Interleukin-1 system in the human fallopian tube - no spatial but a temporal regulation of mRNA and protein expression

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Running title: Interleukin-1 system in human fallopian tube

5 **Interleukin-1 system in the human fallopian tube - no spatial but**
a temporal regulation of mRNA and protein expression

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Abstract

The human fallopian tube provides the environment for the first 5 days of embryonic development *in vivo*. The IL-1 system is involved in human embryo implantation.

25 This study aimed to investigate IL-1 β , IL-1ra and IL-1R t1 expression within the length of the human fallopian tube on mRNA- and protein-level in samples from proliferative versus secretory phase, postmenopause samples (PMP) and samples from intra- (IUP) and extrauterine (EUP) pregnancies to examine possible spatial and hormonal induced changes (fimbrial, ampullary and isthmic tube segments). On

30 mRNA-level, IL-1 β was expressed in all samples except in PMP. IL-1R t1 could be detected in all samples whereas IL-1ra was only expressed in secretory phase and the IUP sample. Immunohistochemically we could detect IL-1 β and IL-1R t1 protein in all proliferative and secretory phase samples with maximum intensity in secretory phase samples whereas IL-1ra was expressed in secretory phase samples only.

35 Overall no spatial but temporal differences possibly due to hormonal changes could be observed suggesting a precise regulation of the IL-1 system, especially for IL-1ra and moreover a stable molecular architecture within the full length of the fallopian tube.

40 **Keywords:** Interleukin-1 (IL-1) system, human fallopian tube, Interleukin-1 receptor type I (IL-1R tI), Interleukin-1 receptor antagonist (IL-1ra), implantation, spatial and temporal expression pattern

1. Introduction

45 The factors involved in the complex embryonic maturation and in the embryo-maternal dialogue resulting in eutopic embryonic implantation are poorly understood. The human fallopian tube is the place of fertilization and also provides the natural environment for the first five days of embryonic development (Hess et al., 2007b). It is, as well, the most common place for ectopic implantation. The preimplantation
50 embryo produces several factors during its development to signal its presence to the maternal organism. Appropriate interaction between the preimplantation embryo and the maternal system is at least partly controlled by paracrine cytokines (Hess et al., 2007a; Popovici et al., 2006). Knowledge of expression patterns of these cytokines could possibly help to gain a better insight of the physiological course of
55 implantation. Although multiple studies have examined the existence and influence of various cytokines at the embryonic-maternal interface (Hess et al. 2007a; Popovici et al., 2006), as yet, little is known about cytokine production by the human fallopian tube (Brabec and Hill, 2000).

 The Interleukin-1-system is composed of two agonists, Interleukin-1 α (IL-1 α)
60 and Interleukin-1 β (IL-1 β), one antagonist, the Interleukin-1 receptor antagonist (IL-1ra) and two membrane-bound receptors, Interleukin-1 receptor type I (IL-1R tI) and II (IL-1R tII) (Dinarello et al., 1994; Dimitriadis et al., 2005). All components of the IL-1 family in humans are located on chromosome 2 and the DNA-, RNA- and protein-structures are all well characterized for many species. Both agonists are
65 initially synthesized as precursor proteins of 31 kDa. The mature proteins have a molecular weight of 17 kDa and although the amino acid sequences have a similarity

of only ~22 %, they induce the same biological responses (Dower et al., 1986). There is also a high similarity between the cDNA-sequences of IL-1 α and - β in mice and human (March et al., 1985). Interleukin-1 receptors type I and II both possess a
 70 transmembraneous domain and their extracellular portions are homologous with similar binding affinities for the agonists and antagonist. There is also a soluble form of the IL-1R tII. The IL-1 receptor type I which is found in low numbers on almost all cell surfaces whereas IL-1R tII is found primarily on white blood cells. Only the binding of either IL-1 α or - β to the IL-1 receptor type I results in signal transduction
 75 (Sims et al., 1993) with receptor type II and the soluble IL-1 receptor acting as competitors of the receptor type I (Colotta et al., 1993). The IL-1 receptor antagonist binds with a high affinity to both receptors and prevents signal transduction by IL-1 α and - β (Hannum et al., 1990).

