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RECEIVED 08 July 2023

ACCEPTED 12 October 2023

PUBLISHED 27 October 2023

CITATION

Urwylker P, Leimbacher M, Charitos P, Moser S, Heijnen IAFM, Trendelenburg M, Thoma R, Sumer J, Camacho-Ortiz A, Bacci MR, Huber LC, Stüssi-Helbling M, Albrich WC, Sendi P and Osthoff M (2023) Recombinant C1 inhibitor in the prevention of severe COVID-19: a randomized, open-label, multi-center phase IIa trial. *Front. Immunol.* 14:1255292. doi: 10.3389/fimmu.2023.1255292

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Recombinant C1 inhibitor in the prevention of severe COVID-19: a randomized, open-label, multi-center phase IIa trial

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Background: Conestat alfa (ConA), a recombinant human C1 inhibitor, may prevent thromboinflammation.

Methods: We conducted a randomized, open-label, multi-national clinical trial in which hospitalized adults at risk for progression to severe COVID-19 were assigned in a 2:1 ratio to receive either 3 days of ConA plus standard of care (SOC) or SOC alone. Primary and secondary endpoints were day 7 disease severity on the WHO Ordinal Scale, time to clinical improvement within 14 days, and safety, respectively.

Results: The trial was prematurely terminated because of futility after randomization of 84 patients, 56 in the ConA and 28 in the control arm. At baseline, higher WHO Ordinal Scale scores were more frequently observed in the ConA than in the control arm. On day 7, no relevant differences in the primary outcome were noted between the two arms ($p = 0.11$). The median time to defervescence was 3 days, and the median time to clinical improvement was 7 days in both arms ($p = 0.22$ and 0.56 , respectively). Activation of plasma cascades and endothelial cells over time was similar in both groups. The incidence of adverse events (AEs) was higher in the intervention arm (any AE, 30% with ConA vs. 19% with SOC alone; serious AE, 27% vs. 15%; death, 11% vs. 0%). None of these were judged as being related to the study drug.

Conclusion: The study results do not support the use of ConA to prevent COVID-19 progression.

Clinical trial registration: <https://clinicaltrials.gov>, identifier NCT04414631.

KEYWORDS

COVID-19, randomized trial, C1 esterase inhibitor, complement system, kallikrein kinin system, contact activation system

1 Introduction

Numerous efforts have been made to develop effective treatments for individuals who have been hospitalized with Coronavirus disease 2019 (COVID-19) (1). The pathogenesis of severe COVID-19 is related to SARS-CoV-2 infection and secondary to a dysregulated host immune response (2). In line with the excessive inflammatory response, coagulation activation plays a critical role (3, 4). Within the coagulation pathways, the importance of targeting the complement system (CS), the kinin-kallikrein system (KKS), and the contact activation system (CAS) to curb the progression of COVID-19 to severe disease has been supported by several observational studies, and the rationale to inhibit the CS with a therapeutic candidate has been outlined by us and others (5–9). Lower complement C1q values have been reported in patients with severe COVID-19, indicating relevant complement activation and consumption of this protein (10, 11). Within the CS, C1 inhibitor (C1INH) is a strong inhibitor of C1r/C1s and MASP-1/-2. In addition, C1INH also serves as an inhibitor of factor XII of the CAS as well as kallikrein of the KKS (12). Hence, C1INH may contribute to the deterrence of COVID-19 disease progression by prevention of microthrombi generation, inhibition of leucocyte activation, and interaction with endothelial cells and microorganisms (13, 14).

Conestat alfa (Ruconest[®], Pharming, Leiden, The Netherlands, abbreviated as ConA in this manuscript) is a recombinant human C1INH and shares an identical protein structure with plasma-derived C1INH (pdC1INH). It has a different glycosylation pattern, which is responsible for a shorter half-life (3 h) than pdC1INH (30 h) (15, 16), and seems to target the activation of the lectin pathway more effectively (17). A pilot study using conestat alfa in five patients with severe COVID-19 showed promising results with immediate defervescence, and decrease of inflammatory markers and oxygen supplementation in four of five patients (18). To evaluate the efficacy and safety of ConA for SARS-CoV-2 infection, we conducted a randomized, open-label clinical trial in adult patients hospitalized with COVID-19 (PROTECT-COVID-19 trial).

2 Materials and methods

2.1 Study design and patients

This was a randomized, parallel-group, open-label, multi-center trial. The study protocol was published elsewhere (19). The study was performed in accordance with the Declaration of Helsinki and approved by the institutional review board and ethics committees for each participating institute. All participants gave written informed consent. An independent data-safety monitoring board (DSMB) committee periodically reviewed the study outcomes. The study was conducted at five sites in three countries (three sites in Switzerland and one site each in Brazil and Mexico) and registered at ClinicalTrials.gov (NCT04414631).

