# Interferon $\beta$ -1a ring prophylaxis to reduce household transmission of SARS-CoV-2: a cluster randomised clinical trial

José A. Castro-Rodriguez,<sup>a</sup> Eleanor N. Fish,<sup>b</sup> Samuel T. Montgomery,<sup>c,d</sup> Tobias R. Kollmann,<sup>c</sup> Carolina Iturriaga,<sup>a</sup> Casey Shannon,<sup>e</sup> Yuliya Karpievitch,<sup>c</sup> Joseph Ho,<sup>c</sup> Virginia Chen,<sup>e</sup> Robert Balshaw,<sup>f</sup> Rym Ben-Othman,<sup>c</sup> Radhouane Aniba,<sup>c</sup> Francisca Gidi-Yunge,<sup>a</sup> Lucy Hartnell,<sup>c</sup> David G. Hancock,<sup>c</sup> Guillermo Pérez-Mateluna,<sup>a</sup> Marcela Urzúa,<sup>a</sup> Scott J. Tebbutt,<sup>e</sup> Diego García-Huidobro,<sup>a,g</sup> Cecilia Perret,<sup>a</sup> Arturo Borzutzky,<sup>a</sup> and Stephen M. Stick<sup>c,\*</sup>

<sup>a</sup>Escuela de Medicina, Pontificia Universidad Católica de Chile, Santiago, Chile <sup>b</sup>Toronto General Hospital Research Institute, University Health Network & Department of Immunology, University of Toronto, Toronto, Ontario, Canada <sup>c</sup>Telethon Kids Institute, Perth, Australia <sup>d</sup>School of Population Health, Curtin University, Bentley, Australia

<sup>e</sup>Prevention of Organ Failure (PROOF) Centre of Excellence, Vancouver, British Columbia, Canada

<sup>f</sup>Centre for Healthcare Innovation, University of Manitoba, Manitoba, Canada

<sup>g</sup>Department of Family and Community Medicine, University of Minnesota, MN, USA

## Summary

**Background** Accumulating evidence indicates that an early, robust type 1 interferon (IFN) response to SARS-CoV-2 is important in determining COVID-19 outcomes, with an inadequate IFN response associated with disease severity. Our objective was to examine the prophylactic potential of IFN administration to limit viral transmission.

Methods A cluster randomised open label clinical trial was undertaken to determine the effects of pegylated IFN $\beta$ -1a administration on SARS-CoV-2 household transmission between December 3rd, 2020 and June 29th, 2021. Index cases were identified from databases of confirmed SARS-CoV-2 individuals in Santiago, Chile. Households were cluster randomised (stratified by household size and age of index cases) to receive 3 doses of 125  $\mu$ g subcutaneous pegylated IFN $\beta$ -1a (172 households, 607 participants), or standard care (169 households, 565 participants). The statistical team was blinded to treatment assignment until the analysis plan was finalised. Analyses were undertaken to determine effects of treatment on viral shedding and viral transmission. Safety analyses included incidence and severity of adverse events in all treatment eligible participants in the standard care arm, or in the treatment arm with at least one dose administered. Clinicaltrials.gov identifier: NCT04552379.

**Findings** 5154 index cases were assessed for eligibility, 1372 index cases invited to participate, and 341 index cases and their household contacts (n = 831) enrolled. 1172 participants in 341 households underwent randomisation, with 607 assigned to receive IFN $\beta$ -1a and 565 to standard care. Based on intention to treat (ITT) and per protocol (PP) analyses for the primary endpoints, IFN $\beta$ -1a treatment did not affect duration of viral shedding in index cases (absolute risk reduction = -0.2%, 95% CI = -8.46% to 8.06%) and transmission of SARS-CoV-2 to household contacts (absolute risk reduction = 3.87%, 95% CI = -3.6% to 11.3%). Treatment with IFN $\beta$ -1a resulted in significantly more treatment-related adverse events, but no increase in overall adverse events or serious adverse events.

Interpretation Based upon the primary analyses, IFNβ-1a treatment did not affect duration of viral shedding or the probability of SARS-CoV-2 transmission to uninfected contacts within a household.

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\*Corresponding author.





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E-mail address: stephen.stick@health.wa.gov.au (S.M. Stick).

#### **Research in context**

#### Evidence before this study

Type I IFNs are critical early mediators of the innate immune response to all virus infections. They achieve viral clearance by directly inhibiting multiple stages of the viral replicative cycle, by protecting uninfected cells from infection and by recruiting and activating immune cell populations to sites of infection. Different clinical studies have identified that early treatment with IFNs- $\alpha/\beta/\lambda$  may accelerate SARS-CoV-2 viral clearance and prevent progression to severe disease. Despite effective vaccines that reduce disease severity, and the emergence of Omicron variants that apparently result in less severe disease, these highly transmissible variants continue to fuel the global pandemic. Prophylactic IFN treatment of 'at risk' individuals in outbreak or high risk settings may offer a solution to blunt transmission and end the pandemic.

