Intravenous immunoglobulin and prednisolone to women with unexplained recurrent pregnancy loss after assisted reproductive technology treatment: a randomised, double-blind, placebo-controlled trial

**STATISTICAL ANALYSIS PLAN**

**Trial registration numbers:**

- Clinicaltrials.gov ID: NCT04701034
- EudraCT number: 2020-000256-35
- Ethical registration number: N-20200066
- Project ID at The North Denmark Region: 2020-156
- WHO unique trial Number: U1111-1273-8585

**Statistical Analysis Plan Version:** 1.0  
**Statistical Analysis Plan Date:** 01.06.2022  
**Number of participants enrolled:** 30

This SAP is based on study protocol version 8.0 dated 01.06.2022.
Statistical analysis plan

01.06.2022

AUTHORS:

Chief Investigator:
Caroline Nørgaard-Pedersen¹ ²*

Sponsor:
Ole B. Christiansen¹ ²

Statistician:
Regitze Gyldenholm Skals³

Affiliations of authors

¹ Centre for Recurrent Pregnancy loss of Western Denmark, Department of Obstetrics and Gynaecology, Aalborg University Hospital, Reberbansgade 15, 9000 Aalborg, Denmark.
² Department of Clinical Medicine, Aalborg University, Søndre Skovvej 15, 9000 Aalborg Denmark
³ Department of Clinical Biostatistics, Aalborg University Hospital, 9000 Aalborg, Denmark

*Corresponding author: Telephone: +45 41120267. E-mail: c.noergaardpedersen@rn.dk. Postal address: Reberbansgade 15, 9000 Aalborg, DK. ORCID: 0000-0001-9609-8396

Contribution:
The SAP was developed and described by CNP and revised by OBC. RGS helped elaborate the SAP description.

Sponsor: Ole Bjarne Christiansen

Chief Investigator: Caroline Nørgaard-Pedersen

Senior Statistician: Regitze Gyldenholm Skals
Statistical analysis plan

01.06.2022

Content

1. Introduction .................................................................................................................. 6

2. Study Methods ............................................................................................................. 7
   2.1 Trial design .............................................................................................................. 7
   2.2 Randomization and Blinding .................................................................................. 7
   2.3 Intervention ............................................................................................................ 8
   2.4 Sample size ............................................................................................................ 8
   2.5 Data collection ........................................................................................................ 9
   2.6 Timing of Interim Analysis and Final Analysis .................................................... 10

3. Study Population ......................................................................................................... 11
   3.1 Study Groups ......................................................................................................... 11
   3.2 Screening Population ............................................................................................. 11
   3.3 Study Sample Inclusion and Exclusion Criteria ..................................................... 11
   3.4 The CONSORT Flow Diagram ............................................................................. 13
   3.5 Baseline Characteristics ....................................................................................... 15

4. Study Objectives, Hypothesis, and Outcomes ............................................................. 16
   4.1. Primary Objective and Outcome ......................................................................... 16
   4.2. Secondary Objectives and Outcomes .................................................................. 17
   4.3. Tertiary, Explorative Objectives and Outcomes ................................................. 17
   4.4. Specification of Endpoints .................................................................................. 22
       4.4.1. Primary Endpoint ....................................................................................... 22
       4.4.2. Secondary Endpoints ................................................................................. 23
       4.4.3. Tertiary Endpoints ..................................................................................... 26

5. Statistical Analysis ....................................................................................................... 29
   5.1 Statistical Principles .............................................................................................. 29
   5.2 The interim analysis .............................................................................................. 29
   5.3 Analysis Methods for Primary Endpoint ............................................................... 30
   5.4. Analysis Methods for Secondary Endpoints ....................................................... 31
   5.5 Analysis Methods for Tertiary outcomes ............................................................... 31
   5.6. Major Protocol Deviations .................................................................................. 36
   5.7 Missing data .......................................................................................................... 37

6. Data Handling and implementation of Statistical Analysis Plan ................................... 38

7. Finances ....................................................................................................................... 39

Page 3 of 44

Statistical analysis plan

01.06.2022

9. References

10. List of abbreviations
The Statistical Analysis Plan (SAP) for "Intravenous immunoglobulin and prednisolone to women with unexplained recurrent pregnancy loss after assisted reproductive technology treatment: a randomised, double-blind, placebo-controlled trial" describes and expands the statistical information presented in the protocol published in BMJ Open.

<table>
<thead>
<tr>
<th>Protocol Number Code</th>
<th>CNPOBC2020</th>
</tr>
</thead>
<tbody>
<tr>
<td>Development Phase</td>
<td>Phase II</td>
</tr>
<tr>
<td>Products</td>
<td>Prednisolone and Intravenous Immunoglobulin Placebo and Human Albumin</td>
</tr>
<tr>
<td>Indication studied</td>
<td>Recurrent pregnancy loss</td>
</tr>
<tr>
<td>Clinical Trial Initiation Date</td>
<td>February 6, 2021</td>
</tr>
<tr>
<td>Clinical Trial Completion Date</td>
<td>Recruiting</td>
</tr>
<tr>
<td>Date of the Analysis Plan</td>
<td>01.06.2022</td>
</tr>
<tr>
<td>Version Number</td>
<td>1.0</td>
</tr>
<tr>
<td>SAP revisions</td>
<td>N/A</td>
</tr>
</tbody>
</table>

This study is conducted according to the Declaration of Helsinki III, local regulations, and Good Clinical Practice (GCP) guidelines and is monitored by the regional GCP board in Northern Denmark. The protocol, the informed consent form, and the participant written information is approved by the Ethics Committee of North Denmark Region, The National Board of Health, and the Data Protection Agency prior to the inclusion of participants. Informed consent from all participants is signed before enrolment.

This SAP will concern analysis of data that will be shared in more than one manuscript.
1. Introduction

Recurrent pregnancy loss (RPL) is defined as 2 consecutive pregnancy losses[1] and affects around 5% of women in reproductive age.[2] It can be divided into primary (pRPL) and secondary RPL (sRPL), and sRPL, in contrast to pRPL, define RPL patients who have had a pregnancy beyond 24 weeks of gestation before the consecutive pregnancy losses (PLs). In Denmark, more than 25% of all RPL cases happen after fertility treatments (in-vitro fertilization (IVF); intracytoplasmatic sperm injection (ICSI) or frozen embryo/blastocyst transfer (FET)), which in the following jointly are called assisted reproductive technologies (ART). The group of patients with RPL is very heterogenic. The underlying cause of RPL is unknown, probably multifactorial as a series of various risk factors is known and can be identified in less than 50% of patients. These include thrombophilia, and a group of endocrine, chromosomal, and anatomic aberrations.[3] In the remaining unexplained RPL (uRPL) cases, immunological aberrations are thought to be at least partly involved in the pathogenesis. This hypothesis is based on studies finding increased frequency of autoantibodies and specific human leukocyte antigen (HLA) alleles associated with other autoimmune diseases as well as unbalanced distribution of lymphocyte subsets, especially natural killer (NK) cells, T regulatory cells, and T-helper (Th) cells, in uRPL patients. Treatment regimens used in autoimmune diseases have therefore been suggested for uRPL patients, including intravenous immunoglobulin (IVIg) and prednisolone. A meta-analysis of RCTs evaluating the efficacy of IVIg treatment for women with RPL found no significant effect on live birth rate; however, the acquired sample size was not obtained. Since subgroup analyses did show a significant effect in sRPL, the authors suggested that further RCTs were required to obtain sufficient evidence.[4] However, a pilot study suggests that a combination of prednisone and IVIg improves the chance of live birth in women with RPL after ART.[5] Some studies suggest that the immunomodulatory treatment is effective in primarily uRPL patients with aberrant distribution of lymphocyte subsets[6–8]; nonetheless, the data is sparse. It is possible, that the lack of a significant effect of IVIg in RPL patients can be explained by a substantial heterogeneity among enrolled participants in such studies obliterating a possible effect that may present in a certain RPL subgroup. At present, we do not have evidence for an effect of the combination of IVIg and prednisolone on reproductive outcome in RPL patients after ART. Simultaneously, no clear characteristic on patients benefitting from immunomodulatory treatment have been identified. If a blood sample analysis of immune cells can identify a characteristic immune profile in patients who are more likely to respond to treatment and benefit from it (i.e., childbirth), it could help
clinicians restrict treatment to the right patients while searching for other treatments in the remaining patients, which consequently would shorten the patients’ harsh fight to get a child.

Research Hypothesis

The null hypothesis is that there is no difference in normal pregnancy rate after embryo transfer assessed at the nuchal translucency (NT) scan between RPL patients in immunomodulatory treatment and RPL patients taking placebo. The alternative hypothesis is that active treatment is a superior to placebo according to the chance of normal pregnancy at NT scan.

Study Objectives

The primary objective of this study is to determine the effectiveness of prednisolone and IVIg (P&IVIg) compared to placebo for increasing normal pregnancy rate at the NT scan.

The secondary objective is to evaluate if P&IVIg is associated with increased rate of pregnancy complications, negative pregnancy outcome, and perinatal outcomes.

