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(As of October 2022)

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# **Original** Article

## Inhaled ciclesonide does not affect production of antibodies or elimination of virus in patients with COVID-19: Subanalysis of a multicenter, open-label randomized trial

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SUMMARY During an earlier multicenter, open-label, randomized controlled trial designed to evaluate the effectiveness of high-dose inhaled ciclesonide in patients with asymptomatic or mild coronavirus disease 2019 (COVID-19), we observed that worsening of shadows on CT without worsening of clinical symptoms was more common with ciclesonide. The present study sought to determine if an association exists between worsening CT shadows and impaired antibody production in patients treated with inhaled ciclesonide. Eighty-nine of the 90 patients in the original study were prospectively enrolled. After exclusions, there were 36 patients each in the ciclesonide and control groups. We analyzed antibody titers against severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) nucleocapsid protein at various time points. Changes in viral load during treatment were compared. There was no significant difference in age, sex, body mass index, background clinical characteristics, or symptoms between the two groups. Although evaluation on day 8 suggested a greater tendency for worsening shadows on CT in the ciclesonide group (p = 0.072), there was no significant difference between them in the ability to produce antibodies (p = 0.379) or the maximum antibody titer during the clinical course. In both groups, worsening CT shadows and higher viral loads were observed on days 1-8, suggesting ciclesonide does not affect clearance of the virus (p = 0.134). High-dose inhaled ciclesonide did not impair production of antibodies against SARS-CoV-2 or affect elimination of the virus, suggesting that this treatment can be used safely in patients with COVID-19 patients who use inhaled steroids for asthma and other diseases.

*Keywords* SARS-CoV-2 nucleocapsid protein antibody, asymptomatic or mild COVID-19, randomized clinical trial, anti-viral effect, viral load

#### 1. Introduction

In 2019, severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) emerged and seriously threatened human health and socioeconomic activity worldwide. Clinical manifestations of SARS-CoV-2 infection vary depending on the presence of risk factors that include obesity, cardiovascular disease, diabetes, older age, and comorbidities (1). The rationale for using steroids in patients who are hospitalized with coronavirus disease 2019 (COVID-19) is based on the finding in a clinical trial that administration of dexamethasone reduced mortality in these patients, suggesting that the anti-

inflammatory effects of the compound overcome its negative immunosuppression effects (2).

Interestingly, ciclesonide is thought to ameliorate inflammation by inhibiting the pathogenic kinase PAK1 (RAC/CDC42-activated kinase 1) (3). Basic research has suggested that ciclesonide may have potent antiviral activity against SARS-CoV-2 (4), and case reports have also shown the effectiveness of ciclesonide (5). Moreover, several lines of evidence suggest that inhaled ciclesonide has anti-viral activity (6,7). Although it is unclear whether asthma is a risk factor for severe SARS-CoV-2 infection (8), several clinical trials of ciclesonide have been conducted in outpatients or hospitalized

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patients based on these results (9-11). However, it has not been confirmed that ciclesonide improves outcomes in patients with severe clinical symptoms. We performed a clinical trial to evaluate the effects of ciclesonide 400 µg inhaled three times daily for 7 consecutive days in patients with mild clinical signs of COVID-19 at the time of enrollment (12). Surprisingly, we found that worsening of shadows on CT was more common in patients treated with ciclesonide (39%, n = 16) than in those who received symptomatic treatment alone (18.8%, n = 9). Given that antibody production has been reported to be impaired after SARS-CoV-2 vaccination in patients on corticosteroids (13), it is necessary to clarify whether ciclesonide causes worsening of CT shadows and impairs antibody production.

The aim of this study was to determine if there is an association of worsening CT shadows with impaired antibody production in patients treated with inhaled ciclesonide.

### 2. Materials and Methods

### 2.1. Study population and protocol

This study is a subanalysis of a multicenter, open-labelled randomized trial (RACCO study). The complete study protocol has been described previously (12). This trial was approved by the Clinical Research Review Board at The University of Tokyo (protocol number: 2019017SP). Written informed consent was obtained from all patients before participation.

### 2.2. Demographic characteristics

Eighty-nine of the 90 patients who participated in our previously reported trial (12) were enrolled in the present study (Table S1 in the Supplementary Data, http://

www.ddtjournal.com/action/getSupplementalData. *php?ID=173*) and were divided into a ciclesonide group (n= 41, 20 men, mean age  $38.7 \pm 17.0$  years) and a control group (n = 48, 24 men, mean age  $42.7 \pm 18.7$  years). One patient was excluded from the analysis due to lack of baseline CT. There was no between-group difference in body mass index, mean age, or duration of hospitalization after onset of clinical signs of SARS-CoV-2 infection. After excluding 2 patients whose condition deteriorated and required additional medications, 3 who were discharged on day 7 (with a medication compliance rate of < 90%), and 12 who were negative on day 1 of the study but positive before hospitalization (ciclesonide group, n = 3; control group, n = 9), this left 72 patients for this study (ciclesonide group, n = 36; control group, n =36) (Figure 1).

2.3. Exacerbation of pneumonia and production of antibodies against SARs-CoV-2 nucleocapsid protein

Antibody titers against SARS-CoV-2 nucleocapsid protein were measured by enzyme-linked immunosorbent assay on hospital days 1, 8, 15, and 22 and then compared between patients with exacerbation of pneumonia in the ciclesonide group and their counterparts in the control group. Exacerbation of pneumonia was judged on CT scans obtained on day 1 (pretreatment) and day 8 (12). The actual dates for antibody measurement were calculated from the date of onset of clinical signs of SARS-CoV-2 infection.

### 2.4. Measurement of antibody titers

A plate coated in recombinant nucleocapsid protein was incubated with a serum/plasma sample diluted to 1/800 for 1 h at 37°C after blocking with 1% BlockAce (KAC, Kyoto, Japan). Following washing with phosphate-



Figure 1. Flow diagram of participant recruitment in this study.

buffered saline containing 0.2% Tween 20, the plate was incubated with anti-human IgG conjugated with horseradish peroxidase (GeneTex, Irvine, CA) for 30 min at 37°C. The captured antibodies were detected with 3,3',5,5'-tetramethylbenzidine substrate solution (Nacalai Tesque, Kyoto, Japan) and measured at a wavelength of 450 nm by a microplate reader (Bio-Rad, Irvine, CA, USA). Samples from a healthy volunteer without SARS-Cov-2 infection were used as a negative control. Antinucleocapsid human IgG (ACROBiosystems, Newark, DE) was used as a positive control and a standard curve was created. Each sample was assayed in triplicate, and all measurements were normalized to the value of the positive control. The positive and negative cut-off values for the antibody titer were set to a value of the mean plus three times the standard deviation of the negative control.

### 2.5. SARS-CoV-2 copy numbers

Nasopharyngeal swab fluid was collected on days 1 and 8. Nucleic acid RNA was extracted from the nasopharyngeal swab fluid using a QIAamp virus RNA mini kit. To determine the amount of SARS-CoV-2 per mL of universal transport medium (14), quantitative realtime reverse transcription-polymerase chain reaction (qRT-PCR) was performed with QuantStudio 12K (Thermo Fisher Scientific, Waltham, MA) using primer/ probe nucleocapsid protein set no. 2 (N2) (15). EDX SARS-CoV-2 standard (BR code: COV019, Bio-Rad Laboratories, Inc. Japan, 200,000 copies/mL of N-region RNA) was used as a quantitation standard to correct the calibration curve in real-time PCR quantification.

### 2.6. Statistical analysis

The anti-nucleocapsid IgG titer was compared between the placebo and ciclesonide groups using the nonparametric Mann-Whitney U test. For statistical analysis of the viral load, the value under the limit of quantification was set to 50 copy/mL and the value under the level detectable by qRT-PCR was set to 1 copy/mL. Categorical variables were compared between the groups using Fisher's exact test. A nonparametric Kruskal-Wallis test was performed to compare more than three groups followed by a post-hoc Dunn's multiple comparisons test to identify significant differences between specific groups. All statistical analyses were performed using Prism software (version 9; GraphPad Software, La Jolla, CA). A *p*-value < 0.05 was considered statistically significant.

### 3. Results

### 3.1. Patient characteristics

The clinical characteristics of the ciclesonide group (n = 36) and the control group (n = 36) are shown in Table 1. Age tended to be younger in the ciclesonide group, with a median age of 44 years in the control group and 32.5 years in the ciclesonide group, but there were no significant differences. The control group consisted

<b>Table 1. Clinical characteristics</b>	of the	study participants	5
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Items	Control group ( $n = 36$ )	Ciclesonide group $(n = 36)$	<i>p</i> -value
Age, years <sup>†</sup>	44.0 [25.3, 60.8]	32.5 [27.0, 46.3]	0.2284ª
Sex			
Female	18	19	$> 0.9999^{b}$
Male	18	17	
$\mathrm{BMI}^\dagger$	22.0 [20.2, 25.7]	22.0 [19.7, 24.4]	$0.448^{a}$
Days from symptom onset to day $1^{\dagger}$	5.0 [3.0, 7.0]	5.0 [3.0, 6.0]	$0.7977^{a}$
Clinical features on day 1 <sup>††</sup>			
Body temperature ( $> 37.5^{\circ}$ C)	13 (36.1) ††	14 (38.9)	$> 0.9999^{b}$
Dry cough	16 (44.4)	22 (61.1)	0.2377 <sup>b</sup>
Productive cough	5 (13.9)	6 (16.7)	$> 0.9999^{b}$
Haemoptysis	0 (0.0)	0 (0.0)	$> 0.9999^{b}$
Sore throat	6 (16.7)	11 (30.6)	0.2668 <sup>b</sup>
Runny nose	7 (19.4)	5 (13.9)	0.7531 <sup>b</sup>
Wheezing	0 (0.0)	0 (0.0)	$> 0.9999^{b}$
Dyspnoea	9 (25.0)	6 (16.7)	0.5628 <sup>b</sup>
Chest pain	3 (8.3)	3 (8.3)	$> 0.9999^{b}$
Muscle pain	6 (16.7)	6 (16.7)	$> 0.9999^{b}$
Joint pain	3 (8.3)	4 (11.1)	$> 0.9999^{b}$
Headache	8 (22.2)	9 (25.0)	$> 0.9999^{b}$
Altered consciousness	0 (0.0)	1 (2.8)	$> 0.9999^{b}$
Fatigue	13 (36.1)	13 (36.1)	$> 0.9999^{b}$
Abdominal pain	3 (8.3)	1 (2.8)	0.6142 <sup>b</sup>
Vomiting/nausea	2 (5.6)	1 (2.8)	$> 0.9999^{b}$
Diarrhoea	7 (19.4)	7 (19.4)	$> 0.9999^{b}$

Data are presented as the <sup>†</sup>median [interquartile range] or <sup>††</sup>number (percentage). <sup>a</sup>Mann-Whitney U test. <sup>b</sup>Fisher's exact test. BMI, body mass index.

of 18 males while the ciclesonide group consisted of 17 males. Both groups had a median body mass index (BMI) of 22.0, and the interval between symptom onset and hospitalization was 5 days for both groups, with no significant differences. Regarding clinical symptoms on the first day of admission, 13 (36.1%) patients in the control group and 14 (38.9%) in the ciclesonide group had fever. Dry cough was more common in the ciclesonide group, 16 (44.4%) in the control group and 22 (61.1%) in the ciclesonide group, but the difference between the two groups was not significant. Other respiratory symptoms such as sore throat, runny nose and dyspnoea were observed in 6 (16.7%), 7 (19.4%) and 9 (25.0%) patients in the control group respectively. In contrast, 11 (30.6%), 5 (13.9%) and 6 (16.7%) patients in the ciclesonide group showed no difference between the two groups. There were also no differences in systemic symptoms such as headache, muscle pain, joint pain and fatigue between the two groups: 8 (22.2%), 6 (16.7%), 3 (8.3%) and 13 (36.1%) patients in the control group and 9 (25.0%), 6 (16.7%), 4 (11.1%) and 13 (36.1%) patients in the ciclesonide group, respectively. In addition, gastrointestinal symptoms such as abdominal pain and diarrhoea occurred in 3 (8.3%) and 7 (19.4%) patients in the control group and 1 (2.8%) and 7 (19.4%) patients in the ciclesonide group, respectively, with no significant differences between the two groups.

3.2. Induction of antibodies against SARS-CoV-2 nucleocapsid protein

Enzyme-linked immunosorbent assay data confirmed that seroconversion was induced in both groups (Figure 2A). The maximum antibody titer was reached in 2-3 weeks (Figure 2B). Although induction of antibodies tended to be weaker in the ciclesonide group (Figure 2A), there was no significant difference in the days taken to reach the maximum antibody titer and the amount of antibodies (Figure 2C left, p = 0.187; right, p = 0.158, respectively). Furthermore, there was no between-group difference in the fold increase of antibody titres on day 8 compared to those detected on day 1 (Figure 2D, p = 0.379). These data indicate that inhaled ciclesonide did not impair seroconversion against SARS-CoV-2.

3.3. Association between exacerbation of pneumonia and impaired antibody production

Worsening CT shadows on day 8 was observed in 15 patients (41.7%) in the ciclesonide group and 7 (19.4%) in the control group, indicating a greater likelihood of worsening CT shadows with ciclesonide (Figure 3A, p = 0.072). However, there was no between-group difference in antibody titers on day 8 among patients with worsening CT shadows (Figure 3B left, p = 0.178 for control; p = 0.505 for ciclesonide). Moreover, there was



Figure 2 Comparison of anti-nucleocapside antibody acquisition and quantity change and comparison of maximum antibody titers between the control and ciclesonide groups. (A) Anti-nucleocapsid antibody titer over time starting at symptom onset. (B) Comparison of number of days from symptom onset to maximum antibody titer. (C) Days until maximum antibody value is reached. (D) Fold increase of antibody titers on day 8.

no significant between-group difference in the maximum antibody titer (Figure 3B right, p = 0.388 for control; p = 0.102 for ciclesonide). These data suggest that the worsening CT shadows observed in the ciclesonide group was not the result of impaired antibody production.

We also analyzed the viral load and frequency of detection of the virus on day 1 and 8. First, in the control group, we observed that the viral load was higher on both days 1 and 8 in cases with worsening CT shadows than in cases with amelioration (Figure 3C, p = 0.007 and p= 0.022, respectively). The frequency of detection on day 8 was significantly higher in cases with worsening CT shadows than in those with amelioration (Figure 3D left, p = 0.034). These data suggest that the worsening CT shadows in the control group were attributable to delayed elimination of the virus. In the ciclesonide group, we observed that the viral load was higher on day 1 in cases with worsening CT shadows than in cases with amelioration (Figure 3C left, p = 0.046), but there were no significant between-group differences on day 8 (Figure 3C, p = 0.568). There were also no betweengroup differences in the frequency of detection on day 8 (Figure 3D right, p = 0.134), and there was no significant difference in the time to discharge between cases with

worsening CT shadows and those with amelioration in the control group or ciclesonide group (Figure 3E, p =0.482 and p = 0.118, respectively). Furthermore, there were no significant between-group differences in clinical observations or laboratory findings on day 8 or at the time of discharge. Although there were more cases with worsening of CT shadows in the ciclesonide group, because there was no delay in viral clearance in the ciclesonide group compared with the control group, these data may indicate a therapeutic effect of ciclesonide.

### 4. Discussion

In this study, we prospectively compared the antibody production capacity between the ciclesonide group and control group in the RACCO study to determine whether administration of ciclesonide affects induction of antibodies against the SARS-CoV-2 nucleocapsid protein. We found that patients in both groups reached their peak antibody titers within 2-3 weeks, with no difference between them in peak values or number of days taken to reach them. There was also no significant difference in antibody production capacity, maximum antibody titer, or change in viral load between worsening



Figure 3 Comparison of the relationships between amelioration and exacerbation of CT images, viral antibody titers, and viral load between the control and ciclesonide groups. (A) Comparison of the amelioration and exacerbation groups: 15 of 46 patients (41.7%) in the ciclesonide group and 7 of 36 (19.4%) in the control group showed worsening CT shadows on day 8. (B) Comparison of antibody titers on day 8 and maximum antibody in the control and ciclesonide groups between the amelioration and exacerbation groups. (C) Viral load and detection frequency of SARS-CoV-2 in the nasopharyngeal swabs at days 1 and 8. (D) Detection frequency of the viruses. (E) Comparison of days from symptom onset to discharge.

CT shadows and improvement. These observations suggest that the influence of inhaled ciclesonide on antibody production capacity was neither statistically nor clinically significant.

Representative CT shadows in each group are compared in Figure 4. Both groups showed worsening CT shadows on day 8, suggesting a possible exacerbation of COVID-19 pneumonia. However, the shadows had disappeared at the 3-month follow-up. Although worsening CT shadows on day 8 were more common in the ciclesonide group, exacerbation of COVID-19 pneumonia during the clinical course was not observed in the ciclesonide group but was observed in 3 controls (Table 2). This discrepancy was addressed in a subsequent study by Inui et al., who found lung abnormalities even in asymptomatic cases, with 54% having pneumonia-like changes (16), suggesting a significant discrepancy between initial CT findings and subjective symptoms. Such imaging changes are reported to be distinct from changes indicative of the progression of severe pneumonia. Therefore, it is hypothesized that

the changes induced by ciclesonide might be different from these imaging-related changes.

Ciclesonide has been demonstrated to inhibit the intracellular enzyme PAK1, which is implicated in the pathogenesis of COVID-19 associated with angiotensinconverting enzyme 2. By inhibiting PAK1, ciclesonide is reported to alleviate immune suppression and reduce

 Table 2. Comparison of exacerbations of COVID-19

 pneumonia between the ciclesonide group and the control group

Adverse event	Control group $(n = 48)$	Ciclesonide group $(n = 41)$	
COVID-19 pneumonia	3	0	
Mild	0	0	
Moderate	2	0	
Severe	1	0	

Exacerbations were more common on computed tomography scans on day 8 in the ciclesonide group. However, exacerbation of COVID-19 pneumonia was not clinically diagnosed in any patients in the ciclesonide group but was diagnosed in 3 patients in the control group. COVID-19, coronavirus disease 2019.



Figure 4. Representative cases of changes in CT image findings in the control and ciclesonide groups. (A) Control group: 57-year-old man. (B) Ciclesonide group: 27-year-old woman. Worsening CT shadows at day 8 in both cases are no longer evident at the 3-month follow-up.

lung inflammation (3). Furthermore, it is believed that ciclesonide targets the non-structural proteins of this novel coronavirus. This target has been shown to hinder viral replication *in vitro* (4, 6, 7).

Song *et al.* performed a randomized non-blinded multicenter phase II clinical trial of inhaled ciclesonide in patients with mild-to-moderate COVID-19 and found a significantly higher SARS-CoV-2 clearance rate on day 14 in their ciclesonide group (p = 0.021). Inhaled ciclesonide was found to shorten the SARS-CoV-2 shedding period and demonstrated the potential to inhibit progression to acute respiratory failure in patients with mild-to-moderate COVID-19 (*17*).

Ciclesonide has been identified as a potential treatment for COVID-19 in several studies, including ours. However, there is still no clear research demonstrating its efficacy. Indeed, Clemency et al. reported that the median time to relief of all COVID-19-related symptoms in outpatients with symptomatic COVID-19 was 19.0 days in both a ciclesonide group and a placebo group. Although there was no significant difference in symptom improvement, patients in the ciclesonide group were less likely to make subsequent visits to the emergency room or be hospitalized for COVID-19-related reasons by day 30 than those in the placebo group (1.0% vs. 5.4%; p = 0.03) (10). Furthermore, in the COVERAGE study, which evaluated ciclesonide in adults at risk of severe progression of COVID-19, there was no significant difference in the primary endpoint of worsening COVID-19 (hospitalization, home oxygen therapy, or death) up to day 14 (9). Studies targeting hospitalized patients have also been conducted, but there was also no reduction in the duration of oxygen administration in patients hospitalized with COVID-19 receiving oxygen therapy (11).

In terms of safety, a study by Lee *et al.* in South Korea indicated that prior use of ICS did not increase the likelihood of SARS-CoV-2 positivity. Moreover, the dosage and type of ICS did not impact the positivity rate (*18*). Meta-analyses have also found that use of ICS, either as monotherapy or in combination with bronchodilators, does not influence the risk of SARS-CoV-2 infection (*19*). Therefore, antibodies are produced in patients with asymptomatic to mild COVID-19, even those on high-dose ICS, with no significant difference in antibody titers according to whether or not these agents are used. This finding suggests that patients with asthma can safely use high-dose ICS containing ciclesonide if they contract COVID-19.

The study has several limitations. The first is its small sample size. When planning the RACCO study, the number of infected individuals in Japan was not particularly high, so the study focused on a smaller number of cases that allowed the results to be delivered more rapidly. The present analysis is based on sera samples that were carefully selected in that study because we believed that there was an association between inhaled ciclesonide and antibody titers. The second limitation is the uncertainty regarding the dosage of ciclesonide needed for patients to achieve intracellular concentrations equal to the concentrations found to induce antiviral effects in previous studies. In basic research, ciclesonide powder is directly dispersed onto infected cells. However, we used inhaled ciclesonide, and there have been no definitive experiments to confirm how much of the drug is absorbed through the respiratory mucosa after inhalation and whether an adequate amount reaches the infected cells. Therefore, in this study, we used high-dose ciclesonide to ensure an adequate dosage. The third limitation is the choice of the primary endpoint. The RACCO study focused on CT images obtained at day 8 rather than clinical symptoms. A relationship between worsening on imaging and progression to severe disease has not been clearly established, making this assessment complex. There have been several studies of ciclesonide but the endpoints have varied.

It is reasonable to conclude that high doses of inhaled ciclesonide do not affect antibody production even in patients with asymptomatic to mild COVID-19. Furthermore, we found no significant difference in antibody titers according to whether or not ciclesonide was inhaled. Therefore, in the event of COVID-19 infection in patients with asthma, it can be inferred that high-dose ICS are safe to use.

