Amotosalen and ultraviolet A light treatment efficiently inactivates severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) in human plasma

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Abstract

**Background and objectives** During the ongoing pandemic of COVID-19, SARS-CoV-2 RNA was detected in plasma and platelet products from asymptomatic blood donors, raising concerns about potential risk of transfusion transmission, also in the context of the current therapeutic approach utilizing plasma from convalescent donors. The objective of this study was to assess the efficacy of amotosalen/UVA light treatment to inactivate SARS-CoV-2 in human plasma to reduce the risk of potential transmission through blood transfusion.

**Methods** Pools of three whole-blood-derived human plasma units (630–650 ml) were inoculated with a clinical SARS-CoV-2 isolate. Spiked units were treated with amotosalen/UVA light (INTERCEPT Blood System™) to inactivate SARS-CoV-2. Infectious titres and genomic viral load were assessed by plaque assay and real-time quantitative PCR. Inactivated samples were subject to three successive passages on permissive tissue culture to exclude the presence of replication-competent viral particles.

**Results** Inactivation of infectious viral particles in spiked plasma units below the limit of detection was achieved by amotosalen/UVA light treatment with a mean log reduction of >3.32 ± 0.2. Passaging of inactivated samples on permissive tissue showed no viral replication even after 9 days of incubation and three passages, confirming complete inactivation. The treatment also inhibited NAT detection by nucleic acid modification with a mean log reduction of 2.92 ± 0.87 PFU genomic equivalents.

**Conclusion** Amotosalen/UVA light treatment of SARS-CoV-2 spiked human plasma units efficiently and completely inactivated >3.32 ± 0.2 log of SARS-CoV-2 infectivity, showing that such treatment could minimize the risk of transfusion-related SARS-CoV-2 transmission.

**Keywords:** amotosalen/UVA, pathogen inactivation, SARS-CoV-2, plasma.
Introduction

A key element of blood safety is the prevention of transfusion-transmitted infections. Multiple safety measures were introduced in the last decades, strongly accelerated by the recognition of HIV transfusion transmission in the early 1980s, significantly decreasing the risk for infectious adverse events. However, there is still a remaining risk for the recipients of labile blood components, considering the limitations of existing safety measures [1]. Newly emerging pathogens, particularly viruses, may pose a challenge to the safety of plasma transfusion, especially if subclinical infection occurs in combination with blood viremia. The majority of individuals infected by one of the recently emerging arboviruses develop no symptoms but may carry a high titre viral load in the blood [1,2]. The emergence and risk for the blood supply caused by emerging infectious diseases are unpredictable [3], as demonstrated by the current COVID-19 pandemic.

The first cases of COVID-19 disease, an infection of the lower respiratory tract, were reported in late 2019, spreading rapidly in and from Wuhan, China [4]. The highly virulent pathogen causing the disease was identified as a beta-Coronavirus (SARS-CoV-2) [5]. On 11 March 2020, the WHO declared a global COVID-19 pandemic, up to date more than 47 million confirmed infections and more than 1.2 million disease-related deaths have been reported globally, with highest disease burden in Europe and the Americas, followed by the Middle East and Asia. In Saudi Arabia, more than 348,000 confirmed cases and more than 5000 disease-related deaths have been reported up to date according to the WHO COVID-19 Dashboard [6]. Respiratory droplets were quickly identified as the main route of infection, other ways of transmission, especially smear infection/oral intake, may also play a role [7]. Not every infected individual develops the COVID-19 disease, the prevalence of asymptomatic infected individuals has been reported by several studies [7,8] including a study from South Korea that reported a prevalence of 19.2% [9].

SARS-CoV-2 is closely related to other human coronaviruses, especially SARS-CoV which caused an outbreak of the Severe Acute Respiratory Syndrome in Singapore and Korea in 2000–2002. By decision of the International Committee for the Taxonomy of Viruses, SARS-CoV and SARS-CoV-2 are different strains of the same species [5]. Transmission by blood transfusion was not shown for SARS-CoV, however, due to the detection of viral genomes in blood in the symptomatic phase, and the detection of SARS-CoV genomes in leucocytes in the convalescent phase together with evidence for subclinical infection, transmission of SARS-CoV is of concern and considered a theoretical risk [10]. Evidence for subclinical infection was also shown for SARS-CoV-2 [7]. Low viral load SARS-CoV-2 RNAemia in blood was detected frequently in symptomatic patients [11,12]. A recent study from China reported SARS-CoV RNAemia also in asymptomatic blood donors; viral nucleic acids were detected retrospectively in frozen plasma products, a platelet product and donor samples for pre-screening [13]. Although transmission of SARS-CoV-2 is not documented to date, the detection of viral nucleic acids in blood products from asymptomatic donors raises concerns about potential transfusion transmission of SARS-CoV-2, especially because blood screening procedures for that pathogen are not established yet. Another potential risk became evident in light of the recent approaches to use convalescent donors’ plasma to treat COVID-19 patients, an idea that seems to be one of the most promising therapeutic approach to date [14].