The tertiary objectives are to explore how P&IVIg treatment affects the lymphocyte subset distribution (LSD) in peripheral blood, and whether the LSD can predict which uRPL patients that benefit from P&IVIg.

2. Study Methods

2.1 Trial design

The present study is a randomized 1:1, double-blinded, placebo-controlled, single-center, phase II therapeutic study conducted at Centre for Recurrent Pregnancy loss of Western Denmark, at which the physicians are specialists in treating patients with RPL. Patents from all over Denmark can be referred without costs. Participants are randomised to treatment with P&IVIg or placebo.

2.2 Randomization and Blinding

Participants are randomized in a 1:1 ratio. The participants, investigators, outcome assessor, and care provider are blinded as the Hospital Pharmacy North Denmark Region (THPNDR) perform the randomisation. Randomization will be in blocks of different sizes ensuring an even distribution after half of the number of (n=38) participants have been enrolled at what time a pre-analysis will be conducted. The randomization code will be disclosed when the last participant has completed her treatment and passed NT scan (gestational week (GW) 11-13).
2.3 Intervention:
The timing for different study interventions including participant enrolment, blood sample collection, and treatment administration is depicted on the flow diagram on Figure 1.

In the menstrual cycle in which the participant expects an embryo transfer (ET) to be carried out, one tablet per day is administered from first day of menstrual bleeding until ET; then, two tablets per day is administered from the day of the ET and until gestational week 8+0, and then gradual discontinuation with one tablet per day from week 8+0 to 8+4. If the participant does not become pregnant after ET or has a pregnancy loss (PL) before week 8+0, gradual discontinuation is initiated right after the negative test or confirmed PL diagnosis.

Intravenous (IV) infusion is administered at the time of the ET (from five working days before to two working days after), and if she has a positive pregnancy test, the IV infusion is repeated in gestational week 5, 6, and 7. Participants with a pre-pregnancy body weight (BW) ≤70 kg will receive 250 ml, participants with BW of 70-85 kg will receive 300 ml, and participants with a BW ≥85 kg will receive 350 ml in each IV infusion.

Active treatment
Prednisolone: 5 mg of prednisolone per tablet. One tablet daily before ET and two tablets daily after ET until GW 8+0.

Intravenous immunoglobulin: Privigen 100 mg/ml (10 %) (CSL Behring) with a dose of approximately 0.4g/kg.

Placebo
Oral placebo: tablets contain 85 mg lactose monohydrate, 86 mg potato starch, 8.1 mg talc, 3 mg gelatine, and 0.9 mg magnesium stearate.

Human albumin: 50 mg/ml (5 %) (CSL Behring) liquid solution for IV infusion.

2.4 Sample size
Based on results from previous studies and a hypothesis that the treatment effect is the same in women with 2 and ≥ 3 consecutive PLs, we expect a minimum live birth rate of 40 % in the P&IVIg group and of 12 % in placebo group.[5,9,10] Based on these expectations, a type I error of 0.05, and type II error of 0.20, the study needs a sample of 74 patients (37 per group) to test if the treatment is effective in uRPL patients after ART.
A dropout rate <2 % and a rate of protocol deviation <10 % (i.e., no ET) is expected. For each participant excluded from the per-protocol (PP) analysis, an extra participant will be included in order to perform a PP analysis of 74 participants.

Based on the number of new patients with ≥2 PLs after ART admitted to The RPL Centre, an inclusion period of approximately 2 years is expected.

2.5 Data collection

Baseline information will be collected before entry to the study, i.e., during the first appointment at the RPL clinic when the patient is also screened for eligibility.

Information about the primary and secondary outcomes is collected in an online questionnaire sent 13 weeks after ET i.e., after the NT scan (GW 11-13) and an online questionnaire sent after due date. If the participant does not become pregnant after ET or miscarries during the treatment, the outcome will be recorded at the time of her next follow-up, and no questionnaire will be sent. The primary outcome (live fetus at NT scan) will be recorded in the first questionnaire. If the participant does not respond on the questionnaire(s), a reminder will be sent to her, and the final attempt is made by calling the participants.

In all randomized participants, a blood sample will be collected just before first IV infusion and again 3-4 weeks later (i.e., right before third infusion is administered if the participant is pregnant. If non-pregnant, the second blood sample will be collected at the same time as it she became pregnant). Noteworthy; non-pregnant participants have only received one albumin/immunoglobulin IV infusion and have discontinued prednisolone/placebo tablet administration at the time of the second blood sample in contrast to pregnant participants who have received two IV infusions and still administer tablets).

On the day the participant is enrolled, she receives a folder containing a check-list with ticking boxes for reporting daily tablet intake and a diary for reporting adverse events/adverse reactions (AEs/ARs). The folder includes a list of known drug-related ARs according to the Summary of Product Characteristics (SmPC) for each drug, and also an explanation about which information is important to describe in detail if an AE/AR occurs. Participants are encouraged to use this folder in between follow-ups. The folder and remaining tablets should be returned to the RPL Clinic shortly after treatment cessation in order to assess treatment compliance (see section 5.6).
Statistical analysis plan

01.06.2022

At each visit, participants are also asked with open-ended questions about AE/ARs by their physician and nurse during and after IV infusion treatment to assess and journalize all AE/ARs and determine whether they are probably, possibly, or definitively drug-related or not.

For each case, it will be considered whether it is serious, unexpected, and possibly, probably, or definitely related to an intervention based on the Summary of Product Characteristics (SmPC) for each drug and the timing. The definitions of these terms are described in the published protocol and in accordance with the definition by EUR-Lex, CT-3.

The first and second online questionnaire are sent right after her NT scan and right after the due date, respectively, and will collect information on primary and secondary outcomes including the participant’s labour, pregnancy complications, AE/ARs occurring after treatment cessation, medical interventions during pregnancy, and the child’s perinatal data. She can also attach medical records on the childbirth in the second questionnaire. Non-pregnant participants are informed to contact the investigators in case of AEs/ARs up to 6 months after treatment cessation. In addition, they are offered to continue as a patient at the RPL Clinic during which follow-up on AE/ARs will be done for 6 months.

Figure 1. Flow diagram of study interventions.

2.6 Timing of the Interim Analysis and Final Analysis:
A interim analysis will be performed after 38 participants who fulfil criteria for per-protocol analysis have passed their NT scan.
After termination of the study and before breaking the blind, the final analysis will take place. It will begin with a pre-analysis blind review on data, carried out by the statisticians, to perform data validation, detect outliers, assess distribution of variables, and measure the primary, secondary, and tertiary outcome. Afterwards, the unblinded data analysis will proceed. The analysis of primary outcome is planned to begin when the last participant has passed NT scan (anticipated to be November 2023) and will be described in one report focusing on treatment effect on reproductive outcome and treatment safety. Publication will therefore await until last pregnant participant has given birth and answered the second questionnaire.

While waiting for the last participants to give birth, the analysis of the tertiary outcomes is planned to be begin, and these results will be described in one or more separate reports focussing on the effect of treatment on LSD and the association between reproductive outcome and LSD.

3. Study Population

3.1 Study Groups

The study will include 74 female patients with uRPL after ART allocated to either active treatment or placebo in a 1:1 ratio.

3.2 Screening Population

All RPL patients admitted to the RPL Center who are also undergoing ART treatment are assessed for eligibility. The number of patients who are assessed for eligibility will be summarised in the manuscript including the total number of screened patients not enrolled divided into the number that did not fulfil criteria after the diagnostic work-up, and the number of patients who did not wish to participate. The number of ineligible patients randomised, if any, will be reported with the reason for ineligibility.

3.3 Study Sample Inclusion and Exclusion Criteria

Inclusion criteria:

- Women with ≥ 2 consecutive pregnancy losses (miscarriages or biochemical pregnancies) before completed gestational week 10 after ART with the same partner or with an egg/semen donor *

Exclusion criteria:
BMI ≥35
- Age ≥41 years
- Significant uterine malformation(s)
- Known parental balanced chromosomal translocations
- ≥2 previous pregnancies with fetuses with known abnormal karyotype
- Patients with IgA deficiency, IgA-autoantibodies or hyperprolinaemia
- AMH <4 pmol/l if the next planned IVF/ICSI cycle does not imply the use of donor eggs. If IVF/ICSI with use of donated eggs donation is planned, then a low AMH value is not an exclusion criterion.
- Treatment with medication interacting with prednisolone
  - CYP3A4-inhibitors (for example erythromycin, itraconazol, ritonavir, and lopinavir), CYP3A4- inducers (for example phenobarbital, phenytoin, and rifampicin), loop diuretics, thiazides, amphotericin B, β2-agonists, antidiabetics (Metformin is acceptable), interleukin-2, somatropins, anticholinergics and regular treatment with NSAIDs.
- Patients with moderate/severe hypertension, diabetes mellitus, heart insufficiency, severe mental disorders, Cushing syndrome, myasthenia gravis, ocular herpes simplex, pheochromocytoma, systemic sclerosis, and moderate/severe renal dysfunction.
- Patients with a clinical or biochemical profile indicating need for heparin or levothyroxine treatment during pregnancy **
- Previous treatment with IVIg
- Allergy to prednisolone and/or IVIg

* The GW of the non-induced pregnancy losses will be based on the date of clinical signs of miscarriage or the fetus’ crown-rump-length of a missed abortion measured on the ultrasonic scan detecting the pregnancy loss. If the participant plans to have egg donation in the study cycle, the previous two pregnancy losses must also have happened with the use of egg donation; however, it is not required that the same egg donor has been used in all three ETs.