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# **Original** Article

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## Elevated indoleamine 2,3-dioxygenase activity is associated with endothelial dysfunction in people living with HIV and ROS production in human aortic endothelial cells *in vitro*

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SUMMARY The precise role of indoleamine 2,3-dioxygenase (IDO) in cardiovascular diseases (CVD) among people living with HIV (PLWH) is still under debate, despite recognized links. This study aimed to investigate the impact of elevated IDO activity on endothelial dysfunction in PLWH. A total of 38 PLWH, who had not previously received anti-retroviral therapy (ART), were enrolled in the study. These participants were monitored for 36 months following the initiation of ART. Measurements including plasma levels of IDO activity, markers of endothelial dysfunction, inflammatory factors, and lipids. In vitro, human aortic endothelial cells (HAEC) were exposed to interferon- $\gamma$ , an IDO inhibitor, a kynurenine 3-hydroxylase (KMO) inhibitor, as well as different concentrations of kynurenine. Pre-ART, PLWH demonstrated notably elevated plasma concentrations of soluble intercellular adhesion molecule 1 (sICAM-1), soluble vascular cell adhesion molecule 1(sVCAM-1), and IDO activity in comparison to healthy controls. Post-ART, both IDO activity and sICAM-1 levels experienced a significant decrease, with IDO activity reaching levels comparable to those observed in healthy controls. Furthermore, a positive correlation was observed between IDO activity and sICAM-1 (p = 0.0002), as well as sVCAM-1 (p < 0.0001) before ART. In vitro, the augmentation of kynurenine concentration in the medium and the induction of IDO expression in HAEC resulted in increased production of reactive oxygen species (ROS), with minimal impact on endothelial dysfunction. From these findings, it can be concluded that long-term ART has the potential to restore the heightened IDO activity observed in PLWH. The overexpression of IDO primarily influences the expression of ROS in HAEC.

*Keywords* Tryptophan metabolism, indoleamine 2,3-dioxygenase, endothelial dysfunction, human immunodeficiency virus (HIV)

### 1. Introduction

Cardiovascular disease (CVD) is a commonly occurring chronic complication in people living with HIV (PLWH) (1,2). The introduction of widespread antiretroviral therapy (ART) has yielded a significant enhancement in the life expectancy of PLWH, which may potentially contribute to an increased prevalence of CVD among this demographic (3). Several factors have been associated with the increased occurrence of CVD (4,5), encompassing hypertension, hyperlipidemia, smoking, alcohol abuse, diabetes, physical inactivity, overweightness, unhealthy lifestyle behaviors, HIV infection, and the use of ART medications, among others.

Tryptophan (Trp), an indispensable amino acid, plays a crucial role in protein synthesis, as well as the production of 5-hydroxytryptamine and melatonin in the human body. The kynurenine (Kyn) synthesis pathway (KP) is responsible for the metabolism of approximately 90% of Trp (6). Indoleamine 2-3 oxygenase (IDO) serves as the catalyst for the initial and rate-limiting step in the degradation of Trp through the KP, resulting in the generation of immunomodulatory tryptophan metabolites and subsequent suppression of immune function.

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Recent research indicates that IDO may play a role in the development of atherosclerosis through multiple mechanisms (7,8). In non-HIV populations, there have been reports of decreased levels of serum Trp and increased Kyn/Trp ratios (K/T ratios) in those with coronary heart disease (9). Additionally, elevated plasma Kyn levels have been linked to acute myocardial infarction in patients with stable angina pectoris (10,11).

Overexpression of IDO activity is frequently observed in PLWH (12), with the HIV-1 Tat protein directly stimulating IDO production in human monocyte-derived dendritic cells (13). The potential correlation between this modification and the increased occurrence of CVD in PLWH is still a matter of discussion, necessitating additional research into the underlying mechanisms. Therefore, this study was conducted to investigate the association between the KP and endothelial dysfunction, a crucial initial step in CVD development.

### 2. Materials and Methods

### 2.1. Study population

In this study, a total of 50 PLWH and 18 HIV-negative healthy individuals as controls were recruited. The inclusion criteria required a positive HIV test, age above 16 years, and no prior ART exposure. Exclusion criteria included a history of opportunistic infections, cancer, pregnancy, hypertension, diabetes, or known CVD. At the time of enrollment, whole blood samples (10 mL) were obtained from both healthy controls and PLWH. After a three-year period of ART, 38 PLWH were effectively monitored and subsequently provided with 10 mL whole blood samples. These samples were subsequently employed to evaluate lipid profiles, HIV RNA, inflammatory markers, tryptophan, and tryptophan metabolites. All subjects gave their informed consent for inclusion before they participated in the study. The study was conducted in accordance with the Declaration of Helsinki, and the protocol was approved by the Ethics Committee of Shanghai Public Health Clinical Center (2016-S028-01).

2.2. Quantification of lipid profiles, HIV RNA, and CD4 and CD8 T-cell counts

Total cholesterol (TCH), high-density lipoprotein cholesterol (HDL-C), and low-density lipoprotein cholesterol (LDL-C) were determined using the ultracentrifugation Abell-Levy-Brodie-Kendall method. Triglyceride (TG) levels were measured using the glycerol lipase oxidase method. HIV-1 RNA loads were quantified using polymerase chain reaction (PCR) with the Cobas Amplicor system (Roche, Basel, Switzerland). CD4 and CD8 T-cell counts were evaluated using flow cytometry (BD Biosciences, Franklin Lakes, New Jersey). 2.3. Quantification of tryptophan and kynurenine levels

Plasma levels of tryptophan and kynurenine were quantified by high performance liquid chromatography as previously described (14). IDO activity was calculated using the plasma K/T ratio.

2.4. Quantification of plasma markers of immune activation and endothelial dysfunction

Plasma levels of markers including angiopoietin-2 (R&D Systems, Minneapolis, Minnesota), sCD14 (R&D Systems, Minneapolis, Minnesota), sCD163 (R&D Systems, Minneapolis, Minnesota), endogenous endotoxin-core antibody (EndoCAb) (Hycult Biotech, Uden, the Netherlands), LPS-binding protein (LBP) (Hycult Biotech, Uden, the Netherlands), IL-6 (Raybiotech, Norcross, America), sICAM-1/ sCD54 (R&D Systems, Minneapolis, Minnesota) and sVCAM-1/sCD106 (R&D Systems, Minneapolis, Minnesota), were determined using enzyme-linked immunosorbent assays in accordance with the manufacturers' instructions.

### 2.5. Cell Culture

Human aortic endothelial cells (HAECs) (Sciencell, Carlsbad, USA) were cultured in Endothelial Cell Medium (ScienCell, Carlsbad, USA) supplemented with 5% FBS (ScienCell, Carlsbad, USA), 1% Endothelial Cell Growth Supplement (ScienCell, Carlsbad, USA), and 1% penicillin/ streptomycin (ScienCell, Carlsbad, USA) as per the manufacturer's instructions. Cells within passages 3 to 8 were utilized and allowed to reach 70-80% confluence before exposure to various agents.

HAECs were subjected to pretreatment with 0.2% dimethyl sulfoxide (DMSO), IDO inhibitor (1-methyl-DL-tryptophan (1-MT) (Sigma-Aldrich, Stockholm, Sweden)), or KMO inhibitor (Ro61-8048) (MedChem Express, Monmouth Junction, USA), for a duration of one hour. Subsequently, the cells were stimulated with Kyn (10  $\mu$ M or 100  $\mu$ M) (Sigma-Aldrich, Stockholm, Sweden) or IFN- $\gamma$  (100 ng/ml) (PEPROTECH, Rocky Hill, USA) for 48 hours. All experiments were conducted in triplicate.

The levels of sICAM-1/sCD54 (R&D Systems, Minneapolis, Minnesota), sVCAM-1/sCD106 (R&D Systems, Minneapolis, Minnesota), Tryptophan (MyBioSource, California, USA), and Kynurenine (MyBioSource, California, USA) were quantified in the cell supernatant using ELISA. Additionally, the levels of caspase-3 and reactive oxygen species (ROS) were quantified in the cell suspension using the PE Active Caspase-3 Apoptosis Kit (BD Biosciences, Franklin Lakes, New Jersey) and the Reactive Oxygen Species Assay Kit (Yisheng, Shanghai, China) respectively, through flow cytometry.

### 2.6. Data analysis

The data analysis was conducted using IBM SPSS version 19.0 (IBM SPSS, Inc., Armonk, NY, USA), while the figures were generated using GraphPad Prism 6.0 software (GraphPad Software Inc., San Diego, CA, USA). Categorical variables were presented as n (%). For continuous data with a normal distribution, means±SD were presented, whereas non-normally distributed data were described using the median with interquartile range (IQR). The statistical tests employed in this study included the t-test and Mann-Whitney rank-sum test to identify significant differences between pre- and post-ART conditions, as well as healthy controls. Additionally, Pearson correlation and Spearman rank correlation tests were utilized to assess the relationship between IDO activity and other variables. Statistical significance was determined by *p*-values less than 0.05.

### 3. Results

3.1. Demographic and clinical characteristics of the study participants

The demographic and clinical characteristics of PLWH who have not yet received ART and healthy controls are presented in Table 1. Out of the 38 PLWH included in the study, a substantial majority of 89.5% (34/38) were male, with a median age of 32. The majority of PLWH were prescribed a regimen consisting of two nucleoside reverse transcriptase inhibitors (NRTIs) in combination with one nonnucleoside reverse transcriptase inhibitor (NNRTI).

3.2. Characteristics changes in PLWH at pre- and on-ART

After ART, all PLWH achieved viral suppression (< 50 copies/ $\mu$ L), accompanied by a significant increase in the CD4/CD8 ratio (0.42 (0.32-0.51) *vs.* 0.74 (0.51-0.97), *p* < 0.001), indicating an improvement in immune function. However, there was no significant change observed

### Table 1. Study participants

Characteristics	PLWH ( <i>n</i> = 38)	HIV-negative controls $(n = 18)$
Age, year, median (year)	32 (26-50)	31 (26-41)
Male sex, No. (%)	34 (89.47)	12 (66.67)
CD4 T-cell count (cells/µL)	357 (264-527)	N/A
CD4/CD8 ratio	0.42 (0.32-0.51)	N/A
ART regimen (No, %)		
TDF+3TC+EFV	32 (84.2)	N/A
TDF+3TC+LPV/r	2 (5.3)	N/A
AZT+3TC+EFV	3 (7.9)	N/A
TDF+3TC+RAL	1 (2.6)	N/A

Abbreviations: PLWH, people living with HIV; TDF, tenofovir disoproxil fumarate; 3TC, lamivudine; EFV, efavirenz; LPV/r, lopinavir/ritonavir; RAL, raltegravir. in the CD4 cell count (357 (264-527) cells/ $\mu$ L vs. 400 (307-530) cells/ $\mu$ L, p = 0.117). Prior to initiating ART, PLWH exhibited higher plasma Kyn concentrations and IDO activity compared to healthy control. However, after three years of ART, these levels were reduced to a similar extent as observed in healthy controls (Figure 1). Additional information pertaining to changes in lipid profiles, markers of endothelial dysfunction, microbial translocation, and chronic inflammation following ART in PLWH can be found in Table 2.

3.3. Plasma Kyn concentration and IDO activity was associated with endothelial dysfunction in PLWH

Prior to ART, PLWH displayed notably higher plasma concentration of sICAM-1 and sVCAM-1 compared to the control group of healthy individuals (217.00 ng/mL (156.80-241.80 ng/mL) vs. 187.90 ng/mL (117.70-209.40 ng/mL), p = 0.0317; 712.40 ng/mL (416.70-1168.00 ng/mL) vs. 467.80 ng/mL (284.40-522.70 ng/mL), p= 0.0052; for sICAM-1 and sVCAM-1 respectively). Post-ART, a significant decrease was observed in plasma sICAM-1 concentration, bringing it in line with the level observed in the healthy control group, while the sVCAM-1 level remained unchanged (Figure 2A). The plasma level of ANG-II in PLWH, both before and after ART, was found to be similar to that of healthy controls. However, a decrease in plasma ANG-II concentration was observed after ART (255.90 pg/mL (206.80-454.60 pg/mL) vs. 241.40 pg/mL (166.60-289.70 pg/mL), p = 0.0075) (Figure 2A).

Prior to ART, IDO activities and plasma Kyn concentration were positively correlated with plasma sVCAM-1 and sICAM-1 concentration in PLWH. However, the correlation between IDO activities and sICAM-1 lost significance after ART. Neither plasma Kyn concentration nor IDO activities showed any correlation with Ang-II in PLWH (Figures 2B and 2C). Furthermore, the changes in IDO activities and plasma Kyn concentration were positively correlated with the concentrations of sVCAM-1, sICAM-1, and Ang-II (Figure S1, *http://www.ddtjournal.com/action/getSupplementalData.php?ID=172*).

3.4. Kyn and IDO activity was not associated with lipids, microbial translocation, and chronic inflammation markers

A significant increase in plasma HDL levels was observed after ART (0.94 mmol/L (0.80-1.08 mmol/L) pre-ART vs. 1.08 mmol/L (0.90-1.30mmol/L) post-ART; p=0.0082). However, these levels remained lower than those observed in healthy controls (1.08 mmol/L (0.90-1.30 mmol/L) vs. 1.24 mmol/L (1.11-1.57 mmol/L); p=0.0148). Plasma levels of TG, TCH, and LDL were comparable to those of the healthy controls, both before and after ART. No significant correlations were found



Figure 1. The impact of ART on plasma concentrations of Trp (A), Kyn (B), and IDO activity (C) in PLWH. ns, not significant; \*p < 0.05; \*\*p < 0.01; \*\*\*p < 0.001; \*\*\*p < 0.001. Abbreviations: Trp, Tryptophan; Kyn, Kynurenine; IDO activity, Kyn/Trp ratio

Table 2. Characteristics ch	anges in PLWI	I pre- and on-ART
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Characteristics	Pre-ART	On-ART	<i>p</i> value
CD4 T-cell count (cell/µL)	357.50 (264.80-527.50)	399.50 (307.80-530.80)	0.117
CD4/CD8 rate	0.42 (0.32-0.51)	0.74 (0.51-0.97)	< 0.001
sICAM-1 (ng/mL)	217.00 (156.80-241.80)	172.00 (135.60-231.10)	0.008
sVCAM-1 (ng/mL)	712.40 (416.70-1168.00)	668.50 (578.90-778.50)	0.278
ANG-II(pg/mL)	255.90 (206.80-454.60)	241.40 (166.60-289.70)	0.008
TG (mmol/L)	1.15 (0.88-1.55)	1.51 (0.90-2.07)	0.086
TCH (mmol/L)	4.21 (3.35-5.11)	4.12 (3.47-4.98)	0.468
LDL (mmol/L)	2.40 (1.77-3.2)	2.73 (2.26-3.45)	0.086
HDL (mmol/L)	0.94 (0.80-1.08)	1.08 (0.90-1.30)	0.009
Tryptophan (µmol/L)	23.68 (23.57-31.87)	26.96 (23.31-29.35)	0.145
Kynurenine (µmol/L)	1.14 (1.01-1.51)	0.84 (0.73-0.92)	< 0.001
IDO Activity (nM/µM)	44.32 (34.72-53.56)	31.10 (28.76-34.81)	< 0.001
Endocab (MMu/mL)	72.30 (34.88-111.50)	169.50 (89.80-293.70)	0.001
LBP (ng/mL)	5623.00 (3362.00-7936.00)	5896.00 (5047.00-7289.00)	0.24
sCD14 (ng/mL)	1450.00 (490.20-2111.00)	1514.00 (1218.00-1829.00)	0.396
sCD163 (ng/mL)	774.30 (578.30-1081.00)	712.10 (579.20-895.90)	0.236
IL-6 (pg/mL)	77.39 (23.93-124.10)	27.67 (14.33-47.08)	0.005

Abbreviations: ART, anti-retroviral therapy; sICAM-1, soluble intercellular adhesion molecule 1; sVCAM-1, soluble vascular cell adhesion molecule 1; ANG-II, angiopoietin-2; TG, triglyceride; TCH, total cholesterol; LDL, low-density lipoprotein; HDL, high-density lipoprotein; IDO, indoleamine 2,3-dioxygenase; EndoCAb, endogenous endotoxin-core antibody; LBP, LPS-binding protein; IL-6, interleukin-6.

between TG, TCH, HDL, LDL, and plasma Kyn levels or IDO activity.

Plasma concentration of sCD14 and sCD163 were found to be significantly elevated, while plasma IL-6 levels were comparable, and plasma EndoCAb and LBP level were lower among untreated PLWH compared to healthy controls. However, apart from sCD14, which demonstrated a significant correlation with IDO activity prior to the initiation of ART ( $R^2 = 0.154$ , p = 0.01), there is no observed correlation between plasma Kyn concentration or IDO activity and plasma levels of LBP, sCD163, EndoCAb, and IL-6, both before and after ART in PLWH. 3.5. Activation of KP induces ROS production in HAEC *in vitro* 

To further elucidate the impact of IDO activity on endothelial cells, HAECs were pretreated with 1-MT or Ro61-8048 for 1 hour prior to stimulation with 100 ng/mL IFN- $\gamma$  for 48 hours. Our observations revealed that IFN- $\gamma$  effectively induced KP activation in HAECs, an effect that could be mitigated by the administration of 1-MT and Ro61-8048 (Figure 3A). Furthermore, these interventions yielded comparable results in terms of ROS production and IDO activity, while exerting minimal impact on sICAM-1, sVCAM-1, and caspase3



Figure 2. The impact of ART on endothelial dysfunction and its correlation with the metabolites of tryptophan. A tripartite panel representation delineating the impact of ART on plasma concentrations of sVCAM-1, sICAM-1, and ANG-II, sequentially from left to right (A). A series of three scatter plots demonstrating the relationships between plasma KYN levels and the respective plasma concentrations of sVCAM-1, sICAM-1, and ANG-II, arranged from left to right (B). A series of three scatter plots demonstrating the associations between IDO activity and the corresponding plasma concentrations of sVCAM-1, and ANG-II, sequentially from left to right (C). ns, not significant; \*p < 0.05; \*\*p < 0.001; \*\*\*p < 0.001; \*\*\*p < 0.001. Abbreviations: sICAM-1, soluble intercellular adhesion molecule 1; sVCAM-1, soluble vascular cell adhesion molecule 1; ANG-II, Angiopoietin-2; Kyn, Kynurenine; IDO, indoleamine 2,3-dioxygenase.



**Figure 3.** The impact of tryptophan and its metabolic products on HAEC in vitro. A set of three bar graphs, arranged from left to right, depicting the effects of IFN- $\gamma$ , 1-MT, and Ro61-8048 on tryptophan concentration, kynurenine concentration, and Kyn/Trp ratio in HAEC *in vitro*, respectively (**A**). A set of four bar graphs, arranged from left to right, depicting the effects of IFN- $\gamma$ , 1-MT, and Ro61-8048 on sVCAM concentration, sICAM concentration, ROS production, and caspase-3 activity in HAEC in vitro, respectively (**B**). A set of four bar graphs, arranged from left to right, depicting the effects of 10  $\mu$ M and 100  $\mu$ M KYN on sVCAM concentration, sICAM concentration, ROS production, and caspase-3 activity in HAEC in vitro, respectively (**C**). ns, not significant; \*p < 0.05; \*\*p < 0.01; \*\*\*p < 0.001; \*\*\*p < 0.001. Abbreviations: Kyn/Trp ratio, kynurenine/tryptophan ratio; sICAM-1, soluble intercellular adhesion molecule 1; sVCAM-1, soluble vascular cell adhesion molecule 1.

(Figure 3B).

In order to provide additional evidence for the hypothesis that KP activation may initiate the generation of ROS, we exposed HAECs to exogenous Kyn at concentrations of 10  $\mu$ M or 100  $\mu$ M for a duration of 48 hours. It was observed that Kyn had the ability to

induce the production of ROS, with the intensity of ROS production increasing proportionally with higher concentrations of Kyn. However, it is worth noting that both interventions had minimal impact on the production of sICAM-1 and sVCAM-1, as well as HAEC apoptosis, as depicted in Figure 3C.

### 4. Discussion

Endothelial dysfunction plays a crucial role in the development of atherosclerosis and other vascular disorders (15). Understanding the factors involved in HIV-related endothelial dysfunction is vital for the effective long-term management of AIDS. Our study revealed a significant association between IDO activity, plasma Kyn concentration and endothelial dysfunction in PLWH. Additionally, we confirmed that activation of KP in increased production of ROS in HAECs *in vitro*. These findings suggest that targeting this pathway could potentially reduce the risk of CVD in individuals with HIV.

In our study, we noted a notable increase in markers of endothelial dysfunction in PLWH who had not received ART, as compared to healthy controls. After ART, only levels of sVCAM-1 remained higher in PLWH compared to the healthy controls. Similarly, prior to ART, there was a significant elevation in IDO activity and plasma KYN concentration, which returned to normal levels after treatment. Consistent with our previous findings and those of Routy et al., it was observed that IDO activity did not fully recover even after antiviral therapy (12,16). The observed disparity between our present findings and prior research can be ascribed to variances in the demographics of the study cohorts. Specifically, the individuals included in this investigation displayed comparatively elevated CD4 levels, which previous studies have established to have an inverse association with IDO activity (12). Interestingly, a previous study has documented that PLWH who exhibited persistently low CD4 levels after receiving treatment exhibited a notably heightened susceptibility to cardiovascular events (17). This finding may partially account for the heightened association observed between pre-treatment IDO activity and plasma Kyn levels, as well as endothelial damage biomarkers in our investigation, which diminished following treatment. Subsequent research endeavors should prioritize the inclusion of populations with lower CD4 levels to gain further insight into these associations.

Both the beneficial and detrimental effects of CVD on Trp and its metabolites have been documented (18). Zhang et al. (8) determined the atheroprotective effects of 3-hydroxybutyrate anthranilic acid, while Kynurenic acid and aryl hydrocarbon receptor were found to inhibit the immune response, thereby enhancing the stability of carotid plaques (19). Conversely, Wang et al. argued that 3-hydroxyanthranilic acid activated the nuclear factor-kappa B transcription factor, leading to increased expression of matrix metallopeptidase 2 in vascular smooth muscle cells and contributing to AngIIinduced abdominal aortic aneurysm *in vivo* (20). K/ ratio demonstrated a positive association with BMI, LDL, TG, carotid intima-media thickness, and abdominal circumference, while exhibiting a negative association with HDL (10,21,22). Furthermore, increased plasma Kyn concentrations were found to be positively associated with the likelihood of experiencing an acute coronary event among older individuals (23). In the context of this study, IDO activity exhibited a positive correlation with plasma sICAM-1 and sVCAM-1 levels in ART-naïve PLWH. These findings are consistent with previous research conducted on PLWH (11,24).