The safety of blood donations has been of major concern for transfusion medicine practitioners and researchers, this has led to the introduction of certain measures to reduce this risk including risk factor questionnaires, screening assay to detect transfusion-transmitted pathogens in blood donations [15]. These measures have led to decreased prevalence of some of the transfusion-transmitted infections such as HIV, HCV and HBV [16]. The increased improvement in the performance of screening assays was very helpful in reducing transmission risks. These assays are pathogen specific and are of no value to detect newly emerging and re-emerging pathogens. Therefore, it is worthwhile to consider treating the donated plasma with a pathogen inactivation technology proven to be efficient against a large panel of pathogens including SARS-CoV-2 [17].

The INTERCEPT technology inactivates a broad spectrum of viruses, bacteria and protozoa in plasma prepared for transfusion [18]. During a photochemical reaction involving amotosalen (the photoactive compound) and UVA light, the pathogens’ genomes are modified in a targeted and specific manner by cross-linking the genomic strands, preventing transcription and replication without affecting the clinical outcome of the plasma transfusion [19,20]. This was also shown in a cohort of thrombotic thrombocytopenic purpura (TTP) patients requiring therapeutic plasma exchange [21,22]. As an additional effect, residual white blood cells of the donor are inactivated more efficiently than by gamma-irradiation [23], reducing the risk for immunological transfusion reactions and transfusion-associated graft-versus-host disease (TA-GvHD). Efficient inactivation by amotosalen/UVA treatment of the very closely related SARS-CoV was reported in human plasma units and platelet concentrates [24,25]. We also recently have shown that amotosalen/UVA

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treatment can efficiently inactivate the closely related Middle East Respiratory Syndrome Coronavirus (MERS-CoV) in human plasma [26] and platelet concentrates [27], and now extending this work to evaluate the inactivation of SARS-CoV-2 in human therapeutic plasma units.

Materials and methods

Cell line and SARS-CoV-2 culture

Vero E6 cells were maintained in Dulbecco’s modified Eagle medium (DMEM) with 10% foetal bovine serum (FBS) as previously described (Al-Amri et al, 2017). A human SARS-CoV-2 clinical patient isolate (SARS-CoV-2/human/SAU/85791C/2020, gene bank accession number: MT630432) was used in all experiments in the Special Infectious Agents Unit (SIAU) Biosafety level 3 facility at King Fahd Medical Research Center (KFMRC), King Abdulaziz University (KAU), Jeddah, Saudi Arabia.

Preparation of SARS-CoV-2 stock

Virus was inoculated on 90%–95% confluent Vero E6 cells (ATCC# CRL-1586) in a T175 tissue culture flask at multiplicity of infection (MOI) of 1 and incubated in a humidified incubator at 5% CO2 and 37°C for one hour with gentle shaking every 15–20 min. Subsequently, the inoculum was replaced by 25 ml of viral inoculation medium (DMEM with 2% FBS, 1% penicillin/streptomycin and 10 mmol/l HEPES [pH 7.2]) and the cells were incubated in a humidified incubator at 5% CO2 and 37°C for one hour at room temperature. Virus was then aliquoted and stored at –80°C, and the titre was determined by plaque assay.

Plasma preparation

Whole blood units (450 ml ± 10%) were collected and prepared at King Abdulaziz University Hospital, Transfusion Services, Jeddah, Kingdom of Saudi Arabia, from voluntary donors. Briefly, blood units were centrifuged at 544 × g for 10 min to separate the platelet (PLT)-rich plasma. The PLT-rich plasma was then centrifuged at 230 × g at 20°C for 10 min. The plasma was then transferred into the plasma bag and kept at –30°C. All blood units were screened routinely for HCV antibody, HBsAg, HBe antibody, HIV (1/2) antibody, HTLV (1/2) antibody, Syphilis as well as HCV, HBV and HIV by NAT.