** Indication for levothyroxine decided by the RPL Center is plasma thyroid stimulating hormone (p-TSH) > 3.5 * 10^-3 IU/l together with presence of thyroid peroxidase (TPO) autoantibodies, and indication for heparin is previous clinical thromboses, ≥2 blood samples >3 weeks apart with IgM/IgG anticardiolipin and/or IgM/IgG β2-glycoprotein-1 antibodies > 35 kU/l, presence of lupus anticoagulant, protein S, C or antithrombin III deficiency, or patients being homozygous for the Factor V Leiden or prothrombin G20210A genetic polymorphisms. If treatment with heparin or levothyroxine is prescribed by another physician for other reasons,
this is respected, and the patient is not eligible for the study. If indication for such treatment occur after cessation of study treatment, i.e., after GW 8+4, the participant is not excluded.

3.4 The CONSORT Flow Diagram
The Consort flow diagram will be included in the manuscript comprising the number of patients who 1) were screened, 2) considered eligible, 3) gave her consent, 4) were randomised, 5) were withdrawn, 5) were not included in PP population due to major protocol deviations, and 6) lost to follow-up. Reason for ineligibility and withdrawal, and also the category of major protocol deviation will be provided. Timing for withdrawal and lost to follow-up will be arranged into four groups: before ET, before cessation of tablet treatment (GW 8), at time for first questionnaire, or at time for second questionnaire. In case of withdrawal, the data collected to date can be used while further data will not be collected.
Enrollment Flow Diagram

Assessed for eligibility (n= )
- Excluded (n= )
  - Not meeting inclusion criteria (n= )
  - Declined to participate (n= )
  - Other reasons (n= )

Randomized (n= )

Allocated to P&IVIg (n= )
- Received allocated intervention (n= )
- Did not receive allocated intervention (reasons) (n= )

Allocated to placebo (n= )
- Received allocated intervention (n= )
- Did not receive allocated intervention (reasons) (n= )

Follow-Up (week 8)

Lost to follow-up (n= )
- Discontinued P&IVIg (n= )
- Withdrawal (n= )
  (reasons)

Lost to follow-up (n= )
- Discontinued placebo (n= )
- Withdrawal (n= )
  (reasons)

Analysis (after nuchal translucency scan)

Included in ITT analysis (n= )
- Included in PP analysis (n= )
  Reason:
  Major protocol deviation category:
  1:
  2:
  3:
  4:
  5:
  6:

Included in ITT analysis (n= )
- Included in PP analysis (n= )
  Reason:
  Major protocol deviation category:
  1:
  2:
  3:
  4:
  5:
  6:
### 3.5 Baseline Characteristics

Each of the following descriptive, baseline and clinical characteristics will be reported for each treatment group in the ITT population. Tests of statistical significance will not be undertaken for baseline characteristics; rather the clinical importance of any imbalance will be noted.

Continuous measures will be summarized by mean and standard deviation (SD) (normally distributed data) or median and interquartile range (non-normally distributed data). Categorical measures will be summarized by frequencies and percentage.

**Table I: Baseline characteristics**

<table>
<thead>
<tr>
<th>Characteristic</th>
<th>Unit</th>
<th>Reporting</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age</td>
<td>Years</td>
<td>Mean (SD)*</td>
</tr>
<tr>
<td>BMI</td>
<td>Kg/m²</td>
<td>Mean (SD)*</td>
</tr>
<tr>
<td>Smoking</td>
<td>Yes/No</td>
<td>Frequency (percentage) of participants who smoke</td>
</tr>
<tr>
<td>Autoantibody*</td>
<td>Above normal range</td>
<td>Cumulative frequency (percentage) having minimum one autoantibody above normal range</td>
</tr>
<tr>
<td>Previous caesarean section</td>
<td>Number of participants</td>
<td>Frequency (percentage)</td>
</tr>
<tr>
<td>Prior reproductive outcomes</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Previous birth (after 22 weeks)</td>
<td>Number with:</td>
<td>Frequency (percentage) of each option</td>
</tr>
<tr>
<td></td>
<td>0 previous birth</td>
<td></td>
</tr>
<tr>
<td></td>
<td>≥1 stillbirth</td>
<td></td>
</tr>
<tr>
<td></td>
<td>≥1 live birth</td>
<td></td>
</tr>
<tr>
<td>Consecutive PLs</td>
<td>Number of events</td>
<td>Mean (SD)*</td>
</tr>
<tr>
<td>Consecutive PLs after ART</td>
<td>Number of events</td>
<td>Mean (SD)*</td>
</tr>
<tr>
<td></td>
<td>Number (percentage) for each value</td>
<td></td>
</tr>
<tr>
<td>Failed ART cycles (no pregnancy after ET)</td>
<td>Number of events</td>
<td>Mean (SD)*</td>
</tr>
<tr>
<td>≥1 ART cycle with prednisolone supplement initiated before ET</td>
<td>Number of participants</td>
<td>Frequency (percentage)</td>
</tr>
<tr>
<td>Indication for ART</td>
<td>Number with:</td>
<td>Frequency (percentage) of each option</td>
</tr>
<tr>
<td></td>
<td>&gt;12 months unexplained infertility</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Male factor</td>
<td>Participants, whose indication is a mix of options, are counted in all the related options</td>
</tr>
<tr>
<td></td>
<td>Tubal factor</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Endometriosis</td>
<td></td>
</tr>
<tr>
<td></td>
<td>PCOS</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Anovulation (non-PCOS)</td>
<td></td>
</tr>
</tbody>
</table>

BMJ Publishing Group Limited (BMJ) disclaims all liability and responsibility arising from any reliance on supplemental material which has been supplied by the author(s).
### Planned ART treatment in the trial

<table>
<thead>
<tr>
<th>ART treatment in study</th>
<th>Number with:</th>
<th>Frequency (percentage) of each option</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>IVF</td>
<td></td>
</tr>
<tr>
<td></td>
<td>ICSI</td>
<td></td>
</tr>
<tr>
<td></td>
<td>FET</td>
<td></td>
</tr>
<tr>
<td></td>
<td>*with PGT-A</td>
<td></td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Preimplantation stage</th>
<th>Number with:</th>
<th>Frequency (percentage) of each option</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Cleavage stage embryo transfer</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Blastocyst transfer</td>
<td></td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>No of embryos transferred</th>
<th>Number</th>
<th>Mean (SD)*</th>
</tr>
</thead>
</table>

<table>
<thead>
<tr>
<th>Gamete donation</th>
<th>Number with:</th>
<th>Frequency (percentage) of each option</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>No gamete donation</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Sperm donation</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Oocyte donation</td>
<td></td>
</tr>
</tbody>
</table>


* Selected autoantibodies include antinuclear autoantibody (ANA), anti-ds-DNA-antibodies, thyroid peroxidase (TPO) antibodies (>60 IU/mL), IgM and IgG β-2-glycoprotein-1 antibodies (20-35 kU/L), and IgM and IgG anti-cardiolipin antibodies (20-35 kU/l). Patients with higher levels of the latter two autoantibodies are not eligible for inclusion in the study.

* The median and interquartile range (IQR) (25th and 75th percentile) for non-normally distributed data

### 4. Study Objectives, Hypothesis, and Outcomes

#### 4.1. Primary Objective and Outcome

The primary objective of the study is to evaluate the clinical efficacy of treatment with P&IVIg relative to placebo in uRPL patients after ART as assessed by the number (percentage) of participants who are pregnant with minimum one apparently normal fetus alive determined at the time of NT scan in GW 11-13 (referred to as a positive reproductive outcome in the following) in the active treatment group and the placebo group. The study hypothesizes that P&IVIg increases the rate of a positive reproductive outcome compared to placebo. The relative difference between groups will be reported as the relative risk (95% CI) between treatment groups, the absolute risk reduction (95% CI) between treatment groups, and the adjusted relative risk (95% CI) for a positive reproductive outcome. A subgroup analysis will be performed on the primary outcome for pRPL and sRPL participants, separately, based on previous findings suggesting a specific treatment effect in sRPL patients[4]. However, this RCT is underpowered
to detect statistical significant differences in analyses including pRPL and sRPL patients, separately, if a true subgroup effect do exist.

An additional analysis of the primary outcome will be performed calculating the number (percentage) of participants with minimum one apparently normal fetus alive at NT scan among participants with a positive pregnancy test after ET (plasma hCG>50 IU/L).

These primary outcome analyses will be undertaken as both a PP and ITT analysis.