Based on the aforementioned studies, it appears that the overexpression of IDO may contribute to endothelial dysfunction and have detrimental effects on CVD. However, an inexplicable phenomenon was observed in vitro, where the activation of IDO and the introduction of exogenous Kyn had minimal impact on endothelial dysfunction and apoptosis in HAEC. Interestingly, in hypercholesterolemic Apo $E^{-7}$ mice, the deficiency of IDO resulted in a significant increase in lesion size and a decrease in blood levels of IL10 (25). Furthermore, the inhibition of IDO led to heightened vascular inflammation and an upregulation of monocyte chemoattractant protein-1 and sVCAM-1 expression in ApoE<sup>-/-</sup> mice (26). Regulatory T cells and KP exhibit reciprocal regulation within the vessel wall, thereby facilitating vascular tolerance mechanisms and mitigating inflammation and atherosclerosis (27).

Our investigation further revealed that the activation of IDO and elevation of Kyn concentration resulted in the generation of ROS in HAEC in vitro, without causing any endothelial dysfunction. Furthermore, the treatment of mouse and human myoblasts with Kyn led to a two-fold increase in ROS levels (28). ROS, functioning as a subcellular messenger in signal transduction pathways, has been found to exert both advantageous and detrimental effects on the cardiovascular system (29). Under normal physiological conditions, the production of ROS at low levels has been observed to play a regulatory role in gene expression, excitation-contraction coupling, cell growth, migration, differentiation, and programmed cell death. However, in pathological scenarios, ROS can lead to oxidative modifications of crucial cellular macromolecules (29). Lubrano's research (30) has demonstrated that elevated levels of ROS can directly enhance the production of lectin-like oxidized low-density lipoprotein. This increase in LOX-1 has been linked to apoptosis induction in various cell types when present in high concentrations, while low concentrations have been associated with cell proliferation.

Our previous research has demonstrated the prevalence of dyslipidemia, microbial translocation, and chronic inflammation in PLWH (31,32), which have been considered to be associated with endothelial dysfunction (15, 33). In Apuet<sup>-/-</sup> mice, inhibition of IDO resulted in an increase in the level of TCH and the ratio of VLDL/HDL (26). The involvement of Kyn metabolites in regulating lipid metabolism, thermogenic

gene expression, and anti-inflammatory gene expression in adipose tissue has been observed through the activation of G protein-coupled receptor Gpr35 or aryl hydrocarbon receptor/signal transducer or transcription 3/interleukin-6 signaling (34,35). Catabolism of tryptophan (Trp) plays a significant role in the precise regulation of intestinal physiology (36), and it has also been linked to CVD (37). Numerous studies have reported the involvement of IDO in the pathophysiology of various inflammatory conditions, such as infection, allergy, autoimmunity, chronic inflammation, and inflammatory neurological diseases (38,39). Based on the aforementioned evidence, it is justifiable to postulate that KP might potentially contribute to the development of endothelial dysfunction through its influence on lipid metabolism, microbial translocation, or chronic inflammatory response. However, this conjecture was promptly refuted as our study revealed minimal association between IDO activity and plasma lipid levels, as well as plasma markers for inflammation or microbial translocation.

Our study is subject to several limitations and deficiencies. Firstly, the majority of participants were young and middle-aged men, thus limiting the generalizability of our findings to women and the elderly. Secondly, the assessment of atherosclerosis was incomplete as we did not measure carotid artery intimamedia thickness, a crucial parameter for evaluating this condition. Thirdly, the utilization of the K/T ratio as a representative measure of IDO activity may lead to erroneous conclusions, as it does not fully capture the complexity of IDO activity. Additionally, the verification of our results through *in vitro* experiments using HAEC may be deemed less persuasive compared to *in vivo* studies.

In summary, long-term ART has the potential to restore normal levels of IDO activity that is elevated due to HIV infection. The activity of IDO is found to be positively associated with endothelial dysfunction in PLWH. The excessive expression of IDO primarily impacts the production of ROS, while its direct influence on endothelial dysfunction in HAEC is limited.

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# **Original** Article

# Effects of plant-based copper nanoparticles on the elimination of ciprofloxacin

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SUMMARY Ciprofloxacin (CIP) is frequently detected in the environment and causes the emergence of drugresistant bacteria. High levels of CIP in the environment are also harmful to humans and animals. Therefore, effective elimination of CIP is required. In this study, plant-based copper nanoparticles (CuNPs) have been fabricated for the purpose of eliminating CIP. Aqueous extracts of 6 plants were compared for their phytochemicals and reducing activity. Among all the extracts, Garcinia mangostana extract (GM) was the most potent with the highest total phenolic compounds, flavonoids, tannins, terpenoids, and reducing activity. CuNPs synthesized using GM (GM-CuNPs) were characterized using UV-VIS spectroscopy and dynamic light scattering. The results showed that the maximum absorption of GM-CuNPs was at 340 nm. The average size of GM-CuNPs is in the nanoscale range of  $159.2 \pm 61$  nm, with a narrow size distribution and a negative zeta potential of  $-4.13 \pm 6.97$  mV. The stability of GM-CuNPs is not solely due to their zeta potential but also phytochemicals in the extract. GM-CuNPs at 25 mM showed the highest efficiency of 95% in removing CIP from aqueous medium pH 6-7 at 25-35°C within 20 min. The results indicated that the electrostatic attraction between the negative charge of GM-CuNPs and the positive charge of CIP controlled the drug adsorption on the nanoparticles. In conclusion, the developed GM-CuNPs have excellent CIP removal efficiency. These synthesized GM-CuNPs are expected to be environmentally friendly for the removal of antibiotics and other drugs.

*Keywords* Garcinia mangostana, eco-friendly, green synthesis, metal nanoparticles, plant extract

### 1. Introduction

It is known that many drugs in humans and animals are eliminated through urine and feces, and released into the wastewater system (1). Ciprofloxacin (CIP), one of the most widely used antibiotics for treatment of bacterial infections in both humans and animals, is secreted in the urine in its original form (2). Therefore, large amounts of CIP are found in wastewater systems. In addition, pharmaceutical manufacturers and hospitals are also the most important sources of contaminated wastewater. This leads to increase contamination in surface and groundwater, and is extremely harmful to the environment and human health (3,4). It has been reported that CIP contamination of daily drinking water can cause mild nausea, vomiting, headaches, and diarrhea. Higher CIP contamination can cause more serious adverse effects, such as thrombocytopenia and acute renal failure. Additionally, the presence of CIP in water supplies has resulted in the development of antibioticresistant bacteria (5), resulting in an increase in health problems. This is because treating drug-resistant bacteria is difficult, high cost, and requires different drug or a higher dose. Therefore, the effective removal of CIP is required. Technology using metal nanoparticles appears to be a convenient alternative (6) and promising for the removal of CIP by an adsorption mechanism (7,8). Several metal-based techniques have been developed to synthesize metal nanoparticles, such as chemical laser ablation, chemical reduction, coprecipitation, and wave-

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assisted processes (9, 10). However, these methods are expensive and not environmentally friendly. This is because many harmful by-products are released into the environment. The limitations of these methods have given rise to new technologies of plant-based synthesis, by biomolecules are used instead. Plant-assisted synthesis is environmentally friendly, cost effective, and can help overcome all limitations. It has been proven to be an effective and highly reliable technique in preparing metal nanoparticles. However, to obtain the desired metal nanoparticles by plant-based synthesis, it is necessary to use biomaterial with high reducing power. Several reports indicate that certain plant-derived phenolic compounds such as flavonoids and tannins possess antioxidant activity and reducing power. In this study, the phytochemical contents of 6 potential plant extracts, Ageratum conyzoides, Emilia sonchifolia, Garcinia mangostana, Sesbania grandiflora, Tridax procumbens, and Xanthium strumarium were compare and the most effective plant extract was selected for synthesis of copper nanoparticles (CuNPs).

A. conyzoides, an allelopathic plant in the family Asteraceae, is a good source of antioxidants due to its high content of total phenolic compounds and flavonoids (11). Different parts of this plant have been used in folk medicine for decades. E. sonchifolia, T. procumbens and X. strumarium are members of the Asteraceae family. E. sonchifolia has previously been reported to inhibit the formation of hydroxyl radicals and superoxide radicals (12). Its major active ingredient is emiline, which is an alkaloid with antioxidant potential and neuroprotective effect against corticosterone-induced apoptosis in cancer cell lines (13). Up to date, there are no reports of using this compound as a reducing agent in the production of metal nanoparticles. Therefore, it remains a challenge as a primary plant in this experiment. T. procumbens contains alkaloids, carotenoids, flavonoids, fatty acids, steroids, phytosterols, tannins, and has high antioxidant activity (14). X. strumarium has been reported to have high antioxidant activity and reducing power (15). The antioxidant properties of this plant are related to the levels of alkaloids, phenolic acids, diterpenes, saponins, glycosides, essential oils and phytosterols (16,17). G. mangostana is a tropical perennial with edible fruits. It belongs to the Guttiferae family. This plant has been used in traditional and folk medicine systems in Southeast Asia to treat cystitis, dysentery, gonorrhea, inflammatory skin and abnormalities of urinary tract and to increase the efficiency of wound healing (18). It has been documented as a reducing agent for the fabrication of metal nanoparticles due to its phytochemical composition in the aqueous extract (19). S. grandiflora, which belongs to the family Leguminoseae, is an important agroforestry as a food source for humans, cattle, and goats in Thailand. Its bark extract possesses strong antibacterial activity (20). Antioxidant activity and the use in the plant-based synthesis of silver nanoparticles

from this plant have also been reported (21). The main objective of this study was to synthesize CuNPs for the elimination of CIP. In this synthesis, a green biosynthesis method was used. The most effective plant extracts from these six potential plants were used as reducing agents. The synthesized CuNPs were studied for their physicochemical characterization and CIP removal efficiency in various water conditions.

### 2. Materials and Methods

### 2.1. Materials

Copper sulfate pentahydrate, ferric chloride, potassium ferricyanide, sodium nitrate and sodium nitrite were purchased from Ajax Finechem (New South Wales, Australia). CIP hydrochloride and gallic acid were from Bio Basic Inc. (Markham, Canada). Epigallocatechin gallate, quercetin, and 2,2-diphenyl-1-picrylhydrazyl hydrate (DPPH) were from Sigma-Aldrich (Steinheim, Germany). Hydrochloric acid (HCl) was from QRëC (Auckland, New Zealand. Folin-Ciocalteu reagent was from Loba Chemie (Mumbai, India). Aluminum chloride hexahydrate, bromophenol blue, and calcium carbonate were from Kemaus (New South Wales, Australia). Trichloroacetic acid and vanillin were from Merck (Damstadt, Germany). Sodium hydroxide (NaOH) was from RCI Lab-scan (Bangkok, Thailand). All other solvents and chemicals were of analytical grade.

### 2.2. Plants and extract preparation

A. conyzoides, E. sonchifolia, G. mangostana, S. grandiflora, T. procumbens and X. strumarium were collected from northern Thailand during July-December 2021. The plants were authenticated by herbalists from Rajamangala University of Technology Lanna, Thailand. All specimens are preserved at the university's herbarium. Fresh leaves of each plant were detached from the stems and washed thoroughly with tap water and deionized water, respectively. After that, the whole leaf material was dried in a hot air oven at 50°C until a constant weight was achieved. Then pulverized with an electric blender. The plant powder sieved through a 20-mesh sieve was vacuum-wrapped in a plastic bag and stored in a deep freezer at  $-20^{\circ}$ C for further use.

To prepare the aqueous extract of each plant, 100 mg of dried plant powder was boiled with 10 mL of distilled water at 85°C for 30 min. Afterwards, the mixture was cooled to room temperature. It was then filtered through a 0.22-micron nylon syringe filter. The obtained filtrate was used for further studies.

### 2.3. Phytochemical studies

Freshly prepared aqueous extracts of each plant were

examined for total phenolic content (TPC), flavonoids and tannins. TPC content was determined by mixing 20  $\mu$ L of the extract with 100  $\mu$ L Folin-Ciocalteu reagent in 1,980  $\mu$ L deionized water. After 5 min of incubation, 300  $\mu$ L of 7% sodium nitrate solution was added and mixed well. The resulting mixture was further incubated for 60 min in the dark at room temperature. It was then measured for the absorbance at 765 nm. TPC was determined using a calibration curve made from various concentrations of gallic acid and expressed as  $\mu$ g of gallic acid equivalent (GAE) per mg of extract.

The total flavonoid content (TFC) of the samples was determined by mixing 20  $\mu$ L of extract with 380  $\mu$ L deionized water, then 100  $\mu$ L of 5% sodium nitrite solution was added. After 5 min of incubation, the mixture was added with 100  $\mu$ L of 10% aluminum chloride solution and allowed to stand for 6 min at room temperature. Finally, 400  $\mu$ L of 1M NaOH solution was added. After completion of 15 min of incubation in the dark, absorbance was measured at 415 nm. TFC was determined using a calibration curve generated from quercetin at various concentrations and expressed as  $\mu$ g quercetin equivalent (QE) per mg of extract.

For determination of total tannin content (TTC), 250  $\mu$ L of extract was vigorously shaken with 450  $\mu$ L of 1% vanillin reagent. After 5 min of incubation, the mixture was mixed with 300  $\mu$ L of concentrated HCl and then was incubated for 30 min at ambient temperature. Finally, the color of the solution changed to red. The absorbance was measured at 500 nm. TTC was determined using a calibration curve generated from epigallocatechin gallate at various concentrations and expressed as  $\mu$ g of epigallocatechin gallate equivalent (EE) per mg of extract.

### 2.4. Determination of reducing power

High reducing power is an important factor in the production of metal nanoparticles. Measurement of the reducing potential of plant extracts was performed using an assay according to a previously described method (22) with some modifications. Briefly, the diluted concentration of each extract (12.5, 25, 50 and 100 µg/ mL) and the standard antioxidant, gallic acid was mixed with 1.0 mL of 0.2 M sodium phosphate buffer (pH 6.6) and 1.0 mL of 1% potassium ferricyanide. After vigorously shaking, the mixture was incubated at 50°C in a dry bath for 20 min, then 1.0 mL of 10% trichloroacetic acid was mixed and centrifuged at 4,000 rpm for 10 min. An aliquot of 0.5 mL of supernatant was diluted with the same volume of distilled water. Finally, 50 µL of 0.1% ferric chloride was added. After incubation for 10 min, the absorbance was determined at 700 nm against a reagent blank. Distilled water was used as a blank. The reducing power was determined using a calibration curve made from gallic acid at various concentrations and expressed as µg of GAE per mg of extract.

### 2.5. Synthesis of CuNPs

The synthesis of CuNPs was conducted according to a previously described method (23) with some modification. Briefly, copper sulfate was dissolved in deionized water to obtain a final concentration of 50 mM. After that, 200  $\mu$ L of extract and 300  $\mu$ L of deionized water were vigorously mixed with 500  $\mu$ L of copper sulphate solution prior to refluxing at 80°C for 30 min in a dry incubator to stimulate the reduction of metal salt. The color of the dispersion gradually changed from colorless to yellow and then to dark orange. This indicates the formation of particles in the nanoscale range (24). The solution was allowed to cool to ambient temperature and used for further studies.

### 2.6. Characterization of CuNPs

The formed CuNPs was characterized using a V-1200 spectrophotometer with UV-Professional analysis software (Dshing Instrument Co., Ltd., ZhuHai, China). The UV-visible spectrum of CuNPs enclosed in 1.5 mL disposable PMMA cuvettes (BRAND GmbH & Co. KG, Wertheim, Germany) was detected in the wavelength range 320 to 800 nm, operating at a resolution of 5 nm. The extracts and copper sulfate solution were also examined under the same conditions.

The lyophilized CuNPs were resuspended in 1 mL deionized water. The size and size distribution (PdI) as well as zeta potential of CuNPs were determined using a Nano-ZS zetasizer (DTS 1060, Malvern Instruments Ltd., Worcestershire, UK) based on the principle of dynamic light scattering (25,26). The results were automatically calculated by the software according to the instrument manufacturer's recommendations.

### 2.7. Removal efficiency of CuNPs

In this experiment, CuNPs synthesized using selected plant extracts were used. The CIP removal efficiency (CRE) of the synthesized CuNPs was examined using the CIP solutions. For the preparation of CIP solution, 25 mg of CIP hydrochloride was dissolved in 100 mL of distilled water. Factors that might affect CRE include: CuNPs concentration, contact time, pH, and temperature were mornitored. To study the effects of pH, the pH of the tested solution was adjusted by adding 0.1 N HCl or 0.1 N NaOH to achieve a pH range of 3-10.

The selected CuNPs colloidal dispersion was mixed with freshly prepared CIP sample solution and kept at room temperature for 60 min. The samples were collected at different time intervals and centrifuged at 500 rpm for 10 min. An aliquot of 250  $\mu$ L of the supernatant was mixed with 100  $\mu$ L of 20 mM acetate buffer, pH 4.1, and 375  $\mu$ L of 0.1% bromophenol blue. After incubation at room temperature for 10 min, 2.5 mL of chloroform was added and the mixture was shaken vigorously for 30 s, which resulted in the separation between chloroform and water. The yellow layer was determined at 416 nm relative to the blank. Calibration curves were generated by diluting CIP to various concentrations ( $R^2 = 0.99$ ). The CIP content was expressed in µg. The system without CuNPs was used as a control. CRE can be calculated using the following equation: CRE (%) = [(Ac - As)/Ac] × 100, where Ac is the absorbance of the control and As is the absorbance of the sample.

### 2.8. Statistical analysis

All experiments were carried out in triplicate. The obtained data were analyzed using Microsoft Excel 2016 for Windows and presented as mean  $\pm$  standard deviation (SD). The data were analyzed by one-way ANOVA and Duncan's mean comparison test at the 5% significance level.

### 3. Results

### 3.1. Phytochemicals and reducing power of extracts

The phytochemical content and reducing power of the extracts of the six plants are shown in Table 1. It can be clearly seen that the *G. Mangostana* extract had the highest TPC, TFC, and TTC with the GAE, QE, and EE values of  $98.86 \pm 1.20 \text{ mg/g}$ ,  $109.22 \pm 4.06 \text{ mg/g}$ , and  $209.67 \pm 7.29 \text{ mg/g}$ , respectively. This extract was selected for further studies.



*G. mangostana* extract was used as a reducing agent in the synthesis of CuNPs, and the obtained CuNPs were named as GM-CuNPs. The blue color of the copper sulfate solution changed to light brown after boiling the mixture at 80°C for 30 min. The different appearance and color of GM-CuNPs colloidal system are shown in Figure 1. After characterization, the results confirmed that the particle size of GM-CuNPs was in the nanoscale range with an average diameter of  $159 \pm 61$  nm and a narrow PdI value of  $0.280 \pm 0.003$ . GM-CuNPs were also found to have a negative surface charge with a zeta potential of  $-4.13 \pm 6.97$  mV.

#### 3.3. Effect of GM-CuNPs concentration on CIP removal

In this study, aqueous medium, pH 7, containing CIP at a concentration 0.25 g/L was treated with GM-CuNPs at concentrations ranging from 0.20 to 100 mM for 60 min at room temperature. The results as shown in Figure 2, indicate that the removal efficiency increased with the increase in the concentration of the synthesized GM-CuNPs. An increase in CIP removal capacity was observed from a CRE value of 24.1% to 98.2% when the GM-CuNPs concentration increased from 0.20 mM to 100 mM. The removal efficiency was highest when the concentration of GM-CuNPs was 25 mM. From these results, GM-CuNPs at a concentration of 25 mM was considered the most effective concentration to remove CIP and was selected for further study.



Figure 1. Appearance and color of *G. mangostana* extract solution (a), GM-CuNPs colloidal system (b), and copper sulfate solution (c).



Figure 2. Effect of GM-CuNPs concentration on CIP removal efficiency. Different letters indicated statistically significant values (p < 0.05).

Table 1. Phytochemical	properties and 1	reducing power	of the extracts

Plants	TPC	TFC	TTC	Reducing power
A. conyzoides	$44.88\pm0.37^{\text{b}}$	$42.19\pm0.45^{\rm b}$	$13.78\pm0.85^{\mathrm{b}}$	$234.61 \pm 4.25^{\text{b}}$
E. sonchifolia	$24.05\pm0.84^{\rm c}$	$21.49\pm0.61^{\circ}$	$2.87\pm0.15^{\rm c}$	$92.98\pm3.92^{\rm d}$
G. mangostana	$98.86 \pm 1.20^{\rm a}$	$109.22\pm4.06^{\mathrm{a}}$	$209.67 \pm 7.29^{\rm a}$	$358.82\pm2.34^{\rm a}$
S. grandiflora	$18.12 \pm 1.20^{\rm e}$	$21.78\pm1.73^{\circ}$	$1.79\pm0.60^{\circ}$	$41.61\pm2.79^{\rm f}$
T. procumbens	$15.73\pm0.21^{\rm f}$	$14.19\pm0.60^{\rm d}$	$6.84\pm2.33^\circ$	$75.45\pm3.21^{\text{e}}$
X. strumarium	$21.40\pm0.16^{\rm d}$	$23.63\pm0.70^{\circ}$	$7.49\pm0.26^{\rm c}$	$121.82\pm0.08^{\rm c}$

Data present mean  $\pm$  standard deviation of triplicate analysis. Different lowercase letters in the same column indicate statistically significant values (p < 0.05).



Figure 3. Effect of exposure time on CIP removal efficiency. Different letters indicate statistically significant values (p < 0.05).



Figure 4. Effect of pH on CIP removal efficiency. Different letters indicate statistically significant values (p < 0.05).

### 3.4. Effect of exposure time

Exposure period between GM-CuNPs and CIP for different time intervals (10, 20, 30, 40, 50, and 60 min) was studied using GM-CuNPs at a concentration of 25 mM in medium at pH 7.0 and at room temperature. CRE values were found to increase with increasing exposure time. As shown in Figure 3, the CRE value at an exposure time of 10 min was  $86.0 \pm 0.7\%$ . Then, as the exposure time increased, the CRE value gradually increased to  $91.8 \pm 1.1\%$  at an exposure time of 60 min. It is seen that CIP was rapidly removed by GM-CuNPs and achieved the highest removal efficiency in the first 20 min and then reached equilibrium.

