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**Intraamniální infekční a zánětlivé komplikace spojené s
předčasným odtokem plodové vody**

**Infection-related intra-amniotic complications in women
with preterm prelabor rupture of membranes**

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Hradec Králové, 2017

Defence on:

Declaration

I declare hereby that this thesis is my own original work and that I indicated by references all used information sources. I also agree with depositing my thesis in the Medical Library of the Charles University in Prague, Faculty of Medicine in Hradec Králové and with making use of it for study and educational purpose provided that anyone who will use it for his/her publication or lectures is obliged to refer to or cite my work properly.

I give my consent to availability of the electronic version of my thesis in the information system of the Charles University in Prague.

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Abbreviations

| | |
|--------|--|
| BDNF | Brain-derived neurotropic factor |
| BMI | Body mass index |
| CAL | Clinical attachment loss |
| CRP | C-reactive protein |
| DMF | Decayed, Missing, and Filled index |
| ELISA | Enzyme-linked immunosorbent assay |
| EOS | Early onset sepsis |
| FIRS | Fetal inflammatory response syndrome |
| GI | Gingival index |
| GM-CSF | Granulocyte macrophage colony stimulating factor |
| HCA | Histological chorioamnionitis |
| HMGB | High mobility group box protein |
| HSP | Heat shock protein |
| IAI | Intra-amniotic inflammation |
| IGFBP | Insulin-like growth factor-binding protein |
| IL | Interleukin |
| IQR | Interquartile range |
| MCP | Monocyte chemotactic protein |
| MIAC | Microbial invasion of the amniotic cavity |
| MIP | Macrophage inhibitory protein |
| MMP | Matrix metalloproteinase |
| PCR | Polymerase Chain Reaction |
| PLI | Plaque index |

| | |
|--------|--|
| PPD | Probing pocket depth |
| PPROM | Preterm prelabor rupture of membranes |
| PTL | Spontaneous preterm delivery with intact membranes |
| rRNA | Ribosomal RNA |
| sTNF-R | Soluble TNF receptor |
| TGF | Tumor growth factor |
| TNF | Tumor necrosis factor |
| WBC | White blood cell |
| WHO | World Health Organization |

1 Introduction

1.1 Preterm delivery

Preterm delivery is defined by the World Health Organization (WHO) as delivery at less than 37 weeks of gestation (1). Preterm delivery results from various disorders related to the mother and/or fetus, some of which are explained, and some that are of unknown etiology (2).

Worldwide, the rate of preterm birth is approximately 11%, and almost 15 million preterm births occur annually (3). In the US, preterm births represent 11-13% of all deliveries; in Europe and other developed countries, that number is approximately 5-9% (4). Despite advancing knowledge about the risk factors and mechanisms leading to preterm delivery, the rate of preterm birth has risen in most developed countries in the last 20 years (5). Despite corrections for difference in different risk factors there are huge differences in preterm delivery between different international countries and also within Europe(6)

Preterm birth is responsible for 75% of perinatal mortality and more than half of the long-term morbidity (7). Improvement in neonatal outcome has occurred since antenatal corticosteroids and magnesium neuroprophylaxes have been implemented into clinical management (8). The crucial point with regard to perinatal adverse outcomes is represented by the gestational age at delivery, when late preterm-born neonates have significantly lower mortality and morbidity rates compared with those born at earlier gestational ages (9, 10).

Preterm birth represents a significant socioeconomic problem, and annual costs of prematurity in the US were 26.2 billion dollars in 2005 (11). The costs do not include just the newborn hospitalization in the neonatal period, but also include maternal delivery costs, medical care up to 5 years for neonates, and needs for special help with handicaps that result from preterm birth (12).

Preterm delivery represents a rather heterogeneous condition that can be subdivided according to various conditions and aspects.

First, based on the phenotypes, it can be divided into the following categories: i) spontaneous preterm delivery with intact membranes (PTL), ii) preterm prelabor

rupture of membranes (PPROM), and iii) iatrogenic preterm delivery for maternal or fetal indications. Approximately 30% to 35% of preterm births are iatrogenic, 40% to 45 % are PTL, and 25% to 30% are PPROM(13).

Second, when gestational age of delivery is taken into consideration, preterm delivery might be divided into the following subgroups:

1. Extreme preterm delivery, when delivery is at less than 28 weeks of gestation (5%).
2. Early preterm delivery, when delivery is at 28-31 weeks of gestation (20%).
3. Moderate preterm delivery, when delivery is at 34-36 weeks of gestation (60-70%).
4. Late preterm delivery, when delivery is at 34-36 weeks of gestation (10%).

1.2 PPROM

PPROM occurs in between 2-8% from all pregnancies (14). PPROM is responsible for approximately one third of preterm births, and it substantially contributes to significant perinatal morbidity and mortality (10, 15).

1.2.1 Definition

PPROM is defined as a leakage of the amniotic fluid before onset of regular uterine contractions before 37 weeks of gestation (10, 16). However, what the interval between leakage of the amniotic fluid and onset of labor, called latency, represents is a controversial issue, as there is no clear consensus regarding the length of latency required to define PPROM. However, in a vast majority of the studies, 1 hour of latency is enough to define PPROM (17, 18).

1.2.2 Pathophysiology

PPROM has a multifactorial etiology and might be diagnosed as one or more etiologic processes in different patients (10, 19, 20). Similarly to PTL, lower socioeconomic status, cigarette smoking, sexually transmitted infection, uterine distension, amniocentesis, and vaginal bleeding in pregnancy are associated with a higher risk of PPROM (1, 20).

Infectious etiology used to be considered to play a crucial role in the pathophysiology of PPRM; however, recent studies have shown that the majority of cases show no presence of bacteria in the amniotic fluid, nor intra-amniotic inflammation (21-24).

Human fetal membranes are composed of a single layer of amnion epithelium that lines the interior of the amniotic cavity and multilayered chorion trophoblast layers connected to maternal decidua (25). These two layers are connected through an extracellular matrix region, which is known for its structural integrity. In terms of PPRM, proteolysis disrupting the extracellular matrix is hypothesized as the dominant factor that weakens the amniotic layers resulting in PPRM (22). Recently, PPRM has been considered to be a disease of fetal membranes, in which multiple risk factors associated with oxidative stress and inflammation might accelerate fetal membrane senescence, apoptosis, and proteolysis, leading to fetal membrane weakness (26). Fetal membrane cells undergoing senescence might produce inflammatory mediators, which leads to the development of a proinflammatory environment in the fetal membranes (24). This condition is related to the activation of various matrix metalloproteinases cleaving fetal membrane collagen. This process results in the reduction of the tensile strength of fetal membranes and leads to their rupture (23, 24, 27).

Based on the pathophysiologic processes leading to PPRM, it might be classified into three major groups: i) PPRM with the absence of cervical change and longer latency to delivery, ii) PPRM involving bleeding disorders or coagulopathies, and iii) PPRM associated with cervical changes and shorter latency to delivery (27).

1.2.3 Diagnosis of PPRM

The diagnosis of PPRM is made primarily on the mother's history, which may have an accuracy of 90% for the diagnosis of PPRM (17). The most objective standard for the diagnosis of PPRM is the confirmation on speculum examination visualizing fluid passage from the cervix or pooling in the posterior vaginal fornix. Although the most common indicator for identifying leakage of the amniotic fluid is represented by positive Temesvary test (pH indicator changes from yellow to green on contact with alkaline amniotic fluid), this approach is associated with a high false-positive rate, related to contamination with vaginal discharge, urine, blood, or semen (10, 16, 28,

29).

The most sensitive and specific commercially available tests are based on the confirmation of insulin-like growth factor-binding protein 1 and placental alpha macroglobulin 1 from amniotic fluid, detected with ActimTM PROM and Amnisure[®] ROM tests, respectively (30, 31).

1.2.4 Infection related and inflammatory complications of PPRM

Pregnancies affected by PPRM might be complicated by microbial invasion of the amniotic cavity (MIAC) and intra-amniotic inflammation (IAI). The presence of these two complications is found in approximately 30% to 60% of cases of PPRM, respectively (32, 33). Both these complications might result in the development of histological chorioamnionitis. The presence of MIAC is thought to activate intra-amniotic immune response through the system of pattern recognition receptors, which results in the development of microbial-associated IAI (33-36). Moreover, the intensity of IAI depends on the type of bacteria and their microbial load (37-39). Therefore, the presence of a small number of bacteria with a low virulent potential, such *Ureaplasma* species, in the amniotic fluid is unlikely to elicit IAI (39, 40). Therefore, this condition is not associated with worse pregnancy and neonatal outcomes. This scenario is characterized as a colonization of amniotic fluid (37, 41).

In contrast, some endogenous mediators called alarmins (e.g., high mobility group box-1 protein) are released from necrotic cells into the amniotic fluid and can trigger IAI through the same system of pattern recognition receptors as in the infectious scenario (23, 42). This scenario leads to the development of sterile IAI (the presence of IAI without any proven microorganisms in the amniotic fluid).

1.2.4.1 MIAC

MIAC is defined as the presence of microorganisms in the amniotic fluid. Microorganisms may invade the amniotic cavity by four potential routes: i) ascension from the vagina and cervix, ii) hematogenous disseminations followed by transplacental invasion, iii) retrograde dissemination through the fallopian tubes, and iii) iatrogenic inoculation by intrauterine procedures (43).

Diagnostic approaches for the identification of microorganisms in the amniotic cavity should consider the fact that a vast majority of micro-organisms found in the amniotic fluid are difficult-to-cultivate bacteria or non-cultivable bacteria. Therefore, non-cultivation methods appear to be an essential tool for the detection of MIAC and dramatically improve the detection rate of MIAC compared with conventional cultivation methods (44-47).

The most commonly found bacteria in PPROM are genital mycoplasmas (e.g., *Ureaplasma* sp., *Mycoplasma hominis*) (48-50). Genital mycoplasmas are the smallest bacteria without a cell wall. Their surface contains lipo-proteins and antigens, which can induce host inflammatory response through pattern recognition receptors (51). Genital mycoplasmas were considered as bacteria with low virulence, but their presence in the amniotic fluid can elicit strong intra-amniotic inflammatory response comparable to other aerobic and anaerobic bacteria (39, 40, 52). The bacterial load of genital mycoplasmas is related to intra-amniotic inflammatory response, whereas no association to fetal inflammatory response was observed (37, 38).

In addition to the genital mycoplasmas, there are other bacteria found in amniotic fluid. Some of them might have an origin in the oral cavity (53-55). Therefore, it has been hypothesized that bacteria might spread from the oral cavity through the bloodstream to the placenta and finally to the amniotic fluid (56, 57). This theory is supported by the facts that oral bacteria such as *Fusobacterium nucleatum*, *Porphyromonas gingivalis*, and group A *Streptococcus* have been found in the amniotic fluid (58-60).

1.2.4.2 IAI

IAI is defined as the elevation of inflammatory marker in the amniotic fluid. Various cytokines, chemokines, proteins, and their combinations have been suggested to be potential markers of IAI (33, 34). However, from a clinical point of view, just two of them have reached the clinics—amniotic fluid interleukin-6 (IL-6) and matrix metalloproteinase-8 (MMP-8) (48, 61, 62).

Despite the fact that IL-6 is a non-specific pleiotropic cytokine, IL-6 has been broadly used as a marker of IAI. Given IL-6 as a marker of IAI, two different cut-off values have been proposed to diagnose IAI with respect to the methods of amniotic fluid IL-

6 evaluation (48, 63). Cut-off values 2600 pg/mL and 745 pg/mL have been suggested for IAI identification when IL-6 is measured by ELISA and bedside test, respectively.

However, our groups have proposed the cut-off value of 1000 pg/mL for amniotic fluid IL-6 measured in a bedside manner to identify the presence of both MIAC and histological chorioamnionitis (HCA) (64). Recently, this cut-off has been shown to have at least a similar diagnostic value for IAI as the aforementioned cut-off value of 745 pg/mL (65).

1.2.4.3 HCA

Acute HCA is defined as diffuse neutrophil infiltration of the fetal membranes, the placenta, and the umbilical cord blood in a response to microbial invasion or due to other pathological processes (66, 67). Different classification systems to describe the intensity and composition of HCA have been suggested; however, the Redline and the Salafia classifications represent the most commonly used classification systems worldwide (68).

Acute HCA might reflect maternal and fetal inflammatory responses (69, 70). The latter is characterized by neutrophil infiltrations of the umbilical cord and/or chorionic plate vessels (68).

Acute HCA might be split into two subgroups based on the cause leading to its development: i) infectious HCA, which occurs when both MIAC and HCA are present and ii) sterile HCA, which is defined as HCA without MIAC.