Based on data from over 400 patients with recurrent early pregnancy losses admitted to our clinic, we expect a low risk of miscarriage after NT scan. Therefore, the primary outcome being assessed at NT scan is expected to represent the difference in live birth rate.

### 4.2. Secondary Objectives and Outcomes

The secondary objective of the study is to evaluate the safety of P&IVIg relative to placebo when administered in early pregnancy according to both the participant and her offspring as assessed by the incidence of serious AEs and ARs (SAE/SAR), and non-serious AEs and ARs censored up until 6 months after the last IV infusion in non-pregnant participants and until after giving birth in pregnant participants. Also, for the pregnant participants, the safety assessment includes information on pregnancy complications, negative pregnancy and perinatal outcomes, and basic perinatal outcome measures censored right after giving birth. The definitions of an AE, AR, SAE/SAR, and unexpected AE/AR are in accordance with EUR-Lex, CT-3. Specific non-serious AEs occurring in ≥2 (5%) participants and any SAE in each treatment group will be presented.

The secondary outcome analysis will be undertaken as an ITT analysis.

AEs/ARs that are rather a cause of early pregnancy than the study treatment will not be recorded as AEs/ARs, including amenorrhea, vaginal hemorrhage, weight gain, backpain, pelvic pain, drowsiness, nycturia, increased hunger, nausea, and vomiting, unless these AE/ARs are serious.

### 4.3. Tertiary, Explorative Objectives and Outcomes

The tertiary objective of the study is to explore the impact of P&IVIg on the immune system in peripheral blood and whether the LSD before ET can predict reproductive outcome after ET. Several analyses will be performed immediately on the fresh blood samples collected before first IV infusion and again approximately 4 weeks later (after ET and 1-2 IV infusions), and the excess biological material will be saved for future research in a biobank.
**Statistical analysis plan**

In this SAP, the statistical analyses for the laboratory analyses performed on fresh blood are described, while future analyses on the biobank material will be described in a separate SAP later.

The tertiary outcome analyses will be undertaken as only PP analyses.

It is important to have an overview of when blood samples are collected in relation to the time when treatment is administered before analysing the tertiary outcomes. Therefore, it is described here in detail:

- **Blood sample 1p**: Outcomes from the first blood sample collected from a participant in the placebo group will represent the status of the investigated immune biomarkers at the time of ET (five working days before to two working days after ET) after administration of 1 placebo tablet daily for 2-3 weeks.

- **Blood sample 2p**: Outcomes from the second blood sample collected from a participant 3-4 weeks after ET in the placebo group will represent the status of the immune biomarkers after treatment continuation with a double dose of placebo tablets and 1-2 IV infusions (depending on the reproductive outcome) with human albumin 5% (no immunomodulatory effects expected).

- **Blood sample 1a**: Outcomes from the first blood sample collected from a participant in the active treatment group will represent the status of the immune biomarkers at the time of ET (five working days before to two working days after ET) after administration of 5 mg prednisolon daily for 2-3 weeks.

- **Blood sample 2a**: Outcomes from the second blood sample collected from a participant 3-4 weeks after ET in the active treatment group will represent the status of the immune biomarkers after treatment continuation with a double dose prednisolone (10mg from the day of ET and until a negative pregnancy test, early pregnancy loss or pregnancy week 8) and 1-2 IV infusions (depending on the reproductive outcome) with mmunoglobulin 10% (approximately 0.4g/kg). Thus, participants with sufficiently rising p-hCG receive the second IV infusion, which is one week before blood sample 2 is collected, while participants with a negative pregnancy test, insufficient p-hCG increase, or a p-hCG decrease, receive only one IV infusion (i.e., before ET = 3-4 weeks before blood sample 2 is collected) and have ceased prednisolone intake 1-2 weeks before the blood sample 2 is collected.
So, when comparing 1p and 1a, the difference will represent the effect of low dose of prednisolone. When comparing 2p and 2a, the difference will represent the effect of double dose prednisolone and IVIg. The difference between blood samples 1 and 2 will represent the effect of IV infusion(s) and natural physiologic changes during a menstrual cycle and/or early pregnancy. Noteworthy, the participants’ latest administration of study drugs before the second blood sample will vary depending on their reproductive outcome and therefore, a sensitivity analysis will decide how subgroups (based on reproductive outcome) should be divided, see section 4.4.3.1.

### 4.3.1 Flow cytometry:

#### Cell concentrations and fractions

This RCT aims to determine how P&IVIg affects the distribution of leukocyte subsets in peripheral blood and whether a difference in distribution of cells between participants with different reproductive outcomes exists.

A difference in distribution of leukocyte subsets between treatment groups will describe how P&IVIg affects the composition of the cellular immune system. By use of the current knowledge of how the different leukocyte subsets function, the results can also to some degree be used to infer the functional effects of P&IVIg.

According to the results from small observational studies, this study expects a reduction of the NK cell, Th1 cell, and Th17 cell population, and an increase of the Th2 cell and Treg cell population after P&IVIg therapy.[8,11–13] The study has no a-priori defined hypothesis on changes in monocytes, neutrophils, CD19+ B cells, and CD8+ T cells after P&IVIg, as this is sparsely explored in RPL patients.

The distribution of leukocyte subsets in participants with different reproductive outcomes will be compared. Although the subgroups within each treatment group will be small, these comparisons may elaborate if a specific immune profile (i.e., distribution of (certain) leukocytes in peripheral blood) is important for a successful implantation. Thus, the RCT is not powered to find statistically significant differences in leukocyte subset distribution between subgroups with different reproductive outcomes, and therefore, only trends for such associations can be expected to be found although a true difference may exist. Nonetheless, if a clear trend or significant difference is observed between such subgroups, it may help characterize the immune profile in peripheral blood associated with a positive reproductive outcome after ET that can be
Statistical analysis plan

01.06.2022

used in the clinic to identify uRPL patients with an immunological genesis expected to benefit from immunomodulatory P&IVIg treatment.

Since a previous study found that the impact of IVIg on Th17 and Treg cell was only significant in patients with a high Th17 and/or low Treg cell fraction before treatment[14], this RCT will evaluate if the effect of IVIg on a given lymphocyte subset is more pronounced when the level before IVIg is high or low for the given lymphocyte subset (percentile subgroup analyses).

These analyses combined may help define which immune cell profile(s) that characterizes uRPL patients in whom the P&IVIg treatment increases the reproductive prognosis.

The outcomes include

- The difference in total white blood cell (WBC) count and fractions of leukocyte subsets between treatment groups, respectively, and the change from blood sample 1 to blood sample 2 in each treatment group.
- The Th1/Th2 cell ratio and the Th17/Treg cell ratio. These cell ratios have been reported as markers to identify patients who might benefit from IVIg treatment previously.[8,15]
- The CD56_{bright} NK/CD56_{dim} NK cell ratio. This cell ratio has been reported to be lower in RPL patients than healthy controls.[16] As the CD56_{bright} NK cell type is predominant in the uterus and has immunoregulatory phenotype, and the CD56_{dim} NK cell type is predominant in peripheral blood and has a cytotoxic phenotype[17], the relative NK cell subset distribution and the treatment effects on this cell ratio may be important for implantation.
- The neutrophil-to-lymphocyte ratio (NLR), monocyte-to-lymphocyte ratio (MLR), and CD4+/CD8+ T cell ratio. These ratios are generally used as biomarkers reflecting the balance between acute and chronic inflammation (neutrophils or monocytes) and immunity (lymphocytes). To the investigators’ knowledge, they have not been reported in uRPL patients before, but the ratios are used in several diseases as a surrogate for disease activity or as a prognostic factor which could possibly also be the case for uRPL.[18–20]

Receiver operating characteristic curve analyses

In normal, uncomplicated pregnancies, studies have found that Th2 and Treg cell fractions increases while Th1 and Th17 cell fractions decrease, which paradigm is often interrupted in uRPL patients.[21–23] Previous small, observational studies have suggested that an elevated

Page 20 of 44
Statistical analysis plan

01.06.2022

Th1/Th2 ratio predicts a beneficial effect of IVIg on reproductive outcome in uRPL patients; however, these studies used Th1/Th2 cytokine ratios[24,25], which cannot be directly converted to cell ratios that will be measured in this RCT. Other studies have focused on Th17 and Treg, and they found that uRPL patients with a high Th17/Treg cell ratio before IVIg more often responded to treatment by decreasing the cell ratio and increasing the successful pregnancy rate compared to patients with normal Th17/Treg cell ratio.[14]

Instead, this RCT will determine the optimal threshold for Th1/Th2 ratio, Th17/Treg ratio, and NK cell fraction at the time for ET to determine if a specific immune profile can help clinicians identify, which patients benefit from immunomodulatory treatment.

In previous studies[6, 17, 24,26,27], an elevated NK cell fraction has been defined as >12% and it predicted a beneficial effect of IVIg on reproductive outcome in uRPL patients. Therefore, this RCT will also examine this specific threshold in a separate analysis; that will evaluate if participants in the active treatment group who have an NK cell fraction ≥12% in blood sample 1 have a higher frequency of a normal viable pregnancy censored at NT scan than participants with a NK cell fraction <12%. The same analysis will be performed in participants in the placebo group to evaluate if participants with a high NK cell fraction not receiving immunomodulatory treatment have a worse prognosis than similar patients in the active treatment group.