The IL-1 system is intimately involved in embryonic implantation. In humans, the IL-
 80 1R tI has been detected in total human endometrium (Simon et al., 1993b) and, more specifically, in endometrial epithelial cells with a maximum mRNA- and protein - expression during the luteal phase (Simon et al., 1993a) - the time of embryonic attachment and implantation. IL-1 β -mRNA was detected in secretory human endometrium beginning on day 23 of the menstrual cycle (Kauma et al., 1990). In the
 85 last years, all major components of the IL-1-system, namely IL-1 β , IL-1ra and IL-1R tI were detected immunohistochemically in single preimplantation embryos (De Los Santos et al., 1996). *In vitro* fertilized, cultured human embryos have been shown to produce both IL-1 α and IL-1 β . High concentrations [>60 pg/ml and >80 pg/ml] of these cytokines in culture media have been correlated with successful implantation

90 after intrauterine transfer of these embryos (Sheth et al., 1991), although other authors could not detect IL-1 α or - β in culture fluids of human embryos (Seifer et al., 1993). In mice, IL-1 α and IL-1 β have been detected and localized at mRNA- and protein-level in endometrial endothelial cells (Tackacs et al., 1988) in increasing levels from day 3 of pregnancy peaking between days 4 and 5 (De et al., 1993). Blastocyst
95 implantation is known to occur late on day 4. It was shown that systemically administered recombinant human IL-1ra given intraperitoneally from day 3 to day 6 of pregnancy inhibited embryonic implantation in mice (Simon et al., 1994a). It was also demonstrated that *in vitro* cultured single mouse embryos at various stages of preimplantation development express different patterns of mRNA for IL-1 β , IL-1ra
100 and IL-1R tI (Simon et al., 1994b). In summary, there is an increasing body of evidence that the IL-1 system is involved in early embryonic development and implantation. The aim of our study was to detect a potentially spatial and temporal IL-1 β -, IL-1ra- and IL-1R tI-expression pattern on mRNA- and protein-level in human fallopian tubes and to describe possible
105 hormonally regulated changes since these changes might influence the time and place of embryonic implantation.

2. Material and Methods

2.1 Patients

110 Patients who underwent hysterectomy or tubal surgery for reasons other than
cancer (except one patient with intrauterine pregnancy diagnosed of cervical cancer)
were asked to participate in this study. Each participating patient signed an informed
consent that was prior approved by the human investigations committee of the
Heinrich-Heine University medical school. Menstrual phase was determined by the
115 patient's history and dating was verified by histological examination of the
endometrium according to Noyes et al. (1950) and serum-levels of estradiol,
progesterone, LH and FSH were measured. The indications for surgery were as
follows: hysterectomy for benign reasons: 19 patients; ectopic tubal pregnancies: 3
patients and one patient received a radical hysterectomy in the 9th week of gestation
120 due to cervical cancer. The mean age of the patients was 39 ± 7.9 yr, range 26 - 51 yr.
Tissue samples from the fimbrial, ampullary and isthmic parts of the fallopian tubes
were collected from both sides. Tissue samples were then split in half: one half was
fixed in 4 % paraformaldehyde and later processed for immunohistochemistry, the
other half was snap-frozen in liquid nitrogen and stored at -80 °C until RNA-
125 extraction.

2.2 RNA isolation

The isolation of RNA from the tissue samples was done as described
previously (Chomczynski and Sacchi, 1987) with the RNA-STAT-60 reagent (Tel-
130 Test "B" Inc., Friedenswood, TX, USA). Briefly, tissue samples were washed three

times in phosphate buffered saline (PBS, GibCo BRL, Grand Island, NY, USA) to remove blood contamination. 100 mg tissue was homogenized in 1 ml of RNA-STAT-60 reagent. Total RNA was separated from DNA and proteins by adding chloroform and precipitated using isopropanol. Precipitate was washed two times in
 135 75 % ethanol, air dried and resuspended in diethylpyrocarbonate- (DEPC) treated dH₂O. The amount and purity of the isolated RNA was quantitated by spectrophotometry in a GenQuant RNA/DNA calculator (Pharmacia Biotech Ltd., Cambridge, UK).