1.3 Inflammatory response according to the presence of MIAC, HCA, IAI

The presence of both MIAC and HCA is associated with the most intensive IAI in PPRM compared to the PPRM pregnancies complicated by HCA alone (sterile HCA), MIAC alone (colonization), and PPRM pregnancies without both MIAC and HCA (64, 69). Similar results were observed in terms of the intensity of the fetal inflammatory response, measured by umbilical cord blood IL-6 concentrations (70). Moreover, pregnancies complicated with MIAC and HCA are prone to subsequent development of early-onset sepsis in newborns (71).

On the other hand, the presence of HCA alone or MIAC alone has similar intra-amniotic and fetal inflammatory responses as in women without either MIAC or HCA. Likewise, the newborns from these pregnancies do not have a higher risk of the development of early-onset sepsis (64, 69).

The presence of MIAC and HCA might be considered to be outcomes only when PPRM is managed actively owing to a short interval between the amniotic fluid and the placenta samplings. Therefore, in PPRM managed expectantly, when a longer latency is expected, the presence of MIAC and HCA cannot be taken as an outcome. In women with PPRM managed expectantly, the knowledge about MIAC and IAI more appropriately describe an intra-amniotic environment.

The presence of both MIAC and IAI (microbial-associated IAI) has been shown to be related to the highest intra-amniotic, cervical, and maternal inflammatory responses (72-74). Interestingly, no difference has been shown between the intensity of fetal inflammatory responses triggered by microbial-associated and sterile IAI (75). It means that the fetal inflammatory response is driven by IAI regardless of the presence or absence of bacteria in amniotic fluid.

1.4 Diagnostics approaches for identifying infection-related and inflammatory intra-amniotic complications

Currently, the evaluation of amniotic fluid obtained by transabdominal amniocentesis is still considered as the gold standard to show infection-related and inflammatory intra-amniotic complications. Despite a solid body of evidence that transabdominal amniocentesis is a safe and reliable procedure in PPRM, clinicians are reluctant to perform amniocentesis due its invasive nature. Therefore, there is an urgent need for surrogate markers from noninvasively obtained body fluids such as cervical and vaginal fluids or maternal blood. Nevertheless, their diagnostic value still has not reached the level of the amniotic fluid markers because of their limited sensitivity and specificity.

Along with the body fluid markers, different ultrasound markers (e.g., the amount of amniotic fluid, fetal adrenal glands, thymus, and spleen) have been suggested to serve as the markers of intra-amniotic infection-related and inflammatory conditions (76-

78). Unfortunately, these ultrasound markers must be further validated before their potential consideration as clinical markers.

1.5 Management

In general, two different management approaches of PPRM are considered, the expectant and the active approaches. Since the gestational age of delivery has been clearly shown to be a main determinant of neonatal morbidity and mortality, the expectant management is recommended in PPRM prior to 34 weeks (79, 80). On the other hand, the management of PPRM beyond 34 weeks remains a controversial issue. However, PPRMEXIL I and II studies have not found any differences in the rates of early-onset sepsis between women with PPRM beyond 34 weeks treated actively and expectantly (81, 82). Moreover, the recent PPRMT trial has shown the reduction of the respiration morbidity and a low risk of the development of early-onset sepsis when expectant management in PPRM beyond 34 weeks was used (83). This suggests that the expectant management might be a method of choice in women with PPRM beyond 34 weeks as well.

However, the presence of IAI and microbial-associated IAI is associated with short-term and long-term consequences (75, 84). Archabald et al have shown that newborns from pregnancies with PPRM complicated by microbial-associated IAI and treated expectantly had the worst outcome (85). Therefore, the individualized management of PPRM based on the knowledge about the presence of MIAC and IAI might be the best option.

1.6 Neonatal outcomes

PPRM might be characterized as a pregnancy complication associated with a relatively high rate of the neonatal morbidity and mortality. The morbidity and mortality are mainly related to gestational age of delivery (13, 15, 20, 86, 87). However, the neonatal outcome from PPRM has been shown to be worse than that from PTL.

Because a majority of PPRM might be complicated by the presence of MIAC, IAI, and HCA, it is possible to anticipate that newborns from these pregnancies are jeopardized by the development of fetal inflammatory response syndrome (FIRS)

(70). This condition is characterized by the elevation of umbilical cord blood IL-6 greater than 11 pg/mL. The presence of funisitis and/or chorionic plate vasculitis represents the histopathological hallmark (88). It is well known that fetuses affected by FIRS are at higher risk of multi-organ disorders and serious neonatal morbidity (89, 90).

However, there is a lack of information about whether and how the presence of MIAC and HCA might affect the short-term neonatal outcome.

2 Periodontal diseases and pregnancy

Periodontal disease represents one of the most frequent chronic inflammatory conditions in the adult population. The reported prevalence of chronic periodontal disease is approximately 10% to 90% in the adult population (91). Periodontal disease is often caused by periopathogenic bacteria such as *Fusobacterium* sp., *Prevotella* species, *Porphyromonas* species, and other anaerobic gram-negative bacilli. It has been suggested that the presence of periodontal disease in pregnancy might cause translocation of periopathogenic bacteria to uteroplacental circulation, damage the placenta, and potentially lead to preterm delivery (92). Periodontal disease is often caused by periopathogenic bacteria such as *Fusobacterium* species, *Prevotella* species, *Porphyromonas* species, and other anaerobic Gram-negative bacilli. It has been suggested that the presence of periodontal disease in pregnancy might cause translocation of periopathogenic bacteria to uteroplacental circulation, damage the placenta, and potentially lead to preterm delivery (93, 94). In addition, periopathogenic bacteria have been demonstrated as a cause of MIAC in women with PPRM (53).

Periodontitis, a chronic local inflammation, might be associated with elevation of pro-inflammatory cytokines in the cervical fluid, such as IL-6 (95-98). These inflammatory mediators may reach the systemic circulation, which could lead to a low-grade systemic inflammation (58, 99). However, there is a lack of knowledge regarding whether this originally local inflammatory response localized in gingival cervical fluid may affect the intra-amniotic compartment via the spreading of bacteria or their products or through the systemic inflammatory response in women with PPRM and lead to the development of sterile IAI.

3 Objective of the thesis

Pregnancies complicated with PPRM may represent clinical dilemma in their management, therefore objectives of the thesis are drafted and can be understood as a review for clinicians,

The first objective of the thesis deals with possibilities of assessment non-invasive tool for prediction of infection-related intra-amniotic complications in PPRM pregnancies, when C-reactive protein (CRP) is used in daily routine work. The second objective clarifies the consequences to neonates in according to each infection-related intra-amniotic subgroups in PPRM pregnancies. The third and fourth objectives evaluate possible relation between periodontal diseases and PPRM, when was evaluate periodontal status and local inflammatory response in gingival fluid.

Therefore, there are four specific aims of the thesis:

1. To evaluate whether maternal serum CRP has a diagnostic value for infection-related intra-amniotic complications in PPRM pregnancies. An additional aim with close relation to the first aim was to evaluate whether there was a correlation between the maternal inflammatory response, measured by maternal serum CRP, and microbial load of *Ureaplasma* species.
2. To identify whether the presence of infection-related intra-amniotic complications is associated with worse short term neonatal outcomes in pregnancies complicated by PPRM in women below 34 weeks of gestation.
3. To investigate if there is an association between the periodontal status and infection-related intra-amniotic complications in pregnancies complicated by PPRM.
4. To identify the association between the local inflammatory response in gingival crevicular fluid, measured by the levels of multiple proteins, and maternal and intra-amniotic inflammatory responses measured by maternal CRP and amniotic fluid IL-6 concentrations, respectively, in women with PPRM.

4 Set of patients, methods and statistical analysis

4.1 Set of patients

4.1.1 Specific aim I.

A prospective cohort study was performed. The study population consisted 386 women with singleton pregnancies at gestational ages between 24+0 and 36+6 weeks who were admitted for PPRM to the Department of Obstetrics and Gynecology in Hradec Kralove between January 2008 to December 2013.

4.1.2 Specific aim II.

Prospective observational cohort study was performed. The study population consisted of 122 women with PPRM at gestation age 24+0-34+0 weeks who were admitted to the Department of Obstetrics and Gynecology, University Hospital Hradec Kralove, Czech Republic between June 2008 and February 2012. All women included in specific aim II. were part of the specific aim I.

4.1.3 Specific aim III. and IV.

A prospective study of pregnant women between 24+0-36+6 weeks gestation with PPRM, who were admitted to the Department of Obstetrics and Gynecology at the University Hospital Hradec Kralove in the Czech Republic between December 2014 and April 2016, was conducted. A total of 78 women with PPRM met inclusion criteria to the study.

4.2 Inclusion criteria

Inclusion criteria were: singleton pregnancies complicated by PPRM between 24+0 to 36+6 weeks of gestation and age of 18 years and more.

4.3 Exclusion criteria

Women were excluded from the study if they had signs of clinical chorioamnionitis, fetuses with an estimated weight <10th percentile, the presence of either congenital or

chromosomal fetal abnormalities, gestational or pre-gestational diabetes, gestational hypertension, preeclampsia, signs of fetal hypoxia, or significant vaginal bleeding.

4.4 Methods

4.4.1 Sample collection

4.4.1.1 Amniotic fluid sampling and assessment

4.4.1.1.1 Amniotic fluid sampling

Ultrasonography-guided transabdominal amniocentesis was carried out, and approximately 5 mL of amniotic fluid was aspirated and divided into three tubes. The first and second tubes with uncentrifuged amniotic fluid were transported immediately to the laboratory for DNA extraction, polymerase chain reaction for *Ureaplasma* species, *Mycoplasma hominis*, and *Chlamydia trachomatis* and 16S rRNA gene. Uncentrifuged amniotic fluid from the third tube was used to the assessment of IL-6 concentrations and after that was centrifuged for 15 minutes at 300 g immediately after collection in order to remove debris and cells, divided into aliquots and stored at -70°C until analysis.

4.4.1.1.2 Amniotic fluid analyses

4.4.1.1.2.1 Detection of *Ureaplasma* species, *Mycoplasma hominis*, and *Chlamydia trachomatis*

DNA was isolated from the amniotic fluid with a QIAamp DNA Mini Kit (QIAGEN, Hilden, Germany) according to the manufacturer's instructions (using the protocol for the isolation of bacterial DNA from biological fluids). Real-time PCR was performed with a Rotor-Gene 6000 instrument (QIAGEN, Hilden, Germany) with a commercial AmpliSens C. trachomatis/Ureaplasma/M. hominis-FRT kit (Federal State Institution of Science, Central Research Institute of Epidemiology, Moscow, Russia) to detect DNA from *Ureaplasma* species, *Mycoplasma hominis*, and *Chlamydia trachomatis* in a common PCR tube. The amount of *Ureaplasma* species DNA in copies/mL was determined using an absolute quantification technique that employs an external calibration curve. Plasmid DNA (pCR4, Invitrogen) was used to prepare the calibration curve. (38, 100)

4.4.1.1.2.2 Non-cultivation detection of other bacteria in amniotic fluid

Bacterial DNA was identified by PCR targeting the 16S rRNA gene with the following primers: 50-CCAGACTCCTACGGGAGGCAG-30 (V3 region) and 50-ACATTTCAACAC-GAGCTGACGA-30 (V6 region) (Greisen et al., 1994). Each individual reaction contained 3 mL of target DNA, 500 nM of forward and reverse primers and Q5 High Fidelity DNA polymerase (NEB, Hitchin, UK) in a total volume of 25 mL. The amplification was performed in a 2720 Thermal Cycler (Applied Biosystems, Foster City, CA). The products were visualized on an agarose gel. Positive reactions yielded products of 950 bp, which were subsequently analyzed by sequencing. The 16S PCR products were cleaned and used in sequencing PCR reactions utilizing the above primers and the BigDye Terminator kit, version 3.1 (Applied Biosystems, Foster City, CA). The bacteria were then typed using the sequences obtained in BLAST and SepsisTest™ BLAST.

4.4.1.1.2.3 IL-6 assessment in amniotic fluid

The IL-6 concentrations were assessed with a lateral flow immunoassay Milenia® QuickLine IL-6 using the Milenia POCScan Reader (Milenia Biotec, GmbH, Giessen, Germany). The measurement range was 50–10,000 pg/mL. The intra-assay and inter-assay variations were 12.1% and 15.5%, respectively(65).