### 3.5. Effect of pH

We hypothesized that pH might be another important factor for controlling the CIP removal process of GM-CuNPs. Therefore, the synthesized GM-CuNPs were added to the CIP aqueous solutions at various pH ranges of 3-10 to obtain a final concentration of 68 mM CIP and 25 mM GM-CuNPs. The results showed that the removal efficiency of GM-CuNPs at different pH was different as shown in Figure 4. It was found that increasing the pH value from 3.0 to 6.0 could increase the percentage of CRE from 60.4% to 92.8%, respectively. The highest CRE value in 20 min contact was obtained in the pH range of 6-7. However, when the pH was further increased, it was found that the efficiency of removal decreased.





Figure 5. Effect of temperature on CIP removal efficiency. Different letters indicate statistically significant values (p < 0.05).

CIP removal efficiency by GM-CuNPs at various temperatures are presented in Figure 5. The results showed that the CIP removal efficiency was highest at the temperature range of 25-35°C (with CRE of 95.1 and 94.5%) and slightly decreased at 15°C and 45°C (with CRE of 85.9 and 89.2%).

### 4. Discussion

Flavonoids and tannins are important phytochemical groups. The principle chemical structure of flavonoids consists of two phenyl rings connected by a linear three-carbon bond (27). Flavonoids are classified according to their substituents and the oxidation state of the heterocyclic ring into flavones, isoflavones, flavanones, flavanols, and anthocyanidin (28). Tannins are bitter nitrogen-free polyphenolic compounds that have a bitter taste. Their molecular weight ranges between 500 to 20,000. Tannins can be divided into two groups: condensed tannins or non-hydrolysable tannins such as proanthocyanidins and hydrolysable tannins such as catechin and gallic acid (29). Both flavonoids and tannins possess various pharmacological effects, such as antioxidant, antiviral, antibacterial, and antiinflammatory properties (28,30). It has been reported that the antioxidant effects of these polyphenolic compounds are due to H-atom transfer or single-electron transfer mechanisms (31,32). Therefore, these phenolic compounds have significant impact on the reducing properties of the sample, and affect the redox reactions of metal nanoparticles formulation (33,34). The results confirm that among the six plant extracts, G. Mangostana extract possessed the highest TPC, TFC, and TTC, and the highest reducing power. Therefore, this extract was selected for nanometal preparation.

During synthesis, the color of copper sulfate solution transformed from blue to light brown, confirming that all Cu<sup>2+</sup> were reduced to Cu<sup>0</sup> and GM-CuNPs were completely formed. The color of the obtained CuNPs colloidal systems is due to the fluctuation of the surface plasmon excitation in CuNPs. A wide range of colors was observed after the formation of CuNPs was completed. This variation can be caused by several factors, including reaction time and temperature. In addition, one of the most important factors is the type of reducing agent. It has been reported that when ascorbic acid (35) and Fortunella margarita extract (36) were used as a reducing agent, the colors of the obtained CuNPs colloidal systems were yellow-brown and yellowish-green, respectively. The current study used G. Mangostana extract, which has a different color from the two reducing agents reported above. Therefore, the color of colloidal GM-CuNPs obtained was light brown. However, these color changes are useful in indicating the complete formation of CuNPs. The maximum absorbance of the synthesized CuNPs is in the range approximately 320-370 nm. High absorbance indicates a high conversion of Cu<sup>+2</sup> to Cu<sup>0</sup> and results in a high concentration of CuNPs (37). Consistent with previously reported (38), they found that an absorption peak of 370 nm was observed when Cissus vitiginea extract was used. In the present study, the absorbance of the synthesized CuNPs was 340 nm, which is consistent with CuNPs synthesized using curry leaf extract (39). The particle size of GM-CuNPs was in the nanoscale range with a narrow size distribution similar to that synthesized using the aqueous extract of Carica papaya leaves as a reducing agent and copper sulfate as a precursor (40). The zeta potential is an important parameter for predicting the stability of particles in colloidal systems. In general, a zeta potential of  $\pm$  30 mV is sufficient to prevent particles from coalescing by electrostatic repulsion (41). Zeta potential values near zero indicate weak repulsive forces between particles. This results in the agglomeration of particles. Interestingly, although the zeta potential values of the synthesized CuNPs using various plant extracts were close to zero (42-44), the obtained metal nanoparticles still did not aggregate. It has been reported that some phytochemicals present in plant extracts play important roles not only as reducing agents but also particle stabilizing agents (45). Therefore, the nanoparticles can be prevented from agglomeration, and the obtained CuNPs are stable despite their low zeta potential.

Our results show that the concentration of GM-CuNPs affects the removal efficiency. It has been reported that the addition of the nanoparticles increases the number of available adsorption sites (46), thus resulting in an increase in CRE. The present study shows that CIP was rapidly eliminated by GM-CuNPs and the highest elimination with CRE of 91.8% was achieved in the first 20 min, and then equilibrium was reached. This result suggests that the increasing time leads to increased contact between the CIP molecules and the active sites of GM-CuNPs (47). After CIP molecules saturate on the active sites of the nanoparticles, the CRE value does not significantly increase because the particles have reached their full capacity. This result is in agreement with previous reports showing that the removal of antibiotics by CuNPs depends on the adsorption mechanism (48).

CIP is a zwitterionic molecule with two  $pK_a$ s;  $pK_{a1}$  of 6.1 and  $pK_{a2}$  of 8.7 (49,50), which is due to the proton predominance of the amine and carboxylic groups, respectively. The removal of CIP from water using GM-CuNPs was a pH-dependent phenomenon. It was observed that the removal process occurred along with the charge interaction adsorption mechanism. The highest CRE values within 20 min were found in the pH range of approximately 6-7, where the CIP molecule was in its zwitterionic form. In this pH range, CIP molecules possess both negative and positive charges. As a result, the interaction between GM-CuNPs and CIP is maximized. At pH lower than 6 or lower than  $pK_{a1}$  of CIP, the solution contained a high amount of H<sup>+</sup> cation, which completely interacted with the anion of GM-CuNPs. This competition resulted in a decrease in the adsorption of CIP by the active site of GM-CuNPs. At pH greater than 7, the negative salt form of CIP at  $pK_{a2}$ was dominant (51, 52) and was repelled by the negatively charged of GM-CuNPs. Therefore, the adsorption of CIP by GM-CuNPs decreased again in alkaline medium. From these results, it was concluded that the highest removal efficiency of CIP by GM-CuNPs could be obtained only in the optimum pH range of 6-7. In studies on the effects of temperature, it was found that the maximum removal efficiency could only be achieved at appropriate temperature. This can be considered that interaction between CIP molecules and GM-CuNPs is a specific reaction that occurs only at specific temperature. The decreased removal efficiency at temperatures above and below 25-35°C is confirmed to be due to the reduced attraction between the active sites of GM-CuNPs and CIP. In addition, the adsorption efficiency at higher temperatures decreased due to the high mobility of both the adsorbent, GM-CuNPs, and the adsorbate, CIP (53). It can be concluded that among the factors studied, pH plays an important role in the CRE of GM-CuNPs. Therefore, it is necessary to adjust the pH to 6-7 before the removal of CIP by this nanoparticle system.

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# **Original** Article

# FASN promotes gallbladder cancer progression and reduces cancer cell sensitivity to gemcitabine through PI3K/AKT signaling

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**SUMMARY** Lipid metabolism plays an important role in the growth and development of tumors. However, the role of lipid metabolism in gallbladder cancer (GBC) has not been clearly clarified. Here, we demonstrated that fatty acid synthase (FASN), a key enzyme in *de novo* fatty acid biosynthesis, had upregulated expression in GBC samples both at protein and mRNA levels. Analysis of clinical data indicated the association between elevated FASN expression and poorer histology grades. Furthermore, FASN activity impairment through FASN knockdown or treatment with orlistat resulted in the inhibition of cell proliferation and migration, as well as increased sensitivity to gemcitabine. Both FASN knockdown and orlistat treatment induced cell apoptosis. Mechanistically, impairment of FASN activity suppressed the activation of the PI3K/AKT signaling pathway, which led to increased cell apoptosis and sensitivity to gemcitabine. These findings were also validated through nude mouse xenograft models, thus highlighting the potential of targeting FASN as a clinical treatment strategy. Collectively, the present study underscores the crucial role of FASN in the progression of gallbladder cancer *via* the PI3K/AKT pathway.

*Keywords* GBC, orlistat, drug combination

### 1. Introduction

Gallbladder cancer (GBC) represents the most frequent malignant tumor within the biliary system (1), with a poor prognosis and a median survival of only 4-7 months (2). It was estimated that there were 115,949 cases of GBC and 84,695 deaths in 2020 (3). Despite surgery being the principal treatment method for gallbladder cancer, most patients are typically diagnosed during advanced stages, and as a result, are not eligible for radical resection (4). As a consequence, chemotherapybased comprehensive treatment represents the primary strategy for extending the survival time of patients suffering from advanced GBC. Currently, gemcitabine (GEM) constitutes one of the principal drugs employed in first-line chemotherapy regimens for gallbladder cancer. Nonetheless, its objective response rate is a mere 30%, largely due to its low sensitivity (5). It is therefore urgent to explore the molecular mechanisms that underlie tumor progression and chemotherapy resistance in GBC to pave the way for the development of novel therapeutic

approaches.

Fatty acid synthase (FASN) is the paramount lipogenic enzyme responsible for the terminal stages of de novo synthesis of fatty acids, specifically palmitate. In contrast to healthy cells, cancerous cells rely on the active production of fatty acids to construct membranes that promote cellular growth and proliferation. Additionally, palmitate can form covalent bonds with certain tumor-promoting growth factors, including Wnt, HRAS, and NRAS. These factors must undergo posttranslational modification through palmitoylation to achieve correct localization and functional efficacy (6). While FASN is known to be overexpressed in a multitude of cancer types, its expression in healthy tissues is comparably meager. FASN activity seems to coincide with the advancement of numerous human malignancies. In human cancers, FASN is excessively upregulated and linked with unfavorable prognoses, as it sustains cancer cell growth and survival. Conversely, in the majority of normal cells, FASN is markedly repressed by adequate exogenous dietary fat supplementation (7). Additionally,

it has been demonstrated that overexpression of FASN contributes to cellular resistance against genotoxic drugs, including doxorubicin and cisplatin (8,9). Another study has indicated that FASN expression is markedly upregulated in pancreatic cancer. Furthermore, this study revealed that knockdown of FASN could reverse the cellular resistance to gemcitabine induced by overexpression of PKM2 (10). Although studies have identified FASN as a potential prognostic and therapeutic target in various cancers, its role in GBC remains unknown. In this study, we will examine the expression level of FASN through tissue microarray (TMA) and quantitative RT-PCR. Furthermore, we will elucidate the functional role of FASN in the progression of GBC and its impact on gemcitabine sensitivity.

Orlistat, a compound with reactive  $\beta$ -lactones, shows irreversible FASN inhibition ability. Currently, orlistat is registered as an obesity treatment drug in several countries. Its mechanism of action involves suppression of gastrointestinal lipases, which impairs the metabolism of lipids in the gastrointestinal lumen and prevents the absorption of up to 30% of the lipids in dietary fat (11). Studies indicate that orlistat shows therapeutic effects on a variety of cancers in vivo and in vitro (12,13). It has been reported that orlistat can reverse resistance to sorafenib in liver cancer (14) and cisplatin in ovarian cancer (15) by regulating the process of fatty acid metabolism. Given its favorable safety profile and affordability, the potential role of orlistat in cancer treatment is an area of significant clinical interest. Specifically, its role in GBC and the underlying mechanisms of action warrants further exploration.

In summary, our study affirms the significance of FASN in GBC through evaluating the correlation between FASN expression in GBC tissues and clinical features. *In vivo* and *in vitro* experiments were performed to ascertain the function of FASN in the progression of GBC and its impact on sensitivity to gemcitabine.

### 2. Materials and Methods

### 2.1. Cell lines and culture

NOZ, and the SGC-996 of human GBC cell lines, obtained from colleagues of Shanghai Key Laboratory of Biliary Tract Disease Research of Xinhua Hospital, Shanghai Jiao Tong University School of Medicine, China. These cells were routinely maintained in DMEM medium (Gibco, NY, USA) supplemented with 10% fetal bovine serum (Yeasen, Shanghai, China) and ampicillin/ streptomycin (1%). All cells were cultured at 37°C, 5% CO<sub>2</sub>. Cells were routinely verified and assured for free of mycoplasma contamination.

### 2.2. Patients and specimens

48 pairs of GBC tissues and adjacent normal tissues were

collected for this study to verify expression of FASN with TMA. The patients received surgery between 2018 and 2021 in Xinhua Hospital of Shanghai Jiao Tong University School of Medicine (Shanghai, China). The complete medical records of 38 cases of these patients were available and the clinicopathologic features were retrospectively analyzed. All procedures followed were in accordance with the Helsinki Declaration (as revised in 2013). Informed consent was obtained from all patients for being included in the study.

### 2.3. Tissue microarray (TMA)

Tissue microarrays were constructed with 2 mm cores to evaluate the expression of FASN. The tissue sections were soaked in xylene for 15 min and then rehydrated through three decreased concentrations of ethanol for 2 min, and boiled in a pH 6.0 antigen retrieval solution for 20 min at 95°C. To remove endogenous peroxidase, the tissue sections were blocked with 3% hydrogen peroxide after cooling for enough time. The slides were then incubated with 10% goat serum for 60 min at room temperature. After incubation with primary and second antibodies, labeling was detected.

FASN expresses cytoplasmically. A total score (IHC score) which is obtained by multiplying proportion score (< 5%, 0; 6 to 25%, 1; 26 to 50%, 2; 51 to 75%, 3; and > 75%, 4) and intensity score (0; no staining, 1; weak intensity, 2; moderate intensity, 3; strong intensity), which represent the average intensity and proportion of positively stained cancer cells relatively. Subsequently, the 48 pairs of samples were designated as absent/low FASN expressers (IHC scores 0 to 4) or high FASN expressers (IHC scores 6 to 12) (16). A ratio was calculated to define the relative FASN expression of cancer (IHC scores) to the adjacent tissue (IHC scores): R = IHC score (cancer)/IHC score (paired normal tissue). A ratio greater than 2 was defined as relatively high expression. The evaluation of the immunohistochemical staining was independently performed by two experienced pathologists without knowledge of clinical data.

### 2.4. Plasmids, lentivirus particles and regents

The sequences of Small Interfering RNA (siRNA) were purchased from Tsingke Biotechnology Co., Ltd. (Beijing, China). siRNAs were transfected using RFect Reagent (Changzhou Biogenerating Biotechnologies corporation, Jiangsu, China) according to provided manufactures. The sequence of the siRNA is as follows: 5'-GCCGAGTACAATGTCAACATT -3'. Cells were used for further experiments at 48 h after transfection.

Short hairpin RNAs (shRNAs) against FASN were cloned into a pLKO.1 vector. HEK293T cells were co-transfected with pLKO.1 shRNA, psPAX2 and pMD2.G plasmid (gifts from Jiaxue Wu, School of Life Sciences Fudan University, Shanghai, China) at a ratio of 4:3:1. The lentiviral particles were harvested 48 h after transfection. GBC cells were infected with recombinant lentivirus transducing units in 4  $\mu$ g/mL polybrene (Beyotime, Shanghai, China). Cells were selected with puromycin for 1 week 48 after infection. The forward sequence of shRNA1 and shRNA2 are 5'-CCGGAAGCCGAGTACAATGTCAACACTCGA GTGTTGACATTGTACTCGGCTTTTTTG -3'and 5'-CCGGAACTGCTAGGTATGGAGTTCTCTCGAG AGAACTCCATACCTAGCAGTT TTTTTG-3'.

### 2.5. CCK-8 assay and colony formation assay

The cell counting kit-8 (CCK-8) (Yeasen, Shanghai, China)was used to assess cell proliferation and viability. The NOZ and SGC-996 cells were seeded at a concentration of 2,000 cells/well in 100  $\mu$ L in 96-well plates. To measure the IC<sub>50</sub> of GEM, gradient concentrations of them were added to cells for 48 or 72 h after cells were seeded for 24 h. The concentration of DMSO in the medium was controlled below 0.1%. CCK-8 (10  $\mu$ L) was added to each well, then the absorbance at 450 nm was measured after incubating at 37°C for 2 h. The IC<sub>50</sub> value was calculated using GraphPad Prism 9.0.

The NOZ and SGC-996 cells were seeded at 1,000 cells/well in 6-well culture plates and grew for about two weeks. 4% paraformaldehyde was then used to fix the cells for 30 min, and 0.5% crystal violet solution was used to stain. The colony numbers (> 50 cells/colony) were manually counted.

### 2.6. Apoptotic assay

GBC cells were plated in 6-well plates  $(2 \times 10^5 \text{ cells/} \text{ well})$  and incubated overnight. After incubation, cells were transfected with control, FASN siRNA or treated with orlistat or gemcitabine for 48 h. Then, the apoptotic state of GBC cells was assessed by the Annexin V-FITC and PI (BD, United States, Cat number: 556547) cell staining following manufacturer's protocol. The samples were detected using flow cytometry (Cytoflex, Beckman, United States) and the results were analyzed using the CytExpert Software.

### 2.7. Western blot analysis

GBC cells were plated in 6-well plates ( $2 \times 10^5$  cells/ well) and incubated overnight. After incubation, cells were transfected with control, FASN siRNA or treated with Orlistat or gemcitabine for 48 h. Then, cells were harvested and protein was extracted with radioimmunoprecipitation assay buffer (Beyotime, Shanghai, China). Protein was separated on 7.5-12.5% sodium dodecyl sulfate (SDS)-polyacrylamide gels and transferred onto 0.22 µm PVDF membrane (Merck Millipore, Burlington, MA, USA). The membranes were blocked with 5% milk (Yeason, Shanghai, China) for 1 h, incubated with the primary antibodies (1:1,000 dilution) at 4°C overnight, and incubated with the secondary antibody (1:1,000 dilution) at room temperature for 1h. Bands were visualized using ECL (Yeason, Shanghai, China). Antibody information: anti-β-actin (Abclonal, Wuhan, China, AC026); anti-FASN (Abclonal, A19050); anti-E-cadherin (Abclonal, A11509); anti-CDK-4 (Raybiotech, Georgia, USA, 144-00366); anti-cyclin-D1 (Cell Signaling, Massachusetts, Danvers, USA, #2978); anti-Bcl-2 (Proteintech, Wuhan, China, 60178-1-Ig); anti-Bax (Proteintech, 50599-2-Ig); anti-vimentin (Proteintech, 10366-1-AP); anti-PI3K (Proteintech, 20584-1-AP); anti-p-PI3K(Tyr458/Tyr199) (Cell Signaling, #4228); anti-AKT (Cell Signaling, 10176-2-AP); anti-p-AKT (Ser473) (Cell Signaling, #9271).

### 2.8. Xenografted animals

Four-week-old BALB/c nude mice were purchased from the Shanghai SLAC Laboratory Animal Co., Ltd. (Shanghai, China), and the study was approved by Laboratory Animal Ethical and Welfare Committee Xin Hua Hospital Affiliated to Shanghai Jiao Tong University School of Medicine. We subcutaneously administered  $1 \times 10^{6}$  NOZ cells (NC, lv-shFASN) resuspended in 0.1 mL PBS to nude mice on the left armpit. The mice were randomly grouped into four groups (n = 6). Two groups (control and shFASN) received an intraperitoneal injection with vehicle (Phosphate buffered saline, PBS) and the others (control and shFASN) with gemcitabine (50 mg/kg, twice a week), and body weight was measured every 2 days. We sacrificed the mice using CO<sub>2</sub>, removed the tumors and weighed them. The tumor volume was calculated (V) (length  $\times$  width<sup>2</sup>)/2.

### 2.8. Statistical analyses

All *in vitro* experiments were conducted in triplicate, and data were presented as mean  $\pm$  standard deviation (SD). Differences between the two groups were evaluated using the Student's *t*-test. Depending on the total sample size and expected frequencies, the chi-square test or Fisher's exact test were chosen to assess the significance of differences of clinicopathological figures between groups. Data analyses were carried out in the GraphPad Prism 9 and SPSS version 25. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001, and \*\*\*\*p < 0.0001 indicated statistical significance.

### 3. Results

3.1. Enhanced FASN expression is detected in GBC and involved with unfavorable clinicopathology of GBC tissue

To evaluate FASN expression in GBC, we collected

a total of 48 human GBC paired tumors and adjacent normal tissues. Tissue microarray data showed that FASN was mainly localized on the cell membrane and the cytoplasm of GBC tissues (Figure 1A). We assessed and scored staining pattern of FASN in both GBC and adjacent paracancerous gland tissue. It revealed that expression levels of FASN protein in cancer and normal tissues are both high. Of note, FASN protein expression was significantly increased in cancers compared to the matched gallbladder tissues (Figure 1B). Specifically, 40 out of 48 cases exhibited high expression of FASN (score > 4) in cancer tissues, while 20 out of 48 cases showed high expression in paracancerous tissues. RT-PCR data from another collected cohort also suggested that FASN mRNA expression levels were notably higher in gallbladder cancer tissues (14 cases) than in the adjacent normal tissues (14 cases) (Figure 1C). To investigate the correlation between FASN protein expression and clinicopathological features, 38 cases of complete medical reports were obtained and analyzed (Table 1). FASN expression was not statistically associated with age, gender, histological types, resection method, and gallstone state, but high FASN expression was significantly correlated with advanced clinical stage and nevin stage (Table 1). These results indicate upregulated FASN expression in GBC tissues and this gene's oncogenic properties in GBC.

3.2. FASN knockdown promotes apoptosis and inhibits proliferation and migration of GBC cells

After evaluating the FASN expression in GBC cell lines by western blot (Figure 2A), we selected NOZ and SGC-996 for further experiments. To further elucidate the oncogenic function of FASN, we knocked down the FASN protein expression in GBC cells by siRNA. The knockdown efficiency was confirmed by western blot (Figure 2B). The CCK8 assay and colony formation assay showed that the proliferative capacity of NOZ and SGC-996 cells were impaired following the knockdown

 Table 1. Summary of patient characteristics according to

 FASN expression

		FASN ex		
Variable	Total number	Relatively low (R < 2)	Relatively high $(R \ge 2)$	P value
Age (y)				1
> 65	19	9	10	
$\leq 65$	19	10	9	
Gender				1
Male	11	5	6	
Female	27	14	13	
Histological types				1
Adenocarcinoma	33	17	16	
Other <sup>a</sup>	5	2	3	
Clinical stage				$0.003^{**}$
1-2	7	7	0	
3-4	31	12	19	
T classification				0.055
T1-T2	11	9	2	
T3-T4	27	10	17	
N classification				0.329
N0	18	10	8	
N1	10	3	7	
N2	10	6	4	
M classification				1
M0	26	13	13	
M1	12	6	6	
Nevin staging				$0.008^{**}$
IV, V	31	12	19	
I, II, III	7	7	0	
Resection method				0.728
R1, R2	12	7	5	
R0	26	12	14	
Gallstone				1
(+)	18	9	9	
(-)	20	10	10	

<sup>a</sup>adenosquamous carcinoma, large cell neuroendocrine carcinoma, sarcomatoid carcinoma;  $\chi^2$  test was performed.