140 2.3 Primers for Reverse Transcription (RT) and Polymerase Chain Reaction (PCR)

Sequences of cDNA-clones for the mRNAs that should be detected in human fallopian tubes (β -actin, IL-1 β , IL-1ra and IL-1R tI) were obtained from the GenBank Database of the National Center for Biotechnology Information (NCBI) of the National Institutes of Health (NIH, Internet address:
 145 <http://www.ncbi.nlm.nih.gov/sites/entrez>). One set of primer-sequences was found with the help of the program OLIGO 4.1 Primer Analysis Software (National Bioscience, Plymouth, MN, USA) and synthesized by MWG Biotech (MWG Biotech AG, Ebersberg, Germany). To ensure that the product detected resulted from amplification of cDNA rather than contaminating genomic DNA, primers were
 150 designed to cross intron/exon boundaries. The human β -actin primers that were used to amplify an internal standard were obtained from Clontech Laboratories Inc., Palo Alto, CA, USA. The primer-sequences, locations on the cDNA and the sizes of the amplified fragments are listed in Table 1.

155 2.4 Reverse transcription (RT)

For each mRNA to be detected, 20 μ l RT-MasterMix were prepared (5 mM MgCl_2 , 1 $\times$ RT Buffer, 1 mM dNTPs (each), 2.5 μ M Oligo d(T)₁₆, 1 U/ μ l RNase Inhibitor, 2.5 U/ μ l MuLV Reverse Transcriptase [all Applied Biosystems, Foster City, CA, USA], and 1 μ g total RNA, DEPC-treated H₂O ad 20 μ l) and filled into a
 160 0.5 ml thin wall PCR-tube (Applied Scientific, South San Francisco, CA, USA). RT-MasterMix in PCR-tubes was covered with 50 μ l of light white mineral oil (Sigma, St. Louis, MO, USA) and kept on ice until the RT.

The RT-reaction was carried out in the DNA Thermal Cycler 480 by using a program with the following parameters: 42 $^{\circ}\text{C}$, 15 min; 99 $^{\circ}\text{C}$, 5 min; 4 $^{\circ}\text{C}$, ∞ . After the
 165 reaction was complete, samples were stored at -20 $^{\circ}\text{C}$ until the PCR.

2.5 Polymerase Chain Reaction (PCR)

2.5 μ l RT-products were mixed with 77.5 μ l PCR-Mastermix containing 1.25 mM MgCl_2 , 1 $\times$ PCR-Buffer, 0.125 μ M 3' and 5' primer-mix for either IL-1 β , IL-1ra,
 170 IL-1R tI or β -actin, 1 mM dNTPs (each), 1.5 U/ μ l AmpliTaq[®] DNA Polymerase, DEPC-treated dH₂O ad 80 μ l (Applied Biosystems) and the reaction-mix was covered with 50 μ l light white mineral oil, put in the DNA Thermal Cycler 480 and heated to 95 $^{\circ}\text{C}$ for 5 min to activate the enzyme. After completion of 30 cycles of 94
 175 $^{\circ}\text{C}$ for 45 sec, 54 $^{\circ}\text{C}$ for 45 sec and 72 $^{\circ}\text{C}$ for 60 sec, the reaction was terminated at 72 $^{\circ}\text{C}$ for 7 min and cooled down to 4 $^{\circ}\text{C}$. PCR-products were stored at -20 $^{\circ}\text{C}$ until 2 % agarose-gel electrophoresis was carried out in the presence of ethidiumbromide. After

completion of electrophoresis, the agarose-gel was analyzed on the GelDoc 1000 system (Bio-Rad Laboratories, Hercules, CA, USA). DNA-size calculation was carried out by using the Molecular Analyst Software (Bio-Rad Laboratories).

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2.6 Immunohistochemical staining procedure

The immunostaining procedure was performed on fallopian tubes sections by the avidin-biotin-peroxidase (ABC) method. Serial sections from each sample were mounted on slides, fixed in formalin and embedded in paraffin. The first section was processed for hematoxylin-eosin staining.

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Sections were deparaffinized in xylene, rehydrated through graded alcohols, and washed in phosphate-buffered saline (PBS; Sigma). Endogenous peroxidases were blocked with 1% H₂O₂ in 96% methanol, and non-specific binding was blocked with non-fat milk 5% in PBS at room temperature. After washing with 0.05% Tween[®]20 in PBS (PBS-T; Sigma), the slides were incubated at room temperature for 30min with the primary antibodies: monoclonal mouse anti-human IL-1 β antibody, monoclonal mouse anti-human IL-1R tI antibody, and polyclonal rabbit anti-human IL-1ra (all from Genzyme Corp., Cambridge, MA, USA) at 20 μ g/ml, 50 μ g/ml, and 20 μ g/ml respectively. Control incubations included deletion of the primary antibody.

