Region-specific expression of nitric oxide synthases in the bovine oviduct during the oestrous cycle and *in vitro*

S E Ulbrich, S Rehfeld², S Bauersachs², E Wolf², R Rottmayer²,
S Hiendleder², M Vermehren³, F Sinowatz³, H H D Meyer
and R Einspanier¹

Physiology-Weihenstephan, Technical University of Munich, Alles Hörsaalgebäude, 85350 Freising, Germany

¹Institute of Veterinary Biochemistry, Free University of Berlin, Oertzenweg 19b, D-14163 Berlin, Germany

²Institute of Molecular Animal Breeding and Biotechnology, Gene Centre of the Ludwig-Maximilians-University Munich, Munich, Germany

³Institute of Veterinary Anatomy, Histology and Embryology, Ludwig-Maximilians-University Munich, Veterinaerstr. 13, 80539 Munich, Germany

(Requests for offprints should be addressed to R Einspanier; Email: einspani@zedat.fu-berlin.de)

Abstract

Nitric oxide synthases (NOS) account for the endogenous production of nitric oxide (NO), a small and permeable bioreactive molecule. NO is known to act as a paracrine mediator during various processes associated with female reproduction. In the present study, the mRNA expression of the endothelial (eNOS) and inducible (iNOS) NO synthases were examined in bovine oviduct epithelial cells (BOEC) during the oestrous cycle. In addition, eNOS and iNOS mRNA and protein were localised by *in situ* hybridisation and immunocytochemistry respectively. Furthermore, the effects of exogenously applied oestradiol-17 β and progesterone on NOS mRNA regulation were studied in a suspension culture of BOEC. The eNOS mRNA abundance was low around ovulation (day 0) and increased significantly until pro-oestrus (day 18) in the ampulla. Immunoreactive protein of eNOS was

detected predominantly in endothelial cells as well as in secretory oviduct epithelial cells at pro-oestrus. The iNOS mRNA concentration was significantly reduced in the isthmus at pro-oestrus (day 18) and oestrus (day 0) compared with persistently high levels in the ampulla. By *in situ* hybridisation, specific iNOS transcripts were additionally demonstrated in the oviduct epithelium. Immunoreactive iNOS protein was localised in secretory epithelial cells as well as in the lamina muscularis. The *in vitro* stimulation showed that both NOS were stimulated by progesterone, but not by oestradiol-17 β . The region-specific modulated expression of eNOS and iNOS provides evidence for an involvement of endogenously produced NO in the regulation of oviductal functions.

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Introduction

The oviduct is responsible for the accommodation of the gametes and the early embryo by providing an optimal environment for successful fertilisation. In addition, it accounts for the transport of the gametes and the embryo into the uterus. The anorganic free radical nitric oxide (NO) is a small, unstable and permeable molecule that can pass through membranes by diffusion. NO is an important intercellular regulatory molecule and a major paracrine mediator, functioning as a vascular, immunological and neuronal signalling molecule (Ignarro *et al.* 2001). NO can react with several targets by exerting unspecific immune responses (Guzik *et al.* 2003). By binding to guanylyl cyclase, cyclic guanosine monophosphate (cGMP)-specific proteinases (protein kinase G), Na⁺ ions and phosphodiesterases can be activated, leading to modulated gene expression (Nathan & Xie 1994). The effect of NO

on vasodilatation has been demonstrated (Furchgott & Vanhoutte 1989). NO is assumed to have a broad range of functions in reproductive processes such as oocyte maturation, ovulation, implantation, pregnancy maintenance, labour and delivery (Shukovski & Tsafiriri 1994, Jablonka-Shariff *et al.* 1999, Sengoku *et al.* 2001, Maul *et al.* 2003).

NO is produced by the conversion of L-arginine to L-citrulline by the enzyme NO synthase (NOS) with several co-factors in a number of different tissues and cell types. To date, three isoforms of NOS, products of separate genes with different molecular weight but apparently similar molecular structure, have been described: neuronal NOS (nNOS) in the brain and peripheral nervous system; endothelial NOS (eNOS) as a constitutive NOS mainly in the endothelium; and inducible NOS (iNOS), synthesised primarily by activated macrophages, hepatocytes and neutrophils in several tissue types and organs and upon inflammatory stimulation (Moncada *et al.* 1997).

Recently, NOS have been identified in the human, bovine and rat oviduct (Bryant *et al.* 1995, Rosselli *et al.* 1996, Ekerhovd *et al.* 1997). Nevertheless, more comprehensive analyses could help to bridge lack of congruency for different species (Chatterjee *et al.* 1996, Gawronska *et al.* 2000) with respect to cellular localisation, regions of the oviduct and cycle-dependent regulation. Therefore, the present study investigated thoroughly the expression and localisation of iNOS and eNOS in the bovine oviduct of days 0, 3·5, 12 and 18 of the oestrous cycle. In addition, the *in vivo* data were compared with NOS expression in bovine oviduct epithelial cells (BOEC) *in vitro*. The obtained data can help to assign a specific regulatory role to NOS in the oviduct.