4.4.1.2 Maternal serum sampling and assessment

Maternal blood was obtained through venipuncture of the cubital vein at admission. Maternal serum CRP was measured using an immunoturbidimetric analysis (Modular PP analyzer, Roche, Basel, Switzerland), with a method sensitivity of 0.3 mg/L. (101)

4.4.1.3 Gingival crevicular fluid sampling and assessment

Gingival crevicular fluid was collected using standard sterile paper strips during a dental examination (MM Absorbent Paper Points, Medin a.s., Czech Republic). The deepest gingival pocket in each individual woman was selected for sampling. The selected tooth was separated and cleaned by cotton rolls and dried by air. The strip was inserted into the pocket using sterile tweezers. The strip was left in situ for 30 seconds. The strips were subsequently placed into the tubes containing 0.5 mL of sterile phosphate-buffered saline. The tube with the strip was shaken at 1400 moves/minute for 20 minutes. Strip was then removed using sterile tweezers. Tube

was subsequently centrifuged $300 \times g$ for 15 minutes at room temperature. The supernatant was aliquoted and stored at -70°C until analyses.

Gingival crevicular fluid levels of the following proteins were analysed at the Statens Serum Institute (Department of Clinical Biochemistry and Immunology, Copenhagen, Denmark) using a multiple sandwich immunoassay based on flowmetric Meso-Scale technology. The gingival fluid levels of IL-1 β , IL-6, IL-8, IL-10, IL-12, IL-17, IL-18, adiponectin, brain-derived neurotrophic factor (BDNF), CRP, heat shock protein 70 (HSP 70), high mobility group box protein-1 (HMGB-1), granulocyte macrophage colony stimulating factor (GM-CSF), insulin-like growth factor-binding protein (IGFBP)-1, macrophage inhibitory protein (MIP) -1 α , MIP-1 β , MMP-8, MMP-9, monocyte chemotactic protein-1 (MCP-1), S100A8, T-cell-specific protein (RANTES), tumor necrosis factor (TNF)- α , TNF- β , tumor growth factor- β 1 (TGF- β 1), soluble TNF receptor-1 (sTNF-R1), and trombospondin-1 were assessed. The samples of gingival fluid were measured undiluted in duplicate. Inter- and intra-assay coefficients and detection levels are as follows: IL-1 β 21%, 9%, 0.26 pg/mL; IL-6 24%, 5%, 0.99 pg/mL; IL-8 9%, 9%, 0.25 pg/mL, IL-10 34%, 130, 0.25 pg/mL; IL-12 39%, 12%, 0.57 pg/mL; IL-17 22%, 12%, 1.47 pg/mL; IL-18 22%, 9%, 2.54 pg/mL; adiponectin 23%, 4%, 0.006 ng/mL; BDNF 25%, 15%, 31.4 pg/mL; CRP 37%, 8%, 0.0002 $\mu\text{g/mL}$; HSP 70 36%, 29%, 0.44 ng/mL; HMGB-1 25%, 7%, 5.67 ng/mL; GM-CSF 30%, 16%, 53.0 pg/mL; IGFBP-1 16%, 4%, 0.035ng/mL; MIP-1 α 19%, 5%, 1.2 pg/mL; MIP-1 β 38%, 3%, 25.7 pg/mL; MMP-8 33%, 3%, 0.005 ng/mL; MMP-9 27%, 6%, 0.004 pg/mL; MCP-1 36%, 12%, 3.84 pg/mL; S100A8 33%, 6%, 10.7 ng/mL; RANTES 26%, 9%, 0.4 pg/mL; TNF- α 18%, 6%, 8.4 pg/mL; TNF- β 14%, 10%, 0.26 pg/mL; TGF- β 1 45%, 5%, 2.03 ng/mL, sTNF-R1 14%, 10%, 9.36 pg/mL; and trombospondin-1 28%, 4%, 0.85 ng/mL. Only proteins with detectable amniotic fluid levels in more than 50% of the samples were included in the analyses (TNF- β and TGF- β 1 were excluded from analyses since their levels were detected just in 48% and 39%, respectively).

4.4.1.4 Placenta and fetal membranes sampling and assessment

After delivery, each placenta was collected and fixed in formalin, and tissue samples with placental membranes were inserted in paraffin. Tissue sections of placentas were stained with hematoxylin-eosin for histological examination. The degree of neutrophil

infiltration was assessed separately in the free fetal membranes (amnion and chorion-decidua), in the placenta (amnion and chorionic plate), and in the umbilical cord according to the criteria given by Salafia et al. (68).

4.4.2 Dental examination

Periodontal examination was performed within 72 hours of admission. A full-mouth recording was used to determine the periodontal and oral hygiene status. Probing pocket depth (PPD), defined as the distance from the marginal gingiva to the bottom of the pockets, and clinical attachment loss (CAL), defined as the distance from the cemento-enamel junction to the bottom of the pockets, were measured at four sites (medial, buccal, oral, and vestibular) on each fully erupted tooth. Third molars and retained roots were not included. Probing depth was measured by a calibrated periodontal probe PCPUNC156 (Hu-Friedy, Chicago, IL) with fine calibration with single millimeter grading. No other probes were used. A Decayed, Missing, and Filled (DMF) index was recorded for each subject. Gingival status was evaluated by gingival index (GI) as adapted from Silness and Løe and the plaque index (PLI) was evaluated according to Løe et al (102, 103).

4.4.3 Neonatal outcomes

Short-term neonatal outcome was evaluated on the presence of the following outcomes: need for tracheal intubation, respiratory distress syndrome (defined by the presence of two or more of the following criteria: evidence of respiratory compromise, a persistent oxygen requirement for >24 h, administration of exogenous surfactant and evidence of hyaline membrane disease on X-ray), intraventricular hemorrhage grade I-IV (diagnosed by cranial ultrasound examinations based on criteria defined by Papile et al.) (104), necrotizing enterocolitis at least grade IIA according to modified Bell's criteria (105), retinopathy of prematurity (identified by retinoscopy), early onset sepsis (EOS - defined as evidence of positive blood culture during the first 72 h of life and either the presence of clinical symptoms or strong clinical suspicions of sepsis [presence of symptoms and elevated CRP and/or affected white blood cell count], late-onset sepsis (defined as evidence of positive blood culture between 4-120 days of life and either the presence of clinical symptoms or strong clinical suspicions of sepsis [presence of symptoms and elevated CRP and/or

affected white blood cell count]), bronchopulmonary dysplasia (defined by the infant's oxygen requirement and/or ventilator support at 28 days of life and in the 36th postmenstrual week), pneumonia (diagnosed by abnormal findings on chest X-rays), or neonatal death, defined as death before hospital discharge(88).

4.4.4 Definitions

4.4.4.1 MIAC

MIAC was determined based on a positive result of the PCR analysis for *Ureaplasma* species, *Mycoplasma hominis* and *Chlamydia trachomatis* and/or positivity for the 16S rRNA gene with subsequent microorganism identification by sequencing.

4.4.4.2 IAI

Diagnosis of IAI was confirmed with an amniotic fluid IL-6 concentration of 745 pg/mL or higher, measured by a point of care test. Sterile IAI was defined as IAI without MIAC (65, 101).

4.4.4.3 HCA

Diagnosis of HCA was based on the presence of neutrophil infiltration in the chorion-decidua (Grades 3-4), the chorionic plate (Grades 3-4), the umbilical cord (Grades 1-4), and/or the amnion (Grades 1-4). The classification of histological chorioamnionitis is presented in Table 1. A single perinatal pathologist, who was blinded to the clinical status of the women, reviewed all of the histopathological samples

4.4.4.4 Periodontal disease

Severe periodontal disease was defined as having two or more sites with ≥ 6 mm CAL (not on the same tooth) or one or more site(s) with ≥ 5 mm PPD. Second, 'other' periodontal disease comprised two lesser levels of disease: moderate periodontal disease, defined as two or more sites with ≥ 4 mm clinical CAL (not on the same tooth) or two or more sites with PPD ≥ 5 mm, also not on the same tooth; and mild periodontal disease, defined as ≥ 2 sites with ≥ 3 mm CAL and ≥ 2 sites with ≥ 4 mm PPD (not on the same tooth), or 1 site with ≥ 5 mm PPD. Subjects not meeting either criterion were considered as having a healthy periodontium. This definition was adopted and simplified for four site probing from the CDC/AAP definition published

by Eke et al. (106). and recommended by the consensual statement published by Holtfreter et al. (107).

4.5 Statistical analyses

All statistical analyses were performed with SPSS 21.0 for Mac OS X (SPSS Inc, Chicago, IL, USA) and GraphPad Prism 6.0 for Mac OS X (GraphPad Software, La Jolla, CA, USA). The differences were considered statistically significant at $p < 0.05$. p -values were from two sided tests.

4.5.1 Specific aim I.

The demographic and clinical characteristics were compared using non-parametric Kruskal-Wallis test and data was presented as the median [interquartile range (IQR)] and using the Chi-square test and presented as number (%). The normality of the data was tested using the D'Agostino and Pearson omnibus normality test. Maternal serum CRP concentrations were compared with using non-parametric Kruskal-Wallis test and data was presented as the median [interquartile range (IQR)]. A Spearman partial correlation was used to adjust the data for gestational age at admission or at delivery. Spearman rank correlation test was used for analysis of correlation between CRP concentrations and the microbial load of *Ureaplasma* species.

4.5.2 Specific aim II.

The demographic and clinical characteristics were compared using non-parametric Kruskal-Wallis or Mann-Whitney tests, as appropriate, and data was presented as the median [interquartile range (IQR)] and using the Chi-square test or Fisher exact test, as appropriate, and presented as number (%). The normality of the data was tested using the D'Agostino and Pearson omnibus normality test. A Spearman partial correlation was used to adjust the data for gestational age at delivery.

4.5.3 Specific aim III.

Continuous variables were compared using the Mann-Whitney U test and were presented as medians [interquartile range (IQR)]. Categorical variables were compared using the Fisher's exact test, and were presented as numbers (%). The

normality of the data was tested using the D'Agostino and Pearson omnibus normality test.

4.5.4 Specific aim IV.

Continuous variables were compared using the Mann-Whitney *U* test and the Kruskal Wallis test, as appropriate, and were presented as medians [interquartile range (IQR)]. Categorical variables were compared using the chi-square test and were presented as numbers (%). The normality of the data was tested using the D'Agostino and Pearson omnibus normality test. Spearman's partial correlation was used to adjust the results for potential confounders (gestational age at sampling, administration of corticosteroids, administration of antibiotics and smoking). To identify an association between levels of proteins in gingival crevicular fluid and maternal serum CRP and amniotic fluid IL-6 concentrations, the Spearman correlations were used. Differences were considered statistically significant at $p < 0.05$. All *p* values were from two-sided tests, and all the statistical analyses were performed using SPSS 19.0 for Mac OS X (SPSS Inc., Chicago, IL).

5 Results

5.1 Specific aim I.

5.1.1 Clinical characteristics of study population.

Maternal and neonatal characteristics according to the presence and absence of MIAC and/or HCA are presented in Table 2. The presence of both MIAC and HCA was observed in 25% (97/386) of women. The presence of HCA alone and MIAC alone was found in 36% (141/386) and 8% (32/386) of women, respectively. The remaining 29% (116/386) of women had neither MIAC nor HCA. The women with both MIAC and HCA had the lowest gestational age at admission and at delivery, as well as the lowest birthweight. Latency from PPROM to delivery, gestational age at delivery and birthweight were not relevant clinical findings in the cohort of women due to the active management above 28 weeks of gestation (108).

5.1.2 Maternal serum CRP concentrations according to the presence or absence of MIAC and or HCA.

Women with both MIAC and HCA exhibited the highest concentration of CRP [median: 9.0 mg/L (IQR: 3.9-20.1)] vs. women with HCA alone [median: 6.4 mg/L (IQR: 3.6–12.1)] or MIAC alone [median: 4.3 mg/L (IQR: 3.6-15.6)] or with neither MIAC nor HCA [median: 4.5 mg/L (IQR: 2.0-8.6)]; (Figure 1) in crude analysis ($p < 0.0001$), as well as after adjustment for gestational age of sampling ($p < 0.0001$).

5.1.3 Maternal CRP concentrations and the bacterial load of *Ureaplasma* species

A positive correlation between the microbial burden of *Ureaplasma* species in the amniotic fluid and maternal CRP concentrations was found (Spearman rho = 0.33, $p = 0.002$; Figure 2).

5.2 Specific aim II.

5.2.1 Clinical characteristics of the study population

The maternal and neonatal characteristics are presented in Table 3 according to the presence and absence of MIAC and/or HCA. The presence of both MIAC and HCA was observed in 36% (45/122) of women. The presence of MIAC alone and HCA alone was found in 5% (6/122) and 34% (41/122) of women, respectively. The remaining 25% (30/122) of women had neither MIAC nor HCA.