4.3.2 TruCulture analysis (pilot study)

A small sample of 25 participants from the RCT population will be used in a pilot study, which aims to explore if a stimulus-specific immune response is affected by immunomodulatory treatment. The analysis will be performed in the middle one-third of participants enrolled; thus, in 25 consecutively, randomized participants. The stimulus-specific immune response will be assessed as the cytokine concentration after exposure to different immune stimuli and compared within and between treatment groups. Despite a small sample size, the outcomes will also be reported for treatment subgroups separating participants into pregnant and non-pregnant patients in GW 8, to test if such subgroups of patients with different reproductive outcomes differ in response to the specific stimuli.

No prior study has reported how the stimulus-specific immune response is affected by immunomodulatory treatment in early pregnancy using the highly standardized TruCulture®
Statistical analysis plan

01.06.2022

analysis. As it is an expensive analysis with unknown relevance in this context, it will be only performed on both blood samples from 25 participants.

The outcome from the TruCulture® analysis include the concentration of ten different cytokines measured after the participant’s whole blood will be exposed to one of four different immune stimuli or no stimuli. These immune stimuli include:

- CD3/CD8: mimicking T cell activation by antigen presenting cells (APC).
- R848: mimicking single stranded RNA viral activation of toll like receptor (TLR) 7/8
- Polynosinic-polycytidylic acid (polyIC): mimicking double stranded (ds) RNA virus activation of TLR3
- Lipopolysaccharide (LPS): mimicking bacterial (E.coli) activation of TLR4
- No-stimulus null control: as a proxy for in vivo activation and baseline circulating cytokine concentration.

4.4. Specification of Endpoints

4.4.1. Primary Endpoint

The primary endpoint is the number (percentage) of participants who are pregnant with ≥1 apparently normal, viable fetus censored at NT scan (GW 11-13) by a sonographer in the ITT and PP population. The relative risk (95% CI) and absolute risk reduction (95% CI) between treatment groups will be reported.

In addition, four other endpoints will be reported:

- The number (percentage) of participants who are pregnant with ≥1 apparently normal, viable fetus censored at NT scan among women with a positive p-hCG after ET.
- The number (percentage) of participants who are pregnant with ≥1 apparently normal, viable fetus censored at NT scan among all women and among women with a positive p-hCG after ET excluding pregnancies with a fetus with confirmed aneuploidy, ectopic pregnancies and partial/complete mola pregnancies.
- The adjusted relative risk (95% CI) of being pregnant with ≥1 normal, viable fetus censored at NT scan for participants allocated to active treatment compared to the placebo group. The confounding variables included are smoking (binary), BMI, and age (continuous).
Statistical analysis plan

- Subgroup analysis: The primary endpoint in subgroups based on diagnosis of pRPL and sRPL, respectively, i.e., no prior birth or a history of ≥1 previous birth after 22 GW.

These analyses will be performed as an ITT and PP analysis. Criteria for ITT and PP analysis are described in the section about Major Protocol Deviations.

4.4.2. Secondary Endpoints

The secondary endpoints are the frequency (percentage) of the conditions or adverse events in the ITT population stratified by treatment listed in table II.

Table II: Secondary endpoints

<table>
<thead>
<tr>
<th>Secondary Endpoints</th>
</tr>
</thead>
<tbody>
<tr>
<td>Non-serious AEs and ARs</td>
</tr>
<tr>
<td>AEs/ARs (non-specific)</td>
</tr>
<tr>
<td>Specific AEs/ARs</td>
</tr>
<tr>
<td>Withdrawal</td>
</tr>
<tr>
<td>Serious AEs and ARs</td>
</tr>
<tr>
<td>SAEs/SARs (non-specific)</td>
</tr>
<tr>
<td>Specific SAEs/SARs</td>
</tr>
<tr>
<td>Withdrawal</td>
</tr>
<tr>
<td>Pregnancy complications</td>
</tr>
<tr>
<td>Preeclampsia</td>
</tr>
<tr>
<td></td>
</tr>
</tbody>
</table>
**Statistical analysis plan**  
01.06.2022

<table>
<thead>
<tr>
<th>Condition</th>
<th>Incidence</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Proteinuria</strong></td>
<td>(urine albumin/creatinine ratio &gt;300mg/g or urine albumin &gt;3g/day or &gt;1+ on urine stix) presenting after GW 20+0.</td>
</tr>
<tr>
<td><strong>Gestational Hypertension</strong></td>
<td>Incidence</td>
</tr>
<tr>
<td></td>
<td>Defined as blood pressure &gt;140 mmHg systolic or &gt;90 mmHg diastolic presenting after GW 20+0</td>
</tr>
<tr>
<td><strong>Gestational Diabetes Mellitus</strong></td>
<td>Incidence</td>
</tr>
<tr>
<td></td>
<td>Defined as an oral glucose challenge test (OGCT) with &gt;9.0 mmol/l 2 hours after oral intake of 75g glucose solution</td>
</tr>
<tr>
<td><strong>Instrumental Delivery</strong></td>
<td>Incidence of:</td>
</tr>
<tr>
<td></td>
<td>Elective caesarean section</td>
</tr>
<tr>
<td></td>
<td>Emergency caesarean section</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Condition</th>
<th>Number of participants with a negative pregnancy test after ET among all participants</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Negative Pregnancy Outcomes</strong></td>
<td></td>
</tr>
<tr>
<td><strong>Negative Pregnancy Test</strong></td>
<td></td>
</tr>
<tr>
<td><strong>Pregnancy Loss</strong></td>
<td>After a hCG measure &gt;50 IU/L</td>
</tr>
<tr>
<td></td>
<td>Specific measures of pregnancy loss:</td>
</tr>
<tr>
<td></td>
<td>The frequency of pregnancy losses with unknown or normal karyotype among all participants and all participants with a positive pregnancy test</td>
</tr>
<tr>
<td></td>
<td>The frequency of pregnancy losses with known aneuploidy among all pregnancy losses tested for aneuploidy</td>
</tr>
<tr>
<td>Statistical analysis plan</td>
<td>01.06.2022</td>
</tr>
<tr>
<td>----------------------------------</td>
<td>------------</td>
</tr>
<tr>
<td><strong>Perinatal outcomes</strong></td>
<td></td>
</tr>
<tr>
<td>Stillbirth</td>
<td>The frequency of participants with a birth after GW 24 which died in utero or within one week after birth</td>
</tr>
<tr>
<td>Congenital deformity</td>
<td>The frequency of a congenital deformity identified during pregnancy or within 1 week after birth</td>
</tr>
<tr>
<td>Abnormal karyotype</td>
<td>The frequency of an abnormal karyotype identified during pregnancy or within 1 week after birth</td>
</tr>
<tr>
<td>Prematurity</td>
<td>The frequency of a birth before 37 weeks of gestation</td>
</tr>
<tr>
<td>Small for gestational age (SGA)</td>
<td>The frequency of a birthweight &lt;10th percentile</td>
</tr>
<tr>
<td>Low birth weight</td>
<td>The frequency of a birth weight &lt;2500g</td>
</tr>
<tr>
<td>Admission to neonatal care unit</td>
<td>The frequency of an admission to the neonatal care unit in &gt;24 hours</td>
</tr>
<tr>
<td>Sex ratio</td>
<td>male:female ratio</td>
</tr>
<tr>
<td>Birth weight of singletons</td>
<td>Mean birth weight (g), SD</td>
</tr>
<tr>
<td>Gestational length</td>
<td>Mean gestational age at birth (days), SD</td>
</tr>
</tbody>
</table>
Continuous variable

(All secondary outcome variables are binary variables unless otherwise stated)

Results on binary variables except from perinatal outcome variables will be presented in a dot plot displaying the frequency, the relative risk (95% CI), and the percentage of participants in each group.[28] The perinatal outcomes will be presented in a separate table.

4.4.3. Tertiary Endpoints

The tertiary endpoints are measured in the PP population. The endpoints will be reported for each two blood samples separately, in the active treatment group and in the placebo group. For further evaluation of treatment on leukocyte subset distribution, subgroup analysis will be performed, in which the participants in each treatment group will be separated into smaller subgroups based on reproductive outcome after ET or WBC percentiles.

Subgroups of participants based on reproductive outcome

If each treatment group has <5 participants with no pregnancy (NP) and an early pregnancy loss (EPL) after ET, respectively, the two treatment groups will be divided based on a negative (NP+EPL) and positive (ongoing pregnancy (OP) at NT scan) reproductive outcome. If each treatment group has ≥5 participants with NP, EPL, and OP, respectively, a sensitivity analyses will be performed to check for differences in total WBC count and total lymphocyte count:

1) Between participants with no pregnancy (NP) and early pregnancy loss (EPL) after ET.
And subsequently,

2) Between participants receiving one and two IVIg infusions before collecting the second blood sample, i.e., a comparison of participants with NP or an EPL, who received only one IV infusion, with participants with an EPL who received two IV infusions.