Male and non-pregnant female patients (18–85 years of age) were eligible and approached by study physicians of the participating centers if they had a diagnostic specimen that was positive for SARS-CoV-2 on reverse-transcriptase-polymerase-chain-reaction (RT-PCR), had pneumonia confirmed by chest imaging, and onset of symptoms ≤ 10 days or shortness of breath ≤ 5 days. The latter criterion was adapted/added with an amendment 4 months after approval of the first study protocol, because of the changing patient behaviors over time when seeking for medical help after a positive SARS-CoV-2 test. In addition, at least one of the following risk factors for disease progression was required for enrollment: arterial hypertension, age ≥ 50 years, obesity (BMI ≥ 30.0 kg/m²), cardiovascular disease, chronic pulmonary disease, chronic renal disease, C-reactive protein (CRP) of >35 mg/L, or oxygen saturation of $\leq 94\%$.

Exclusion criteria included refusal to participate, history or suspicion of allergy to rabbits or the study drug, pregnant or breast-feeding individuals, known severe liver disease, and renal disease requiring dialysis. Patients who received tocilizumab or another IL-6 or IL-6 receptor inhibitor prior to enrollment were not eligible for this study. Patients currently requiring intensive care (or expected to do so) within 24 h after initial presentation were not eligible for this study, because “prevention of disease progression” was investigated. Hospitalized patients requiring high-flow oxygen therapy on medical wards were eligible for inclusion.

2.2 Standard-of-care therapy, randomization, and study medication

Standard of care (SOC) therapy: All patients received SOC therapy. It was defined as any of the following treatment options, in addition to oxygen and fluid supply: dexamethasone, remdesivir, empiric antimicrobial treatment when bacterial co-infection was suspected, and anti-thrombotic treatment, according to in-house guidelines, aligned to published recommendations (20, 21). Other treatments such as tocilizumab, baricitinib, anti-SARS-CoV-2 monoclonal antibodies, and nirmatrelvir/ritonavir were not available or recommended in national guidelines during the conduct of the study.

2.2.1 Randomization

Study participants were randomized in a 2:1 ratio using permuted-block allocation with varying block sizes. Randomization was stratified by the study site, generated by an independent statistician, and implemented via the electronic data capture system software SecuTrial[®]. There was no blinding of study medications.

2.2.2 Intervention arm

Patients allocated to the intervention arm received a total of nine intravenous ConA doses over 72 h (8,400 U followed by 4,200 U every 8 h), in addition to SOC treatment.

2.2.3 Control arm

Patients allocated to the control arm received SOC only.

2.3 Monitoring, outcomes, and follow-ups

Data on vital signs, disease severity according to WHO scale, clinical improvement, admission to ICU, and requirement for non-invasive or invasive ventilation were prospectively obtained. Virological clearance (time from positive to negative PCR test results in nasopharyngeal samples), hematological, liver, renal, and inflammatory parameters in laboratory examinations were monitored during the first 14 days of hospitalization or until discharge. C1INH antigen and C4 concentrations were determined on semi-automated platforms in the clinical laboratories in all except two study centers (not monitored in Monterrey, Mexico and Zurich, Switzerland). Additional EDTA- and citrate-plasma as well as serum samples were frozen at -80°C and subsequently analyzed for concentrations of C4d (Svar Life Sciences, Malmö, Sweden), sC5b-9 (BD Biosciences Pharmingen, San Diego, CA, USA), VCAM-1 and E-selectin (ProteinSimple, Biotechne, Minneapolis, MN, USA), and kallikrein-like activity (Unitest, Haemochrome Diagnostica, Essen, Germany) according to the manufacturer's instruction.

Final follow-up included a structured telephone interview or study visit (if still hospitalized) 28 days after enrollment to assess adverse events (AEs) and outcome. Afterwards, serious adverse events (SAEs) were followed until resolution, stabilization, or death, whichever occurred first.

The primary endpoint was disease severity on the WHO Ordinal Scale on day 7. For statistical analysis, the scores 6 and 7 were combined into one score. Secondary endpoint included time to clinical improvement within 14 days after enrollment. It was defined as improvement of two points on the WHO Ordinal Scale or live hospital discharge, whatever occurred first. Other secondary endpoints included the proportion of participants alive and not having required invasive or non-invasive ventilation at day 14 day, and the proportion of subjects with an acute lung injury. The latter was defined by the presence of $\text{PaO}_2/\text{FiO}_2$ ratio ≤ 300 . Further endpoints included the length of hospital stay until day 28 in survivors, and all-cause mortality.