Figure 1. Enhanced FASN expression is detected in GBC and involved with unfavorable clinicopathology of GBC tissue. (A), Representative images of FASN expression in GBC tissues as visualized by TMA(Bar,100 $\mu$ m). (B), Expression of protein FASN in 48 paired GBC and adjacent normal tissues analyzed by TMA. (C), qPCR analysis of FASN mRNA expression. Data information: In (B), data are presented as mean ± SD. \*p < 0.05, \*\*\*\*p < 0.0001 (Mann–Whitney U test for B, Student's-*t* test for C).

of FASN (Figures 2C and 2D). This was further proved by impaired protein levels of cyclin-D1 and CDK-4 in FASN knocking down cells (Figure 2G). Moreover, FASN knockdown significantly enhanced apoptosis in NOZ and SGC-996 cells (Figure 2E). Increased levels of apoptosis associated proteins Bax and decreased levels of Bcl-2 in FASN knockdown cells supported the results above (Figure 2G). Transwell assay and western blot assay were performed to evaluate the effect of FASN on the migration ability of GBC cells. The migrated cell numbers were significantly reduced after the knocking down of FASN (Figure 2F). Enhanced E-cadherin and reduced vimentin protein expression after knocking down FASN supported that FASN indeed promoted migration in GBC cells (Figure 2G). Taken together, these results demonstrate that FASN might promote proliferation and migration while impairing apoptosis of GBC cells.

3.3. Orlistat promotes apoptosis and inhibits proliferation and migration of GBC cells

Orlistat is a classical inhibitor of the thioesterase domain of FASN approved by the Food and Drug Administration for treating obesity, recently it has been proven to halt cell proliferation, and induce cancer cell apoptosis in various types of tumors (12,17). However, the role of orlistat in GBC remains unknown. To characterize the effects of orlistat on GBC and pave a way for clinical application, we first measure cell viability of NOZ/SGC-996 incubated with different concentrations of orlistat for 48 h via CCK-8 assay (supplementary Figure S1, http://www.ddtjournal. com/action/getSupplementalData.php?ID=167). Cell viability assay showed orlistat suppressed the growth of both two cell lines dose-dependently (supplementary Figure S1, http://www.ddtjournal.com/action/ getSupplementalData.php?ID=167, Figure 3A). Also, the



**Figure 2. FASN knockdown promotes apoptosis and inhibits proliferation and migration of GBC cells.** (A), Relative expression of FASN in three GBC cell lines. (B), Knockdown efficiency in NOZ and SGC-996. (C), Cell viability assay of FASN knockdown cell lines. (D), Colony formation assays and statistical analysis of two cell lines transfected with si-FASN. (E), Apoptosis assays and quantification of the apoptotic cell population. (F), Transwell assays and statistical results (Bar,100µm). (G), Western blotting of cell fractions in FASN knockdown cell lines of NOZ and SGC-996. Data are presented as mean  $\pm$  SD. \*p < 0.05, \*\*p < 0.01,\*\*\*p < 0.001.

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number of colonies formed by GBC cells was obviously reduced after treatment by different concentrations of orlistat for 48 h (Figure 3B). Orlistat also induced NOZ and SGC-996 cell apoptosis to a degree (Figure 3C). Besides, the transwell assay for cell migration showed that the number of migrated NOZ (0, 10, 100  $\mu$ M) and SGC-996 (0, 50, 100  $\mu$ M) cells treated with orlistat were decreased in a dose-dependent manner (Figure 3D). We observed an increase in protein expression of Bax and E-cadherin, and a decrease in the expression of Bcl-2, cyclin-D1, CDK-4, and vimentin in NOZ/SGC-996 cells (Figure 3E), which supported and verified that orlistat treatment could induce cell apoptosis, inhibit GBC cell proliferation and migration.

3.4. Impaired activity of FASN enhances gemcitabine sensitivity in GBC

FASN was reported to mediate chemoresistance of various drugs like gemcitabine (10), cisplatin, and doxorubicin (18). Considering that orlistat indeed showed anti-cancer activity in GBC cells without killing normal cells (18), we further wondered whether FASN inhibition by orlistat could show a positive effect on gemcitabine sensitivity *in vitro*. Then we evaluated sensitivity changes to gemcitabine in both NOZ and SGC-996 cells in the presence of 10  $\mu$ M orlistat. Compared to gemcitabine alone, combined treatment with orlistat for 72 h significantly lowered the IC<sub>50</sub> of gemcitabine in NOZ (from 56.14 nM to 15.18n M) and SGC-996 (from 1,206 nM to 406.6 nM) (Figures 4A and 4B).

We then examined the effect of the combined use of gemcitabine (1  $\mu$ M) and orlistat (10  $\mu$ M) on protein expression associated with cancer progression. Western



Figure 3. Orlistat promotes apoptosis and inhibits proliferation and migration of GBC cells. (A), Cell viability assay of cells pretreated with orlistat for 48h. (B), Colony formation assays and statistical analysis of two cell lines incubated with orlistat. (C), Apoptosis assays and quantification of the apoptotic cell population. (D), Transwell assays and statistical results (Bar,100µm). (E-F), Western blotting of cell fractions in cell lines of NOZ and SGC-996 treated with orlistat for 48h. Data are presented as mean  $\pm$  SD. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001.
blot results demonstrate that the addition of orlistat could enhance the effect of gemcitabine on the expression of apoptosis-associated protein (Bax and Bcl-2), which indicated that orlistat could promote gemcitabine-induced cell death (Figures 4C-4E). Also, the combination of gemcitabine with orlistat showed synergistic decreasing expression of cyclin-D1 and CDK-4 (Figures 4C-4E). Additionally, the combination of the two agents enhanced the suppression of migration compared to treatment with gemcitabine alone (Figures 4C-4E).

Apart from that, flow cytometry apoptosis assays were conducted to evaluate the effects of FASN knockdown on gemcitabine sensitivity in GBC cells, revealing that FASN knockdown cells had a higher proportion of apoptosis (Figure 4F). Taken together, these results revealed that impairment of FASN activity could enhance the gemcitabine sensitivity in GBC cells. 3.5. FASN mediated PI3K/AKT activation involves in GBC cells sensitivity to gemcitabine

Accumulating evidence has revealed that PI3K/AKT activation participates in multi-drug resistance and neoplastic lipogenesis of various cancers. Therefore, we examined the phosphorylated forms of PI3K and AKT in NOZ and SGC-996 with downregulated FASN expression or orlistat treatment by western blot assay. It suggested that activation levels of PI3K/AKT pathway was significantly impaired by FASN activity repression, with relatively slight or no changes in the total protein expression levels (Figures 5A and 5B). Furthermore, treatment with gemcitabine or orlistat independently or in combination was conducted to assess the modulatory effect on PI3K/AKT cascade reaction. Compared to separately application or vehicle control (DMSO), the



Figure 4. Impaired activity of FASN enhances gemcitabine sensitivity in GBC. (A-B), CCK8 assay showing the IC<sub>50</sub> of gemcitabine with or without orlistat (10  $\mu$ M) addition in NOZ/SGC-996 cells. (C-E), Western blot analysis of bax, bcl-2, cyclin D1, CDK-4, vimentin and E-cadherin in cells treated with gemcitabine (1  $\mu$ M) or orlistat (10  $\mu$ M) alone or in combination. (F), Apoptosis assays and quantification of the apoptotic cell population in GBC cells treated with gemcitabine (1  $\mu$ M), siFASN, or a combination of both. Data are presented as mean ± SD. The GEM + orlistat group was compared to the GEM group, and the other groups were compared to the control group. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001. GEM: gemcitabine.

gemcitabine/orlistat simultaneous administration showed more efficiency on inhibition of PI3K/AKT signaling (Figure 5C). Taken together, FASN mediates GBC progression and gemcitabine resistance, at least, in part, through PI3K/AKT pathway.

To further validate the role of the PI3K/AKT pathway in GBC cells, a rescue experiment was conducted using SC79, an AKT agonist. The apoptotic proportions of NOZ and SGC-996 cells treated with gemcitabine or gemcitabine in combination with orlistat (100  $\mu$ M) were assessed. As depicted in Figure 5D, orlistat augmented the apoptotic effect induced by gemcitabine, and pre-treatment with SC79 for one hour partially reversed the trend.

3.6. FASN knockdown inhibits tumor growth and enhances gemcitabine sensitivity in mice

To further explore the role of FASN in GBC progression

and gemcitabine sensitivity in vivo, we developed nude mouse xenograft models to perform following experiments. FASN-silenced and matching control (lvshFASN and lv-control) NOZ cells (Figure 6A) were subcutaneously injected into the right armpit of nude mice. These two kinds of mice were randomly divided into two groups separately and administered either gemcitabine (50 mg/kg) or PBS intraperitoneally. The tumor size and weight of mice were measured twice a week. The difference in the tumor volumes and weights between the lv-control and FASN-silenced, the lv-control treated with gemcitabine and the FASN-silenced treated with gemcitabine grouped mice were calculated. As exhibited by Figures 6B-6E, FASN knockdown could significantly abrogate tumor growth and the tumor abrogation effect of gemcitabine was significantly more evident in mice transfected with FASN-impaired cells. Immunohistochemical analyses of xenograft



Figure 5. FASN mediated PI3K/AKT activation involves in GBC cells sensitivity to gencitabine. (A-B), Western blot assay showing the expression levels of PI3K, p-PI3K, AKT, and p-AKT in FASN knockdown (A) and orlistat treatment (B) NOZ/SGC-996 cells. (C),Western blot analysis of PI3K, p-PI3K, AKT, and p-AKT in cells treated with gencitabine  $(1\mu M)$ /orlistat  $(10\mu M)$  alone or in combination. (D), Apoptosis assays and quantification of the apoptotic cell population in cells treated with gencitabine (500 nM) or gencitabine in combination with orlistat (100  $\mu$ M). Pretreatment with 10  $\mu$ M SC79 was used to rescue the effect of orlistat. Data are presented as mean ± SD. The GEM + orlistat group was compared to the GEM group, and the other groups were compared to the control group. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001.

tumor tissues demonstrated that the expression level of apoptosis marker cleaved caspase 3 was elevated in the gemcitabine treatment group, and this was further enhanced by FASN knockdown (Figure 6F). Additionally, gemcitabine treatment, FASN knockdown, and their combination decreased the expression of Ki-67 (Figure 6F). Western blot analysis also showed that stable FASN knockdown downregulated p-AKT protein levels in xenograft tumor tissues (Figure 6G). Collectively, these results suggest that FASN knockdown can inhibit GBC tumor growth and improve gemcitabine sensitivity *in vivo via* the PI3K/AKT signaling pathway.

#### 4. Discussion

Dysregulation of lipid metabolism is closely associated with the occurrence and progression of various tumors (19-21). Current clinical research indicates that lipid metabolism-targeted therapies exhibit promising anticancer effects (19, 22). Nevertheless, there is still a relatively limited amount of research regarding lipid metabolism in GBC. Epidemiological and clinical studies have demonstrated a positive correlation between obesity and overweight status with an elevated risk of GBC (23-25). Bile lipidomics experiments have revealed that the abnormal composition of lipids in the bile of GBC patients is correlated with disease severity (26), suggesting that lipid metabolism dysfunction may serve as a hallmark for early diagnosis of GBC (27). a-Mangostin, a dietary xanthone, has been shown to augment the susceptibility of gallbladder cancer to gemcitabine treatment by repressing lipid biosynthesis through the targeting of AMPK/SREBP1 signaling pathways (28). These findings suggest that aberrant lipid metabolism may also be involved in the tumorigenesis and malignant progression of GBC. To further clarify the relationship between lipid metabolism and GBC progression, we focused on a potential target, FASN, by



Figure 6. FASN knockdown inhibits tumor growth and enhances gemcitabine sensitivity in mice. (A), Validation of FASN knockdown efficiency. (B-C), Xenograft tumors in BALB/c nude mice. NOZ (lv-control, lv-shFASN) cells were inoculated into BALB/C nude mice. One week later, the mice were intraperitoneally injected with gemcitabine(50mg/kg) or PBS. The volumes (D) and weight (E) of tumors were measured. (F), IHC of Ki-67 and cleaved caspase 3 (c-caspase 3) in the tumors (Bar,100  $\mu$ m). (G), Western blot analysis of AKT and p-AKT in four groups of xenograft tumor tissues. Data are presented as mean  $\pm$  SD. The lv-shFASN + GEM group was compared to the lv-control + GEM group, and the other groups were compared to the lv-control group. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001, \*\*\*\*p < 0.0001.

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exploring its expression and function in GBC.

Fatty acids are a primary source of energy for rapidly dividing tumor cells and also serve as the fundamental building blocks for cell membrane synthesis. FASN is an essential enzyme in fatty acid synthesis and plays a pivotal role in this process. Extensive research has been conducted on the role of FASN in cancer progression and carcinogenesis (8,9,12), but its involvement in GBC has not been reported. In this study, we demonstrate that FASN is upregulated in GBC and facilitates malignant progression and gemcitabine insensitivity. Additionally, we show that inhibition of FASN leads to cell apoptosis by regulating the levels of PI3K and AKT phosphorylation, which are crucial for the oncogenic process. Further, we reveal that apoptosis induced by FASN inhibition can be rescued by the application of an AKT activator, SC79.

Studies have shown that various types of tumors, including prostate cancer, breast cancer, ovarian cancer, colon cancer, and gastric cancer, exhibit high FASN expression and elevated FASN activity (12). According to a previous study, ovarian cancer tissue overexpressed FASN protein by an average of 1.8 times relative to healthy fallopian tube tissue (29). The expression levels of FASN were found to be associated with different grades of ovarian cancer tissues (29). Additionally, FASN in serum was absent or present at lower levels in both normal and non-malignant conditions, suggesting that it may be useful as a marker for early cancer detection (30). In our results, both FASN protein and mRNA are overexpressed in GBC tissues compared to adjacent normal tissues. Moreover, increased FASN expression is positively correlated with more advanced GBC pathological stages, which are associated with poorer clinical outcomes for patients. Since some patients are lost to follow-up and the samples collected were from 2018 to 2020, the survival data is incomplete. Additional cohorts including more patients are needed to reaffirm these conclusions.

A previous study demonstrated that knockdown or knock-out of FASN inhibited esophageal squamous cell carcinoma cell proliferation (31). Exogenous expression of FASN enhanced the proliferation, migration, and cell motility of SK-UT-1 via promoting trimethylation of H3K9 (H3K9me3) and acetylation of H3K27 (H3K27ac) in uterine leiomyosarcomas cells (32). In addition, the reduction of palmitate synthesis by FASN inhibition contributed to cancer cell apoptosis by disrupting cell membranes formation, repressing signaling transduction such as PI3K-AKT-mTOR and lipid biosynthesis, and suppressing gene expression such as c-Myc (33). Interestingly, FASN is also dispensable for the function of Treg cells and FASN knock-out in Treg inhibits tumor growth (34). But no documentation uncovers the role of FASN playing in GBC. The results of our in vitro and in vivo experiments clearly demonstrate that inhibition of FASN by knockdown leads to a marked reduction in

GBC cell proliferation and migration while increasing apoptotic rates. Unfortunately, we encountered challenges in performing functional experiments by overexpressing FASN in GBC cells due to the large size of the protein, which can reach up to 273 kD, making it difficult to overexpress using current technologies. Therefore, we selected the FASN inhibitor orlistat to support our knockdown experiment. In the future, this part of the experiment could potentially be completed through supplementation with FASN catalytic products or the use of mutation techniques to enhance the enzyme activity of FASN (*31*).

Resistance is one of the key contributing factors to low response rate to chemotherapy in GBC. Lipid metabolism has been reported to be associated with drug resistance (20), FASN activity inhibition was also reported to reverse chemotherapy resistance in pancreatic cancer (10,35) and ovarian cancer (29). In our study, gemcitabine treated GBC cells show increased apoptotic rates after FASN knockdown. And in our xenograft animal experiment, gemcitabine shows more excellent therapeutic efficiency on tumor growth in mice transfected with FASN-knockdown NOZ cells. These findings suggest a possible target for GBC treatment and warrent further investigation to uncover the underlying mechanisms.

Orlistat irreversibly inhibits FASN through binding to the thioesterase domain of this enzyme. It is widely used for obesity treatment with considerable advantages in clinical safety over other inhibitors. Moreover, scores of studies have sought to verify the anticancer potential of this agent through in vivo and in vitro experiments aimed at evaluating its efficacy (12). Therefore, we conducted further investigations into the potential role of orlistat in GBC cells. Our findings reveal that orlistat effectively inhibits GBC cell progression, particularly in NOZ cells expressing higher levels of FASN. These results suggest that inhibitors may exhibit greater efficacy against tumors with elevated FASN activity. Thus, cancer patients exhibiting abnormally high FASN expression may represent appropriate targets for therapy. We further explored whether FASN inhibition by orlistat could enhance gemcitabine sensitivity. The CCK-8 assay demonstrates that administration of orlistat (10  $\mu$ M) can indeed reduce the IC<sub>50</sub> in GBC cell lines. Additionally, apoptotic assays indicate that orlistat can amplify gemcitabine-induced cell death. Nonetheless, we must acknowledge that the synergistic effect of orlistat and gemcitabine is not particularly pronounced, although we should not discount its modest effect. Considering that the side effects of orlistat are manageable, higher doses of the drug could be administered to enhance its effectiveness. In the future, a combination of orlistat and gemcitabine may improve the efficacy and safety of chemotherapy regimens for GBC. Further animal and clinical studies should be conducted to pave the way for GBC treatment.

PI3K/AKT signaling transduction is considered one of the causes of chemoresistance in various tumors. Aberrant activation of this pathway inhibits chemotherapy-induced apoptosis via multiple mechanisms (36). Phosphorylation activated AKT promotes cancer cells survival through mediating Bcl-2 and Bax expression (37). Cancer cells escape from apoptosis depending on Bcl-2 in a variety of tumors (38). In gastrointestinal stromal tumor cells, FASN knockdown resensitized drug-resistant cells to imatinib by inactivating the PI3K/AKT/mTOR signaling pathway (39). The present study validates that FASN knockdown and inhibition downregulate the phosphorylation level of PI3K and AKT protein without affecting expression levels of total protein. Both FASN downregulation and inhibition also induce apoptosis via enhancing Bax and suppressing Bcl-2 protein expression. Similarly, orlistat enhances gemcitabine-induced GBC cell death through boosting regulation of Bcl-2/Bax expression. Additionally, pretreatment with SC79, an AKT agonist, was shown to reverse the increased apoptosis of GBC cells caused by the combination treatment of orlistat and gemcitabine. Although we have conducted different experiments to confirm the role PI3K/AKT pathway plays in the function of FASN, we do not know how FASN interact with PI3K/AKT. Further mechanisms should be explored in following studies.

In conclusion, our study provides compelling evidence that FASN inactivation exerts inhibitory effects on GBC cell proliferation and migration, while promoting cell apoptosis and gemcitabine sensitivity through the PI3K/AKT pathway. This work not only identifies FASN as a promising therapeutic biomarker for GBC patients, but also highlights the potential clinical utility of orlistat, a lipid metabolism regulator, in improving the clinical outcomes of GBC patients through enhancing gemcitabine sensitivity. These findings open up a promising avenue for further research into targeting lipid metabolism in the treatment of GBC.

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# **Brief Report**

# Efficacy of *Andrographis paniculata* spray in acute pharyngitis: A randomized controlled trial

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SUMMARY Acute viral pharyngitis is a self-limited disease but the symptoms, a sore throat in particular, can affect one's quality of life. Medicine for symptom relief is the main treatment. Recently, many studies have shown that Andrographis paniculata was efficacious in treating many diseases, including upper respiratory infections. However, adverse reactions to systemic intake are a concern. Therefore, A. paniculata spray is intended to reduce systemic adverse reactions and provide patients with more comfort as its local use. This randomized, double-blind study enrolled 60 adult patients with acute viral pharyngitis. All patients were asked to score the severity of symptoms including a sore throat, difficulty swallowing, and coughing using an 11-point numeric rating scale from 0 to 10. A physical examination was performed to score the severity of erythematous and swollen mucosa using a 0-3 score ( $0 = n_0$ , 1) = mild, 2 = moderate, and 3 = severe). The patients were randomized to receive treatment with either an A. paniculata spray or a positive control chamomile spray. Results revealed a significant reduction in the severity of all signs and symptoms in both groups (p < 0.05). The duration of treatment response in the A. paniculata spray group was  $1.9 \pm 0.7$  days compared to  $2.5 \pm 1.2$  days in the chamomile spray group (p = 0.049). No adverse events were noted in either group. A. paniculata spray is safe and highly efficacious in treating acute viral pharyngitis and can reduce symptoms more rapidly than a positive control spray.

*Keywords* Andrographis paniculata, acute viral pharyngitis, sore throat, plant extract, spray

#### 1. Introduction

Acute pharyngitis is the inflammation of the mucous membrane of the oropharynx located in the upper part of the respiratory system. It is mostly caused by viruses and bacteria (1). Other less common causes of pharyngitis are allergies, trauma, cancer, reflux, and certain toxins (2). Patients with acute pharyngitis have a sore throat, cough, and difficulty swallowing. The disease can progress rapidly and may cause severe throat edema leading to dyspnea and suffocation. In adults, approximately 80%-90% of pharyngitis cases are caused by viruses (3)and in children, up to about 95% of cases of an acute sore throat are associated with viral infection (4). This indicates that the primary cause of acute pharyngitis is viruses. Even though viral pharyngitis can self-resolve in less than a week (5), many patients suffer from a severe sore throat that affects their quality of life. Thus far, there are no specific drugs for the treatment of viral pharyngitis. However, antibiotics are sometimes

prescribed without being indicated. A point that must be emphasized is that excessive and inappropriate use of antibiotics can lead to the development of new resistant strains of microorganisms ( $\delta$ ). Therefore, treatment or medication to relieve symptoms and avoid unnecessary use of antibiotics might be the best alternative. Many herbs have been used for centuries as remedies for human diseases. Recently, interest in searching for high efficacious and safe, new bioactive components from plants has increased. Many types of plants have been reported to have various pharmacological activities (7–9).