If the sensitivity analysis finds no significant differences, these participants will be unified in one group representing those with a negative reproductive outcome. If the first sensitivity analysis finds a significant difference, the subgroups will be divided into three groups: NP, EPL, and ongoing pregnancy (OP) at NT scan. If the second sensitivity analysis finds a significant difference, the subgroups will be divided into the following three groups: 1) participants with NP or EPL receiving one IV infusions, 2) participants with an EPL (GA week 5-12) receiving two infusions, and 3) participants with an ongoing pregnancy (OP) at NT scan. Thus, depending on the sensitivity analyses, the subsequent analyses stratified for reproductive outcome will be in two or three subgroups.
**Flow cytometry: cell concentration and cell fraction**

Flow cytometry endpoints include those listed in table III.

<table>
<thead>
<tr>
<th>White blood cell count and differential</th>
<th>Mean total WBC count (x10⁹/L) (SD) (continuous)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean concentration (x10⁹/L) (SD) and relative percentage (SD) of the following WBC differential (continuous):</td>
</tr>
<tr>
<td></td>
<td>1) Monocytes</td>
</tr>
<tr>
<td></td>
<td>2) Neutrophils</td>
</tr>
<tr>
<td></td>
<td>3) Lymphocytes</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Lymphocyte subsets</th>
<th>The mean percentage (SD) of the following lymphocytes subsets in the total lymphocyte population (continuous).</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>1) CD19⁺ B cells</td>
</tr>
<tr>
<td></td>
<td>2) CD8⁺ Tc cells</td>
</tr>
<tr>
<td></td>
<td>3) CD3⁺CD4⁺ CCR4CCR10 Th1 cells</td>
</tr>
<tr>
<td></td>
<td>4) CD3⁺CD4⁺ CCR4CCR10CCR6 Th2 cells</td>
</tr>
<tr>
<td></td>
<td>5) CD3⁺CD4⁺ CCR4CCR10CCR6⁻ Th17 cells</td>
</tr>
<tr>
<td></td>
<td>6) CD3⁺CD4⁺ CD25⁺ CD127low Treg cells</td>
</tr>
<tr>
<td></td>
<td>7) CD3⁺CD56⁺ NK Cell (total)</td>
</tr>
<tr>
<td></td>
<td>8) CD16⁺CD56dim NK cells</td>
</tr>
<tr>
<td></td>
<td>9) CD16⁺CD56bright NK cells</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Cell ratios</th>
<th>Mean cell ratios (SD) (continuous), calculated based on cell fractions, include the following:</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>1) Neutrophil-to-lymphocyte ratio (NLR)</td>
</tr>
<tr>
<td></td>
<td>2) Monocyte-to-lymphocyte ratio (MLR)</td>
</tr>
<tr>
<td></td>
<td>3) CD8⁺Tc/CD4⁺Th ratio</td>
</tr>
<tr>
<td></td>
<td>4) Th1/Th2 ratio</td>
</tr>
<tr>
<td></td>
<td>5) Th17/Treg ratio</td>
</tr>
<tr>
<td></td>
<td>6) CD56bright NK/CD56dim NK ratio</td>
</tr>
</tbody>
</table>

If the variable is not normally distributed, the median and 25-75% interquartile range (IQR) will be reported.

These endpoints will be reported for both treatment groups, separately, and subsequently for subgroups based on the reproductive outcome.
The mean change of leukocyte fractions (absolute concentration (x10⁹/L) and/or percentage point) from blood sample 1 to 2 will be reported in each treatment group based on the formula: 
\[ \Delta \text{Leukocyte} = \text{Fraction}_{\text{before}} - \text{Fraction}_{\text{after}} \]

**Subgroup analysis of participants based on cell level percentiles**

Participants in the active treatment group will be divided into three subgroups based on ascending order of the cell fraction level of each lymphocyte subset: the lowest percentile subgroup (PS1) counting 1/3 of participants with the lowest cell fraction levels; the medium percentile subgroup (PS2) counting participants with the cell fraction levels between the 33\textsuperscript{th} and 66\textsuperscript{th} percentile; and the highest percentile subgroup (PS3) counting participants with the highest cell fraction levels above 66\textsuperscript{th} percentile.

The mean (SD) cell fraction in blood sample 1 and 2, respectively, will be presented in each percentile subgroup (PS) and the fraction in blood sample 1 will be compared to the fraction in blood sample 2. Subsequently, for each lymphocyte subset, the change (percentage points (SD)) in each PS will be compared.

**Receiver operating characteristic curve analysis**

Endpoints from the receiver operating characteristic (ROC) curve analyses include:

- The optimal threshold for Th1/Th2 cell ratio, Th17/Treg cell ratio, and NK cell fraction in the active treatment group for identification of uRPL patients at high chance of an ongoing pregnancy at NT scan.
- In active treatment group, the number (percentage) of participants with 0 or ≥1 abnormal value (i.e., abnormal immune profile), respectively, based on the three optimal thresholds calculated using data from the active treatment group.
- The number (percentage) of participants with ongoing pregnancy at NT scan in the group with 0 and with ≥1 abnormal value before ET, respectively, in each treatment group, separately.

The same endpoints will be provided only using NK cell fraction >12 % (and not using any Th1/Th2 or Th17/Treg ratio) to divide participants in each treatment group into those with or without an abnormal immune profile.

**TruCulture**

The mean (SD) or median (25-75 IQR) concentration (pg/mL) (continuous) of IFN-α2, IFN-γ, TNF-α, IL-2, IL-4, IL-6, IL-12 (p40), IL-13, IL-17, and Monocyte Chemoattractant Protein-1 (MCP-1) will be reported for each of the four different immune stimuli and the non-stimulation analysis in each blood sample, separately.
Statistical analysis plan 01.06.2022

These endpoints will be illustrated in boxplots for the active treatment group relative to placebo group. If ≥5 participants in active treatment group have a positive reproductive outcome, subgroup analysis for these endpoints in participants with OP relative to NP&EPL will be performed.

5. Statistical Analysis

5.1 Statistical Principles

All the applicable statistical tests used will be two-sided and performed using a 5% significance level unless otherwise stated.

Baseline characteristics will for continuous variables be described with mean and standard deviation (SD) (normally distributed data) or median and the interquartile range (25th and 75th percentile) (non-normally distributed data). For categorical variables, the frequency and percentage will be provided. Data distribution of continuous variables will be determined using QQ-plots and histograms.

5.2 The interim analysis

When 38 patients, who fulfill criteria for entering the PP-analysis, have been included in the study, an interim analysis will be conducted by independent statisticians in order to remain all investigators blinded. Blinded results will be shared with investigators. Differences between treatment groups regarding the primary outcome as well as the number of women with SARs/SAEs and SUSARs in the two groups (denoted A and B) will be analysed with the objective of early evaluation of unacceptable side effects of the treatments to continue the study.

The interim analysis will analyse the primary outcome and compare the incidence of SARs/SAEs and SUSARs between groups. A between group difference of ≥3 SUSARs or SARs/SAEs will lead to early termination of the trial. The alpha spending method used for adjusting significance levels for the analysis of SARs/SAEs and SUSARs between groups will be explicitly defined.[29] Analysis of the primary outcome will not lead to early termination in case of futility nor any modifications to trial procedures. The sponsor has the ultimate authority to terminate the trial.
A correlation between occurrence of SARS/SAEs and SUSARs and the primary outcome may exist, however, one such correlation will be discussed for clinical relevance and no adjustments of significance levels on the analysis of the primary outcome will be made.

5.3 Analysis Methods for Primary Endpoint

In two separate analyses in all participants in the ITT and PP population, respectively, the number (percentage) of participants in each treatment group who had an apparently normal viable fetus at NT scan will be compared using Fisher’s Exact Test (0=no pregnancy or early pregnancy loss, 1= normal pregnancy at NT scan). Also, the corresponding relative risk (95% CI) and absolute risk reduction (95% CI) will be reported.

Additional analyses on the primary outcome include:

1) The number (percentage) of participants who had an apparently normal viable fetus at NT scan among all participants with a positive pregnancy test (p-hCG >50) in each treatment group will be compared using Fisher’s Exact Test (0= pregnancy loss before NT scan; 1= normal viable fetus at NT scan).

2) The number (percentage) of participants who had an apparently normal viable fetus at NT scan among a) all participants and b) all participants with a positive pregnancy test (p-hCG >50) in each treatment group after exclusion of pregnancies with a fetus with confirmed aneuploidy, ectopic pregnancies and partial/complete mola pregnancies, will be compared using Fisher’s Exact Test (0\text{a} = no pregnancy or pregnancy loss before NT scan; 0\text{b} = pregnancy loss before NT scan; 1= normal viable fetus at NT scan).

3) A modified poisson regression with robust variance will estimate the adjusted relative risk (RR) (95% CI) for an apparently normal viable fetus at NT scan adjusted for baseline characteristics including age at enrolment (continuous), BMI at enrolment (continuous), and smoking habits at enrolment (binary). No alternative method will be applied if assumptions are not completely fulfilled, since the modified poisson regression uses a robust error variance, which is robust to violations of model assumptions.