2.4 Safety evaluations

The overall incidence of AEs and SAEs were assessed during a 4-week follow-up and documented by the investigators according to seriousness, intensity, causal relationship with study treatment, action taken with study treatment (e.g., withdrawal), specific treatment for AE, and outcome.

AE and SAEs were grouped by the organ class system (22). There were no prespecified futility margins for the DSMB; however, the DSMB was entitled to recommend termination of the study.

2.5 Statistical plan and analysis

Full analysis set and intent-to-treat population (FAS/ITTP) was defined as all patients who were randomly allocated to a study arm. The primary efficacy analysis was based on the FAS/ITT population. The primary endpoint was analyzed by Wilcoxon test stratified by its baseline values with a two-sided α -level of 5%. The secondary endpoints were tested after a significant test of the primary endpoint (*a priori* ordered hypotheses); therefore, no alpha adjustment was necessary.

Quantitative secondary study parameters were described based on their mean, standard deviation (SD), median, interquartile range (IQR), minima, and maxima and portrayed by Kaplan–Meier plots and compared with the log rank test. Qualitative secondary study parameters were analyzed by means of absolute and relative frequencies: Chi-square tests were carried out in order to compare the active treatments to SOC. With a two-sided significance level of $\alpha = 0.05$ and a power of $1 - \beta = 0.80$, a sample size of 76 (2×38) was calculated for a fixed sample size design and an advantage of at least 1 point on the WHO Ordinal Scale (SD 1.5 points) (19). For a 2:1 randomization, a nonparametric analysis by the stratified Wilcoxon test, and an adaptive group sequential analysis, the overall sample size was estimated as $120 = 80 + 40$. Two adaptive interim analyses after 40 and 80 patients were planned according to the Pocock adjusted levels $\alpha_p = 0.0221$. The analyses were carried out by the contract research organization CRM Biometrics GmbH (Bornheim, Germany).

The independent clinical trial statistician of CRM Biometrics GmbH, the first author (PU), and last author (MO) analyzed the data and had access to all data. All authors had access to primary clinical trial data only related to their institution. Study participants,

treating physicians, nurses, investigators, and DSMB members were not blinded.

3 Results

3.1 Patient characteristics

Between August 2020 and March 2021, 621 patients were screened and 84 were enrolled (Supplementary Figure S1). Of these, 56 were randomized to the intervention and 28 to the control arm. One patient of the latter group withdrew consent after randomization (i.e., 27 for final analysis in the control arm). Screening failure reasons are cited in Supplementary Table S1. Fifty percent of the patients were recruited at the principal investigator’s institute (Supplementary Table S2). The baseline characteristics are presented in Table 1. Participants in the intervention arm presented with more severe disease at enrollment as assessed by the WHO scale score and the requirement of oxygen supplementation than the ones in the control arm, although the differences were not statistically significant. Corticosteroids such as dexamethasone were administered during the study in 42/56 (75%) and 19/27 (70%) of patients in the intervention and control arm, respectively.

TABLE 1 Baseline characteristics.

| | Intervention arm (n = 56) | Control arm (n = 27) |
|---|---------------------------|----------------------|
| Age (years), mean (SD) | 61.1 (13.9) | 64.4 (15.3) |
| Female gender, n (%) | 17 (30) | 10 (36) |
| BMI (kg/m ²), mean (SD) | 29.1 (6.1) | 26.4 (4.5) |
| Shortness of breath, n (%) | 42 (75) | 15 (54) |
| Fever, n (%) | 36 (64) | 13 (46) |
| Cough, n (%) | 43 (77) | 18 (64) |
| Arterial hypertension, n (%) | 27 (49) | 15 (54) |
| COPD/chronic lung disease, n (%) | 16 (29) | 6 (22) |
| Cardiovascular disease, n (%) | 12 (21) | 6 (22) |
| Diabetes mellitus, n (%) | 15 (27) | 8 (29) |
| Heart failure, n (%) | 5 (9) | 4 (14) |
| Smoking (past and present), n (%) | 27 (48) | 8 (30) |
| Anticoagulation, n (%) | 10 (18) | 4 (14) |
| ACE/ARB, n (%) | 23 (41) | 13 (48) |
| Oxygen supplementation (L), mean (SD) | 4.6 (8.4) | 2.1 (2.1) |
| WHO Ordinal Scale score at enrolment | | |
| Score 3 | 12 (21) | 10 (37) |
| Score 4 | 41 (73) | 16 (59) |
| Score 5 | 3 (5) | 2 (4) |

ACE, angiotension-converting enzyme; ARB, angiotension receptor blocker; BMI, body mass index; COPD, chronic obstructive pulmonary disease; SD, standard deviation.