Andrographis paniculata, known as Fah Talay Jone (in Thai), is an annual herb in the family Acanthaceae. The plant is commonly found in many countries in Asia, and particularly India, China, and Thailand. It is one of the principal herbs in traditional medicinal remedies for the treatment of acute upper respiratory tract infection and diarrhea in India, China, and Thailand (10). It has been reported to have medicinal and pharmacological properties and to be effective for the treatment of various 341

diseases such as cancer, diabetes, high blood pressure, ulcer, leprosy, bronchitis, skin diseases, flatulence, colic, influenza, dysentery, dyspepsia, and malaria (11). The anti-inflammatory and antimicrobial activities of this plant have been reported (12,13). Its extract was prescribed in oral form to treat upper respiratory infections, including acute pharyngitis, during the coronavirus disease 2019 (COVID-19) pandemic (14). However, adverse reactions have been reported after systemic administration (15). Therefore, local administration using throat spray containing *A. paniculata* extract has been considered.

In many countries including Thailand, the herbal sprays that are commercially available are those containing chamomile extract. Due to its antiinflammatory action, chamomile extract has been reported to relieve symptoms in patients with acute nasopharyngitis (16). Since the COVID-19 outbreak, the sprays have been used more widely and are in short supply. A. paniculata extract has therefore been put forward as a local herb that can help to relieve COVID-19 symptoms. To the extent known, no study has reported on A. paniculata extract spray to treat pharyngitis. Therefore, this pilot study was conducted with the aim of highlighting the first ever report on the efficacy of A. paniculata spray in acute pharyngitis. The clinical effectiveness and safety of A. paniculata spray in relieving symptoms of acute pharyngitis were investigated in comparison to chamomile spray.

#### 2. Materials and Methods

#### 2.1. Materials

*A. paniculata* extract was kindly provided by the Chao Phya Abhaibhubejhr Hospital Foundation, Prachinburi Province, Thailand. All chemicals and solvents used were of the highest pharmaceutical grade available. A commercial herbal spray containing mainly 37% chamomile extract, namely chamomile spray, was used as a positive control and purchased from a drug store in Thailand.

#### 2.2. Preparation of A. paniculata spray

*A. paniculata* spray was prepared by the Chao Phraya Abhaibhubejhr Hospital Foundation using a cosolvent method. The formula contains 0.3% of *A. paniculata* extract, glycerine, ethanol, and flavoring agents. The mixture was gently mixed until a homogeneous solution formed. The resulting solution was adjusted to volume with sterile water and gently shaken to avoid air bubbles. The final solution was transferred into a light-protected container with a spray nozzle.

#### 2.3. Study design and sample size

This study was approved by Chiang Rai Prachanukroh

Hospital's Ethics Committee in Human Research (No. CR 0033.102/EC.66-025). All study procedures were in accordance with the provisions of the Declaration of Helsinki. The study protocol was registered with the Thai Clinical Trials Registry (TCTR 20230915005).

A randomized, double-blind study was conducted by the Department of Otolaryngology at Chiang Rai Prachanukroh Hospital in Thailand from February 1 to April 30, 2023. Sample sizes were calculated using the ANCOVA method and based on a previous study (17). The estimated sample sizes for two groups with repeated measures were used. From this calculation, the sample size (n) with an alpha error of 5% and a power of 80% is 28 patients/group. To compensate for unexpected loss during follow-up, 30 patients were included in each group. Therefore, a total of 60 patients were recruited.

#### 2.4. Participant selection and eligibility criteria

Adult patients (aged from 18 years old) treated at Chiang Rai Hospital with an acute sore throat for less than 4 days were enrolled. Patients with a pain score greater than 5 out of 10 and a diagnosis of acute pharyngitis confirmed by an otolaryngologist were included in this study. The exclusion criteria were a positive COVID-19 test, a history of allergies, intolerance or sensitivity to any of the study medications, pregnant or lactating women, and patients who had taken antibiotics or corticosteroids less than 14 days prior.

#### 2.5. Randomization and blinding

The patients in each group were randomized using a computer-generated blocked randomization schedule. Both groups received throat spray bottles and were blinded to treatment. Both sprays were packaged in bottles with the same appearance labeled as either A or B. Codes A and B were announced as either *A. paniculata* spray or chamomile spray after the end of the study analysis.

#### 2.6. Study assessment

On the first day of study participation (day 0), the patients were asked to rate the severity of their symptoms, including a sore throat, difficulty swallowing, and coughing using an 11-point numeric rating scale from 0 to 10 (0 = no symptoms and 10 = the worst symptom that you can imagine). An otolaryngologist then performed a physical examination to assess the throat mucosa and score the severity of erythema of the pharyngeal mucosa (0 = no, 1 = mild, 2 = moderate, and 3 = severe erythema) and swelling of the pharyngeal mucosa (0 = no, 1 = mild, 2 = moderate and 3 = severe swelling). After the history was taken and a physical examination was performed, the patients were advised to spray the throat 2 spritzes/time and 3 times a day for

5 days. Symptoms were assessed for another 5 days (day 1 to day 5 = 1 to 5 days after using the sprays). The patients were asked to record the severity of their daily symptoms. These included a sore throat, difficulty swallowing, and coughing on an 11-point numeric rating scale from 0 to 10. A 20% or greater reduction in symptoms from the pretreatment score was considered acceptable (*18*). In addition, the patients were examined twice by an otolaryngologist on day 3 and day 5 of the study. All patients were asked to score their satisfaction with the received spray at the end of day 5.

#### 2.7. Safety assessment

Adverse events were monitored throughout the study. All participants were asked to promptly report any severe adverse events or adverse reactions that occurred. They would then be asked to discontinue the study and receive immediate medical treatment from an expert physician.

#### 2.8. Statistical analysis

Baseline demographic characteristics were described using descriptive statistics. Scores from the numeric rating scale and evaluations of the pharyngeal mucosa

Table 1. Patient demographics and clinical characteristic
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were expressed as the mean  $\pm$  S.D. Differences between groups were assessed using the Chi-squared test for categorical variables and the Mann-Whitney U test for continuous variables. The level of statistical significance was set at p < 0.05.

#### 3. Results and Discussion

The aim of the current study was to investigate the efficacy of the developed throat spray containing an extract of A. paniculata in patients with acute viral pharyngitis. A commercial herbal spray, namely chamomile spray, was used as a positive control. Although acute viral pharyngitis can resolve by itself, supportive treatment is also an important option for relieving a sore throat (5). Many studies have proven that A. paniculata extract has the potential to inhibit inflammation of the upper respiratory tract (URI) including pharyngitis (19,20). However, some systemic adverse effects, e.g., induction of acute renal injury (15) and gastrointestinal disorders (21) after parenteral and oral administration, have been reported. In the current study, local administration of the developed A. paniculata spray was used to reduce systemic adverse reactions. A total of 60 patients were enrolled and remained in the

Items	A. paniculata spray		Chamomile spray		1
	n	%	n	%	<i>p</i> -value
Gender					0.596
Female	17	56.67	20	66.67	
Male	13	43.33	10	33.33	
Age (years)	$42.6 \pm 13.9^*$		$44.1 \pm 15.2^*$		0.685
Occupation					0.638
None	2	6.67	0	0.00	
Government employee	5	16.67	6	20.00	
Office worker	3	10.00	4	13.33	
Company employee	2	6.67	2	6.67	
Farmer	1	3.33	4	13.33	
Nurse	2	6.67	2	6.67	
Housewife	3	10.00	6	20.00	
Freelance	8	26.67	4	13.33	
Other	4	13.33	2	6.67	
Underlying illness	16	53.33	3	36.67	0.299
Currently using medication	12	40.00	9	30.00	0.589
Duration of sore throat (days)	$2.4 \pm 1.1*$		$2.6 \pm 0.9*$		0.508
Difficulty swallowing	21	70.00	25	83.33	0.360
Cough	22	73.33	26	86.67	0.333
Fever	1	3.33	3	10.00	0.612
Headache	3	10.00	3	10.00	1.000
Hoarseness	6	20.00	7	23.33	1.000
Dyspnea	1	3.33	0	0.00	1.000
Nasal congestion/rhinorrhea	13	43.33	17	56.67	0.439
Otalgia	1	3.33	1	3.33	1.000
Throat soreness score (0-10)	$5.7 \pm 0.9$ *		$6.5 \pm 1.7*$		0.148
Difficulty swallowing scale (0-10)	$4.6 \pm 1.9*$		$5.5 \pm 2.1*$		0.202
Cough severity (VAS 0-10)	$4.0 \pm 3.0*$		$5.1 \pm 2.7*$		0.065
Body temperature (°C)	$36.41 \pm 0.17*$		$36.42 \pm 0.24*$		0.759
Erythematous of the pharynx (0-3)	2.4	$\pm 0.7*$	$2.3 \pm 0.7*$		0.469
Swollen mucosa of the pharynx (0-3)	2.0	$\pm 0.5*$	2.1	$\pm 0.6*$	0.812

\* Data express the mean  $\pm$  S.D.



Figure 1. Mean score for the severity, as determined by patients using an 11-point numeric rating scale from 0 to 10, of throat pain (A), difficulty swallowing (B), and coughing (C) after using *A. paniculata* spray (black column) and chamomile spray (white column).

final analysis (*A. paniculata* spray: n = 30, chamomile spray: n = 30). The two groups did not differ in baseline demographics and clinical characteristics as shown in Table 1. The mean duration of the clinical sore throat was 2.4 days in the *A. paniculata* spray group and 2.6 days in chamomile spray group. Common comorbid symptoms for each group were difficulty swallowing and coughing. The baseline signs and symptoms of pharyngeal inflammation did not differ significantly between the two groups.

Results indicated that the severity of a sore throat, difficulty swallowing, and coughing decreased significantly from day 1 after use of the spray in both groups and the score continued to decrease to nearly 0 at the end of the assessment as shown in Figure 1. The percentage of responders on day 1 was 80% in the *A. paniculata* spray group and 66.7% in the chamomile spray group. The mean ( $\pm$  SD) of the response duration was 1.9 ( $\pm$  0.7) days and 2.5 ( $\pm$  1.2) days in the *A. paniculata* spray group and chamomile spray groups, respectively (p = 0.049). Twenty-two patients in each group (73.3%) had no sore throat (score = 0) at the end of the study. The mean ( $\pm$  SD) duration to absence of pain was 3.4 ( $\pm$  1.1) days and 3.7 ( $\pm$  1.1) days in the *A. paniculata* spray and chamomile spray groups,



Figure 2. Mean score for the severity, as determined by otolaryngologists, of erythema (A) and swelling (B) of the pharyngeal mucosa after using *A. paniculata* spray (black column) and chamomile spray (white column).

respectively (p = 0.468). There was no difference in the overall satisfaction of using sprays between the two groups. In addition, there were no adverse events during this study. Other assessments scored by an otolaryngologist including erythema and swelling of the pharyngeal mucosa also decreased significantly in both groups as shown in Figure 2. Daily symptom scores between the two groups were compared, revealing no statistical difference.

The current results indicate that the sprays significantly alleviated clinical symptoms. Most patients were pain-free after less than 5 days of using the sprays, while the disease resolved on its own within 10 days. The duration to becoming pain-free did not differ between groups, but the duration of a clinical response differed. The A. paniculata group had a slightly faster recovery than the chamomile group. The current results on the effectiveness of an A. paniculata spray in reducing the severity of a sore throat are consistent with those reported by other studies involving A. paniculata in tablet form (22-24). However, the use of tablets results in significantly more adverse reactions than a topical spray. Adverse effects from an A. paniculata extract as reported earlier were mostly mild gastrointestinal symptoms such as constipation, nausea, vomiting, diarrhea, or dyspeptic symptoms (25). In contrast, there were no adverse events during the current study. This confirms that that topical application of an A. paniculata spray can be highly efficacious without causing systemic adverse reactions and that the study objective was achieved. The A. paniculata spray had anti-inflammatory action in terms of reducing pain without causing adverse reactions in this study, but the medicine should only be recommended when appropriate as determined by a physician.

Most patients were similarly satisfied with each

spray, but some participants mentioned the intense taste of the *A. paniculata* spray. Patients complaining about the taste of pharmaceutical products is one of the most important problems. *A. paniculata* is known as the king of bitters (*11*), so a challenge for further development is to use pharmaceutical technology to mask or minimize the bitter taste of an *A. paniculata* spray while maintaining its powerful anti-inflammatory action. In addition, the current work is a pilot study. The sample size for this study was calculated based on previous studies, assuming similar efficiency of the spray. Further studies with larger populations will help with the use of an *A. paniculata* spray in the future.

#### 4. Conclusion

The current study yielded positive results with an *A. paniculata* spray that can be used as an alternative treatment to relieve the symptoms of pharyngitis while causing fewer systemic adverse reactions. It is a safe and convenient form of administration that capitalizes on the beneficial properties of Asian herbs. It has anti-inflammatory action in patients with acute viral pharyngitis. It can effectively relieve symptoms faster than a chamomile extract spray.

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# **Brief Report**

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## Hypofibrinolytic phenotype in Tsumura Suzuki Obese Diabetes (TSOD) mice unrelated to hyperglycemia

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*Keywords* type 2 diabetes, plasma, obesity, coagulation, hypofibrinolytic state

#### 1. Introduction

Obesity and diabetes are significant risk factors for the development of cardiac diseases, arterial thrombosis and venous thromboembolism. The status of these illnesses can be determined as altered levels of blood lipids, adipocytokines, coagulative, and fibrinolytic factors in plasma (1). Various mouse models of type 2 diabetes have contributed to current understanding of metabolic syndrome and its related diseases (2). The Tsumura Suzuki Obese Diabetes (TSOD) polygenetic mouse model of spontaneous obese Type 2 diabetes mellitus (DM) is an inbred line in which males are obese and have urinary glucose, hyperglycemia, hyperinsulinemia, increased food intake, body and fat weight (3). Obesity and DM are associated with increased risk of arterial thrombosis and venous thromboembolism. Effectively preventing thrombotic complications requires better understating of how prothrombotic states develop in patients with metabolic disorders. Significantly higher plasminogen activation inhibitor-1 (PAI-1) levels in prediabetic patients than in healthy persons might function as predictors of progressive diabetes (4). Although animal models are useful for investigating mechanisms, detailed studies of pro-thrombotic and hypofibrinolytic states are hampered by a lack of appropriate prediabetic mouse models.

Here we evaluated metabolic, coagulative and fibrinolytic parameters in normoglycemic [Ng]-TSOD mice with a metabolic abnormality as well as coagulative and fibrinolytic states to determine whether they could function as prediabetic models.

#### 2. Materials and Methods

#### 2.1. Animals

Sixteen-week-old Male TSOD, TSOD with undetectable urinary sugar and normoglycemia (Ng-TSOD) and nondiabetic Tsumura Suzuki Non-Obesity (TSNO) mice (Institute of Animal Reproduction, Kasumigaura, Japan) were housed at  $24 \pm 2$ °C and provided with food and water *ad libitum* for one week. Casual blood glucose levels were measured using a StatStrip Express 900

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**SUMMARY** Obesity and diabetes mellitus are associated with increased risk of arterial thrombosis and venous thromboembolism. Tsumura Suzuki Obese Diabetes (TSOD) mice are useful models for elucidating the molecular mechanisms of these diseases. We investigated normoglycemic [Ng]-TSOD mice with a metabolic abnormality that was accompanied by a coagulative and fibrinolytic state with a phenotype that distinctly differed from that of standard TSOD mice. As in TSOD mice, plasminogen activation inhibitor-1 (PAI-1) that inhibits fibrinolysis was substantially augmented in Ng-TSOD mice, suggesting that they are hypofibrinolytic. However, blood clotting parameters were within the normal range in Ng-TSOD mice. These findings indicated that Ng-TSOD mice are novel models with a hypofibrinolytic phenotype that is not associated with hyperglycemia.

(Siemens Healthineers, Munich, Germany). Urinary sugar was evaluated using Uropaper IIIG (Eiken Chemical Co., Ltd., Tokyo, Japan). Table 1 shows renal and blood glucose levels in 16-week-old TSOD and Ng-TSOD mice. Blood specimens were collected from the inferior vena cava of 17-week-old mice as described (5). The Animal Care and Use Committee at Teikyo University approved all experiments involving mice (Approval No: 15-046).

2.2. Measurements of biochemical parameters and blood coagulation and fibrinolytic factors

Levels of glucose, triglyceride, and total cholesterol, free fatty acid, total protein, albumin, phospholipids, aspartate aminotransferase (AST) and alanine aminotransferase (AST), were determined using Biochemical Test Kits (Fujifilm Wako Chemicals, Osaka, Japan). Insulin,

Table 1. Body weight, blood and urinary glucose levels in TSOD and Ng-TSOD mice (n = 6 each)

Mouse no.	Body weight (g)	Blood glucose (mg/dL)	Urinary glucose
TSOD			
1	63.2	525	+
2	62.0	470	++
3	65.6	458	+
4	64.5	501	++
5	60.3	451	+
6	64.0	477	++
$Average \pm SD$	$63.1\pm1.9$	$480\pm21$	
Ng-TSOD			
1	58.1	203	-
2	56.1	183	-
3	55.8	188	-
4	58.2	175	-
5	48.7	168	-
6	57.6	172	-
$Average \pm SD$	$55.8\pm3.6$	$182\pm13$	

Glucose in blood from tail veins was measured using StatStrip Express 900 blood glucose meter (Nova Biochemical Corp., Waltham, MA, USA). Urinary glucose was measured using Uropaper III (Eiken Chemical Co., Ltd., Tokyo, Japan). –, undetectable; +, 500 mg/dL; ++, 2,000 mg/dL. adiponectin and leptin were measured using ELISA kits from their respective suppliers (Mercodia, Uppsala, Sweden, Otsuka Pharmaceutical Co., Ltd., Tokyo, Japan, and Morinaga Institute of Biological Science, Inc. Kanagawa, Japan). We analyzed tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) and C-reactive protein (CRP) using the respective ELISA kits (Chondrex Inc., Woodinville, WA, USA and Assaypro, St. Charles, MO, USA).

Levels of PAI-1 antigen and active PAI-1 were measured using Total and Active Murine PAI-1 ELISA kits (Molecular Innovations Inc., Novi, MI, USA) with slight modification. Fibrinogen and antithrombin levels were measured using ELISA kits (Assaypro). Prothrombin and activated partial thrombin (APTT) times were measured using Thromborel S (Dade Behring, Liederbach, Germany) and APTT reagent (Sysmex, Kobe, Japan), respectively, as described with slight modification (5).

#### 2.3. Statistics

Values are expressed as means  $\pm$  standard deviations (SD). Differences among the three groups were analyzed using one-way ANOVA followed by Tukey post-hoc multiple comparison tests.

#### 3. Results and Discussion

Glucose levels were significantly lower in Ng-TSOD, than in TSOD (p < 0.001), but the same as those in TSNO mice (Figure 1A). Mean body weight was slightly, but significantly lower in Ng-TSOD, than in TSOD mice (p < 0.01; Figure 1B). Insulin levels were significantly elevated in TSOD and Ng-TSOD, compared with TSNO (p < 0.005; p < 0.01), and those of Ng-TSOD and TSNO were indistinguishable (Figure 1C).

Figure 2 shows the plasma biochemical parameters in TSOD, Ng-TSOD and TSNO mice. Triglyceride levels were significantly lower in Ng-TSOD, than in TSOD mice (p < 0.05). Total cholesterol levels did not significantly differ between the Ng-TSOD and TSOD



Figure 1. Body weight, glucose and insulin levels in Ng-TSOD, TSOD and TSNO mice (n = 6 each). Data are shown as means ± standard deviation (SD). Significantly different at \*p < 0.01,  $^{\dagger}p < 0.005$ ,  $^{\ddagger}p < 0.001$ .



Figure 2. Plasma biochemical parameters in TSOD, Ng-TSOD and TSNO mice (n = 6 each). Data are shown as means  $\pm$  standard deviation (SD). Significant differences at \*p < 0.05,  $^{\dagger}p < 0.005$ ,  $^{\dagger}p < 0.001$ .

mice, but both were significantly higher than in TSNO mice (p < 0.01 for both). Levels of free fatty acids did not significantly differ between Ng-TSOD and TSOD mice, and neither did adiponectin levels, both of which were significantly lower than in TSNO mice (p < 0.001for both). Leptin levels did not significantly differ between Ng-TSOD and TSOD mice, but both were significantly higher than that in TSNO mice (p < 0.001for both). Levels of AST did not significantly differ between Ng-TSOD and TSOD mice, but both were significantly higher than in TSNO mice (p < 0.001; p <0.01). Levels of ALT did not significantly differ between Ng-TSOD and TSOD mice, but were significantly higher in TSOD, than in TSNO mice (p < 0.05) and similar between Ng-TSOD and TSNO mice. Levels of CRP did not significantly differ between Ng-TSOD and TSOD mice, but both were significantly higher than in TSNO mice (p < 0.01 for both). Values for TNF- $\alpha$  did not significantly differ between Ng-TSOD and TSOD mice, but both were significantly higher than in TSNO mice. Total protein, albumin, globulin and PL did not significantly differ among all three groups.

Figure 3 shows values for coagulation parameters that are usually measured in humans at clinical laboratories and PAI-1 in TSOD, Ng-TSOD, and TSNO mice. Levels of PAI-1 and active PAI-1 were significantly higher in Ng-TSOD and TSOD, than in TSNO mice (p < 0.005 and p < 0.05, respectively), but the difference between Ng-TSOD and TSOD mice did not reach significance. Plasminogen levels



Figure 3. Plasma coagulation and fibrinolytic parameters in TSOD, Ng-TSOD and TSNO mice (n = 6 each). Data are shown as means  $\pm$  standard deviation (SD). Significantly different at \*p < 0.05,  $^{\dagger}p < 0.001$ .

were significantly lower and higher in Ng-TSOD, than in TSOD (p < 0.01) and TSNO mice (p < 0.05), respectively. None of prothrombin, APTT, fibrinogen and antithrombin significantly differed among the three groups.