#### Added value of this study

This randomized cluster trial of 1172 participants in 341 households demonstrated that 3 subcutaneous injections of

Introduction

The SARS-CoV-2 pandemic has claimed over six million lives. Despite the rapid development and deployment of vaccines in many countries, the number of new cases worldwide is approximately 500,000 daily (https:// covid19.who.int). With each wave of the pandemic, health systems have been challenged, complicated with emergent mutant strains of the virus. Mutated strains may be more transmissible,<sup>1,2</sup> cause more severe disease than the original pandemic strain of SARS-CoV-2<sup>3</sup> and have the potential to evade available vaccines.4,5 Whilst widespread vaccination has had success limiting the trajectory of the pandemic, the emergence of the Omicron variants demonstrates that even with mutations that appear to cause less severe disease,6 high transmission despite immunization nonetheless results in significant pressures on health services.7 The solution to halting any pandemic is ending community transmission. During the current pandemic, measures such as healthy hygiene, self-isolation when sick, physical distancing and use of face masks have all been effective.8 Moreover, expedited public health responses such as extensive contact tracing, testing for infection and community lockdowns have all been effective in limiting transmission.9 International and local border closures plus strict quarantine measures have reduced community transmission to zero for periods in countries such as Australia and New Zealand.<sup>10,11</sup> However, in these countries and elsewhere, as restrictions are relaxed, localized outbreaks have occurred<sup>12</sup> that have required rapid, community-wide responses to again supress transmission. Importantly, these community constraints cause unprecedented civil disruption and come at enormous economic<sup>13,14</sup> and social costs.<sup>15,16</sup>

125 µg each of pegylated IFNβ-1a had no effect on SARS-CoV-2 viral shedding or transmission from an infected index cases in a household to uninfected household contacts. However, post-hoc sensitivity and exploratory analyses identified a reduction in transmission to household contacts where the index case had a high (>10<sup>6</sup> copies/mL) viral load. Pegylated IFNβ-1a was safe and well tolerated, with no concerning adverse laboratory events.

#### Implications of all the available evidence

There is no approved therapy to prevent SARS-CoV-2 transmission. This unique human challenge study demonstrates that prophylactic treatment with IFN $\beta$ -1a reduced transmission of SARS-CoV-2 within households. Whilst IFN $\beta$ -1a prophylaxis cannot be recommended as a useful intervention given the available evidence, our observations should be considered when designing future clinical trials aimed at preventing the transmission of highly contagious viruses.

Since the evolution of dominant SARS-CoV-2 virus cannot be easily predicted,<sup>17</sup> there remains a need to identify interventions that can be rapidly deployed should highly pathogenic strains emerge despite high levels of community immunization. Furthermore, preparations for the next pandemic must include strategies to limit the potential for infection and transmission on first contact with pathogenic respiratory viruses.

One of the many therapeutic approaches investigated that appeared clinically useful early in the course of the pandemic was treatment with interferons (IFNs), namely IFNs- $\alpha/\beta$ . Randomised, controlled studies suggest that IFNs- $\alpha/\beta$  offer clinical benefits in moderate<sup>18</sup> and severe disease,<sup>19,20</sup> prevent infection in front-line hospital workers,<sup>21</sup> and recent data indicate that IFN- $\lambda$  reduces hospitalization and duration of viral shedding.<sup>22–24</sup> Nonetheless, IFNs are not generally recommended for treatment of proven cases of COVID-19<sup>25</sup> and other clinical trials have failed to demonstrate efficacy.<sup>26,27</sup>

IFNs are sentinel innate immune signalling molecules produced early after first contact with viral pathogens, that mediate their antiviral effects by direct inhibition of viral replication, protection of uninfected cells and also recruitment and activation of immune cells involved in viral clearance.<sup>26-30</sup> Accordingly, we postulated that prophylactic IFN administration might reduce susceptibility to infection of uninfected contacts of individuals with SARS-CoV-2 infection. Such ring prophylaxis could provide non-specific, antiviral protection to curb episodic viral outbreaks,<sup>31</sup> help suppress community transmission, even in vaccinated populations, and therefore reduce the risk of emergence of dangerous mutations.<sup>4,32</sup> We therefore undertook a cluster, randomised, controlled study of sub-cutaneous pegylated IFN $\beta$ -1a (Plegridy. Biogen Inc, Cambridge MA) administration, to determine whether IFN $\beta$ -1a given to index cases and household contacts might reduce transmission of SARS-CoV-2.

## Methods

The Containing Coronavirus Disease-19 (ConCorD-19) trial was a cluster randomised open label clinical trial of subcutaneous administration of pegylated IFN $\beta$ -1a (IFN $\beta$ -1a) versus standard care (control),<sup>33</sup> completed between December 3rd, 2020 and June 29th, 2021. Each household of an index case (IC) was randomly assigned to either the IFN $\beta$ -1a or control arm. The study was approved by the Institutional Review Board of the Pontificia Universidad Católica de Chile and was registered with clinicaltrials.gov (NCT04552379). The published trial protocol is available online.<sup>33</sup> All participants provided written informed consent.

## Trial population

ICs were identified from databases of those with confirmed SARS-CoV-2 from COVID-19 clinics and emergency room visits in Santiago, Chile. Households were contacted by telephone to determine eligibility prior to enrolment (Supplementary Table S5). Household contacts aged between 18 and 80 years who met inclusion and exclusion criteria were deemed as 'eligible' household contacts, with households only enrolled if there was at least one eligible contact.33 Inclusion and exclusion criteria for each participant type (index case, eligible household contact, ineligible household contact) are listed in the full study protocol in the Supplementary Appendices. Household characteristics were captured consistent with recommendations of the World Health Organization for assessing household transmission.<sup>34</sup> All participants implemented quarantine measures as mandated by local authorities and maintained a daily symptom diary which was collected and reviewed at each study visit by the study team (see Supplementary Appendices). Index cases were instructed to remain in isolation/quarantine for 11 days from onset of symptoms or, if asymptomatic, 11 days from the sample collection date that resulted in the COVID-19 diagnosis. Household contacts remained in isolation/quarantine for 11 days from date of the sample resulting in the diagnosis of the IC or of a newly diagnosed household member as per recommendations/ rulings by local authorities.