3.2 Endpoints

Disease severity on day 7, the primary endpoint, was not different in the two groups ($p = 0.11$) (Table 2), which was also the case after stratification by the baseline disease severity score (Table 3), and when the change of the WHO Ordinal Scale from baseline to day 7 was analyzed ($p = 0.96$, data not shown). Consistent with the observations at baseline, severe disease presentations were more frequently observed at day 7 in the intervention than in the control arm (19.7% vs. 7.4%, $p = 0.21$). The median time to clinical improvement was 7 days in both arms ($p = 0.57$) (Figure 1). Patients in the intervention arm more frequently required mechanical ventilation within 14 days ($n = 8$, 14.3%) and developed more often acute lung injury ($n = 3$, 5.4%) than the ones in the control arm ($n = 1$, 3.7% and $n = 0$, respectively, $p = 0.26$ for the comparison of mechanical ventilation). The median time to defervescence was 3.0 days in both arms ($p = 0.22$). The analysis of disease severity on day 14 mirrored the results on day 7 with no differences in the two groups ($p = 0.40$). Admission to the ICU until day 14 was required in 10 (17.9%) and 2 (7.4%) patients in the intervention and control arm, respectively ($p = 0.32$). The proportions of patients discharged at day 14 were 38 (67.9%) and 23 (85.2%) in the intervention and control arm, respectively. The length of hospital stay in survivors was longer in the intervention arm (mean 11.4 days, SD 8.0) in comparison to the one in the control arm (mean 7.6 days, SD 5.2, $p = 0.04$). During the 28-day follow-up, death was only observed in the intervention arm ($n = 6$; three, two, and one case in Switzerland, Brazil, and Mexico, respectively). Two additional deaths in the intervention arm and one in the control arm were noted during the subsequent follow-up period of SAEs.

3.3 Laboratory analyses

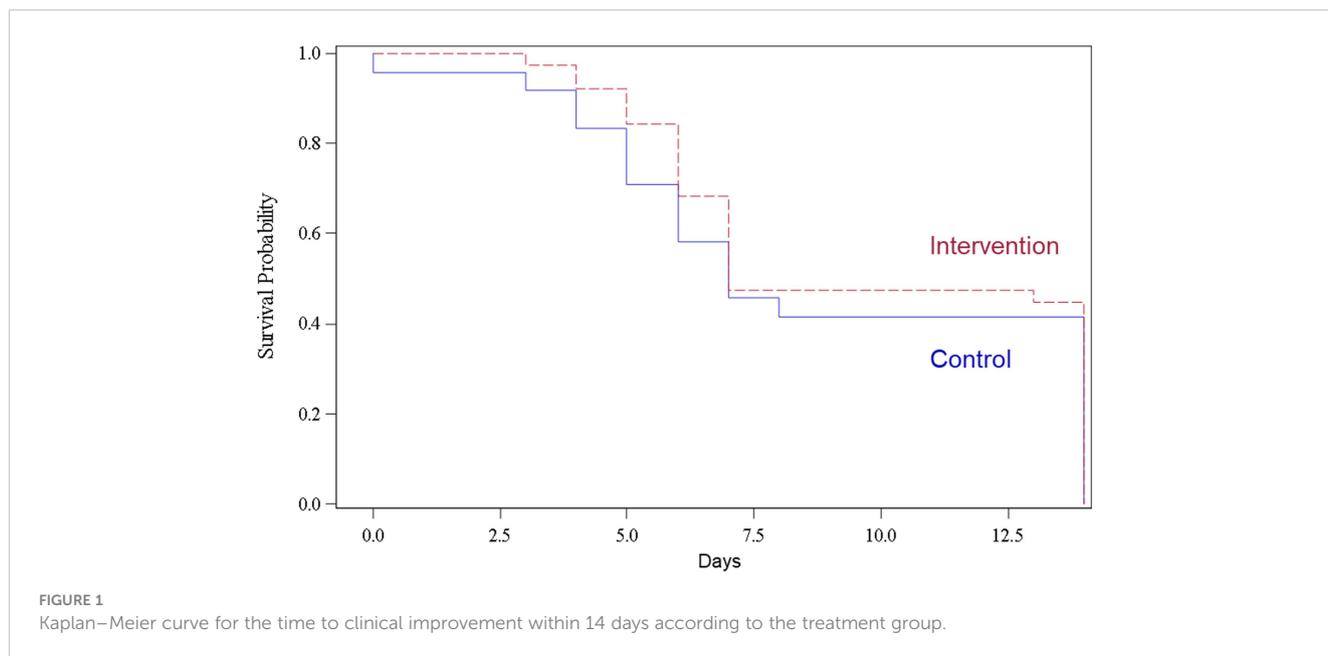
Analysis of C1INH concentrations revealed elevated values in both groups at baseline (Figure 2A). They nearly doubled after the first ConA administration (8,400 U) in the intervention arm, but