Materials and Methods

Synchronisation of the oestrous cycle and collection of oviduct samples

Twelve cyclic Simmental heifers 18–24 months old were cycle synchronised by injecting intramuscularly a single dose of 500 µg cloprostenol (Estrumate; Essex Tierarznei, Munich, Germany) at dioestrus. Animals were intensively observed for sexual behaviour (i.e. toleration, sweating, vaginal mucus) to determine standing heat, which occurred around 60 h after Estrumate injection.

Ovary function was monitored by ultrasonographic examination to locate the time frame of ovulation, and blood samples were taken to determine serum progesterone (P4) levels, which were executed in 6-h rhythms. Three animals were slaughtered the morning after standing heat occurred (day 0) and three animals each at days 3·5, 12 and 18 after oestrus respectively. Final blood samples were taken just before slaughter to determine serum P4 levels. Animals slaughtered at oestrus (day 0) displayed low serum P4 levels (<1·0 ng/ml) and animals slaughtered at dioestrus (day 12) had high serum P4 levels (>6 ng/ml). Oviducts were trimmed free of surrounding tissue, subdivided into ampulla and isthmus, and dissected in 1-cm-long sections. To isolate the epithelial cells, the sections were stripped with two tweezers, and the epithelia were transferred directly in cryotubes. These tubes were dropped immediately into dry ice, transported in liquid nitrogen and stored at –80 °C until further processing. All samples were taken and frozen within 20 min after death.

In vitro cell suspension culture

Four heifers were slaughtered on day 3·5 of the oestrous cycle, and BOEC were obtained as described previously (Rottmayer *et al.* 2005). Briefly, the oviducts were squeezed along the ampulla with forceps. The cell sheets were separated mechanically by repeated passages

through syringes and pipetting, and recovered by sedimentation. Cells were cultured in 24-well plates with 800 µl TCM-199 supplemented with 2% serum obtained from heifers on day 3·5 of the oestrous cycle and 0·25 mg/ml gentamicin at a density of 10⁶ cells per well at 38 °C in a humidified atmosphere of 5% CO₂ in air. BOEC were stimulated with oestradiol-17β (10 pg/ml) or progesterone (10 ng/ml) (both purchased from Sigma) for 6 and 12 h respectively. Both experiments were carried out with appropriate negative controls diluting the carrier of the steroid (ethanol and water respectively) in the same way as the steroid itself. Two animals were also used for a short-time stimulation of 2 h. Cells were collected by centrifugation, washed in buffer solution, snap-frozen in liquid nitrogen and stored at –80 °C until further investigation.

Reverse transcription and real-time RT-PCR

Total RNA from BOEC *in vivo* and *in vitro* was isolated with TRIzol reagent (Invitrogen) according to the manufacturer's instructions. Two-step quantitative real-time RT-PCR with the LightCycler DNA Master SYBR Green I protocol (Roche) was done as described previously (Ulbrich *et al.* 2004). Briefly, 1 µg of each sample of RNA was reverse transcribed in a total volume of 60 µl: 5X Buffer (Promega), 10 mM dNTPs (Roche), 50 µM hexameres (Gibco BRL), 200 U Superscript RT enzyme (Promega). The conventional PCR was performed in a thermal cycler (Biometra, Göttingen, Germany) as previously described (Berisha *et al.* 2002). A 7-µl volume of each reaction was subsequently subjected to agarose gel electrophoresis followed by ethidium bromide staining. For each of the following real-time PCR reactions, 1 µl cDNA was used to amplify specific target genes. In each PCR reaction, 17 ng/µl cDNA were introduced and amplified in a 10 µl reaction mixture (3 mM MgCl₂, 0·4 µM primer forward and reverse each, 1 LightCycler DNA Master SYBR Green I; Roche) and compared with a standard curve based on a specific PCR product. Primers were adapted to amplify specific PCR products for 18S rRNA (forward: 5'-AAGTCTTTGGGTTCGCGG-3'; reverse: 5'-GGACATCTAAGGGCATCACA-3' [365 bp]), eNOS (forward: 5'-AGGAGTGAAGTGTTCCG-3'; reverse: 5'-GCCCCGGTACTACTCTGTCA-3' [126 bp]) and iNOS (forward: 5'-ACCTACCAGCTGACGGGAGAT-3'; reverse: 5'-TGGCAGGGTCCCCTGTGATG-3' [316 bp]). The predicted size of each PCR product is assigned in brackets. The amplified PCR fragments were sequenced once (MWG, Ebersberg, Germany) to verify the resulting PCR product (Einspanier *et al.* 2002). Thereafter, the specific melting point (MP) of the amplified products served as verification of the product identity (18S (MP 88 °C, fluorescence acquisition at 80 °C), eNOS (MP 93 °C, fluorescence acquisition at 87 °C) and iNOS (MP 90 °C, fluorescence acquisition at 86 °C)).