A sensitivity analysis will be conducted with the following covariates included: low plasma mannose binding lectin (p-MBL) level (binary), previous pregnancy > 22 weeks (binary), and presence of minimum one of the measured autoantibodies (binary).

4) The primary outcome will be analysed in subgroups based on whether the participants were diagnosed with pRPL or sRPL, i.e., had no previous birth after 22 GW or had ≥1
Statistical analysis plan 01.06.2022

Previous birth after 22 GW, respectively. The relative risk (95% CI) between treatment groups will be provided.

Table IV: Definition of binary covariates

<table>
<thead>
<tr>
<th>Definition of binary variables used as covariates in regression analyses:</th>
<th>0</th>
<th>1</th>
</tr>
</thead>
<tbody>
<tr>
<td>Smoking</td>
<td>No smoking within the last 3 months before enrolment</td>
<td>Smoking minimum 1 cigarette daily during the last 3 months before enrolment.</td>
</tr>
<tr>
<td>Low MBL level</td>
<td>&gt;500 ug/l</td>
<td>≤500 ug/l</td>
</tr>
<tr>
<td>Previous liveborn</td>
<td>0 liveborn (=pRPL)</td>
<td>≥1 pregnancy &gt;22 weeks (=sRPL).</td>
</tr>
<tr>
<td>Autoantibody</td>
<td>No autoantibody</td>
<td>Minimum 1 of the following autoantibodies above normal range: TPO-Ab (&gt;60kU/l), anti-ds-DNA-Ab (&gt;10kIU/l), ANA (≥1), anti-cardiolipin and/or anti-β2-glycoprotein antibody (20-35 kU/l in two measurements with &gt;3 weeks interval).</td>
</tr>
</tbody>
</table>

5.4 Analysis Methods for Secondary Endpoints

The secondary endpoints including all binary outcome variables will be compared between active treatment group and placebo group using Fisher’s Exact Test. Furthermore, the relative risk (95% CI) will be reported. The continuous outcome variables will be compared using unpaired t-test for parametric variables and unpaired Mann-Whitney U Test for non-parametric variables.

5.5 Analysis Methods for Tertiary outcomes

5.5.1 Flow cytometry: cell concentration and fraction
The mean absolute cell count (ACC) (SD) of WBC differentials, and the mean percentage (SD) of lymphocyte subsets of the total lymphocyte population in peripheral blood in blood sample 1 and 2, separately, will be reported. These endpoints will be used in the following statistical analyses.

**Treatment effects on Lymphocytes Subsets**

Tertiary endpoints from flow cytometry analyses will be reported in a contingency table or bar/dot plot (example in Table V). A comparison within the treatment group, i.e., comparing the endpoints in blood sample 1 (before) with blood sample 2 (after IV infusion), will be performed using paired t-test for parametric variables and paired Wilcoxon signed-rank test for non-parametric variables. Between-group comparisons, i.e., comparing endpoints from each two blood samples between the two treatment groups, will be performed using unpaired t-test for parametric variables and unpaired Mann-Whitney U Test for non-parametric variables. The mean change will be calculated by subtracting the value in blood sample 1 (before IV infusion) with the value blood sample 2 (after IV infusion), and the mean change will be compared between treatment groups using the same unpaired statistical tests.

**Table V: Results from immunological analyses on blood samples collected before first IV infusion (blood sample 1) and after IV infusion(s) (blood sample 2) in uRPL patients.** The table will contain a row for each the leukocyte subsets and the cell ratios described in the section on “Tertiary Endpoints”. \( P^1 \): comparison of blood sample 1 and 2 within the treatment group. \( P^2 \): comparison of endpoints from either 1) before IV infusion or 2) after IV infusion or 3) the mean change between the sample collected before and after IV infusion within a treatment group, respectively, between the two treatment groups. ACC: absolute cell count. SD: standard deviation.

<table>
<thead>
<tr>
<th>Leukocyte subset distribution</th>
<th>Placebo</th>
<th>Active treatment</th>
<th>Comparisons between treatment groups</th>
</tr>
</thead>
<tbody>
<tr>
<td>Before IV</td>
<td>After IV</td>
<td>Mean change</td>
<td>Before IV</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
Subgroup analysis: Association between the lymphocyte subset distribution and the reproductive outcome

The participants in each two treatment groups will be divided into subgroups based on their reproductive outcome. The number of groups depend on the sensitivity analysis.

The sensitivity analysis made before dividing participants into subgroups based on reproductive outcome will use an unpaired t-test or Mann-Whitney U test to test for differences in total WBC count and total lymphocyte count when ≥5 participants are expected in each group, i.e., NP and EPL group.

Flow cytometry endpoints will be reported in a contingency table (See table II). To test for differences in ACC and/or percentage of each lymphocyte subset between subgroups based on reproductive outcome, an unpaired t-test (two subgroups) or One-Way Anova analysis (three subgroups) will be used if the variable of interest is normally distributed, and an unpaired Mann-Whitney U Test or Kruskal-Wallis One-Way ANOVA will be used if the variable of interest is skewed. To test if the ACC and/or percentages differ between participants with the same reproductive outcome but in different treatment groups, an unpaired t-test or Mann-Whitney U Test will be used depending on the data distribution of the variable of interest.

Table VI: Immune cell levels in patients with different treatments and reproductive outcomes.
The table will contain a row for each of the leukocyte subsets measured as well as the cell ratios in blood sample 1 and 2, respectively. The cell types and ratios are described in the section on “Tertiary Endpoints”. There will be one column for each subgroup based on the reproductive outcome in each treatment group (NP: no pregnancy, EPL: early pregnancy loss, OP: ongoing pregnancy at NT scan). P₁: comparison of NP&EPL with OP. P²: comparison of subgroups with the same reproductive outcome between treatment groups.
Statistical analysis plan

<table>
<thead>
<tr>
<th>Cell type</th>
<th>Placebo</th>
<th></th>
<th>Active treatment</th>
<th></th>
<th>Between treatment groups</th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>NP&amp;EPL</td>
<td>OP</td>
<td>P₁</td>
<td>NP&amp;EPL</td>
<td>OP</td>
<td>P² (NP&amp;EPL)</td>
</tr>
<tr>
<td>Total Leukocyte count</td>
<td>Before IV</td>
<td>After IV</td>
<td>Mean change</td>
<td>Before IV</td>
<td>After IV</td>
<td>Mean change</td>
</tr>
<tr>
<td>ACC, Mean (SD)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

If no differences are observed between treatment groups with the same reproductive outcome in either blood sample 1 or 2, the flow cytometry endpoints will be reported for all participants only stratified for the reproductive outcome (but not stratified for treatment) (NP and EPL combined or separated according to the sensitivity analysis for reproductive treatment subgroups) and compared (See table VII). The tests used for comparing data will be same as described for results reported in table VI.

Table VII: Immune cell levels in patients with different reproductive outcomes. The table will contain a row for each of the leukocyte subsets measured as well as the cell ratios in blood sample 1 and 2, respectively. The cell types and ratios are described in the section on “Tertiary Endpoints”. There will be one column for each subgroup based on the reproductive outcome (NP: no pregnancy, EPL: early pregnancy loss, OP: ongoing pregnancy at NT scan).

Percentile subgroup analysis: Association between the immune cell level before IV infusion and the treatment response
Three percentile subgroups (PS) will be made based on ascending leukocyte cell level in each leukocyte subset. Thus, when comparing e.g., Th1 cell fraction, participants will be divided into three groups based on the 33rd and 67th percentile of Th1 cell fraction. The same way is PSs made for each leukocyte subset (see Table VIII).

The cell fraction level in the blood sample 1 will be compared with blood sample 2 in each PS, respectively, using paired t-test for parametric variables and paired Wilcoxon signed-rank test for non-parametric variables in active treatment group. The change in cell fraction (ACC or percentage point) observed in each three PS will be compared using One-Way ANOVA for parametric variables and Kruskal-Wallis One-Way ANOVA for non-parametric variables. These analyses will be performed for Th1, Th2, Th17, Treg, CD8+ T cell, B cells, total NK cells, and the two NK subsets.

Table VIII: Lymphocyte subset level in percentile subgroups (PS) before and after intravenous (IV) infusion (blood sample 1 and 2). The level before is compared with the level after IV. The mean change in each PS is compared. The will one row for each lymphocyte subset described in section “Tertiary endpoints”.

<table>
<thead>
<tr>
<th></th>
<th>PS1</th>
<th>PS2</th>
<th>PS3</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Before IV</td>
<td>After IV</td>
<td>Mean change</td>
</tr>
<tr>
<td>Th1 cells, mean percentage (SD)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>CD3:CD4:CCR4:CCR10</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

**Receiver operating characteristic curve analyses**

ROC curve analyses will be performed to determine the performance of Th1/Th2 ratio, Th17/Treg ratio, and the NK cell fraction in blood sample 1 to predict a positive reproductive outcome (i.e., pregnancy at NT scan) in the active treatment group. The optimal cut-off for each of the three immune markers will be defined as the value with maximum sensitivity and specificity in the ROC curve analysis.
The AUC value and the Youden’s index ($J_{\text{max}}$) together with the corresponding sensitivity, specificity, positive predictive value (PPV), and negative predictive value (NPV) will be reported for each ROC curve analysis.