TABLE 2 Disease severity at day 7.

| | WHO Ordinal Scale (score points) | Intervention arm (n = 56) | Control arm (n = 27) |
|--|----------------------------------|---------------------------|----------------------|
| Outpatient, n (%) | 1 | 8 (14.3) | 7 (25.9) |
| | 2 | 12 (21.4) | 5 (18.5) |
| Hospitalized—mild disease, n (%) | 3 | 13 (23.2) | 5 (18.5) |
| | 4 | 12 (21.4) | 8 (29.6) |
| Hospitalized—severe disease, n (%) | 5 | 3 (5.4) | 1 (3.7) |
| | 6 | 6 (10.7) | 1 (3.7) |
| Death, n (%) | 7 | 2 (3.6) | – |
| <i>p</i> -value (Wilcoxon test stratified by baseline) | | 0.11 | |

TABLE 3 Disease severity on day 7 stratified by baseline scores.

| WHO Ordinal Scale | | Intervention arm (n = 56) | Control arm (n = 27) | |
|-------------------------|----------------------|---------------------------|----------------------|----------|
| Baseline (score points) | Day 7 (score points) | | | |
| 3 | n | 12 | 10 | |
| | 1 | n (%) | 3 (25.0) | 6 (60.0) |
| | 2 | | 6 (50.0) | 1 (10.0) |
| | 3 | | 2 (16.7) | 2 (20.0) |
| | 4 | | 4 (8.3) | 1 (10.0) |
| | p-value | | 0.38 | |
| ≥4 | n | 44 | 17 | |
| | 1 | n (%) | 5 (11.4) | 1 (5.9) |
| | 2 | | 6 (13.6) | 4 (23.5) |
| | 3 | | 11 (25.0) | 3 (17.7) |
| | 4 | | 11 (25.0) | 7 (41.2) |
| | 5 | | 3 (6.8) | 1 (5.9) |
| | 6 | | 6 (13.6) | 1 (5.9) |
| | 7 | | 2 (4.6) | - |
| | p-value | | 0.70 | |



returned close to baseline before the 4th administration (4,200 U). Subsequent dose administrations increased the C1INH concentrations each by approximately 50%. The mean time until SARS-CoV-2 clearance was 6.3 (SD 4.1) and 6.4 (SD 2.4) days in the intervention and control arm, respectively ($p = 0.74$). Similarly, the

course of CRP (Figure 2B), ferritin, D-dimer, creatinine (Supplementary Figure S2), and lymphocyte values (data not shown) was similar in both groups. There was no evidence of relevant complement C4 consumption (Supplementary Figure S3) and no difference in the activation of the CS (as measured by C4d