SARS-CoV-2 inactivation

Pools of three single-donor plasma units were used in this study (volume 630–650 ml). All plasma pools were tested for the presence of anti-SARS-CoV-2 neutralizing antibodies using an in-house neutralization assay. Each plasma pool was inoculated with SARS-CoV-2 stock in a 1:100 dilution. Plasma units were treated with amotosalen/UVA using the INTERCEPT Processing Set for Plasma and an INTERCEPT Illuminator INT-100 (Cerus Corporation, U.S.A.). The following samples were collected for testing: a positive control sample from the virus stock; a negative control sample from the plasma before inoculation with the virus; a pretreatment sample after the addition of amotosalen and a sample from the treated plasma units after INTERCEPT illumination (Fig. 1). All samples were stored at –80°C until testing.

Detection of replicating SARS-CoV-2

Detection of replicating SARS-CoV-2 in amotosalen/UVA-treated plasma was performed as previously described for SARS-CoV-1 and MERS-CoV with minor modifications [25,26]. Briefly, collected pretreatment and inactivated samples were diluted at 1:10 dilution in DMEM with 10% FBS, inoculated on Vero E6 cells in 6 well plates in duplicates and incubated for 1 h at 37°C. Inoculum was then removed and replaced with 2 ml DMEM with 10% FBS and incubated for 3 days at 37°C. Then, supernatants were collected, diluted 1:100 in DMEM with 10% FBS and re-inoculated on Vero E6 cells for two more successive passages. Supernatants collected from each passage were also used for viral load quantification by RT-qPCR.

Plaque assay

Plaque assays were performed as previously described [26] with minor modifications. Briefly, samples were serially diluted in DMEM with 10% FBS starting from 1:10 and 1 ml from each dilution was inoculated on confluent Vero E6 cell monolayers and incubated for 1 h at 37°C. Then, the inoculum was removed and overlaid with DMEM containing 0.8% agarose and incubated for 3 days at 37°C. Cells were then stained with crystal violet for 4 h at 37°C, and plaques were counted to determine the viral titre as plaque forming unit (PFU)/ml.

RT-qPCR quantitation

Viral RNA was extracted from all samples collected directly from the plasma units (positive, negative,
pretreatment and post-treatment samples) using the QIAamp Viral RNA Mini Kit (Qiagen, Germany) according to the manufacturer’s instructions. Relative quantification of the SARS-CoV-2 viral load was performed by one-step dual-target real-time RT-PCR (RealStar SARS-CoV-2 RT-PCR Kit 1.0, Altona Diagnostics, Germany) according to the manufacturer’s instructions using a 7500 Fast Real-Time PCR System (Applied Biosystems, U.S.A.). The PCR detects a beta-coronavirus specific target (E-gene), a SARS-CoV-2 specific target (S-gene) and an internal control. The decrease in viral load was expressed by comparing the cycle threshold (CT) values from each sample relative to the CT values of the pretreatment inoculated sample (with the SARS-CoV-2 specific primers). The SARS-CoV-2 titres were expressed as PFU equivalents per ml (PEq/ml) using a standard curve (standard: serial dilutions of the viral stock) and choosing dilutions of the original sample (10^{-1} to 10^{-8}) with CT values in the exponential phase. Each run included a positive viral template control and no-template negative control. Each sample was tested in duplicate, and the mean is reported as PEq/ml.

Figure 1 Experimental design. Schematic view of the experimental design. Briefly, pools of 3 plasma units were spiked in a 1:100 dilution with a SARS-CoV-2 viral stock. The pools were treated with amotosalen/UVA pathogen inactivation and post-inactivation subject to 3 consecutive passages of 3 days respectively on Vero E6 cells. Samples were taken at various stages and assessed for replicating virus by plaque assay and viral genomes by RT-qPCR.
IRB approval

The study was approved by the Unit of Biomedical Ethics of the King Abdulaziz University Hospital (approval # 285-20).