## Intervention

A mobile health team conducted home visits of all participant households on study days 1 (enrolment), 6, 11, 16, 21 and 29. ICs and eligible HCs in the IFN $\beta$ -1a arm received three subcutaneous doses of IFN $\beta$ -1a (125)

 $\mu$ g/0.5 ml × 0.5 ml) on study days 1, 6 and 11. Ineligible contacts in the IFNβ-1a arm, as well as ICs and all HCs in the control arm, received standard care. All participating households received information regarding hygiene, isolation, social distancing and wearing of face masks as per public health advice at the time of enrolment. The IFNβ-1a injection was given by a trained member of a mobile health team and participants were recommended to take paracetamol (1000 mg, 6 hourly) commencing at the same time as the IFNβ-1a for up to 24 h, in order to mitigate predictable flu-like symptoms.<sup>35</sup>

## Outcomes

The primary outcomes were (i) the proportion of participants in the IC-INF population shedding SARS-CoV-2 at study day 11 in the IFNβ-1a compared to control arm and (ii) the proportion of treatment eligible household contacts shedding SARS-CoV-2 at study day 11 in the IFNβ-1a compared to control arm. Secondary and exploratory outcomes in the trial protocol are listed in Supplementary Methods (Supplementary Table S1). Briefly, secondary outcomes included examination of the effects of IFNB-1a treatment on SARS-CoV-2 IgG serological conversion, the incidence of adverse events and hospitalizations. Shedding was determined by the presence of SARS-CoV-2 by PCR in saliva collected on days 1, 6, 11, 16, 21, and 29 (see Supplementary Methods). A SARS-CoV-2 PCR Ct value ≥37 was considered negative. Viral load (copies/mL) was estimated from the Ct value using a standard curve of known viral titre (Supplementary Figure S1). Anti-SARS-CoV-2 IgG antibodies to nucleocapsid protein were measured on day 29 (using the Liaison assay according to the manufacturer's instructions [DiaSorin, Saluggia, Italy]).

## **Biospecimen collections**

The full schedule of biospecimen collection is provided in the Supplementary Methods (Supplementary Tables S2 and S3). All consenting, non-eligible HCs also provided biospecimens according to the schedule collection for non-eligible HCs.

## Adverse events

These were classified in accordance with Good Clinical Practice,<sup>36</sup> as non-serious or serious and as related or unrelated to the trial medication. Because of overlap between symptoms of COVID-19 and potential IFN $\beta$ -1a-related adverse events, all symptoms were recorded and categorized, and any symptoms outside of the directed symptom assessment were considered adverse events. An independent data and safety monitoring committee reviewed safety data.

#### **Power calculations**

The study was designed at the start of the pandemic and there were few data to guide sample size calculations. Sample size calculations assumed a twosided alpha level of 0.025 and power >90% to ensure, assuming an extremely conservative correlation between outcome measures of zero, the familywise type 1 error rate remains below 0.05 and power remains above 80%. To estimate the required sample size, we used available transmission data reported at the start of the pandemic, census data to estimate household size in Santiago and an effect size based on a pilot study undertaken in Wuhan.37 Data from Wuhan suggested that the proportion of untreated index cases still shedding virus on study day 11 would be ~85%. Based on a two tailed Fisher's exact test, a sample size of 278 ICs (310 allowing for a 10% drop-out rate) would have >90% power, at  $\alpha$  = 0.025, to detect a difference in the proportion of index cases shedding SARS-CoV-2 at study day 11, if the proportion in the IFN $\beta$ -1a arm was 65%. The estimated average household size was 4, based on available census data,<sup>38</sup> and we estimated the secondary infection rate (transmission within the household) where there is an untreated IC would be 28%.<sup>39,40</sup> A sample of 278 households, providing 834 household contacts, would have >90% power, at alpha 0.025, to detect an odds ratio of 0.5 for a reduction in transmission to a household contact, based on a stratified Cochran-Mantel-Haenszel test (two-tailed with intra-class correlation of 0.15). Based on the uncertainty regarding the effect size in the trial setting, we also planned a Bayesian analysis to estimate the probabilities of transmission for even small effects of therapy.<sup>33</sup>

## Randomisation

Households were randomised as individual clusters using a minimization technique (biased coin, p = 0.7) in order to achieve balance in the total number of people within the household between treatment arms.<sup>41</sup> Households were randomized as clusters to receive either IFN  $\beta$ -1a treatment or standard of care at a 1:1 ratio. Households were randomized during the first home visit using minimization software,<sup>42</sup> once eligibility was confirmed, participants signed informed consent, and baseline data collection procedures were completed. Participants and study staff were not blinded to randomization of treatment or standard of care, but the statistical team was blinded to treatment assignment until the analysis plan was finalised.

#### Statistical analyses

We undertook frequentist analyses based upon the primary and secondary outcomes as stated in the study protocol and described in detail in the statistical analysis plan (available in the Supplementary Appendices) which included intent-to-treat and per protocol approaches. Participants were excluded from the populations used in the primary analyses if the study visit was not performed or was performed outside the 1-day window either side of the scheduled visit date as outlined in the Statistical Analysis Plan. Participants who didn't complete the full course of the treatment were excluded from the per protocol populations used in sensitivity analyses. Briefly, the analysis for primary outcome 1 used a generalized linear model (binomial, with a logit link for dichotomous outcomes) and was adjusted for age and sex, reporting an adjusted odds ratio (OR) and its corresponding 95% confidence interval. The analysis for primary outcome 2 used a generalized linear mixed model (binomial with a logit link) adjusted for age and sex and used a random intercept per household with an assumed normal distribution, reporting an adjusted OR and its corresponding 95% confidence interval. Sensitivity analyses were included with additional covariate adjustments and subgroup analyses.