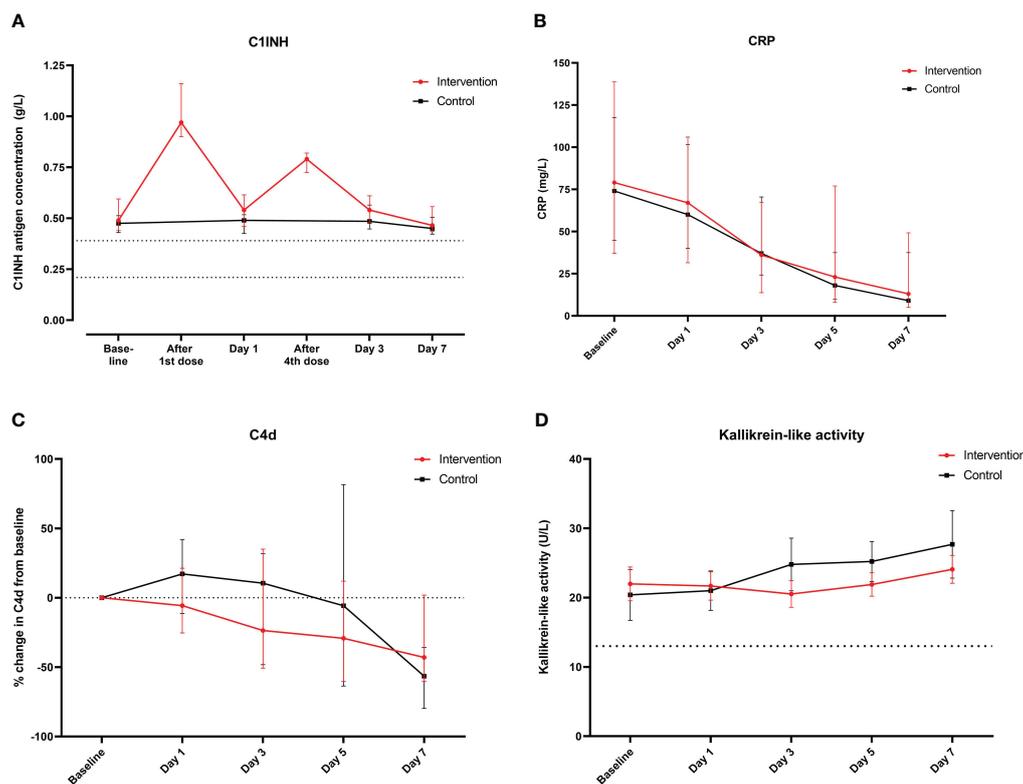


FIGURE 2
 Course of main laboratory parameters from baseline until day 7 according to the treatment group. Course of (A) C1INH concentrations (g/L, medians and interquartile ranges; dashed lines indicate the normal range in healthy adults), (B) CRP (mg/L, medians and interquartile ranges), and (C) relative change in C4d (% change from baseline, medians and interquartile ranges; dashed line indicates baseline) and (D) course of kallikrein-like activity (U/L, medians and interquartile ranges; dashed line indicates activity of control plasma).

and sC5b-9, respectively), endothelial cells (as measured by VCAM-1 and E-selectin), or the KSS (as measured by kallikrein-like activity) in both groups (Figure 2; Supplementary Figure S3).

[6 (10.7%) and 1 (3.7%)], and “cardiac disorders” [3 (5.4%) and 0]. Details are listed in Supplementary Table S5.

None of the AEs or SAEs were judged as being related to the study drug by the investigators.

3.4 Safety

3.4.1 Adverse events

Twenty-two subjects had at least one AE: 17 (30%) in the intervention arm and 5 (19%) in the control arm. Overall, 35 AEs were reported: 30 (85.7%) in the intervention arm and 5 (14.3%) in the control arm. Ten of these AEs were attributed to investigations [i.e., laboratory parameters: eight (80.0%) in the intervention arm and two (20.0%) in the control arm]. Details are listed in Supplementary Tables S3 and S4.

3.4.2 Serious adverse events

These occurred in 15 (27%) patients in the intervention arm and 4 (15%) in the control arm. The most common system organ classes of the SAEs were “respiratory, thoracic and mediastinal disorders” [10 (17.9%) and 2 (7.4%)], “infections and infestations”

3.5 Premature trial determination

While SAEs and deaths were judged as not related to the trial medication, their increased occurrence in the intervention arm led the DSMB to recommend an interruption of the trial after the second interim analysis. In addition, the DSMB recommended to await the outcome data of this analysis, including analyses of the activity of the plasma cascades prior to further decision-making on cessation or continuation of the trial. Subsequently, the trial was terminated prematurely in September 2021 for the following reasons. First, the intervention regimen was not associated with a significant inhibition of the CS and activation of other plasma cascades or endothelial cells. Second, considering the results in the primary outcome not favoring the intervention and the observed imbalances in the baseline characteristics, a larger sample size

would have been required to show a difference if there was a true difference. Third, during the second half of the year 2021, additional treatment options such as anti-SARS-CoV-2 monoclonal antibodies and tocilizumab were introduced and recommended, making the study groups even more heterogeneous.

4 Discussion

This randomized trial found that C1INH treatment added to SOC was not associated with clinical improvement or faster recovery from COVID-19 in comparison with SOC alone in hospitalized, non-critically ill patients. The overall mortality in this trial (7%) was substantially lower than the one previously reported in the dexamethasone trial (23%), published by the RECOVERY collaborative group (23). The current trial was terminated prematurely because of a lack of efficacy. The power of the study is low because the sample size needed for adjustment after the interim analyses could not be realized. Although there were more AEs and SAEs in the intervention arm than in the control arm, the trial was not terminated because of safety concerns.

The conduction of this trial was justified by scientific rationales and a previous pilot study (18). The former included the idea that C1INH may dampen uncontrolled complement activation and collateral lung damage, may reduce capillary leakage and subsequent pulmonary edema and the generation of microthrombi by inhibition of the KKS and the CAS, and may preserve the regulatory role of endothelial cells. Despite these arguments, we did not observe a clinical benefit in preventing COVID-19 progression when using ConA plus SOC. The following reasons may explain the study outcomes observed.