We found that levels of insulin and body weight were significantly increased in Ng-TSOD, compared with TSNO mice, but were the same between Ng-TSOD and TSOD mice. In contrast, glucose levels were the same between Ng-TSOD and TSNO mice. We also identified levels of adiponectin, leptin and CRP in Ng-TSOD and TSOD mice, which are abnormal in nascent metabolic syndrome in humans (*6*). The genetics of Ng-TSOD and normal TSOD mice might differ because TSOD mouse models of multifactorial genetic diseases spontaneously develop various metabolic diseases (*3*, *7*).

Patients with diabetes are likely to develop thrombophilia due to abnormal blood coagulation and fibrinolytic factors (8). The primary inhibitor of tissuetype plasminogen activator (tPA) in plasma is PAI-1, which plays a key role in fibrinolysis (9). Increased PAI-1 is one of the main coagulation abnormalities associated with obesity and diabetes (9). Many factors such as insulin, glucose, lipids affect increases in blood PAI-1 activities in patients with diabetes (10). The present study found the same levels of total PAI-1 and active plasma PAI-1 that reflect PAI-1 activity in Ng-TSOD and TSOD mice. Nonetheless, blood clotting parameters in Ng-TSOD mice were within the normal range. Although blood glucose levels might increase with advancing age in this phenotype, these mice could serve as models of hypofibrinolysis unrelated to hyperglycemia at least until they reach the age of  $\sim 17$  weeks.

In conclusion, we found that PAI-1 was substantially augmented in Ng-TSOD, as in TSOD mice. This finding suggested that Ng-TSOD mice are hypofibrinolytic and that Ng-TSOD mice might be novel models that could provide insight into the association of prediabetes with hypofibrinolytic states. We believe that Ng-TSOD mice have potential applications for investigating the diagnosis and treatment of prediabetes.

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*Conflict of Interest*: The authors have no conflicts of interest to disclose.

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# Brief Report

## Effectiveness of an artificial intelligence-based training and monitoring system in prevention of nosocomial infections: A pilot study of hospital-based data

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**SUMMARY** This work describes a novel artificial intelligence-based training and monitoring system (AITMS) that was used to control and prevent nosocomial infections (NIs) by improving the skills of donning/ removing personal protective equipment (PPE). The AITMS has two working modes, namely an AI-based protective equipment surveillance mode and an AI-based training mode, that were used for routine surveillance and training, respectively. Data revealed that the accuracy rate of donning/ removing PPE improved as a result of the AITMS. Interestingly, the frequency of NIs decreased with the use of the AITMS. This study suggested the key role of using PPE in controlling and preventing NIs. Data preliminarily proved that appropriate donning/removing PPE may help to reduce the risk of NIs. In addition, the newest computerized technologies, such as AI, have proven to be useful in controlling and preventing NIs. These findings should helpful to formulate a better strategy against NIs in the future.

*Keywords* artificial intelligence, nosocomial infection, surveillance, training, artificial intelligence-based training and monitoring system (AITMS)

#### 1. Introduction

The term "nosocomial infections (NIs)" is used to describe all infections acquired in a hospital. NIs have several characteristics that differ from communityacquired infections: i) NIs are healthcare-associated infections that are acquired after hospitalization and that manifest 48 hours after admission (1). Due to their considerable impact on clinical outcomes, some authors have proposed that the period for NIs should be extended to within 30 days of hospital discharge or 90 days after undergoing surgery (2). ii) NIs can be systemic; common NIs are pneumonia (hospitalacquired or ventilator-associated), urinary tract infections (catheter-associated), surgical site infections, central line-associated bloodstream infections, and Clostridium difficile infections (3). iii) Patients, relatives of the patient, caregivers, and medical personnel may be involved. iv) NIs are often associated with medical interventions, such as surgery, medication,

or invasive examinations. v) NIs readily occur in departments dealing with critical illnesses, such as the transplantation unit, chemotherapy unit, burn unit, or intensive care unit. vi) NIs can often be monitored and dealt with. NIs have been the sixth leading cause of death in United States (2). As a notable but intervenable cause of death in the hospital, NIs are of great concerned to clinicians and governments around the world. Indeed, NIs are commonly problematic because most NI-related pathogens, be they bacteria (4) or fungi (5), have natural resistance to empirical antibiotics and thus cannot be treated with commonly used antibiotics. Moreover, most people who contract an NI are older, neonates, those who are immunocompromised, or those who are suffering from chronic diseases. The clinical outcomes and quality of life (QOL) of patients markedly worsen once they develop an NI. Fortunately, however, NIs can be dealt with and prevented since they are hospital-related infections. In line with World Health Organization guidelines, the incidence of NIs

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can be reduced approximately 70% by satisfactory infection prevention and control programs (6). The vast majority of countries have set up surveillance and information sharing systems to fight against NIs (7). To the extent known, however, merely creating such systems without specifying their detailed operation in clinical practice might yield limited achievements in terms of reducing the frequency of NIs. Hence, many practical measures are taken to reduce the risk of NIs in routine clinical tasks, such as using personal protective equipment (PPE). PPE is used to reduce the exposure risk of medical personnel when they are treating infected patients or making contact with contaminated surfaces. Typical PPE commonly consists of gloves, goggles, a gown, a head cover, a face mask and respirator, shoe covers, and a face shield, which are regarded as a general safety precaution for infection prevention. The effectiveness of PPE is influenced by proper selection of PPE (selection of the type of mask, for example), appropriate usage of PPE (donning/ removal), and proper disposal of PPE (8). Although recommendations regarding PPE are sometimes controversial, PPE is undoubtedly emphasized and recommended during pandemics involving infectious respiratory diseases. During the COVID-19 pandemic, all frontline medical personnel were required to know appropriate methods of using PPE in clinical settings. Nonetheless, appropriately donning/removing PPE is quite complex and is poorly understood by most new medical personnel. John et al. reported that only 39% of untrained medical students can select the correct sequence of donning/removing PPE (9). Wilder-Smith et al. reported only 34% of medical personnel could correctly use PPE (10). Mitchell et al. reported that inappropriate donning/removing PPE is a significant risk factor for development of NIs (11). Conversely, proper use of PPE may help to reduce NIs (12). A point worth noting is that during an infectious respiratory disease pandemic, and especially in the early stages of the pandemic, inappropriate use of PPE might be dangerous, or even life-threatening. Flaws in donning/ removing PPE can cause pathogen transmission and lead to NIs (13). Accordingly, effective and evaluable training in using PPE is indispensable for medical personnel. The conventional centralized training model has many limitations, such as the time and effort it takes and limited efficiency, so it cannot ensure all trainees learn/understand proper skills of donning/ removing PPE. In this regard, many optimized training models have been proposed, such as the informationmotivation-behavioral skills (IMB) model. Recently, Song et al. used the IMB model to improve the skills of donning/removing PPE. They found that a group receiving an IMB model-based intervention had better performance in terms of self-efficacy and qualified usage of PPE among medical personnel during the COVID-19 pandemic (12). However, this model

also has limitations. Noticeably, the entire process of donning/removing PPE cannot be monitored and supervised in real time. Flaws in the process could not be identified and corrected in a timely manner. This might serve as a Trojan horse for NIs. To ensure all personnel can properly don/remove PPE both in training and also in clinical practice, the current authors developed a novel artificial intelligence-based training and monitoring system (AITMS). The aim was to develop a newer system that would help to improve the donning/removing of PPE and reduce the frequency of NIs, ultimately helping to improve the clinical outcomes and QOL of inpatients. The current pilot study was designed to verify the effectiveness of the AITMS in preventing NIs.

#### 2. Materials and Methods

#### 2.1. Components of the AITMS

The AITMS consists of a camera with a voice broadcasting feature, a backend administration system on the cloud, a smartphone app, and an AI-based surveillance/training terminal for donning/removing PPE and hand hygiene (these are denoted as "behaviors" in the following sections). The terminal consists of an AI on a local computer, along with a large screen (for displaying correct "behaviors" or serving as a humancomputer interface) with voice packets (for voice guidance and error prompts) (Figure 1). The AITMS has two working modes, namely, AI-based protective equipment surveillance mode (AIPESM) and AI-based training mode (AITM), which can be simply selected via the smartphone app or the terminal (Figure 1). The AIPESM is used to monitor a member of medical staff's "behaviors" and detect flaws in practice, whereas the AITM is used to training personnel if they need training. The AITM can certainly be used to solely train new personnel.

# 2.2. Definition of "standardized target behaviors (STBs)"

STBs were defined as per China's Technical Guidelines on Prevention and Control of Novel Coronavirus Infection in Medical Institutions (third edition) (14), including 7 steps of donning and 15 steps of removing protective equipment.

#### 2.3. Generation of a dataset

Data collection sites were established where medical personnel don/remove PPE. All the normative "behaviors" performed by a group of skilled personnel were recorded by a camera based on the Kinect technology. Every frame of the video was recorded and preprocessed. The size of the video frame was



Figure 1. A diagram of the artificial intelligence-based training and monitoring system (AITMS). AI: Artificial intelligence; AITMS: artificial intelligence-based training and monitoring system; AIPESM: AI-based protective equipment surveillance mode; AITS: AI-based training mode (AITM); APP: application; PPE: personal protective equipment.

normalized. Then, the types of behaviors involved in donning/removing PPE and hand hygiene that need to be marked (target behaviors) and the marking rules were determined. Behaviors in all frames of preprocessed images were subsequently marked. The marked data were approved both by the AI and manually to avoid missing or wrong marks.

2.4. Training and creating an AI model to evaluate STBs in images with machine learning

After all of the images were marked, the system automatically identified marked images and label files, which were subsequently divided into a training set and a test set at a ratio of 9:1. The AI used the training set to train the model, and the model performance was checked using the test set. The models were continuously optimized by adjusting the parameters and tests. On the basis of the results of model testing, samples were added when target behaviors or backgrounds were readily misdetected. These processes helped to improve the accuracy to reach the anticipated standard. Ultimately, an AI model to evaluate STBs in images was successfully created.

#### 2.5. Application to actual processes

Once the data accuracy reached the standard, the AI model was applied to actual processes to make precise adjustments and to facilitate optimization. The system was used in locations where PPE is donned/removed. We ensured that all the "behaviors" of the testees (hand washing, donning/removing one's mask and protective suit, *etc.*) could be clearly detected. The current AI model to evaluate STBs in images had satisfactory accuracy, particularly at detecting the behaviors of donning/removing one's protective suit. In addition, this system enabled the AI model to improve its accuracy by continuous self-learning during actual use.

#### 2.6. Scenarios for application

The AIPESM is activated to monitor clinical routine tasks of medical personnel. When medical personnel step into the designated area, he or she should don/ remove PPE and perform hand hygiene following the guidance of the voice prompt. The surveillance camera is activated and it performs real-time surveillance while the "behaviors" are performed. All the human postures and limb motion trajectories are captured and analyzed based on Kinect technology. All the "behaviors" of the testees are compared to the STBs in images evaluated by the AI model, including the sequence of behaviors, range, and angle. A "correct/incorrect" behavior is defined by the AI model according to its conformance to the STBs. Once an "incorrect behavior" occurs during the process, the camera captures this flaw and then submits it to the AI on the local computer. The AI system will judge whether this behavior is "incorrect". Once a flaw is confirmed, the system will remind the person to correct his or her behavior via the voice prompt. Only when a behavior is in line with the "STB" is it deemed to be a "correct behavior", and the person can continue with the process. If needed, experienced backend staff can also help personnel. Once a person is identified as needing training, he or she may choose to use the AITM. The required training course (donning a gown, for example) can be selected via the smartphone app. The theoretical knowledge and a video of the STB video will be displayed via the large screen to guide the person so that he or she can perform the "correct" behavior. The camera captures the person's movements during training, the system corrects his or her mistakes via the voice prompt, and it evaluates the person's performance during training. Other than in a clinical setting, the AITM can also

be used to train new personnel. Training, real-time evaluation/corrections, and administration of a final examination can be performed. Certificates will be issued along with the complete training files after the trainee passes the theoretical and practical exams. The files can be synchronized in the main computer and mobile terminal. Analysis of the obtained data will help to continuously optimize the training programs as well as the algorithms.

#### 2.7. Data acquisition and analysis

The AITMS was introduced at this Hospital starting in February 2020. When surveillance started (from February 7, 2020 to February 13, 2022), none of the personnel had ever been trained by the AITMS. The accuracy rate automatically recorded by the system was recorded as the non-AITMS rate. From November 20 to 26, 2022, almost all of the personnel were trained with the AITHM. The accuracy rate at that time was recorded as the AITMS rate. Data on the frequency of NIs from 2019 to 2022 at this Hospital are available on the National Medical Institution Infection Surveillance System of China (https://cniss.yygr.cn/). The software SPSS (v24.0.0, IBM, USA) was used for statistical analysis. A chi-square test was used to compare rates. p < 0.05 was considered to be a statistically significant difference.

#### 3. Results and Discussion

Figure 2A shows the accuracy rate before and after training with the AITMS. From February 7, 2020 to February 13, 2022, a total of 163 personnel had finished the evaluation, and their accuracy rate was 52.15% (85/163, non-AITMS rate). From November 20 to 26, 2022, the accuracy rate was 98.14% (3,159/3,219, AITMS rate). These data indicate that the accuracy rate of donning/removing PPE significantly increased as a result of using the AITMS (98.14% vs. 52.15%,  $\chi^2 = 834.35, p < 0.001$ ). Indeed, after training with the AITMS, most of the medical personnel could properly don/remove PPE (Figure 2A). Interestingly, the frequency of NIs also decreased accordingly. As shown in Figure 2B, the frequency of NIs was 1.31% in 2019 and 1.39% in 2020 before use of the AITMS; after use of the AITMS, the frequency decreased sharply to 0.58% in 2021 and 0.38% in 2022. Hence, use of the AITMS helped to reduce the frequency of NIs even though the COVID-19 pandemic struck in 2021 and 2022.

This pilot study preliminarily verified the effectiveness of the AITMS. Use of the AITMS eventually reduced the frequency of NIs as the accuracy rate of donning/removing PPE improved. This study suggested the key role of using PPE in controlling and preventing NIs. Data preliminarily proved that appropriate donning/ removing PPE may help to prevent the development of



Figure 2. Effectiveness of using the AITMS. (A). Using the AITMS significantly improved the accuracy rate of donning/removing PPE. (B). Using the AITMS markedly reduced the frequency of NIs. The AITMS was not used during the two years on the left while it was used during the two years on the right. The decline in nosocomial infections is evident despite the COVID-19 pandemic in 2021 and 2022. AITMS: artificial intelligence-based training and monitoring system; PPE: personal protective equipment. \*\*\* means p < 0.001

NIs In addition, the newest computerized technologies, such as AI, machine learning, big data, and mobile Internet (15), have proven to be useful in controlling and preventing NIs. The current results are in line with those of previous analogous studies (16,17). These findings should help to formulate a strategy against NIs in the future. In the future, more rigorous verification will be performed to provide robust evidence regarding use of the AITMS to reduce NIs.

Using AI-based technology to control and prevent NIs is a novel idea. Here, this novel AITMS has proven to be effective in both routine clinical practice and training. The two-mode design allows this system to be applied in multiple scenarios, such as routine surveillance, real-time correction of inappropriate "behaviors", timely training on demand, systematic training for new personnel, and evaluation and examination of trainees. Using this AI-based system markedly improved the efficiency of surveillance and training. This system can ensure that medical personnel and trainees actually understand the skills of donning/removing PPE. Compared to conventional methods, the SITMS can save manpower and time, so it is particularly good in scenarios involving the outbreak of a certain infectious disease when numerous new personnel need to be rigorously trained in a short amount of time. It can also serve as a routine surveillance system in a hospital or department when PPE is often used. However, AI technology is far from flawless. There are still technological limitations so far: The recognition accuracy is affected by many factors, such as the camera angle, speed of human movement, and the degree to which movement is standardized (18), and this is particularly true for the recognition of fine movements (19). The efficiency of self-learning is still too limited (20). These limitations might potentially cause errors in the recognition of movements, so they should be addressed in future research and development. Even though AI cannot completely take the place of clinicians right now, the value of AI technology cannot be ignored. It can help to continuously optimize the strategies against NIs, including devising more reasonable "STBs", simpler but efficient training processes, and a more friendly human-computer interface. The current finding should help to achieve medical care with "no harm and no infections" in an era of constantly emerging infectious diseases (21).

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## **Brief Report**

# A pilot study comparing the disinfecting effects of commercialized stable $ClO_2$ solution (free of activation) with conventional $H_2O_2$ on dental unit waterlines in the dental practice setting

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SUMMARY Disinfection of dental unit waterlines (DUWLs) plays a key role in control and prevention of nosocomial infection in a dental clinic. The most conventional disinfectant is hydrogen peroxide  $(H_2O_2)$ , while chlorine dioxide  $(ClO_2)$  has been considered however was limited by the "activation" procedures. With the availability of commercialized stable ClO<sub>2</sub> solution (free of activation), direct application of  $ClO_2$  in the dental practice became possible. This study was designed to compare the disinfecting effects of stable 5 ppm of  $ClO_2$  solution with conventional 0.24% of  $H_2O_2$  on DUWLs in dental practice. Studies of colony-forming units (CFUs), confocal laser scanning microscopy (CLSM) and scanning electron microscope (SEM) were employed for evaluation. In CFUs studies, we found that the efficiency of  $ClO_2$  was no less than those of  $H_2O_2$ . In the morphological studies, the stronger disinfecting effects of ClO<sub>2</sub> was verified by both CLSM and SEM studies for removal and prevention of biofilm. Importantly, ClO<sub>2</sub> solution achieved a better disinfecting efficiency not only at the surface of bacterial biofilm, but also, it has penetrating effects, presented disinfecting effects from the surface to the bottom of the biofilm. This pilot study provided evidence regarding the efficiency of stable  $ClO_2$ solution on disinfection of DUWLs in the dental practice setting. Application of stable ClO<sub>2</sub> solution in dental practice is therefore become possible.

*Keywords* dental unit waterlines, hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), chlorine dioxide (ClO<sub>2</sub>), disinfection, biofilm

#### 1. Introduction

Dental unit waterlines (DUWLs) are a piping system providing pure water for dental treatment. This system is comprised of several narrow, long pipelines, which is often intermittent used with unbalanced and slow flows. Accordingly, DUWLs are easily contaminated by bacteria and then induced bacterial biofilm formation. Frequently used positions, such as air/water syringe, dental hand piece, and cuspidor faucet usually have more chance to be contaminated, potentially conduct bacteria to the waterline, and promote biofilm formation (1). It has been documented that bacterial biofilm on DUWLs is widely distributed, with the approximately 30-50  $\mu$ m thickness, which is believed to potentially cause serious waterline contamination (2). If such contamination is neglected, the floating microorganisms or dissociative biofilms might be transferred to the patient, or come to the air through a handpiece, thereby increasing the infectious risks to patients and dental staffs (3). Hence, surveillance and prevention of DUWLs-related contamination are routine works of a dental clinic. A battery of disinfectants and disinfecting methods, such as hydrogen peroxide (H2O2) (4), chlorine dioxide (ClO2) (5), chlorhexidine gluconate (6), sodium hypochlorite (7), peracetic acid (8), intermittent sterilization with peracetic acid/ $H_2O_2$  (9), continuous disinfection with hydrogen peroxide/silver ions (6) were investigated for use in DUWLs. Nonetheless, only few of them are actually used in a dental clinical setting for various reasons. An ideal disinfectant for using in the dental practice setting should have several characteristics, such as effective, safe, appropriate priced, convenient, and easily available. Accordingly, H<sub>2</sub>O<sub>2</sub> is the most

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commonly used disinfectant for DUWLs clinically.

ClO<sub>2</sub> is an effective, safe high-level disinfectant, which is widely used for disinfection of environments, surface of articles, and human. It has been reported using  $ClO_2$  for oral cleaning (10-12) and wound cleaning (13). However,  $ClO_2$  is not commonly used for dental clinical setting because it is difficult to obtain a stable ClO<sub>2</sub> solution, and store it for a long time. Hence, the ClO<sub>2</sub> solution usually has to be prepared before using by a chemical reaction of precursors, which is termed as "activation", that is inconvenient and unsafe for DUWLs in the actual clinical setting (14) because the reaction concentration is not easily controlled. Our previous studies mentioned availability of a commercialized stable  $ClO_2$  solution that was free of activation (14,15), that make it possible for convenient use of ClO<sub>2</sub> in clinical setting since we can purchase the stable solution with a certain concentration. On the other hand, colonyforming units (CFUs) have been used as a standard index for evaluating the efficiency of disinfection in DUWLs scenario. Conversely, remove/control of bacterial biofilm during the disinfection in DUWLs has never been a standard index, even though it plays a key role in prevention and intervention of the DUWLs contamination.

Based on the aforementioned contexts, we designed this pilot study to compare the efficiency of disinfection in DUWLs between the conventional  $H_2O_2$  and the commercialized stable  $ClO_2$  solution (free of activation) in the clinical practice. Meanwhile, we also attempted to observe the changes of bacterial biofilm along with the CFUs affected by  $ClO_2$  solution and  $H_2O_2$ . We believe that the findings of this study will be useful for better understanding the efficiency of the commercialized stable  $ClO_2$  solution (free of activation) as well as changes of bacterial biofilm affected by  $ClO_2$  and  $H_2O_2$ , that is useful for selection of an appropriate disinfectant for DUWLs in the dental practice setting.

#### 2. Materials and Methods

2.1. Preparation of DUWLs and collection of the water samples

Experimental DUWLs in the present study were derived from the dental chair units (DCUs, UTTG27959, Planmeca, Helsinki, Finland), which had been normally used for the routine clinical practice for three years. Total 18 DCUs were involved in this study, where 12 DCUs were allocated to the ClO<sub>2</sub> group and 6 were allocated to the H<sub>2</sub>O<sub>2</sub> group using a simple coin toss randomized method. Two sorts of disinfectants were prepared in the present study, namely 5 ppm of commercialized stable ClO<sub>2</sub> solution (free of activation) which was purchased from the manufacturer (Shenzhen Caseche Biotech Co., Ltd., Shenzhen, Guangdong, China) and 0.24% H<sub>2</sub>O<sub>2</sub>(4). Concentrations of the agents were determined according to the previous studies using  $ClO_2$  (16,17) and  $H_2O_2$  (4) for disinfection.