5.2.2 Neonatal outcomes according to subgroups of PPRM.

The short-term neonatal outcome according to the presence or absence of MIAC and/or HCA is presented in Table 4. In the unadjusted analysis, a higher need to administrate surfactant and a higher incidence of EOS were observed in women with both MIAC and HCA. However, after adjusting the analysis for gestational age at delivery, only EOS remained significant ($p=0.001$). No other significant differences were observed for short-term neonatal outcomes among the subgroups.

5.3 Specific aim III.

5.3.1 Clinical and periodontal characteristics of study population

The demographic and clinical data, which are the same for the aims III and IV, are shown in Table 5. A total of 45% (35/78) women had periodontal disease. Mild, moderate, and severe periodontal disease was present in 19% (15/78), 19% (15/78), and 6% (5/78) of women, respectively. In women with mild periodontal disease, 67% (10/15) had MIAC and 67% (10/15) had IAI. In women with moderate periodontal disease, 26% (4/15) had MIAC and 33% (5/15) had IAI. In women with severe periodontal disease, 20% (1/5) had MIAC and 40% (2/5) had IAI. The median tooth count of the whole cohort was 28 teeth per mouth (IQR 27-28). The median gingival index value was 0.75 (0.39-1.72) and the median plaque index was 0.60 (0.23-1.15). The median DMF value was 10 (7-15) for the whole cohort.

5.3.2 Periodontal status according to the presence or absence of MIAC

The presence of MIAC was found in 28% (22/78) of women. The presence of *Ureaplasma* species was the most common cause of MIAC (n=16). In women with MIAC, other bacteria were found in the amniotic fluid: *Streptococcus intermedius* (n=2), *Sneathia sanguinegens* (n=1), *Fusobacterium nucleatum* (n=1), *Peptoniphilus* species (n=1), and one non-identifiable bacteria by sequencing (n=1). None of the women had a polymicrobial finding in the amniotic fluid. The presence of *Streptococcus intermedius* and *Fusobacterium nucleatum* was considered as positive for periopathogenic bacteria in amniotic fluid. There were no differences in the evaluated parameters of periodontal status in women with and without MIAC. The results are presented in Table 6.

5.3.3 Periodontal status according to the presence or absence of IAI

The presence of IAI was revealed in 26% (20/78) of women. Women with IAI did not have worse periodontal statuses compared to women without IAI. The results are shown in Table 7.

5.3.4 Periodontal status according to the presence of MIAC alone, IAI alone, and MIAC with IAI

Based on the presence of MIAC and or IAI, women were split in four subgroups: 1) with both MIAC and IAI (microbial-associated IAI; 15% [12/78]), 2) with IAI alone (sterile IAI; 10% [8/78]), with MIAC alone (colonization; 13% [10/78]), and without both MIAC and IAI (62% [48/78]). No associations between periodontal status and any particular subgroup of women were found. The results are presented in the Table 8.

5.4 Specific aim IV.

5.4.1 Association between the levels of gingival crevicular fluid proteins and maternal serum CRP and amniotic fluid IL-6 concentrations

No correlations between the levels of proteins in the gingival crevicular fluid and maternal serum CRP and amniotic fluid IL-6 concentrations were observed, except for a weak positive correlation between the GM-CSF and CRP (Spearman rho = 0.26, $p = 0.02$; Table 9).

5.4.2 The association between the levels of gingival crevicular fluid proteins and amniotic fluid IL-6 concentrations in the subgroup of women with sterile IAI

No correlations between levels of proteins in gingival crevicular fluid and amniotic fluid IL-6 concentrations were observed, except the correlation with BDNF (rho= 0.73; $p=0.05$; Table 10).

5.4.3 The presence of sterile IAI and levels of proteins in gingival crevicular fluid

Women with sterile IAI had lower levels of IL-1 β (with: median 10.7 pg/mL, range 1.4-597.7 vs. without: median 112.6 pg/mL, range 1.3-1437.0; $p=0.04$), IL-18 (with: median 10.3 pg/mL, range 3.8-248.7 vs. without: median 67.8 pg/mL, range 4.0-663.1; $p=0.04$), IGFBP-1 (with: median 0.09 ng/mL, range 0.01-1.1 vs. without: median 0.41 ng/mL, range 0.01-24.7; $p=0.03$), adiponectin (with: median 0.6 ng/mL, range 0-21.3 vs. without: median 4.4 ng/mL, range 0-35.9; $p=0.02$), and MMP-8 (with: median 4.4 ng/mL, range 0-106.3 vs. without: median 64.5 ng/mL, range 0.1-1267.3; $p=0.03$) in crude analysis. No differences were identified after the adjustment for gestational age of sampling, administration of corticosteroids and antibiotics and smoking (Table 11).

6 Discussion

6.1 Specific aim I

The main findings of the specific aim I are as follows: i) the presence MIAC and HCA was associated with the highest maternal inflammatory response measured by maternal serum CRP; ii) microbial load of *Ureaplasma* species in the amniotic fluid was related to the intensity of maternal inflammatory response.

The worst scenario in PPRM is the presence of MIAC and HCA (33, 62, 64). Previously was demonstrated, the presence of MIAC and HCA represented the condition with the highest inflammatory response in the amniotic cavity and fetal compartment (69, 70). However, the only method to diagnose MIAC is an invasive procedure, and histopathological information of HCA is not available antenatally. Therefore, the present study was focused on the prediction of MIAC and HCA using a non-invasive approach, which involved the determination of a classical marker of infection - maternal serum CRP.

Previous studies reported the influence of CRP in pregnancies complicated with HCA (109-111), but few studies focused on the relationship with MIAC (32). Howman et al. observed a significantly higher maternal inflammatory response when evaluating CRP in women with HCA (109). The study population also included women with clinical chorioamnionitis. Therefore, the differences in CRP could partially reflect the inclusion of women with severe and late stage infections, when the maternal inflammatory response is primarily activated (112). Contrary to Howman et al., Laar et al. found that the inclusion of women with clinical chorioamnionitis did not exhibit differences when the role of CRP was evaluated to detect HCA (110). Similarly, Martinez et al. conducted a review based on the evaluation of eight studies and did not observe data to support an influence of HCA on CRP (111). A lack of association was also observed when evaluating maternal serum CRP between PPRM women with or without MIAC (32).

The data obtained in the present study revealed differences in CRP levels between different infectious subgroups of women with PPRM. CRP concentrations are gestational age-dependent (113), and differences in gestational age at sampling were observed between groups. Therefore, CRP concentrations were adjusted for this

confounding factor. CRP remained significantly higher in the subgroup of women with MIAC and HCA after adjusting by gestational age at sampling. However, the predictive value of CRP to identify the worst infectious scenario was weak, even at early gestational ages. These findings revealed that assessment of CRP levels was a poor predictor of MIAC and HCA, which highlights the importance of considering the amniocentesis as the only accurate method to identify this infectious condition. Therefore, these findings are interesting and clinically relevant.

The previous studies clearly demonstrated that the intraamniotic and maternal inflammatory responses were depended on the microbial load of *Ureaplasma* species (37, 39, 40). The present study confirmed an association between maternal CRP and the amount of *Ureaplasma* species in the amniotic fluid.

The present study also has some limitations. We analyzed only the association between maternal CRP at the time of admission and HCA, but not the association between the trend of CRP during latency and HCA. Information on IL-6 levels in amniotic fluid was not available. Therefore, the evaluation of the influence of maternal CRP on the intra-amniotic inflammatory response could not be performed.

6.2 Specific aim II.

The main finding of the specific aim II was the fact that PPRM pregnancies before 34 weeks of gestation complicated by the presence of both MIAC and HCA are at increased risk of subsequent development of EOS.

There are few studies evaluating the influence of MIAC on short-term neonatal outcome in women with PPRM, and they report a negative impact on neonatal morbidity (114-116). In this regard, Shim et al. reported a lower 1-min Apgar score and a higher risk of respiratory distress syndrome, intraventricular hemorrhage and bronchopulmonary dysplasia in newborns from women with MIAC (115). A higher risk of EOS in newborns exposed to intraamniotic inflammation and MIAC was also observed by Buhimschi et al. (114). Finally, Cobo et al. found that the presence of specific proteomic biomarkers (neutrophil defensins and calgranulins A, C) were found to be independent predictors of MIAC and neonatal composite morbidity in women with PPRM (116).

Similar results have been reporting regarding the occurrence of HCA (117-119). In this regard, Rusell et al. observed higher rates of perinatal death and EOS in the first 48 hours when HCA was present (119). Moreover, Tsiartas et al. observed an association between HCA and the occurrence of EOS (118). Erdemir et al. observed higher rates of EOS, surfactant requirements, bronchopulmonary dysplasia and mortality when pregnancies exhibited HCA. None of these studies, except that of Erdemir et al., considered gestational age at delivery when evaluating short-term neonatal outcome. Erdemir et al. suggested that the effect of HCA on neonatal morbidity and mortality was more prominent than the effect of low gestational age alone (117).

The fact that in the current study was used active management allowed to explore clearly defined subgroups of women with PPROM according to the presence of MIAC and/or HCA. Thus, in the current study, a significantly higher risk of EOS when both MIAC and HCA were present in women with PPROM prior to 34 weeks was observed.

Similar results were reported in women with PPROM after 34 weeks of gestation; however, gestational age was not considered as a potential confounder factor (62). In the current study, not only in the crude analysis but also after adjusting the analysis for gestational age at delivery, the risk of EOS remained significantly higher in women with both MIAC and HCA. This highlights the importance of considering the presence of both MIAC and HCA as the worst-case scenario for the fetus. This infectious exposition increased the risk of EOS regardless of the gestational age at which newborns are born.

Finally, no significant differences in the short-term neonatal outcome were observed when compared women with HCA alone, MIAC alone or with women without either MIAC or HCA. Similarly, although in women with PTL, Combs et al. reported that women with MIAC alone with low inflammatory response in the amniotic cavity presented a similar short-term neonatal outcome compared to women without infection (41). Altogether, it seems that it is the infectious nature of the combined MIAC and HCA condition that is responsible for the increased risk of EONS and not the activation of the intra-amniotic inflammatory response given that there was no difference on the composite neonatal outcome.

The study has some limitations. The main limitation was that the number of women in the MIAC subgroup was small, which reduced the statistical power to compare outcomes between subgroups. Secondly, no information about blood cultures of neonates were not available, so the bacterial etiology of neonatal sepsis and its link to cultivation from the amniotic fluid could not be assessed. Finally, the influence of inflammation of either the amniotic cavity or the fetal compartment on the short-term neonatal outcome was evaluated.

Altogether, it seems crucial an antenatal prediction of those women with both MIAC and HCA. Whether MIAC and HCA are associated with the highest intra-amniotic inflammatory response (mediated by not only a multiplex approach (116) but also by bedside IL-6 (64) and whether the presence of MIAC and HCA is associated with a significant risk of EOS, for further clinical perspective, defining the best cut-off of IL-6 for prediction of the occurrence of EOS in women with MIAC and HCA should be given.

6.3 Specific aim III.

The principal findings of this specific aim are as follows: i) periodontal disease was found in 45% women with PPRM; ii) women with MIAC did not have a worse periodontal status than women without MIAC; iii) women with IAI did not have a worse periodontal status than women without IAI; iv) the presence MIAC and/or IAI was not associated with a different periodontal status in women with PPRM.

In this cohort of women with PPRM, the prevalence of periodontal disease reached 45%. Given that the prevalence of periodontal disease in women in PPRM has been rarely studied, observed results are unique. An epidemiological study from Italy comparing the periodontal status of healthy pregnant women and women with selected pregnancy pathologies included women with PPRM (74 out of 230 women with pregnancy pathologies). Overall, periodontal disease was diagnosed in 82% of women with pregnancy pathologies (120). Nevertheless, this study did not include specific details of periodontal health in the women with PPRM, so it remains unclear how many women with PPRM had periodontal disease (120). Another study from Croatia found periodontal disease in 62% of women with spontaneous preterm delivery. Even though women with PPRM were included, the rates of periodontal disease in women with PPRM were not provided (121). The only study specifying

periodontal health of women with PPROM is a study from Chile (122). However, only eight women with PPROM were included and therefore it is impossible to draw a conclusion about the prevalence of periodontal disease in PPROM from such a small cohort.