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After rinsing with PBS-T, sections were incubated with a secondary antibody at room-temperature for 30min: anti-rabbit IgG (whole molecule) biotin conjugate (200 μ g/mL; Sigma) and anti-mouse IgG biotin conjugate (Vector Laboratories INC, Burlingame, CA, USA) at 150 μ g/ml. Histological sections were incubated with Vectastain ABC Kit (Vector Laboratories) reagents for 60 min at room temperature.

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200 Immunoreaction products were visualized by incubating sections with the substrate
solution in 0.1 M Tris-HCl pH 8.2 buffer blocking alkaline phosphatase activity with
levamisole. Slides were counterstained with 25 % of hematoxylin, then cleared, cover
slips applied and finally examined by light microscopy. Positive staining by the
primary antibody was indicated by a pink precipitate. The relative intensity of the
205 immunostaining was evaluated by two of the authors in a double-blind manner as
absent (0), weakly positive (+), moderate (++) or intense (+++) for biopsy specimens
from each portion and each phase of the menstrual cycle.

210 3. Results

3.1 RT-PCR of spatial expression pattern throughout the menstrual cycle

All 19 collected oviduct samples from the proliferative (n=8) and secretory (n=9) phase of the cycle and postmenopausal (n=2) women were divided into fimbrial, ampullary and isthmic parts and investigated separately by RT-PCR. From 215 the total of 19 samples all were positive for β -actin mRNA expression and therefore considered for further investigation. The following investigation of IL-1 β , IL-1ra and IL-1R tI expression showed that the presence or absence of the respective gene was constant regarding the spatial distribution. This shows that the gene expression of IL-1 β , IL-1ra and IL-1R tI within the length of the fallopian tube has no spatial difference 220 (Fig. 1).

3.2 RT-PCR of temporal expression pattern throughout the menstrual cycle

All 23 collected oviduct samples from the proliferative (n=8) and secretory (n=9) phase of the cycle and from postmenopausal (n=2) women and extra- (n=3) and 225 intrauterine (n=1) pregnancies were separately investigated by RT-PCR for temporal expression patterns of IL-1 β , IL-1ra and IL-1R tI. From the total of 23 samples, all were positive for β -actin mRNA expression and therefore considered for further investigation. The following investigation of IL-1 β , IL-1ra and IL-1R tI expression showed that there was a temporal and therefore most likely hormonal regulation of 230 two of the target mRNAs. IL-1R tI was expressed in all samples regardless the hormonal situation. IL-1ra mRNA could be detected in samples of the secretory phase and in intrauterine pregnancy, whereas IL-1 β could be detected in proliferative and

secretory phase fallopian tubes and in samples of intra- and extra-uterine pregnancy. This shows that the gene expression of IL-1 β , IL-1ra and IL-1R tI is temporally regulated within the human cycle (Fig. 1 and 2). Additionally IL-1 β is expressed not only in all different hormonal situations investigated but also in oviducts from intra- and extrauterine pregnancies.

3.3 Immunohistochemistry

The protein expression of IL-1 β , IL-1ra and IL-1R tI of proliferative phase versus secretory phase samples was evaluated using immunohistochemistry. Figure 3 A-H shows representative samples from fallopian tubes of the secretory phase whereas Fig. 3 I-P shows representative samples from proliferative phase fallopian tubes. Panel A-G and I-O are magnified 200 $\times$, panel B-H and J-P 400 $\times$. Panels A and B represent the control of secretory phase oviducts, panel I and J represent the negative control (omission of first antibody) of proliferative phase oviducts. We could show a strong expression of IL-1 β foremost in the luminal epithelium and also in the small vessels endothelium of secretory phase oviducts (Fig. 3 C, D), whereas the staining for IL-1 β in the luminal epithelium of proliferative phase oviducts was less intense and only a weak staining was localized in the small vessels endothelium (Fig. 3 K, L). The IL-1R tI was shown to be expressed in the luminal epithelium, small vessels endothelium and glandular epithelium of secretory phase samples (Fig. 3 E, F). In comparison the proliferative sample showed only a sporadic staining for the IL-1R tI in the luminal epithelium and no staining either in the vessel endothelium or in the glandular epithelium (Fig. 3 M, N). Fig. 3 G and H show an intense staining of IL-

1ra in the luminal epithelium and the small vessel endothelium of secretory phase oviducts whereas a staining for IL-1ra in proliferative oviducts was completely absent revealing a strict temporal expression of IL-1ra.