While the frequentist approach allowed us to estimate the effects of IFN $\beta$ -1a administration on the risk of an individual becoming infected, the planned Bayesian analysis allowed us to determine whether the ring prophylaxis strategy using IFN $\beta$ -1a reduces the probability of transmission within the household of an infected index case. The analysis utilised the household contact population as defined in the Statistical Analysis Plan. Detailed methods are provided in the Supplementary Methods. Briefly, a generalized linear mixed effects model with a binomial logit function was developed to estimate the probability of infection that is influenced by explanatory variables for each contact case using the rstanarm R package.<sup>43,44</sup>

The analyses of secondary outcomes included duration of SARS-CoV-2 shedding assessed with discrete time-to-survival analysis using a generalized linear model with a complementary log–log link, proportion of participants positive for SARS-CoV-2 at study days 1 and 11 and seroconversion at study day 29 were assessed using Cochran-Mantel-Haenszel tests stratified by baseline immunity, incidence of hospitalization, death, or hospitalization and/or death were assessed using Cochran-Mantel-Haenszel tests stratified by household size, and duration of hospitalization was assessed using Fine-Gray competing risks regression with in-hospital death as a competing risk. Safety analyses comparing incidence of adverse events were performed using Fisher's exact test.

We also undertook post hoc exploratory frequentist analyses based on a modified subgroup analysis that accounted for households where at time of enrolment, it was unknown that all household contacts were SARS-CoV-2 positive or where index cases were no longer shedding virus, and for effects that may only be associated with active treatment. The analysis plan and statistical report for the exploratory analyses are available in the Supplementary Appendices. Briefly, both frequentist analyses fitted generalized linear mixed effects models using lme4<sup>45</sup> using a random intercept per household with an assumed normal distribution.

### Role of funder

The funding bodies for this study had no role in data collection, analysis, or interpretation of data, writing of the manuscript, or decision to submit.

## Results

Recruitment, participation, and completion data are shown in Fig. 1. Household characteristics and participant demographics are shown in Tables 1 and 2 respectively. Between December 2020 and May 2021, 5154 index cases were assessed for eligibility, 1372 index cases invited to participate. 341 households were enrolled and randomised, of which 137 (IFNβ-1a arm) and 151 (control arm) completed the study. Of the 1172 individuals randomised (IFNβ-1a arm = 607; standard of care arm = 565), 53 individuals withdrew from the study, of which 15 were ICs: 35 (14 ICs) in the IFNβ-1a arm and 18 (1 IC) in the control arm. The reasons for withdrawal are summarized in the Supplementary Results (Supplementary Table S4). One IC withdrew before randomisation. Eighty-two households where the IC had a negative salivary PCR on Days 1 and 6, or where there were no eligible contacts who tested negative at recruitment, were excluded for the exploratory analyses. (36 households in the treatment arm, 46 households in the standard care arm): 259 households were considered as the 'at risk' population.

When the study population was assessed for descriptive differences, the baseline (study day 1) viral load of ICs was significantly higher in the IFN treatment arm compared with ICs in the standard care arm (p = 0.034) (Fig. 2A). We then assessed whether IFN $\beta$ -1a treatment reduced median viral load when compared to the standard care arm at each study day (1, 6, 11, 16, 21, & 29) including all household contacts (treatment eligible and ineligible). Treatment with IFN $\beta$ -1a significantly reduced median viral load at study day 6 in household contacts (p = 0.034; Fig. 2B) but had no effect on median viral load at study days 1, 11, 16, 21, or 29.

All point estimates and 95% confidence intervals for primary and secondary analyses are available in Table 3. The Statistical Analysis Report with all analyses undertaken as outlined in the Statistical Analysis Plan are included in the Supplementary Appendices. There was no evidence of an effect of IFNβ-1a administration (vs. standard of care) on the probability of viral shedding at study day 11 in the IC-INF population (ICs and treatment eligible HCs infected at study day 1) (Primary outcome 1; OR = 0.979, 95% CI = 0.647–1.479; absolute risk reduction = -0.2%, 95% CI = -8.46% to 8.06%) or in the subsequent sensitivity analyses (Figs. 3 and 4). Similarly, there was no evidence of an effect of IFN<sub>β</sub>-1a on the probability of viral shedding in (i.e. transmission to) treatment eligible HCs (EHC-ITT) associated with IFN $\beta$ -1a administration (vs. standard of care) at study day 11 (Primary outcome 2: OR = 0.582, 95% CI = 0.271–1.247; absolute risk reduction = 3.87%, 95% CI = -3.6% to 11.3%), however where the day 1 IC viral load was >10<sup>6</sup> copies/mL, there was evidence that treatment with IFNβ-1a significantly reduced transmission to treatment eligible HCs (OR = 0.121, 95% CI = 0.025–0.524) (Figs. 3 and 4). Additionally, in an unplanned sensitivity analysis with the outcome measure changed to the probability of viral shedding at study day 6, there was a significant reduction in the probability of viral shedding in treatment eligible HCs (EHC-ITT): OR = 0.493, 95% CI = 0.256–0.949.

Analysis of the secondary outcomes revealed no evidence of an effect of IFN $\beta$ -1a treatment (vs. standard of care) on duration of viral shedding in the IC-INF population using a discrete time survival analysis or in the subsequent sensitivity analyses. Similarly, there was no evidence of an effect of IFN $\beta$ -1a treatment (vs. standard of care) on incidence of saliva PCR positive for SARS-CoV-2 in HCs at day 11 or on seroconversion in HCs at day 29 stratified by seroconversion at baseline.