First, the administration of ConA may have been too late in the course of COVID-19. Because of the short half-life of ConA, repeated administration of ConA over 3 days was chosen. The dosage chosen was supposed to increase the C1INH concentrations by at least 50%. Pharmacokinetic analyses revealed that C1INH concentrations were already markedly elevated at baseline in both arms, which is in agreement with our previous data (24). It is also a potential hint that activation of plasma cascades is already significant and potentially beyond the “point of no return” for inhibition by C1INH, when patients with COVID-19 seek for medical help because of their symptoms. C1INH targets plasma cascades at an early activation level.

Second, the dosing regimen may have been unable to inhibit the CS in COVID-19 sufficiently and for a prolonged period of time. C1INH concentrations doubled initially after the first administration of ConA but returned to almost baseline before the 4th administration. The exact concentration required to inhibit the complement and other cascades and the optimal target are unknown both in the specific context of COVID-19 and in hyperinflammation secondary to infection in general. Of note, COVID-19 is characterized by a local activation of plasma cascades that progresses to a systemic activation early in the disease (25). As such, our data demonstrate that even a twofold increase in C1INH concentration is not sufficient to significantly

inhibit all potential downstream effects of the CS, KKS, CAS, and endothelial cells (26, 27).

Third, in COVID-19, the potential inhibitory role of C1INH may have been overestimated when considering activation through the classical and alternative pathway (11, 24, 28, 29). The lectin pathway of complement has been initially implicated as a major driver of complement activation, and binding of MBL or MASP-2 to SARS-CoV-2 was demonstrated with subsequent complement activation (30–32). Subsequently, it became clear that the S protein of SARS-CoV-2 directly activates the alternative pathway (33), which is not efficiently inhibited by C1INH, and circulating immune complexes may activate the classical pathway (28), in line with results from autopsy studies (34).

Fourth, cross-activation of the CS downstream of the lectin and classical pathway proteases may play a considerable role in COVID-19 disease progression. For example, factor XIa inhibited the regulatory complement factor H of the alternative pathway enhancing alternative pathway activity (35). Activated platelets may trigger alternative pathway activation (36–38), and the proteases of the coagulation and fibrinolysis cascades (both activated during COVID-19 but not or only poorly influenced by C1INH) are capable of cross-activating the CS at the level of C3 or C5 (39, 40). In line, the course of serum markers of inflammation and activation of the coagulation system was similar in both arms. Last, it is unclear if different SARS-CoV-2 variants circulating during the study period may have differential effects on the activation of plasma cascade and hence the inhibitory effect of C1INH.

The CS has been a target of a number of randomized controlled trials in COVID-19. However, only retrospective data from patients requiring continuous positive airway pressure support (41) and a single phase 3 trial reported a survival benefit when administering an anti-C5a antibody, albeit in mechanically ventilated COVID-19 patients for the latter (42), whereas another large phase 3 trial in the same population using a C5 inhibitor was futile (43). Clinical trials targeting the CAS and KKS are scarce, and consist of small sample sizes (44, 45). In the only published randomized controlled trial of C1INH in COVID-19, C1INH did not influence clinical improvement and length of stay compared to the SOC arm, in line with our study results (46). The present trial, despite reporting negative findings, is important to direct future clinical interventions targeting the CS, KKS, and CAS in other respiratory viral diseases.

This trial was designed early during the pandemic. As such, it was decided that instead of defining futility margins for a rather small-scale trial including a diverse patient population, a regular review of the DSMB might be more appropriate, as the included patients would potentially receive various new treatments with poorly known safety profiles as time evolved. Instead, two adaptive interim analyses after 40 and 80 patients were carried out. Although more SAEs occurred in the intervention arm, the majority of them were noted after the intervention period. In agreement with external evaluations, we attributed most SAEs to COVID-19 or a complication of it (e.g., pneumonia or sepsis). This interpretation is also supported by the observation of more severe COVID-19 at baseline and throughout the study period in the intervention arm. Of note and in contrast to C5 inhibitory

strategies, CIINH treatment has not been associated with an increased risk for bacterial or viral infections or any effect on viral replication (47). Despite being implemented in the electronic case report form and using variable block sizes, which impairs correct forecasting of the allocation, randomization was not associated with a balanced distribution of severity. Reasons include the 2:1 randomization ratio associated with a rather small number of individuals included in the control arm. Moreover, stratification by center may have contributed to the imbalances observed given that some centers only enrolled a limited number of patients. Indeed, we observed a much larger imbalance after the first interim analysis (40 patients included), which was reduced after including more patients. The DSMB reviewed all safety data and deaths in detail during three meetings. For several reasons including the imbalances in disease severity at baseline favoring the control group and the overall lower-than-expected mortality in the intervention arm in the Swiss centers with the majority of enrolled patients, termination of the study for safety was not recommended.