No staining was observed in the absence of either primary or secondary antibodies.

260 These protein data support the previous mRNA data showing an expression of IL-1 β in fallopian tubes of both cycle phases with an increase in the secretory phase as well as a pronounced expression of IL-1R tI in secretory phase oviducts. However, IL-1ra expression was seen in secretory phase samples only.

265 4. Discussion

Factors of the embryo-maternal communication that have an influence on maturation and transport of an embryo as it transverses the oviducts are crucial for a successful implantation (Hess et al., 2007a). The fallopian tube on the maternal side plays an important role in providing the place for fertilization as well as an active
270 component for transport and early development of the embryo. Within the oviduct, remarkable changes occur in dependence to the sex steroids estradiol in the proliferative phase and progesterone in the post-ovulatory secretory phase indicating a hormonal cyclicity of the oviduct comparable to the endometrium (Gardner et al., 1996). A growing body of evidence suggests that chemokines of the IL-1 family play
275 an important role at the feto-maternal interface (Krussel et al., 2003). Therefore, the knowledge of normal expression patterns of the IL-1 family members could help in understanding the physiological processes leading to proper maturation and transport of the blastocyst within the fallopian tube and subsequently to a successful intrauterine implantation. Since we have shown before that the human embryo
280 produces IL-1 β when cultured *in vitro* (Krussel et al., 1998), it is reasonable that it also produces IL-1 β during its passage through the fallopian tube. Consequently, it is very likely that the fallopian tube as the early communication partner for the developing embryo itself expresses components of the IL-1 family. Therefore, our objective was to first investigate the spatial expression pattern of IL-1 β , IL-1ra and IL-
285 1R tI in the human fallopian tube and second to elucidate a possible hormonal influence. Furthermore we wanted to explore whether we could find a hint for an

abnormal expression pattern or an imbalance of gene expression contributing to a risk of an extrauterine tubal pregnancy.

All collected oviduct samples from the proliferative and secretory phase of the cycle
290 as well as the postmenopausal samples were divided into fimbrial, ampullary and
isthmic parts before investigation and prepared separately. This led us to the
observation that regardless of the original spatial belonging of the oviduct part, the
mRNA expression of IL-1 β , IL-1ra and IL-1R tI showed no differences at all. These
findings on the mRNA level were also proven on the protein level by
295 immunohistochemistry (data not shown). The data support the notion that although
the oviduct is compromised of a changing architecture within its full length, the
molecular base of gene expression remains the same in all parts of the fallopian tube.
Therefore, possible differences in gene expression regarding to the biopsy technique
are erased which makes studies involving different parts of the oviduct comparable.
300 Furthermore, our aim was to elucidate whether the sex steroids estrogen and
progesterone have an influence on the temporal expression of IL-1 β , IL-1ra and IL-1R
tI during the reproductive phase. Therefore, fallopian tubes from the proliferative and
secretory phase of the menstrual cycle as well as oviducts from postmenopausal
women were investigated. This showed that IL-1 β was only present in samples from
305 the proliferative and secretory phase but in none of the postmenopausal oviducts
suggesting a regulation by sex steroids and a role in the process of human
reproduction. These data highlight that the oviduct and the uterus underlie partially
common most likely hormonally driven regulatory expression phenotypes since a
study has shown that IL-1 β is expressed as well by epithelial and stromal

310 endometrium cells throughout the cycle (Simon et al., 1993a). In unison with our immunohistochemistry results they detected an increase of IL-1 β immunostaining from proliferative to secretory endometrium suggesting an additional role in supporting the implantation within the endometrium during the window of implantation. On the contrary, another study primarily focused on the expression of
315 IL-1 α by endometrial stromal cells reports a hormonally controlled expression of IL-1 α but not for IL-1 β or IL-1ra (Pretto et al., 2008). Furthermore, it was shown that incubation of endometrial stromal cells with IL-1 β upregulates the expression of IL-1 β and IL-1ra (Huang et al., 2001) suggesting that besides a hormonal regulation chemokines seem to influence the expression of IL-1 family members in endometrial
320 stromal cells which could be likewise for the tube. In addition an immunohistochemical approach revealed IL-1 β staining in the villous cytotrophoblast, syncytiotrophoblast and maternal decidual cells, suggesting an autocrine/paracrine role during the human implantation (Simon et al., 1994b). In contrast to the ligand, we detected mRNA expression of its receptor IL-1R tI in all
325 samples investigated. The immunostaining showed a less intense staining in proliferative phase samples compared to secretory phase oviducts suggesting at least a moderate influence of the sex hormones. Furthermore it provides new evidence that the preimplantation embryo is able to communicate with the maternal side already during its transit through the fallopian tube by its own IL-1 β secretion in consent to a
330 study by Strakova et al. (2005) simulating endometrial changes of an early pregnancy in baboons by *in vivo* infusion of hCG and IL-1 β .