There were 58 serious adverse events in index cases, 57 hospitalizations due to COVID-19 (25 in the IFNβ-1a treatment arm and 32 in the control arm) and one death in the standard care arm due to COVID-19 in an individual with significant co-morbidities and uncompensated diabetes mellitus. Twenty-six household contacts were hospitalized, 10 in the IFNβ-1a treatment arm and 16 in the control arm. Analysis of safety outcomes (Secondary outcome 3 and 4) found no evidence of an effect of IFNB-1a treatment on incidence or duration of hospitalization due to COVID-19 in the IC-ITT population (Secondary outcome 3) or on incidence of hospitalization in the IC-INF population. However, in sensitivity analyses for secondary outcome 3, there was evidence of a statistically significant effect of IFNβ-1a treatment (vs. standard of care) on reducing the incidence of hospitalization in the IC-PP population (p = 0.0312), and evidence of a statistically significant effect of IFNβ-1a treatment (vs. standard of care) on increasing the duration of hospitalization in the IC-INF population (HR = 0.452, 95% CI 0.233-0.877). For analysis of safety outcomes of IFN<sub>β</sub>-1a treatment (Secondary outcome 4), there was a significant increase in treatment-related adverse events in the IFNβ-1a arm, but no effect on overall adverse events or incidence of serious adverse events.

The Bayesian analysis (see Supplementary Methods) identified a 95% probability of reduction of infection within a household by IFN $\beta$ -1a treatment and the credible interval for the reduction in transmission probability was in the order of 0.5% to 15.3% during the active treatment period (study days 1–11). During the active treatment period, there was a significant reduction in the odds of transmission (OR = 0.43, 95% credible interval = 0.21–0.86) with a posterior probability that IFN $\beta$ -1a treatment is superior to standard care of 97.5%. The estimated Bayes factor for period 1 was

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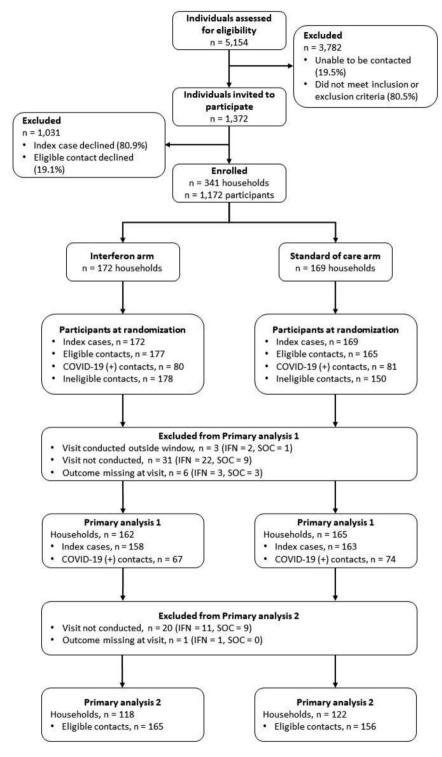


Fig. 1: CONSORT diagram describing participant screening, enrolment, randomisation, and analysis.

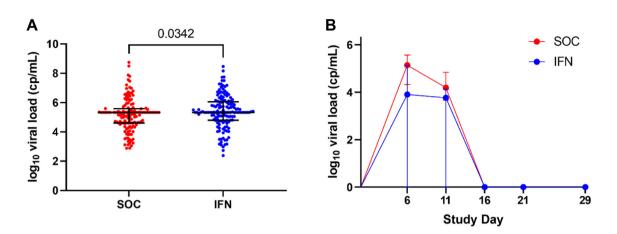
	Household demographics	
	SOC (n = 169)	IFN (n = 172)
Household occupants		
Mean (SD)	3.74 (1.29)	3.90 (1.64)
Infected household occupants at study start		
Median (IQR)	1 (1-2)	1 (1-2)
At risk household occupants at study start		
Median (IQR)	2 (1-3)	2 (1-3)
Household occupants fully vaccinated for COVID-19		
Median (IQR)	1 (0-2)	1 (0-2)
$Log_{10}$ viral load of household index case at study start		
Mean (SD)	5.25 (1.08)	5.42 (1.11)
SOC: Standard of care, IFN: IFNβ-1a treatment arm.		
Table 1: Household demographics at baseline.		

38.80, indicating strong support for the hypothesis that IFN $\beta$ -1a treatment reduces infection rate (~38 times more likely that observed data agrees with the hypothesis that IFN $\beta$ -1a treatment reduces infection than the hypothesis that treatment increases infection rate).<sup>46</sup> By contrast, in period 2 (day 12–29), the 95% credible interval for the reduction in transmission probability

	Full study population		
	SOC (n = 565)	IFN (n = 607)	
Age (years)			
Mean (SD)	35.0 (19.4)	33.2 (18.7)	
Sex			
Female, n (%)	294 (52.0%)	323 (53.2%)	
Household occupants			
Mean (SD)	3.74 (1.29)	3.90 (1.64)	
Fully vaccinated for SARS-CoV-2			
No, n (%)	380 (67.3%)	438 (72.2%)	
Cancer			
No, n (%)	559 (98.9%)	598 (98.5%)	
Diabetes			
No, n (%)	542 (95.9%)	583 (96.0%)	
Heart disease			
No, n (%)	533 (94.3%)	578 (95.2%)	
Asthma			
No, n (%)	524 (92.7%)	581 (95.7%)	
Chronic lung disease			
No, n (%)	564 (99.8%)	606 (99.8%)	
Chronic kidney disease			
No, n (%)	564 (99.8%)	604 (99.5%)	
Chronic neurological disease			
No, n (%)	562 (99.5%)	602 (99.2%)	
Smoker			
No, n (%)	502 (88.8%)	519 (85.5%)	
SOC: Standard of care, IFN: IFN $\beta$ -1a treatment arm.			
Table 2: Full study population demographics at baseline.			
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includes zero and therefore treatment was not effective (Supplementary Table S7). The estimated Bayes factor indicates that the data are only 2 times more likely to be observed under the hypothesis that IFN $\beta$ -1a treatment increases infection rate, but the evidence is weak and therefore considered non-significant.<sup>46</sup> The effect of IFN $\beta$ -1a on transmission was independent of household size (Supplementary Table S8). When the infection reduction is stratified by household size, only households of size 4 in period 1 had a 95% credible interval without zero.