Noteworthy, the sample size in this RCT has not been calculated based on this analysis and with only this small sample available, no post validation cohort will be separated from the treatment group before the ROC curve analysis will be performed. Bootstrapping with 1000 bootstrap samples will be applied to estimate 95% confidence limits on the estimates.

Using the three thresholds, the number of participants in active treatment group with $\geq 1$ “abnormal” value will be determined. An abnormal value will be defined as a value higher than the threshold for Th1/Th2 ratio, Th17/Treg ratio, and the NK cell fraction. The number of participants with an ongoing pregnancy at NT scan will be compared between participants with 0 and with $\geq 1$ abnormal value in the treatment groups, separately, using $\chi^2$ Test or Fisher’s Exact Test.

The number (percentage) of participants with a total NK cell fraction $>12\%$ (including both CD3$^{-}$CD16$^{+}$CD56$^{\text{dim}}$ and CD3$^{-}$CD16$^{-}$CD56$^{\text{bright}}$) of the total lymphocyte population in the sample before and after IV infusion, respectively, will be reported. The number of participants with elevated NK cell fraction before IV infusion will be compared with the number after IV infusion using $\chi^2$ Test or Fisher’s Exact Test.

The number of participants with an ongoing pregnancy at NT scan among all participants with NK cell fraction $\geq 12\%$ and $<12\%$, respectively, will be compared using $\chi^2$ Test or Fisher’s Exact Test in each treatment group, respectively.

5.5.2 TruCulture pilot study: Immune response ex vivo after induced stimulation

The concentration of cytokines before and after IV infusion will be compared in each treatment group, separately, using unpaired t-test or Mann-Whitney U Test depending on the distribution of the variable of interest. Results will be presented in boxplots for each of the four stimuli-specific protocols and for the no-stimuli protocol. The boxplot will include the information described in Table IX.
Statistical analysis plan

Table IX: For each stimulator, mean cytokine concentration (pg/ml) will be presented in a box plot containing data stratified and compared as described in this table, evaluating the treatment effect on the stimulus-specific immune cell response by in-group and in-between group comparisons. Information on each cytokine will be reported. P\textsuperscript{1}: comparison of blood sample 1 and 2 within the treatment group. P\textsuperscript{2}: comparison of endpoints from either 1) before IV infusion or 2) after IV infusion or 3) the mean change between the sample collected before and after IV infusion within a treatment group, respectively, between the two treatment groups.

<table>
<thead>
<tr>
<th>Stimulator</th>
<th>Placebo</th>
<th>Active treatment</th>
<th>Comparison between groups</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Before IV</td>
<td>After IV</td>
<td>Mean change</td>
</tr>
<tr>
<td>IFN-α2</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

5.6. Major Protocol Deviations

The predefined major protocol violations are divided into six categories described in Figure 2.

---

Figure 2: description of major protocol deviations excluding participants from per-protocol population.
Statistical analysis plan

01.06.2022

All randomised participants will be included in ITT analysis population. Participants who completed the trial with no major protocol deviations will be included in the PP analysis. The reason for not fulfilling eligibility criteria will be reported. In case of category 4 violations, the participant will be counted in according to the treatment she was randomised to receive. All protocol deviations will be recorded in the eCRF and summarised by the treatment group and the category with frequency and percentage of all included participants.

5.7 Missing data

To prevent missing data, the eCRF has a record status dashboard with an overview of instruments to be filled in after follow-up appointments including instruments for the investigator to fill in and questionnaires for the participant to fill in. When the instrument is changed from incomplete to complete, i.e., all data items are filled in, the instrument on the record status dashboard changes colour from red to green. Also, key data items on primary and secondary outcomes collected at follow-ups before gestational week 9 are described in the participant’s medical records. In addition, AEs and treatment compliance are described in the information folder returned to the RPL Clinic after treatment cessation. Thus, these key data items are readily obtainable if missing in the eCRF by the end of the study. For this reason, the degree of missing data on key data items is expected to be minor or not existing. However, in case of missing data, the number missing will be reported.

For the analysis of the primary and secondary outcome, complete case analysis will be performed. In analysis tertiary outcomes, the management of missing data will follow the guideline presented by Jakobsen et al. and include complete case analysis and single or multiple imputation.[30] A sensitivity analysis will be performed if MI or other methods to account for missing data has been used in accordance with the description by Jakobsen et al.[30]

6. Data Handling and implementation of Statistical Analysis Plan

All data regarding primary and secondary (except data on blood samples) outcome will be collected and stored in an electronic case report form (e-CRF) in a REDCap database (Vanderbilt University, Nashville, TN, USA) which only the sponsor and chief investigator have full access to. GCP board members have rights to “see only”. Information will be obtained from the participant’s electronic medical records, information folder, and directly from the participant. REDCap makes audit trail logging and back-ups. The eCRF will be used to collect
Statistical analysis plan 01.06.2022

all clinical data about the participant, and also to automatically send out and collect questionnaires replies after NT scan and after due date. Personal identifiers are marked, and they will not be exported from the database when the study terminates. Data will be exported to Microsoft Excel (Microsoft Corporation, Washington, USA) and prepared for analyses performed in Stata/MP 15.0 (TX, US) for Mac.

Data regarding tertiary outcomes will be collected in a secured database at the Department of Clinical Immunology (DCI) until all participants have been enrolled and the last pregnant participant has passed her NT scan. At this time, blinded data will be shared with the sponsor and chief investigator at the RPL clinic and not at any time before, since sharing results from these analyses at any time before may reveal enough information for investigators to speculate and predict treatment allocation. Thus, in order to remain fully blinded, data on peripheral blood immunological analyses will only be handled by the DCI personnel, who have no contact to study participants, until all data has been collected.

When the last pregnant patient has passed her NT scan, we will start data analysis. Before data is unblinded, a blind review pre-analysis on data will be carried out to check for outliers in continuous outcome variables, distribution of continuous and ordered data (using Q-Q plots and histograms to assess data normality), missing data points (and if possible, collecting these missing data points) and important potential covariates identified in other recent research may be added to statistical models. This final pre-analysis will be handled by statisticians who had not been involved in the study. They will present the investigators for any uncertainties as well as blinded interpretations of the primary endpoint results. The pre-analysis may add modifications to the SAP, which will all be described in an amendment to the SAP before performing unblinded analyses. After data validation is complete, unblinded data analysis will proceed. Unblinded analyses will be conducted by the chief investigator supervised by statisticians.

This SAP will be used by all statistician and investigators analysing study data. Analysis of primary, secondary, and tertiary endpoints will be carried out in continuous order in the pre-analysis.

7. Finances

The investigators gratefully acknowledge The Svend Andersen Fund for funds to help defray medicinal expenditure, “Grosserer L.F. Foght’s” fund and “Beckett-Fonden” for funds for expenditure for immunological analyses. The funding sources had no role in the design of the

Page 39 of 44
Statistical analysis plan

01.06.2022

study and will not have any role during its execution, analyses, interpretation of the data, or
decision to submit results. The investigators also thank the DCI for performing all
immunological analyses and preparing infusion medicine for all participants.
9. References

Statistical analysis plan


10. Abbreviations

AaUH: Aalborg University Hospital

ACC: Absolute cell count

AE: adverse events

ANA: antinuclear autoantibody

AR: adverse reactions

ART: assisted reproductive technologies

BMI: body mass index
Statistical analysis plan

DCI: the Department of Clinical Immunology
e-CRF: electronic case report form
EPL: early pregnancy loss
FET: frozen embryo/blastocyst transfer
GCP: Good Clinical Practice
HLA: Human leukocyte antigen
ICSI: intracytoplasmatic sperm injection
Ig: Immunoglobulin
ITT: Intention to treat
IVF: in-vitro fertilization
IVIG: intravenous immunoglobulin
Jmax: Youden index
LR: likelihood ratio
NK: natural killer
NP: no pregnancy
NPV: negative predictive value
OGCT: oral glucose challenge test
OP: ongoing pregnancy
OR: odds ratio
p-MBL: plasma mannose binding lectin
PGT-A: pre-implantation genetic testing for aneuploidy
PP: Per protocol
PPV: positive predictive value
pRPL: primary recurrent pregnancy loss
PS: percentile subgroup
ROC: receiver operating characteristic
RPL: recurrent pregnancy loss
RR: relative risk
SAE: serious adverse event
SAP: statistical analysis plan
SAR: serious adverse reaction
SD: standard deviation
SGA: small for gestational age
sRPL: secondary recurrent pregnancy loss
Statistical analysis plan

SUSAR: suspected unexpected serious adverse reaction
Tc: cytotoxic T
Th: T-helper
TPO: thyroid peroxidase
Treg: regulatory T
uRPL: Unexplained recurrent pregnancy loss