In contrast, IL-1ra mRNA was only detected in progesterone dominated secretory phase samples and in the intrauterine pregnancy sample suggesting a hormonal influenced regulation and further a possible protection mechanism to avoid a false
335 implantation of the embryo within the fallopian tube. The IL-1ra expression seems to be a rather systemic effect independent of the embryos presence since it was equally expressed in both sides of the fallopian tubes of each patient in the secretory phase. IL-1ra's possible protective effect becomes even more likely in the context that IL-1ra was not only present in all secretory phase samples and the one investigated
340 sample with an intrauterine pregnancy but did not occur in anyone of the 3 tested fallopian tubes with an extrauterine pregnancy. This hypothesis is furthermore supported by findings of other groups describing a decrease in IL-1ra expression in the human endometrium at the time of implantation to possibly facilitate the embryos invasion into the maternal endometrium (Simon et al., 1995). Another study showed
345 increased levels of IL-1ra in endometrial stromal cells of the secretory phase (Fukuda et al., 1995). Those differences though might be explained by the fact that the first study investigated the mRNA and protein expression for secreted IL-1ra and intracellular IL-1ra whereas the second study did not differentiate between those two forms. The findings of the first study positively correlate with a study in mice where
350 systemically administered IL-1ra prevented embryonic implantation (Simon, 1994a). However, this distinct inhibition of implantation might have been a mouse strain specific effect as a study by Abbondanzo et al. (1996) disproved the latter finding in IL-1R tI female knockout mice showing no deficiency in fertility besides a slightly smaller litter size as well as Cohen and Pollard could not show noticeable fertility

355 abnormalities in male IL-1R tI knockout mice (1998). On the other hand a recent
knock out study with male^{IL-1ra/-} mice showed a reduced capacity to fertilize oocytes
of wild type females *in vitro* and a reduced litter size when mated with wild type
females (Genaïem et al., 2009). The discrepancy of those studies might have been
evolved due to the fact that Abbondanzo et al. e.g. have used an outbred strain with a
360 mutated gene which are known to possibly have an influenced phenotype. Otherwise
it might be conceivable that subsequent events in the signaling cascade after blocking
a receptor with the receptor antagonist compared to a knock out of the receptor itself
may vary and possibly lead to inconsistent results as shown above when data of IL-
1R tI knock out studies are compared with IL-1ra knock out studies. With regards to
365 the event of extrauterine tubal pregnancy in human it could be possible that not only
the absence of IL-1ra but furthermore an inappropriate ratio of IL-1 β and IL-1ra
favoring IL-1 β may serve as one underlying molecular mechanism.

So far most investigations regarding the molecular base of extrauterine pregnancies
were conducted in the mouse or baboon model. Therefore little is known in the
370 human system. Huang et al. (2005) investigated fallopian tubes from extrauterine
pregnancies versus normal healthy control tubes from tubal ligations. The PCR
approach showed partially varying results from our data with an upregulation of IL-
1R tI and IL-1ra mRNA simultaneous to an IL-1 β downregulation in tubes with
ectopic pregnancies. The study by Huang et al. applied a quantitative PCR approach
375 using a competitor compared to the qualitative method which was used in this study
showing no upregulation of IL-1RtI and a downregulation of IL-1 β but rather the
presence of both in fallopian tubes of extrauterine pregnancies. Presumptions about a

regulation can not be drawn therefore. However the upregulation of IL-1ra in extrauterine tubes is opposed to the absence of IL-1ra in extrauterine and healthy proliferative phase tubes as occurred in the present study. Both studies have investigated a relatively small number of fallopian tube samples most likely due to low availability which might explain the partly varying results.