Subsequent exploratory analyses focused on a population of household contacts deemed "at risk". The at risk population is specifically defined in the Supplementary Appendices, with the significant distinction being that households were excluded from analyses where the index case subsequently tested negative for SARS-CoV-2 on study days 1 and 6, and where no SARS-CoV-2 negative household contacts remain on study day 1, as transmission of SARS-CoV-2 would be biologically implausible. In the at-risk population, 55/293 (absolute risk: 18.7%, 95% CI = 13.6% to 25.9%) household contacts in the IFNβ-1a arm became infected during active treatment (days 1-11) of the study compared to 60/246 (absolute risk: 24.4%, 95% CI = 17.6% to 33.7%) household contacts in the control arm (Fig. 4), indicative of an relative risk reduction in the IFN arm equivalent to 23% (95% CI = -6.4% to 44.4%). When the effect of IFN $\beta$ -1a treatment on the probability of a SARS-CoV-2 positive saliva PCR in the at risk population was assessed post-hoc using a binomial generalized linear mixed model with a logit link function (Supplementary Appendices), treatment with IFNβ-1a was associated with a significant reduction in the odds of a SARS-CoV-2 positive saliva PCR for all household contacts compared to standard of care during the treatment period (study days 1-11) (p = 0.033; OR = 0.55, 95% CI = 0.36-0.99; Fig. 5). The treatment effect was not significant in the subsequent period (days



**Fig. 2: Descriptive differences in the household contact population. (A)** Baseline (study day 1) viral load of index cases in the standard of care (SOC) and IFN $\beta$ -1a treatment (IFN) arms in the household contact (HC) (SOC: n = 142. IFN: n = 147) population, with data present as median (circle) with interquartile range (whiskers). **(B)** Viral load of household contacts who test positive for SARS-CoV-2 in the SOC and IFN treatment arms in the (SOC: n = 66, IFN: n = 64) with data presented as median (circle) and interquartile range at each study visit. Zero values represent data below the limit of quantification, reported as zero per the statistical analysis plan. cp/mL = copies/mL.

12-29) i.e. after treatment and isolation had ended. Testing positive to SARS-CoV-2 was significant more common in the first period when compared with the second period (OR = 7.08, 95% CI = 3.76-9.29; Fig. 5). To assess whether the protective effect of treatment was more pronounced in households where the index case had a high viral load at the start of the study, we included a covariate dichotomizing index case viral load >10<sup>6</sup> copies/mL and compared generalized linear mixed effect models with and without an interaction term between this covariate and treatment arm. The treatment by viral load interaction term was significant (p = 0.005), indicating that the treatment effect is significantly different in households with IC with low (<10<sup>6</sup>) vs. high (>10<sup>6</sup>) viral loads. The odds of positive PCR result were significantly higher in households with index cases with high viral load (>10<sup>6</sup>) compared to those with low viral load (p = 0.02; OR = 3.29, 95% CI = 1.14-6.1). In households with index cases with high viral load (> $10^6$ ), treatment with IFNβ-1a significantly reduced the odds of a positive PCR result during the study period (p = 0.0028; OR = 0.17, 95% CI = 0.084-0.56; Fig. 5).Conversely, in households with index cases with low viral load (<10<sup>6</sup>), treatment with IFNβ-1a did not reduce the odds of a positive PCR result during the study period (p = 0.84; OR = 0.93, 95% CI = 0.54-1.68; Fig. 5).

## Discussion

Pegylated IFN $\beta$ -1a is an FDA approved therapy for multiple sclerosis for which the pharmacokinetics and safety profile are well-characterized.<sup>42</sup> The primary analyses from this prospective, cluster randomised, ring prophylaxis trial demonstrated that IFN $\beta$ -1a administration was ineffective as a public health measure to reduce the household transmission of SARS-CoV-2. The chosen formulation allowed us to predict the likely duration of activity of the three-dose regimen to cover the period of peak transmissibility of the virus. Using a mobile medical team to administer doses at home, optimized adherence to therapy and allowed reasons for non-adherence and withdrawal to be accurately documented.

Although the intention to treat and per protocol analyses failed to demonstrate a significant effect on primary outcomes, the sensitivity analyses for primary outcome 2 indicated an effect was observable using day 6 as an outcome measure, and that treatment was only effective in households where the initial viral load of index cases was high. Additionally, these analyses failed to account for index cases not shedding virus at randomization nor an absence of eligible contacts within a household. Using plausible filters to define an at-risk population provided additional insight regarding the biological effects of IFNB-1a administration on viral transmission. These exploratory analyses indicated that there was a high probability of a small reduction in household transmission. Since point-of-care diagnostics were unavailable at the time of the study, enrichment of the population with highrisk eligible household contacts was not possible. This is perhaps an important point to consider in the design of future ring prophylaxis studies, particularly if a small reduction in viral transmission at the start of a pandemic could translate into large heath and socioeconomic benefits.

To our knowledge, only one previous study assessed ring prophylaxis in COVID-19: Labhardt et al. observed that a combination of lopinavir/ritonavir for 5 days as post-exposure prophylaxis was not effective at