The mechanisms by which periodontal disease can affect pregnancy and trigger preterm delivery are still poorly understood. It has been hypothesized that cytokines produced by periodontal inflammation are released into systemic circulation and may affect the placenta and fetus, and may be a cause of sterile IAI. Alternatively, oral organisms can be disseminated into amniotic fluid, placental circulation, and the fetus itself, causing MIAC. The presence of MIAC may trigger an intra-amniotic inflammatory response and lead to preterm delivery. This hypothesis is supported by data from an animal model. Mice infected with *Porphyromonas gingivalis* (periopathogenic bacteria) had increased IL-6 levels and exhibited preterm delivery. Moreover, *Porphyromonas gingivalis* was found in the placental tissue in situ (123). A clear translation of these data to humans is a challenging issue. In this study, no associations between periodontal disease and MIAC were found. Given MIAC is a heterogeneous group containing different bacterial species having different microbial loads in amniotic fluid, it would be important to know whether some periopathogenic bacteria were found in women with PPROM. In this study, periopathogenic bacteria in amniotic fluid was found in 4% (3/78) of women (2x *Streptococcus intermedius* and 1x *Fusobacterium nucleatum*). In two out of these three women, the presence of periopathogenic bacteria in amniotic fluid led to the development of IAI. Interestingly, none of the women with periopathogenic bacteria in amniotic fluid had severe periodontal disease (one had none, one mild, one was moderate). The rate of amniotic fluid periopathogenic bacteria identified in this study appears to be lower than the rates presented in previous studies by Gauthier et al. (11% [3/27] of women with spontaneous preterm delivery) and by León et al. (31% [8/28] of women with spontaneous preterm delivery) (56, 122). However, the fact that the above-mentioned studies evaluated the presence of periopathogenic bacteria in cohorts of women containing both phenotypes of preterm delivery, as well as with gestational ages less than 34 weeks must be taken in the consideration. The latter is important information since the rate of MIAC is much higher in spontaneous preterm delivery prior to 34 weeks than beyond 34 weeks. In this cohort, women with gestational age of less than

34 weeks represented just 60% (47/78) of women. Moreover, the above-mentioned studies evaluated the presence of periopathogenic bacteria in amniotic fluid with genus-specific PCR. This specific approach is able to reveal lower amniotic microbial loads of bacteria than standard PCR evaluation of 16S rRNA followed by Sanger sequencing.

The increase in systemic proinflammatory cytokines produced by periodontal inflammation is possibly the likely event (98) that leads to the development of IAI. It has been shown that IL-6 levels tend to be higher in patients with periodontal disease in serum, as well as in gingival crevicular fluid (124). In this study, no differences in periodontal status between women with and without IAI were observed. Moreover, no differences were revealed when women were split into four subgroups based on the presence of MIAC and/or IAI in order to more precisely specify the background of IAI (microbial-associated vs. sterile). On the other hand, in this study results were in line with the recent study conducted by Soucy-Giguère et al., where no differences in amniotic fluid IL-6 and matrix metalloproteinase 8 concentrations were identified in women with and without periodontal disease (125). In addition, they observed no association between spontaneous preterm delivery and periodontal disease.

The other issue with comparing data from other cohorts previously published is a lack of standardization in periodontal disease reporting. There has been a secondary analysis published by Manau et al. comparing 14 individual periodontal disease definitions and more than 50 continuous measurements on 1296 pregnant women(126). The conclusion of this study was quite simple. Out of those 14 definitions applied on the same sample, 14 different prevalences of periodontal disease were reported, ranging from 14.5 to 70.8%. This fact clearly complicates the comparison of the data. The first large consensual statement for periodontal disease reporting was published in 2015 by Holtfreter et al. (107). This statement suggested the standardized use of periodontal disease reporting according to CDC/AAP definitions published previously by Eke et al. (106). Even though these recommendations were published after the initiation of this study, these definitions with negligible changes were adopted.

This study was subject to some limitations. One of the limitations of this study was that it was performed at a single institution, which prevented of larger sample size. The second limitation was, evaluation of periodontal status in women with

spontaneous preterm delivery with intact membranes was not performed, which prevented from making of the periodontal status comparison between these two phenotypes of spontaneous preterm delivery. Third, genus-specific PCRs for the evaluation of the presence of periopathogenic bacteria in the amniotic fluid was not used. This prevented of possibly identifying very low microbial burdens of periopathogenic bacteria in amniotic fluid.

6.4 Specific aim IV.

The main findings of the specific aim IV are as follows: i) no correlations between local inflammatory response in gingival crevicular fluid, characterized by levels of multiple proteins, and maternal and intra-amniotic inflammatory responses, measured by maternal serum and amniotic fluid IL-6 concentrations, respectively, were found; ii) no correlations between local inflammatory response in gingival crevicular fluid and intra-amniotic inflammatory response in women with sterile IAI was revealed; and iii) the presence of sterile IAI was not related to different levels of mediators in gingival crevicular fluid.

Gingival crevicular fluid represents complex fluid derived from a variety of sources, mainly from a maternal serum transudate and an inflammatory exudate (127-130). Therefore, gingival crevicular fluid in women with PPRM contains constituents from the host, from the bacteria colonizing subgingival and supragingival plaque, as well as from the cells and tissues of periodontium (127-130). A plethora of various protein markers, including cytokines, chemokines, neuropeptides, and enzymes, has been found in this fluid (127, 131). Dozens of them have been suggested to be potential markers of periodontal disease so far (131, 132). In addition, analysis of gingival crevicular fluid might be used to assess the process whether and how periodontal disease influences other specific systemic disease and vice versa (127, 133, 134).

In this study, the associations between local inflammatory response in gingival crevicular fluid, measured by levels of multiple proteins and maternal and intra-amniotic inflammatory response in women with PPRM, were evaluated. To assess the intensity of the local inflammatory response in gingival crevicular fluid, the Meso-Scale technology was employed.

Severity of periodontal disease has been proposed to be a main reason for the elevation of inflammatory mediator levels in gingival crevicular fluid. (96, 135, 136) These locally produced inflammatory mediators in gingival crevicular fluid are able enter systemic circulation and induce an acute-phase response in the liver leading to an elevation of maternal serum CRP concentrations. (137, 138) Given this information as basic, the correlation between proteins levels in gingival crevicular fluid and maternal serum CRP concentrations was evaluated. In this study, no association was observed. Likewise, no association between proteins levels in gingival crevicular fluid and amniotic fluid IL-6 concentrations was observed. These results confirm that intra-amniotic complications in PPRM, but not periodontal disease are responsible for systemic and intra-amniotic inflammatory response in women with PPRM. This is supported by the fact, that the subgroup of women with microbial-associated IAI, where the systemic (maternal serum CRP concentrations) and intra-amniotic (amniotic fluid IL-6 concentrations) inflammatory responses were the highest, did not have different levels of inflammatory mediators in gingival crevicular fluid than women in the others subgroups.

The pathophysiology of sterile IAI has not been fully explained yet (73, 139-141). Therefore, it is hypothesized that a low-grade systemic inflammation, which may occur due to high inflammatory mediator levels in gingival crevicular fluid, might led to the presence of sterile IAI. Nevertheless, this theory has not been proven in this study, since there was no association between gingival crevicular fluid levels and amniotic fluid IL-6 concentrations in women with sterile IAI. In addition the subgroup of women with sterile IAI did not have different levels than remaining women groups.

The information from this study might be a clinically relevant, because it clearly shown local inflammation in gingival crevicular fluid doesn't lead to the development of sterile IAI in women PPRM. Therefore the information from gingival crevicular fluid would not be helpful in the process of selection of women who are at high or at low risk of sterile IAI.

There are some limitations of this study. First, the samples of gingival crevicular fluid were not obtained at the same time when amniocentesis was performed (at the time of admission) but within 72 hours from admission. This approach did not allow the possibility to obtain the samples of gingival and crevicular fluid before the antibiotics

and corticosteroids administrations. Considering the antibiotics and corticosteroids administration as the potential confounders, the results were controlled for them. Second, this study is limited by the relatively small sample sizes of the subgroups (mainly the presence microbial associated IAI, sterile IAI, and colonization), which increases the potential for type II error.

7 Conclusion

The presence of both MIAC and HCA is associated with the highest maternal inflammatory response in women with PPROM. The microbial load of *Ureaplasma* species affects the intensity of maternal inflammatory response.

The presence of both MIAC and HCA increases the risk of early onset sepsis in pregnancies complicated by preterm prelabor rupture of membranes before 34 weeks of gestation.

The presence of MIAC and IAI is not related to the periodontal status of women with PPROM.

The local inflammatory response in the gingival crevicular fluid is not related to the maternal and intra-amniotic inflammatory responses in women with PPROM.

8 Appendices

Table 1. Grading system for acute chorioamnionitis, based on Salafia et al. (68).

| | Grade 1 | Grade 2 | Grade 3 | Grade 4 |
|------------------------|--|---|---|---|
| Chorion-decidua | One focus of at least 5 neutrophils | More than one grade-1 focus or at least one focus of 5-20 neutrophils | Multiple and/or confluent grade-2 foci | Diffuse and dense acute inflammation |
| Amnion | One focus of at least 5 neutrophils | More than one grade-1 focus or at least one focus of 5-20 neutrophils | Multiple and/or confluent grade-2 foci | Diffuse and dense acute inflammation |
| Chorionic plate | One focus of at least 5 neutrophils in subchorionic fibrin | Multiple grade-1 foci in subchorionic fibrin | Few neutrophils in connective tissue or subchorionic fibrin | Numerous neutrophils in chorionic plate, and chorionic vasculitis |
| Umbilical cord | Neutrophils in inner third of umbilical vein wall | Neutrophils in inner third of at least two umbilical vessel walls | Neutrophils in the perivascular Wharton's jelly | Panvasculitis and funisitis extending deep into Wharton's jelly |

Table 2. Maternal and neonatal characteristics in the different subgroups of PPRoM pregnancies from 24+4 to 36+6 weeks of gestation.

| | The presence of MIAc and HCA (n=97) | The presence of HCA alone (n=141) | The presence of MIAc alone (n=32) | The absence of MIAc and HCA (n=116) | p-value |
|--|-------------------------------------|-----------------------------------|-----------------------------------|-------------------------------------|---------|
| Maternal age | 31 (26-35) | 32 (28-35) | 31 (26-34) | 30 (27-34) | 0.13 |
| Primiparous | 33 (34%) | 69 (49%) | 17 (53%) | 64 (55%) | 0.02 |
| Prepregnancy BMI | 22.4 (20.1-25.1) | 23.4 (21.1-26.3) | 22.2 (19.6-24.2) | 21.8(19.9-24.9) | 0.01 |
| Smoking | 29 (30%) | 21 (15%) | 10 (31%) | 15 (12%) | 0.002 |
| Gestational age at admission (week+days) | 30+5 (27+6-33+6) | 32+4 (31+1-35+0) | 34+0 (33+0-35+6) | 33+1 (31+5-35+1) | <0.0001 |
| Gestational age at delivery (week+days) | 31+1 (28+1-33+6) | 32+6 (31+4-35+1) | 34+1 (33+0-35+6) | 33+2 (32+0-35+2) | <0.0001 |
| Latency from PPRoM to amniocentesis (hours) | 7 (4-12) | 6 (4-11) | 4 (3-6) | 6 (3-10) | 0.002 |
| Latency from PPRoM to delivery (hours) | 46 (24-83) | 40 (22-76) | 17 (9-51) | 31 (14-63) | 0.001 |
| CRP levels at admission (mg/L) | 9.0 (3.9-20.1) | 6.4 (3.5-9.9) | 4.3 (2.1-9.3) | 4.5 (2-8.5) | <0.0001 |
| WBC count at admission (x10 ⁹ /L) | 13.1 (11.0-15.5) | 11.7 (9.0-13.0) | 11.7 (10.1-14.8) | 12.0 (10.0-14.6) | 0.46 |
| Induction of labor | 42 (43%) | 37 (26%) | 14 (44%) | 29 (25%) | 0.006 |
| Vaginal delivery | 69 (71%) | 91 (65%) | 26 (81%) | 81 (70%) | 0.29 |
| Birth weight (grams) | 1720 (1155-2125) | 2080 (1640-2455) | 2235 (1838-2518) | 2280 (1748-2565) | <0.0001 |
| 5 min Apgar score < 7 | 10 (10%) | 8 (6%) | 0 (0%) | 3 (3%) | 0.06 |
| 10 min Apgar score < 7 | 6 (6%) | 3 (2%) | 0 (0%) | 2 (2%) | 0.13 |

Abbreviations:

PPROM: preterm prelabor rupture of membranes

MIAC: microbial invasion of amniotic cavity

HCA: histological chorioamnionitis

BMI: body mass index

CRP: C-reactive protein

WBC: white blood cell

Continuous variables were compared using a non-parametric Kruskal Wallis test and presented as medians (IQR). Categorical variables were compared using Chi-square test and presents as numbers (%). Statistically significant differences are marked in bold.