Once the investigation initiated, 500 mL ClO<sub>2</sub> and H<sub>2</sub>O<sub>2</sub> solutions were put into the sterilizing bottle of DCUs respectively after the daily dental clinical work was finished. The disinfection procedures were opened for 4 min (wash with disinfectant for 2 min and then wash with pure water for 2 min); then the power switch was turned off overnight. Water samples were collected before the clinic work at the next morning. Sampling was performed as per the 2023 Guidelines for Infection Control and Management in Dental Unit Waterlines (18). Sampling was implemented at three positions, namely air/water syringe, dental hand piece, and cuspidor faucet following the principles of aseptic operation. Experiments were performed for 45 weeks, except the previous day for the baseline test. Water samples were measured once per week for the first 29 weeks, and once per two weeks for the last 16 weeks.

#### 2.2. Detection of the CFUs in water samples

In terms of CFUs test, 200  $\mu$ L sample water was put into a sterile petri dish, and mixed with medium, subsequently cultured at 37°C for 48 hours. CFUs were calculated as the numbers of bacterial colonies divided by the volume of diluent. Less than 100 CFUs/mL is considered as negative.

#### 2.3. Confocal laser scanning microscopy (CLSM) study

After 12 weeks of disinfection (ClO<sub>2</sub> or H<sub>2</sub>O<sub>2</sub>), Waterline samples of DUWLs were cut into rings (0.2-0.5 mm length), which were immediately exposed to a LIVE/ DEAD BacLight Bacterial Viability Kit (Cat. No. L7012, Thermo Fisher Scientific Inc., Waltham, MA, America) for 10 min, washed with PBS for 1 min, and then rinsed twice. Non-invasive CLSM images were acquired on the complete biofilm at the inner wall of DUWLs using a CLSM (FV3000, Olympus, Tokyo, Japan) (excitation light wavelength = 510/480 nm). Vital fluorescence staining (VFS) was performed as per the manufacturer's manual. Bright green staining displays live bacteria, red staining shows dead bacteria, and the yellow staining is the overlap (coexistence) of dead and live bacteria.

Image analysis was performed using an ImageJ 1.34p software (National Institutes of Health, Bethesda, MD, USA; *http://rsb.info.nih.gov/ij/*). Images of each color channel were assembled into stacked images, and the areas occupied by live bacteria and dead bacteria were calculated respectively. The ratio of live bacteria to dead bacteria was calculated and submitted to statistical analysis.

#### 2.4. Scanning electron microscope (SEM)

Remaining waterline samples of DUWLs undergone

a 12-week sterilization were cut into a 1 cm section, then cut vertically from the middle line. All samples were placed into 2.5% glutaraldehyde for overnight fixation. After being dehydrated by ethanol gradient (30%, 50%, 70%, 80%, 85%, 90%, 95%, and anhydrous ethanol for 0.5 h at each concentration), the tubes were fixed on a special aluminum base. After spraying gold nanoparticles, they were observed and photographed using a scanning electron microscope (Su8220, Hitachi, Tokyo, Japan).

#### 2.5. Statistics

A SPSS soft (V26.0.0, IBM, Armonk, NY, USA) was used for statistical analyzes. Comparisons of proportion were performed with a Chi-square test. The quantitative VFS data were compared using a Mann-Whitney U test. p < 0.05 was considered as the statistical significance.

#### 3. Results and Discussion

In the present study, we compared the disinfecting effects of commercialized stable  $ClO_2$  solution (free of activation) with conventional  $H_2O_2$  for DUWLs by observing the states of biofilm. Our data suggest a better disinfecting efficiency of this  $ClO_2$  solution than that of conventional  $H_2O_2$  in terms of DUWLs disinfection. To the best of our knowledge, this study is the first study to evaluate the efficiency of commercialized stable  $ClO_2$  solution (free of activation) using in disinfection of DUWLs. We believe that the findings of this study are helpful to select an appropriate disinfectant for DUWLs in the routine dental practice.

#### 3.1. Analysis of the CFUs in the DUWLs

As shown as in Table 1, total 1,998 water samples were tested, of those, 1,332 were in ClO<sub>2</sub> group, and 666 were in H<sub>2</sub>O<sub>2</sub> group. In the ClO<sub>2</sub> group, total 1,312 samples were identified as "-" once their detection values < 100 CFU/mL, the pass rate was 98.48%. In the H<sub>2</sub>O<sub>2</sub> group, total 648 samples were identified as "-", the pass rate was 97.30%. No significant difference was found between groups in total ( $\chi^2$  = 3.434, *p* = 0.064). In terms of different positions, no signification difference was found between two groups (Table 1). These data indicated that the disinfecting efficiency of this stable

 
 Table 1. Analysis of the colony-forming units in the dental unit waterlines

Positions	ClO <sub>2</sub> -/+	$\mathrm{H_2O_2}$ -/+	$X^2$	<i>p</i> value
Air/water syringe	440/4	220/2	0	1.000
Dental hand piece	440/4	216/6	3.249	0.071
Cuspidor faucet	432/12	212/10	1.504	0.220
Total	1312/20	648/18	3.434	0.064

"-" means the detection value < 100 CFUs/mL; "+" means the detection value  $\geq$  100 CFUs/mL, CFUs = colony-forming units.

 $ClO_2$  solution was no weaker than those of conventional  $H_2O_2$ 

3.2. Comparison of the disinfecting effects between  $ClO_2$  and  $H_2O_2$  by observing the changes of the biofilms

As shown as in Figure 1, the disinfecting effects between  $ClO_2$  and  $H_2O_2$  were compared with a CLSM along with a SEM. Biofilm is a thin layer at the surface of waterline. The results of the CLSM displayed that multitudes bacteria in the biofilm were



Figure 1. Comparison of the disinfecting effects between ClO<sub>2</sub> and H<sub>2</sub>O<sub>2</sub> by observation of the changes of biofilms. A. Representative images of CLSM displaying the disinfecting effects on the biofilm (ClO2 vs. H2O2). Green background is the wall of the waterline ("Wall" in the figure). Biofilm is a thin layer at the surface of waterline, where bright green patches represent live bacteria ("LB" in the figure); red patches represent dead bacteria ("DB" in the figure); yellow patches represent coexistence dead and living bacteria ("CDL" in the figure). In the ClO<sub>2</sub> group (left column), bright green patches could not be observed, the intermittent biofilm included large red patches and underneath linear yellow stripes indicating bacteria in the biofilm were disinfected. Moreover, ClO2 had effects of infiltrating into the biofilm (penetrating effects). Whereas in the H2O2 group (right column), bright green patches were still visible indicating the survival of numerous live bacteria indicating the disinfecting effects were not satisfactory. No more penetrating effects were found here, hence, H2O2 exhibited a modest effect on killing the bacteria at the bottom of the biofilm. These data suggested a better efficacy of ClO2 in removal of biofilm. B. Representative images of SEM. In the ClO<sub>2</sub> group (left column), the matrix of biofilm disappeared, indicating its integrity was destroyed. Whereas in the H<sub>2</sub>O<sub>2</sub> group (right column), the biofilm structure was partially damaged. The damaged matrix structure along with the undamaged matrix structure were observed attaching to the surface of waterline. C. Quantitative results of VFS. The ratio of average live/ dead ration of the ClO2 group was significantly lower than that of H2O2 group. Data were presented as mean  $\pm$  standard error, \* means p < 0.05. CLSM: confocal laser scanning microscopy, SEM: scanning electron microscope, VFS: vital fluorescence staining.

killed (red patches representing dead bacteria, and/ or the underneath linear yellow stripes representation coexistence of live and dead bacteria) in the ClO<sub>2</sub> group, whereas they were partially killed as uneven red patches (dead bacteria) over the bright green stripes (representing alive bacteria) in the H<sub>2</sub>O<sub>2</sub> group. Active bacteria (bright green patches) almost could not be observed in the ClO2 group, whereas could be still found in the  $H_2O_2$  group. These findings demonstrated a better disinfecting efficiency of ClO<sub>2</sub> than H<sub>2</sub>O<sub>2</sub> (Figure 1A). Meanwhile, the results of the SEM showed that the matrix structure of biofilms was disrupted from surface to deep layers, thereby the matrix could not be found till the bottom of the biofilm (close to the basal layer), and the bacterial body were exposed in the ClO<sub>2</sub> group. Whereas in the  $H_2O_2$  group, the damage of the matrix structure was slighter, only partial surface layer and matrix were damaged (Figure 1B). Findings of CLSM were in agreements with those of SEM, indicating  $ClO_2$  in comparison to  $H_2O_2$ , can markedly damage the surface (including the matrix) structures and infiltrate into the biofilm, thereby achieves better disinfecting effects (referred to as "penetrating effects"). Quantitative data of VFS also verified the better efficiency of ClO<sub>2</sub> The average live/dead bacteria ratio in ClO<sub>2</sub> group were significantly lower than that of  $H_2O_2$  group (38.41% vs. 66.36%, U = 19/00, p = 0.032) (Figure 1C).

3.3. What is special issue of the  $ClO_2$  solution used in this study?

Disinfection of DUWLs plays a key role in control and prevention of nosocomial infection in a dental clinic. It has been documented that contaminated DUWLs are risky for the patients' health (19), even life-threatening in some extreme cases (20, 21). Removal and control biofilm and planktonic microbes developed in DUWLs are undoubtedly the most important tasks in terms of prevention of DUWLs contamination-related nosocomial infections (22). In this regard, many disinfectants were evaluated. But only several disinfectants were actually applied in the clinical setting. H<sub>2</sub>O<sub>2</sub> as a high-level disinfectant, acts as the most conventional disinfectant using in the DUWLs scenario (23), that is recommended by the manufacturer's manual of many DUWLs makers. However, H<sub>2</sub>O<sub>2</sub> is far from a faultless disinfectant in the context of a dental practice. Its unstable and irritant nature limits its further application for dental practice setting. ClO<sub>2</sub> is another high-level disinfectant which has been considered for using in the dental practice due to its nontoxicity and nonirritant. The limitation of ClO<sub>2</sub> lies in difficulties of availability of a stable and storable  $ClO_2$  solution (14). The aforementioned "activation" processes are quite inconvenient and inoperable in a dental scenario because the activation concentration sometimes is difficult to control. Fortunately, a novel stable ClO<sub>2</sub> solution (free of activation) recently became available. Hence, using ClO<sub>2</sub> solution in the dental practice setting, even directly using it in human body (14)are becoming possible. Here, first, our CFUs study found that the disinfecting efficiency of 5 ppm of stable ClO<sub>2</sub> solution (free of activation) was no weaker than those of conventional 0.24% of H2O2 on DUWLs in actual dental practice (Table 1). During the subsequent morphological studies, we found that the 5 ppm of stable ClO<sub>2</sub> solution exhibited stronger disinfecting effects to biofilm at the surface of the waterline. Results of CLSM indicated that almost all patches representing live bacteria (bright green) were disappeared. Only patches representing dead bacteria (red) and coexistence of live/dead bacteria (yellow) were residual. By contrast, patches of live bacteria remained visible after disinfection with 0.24% of H<sub>2</sub>O<sub>2</sub> (Figure 1A). Importantly, our CLSM data implied that ClO<sub>2</sub> may infiltrate into the biofilm (penetrating effects) and exhibit a better disinfection. The SEM data were in line with the CLSM data, namely 5 ppm of ClO<sub>2</sub> solution could completely destroy the integrity of biofilm, whereas 0.24% of H<sub>2</sub>O<sub>2</sub> could only achieve a partial destroy (Figure 1B). Our data suggested that 5 ppm of ClO<sub>2</sub> solution displayed a stronger effect than 0.24% H<sub>2</sub>O<sub>2</sub> in terms of removal/ control of biofilm. The quantitative results of VFS also confirmed this finding (Figure 1C). Accordingly, the disinfecting efficiency of this stable ClO<sub>2</sub> solution (free of activation) was verified.

Another important issue is regarding the safety. In terms of the application scenarios of ClO<sub>2</sub>, 5 ppm is indeed a very low dose, which is commonly used for disinfection of the fresh fruits and vegetables (24). As early in 1984, a human study by Lubbers et al. documented that no toxic reactions were found after oral intake of 5 ppm of ClO<sub>2</sub> (containing in the tap water) for 12 weeks (25). A later animal study found that no toxic effects were observed in the main organs in mice after oral administration of  $ClO_2$  at 0-40 ppm for 90 days (26). By contrast, the doses of application of ClO<sub>2</sub> for the other scenarios commonly larger, for example, 300 ppm for disinfection of wounds with deep venous thrombosis or diabetic foot (13), 1,000 ppm for dental disinfection (12). These doses of  $ClO_2$  directly used in human body are much greater than 5 ppm, however, are still safe. In this regard, 5 ppm of ClO<sub>2</sub> for DUWLs disinfection is undoubtedly safe.

#### 3.4. Limitations and future prospects

Because the present study was designed in the scenario of dental practice setting, that means all the DUWLs were in practice every day, which required to be disinfected every day. Thus, we could not set up a "blank" control. This might be a limitation of this study. In addition, gradient experiments in different concentrations of  $ClO_2$  and  $H_2O_2$  are also indispensable to elucidate their destroying effects on biofilm, which should be addressed in our future investigation.

Taken together, this pilot study conducted a comparison of the disinfecting effects on DUWLs between a commercialized stable  $ClO_2$  solution (free of activation) and conventional  $H_2O_2$ . The present study verified the satisfactory efficiency of this stable  $ClO_2$  solution in a low dose (5 ppm). The safe and effective nature of stable  $ClO_2$  solution (free of activation) to biofilm indicates that it is suitable for disinfection and sterilization of DUWLs in actual dental practice.

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### Correspondence

# **EQUIBIND:** A geometric deep learning-based protein-ligand binding prediction method

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**SUMMARY** Structure-based virtual screening plays a critical role in drug discovery. However, numerous docking programs, such as AutoDock Vina and Glide, are time-consuming due to the necessity of generating numerous molecular conformations and executing steps like scoring, ranking, and refinement for the ligand-receptor complexes. Consequently, achieving rapid and reliable virtual screening remains a noteworthy challenge. Recently, a team of researchers from Massachusetts Institute of Technology, led by Stärk et al., developed an SE(3)-equivariant geometric deep learning based protein-ligand binding prediction approach, EQUIBIND. In comparison to conventional docking methods, EQUIBIND has the capacity to predict the binding modes of small molecules with target proteins rapidly and precisely. It presents an innovative resolution for high-throughput screening of drug-like compounds.

*Keywords* EQUIBIND, deep learning, virtual screening, protein-ligand binding prediction

Drug discovery is a highly costly and time-intensive process, requiring several years and billions of dollars to discover new drugs that target either newly identified or well-established receptors. Throughout this protracted timeline, issues like drug-induced side effects and adverse reactions can lead to the failure of drug development endeavors (1). The central challenge in drug discovery revolves around deciphering the complex binding interactions between drug-like compounds and proteins. As a result, the accurate and efficient identification of binding sites within proteins for compounds, the characterization of binding conformations, and the evaluation of ligand-protein interactions have become critical aspects of the drug discovery process. In recent decades, computational methodologies have gained significant traction in predicting drug binding sites and conducting virtual screenings to identify potential therapeutic agents (2-4).

In the field of drug design and development, widely used molecular docking approaches such as AutoDock Vina and Glide usually employ strategies based on geometric and energy-based matching principles (5,6). The protein-ligand binding complex undergoes a series of conformational sampling, score ranking and energy-driven refinement (7). Nowadays, driven by the advancement of computational technology, artificial intelligence-driven drug design has emerged as a prominent approach in drug development. Artificial intelligence approach like machine learning, deep learning, and related methodologies have been integrated in traditional virtual screening area. Recently, EQUIBIND, an SE(3)-equivariant geometric deep learning model was reported, which integrates Graph Matching Networks (GMN) (8) and E(3)-equivariant graph neural networks (E(3)-GNN) (9) to predict the binding conformations of ligand-receptor complexes (10). EQUIBIND significantly enhances predictive efficiency by bypassing the need for extensive sampling procedures. It provides direct predictions of binding sites on receptors (blind docking) and ligand binding conformations. A benchmark test demonstrates that EQUIBIND exhibits a processing efficiency of about nine times surpasses than currently prevalent commercial binding predicting programs (10,11).

Moreover, EQUIBIND capitalizes on a K-NN graph (k-nearest neighbor graph) to delineate threedimensional structures. The input parameters predominantly encompass the rigid three-dimensional arrangement of the protein and the adaptable, optimizable conformations of small molecules, which are generated at random using RDKit. Subsequently, EQUIBIND employs Invariant Equivariant Graph Matching Networks (IEGMN) to transform the threedimensional coordinates and predict the binding conformation of the ligand. To ensure accuracy, EQUIBIND employs the SE(3)-equivariant mechanism to calculate the binding sites of receptors, in which the predictive sites are trained to match the actual binding sites. However, EQUIBIND does not guarantee the absolute correctness of complex structures, often resulting in bond lengths and bond angles that deviate from reasonable ranges. To address this issue, EQUIBIND aligns the initial conformations generated by RDKit with those generated by IEGMN. As this process involves only rotatable single bonds, it ensures the local structural integrity. The algorithm primarily focuses on aligning atomic positions around rotatable single bonds, each of which can be manipulated independently. In comparison to traditional docking methods, the EQUIBIND model demonstrates commendable predictive accuracy, enabling precise forecasts of ligand-target binding modes (10).

However, EQUIBIND is not devoid of its limitations. It indirectly models the atomic positions of side chains by encoding features within the receptor  $\alpha$ -carbon graph utilizing localized frameworks, as a result, some predictions may manifest notable disparities. Stärk *et al.* have attempted surface atom refinements and fine-tuning approaches by incorporating atomic subgraphs of the receptor. Regrettably, these endeavors failed to yield significant enhancements to the performance and instead contributed to extended runtimes. Consequently, there remains efforts for the future advancement and refinement of EQUIBIND.

In conclusion, EQUIBIND serves as a novel deep learning-based protein-ligand binding prediction tool. By leveraging SE(3)-equivariant transformations and utilizing minimal input information, it can directly predict the binding sites on receptors and the binding modes of ligands. This capability ensures both high prediction accuracy and a substantial enhancement in prediction efficiency. In the future, whether employed in conjunction with existing virtual screening tools or utilized independently for protein-ligand binding prediction, EQUIBIND holds the potential to emerge as a crucial asset in the drug molecule screening and discovery process.

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# Letter to the Editor

# **Complete trisomy 9 detected by noninvasive prenatal testing and confirmed by amniocentesis**

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**SUMMARY** Complete chromosome 9 trisomy (T9) is a rare and fatal chromosomal disorder. We performed noninvasive prenatal testing (NIPT) in a patient with threatened abortion symptoms and found that the fetal was at risk for complete chromosome 9 trisomy. This shows that NIPT has certain accuracy in detecting trisomy of chromosome 9, which provide options for prenatal diagnosis of rare chromosomal abnormalities.

*Keywords* noninvasive prenatal testing, amniocentesis, complete trisomy 9, early threatened abortion

Letter to the Editor,

Complete T9 which means trisomy of the whole chromosome 9 with no evidence of mosaicism is rarely been reported. This rare an euploidy, accounts for only 2.7 percent of all trisomy, usually leads to early pregnancy miscarriage (1). In rare cases, when the babies are born alive, they usually die during the neonatal period (2,3).

On September 16, 2019, a 26-year old female (gravida 1, para 0) patient at 19 week of pregnancy was admitted to our hospital due to abnormality of chromosome 9. There was no history of present disease and the family history was unremarkable. Ultrasonography performed at 7, 3/7 and 12, 1/7 week of pregnancy showed normal results. Early serological screening at 12,3/7 week of pregnancy showed the risk of trisomy 18 was high at 1: 119 (Greater than or equal to 1/460 is high risk), and the risks of trisomy 21 and trisomy 13 were low. Non-invasive prenatal testing (NIPT) at 14,5/7 wk of pregnancy suggested an abnormal Z-score of fetal chromosome 9 at 15.340 (the normal range was Z-score > 3 or < 3). The result of amniocentesis performed at 19 week of pregnancy found the final karyotyping of the fetus was 47, XX, +9, and complete T 9 was confirmed. Transabdominal level II ultrasound performed at 21, 6/7 wk of gestation showed intrauterine growth restriction (IUGR), enlargement of anterior fontanelle (Figure 1A), widened eye distance (Figure 1B), cleft lip and palate (Figures 1C and 1D), high echo of left kidney has, widened renal collecting system and missing image of right renal echo (Figure 1E). Transverse section of the bladder, with only one umbilical artery visible on the right side of the fetus. In transverse view of the bladder, only one umbilical artery was observed at the right (Figure 1F). The four-chamber view revealed heart enlargement (Figure 1G) and pulmonary artery broadening (Figure 1H). No gallbladder echo was found in this test, that means absence of the gallbladder.

The patient requested termination of the pregnancy. At 22,1/7 week of pregnancy, the patient received ultrasound-guided amniotic cavity 6, 9-diamino-2ethoxylacridine lactate hydrate (Rivanol, Guangxi Hefeng Pharmaceutical Co., Ltd., China) injection for labor induction. The fetus was stillborn 1 day after Rivanol injection.

NIPT for fetal chromosome detection has been widely used in recent years for further prenatal confirmation (4). It is mainly used for screening T21, T18, and T13 (5) and the use of NIPT in detecting T9 is rarely reported. The chromosome 9 abnormality suggested by NIPT in our case was consistent with the results of late amniocentesis. It is suggested that NIPT has accuracy in the early detection of complete T9. In addition, compared with amniocentesis which is suitable at 15-22 weeks of gestation, and umbilical vein puncture which is suitable at late than18 weeks of gestation (6), NIPT can be used in early gestational weeks with higher security for it doesn't increase the rate of miscarriage and has no contraindications (7).

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Figure 1 Imaging findings. (A) The arrow denotes the site of large anterior fontanelle, (B) widened eye distance, (C) cleft lip, (D) cleft palate, (E) widened renal collecting system, (F) Single umbilical artery, (G) large heart and (H) Widened pulmonary artery.

It has been reported that the fetal free DNA can be detected as early as 5 weeks of gestation in peripheric blood of pregnant women, and the detection rate is higher in the second trimester (8). Compared with transabdominal chorionic villus sampling (TA-CVS) that can be performed in the earlier gestational weeks (10-14 weeks) (9), NIPT is a non-invasive operation, while the TA-CVS has certain effects on the fetus such as fetal extremity developmental impairment, abortion, *etc.* (10). Our case reported a complete T9 with signs of threatened abortion, detected by NIPT, finally confirmed by amniocentesis and ultrasonic testing. To our knowledge, there is no similar literature at present. And it indicates that NIPT can be used for early detection of complete T9.

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