Our study has limitations. These include the premature closure of the trial, the lack of prespecified futility margins, the small number of patients in the control arm (due to the 2:1 randomization), and the imbalances in baseline characteristics and disease severity observed between the groups at baseline (potentially due to the 2:1 randomization and stratification in the setting of a limited number of enrolled patients). Almost 50% of patients were recruited in one center, reflecting an uneven distribution between the institutions. It was an open-label trial, and as such, it may have been prone to bias. The SOC treatment allowed the use of remdesivir and dexamethasone, which may have diminished the possible effect of ConA. Last but not least, follow-up phone calls might potentially have been less accurate than in-person follow-up visits when assessing for AEs. Despite these limitations, this is the largest study to investigate a treatment strategy that interferes with the CS, KKS, and CAS in hospitalized COVID-19 patients. In addition, we provide patient-matched laboratory data regarding the inability of ConA to limit the activation of these plasma cascades in COVID-19.

In conclusion, ConA did not significantly accelerate clinical improvement or reduce mortality in hospitalized patients with non-critically ill COVID-19 infection. The chosen dose and regimen of ConA used in this study are insufficient to limit complement activation in COVID-19. It remains unclear whether or not higher dosages, a longer treatment regimen, or an earlier intervention time point with ConA in COVID-19 could have prevented disease progression.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, on reasonable request.

Ethics statement

The studies involving humans were approved by the ethics committees «Ethikkommission Nordwest- und Zentralschweiz», «Ethikkommission Ostschweiz», and “Kantonale Ethikkommission Zürich» in Switzerland, “Comissão Nacional de Ética Em Pesquisa” (CONEP) in Brazil and “Comisión Federal Para La Protección Contra Riesgos Sanitarios” (COFEPRIS) in Mexico. The studies were conducted in accordance with the local legislation and institutional requirements. The participants provided their written informed consent to participate in this study.

Author contributions

PU: Conceptualization, Data curation, Formal Analysis, Investigation, Methodology, Project administration, Supervision, Visualization, Writing – original draft, Writing – review & editing. ML: Investigation, Project administration, Writing – review & editing. PC: Investigation, Project administration, Writing – review & editing. SM: Investigation, Project administration, Writing – review & editing. IH: Conceptualization, Investigation, Methodology, Writing – review & editing. MT: Investigation, Project administration, Supervision, Writing – review & editing. RT: Investigation, Project administration, Writing – review & editing. JS: Investigation, Project administration, Writing – review & editing. AC: Investigation, Project administration, Supervision, Writing – review & editing. MB: Investigation, Project administration, Supervision, Writing – review & editing. LH: Investigation, Project administration, Supervision, Writing – review & editing. MS: Investigation, Project administration, Writing – review & editing. WA: Investigation, Project administration, Supervision, Writing – review & editing. PS: Conceptualization, Formal Analysis, Investigation, Methodology, Project administration, Supervision, Validation, Writing – original draft, Writing – review & editing. MO: Conceptualization, Data curation, Formal Analysis, Funding acquisition, Investigation, Methodology, Project administration, Resources, Supervision, Validation, Visualization, Writing – original draft, Writing – review & editing.

Funding

This work was supported by the Swiss National Science Foundation (SNSF) within the framework of the National Research Program “Covid-19” (NRP 78) Grant-N°40780_198403, an unconditional research grant including free investigational medicinal product of Pharming Biotechnologies B.V., Leiden, The Netherlands, and departmental funds of MO (University Hospital Basel).

Acknowledgments

We thank the trial participants, caregivers, investigators, and research staff who contributed to the trial. We are particularly thankful to the nurses and physicians who cared for the COVID-19 patients during stressful times of the pandemic. We thank Pharming Biotechnologies B.V., Leiden, The Netherlands, for providing ConA (Ruconest®) free of charge.

Conflict of interest

MT reports receiving grants from the Swiss National Science Foundation, and having research collaborations with Roche, Novartis, and Idorsia outside the submitted work. WA reports receiving fees and research grants from A. Vogel AG, Gilead, and OM Pharma and fees for attendance of advisory boards to Pfizer, MSD Vifor Pharma, GSK, Sanofi, OM Pharma, and Janssen that were paid to his institution outside the submitted work. MO reports receiving grants from the Swiss National Science Foundation, consulting fees from Pharming Biotechnologies B.V. during the conduct of the study and grants from Pharming Biotechnologies B.V. outside the submitted work. LH reports receiving consulting fees from GlaxoSmithKline and Novartis during the conduct of the study but unrelated to this trial.

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Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fimmu.2023.1255292/full#supplementary-material>

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