Results

Inactivation of SARS-CoV-2 in human plasma units

Five pools (A-E) of human plasma collected from healthy donors were spiked with SARS-CoV-2. Spiked units were then treated with amotosalen and UVA light. The mean infectious viral titre in pretreatment samples was $3.32 \pm 0.19 \log_{10} \text{PFU/ml}$ (range: $3.1-3.6 \log_{10} \text{PFU/ml}$) (Table 1). Treatment of spiked units resulted in a mean reduction of $>3.32 \pm 0.19 \log_{10} \text{PFU/ml}$ as no infectious virus was detected by plaque assay (Table 1). Figure 2 shows a representative plaque assay result for all tested units. As expected, testing of negative control samples collected before spiking with SARS-CoV-2 showed no replication-competent virus. Of note, the viral infectivity titre in viral stock was $5.6 \pm 0.2 \log_{10} \text{PFU/ml}$ and it was reduced to a mean infectivity of $3.32 \pm 0.19 \log_{10} \text{PFU/ml}$ after spiking in a 1:100 dilution, which is close to an expected post-dilution titre of $3.6 \log_{10} \text{PFU/ml}$. Considering additional dilution of the spiked plasma by the addition of amotosalen solution, the dilution of the viral stock in plasma did not lead to unexpected loss of viral infectivity.

The impact of pathogen inactivation treatment on the genomic viral load

To further confirm these results, the viral genomic viral load was determined for all collected samples. The median CT preinactivation was 25-6 (18-3-29-1). The pretreatment samples mean genomic viral load was $2.92 \pm 0.87 \log_{10} \text{PEq/ml}$, which was in the same range as the infectious titre ($3.32 \pm 0.19 \log_{10} \text{PFU/ml}$). After treatment, no viral genomes were detectable by PCR (Table 2). The internal PCR control was always positive indicating no PCR inhibition and confirming the decrease in the signal from the SARS-CoV-2 specific primers is due to a mean minimum inactivation of $2.92 \pm 0.87 \log_{10} \text{PEq/ml}$.

Passaging of INTERCEPT treated plasma to confirm complete inactivation

To exclude the possibility of any remaining replicating SARS-CoV-2 associated with the presence of viral genomic load in the treated plasma units (Table 2), we inoculated the collected samples on Vero E6 cells and evaluated infectivity over three successive passages. While culture of all pretreatment samples showed viral replication and complete CPE within 3 days post-inoculation similar to that of positive control, neither viral replication nor CPE was observed in cells inoculated with inactivated samples similar to negative controls (Fig. 3) even after 9 days of incubation in all three passages. For further confirmation, we determined the genomic viral load from supernatants collected from all passages inoculated with either pretreatment or post-treatment samples. As shown in Table 3, passaging of pretreatment samples showed viral replication as evident by CPE. On the other hand, viral genomes in cells inoculated with inactivated samples in culture supernatants were not detectable. Together, these data confirm the complete inactivation of SARS-CoV-2 in the tested platelets units and the absence of replication-competent virus post-inactivation.

Discussion

Newly emerging pathogens are an ongoing potential threat for safety of the blood supply. Pathogen inactivation technology may provide immediate enhanced safety in an outbreak setting, in contrast to diagnostic tools which, even if they already exist, need to be developed and validated for multiple weeks to months [28].

We were able to show complete inactivation of $>3.32 \pm 0.19 \log \text{PFU/ml}$ and $2.92 \pm 0.87 \log \text{PEq/ml}$ SARS-CoV-2 in human plasma pools of 630–650 ml in the present study, corresponding to volumes usually collected by plasmapheresis for the production of convalescent plasma (CP). Since a clinical isolate was used, the maximum titre obtained in vitro was not very high. Higher in vitro infectivity titres could likely show higher minimum inactivation efficacy. No CPE and no genomic viral load were detectable after 3 consecutive rounds of passaging on Vero E6 cells, pointing towards complete inactivation. The reported genomic viral load in the blood of symptomatic patients and asymptomatic donors is relatively low, often close to the limit of detection [11–13,29], pointing towards sufficient inactivation efficacy of amotosalen/UVAs to prevent potential transfusion transmission (Table 2).

Recently, the inactivation of a different SARS-CoV-2 isolate (USA-WA1-2020) from the USA with Riboflavin/UVB in single-donor plasma units (>3-4 log PFU/ml) and platelets (>4-53 log PFU/ml) was reported [30]. Another study using Riboflavin/UVB technology with the same isolate reported an inactivation efficacy of $>4.59 \pm 0.15 \log \text{PFU}$ in single-donor plasma units and $>3.30 \pm 0.26 \log \text{PFU}$. © 2020 The Authors. Vox Sanguinis published by John Wiley & Sons Ltd on behalf of International Society of Blood Transfusion.
Table 1 Reduction of infectious SARS-CoV-2 titres in human plasma units after amotosalen/UVA treatment

| Experiment | Positive control | Negative control | Pretreatment sample | Inactivated sample | Log reduction |
|------------|------------------|------------------|---------------------|-------------------|---------------|
| A          | 5.5              | ND               | 3.4                 | ND                | >3.4          |
| B          | 5.7              | ND               | 3.2                 | ND                | >3.2          |
| C          | 5.6              | ND               | 3.6                 | ND                | >3.6          |
| D          | 5.9              | ND               | 3.3                 | ND                | >3.3          |
| E          | 5.3              | ND               | 3.1                 | ND                | >3.1          |
| Mean ± SD  | 5.6 ± 0.20       | ND               | 3.32 ± 0.19         | ND                | >3.32 ± 0.19  |