Outcome(s)	IFNβ-1a n (%)	Standard of care n (%)	Estimated treatment effect (95% ${\rm CI)}^{\rm a}$
Primary outcome 1 (n = 462)			
Proportion of IC-INF population shedding SARS-CoV-2 at study day 11	65 (28.9)	68 (28.7)	0.979 (0.647-1.479) ARR = -0.197% (-8.5 to 8.1%)
Sensitivity analyses			
Adjusted for vaccination status (n = 462)	-	-	0.963 (0.635–1.459)
Intent-to-treat index cases $(n = 321)$	-	-	1.106 (0.657–1.863)
Per-protocol index cases $(n = 309)$	-	-	1.062 (0.622–1.810)
Household index case viral load (> $10^6$ copies/mL) as a covariate (n = 309)	-	-	0.55 (0.274–1.082)
Primary outcome 2 (n = 321)			
Proportion of eligible household contacts shedding SARS-CoV-2 at study day 11	19 (11.5)	24 (15.4)	0.582 (0.271-1.247) ARR = 3.87% (-3.6 to 11.3%)
Sensitivity analyses			
Adjusted for risk factors (n = 321)	-	-	0.492 (0.202–1.199)
Household contact population ( $n = 518$ )	-	-	0.468 (0.178-1.231)
Per protocol eligible household contacts (n = 311)	-	-	0.589 (0.272-1.276)
Excluding eligible household contacts seropositive on study day 1 (n = 266)	-	-	0.640 (0.287-1.429)
Outcome at study day 6 (n = 329)	20 (11.8)	31 (19.5)	<b>0.493 (0.256–0.949)</b> ARR = 7.7% (–0.1 to 15.6%)
Index case viral load (<10 <sup>6</sup> copies/mL) (n = 321)	-	-	0.552 (0.306-1.12)
Index case viral load (> $10^6$ copies/mL) (n = 321)	-	-	0.113 (0.021-0.581)
Secondary outcome 1 (n = 502)			
Effect of treatment on duration of viral shedding	-	-	HR = 1.083 (0.964-1.217)
Secondary outcome 2			
Incidence of SARS-CoV-2 shedding in household contacts at study day 11 (n = 518)	37 (13.6)	43 (17.6)	0.680 (0.353-1.309)
Seroconversion in household contacts at day 29 (n = 462)	86 (34.5)	101 (47.4)	0.580 (0.336-1.002)
Secondary outcome 3			
Incidence of hospitalisation in IC-ITT (n = 299)	15 (9.38)	14 (10.1)	0.842 (0.385-1.842)
Sensitivity analyses			
IC-INF population (n = 437)	21 (9.29)	23 (10.9)	0.789 (0.418-1.489)
Per protocol index cases (n = 275)	4 (2.94)	14 (10.1)	0.272 (0.088–0.844)
Secondary outcome 3			
Duration of hospital stay due to COVID-19 (n = 29)	-	-	HR = 0.522 (0.237–1.152)
Sensitivity analyses			
IC-INF population ( $n = 43$ )	-	-	HR = 0.452 (0.233–0.877)
Per protocol index cases (n = 18)	-	-	HR = 2.006 (0.601–6.690)
Secondary outcome 4 (n = 844)			
Incidence of adverse events—rate of AE per participant (incidence of any AE)	2.117 (0.235)	1.824 (0.280)	p = 0.156 (Fisher's exact test)
Incidence of adverse events related to treatment events—rate of related AE per participant (incidence of any related AE)	0.287 (0.235)	0.00964 (0.00964)	p < 0.0001 (Fisher's exact test)
Incidence of serious adverse events—rate of SAE per participant (incidence of any SAE)	0.0559 (0.0559)	0.0627 (0.0578)	p = 1.00 (Fisher's exact test)
HR: Hazard ratio, ARR: Absolute risk reduction. Bold text indicates a statistically significant effect. <sup>a</sup> The estimated treatme	ent effect is an o	dds ratio unless stat	ed otherwise.

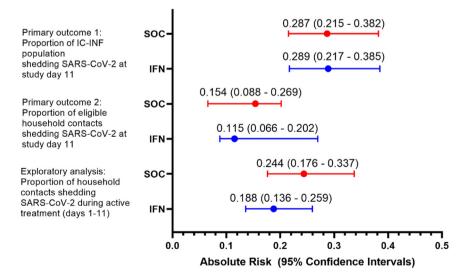
preventing infection in close contacts of ICs.<sup>43</sup> However, a trial of ring vaccination against the Alpha variant demonstrated reduced risk of contracting COVID-19 in areas with high local transmission.<sup>47</sup>

The overall viral load trajectory we observed was somewhat similar to that observed in a laboratory human challenge experiment using wild-type SARS-CoV-2 virus in healthy volunteers.<sup>44</sup> Treatment with IFN $\beta$ -1a had no effect on viral load trajectory; therefore, reduced viral shedding by infected individuals is unlikely to

explain the protective effect on household transmission of IFN $\beta$ -1a treatment that we report. Given that treatment affected the probability of transmission only in the active treatment phase and appears unrelated to viral load, we speculate that the observed effects of IFN $\beta$ -1a were direct through protection of the at risk, exposed individual rather than indirectly though effects on the index case. The ConCoRD-19 biorepository will allow further examination of the mechanisms of action of IFN $\beta$ -1a on resistance to infection.

Outcome(s)	Estimated treatment effect (95% Cl)	
Primary outcome 1		
Proportion of IC-INF population shedding SARS-CoV- 2 at study day 11	0.979 (0.647–1.479)	·
Sensitivity analyses		
Adjusted for vaccination status	0.963 (0.635–1.459)	<b>⊢</b>
Intent-to-treat index cases	1.106 (0.657–1.863)	<b>⊢</b>
Per-protocol index cases	1.062 (0.622–1.810)	· · · · · · · · · · · · · · · · · · ·
Household index case viral load (>10 <sup>6</sup> copies/mL	0.55 (0.274–1.082)	
Primary outcome 2		
Proportion of eligible household contacts shedding SARS-CoV-2 at study day 11	0.582 (0.271–1.247)	· • • • • • • • • • • • • • • • • • • •
Sensitivity analyses		
Adjusted for risk factors	0.492 (0.202–1.199)	
Household contact population	0.468 (0.178–1.231)	
Per protocol eligible household contacts	0.589 (0.272–1.276)	· • •
Excluding eligible household contacts seropositive on study day	0.640 (0.287–1.429)	
Outcome at study day 6	0.493 (0.256–0.949)	·
Index case viral load (<10 <sup>6</sup> copies/mL)	0.552 (0.306–1.12)	
Index case viral load (>10 <sup>6</sup> copies/mL)	0.113 (0.021–0.581)	<b>⊢</b> ●−−−−−1
		0.0 0.5 1.0 1.5 2.0
		Odds Ratio

**Fig. 3: Forrest plots describing primary outcomes.** The estimated treatment effect is an odds ratio unless stated otherwise. Effects were considered statistically significant if the interval estimates exclude the no-effect value of 1. Data is presented as point estimate (circle) with 95% confidence intervals (whiskers).