Table 3. Maternal and neonatal characteristics in the different subgroups of PPRM pregnancies prior to 34 weeks.

| | The presence of MIAC and HCA (n=45) | The presence of HCA alone (n=41) | The presence of MIAC alone (n=6) | The absence of MIAC and HCA (n=30) | P value |
|--|-------------------------------------|----------------------------------|----------------------------------|------------------------------------|--------------------|
| Maternal age | 31 (26-36) | 33 (28-38) | 31 (24.5-37.5) | 31 (28.5-33.5) | 0.58 |
| Primiparous | 14 (31%) | 20 (49%) | 3 (50%) | 20 (67%) | 0.001 |
| Prepregnancy BMI | 21.3 (19.4-23.2) | 24.9 (21.4-28.4) | 20.7 (18.1-23.3) | 21.7 (19.4-24.0) | 0.003 |
| Smoking | 14 (31%) | 6 (15%) | 2 (33%) | 3 (10%) | 0.05 |
| Gestational age at admission (week+days) | 28+3 (25+3-31+3) | 31+3 (28+5-34+1) | 32+4 (31+1-34+0) | 31+2 (29+2-33+2) | <0.00001 |
| Gestational age at delivery (week+days) | 28+6 (26+1-31+4) | 31+6 (29+2-34+3) | 33+0 (31+5-34+2) | 31+6 (29+6-33+6) | <0.00001 |
| Latency from PPRM to amniocentesis (hours) | 8 (4-12) | 9 (4.5-13.5) | 4 (2-6) | 7 (3.5-10.5) | 0.48 |
| Latency from PPRM to delivery (hours) | 76 (45-87) | 71 (49-93) | 56 (20-92) | 61 (16-106) | 0.11 |
| Vaginal delivery | 31 (69%) | 22 (54%) | 4 (67%) | 16 (53%) | 0.23 |
| Birth weight (grams) | 1310 (912-1708) | 1770 (1331-2209) | 1980 (1663-2297) | 1780 (1331-2229) | <0.00001 |
| 5 min Apgar score < 7 | 9 (20%) | 3 (7%) | 0 (0%) | 1 (3%) | 0.02 |
| 10 min Apgar score < 7 | 6 (13%) | 1 (2%) | 0 (0%) | 0 (0%) | 0.02 |
| Perinatal death | 6 (13%) | 3 (7%) | 0 (0%) | 1 (3%) | 0.11 |

Abbreviations:

PPROM: Preterm prelabor rupture of membranes

MIAC: Microbial invasion of amniotic cavity

HCA: Histological chorioamnionitis

CRP: C-reactive protein

WBC: White blood cells

Continuous variables were compared using a non-parametric Kruskal Wallis test and presented as medians (IQR). Categorical variables were compared using Chi-square test and presented as numbers (%). Statistically significant differences are marked in bold.

Table 4. Short-term neonatal morbidity in the different subgroups of pregnancies complicated by preterm prelabor rupture of membranes prior to 34 weeks.

| | The presence of MIAC and HCA (n=45) | The presence of HCA alone (n=41) | The presence of MIAC alone (n=6) | The absence of MIAC and HCA (n=30) | P value | P value* |
|--|-------------------------------------|----------------------------------|----------------------------------|------------------------------------|---------------|--------------|
| Respiratory distress syndrome | 24 (53%) | 20 (49%) | 1 (17%) | 12 (40%) | 0.17 | 0.94 |
| Administration of surfactant | 16 (36%) | 9 (22%) | 1 (17%) | 3 (10%) | 0.01 | 0.26 |
| Early onset sepsis | 15 (33%) | 5 (12%) | 0 (0%) | 0 (0%) | 0.0007 | 0.001 |
| Late onset sepsis | 5 (11%) | 1 (2%) | 0 (0%) | 1 (3%) | 0.16 | 0.66 |
| Intraventricular hemorrhage grade III-IV | 2 (4%) | 0 (0%) | 0 (0%) | 1 (3%) | 0.81 | 0.74 |
| Bronchopulmonary dysplasia | 1 (2%) | 1 (2%) | 0 (0%) | 0 (0%) | 0.41 | 0.80 |
| Necrotizing enterocolitis | 1 (2%) | 0 (0%) | 0 (0%) | 1 (3%) | 0.69 | 0.59 |
| Pneumothorax | 1 (2%) | 0 (0%) | 0 (0%) | 1 (3%) | 0.69 | 0.27 |
| Retinopathy of prematurity | 1 (2%) | 0 (0%) | 0 (0%) | 1 (3%) | 0.69 | 0.41 |
| Severe neonatal morbidity | 28 (62%) | 21 (51%) | 1 (17%) | 13 (43%) | 0.78 | 0.18 |

Variables were compared using Chi-square test and presented as numbers (%).
Statistically significant differences are marked in bold.

p-value* - adjusted for gestational age at delivery

Table 5. Maternal and neonatal characteristics of PPRoM pregnancies according to the presence or absence of MIAC and/or IAI

| | The presence of MIAC (n=22) | The absence of MIAC (n=56) | p-value ¹ | The presence of IAI (n=20) | The absence of IAI (n=58) | p-value ² |
|--|-----------------------------|----------------------------|----------------------|----------------------------|---------------------------|----------------------|
| Maternal age | 31 (28-35) | 32 (29-36) | 0.35 | 31 (29-31) | 33 (29-36) | 0.08 |
| Primiparous | 13 (59%) | 25 (45%) | 0.32 | 11 (55%) | 27 (47%) | 0.61 |
| Prepregnancy body mass index | 23.5 (20.2-27.0) | 24.0 (21.0-27.3) | 0.42 | 26.0 (21.4-29.3) | 23.4 (20.4-26.7) | 0.20 |
| Gestational age at admission (weeks+days) | 33+0 (29+3-34+3) | 33+3 (30+3-35+2) | 0.48 | 29+5 (25+6-32+6) | 33+6 (31+4-35+3) | <0.0001 |
| Gestational age at delivery (weeks+days) | 33+3 (29+5-34+5) | 33+6 (30+5-36+2) | 0.36 | 30+1 (27+2-33+1) | 34+0 (32+2-35+5) | <0.0001 |
| Latency from PPRoM to amniocentesis (hours) | 5 (3-8) | 5 (3-10) | 0.89 | 4 (4-9) | 5 (3-7) | 0.76 |
| Latency from amniocentesis to delivery (hours) | 45 (23-121) | 62 (19-113) | 0.64 | 103 (27-168) | 50 (17-101) | 0.05 |
| CRP levels at admission (mg/L) | 9.8 (13.8-17.4) | 4.9 (2.3-8.6) | 0.02 | 9.8 (3.7-38.2) | 4.9 (2.4-8.9) | 0.01 |
| WBC count at admission (x10 ⁹ /L) | 13.6 (10.4-17.6) | 11.8 (9.8-14.7) | 0.13 | 13.5 (12.0-17.9) | 11.7 (9.7-14.3) | 0.03 |
| Amniotic fluid IL-6 at admission (pg/mL) | 1058 (229-10000) | 264 (148-450) | 0.001 | 2649 (1188-10000) | 225 (132-348) | <0.0001 |
| Administration of corticosteroids | 18 (82%) | 47 (84%) | 1.00 | 18 (90%) | 47 (81%) | 0.50 |
| Administration of antibiotics | 22 (100%) | 54 (96%) | 1.00 | 20 (100%) | 56 (97%) | 1.00 |

| | | | | | | |
|------------------------------|------------------|------------------|------|-----------------|------------------|-------------------|
| Spontaneous vaginal delivery | 18 (82%) | 35 (63%) | 0.12 | 16 (80%) | 37 (64%) | 0.27 |
| Cesarean section | 4 (18%) | 20 (36%) | 0.27 | 4 (20%) | 20 (35%) | 0.67 |
| Forceps delivery | 0 (0%) | 1 (1%) | 0.28 | 0 (0%) | 1 (1%) | 1.00 |
| Birth weight (grams) | 2015 (1323-2525) | 1970 (1483-2465) | 0.89 | 1335 (900-2015) | 2170 (1750-2573) | <0.0001 |
| 5 min Apgar score < 7 | 2 (9%) | 2 (4%) | 0.32 | 2 (10%) | 2 (3%) | 0.27 |
| 10 min Apgar score < 7 | 0 (0%) | 2 (4%) | 1.00 | 1 (5%) | 1 (2%) | 0.45 |

Abbreviations:

PPROM: Preterm prelabor rupture of membranes

MIAC: Microbial invasion of amniotic cavity

IAI: Intra-amniotic inflammation

CRP: C-reactive protein

WBC: White blood cells

IL: Interleukin

Continuous variables were compared using a nonparametric Mann-Whitney *U* test. Categorical variables were compared using Fisher's exact test. Continuous variables are presented as median (IQR) and categorical as number (%). Statistically significant results are marked in bold.

p-value¹ – a comparison between women with and without MIAC, *p*-value² – a comparison between women with and without IAI.

Table 6. Periodontal status of PPRM pregnancies according to the presence or absence of MIAC.

| Characteristics | The presence of MIAC (n=22) | The absence of MIAC (n=56) | <i>p</i> -value |
|---|-----------------------------|----------------------------|-----------------|
| Never smokers | 11 (50%) | 33 (59%) | 0.64 |
| Former smokers | 7 (32%) | 12 (21%) | 0.50 |
| Current smokers | 4 (18%) | 11 (20%) | 0.58 |
| Tooth count* | 28 (27-28) | 28 (27-28) | 0.23 |
| Gingival Index | 0.9 (0.4-2.1) | 0.7(0.4-1.5) | 0.45 |
| Plaque Index | 0.9 (0.3-1.3) | 0.5 (0.2-1.0) | 0.14 |
| DMF* | 12 (8-14) | 10 (6-15) | 0.18 |
| No periodontal disease | 10 (45%) | 33 (59%) | 0.45 |
| Mild periodontal disease | 4 (18%) | 11 (19%) | 0.86 |
| Moderate periodontal disease | 7 (32%) | 8 (14%) | 0.15 |
| Severe periodontal disease | 1 (5%) | 4 (7%) | 0.93 |
| Mean CAL (mm) | 2.5 (2.3-2.8) | 2.2 (1.9-2.7) | 0.45 |
| Mean PPD (mm) | 2.5 (2.3-2.9) | 2.2 (1.9-2.7) | 0.17 |
| Proportion of sites/mouth CAL \geq 3 mm | 45.5 (27.7-56.7) | 27.4 (16.5-53.0) | 0.18 |
| Proportion of sites/mouth CAL \geq 4 mm | 11.4 (4.9-20.5) | 7.7 (1.8-17.4) | 0.31 |
| Proportion of sites/mouth CAL \geq 5 mm | 1.0 (0.0-5.8) | 0.9 (0.0-3.0) | 0.60 |
| Proportion of teeth/mouth CAL \geq 3 mm | 57.1 (29.8-68.0) | 25.6 (17.0-63.3) | 0.16 |
| Proportion of teeth/mouth CAL \geq 4 mm | 21.8 (8.1-35.9) | 12.7 (0.0-33.9) | 0.23 |
| Proportion of teeth/mouth CAL \geq 5 mm | 3.6 (0.0-12.5) | 0.0 (0.0-7.2) | 0.23 |
| Proportion of sites/mouth PPD \geq 4 mm | 45.5 (26.9-58.5) | 25.0 (14.2-56.6) | 0.39 |
| Proportion of sites/mouth PPD \geq 5 mm | 11.4 (4.1-22.5) | 7.3 (1.8-18.8) | 0.38 |
| Proportion of sites/mouth PPD \geq 6 mm | 1.0 (0.0-5.8) | 0.9 (0.0-3.2) | 0.54 |
| Proportion of teeth/mouth PPD \geq 4 mm | 14.8 (7.1-30.6) | 7.7 (0.0-22.3) | 0.29 |

| | | | |
|---|----------------|----------------|------|
| Proportion of teeth/mouth PPD ≥ 5 mm | 3.7 (0.0-17.1) | 3.6 (0.0-11.1) | 0.40 |
| Proportion of teeth/mouth PPD ≥ 6 mm | 0.0 (0.0-2.7) | 0.0 (0.0-0.0) | 0.79 |

* without 3rd molars

Abbreviations:

PPROM: preterm prelabor rupture of membranes

MIAC: microbial invasion of the amniotic cavity

IQR: interquartile range

DMF: decayed, missing, filled index

CAL: clinical attachment loss

PPD: probing pocket depth

Continuous variables were compared using a Mann-Whitney *U* test. Categorical variables were compared using Fisher's exact test. Continuous variables are presented as median (IQR) and categorical as number (%).