(Ulbrich *et al.* 2004). The annealing temperature was 60 °C for 18S, 61 °C for iNOS and 66 °C for eNOS. As negative control, water instead of cDNA was used. The nucleotide sequence for the partial bovine iNOS cDNA was subsequently submitted to the EMBL database (accession no. AJ699400). The cycle number (CP) required to achieve a definite SYBR Green fluorescence signal was calculated by the second derivative maximum method (LightCycler software, Version 3.5.28). The CP was correlated inversely with the logarithm of the initial template concentration.

Data analysis of real-time RT-PCR

The CP determined for the target genes were normalised against the housekeeping gene 18S. Resulting data are presented as means of CP ($n=3$) $\pm$ S.E.M. For statistical analysis, the SAS program package release 9.1.3 (2002, SAS Institute, Cary, NC, USA) was used. All data were analysed by one-way ANOVA (analysis of variance). In case of significant different groups, a multiple *t*-test analysis was done with the Bonferroni correction. Results were considered statistically significant at $P<0.05$.

In situ hybridisation

The detailed *in situ* hybridisation procedure has been described previously (Bauersachs *et al.* 2005). Briefly, formalin-fixed, paraffin wax-embedded samples were used. Sections were deparaffinised with xylene and immersed in isopropanol. Dried sections were submerged in 2 saline sodium citrate buffer and preheated at 80 °C. Slides were then washed in distilled water and TBS and permeabilised with 0.05% proteinase K (VWR, Ismaning, Germany) in Tris-buffered saline (TBS) at room temperature. Sections were relocated in TBS followed by distilled water and post-fixed for 10 min in 4% paraformaldehyde/PBS. After washing in PBS and distilled water, slides were dehydrated and air-dried. Hybridisation was carried out by overlaying the dried sections with the corresponding biotinylated oligonucleotide probe (100 pmol/ μ l), diluted 1:20 in *in situ* hybridisation solution (DAKO, Munich, Germany), and incubating them in a humidified chamber at 38 °C overnight. The sequence of the iNOS antisense oligonucleotides was 5'-TCCAGCATCTCCTCCCAGTA-3'. RNase-free hybridisation solution (DAKO, Munich, Germany) contained 60% formamide, 5 $\times$ SSC, hybridisation accelerator, RNase inhibitor and blocking reagents. Subsequently, slides were washed in 2 $\times$ SSC (2 h 15 min, preheated to 38 °C), distilled water (25 min) and TBS (25 min). Hybridised probes were detected with HRP-labelled ABC kit reagents developed by DAB (DAKO, Munich, Germany) according to the manufacturer's instructions. Negative controls were done by

exchanging the oligonucleotide probe with the corresponding sense oligonucleotide.

Immunohistochemistry

For the immunohistochemical demonstration of eNOS and iNOS, tissue samples were fixed in Bouin's solution for 12 h, as described previously (Berisha *et al.* 2004). The specimens were dehydrated and embedded in paraffin wax. Serial sections (5 μ m) were cut on a Leitz microtome and collected on gelatin/chrom alum-coated slides. To expose antigenic sites, dewaxed sections were heated four times to 95 °C in a 600 W microwave oven in citrate buffer for 5 min. Endogenous peroxidase activity was then eliminated by incubation with 0.5% (v/v) H₂O₂ solution in absolute methanol for 15 min at 20 °C. Non-specific protein binding was eliminated by incubation with 10% normal goat serum in PBS for 1 h at 20 °C. Sections were then incubated with either polyclonal rabbit antibody against iNOS (Upstate, Lake Placid, NY, USA) (Sherman *et al.* 1999), which is known to cross-react with bovine iNOS, or monoclonal rabbit antibody against eNOS (Alpha Diagnostic, San Antonio, TX, USA), from which the human peptide sequence is 100% conserved in the bovine (Marsden *et al.* 1992, Welter *et al.* 2004). Each was used at a dilution of 1:200. Incubation was performed at 18 h at 4 °C in a humidified chamber. This was followed by incubating the sections with biotinylated anti-rabbit IgG 1:400 (Amersham-Pharmacia) for 1 h. The sections were then incubated with ABC reagent from a commercial kit (Vector Laboratories, Burlingame, CA, USA). The bound complex was made visible by reaction with 0.05% 3,3-DAB and 0.0006% H₂O₂ in 0.1 $\times$ PBS. Sections were viewed unstained or counterstained in Mayer's haematoxylin, dehydrated, cleared and mounted. Controls were performed by either replacing primary antibody with buffer or non-immune serum, or incubating with DAB reagent alone to exclude the possibility of non-suppressed endogenous peroxidase activity. Lack of detectable staining in the controls demonstrated the specificity of the reactions.