**Fig. 4:** Absolute risk of positive SARS-CoV-2 test for primary and exploratory outcomes in treatment and standard care arms. SOC (red): Standard of care, IFN (blue): IFNβ-1a treatment. Data is presented as point estimate (circle) with 95% confidence intervals (whiskers).

Outcome(s)	Estimated treatment effect (95% CI)			
Exploratory outcomes				
Incidence of SARS-CoV-2 shedding during the active treatment period (study days 1-11)	0.550 (0.364–0.989)		Н	-
Additional analyses				
Incidence of SARS-CoV-2 shedding during the active treatment period (study days 1-11) compared to following the active treatment period (study days 16-29)	7.076 (3.759–9.294)			<b>⊢</b> •+
Outcome where index case viral load (<10 <sup>6</sup> copies/mL)	0.930 (0.537–1.678)		1	<b></b>
Outcome where index case viral load (>10 <sup>6</sup> copies/mL)	0.168 (0.084–0.562)		⊢	
		0.01	0.1 Odds Ra	i 10

**Fig. 5: Forrest plots describing exploratory outcomes.** The estimated treatment effect is an odds ratio unless stated otherwise. Effects were considered statistically significant if the interval estimates exclude the no-effect value of **1**. Data is presented as point estimate (circle) with 95% confidence intervals (whiskers).

This study had several clear limitations due to its setting early in the course of the pandemic that may have impacted the observed outcomes. The trial was undertaken prior to the emergence of the Delta and Omicron strains (Supplementary Figure S3) and prior to widespread vaccination. The effect size observed in this study could be greater for variants with higher transmissibility such as the Omicron strains and derivatives that are now the dominant worldwide. Simulation data suggests antiviral therapy may be more effective against highly transmissible strains,48 and subgroup analysis of a clinical trial of IFN- $\lambda$  found the treatment effect was only significant against Omicron, not earlier variants.24 Additionally, early intervention is important for effective prophylaxis-this study administered the first dose within 72 h of a positive COVID-19 test or symptom onset. Based upon other recent studies, this appears to likely be within the therapeutic window for antiviral therapies.<sup>24,49</sup> Additionally, we identified two sources of potential bias in the study. Bias in primary analyses utilising intent-to-treat analysis cannot be excluded due to missing data at visit 11 for primary analysis 1 (7.9%) and primary analysis 2 (6.1%). However, there is little risk of selection bias in per-protocol analyses due to the minimal amount of non-compliance or missing data for primary outcome 1 (3.7%) and primary outcome 2 (3.1%). We calculated the sample size based upon known household transmission characteristics of the Alpha strain which was the dominant strain of the virus early in the pandemic. Simple hygiene measures and quarantine of affected individuals within households could have contributed to lower rates of transmission than expected. Together with a smaller number of eligible household contacts than anticipated from census data, these factors may have reduced the power of the study for the primary outcomes. Finally, our posthoc analysis demonstrated that certain assumptions regarding viral shedding by index cases and the likelihood of SARS-CoV-2 negative eligible household contacts at randomisation were incorrect.

In summary, intention to treat and per protocol analyses failed to demonstrate significant effects on primary outcomes. In a sub-population biologically defined as at-risk of infection, IFN $\beta$ -1a significantly reduced the probability of household transmission of SARS-CoV-2. These observations suggest that ring prophylaxis with therapies that can interrupt transmission is a strategy worth pursuing and that point of care diagnostics to identify those at highest risk for infection can increase the likelihood of success. These factors should be considered when designing future clinical trials to address transmission of highly contagious viruses.

#### Contributors

Authors Castro-Rodriguez, Fish, Montgomery, Kollmann, Iturriaga, Karpievitch, and Shannon contributed equally to the conduct of the study and manuscript preparation as first authors. Authors Montgomery, Karpievitch, Shannon, and Chen have directly accessed and verified the underlying data. Authors Tebbutt, Diego García-Huidobro, Perret, Borzutsky and Stick contributed equally to the conduct of the study and manuscript preparation as senior authors.

#### Data sharing statement

The data that support the findings of this study are available from the corresponding author, SMS, upon reasonable request.

#### Declaration of interests

Authors Castro-Rodriguez, Kollman, and Stick received funding from BHP Holdings Pty Ltd and Biogen Pty Ltd to conduct this study. Balshaw was paid consulting fees to advise on this study. Castro-Rodriguez received honoraria from AstraZeneca and GlaxoSmithKline to speak at symposia and Eurofarma for participation on advisory board. Kollmann has received research funding from NIH, Medical Research Future Fund, and Bill and Melinda Gates Foundation, honoraria from the Human Immunome Project to attend a conference, and holds two patents unrelated to this study. Montgomery received funding from NHMRC and TSANZ to present findings from this study at conferences. Authors Aniba, Borzutzky, Chen, Fish, Garcia-Huidobro, Gigi-Yunge, Hancock, Hartnell, Ho, Iturriaga, Karpievitch, Othman, Perret, Shannon, Tebbutt, and Urzua have no declared conflicts of interest.

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#### Appendix A. Supplementary data

Supplementary data related to this article can be found at https://doi. org/10.1016/j.eclinm.2023.102082.

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