Table 7. Periodontal status of PPRM pregnancies according to the presence or absence of IAI.

| Characteristics | The presence of IAI (n=20) | The absence of IAI (n=58) | <i>p</i> -value |
|---|-------------------------------|------------------------------|-----------------|
| Never smokers | 10 (50%) | 34 (59%) | 0.68 |
| Former smokers | 7 (35%) | 16 (28%) | 0.73 |
| Current smokers | 3 (15%) | 8 (13%) | 0.81 |
| Tooth count* | 28 (27-28) | 28 (27-28) | 0.42 |
| Gingival Index | 0.8 (0.4-1.8) | 0.7 (0.4-1.6) | 0.89 |
| Plaque Index | 0.8 (0.3-1.7) | 0.6 (0.2-1.1) | 0.23 |
| DMF* | 9 (6-15) | 11 (7-15) | 0.70 |
| No periodontal disease | 10 (50%) | 33 (57%) | 0.78 |
| Mild periodontal disease | 5 (25%) | 10 (17%) | 0.67 |
| Moderate periodontal disease | 3 (15%) | 12 (20%) | 0.82 |
| Severe periodontal disease | 2 (10%) | 3 (5%) | 0.82 |
| Mean CAL (mm) | 2.4 (2.0-3.1) | 2.2 (2.0-2.7) | 0.45 |
| Mean PPD (mm) | 2.4 (0.0-3.1) | 2.2 (1.9-2.8) | 0.42 |
| Proportion of sites/mouth CAL \geq 3 mm | 39.1 (22.0-75.4) | 29.1 (18.0-53.3) | 0.43 |
| Proportion of sites/mouth CAL \geq 4 mm | 11.6 (4.3-27.0) | 7.7 (2.7-14.4) | 0.43 |
| Proportion of sites/mouth CAL \geq 5 mm | 0.9 (0.0-3.2) | 0.9 (0.0-4.3) | 1.00 |
| Proportion of teeth/mouth CAL \geq 3 mm | 45.4 (27.6-90.2) | 27.4 (17.9-64.1) | 0.24 |
| Proportion of teeth/mouth CAL \geq 4 mm | 23.6 (7.2-50.9) | 12.7 (3.6-31.5) | 0.17 |
| Proportion of teeth/mouth CAL \geq 5 mm | 1.8 (0.0-14.3) | 0.0 (0.0-7.1) | 0.51 |
| Proportion of sites/mouth PPD \geq 4 mm | 37.3 (22.0-75.2) | 26.4 (14.5-57.8) | 0.39 |
| Proportion of sites/mouth PPD \geq 5 mm | 10.9 (3.6-27.0) | 7.3 (1.8-19.4) | 0.45 |
| Proportion of sites/mouth PPD \geq 6 mm | 0.9 (0.0-3.2) | 0.9 (0.0-4.6) | 0.99 |
| Proportion of teeth/mouth PPD \geq 4 mm | 12.5 (7.1-33.9) | 7.7 (0.0-21.1) | 0.24 |
| Proportion of teeth/mouth PPD \geq 5 mm | 3.6 (0.0-10.7) | 3.6 (0.0-13.7) | 0.82 |
| Proportion of teeth/mouth PPD \geq 6 mm | 0.0 (0.0-3.6) | 0.0 (0.0-0.0) | 0.60 |

* without 3rd molars

Abbreviations:

PPROM: preterm prelabor rupture of membranes IAI: intra-amniotic inflammation

IQR: interquartile range DMF: decayed, missing, filled index

CAL: clinical attachment loss PPD: probing pocket depth.

Continuous variables were compared using a Mann-Whitney U test. Categorical variables were compared using Fisher's exact test. Continuous variables are presented

Table 8. Periodontal status of PPRM pregnancies according to the presence or absence of MIAC and/or IAI.

| Characteristics | The presence of MIAC and IAI (n=12) | The presence of IAI alone (n=8) | The presence of MIAC alone (n=10) | The absence of MIAC and IAI (n=48) | p-value |
|------------------------------|-------------------------------------|---------------------------------|-----------------------------------|------------------------------------|---------|
| Never smokers | 7 (58%) | 3 (38%) | 4 (40%) | 30 (63%) | 0.39 |
| Former smokers | 2 (17%) | 1 (13%) | 5 (50%) | 11 (23%) | 0.20 |
| Current smokers | 3 (25%) | 4 (50%) | 0 (0%) | 7 (15%) | 0.18 |
| Tooth count* | 28 (27-28) | 28 (28-28) | 27 (25-28) | 28 (27-28) | 0.10 |
| Gingival Index | 0.9 (0.3-2.1) | 0.7 (0.5-1.3) | 0.9 (0.7-2.3) | 0.7 (0.4-1.5) | 0.79 |
| Plaque Index | 0.6 (0.3-1.2) | 0.9 (0.7-1.8) | 1.1 (0.7-1.3) | 0.4 (0.1-1.0) | 0.05 |
| DMF* | 12 (8-17) | 6 (5-7) | 12 (8-14) | 11 (7-15) | 0.11 |
| No periodontal disease | 7 (58.3) | 3 (37.5) | 3 (30.0) | 30 (62.5) | 0.40 |
| Mild periodontal disease | 2 (16.7) | 3 (37.5) | 2 (20.0) | 8 (16.7) | 0.40 |
| Moderate periodontal disease | 2 (16.7) | 1 (12.5) | 5 (50.0) | 7 (14.6) | 0.63 |
| Severe periodontal disease | 1 (8.3) | 1 (12.5) | 0 (0.0) | 3 (6.25) | 0.68 |
| Mean CAL (mm) | 2.3 (1.9-2.8) | 2.5 (2.2-3.1) | 2.6 (2.4-2.8) | 2.2 (1.9-2.6) | 0.09 |
| Mean PPD (mm) | 2.3 (1.8-2.8) | 2.4 (2.2-3.1) | 2.6 (2.4-2.9) | 2.1 (1.8-2.6) | 0.10 |

| | | | | | |
|---|------------------|------------------|------------------|------------------|------|
| Proportion of sites/mouth CAL ≥ 3 mm (%) | 34.6 (17.0-60.3) | 43.6 (34.6-76.8) | 51.8 (41.7-56.2) | 25.9 (15.2-52.2) | 0.11 |
| Proportion of sites/mouth CAL ≥ 4 mm | 6.4 (3.9-17.8) | 14.1 (10.3-31.7) | 16.2 (7.9-20.5) | 7.1 (1.8-15.5) | 0.20 |
| Proportion of sites/mouth CAL ≥ 5 mm | 0.4 (0.0-3.1) | 1.3 (0.7-4.7) | 1.9 (0.9-5.8) | 0.9 (0.0-3.0) | 0.45 |
| Proportion of teeth/mouth CAL ≥ 3 mm | 38.4 (16.6-70.5) | 55.4 (34.2-93.8) | 62.2 (50.4-68.0) | 22.2 (14.3-61.3) | 0.05 |
| Proportion of teeth/mouth CAL ≥ 4 mm | 16.7 (6.4-48.2) | 35.7 (15.2-53.6) | 23.2 (13.0-33.4) | 10.7 (0.0-26.8) | 0.20 |
| Proportion of teeth/mouth CAL ≥ 5 mm | 0.0 (0.0-14.3) | 3.6 (0.0-11.6) | 3.7 (0.9-7.1) | 0.0 (0.0-4.6) | 0.33 |
| Proportion of sites/mouth PPD ≥ 4 mm | 33.6 (16.6-60.3) | 42.9 (32.9-76.1) | 54.5 (41.0-58.5) | 21.8 (13.2-52.4) | 0.1 |
| Proportion of sites/mouth PPD ≥ 5 mm | 6.4 (3.2-17.8) | 13.8 (9.2-31.7) | 16.2 (7.2-22.5) | 6.2 (1.6-15.9) | 0.29 |
| Proportion of sites/mouth PPD ≥ 6 mm | 0.4 (0.0-3.1) | 1.3 (0.7-4.7) | 3.2 (0.9-5.8) | 0.4 (0.0-3.2) | 0.41 |
| Proportion of teeth/mouth PPD ≥ 4 mm | 7.7 (5.4-26.5) | 21.8 (9.8) | 17.9 (9.4) | 5.4 (0.0-20.4) | 0.19 |
| Proportion of teeth/mouth PPD ≥ 5 mm | 3.6 (0.0-10.7) | 3.6 (2.7-15.2) | 9.0 (3.6-17.1) | 1.8 (0.0-11.1) | 0.46 |
| Proportion of teeth/mouth PPD ≥ 6 mm | 0.0 (0.0-3.6) | 0.0 (0.0-1.8) | 0.0 (0.0-0.3) | 0.0 (0.0-0.0) | 0.91 |

* without 3rd molars

Abbreviations:

PPROM: preterm prelabor rupture of membranes

MIAC: microbial invasion of range

DMF: decayed, missing, filled index

CAL: clinical attachment loss

PPD: probing pocket depth

Continuous variables were compared using a Kruskal–Wallis test. Categorical variables were compared using Chi-square test. Continuous variables are presented as median (IQR) and categorical as number (%).

Table 9. The correlations between levels of proteins in gingival crevicular fluid and maternal serum C-reactive protein and amniotic fluid interleukin-6 and concentrations.

| | maternal serum C-reactive protein | | amniotic fluid interleukin -6 | |
|---------------------|-----------------------------------|-----------------|-------------------------------|-----------------|
| | rho | <i>p</i> -value | rho | <i>p</i> -value |
| IL-1b (pg/mL) | -0.04 | 0.73 | 0.04 | 0.76 |
| IL-6 (pg/mL) | 0.01 | 0.93 | 0.02 | 0.85 |
| IL-8 (pg/mL) | -0.07 | 0.56 | -0.12 | 0.29 |
| IL-10 (pg/mL) | 0.13 | 0.25 | 0.06 | 0.63 |
| IL-12 (pg/mL) | 0.09 | 0.44 | 0.11 | 0.33 |
| IL-17 (pg/mL) | -0.09 | 0.45 | -0.07 | 0.54 |
| IL-18 (pg/mL) | -0.0007 | 0.99 | -0.18 | 0.76 |
| GM-CSF (pg/mL) | 0.27 | 0.02 | 0.16 | 0.16 |
| IGFBP-1 (ng/mL) | 0.14 | 0.22 | -0.05 | 0.67 |
| Adiponectin (ng/mL) | -0.11 | 0.36 | -0.18 | 0.12 |
| BDNF (pg/mL) | -0.04 | 0.76 | 0.05 | 0.66 |
| RANTES (pg/mL) | -0.11 | 0.32 | -0.13 | 0.25 |
| CRP (µg/mL) | 0.04 | 0.70 | -0.02 | 0.85 |
| HMGB-1 (ng/mL) | 0.02 | 0.87 | 0.02 | 0.87 |
| HSP70 (ng/mL) | -0.10 | 0.37 | -0.04 | 0.76 |
| S100A8 (ng/mL) | 0.09 | 0.45 | 0.05 | 0.68 |
| MCP-1 (pg/mL) | 0.18 | 0.12 | 0.05 | 0.64 |
| MIP-1α (pg/mL) | -0.18 | 0.11 | -0.15 | 0.20 |
| MIP-1b (pg/mL) | 0.05 | 0.63 | 0.06 | 0.58 |
| MMP-8 (ng/mL) | -0.06 | 0.61 | -0.15 | 0.89 |
| MMP-9 (ng/mL) | -0.11 | 0.35 | -0.08 | 0.50 |
| TNFα (pg/mL) | 0.02 | 0.87 | 0.03 | 0.79 |

| | | | | |
|-----------------------|-------|------|-------|------|
| TNF-R1 (ng/mL) | -0.13 | 0.24 | -0.04 | 0.75 |
| Trombospondin (ng/mL) | -0.04 | 0.74 | -0.02 | 0.84 |

The variables were compared using Spearman correlation. Statistically significant result is marked in bold.