ND indicates not detected.
*After addition of amotosalen.
†This indicates complete inactivation of the virus.

Figure 2 Inactivation of SARS-CoV-2 in plasma by amotosalen and UVA treatment assessed by a plaque assay. Vero E6 cells were inoculated for 1 h with the following samples in a 1:10 dilution in DMEM: the SARS-CoV-2 viral stock (Positive Control), human plasma (Negative Control), plasma from a SARS-CoV-2 spiked pretreatment sample (Spiked Plasma Unit) and amotosalen/UVA-treated, SARS-CoV-2 spiked plasma (Post-iInactivation). The cells were overlaid with agarose, incubated for three more days followed by neutral red staining. Experiments were conducted in serial dilutions. Photographs (4×) are shown from one of five representative experiments. [Colour figure can be viewed at wileyonlinelibrary.com]

Table 2 SARS-CoV-2 genomic load in plasma before and after amotosalen/UVA treatment*,†

| Experiment | Positive control | Negative control | Pretreatment sample | Inactivated sample |
|------------|------------------|------------------|---------------------|-------------------|
| A          | 5.99             | ND               | 3.60                | ND                |
| B          | 5.88             | ND               | 2.82                | ND                |
| C          | 5.53             | ND               | 3.96                | ND                |
| D          | 5.36             | ND               | 2.44                | ND                |
| E          | 5.18             | ND               | 1.79                | 0.24              |
| Mean ± SD  | 5.59 ± 0.34      | ND               | 2.92 ± 0.87         | 0.04 ± 0.11       |

ND indicates not detected.
*Data are shown as log_{10} PEq/ml.
†Titres were determined from the same samples used in Table 1.
log PFU/ml in whole blood [31]. Both Riboflavin/UVB studies did not report passaging experiments or used sensitive NAT testing to confirm the findings of the plaque assay used in these studies.

The current study also supports the use of amotosalen/UVA for mitigating the risk of potential superinfection with SARS-CoV-2 through convalescent plasma.

The use of pathogen inactivation technology for CP also raises the question of the impact on product quality, hence the neutralizing activity. A study with Ebolavirus CP (EBOV CP) showed no significant impact on the neutralizing activity by amotosalen/UVA treatment [32]. A single case study showed a loss of 2–4% total IgG after amotosalen/UVA treatment of EBOV CP without assessing the consequences for the neutralizing activity [33]. Recently, no significant impact of amotosalen/UVA treatment on the neutralizing activity and neutralizing antibody quantity of COVID-19 convalescent plasma (CCP) was reported [34]. These reports point towards the preservation of neutralizing activity during amotosalen/UVA treatment.

**Conclusion**

In the present study, we showed efficient, complete inactivation of SARS-CoV-2 in large human plasma units by amotosalen/UVA using a local clinical isolate, which may serve as an additional layer of safety in plasma transfusion during the COVID-19 pandemic. These findings are in line with former findings of efficient inactivation of human coronaviruses in plasma by amotosalen/UVA pathogen inactivation [24,26].

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**Conflict of interest**

There is no conflict of interest identified.

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Table 3 Results of passaging experiments of SARS-CoV-2 in Vero E6 cells before and after inactivation of spiked plasma*,†

| Experiment | Passage 1 | Passage 2 | Passage 3 |
|------------|-----------|-----------|-----------|
| A          | 5.05      | 4.54      | 4.14      |
| B          | 5.12      | 4.61      | 4.31      |
| C          | 5.24      | 4.05      | 4.01      |
| D          | 4.96      | 4.01      | 3.95      |
| E          | 5.16      | 4.21      | 3.97      |

*Data are shown as log10 PFU/ml.

Samples in Table 1 were used in this experiment. Samples were used at 1:10 dilution, and titre was determined on day 3 post-inoculation.

ND indicates not detected.
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