Based on those findings in human as well as in the animal studies the IL-1 β system appears to be involved in the establishment of pregnancy although the exact magnitude is still questionable due to the conflicting data in the literature. Furthermore, rigorous species specific differences might attribute to an inconsistency in the importance of the IL-1 system in implantation as well.

To our knowledge this is the first study to report that the gene expression profile is not altered within the different anatomical parts of the fallopian tube showing a constant gene expression pattern of all IL-1 β system factors throughout the complete length of the oviduct investigated. Additionally, the restricted IL-1ra expression in secretory phase fallopian tubes and absence in extrauterine pregnancy tube samples might be an important mechanism on the molecular base to prevent ectopic pregnancies within the oviduct which needs further investigation. The finding that IL-1R tI is expressed in both cycle phases independent of a pregnancy in addition to the embryos own IL-1 β expression supports the concept of an early embryo-maternal dialogue facilitating a proper implantation process.

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Figure Legends**Figure 1**

530 Fig. 1 Representative RT-PCR-gelelectrophoresis for IL-1 β (549 bp), IL-1ra (424 bp) and IL-1R t1 (284 bp) for 3 samples each showing that the gene expression of IL-1 β , IL-1ra and IL-1R t1 within the length of the fallopian tube has no spatial difference (β -actin with 838 bp serves as control). All oviduct samples were divided into fimbrial, ampullary and isthmic parts and investigated separately by RT-PCR. From the total of

535 19 samples, all were positive for β -actin mRNA expression (M - DNA size marker, p - proliferative, s - secretory, f - fimbrial, a - ampullary, i - isthmic).

Figure 2

Fig. 2 Representative RT-PCR gelelectrophoresis for temporal expression patterns of

540 IL-1 β , IL-1ra and IL-1R t1 showing a temporal and therefore most likely hormonal regulation of IL-1 β and IL-1ra, whereas the receptor was expressed in all samples. From the total of 23 samples, all were positive for β -actin mRNA expression (M - DNA size marker, p - proliferative, s - secretory, m - menopausal, iup - intrauterine pregnancy, eup - extrauterine pregnancy).

545

Figure 3.

Fig. 3 Immunohistochemistry staining of representative samples from fallopian tubes of the secretory (A-H) and proliferative phase (I-P) (panel A-G and I-O 200 $\times$ magnification, panel B-H and J-P 400 $\times$ magnification). Panels A and B represent the

550 negative control of secretory phase oviducts, panel I and J represent the negative

control (no first antibody) of proliferative phase oviducts. Note that IL-1 β is strong expressed in the luminal epithelium and also in the small vessel endothelium of secretory phase oviducts (C,D), whereas the staining for IL-1 β in the luminal epithelium of proliferative phase oviducts was less intense and only a weak staining was localized in the small vessel endothelium (3 K,L). IL-1R t1 is expressed in the luminal epithelium, small vessel endothelium and glandular epithelium of secretory phase samples (E, F). In comparison, the proliferative sample showed only a sporadic staining for the IL-1R t1 in the luminal epithelium and no staining either in the vessel endothelium or in the glandular epithelium (M, N). G and H show an intense staining of IL-1 ra in the luminal epithelium and the small vessel endothelium of secretory phase oviducts whereas a staining for IL-1ra in proliferative oviducts was completely absent.

Table 1

mRNA	size of amplified fragment [bp]	position of primers on cDNA	3' / 5' - end	sequence of oligonucleotide (5'→3')
β-actin	838	294-325	5'-end	ATC TGG CAC CAC ACC TTC TAC AAT GAG CTG CG
		1100-1131	3'-end	CGT CAT ACT CCT GCT TGC TGA TCC ACA TCT GC
IL-1β	549	101-122	5'-end	CAG TGA AAT GAT GGC TTA TTA G
		628-648	3'-end	CTT TCA ACA CGC AGG ACA GGT
IL-1ra	424	91-114	5'-end	CAG AAG ACC TCC TGT CCT ATG AGG
		498-514	3'-end	GCTT GTG CAG AGG AAC CA
IL-1R tI	284	341-361	5'-end	AAG GTG GAG GAT TCA GGA CAT
		604-624	3'-end	AGC CTA TCT TTG ACT CCA CTA

Tab. 1: primers for synthesis of cDNA clones of human β-actin, IL-1β, IL-1ra and IL-1R tI

Figures

Figure 1

Figure 2

Figure 3