferase enzymes; KMO: Kynurenine-3-Monooxygenase; 3-HAO: 3-Hydroxyanthranilic Acid Oxygenase.

ers. We propose that the attribute of tryptophan metabolites in MA-induced cognitive deficits may gain further insights into their potential for developing companion diagnostics and more targeted drug interventions in neurological diseases.

Conclusion

The key findings in this study suggest that blood biomarkers of anemia of inflammation and tryptophan metabolites might be the alternative markers to demonstrate the induction of inflammatory reactions in MA abusers. Moreover, these surrogate blood inflammatory markers might utilize as an indicator to represent neuroinflammation-related neurological dysfunction in MA-induced neurotoxicity and cognitive deficits.

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Disclosure of conflict of interest

None.

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