Table 10. The correlations between levels of proteins in gingival crevicular fluid and maternal serum C-reactive protein and amniotic fluid interleukin-6 and concentrations in the subgroup of women with sterile IAI.

| | amniotic fluid interleukin -6 | |
|---------------------|-------------------------------|-----------------|
| | rho | <i>p</i> -value |
| IL-1b (pg/mL) | 0.19 | 0.66 |
| IL-6 (pg/mL) | 0.21 | 0.62 |
| IL-8 (pg/mL) | 0.19 | 0.66 |
| IL-10 (pg/mL) | 0.47 | 0.24 |
| IL-12 (pg/mL) | -0.10 | 0.84 |
| IL-17 (pg/mL) | 0.21 | 0.62 |
| IL-18 (pg/mL) | 0.19 | 0.66 |
| GM-CSF (pg/mL) | -0.29 | 0.50 |
| IGFBP-1 (ng/mL) | 0.26 | 0.54 |
| Adiponectin (ng/mL) | 0.32 | 0.43 |
| BDNF (pg/mL) | 0.73 | 0.05 |
| RANTES (pg/mL) | 0.40 | 0.46 |
| CRP (µg/mL) | -0.11 | 0.79 |
| HMGB-1 (ng/mL) | 0.21 | 0.62 |
| HSP70 (ng/mL) | 0.45 | 0.27 |
| S100A8 (ng/mL) | 0.07 | 0.88 |
| MCP-1 (pg/mL) | -0.07 | 0.87 |
| MIP-1α (pg/mL) | 0.12 | 0.79 |
| MIP-1b (pg/mL) | 0.50 | 0.22 |
| MMP-8 (ng/mL) | 0.18 | 0.67 |

| | | |
|-----------------------|------|------|
| MMP-9 (ng/mL) | 0.31 | 0.46 |
| TNF α (pg/mL) | 0.21 | 0.62 |
| TNF-R1 (ng/mL) | 0.06 | 0.90 |
| Trombospondin (ng/mL) | 0.61 | 0.12 |

Abbreviations:

IAI: Intra-amniotic inflammation

The variables were compared using Spearman correlation. Statistically significant result is marked in bold.

Table 11. Gingival crevicular fluid levels of selected proteins in women with PPRM with respect to the presence of sterile IAI.

| | Women with sterile IAI (n=8) | The other women (n=70) | <i>p</i> -value | <i>p</i> -value [#] |
|--------------------------|---------------------------------|---------------------------|-----------------|------------------------------|
| IL-1 β (pg/mL) | 10.7 (1.4-597.7) | 112.6 (1.3-1437.0) | 0.04 | 0.19 |
| IL-6 (pg/mL) | 17.8 (3.4-88.0) | 34.9 (2.4-694.8) | 0.07 | 0.27 |
| IL-8 (pg/mL) | 56.7 (9.6-387.2) | 176.5 (9.2-1586.3) | 0.13 | 0.13 |
| IL-10 (pg/mL) | 1.0 (0.3-1.4) | 0.90 (0.3-2.4) | 0.55 | 0.71 |
| IL-12 (pg/mL) | 19.8 (10.1-25.6) | 20.6 (3.5-46.8) | 0.62 | 0.68 |
| IL-17 (pg/mL) | 10.2 (5.1-36.8) | 17.2 (2.7-44.2) | 0.26 | 0.35 |
| IL-18 (pg/mL) | 10.3 (3.8-248.7) | 67.8 (4.0-663.1) | 0.04 | 0.58 |
| GM-CSF (pg/mL) | 164.7 (76.4-304.6) | 128.8 (53.0-382.6) | 0.66 | 0.93 |
| IGFBP-1 (ng/mL) | 0.09 (0.01-1.1) | 0.41 (0.01-24.7) | 0.03 | 0.65 |
| Adiponectin (ng/mL) | 0.6 (0-21.3) | 4.4 (0-35.9) | 0.02 | 0.09 |
| BDNF (pg/mL) | 143.1 (31.4-220.3) | 163.6 (31.4-1090.9) | 0.16 | 0.19 |
| RANTES (pg/mL) | 66.0 (38.8-1379.9) | 90.8 (0.4-10000.0) | 0.32 | 0.70 |
| CRP (μ g/mL) | 0.0002 (0.0002-0.01) | 0.002 (0.0002-12.6) | 0.07 | 0.44 |
| HMGB-1 (ng/mL) | 96.8 (17.2-642.3) | 242.8 (6.1-8448.6) | 0.08 | 0.41 |
| HSP70 (ng/mL) | 4.1 (0.8-9.7) | 6.1 (0.4-129.8) | 0.25 | 0.58 |
| S100A8 (ng/mL) | 574.9 (349.1-1740.2) | 501.4 (72.2-5469.3) | 0.38 | 0.94 |
| MCP-1 (pg/mL) | 4.8 (3.8-15.6) | 10.9 (3.8-37.6) | 0.14 | 0.25 |
| MIP-1 α (pg/mL) | 10.0 (1.2-44.9) | 24.8 (1.2-788.1) | 0.14 | 0.50 |
| MIP-1b (pg/mL) | 112.8 (50.7-207.6) | 126.4 (53.3-561.6) | 0.31 | 0.39 |
| MMP-8 (ng/mL) | 4.4 (0-106.3) | 64.5 (0.1-1267.3) | 0.03 | 0.10 |
| MMP-9 (ng/mL) | 13.2 (0.1-83.1) | 37.1 (0-473.6) | 0.08 | 0.21 |
| TNF α (pg/mL) | 90.2 (17.8-357.3) | 165.1 (20.4-2318.0) | 0.13 | 0.25 |
| TNF-R1 (ng/mL) | 0.06 (0.02-0.35) | 0.17 (0.02-1.5) | 0.07 | 0.06 |
| Trombospondin (ng/mL) | 1.6 (0.9-210.6) | 11.7 (0.9-1000.0) | 0.09 | 0.55 |

Abbreviations:

PPROM – preterm prelabor rupture of membranes

MIAC – microbial invasion of the amniotic cavity

IAI – intra-amniotic inflammation

Continuous variables were compared using a nonparametric Mann-Whitney *U* test and presented as median (range). Statistically significant differences are marked in bold.

p-value[#] - adjusted for gestation age of sampling, administration of corticosteroids, administration of antibiotics and smoking

Figure 1. Maternal CRP concentrations according to subgroups of women with PPROM.

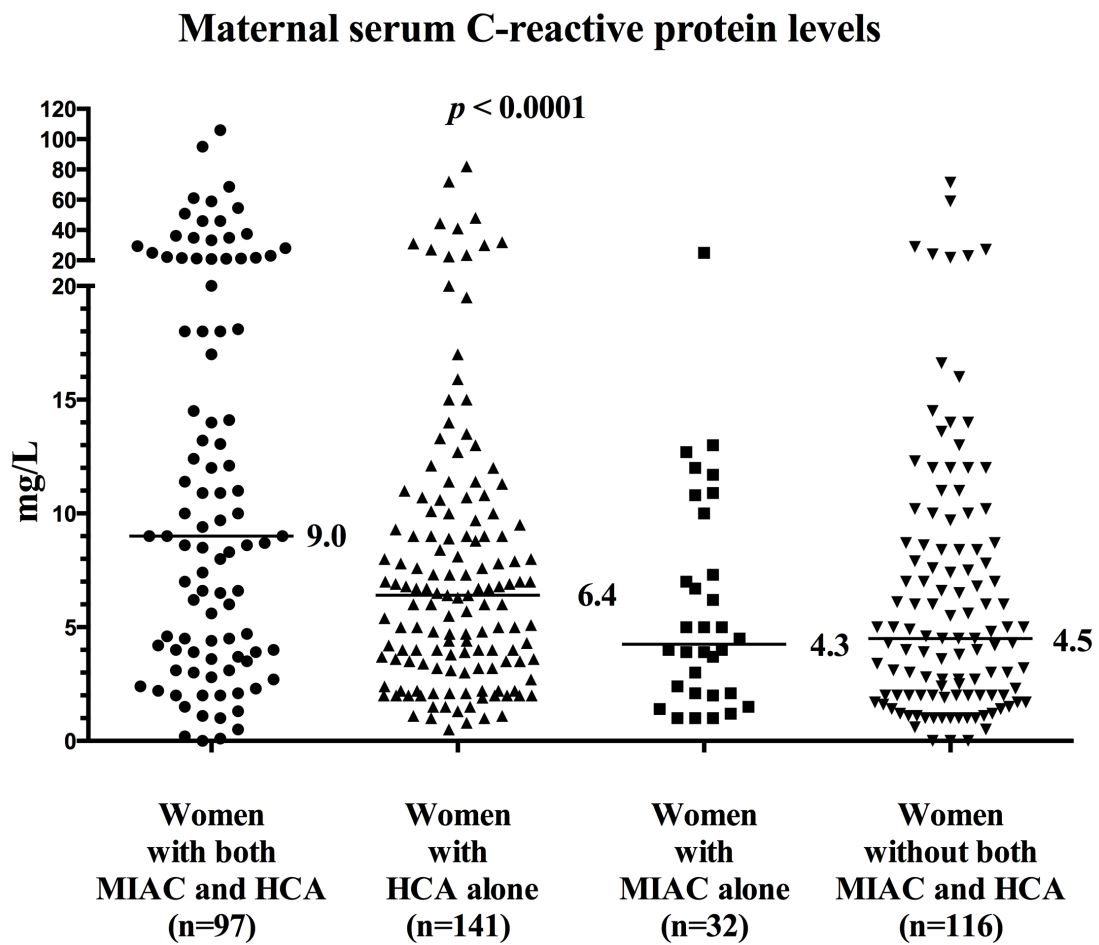
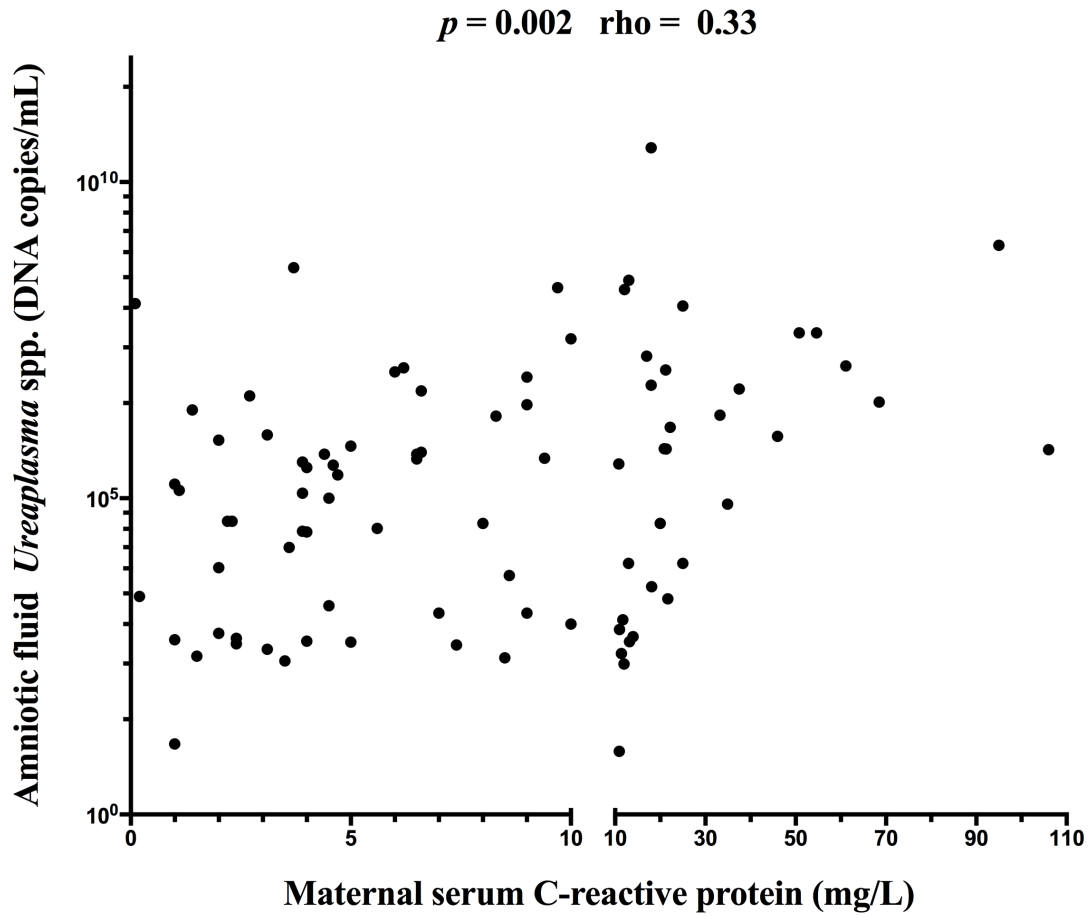


Figure 2. Correlation between microbial burden of *Ureaplasma species* in the amniotic fluid (copies DNA/mL) and maternal CRP concentrations.



9 Literature review

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10 Supplements