Results

A first screening of eNOS and iNOS mRNA in bovine oviduct total RNA by conventional RT-PCR led to different amplification signal intensities, as shown in Fig. 1. To further characterise these preliminary results, a real-time RT-PCR approach was applied to quantify the gene expression patterns in oviduct epithelial cells.

Transcripts of eNOS and iNOS were detected in BOEC throughout the oestrous cycle. A statistical analysis of eNOS/iNOS transcript concentrations from the ipsi-versus contralateral side revealed no significant differences at any time point ($P=0.07$) in either ampulla or isthmus.

Figure 1 Messenger RNA transcripts of the housekeeping gene 18S and the target genes eNOS and iNOS in the bovine oviduct during the oestrus cycle, exemplified by conventional RT–PCR demonstrating differential expression between ampulla and isthmus as well as between the days of the oestrous cycle. One representative experiment is shown.

Therefore ipsi- and contralateral oviducts were grouped, so that each bar in Fig. 2 represents a data set of six individual oviducts from three different animals.

The eNOS mRNA expression in the ampulla was low at oestrus (day 0) and increased significantly, more than threefold, until reaching the highest levels at pro-oestrus (day 18) (70.7 and 245.4 fg mRNA/μg total RNA respectively) (Fig. 2A). The same tendency was observed in the isthmus, yet here the differences between days 0 and 18 were not significant. Immunoreactive protein was observed in endothelial cells of blood vessels (data not shown). Additionally, supranuclear staining in secretory epithelial cells of the oviduct epithelium was visible at day 18 (Fig. 3A). Nuclear and cytoplasmic staining of both smooth muscle and endothelial cells of small blood vessels could be detected also in the lamina muscularis (Fig. 3B).

The highest transcript amounts for iNOS were detected in the ampulla (3680 fg mRNA/μg total RNA at day 18), high levels being retained throughout the oestrous cycle (Fig. 2B). At oestrus (day 0), transcript levels in the isthmus were almost 10-fold lower than in the ampulla (333.8 and 2950 fg mRNA/μg total RNA respectively). At day 3.5 the iNOS expression in the isthmus was already again as high as in the ampulla (3246 and 2657 fg mRNA/μg total RNA respectively). It declined to an intermediate level at day 12, which was significantly lower than at day 3.5 (810.7 fg mRNA/μg total RNA in the isthmus). At pro-oestrus (day 18), there was a significantly lower iNOS mRNA concentration (threefold) in the isthmus than the ampulla (1222 and 3676 fg mRNA/μg total RNA respectively). *In situ* hybridisation revealed transcripts in the epithelial cells mainly in the oviductal ampulla at pro-oestrus and oestrus (Fig. 3 G and H), in accordance with the high mRNA transcript levels measured by real-time RT–PCR (Fig. 2). Immunohistochemical analysis revealed a conspicuous supranuclear staining in secretory epithelial cells mainly toward

the lumen (Fig. 3C). The staining appeared mostly at the apical, and not in the basal parts of the luminal branching folds (Fig. 3E). Additionally, pronounced nuclear staining of the lamina muscularis was observed (Fig. 3D and F). The stroma was consistently devoid of iNOS protein (Fig. 3C and E).

The stimulation of BOEC with oestradiol-17β had no effect on the transcript regulation of eNOS (Fig. 4A). However, progesterone significantly stimulated eNOS transcript levels more than threefold only 2 h after application (36.7 and 112 fg mRNA/μg total RNA respectively) (Fig. 4A).

Nor did oestradiol-17β have an effect on the transcript regulation of iNOS in cultured BOEC (Fig. 4B). But a significant twofold upregulation of iNOS transcript abundance was observed 6 and 12 h after progesterone stimulation (399 and 904 fg mRNA/μg total RNA, and 368 and 883 fg mRNA/μg total RNA respectively) (Fig. 4B).

Discussion

This study demonstrates the presence of the two iso-enzymes eNOS and iNOS in the bovine oviduct during the oestrous cycle. Moreover, immunoreactive protein of eNOS and iNOS could be located in distinct cell types of the oviduct.

In agreement with our mRNA data for iNOS, the lowest expression of NADPH diaphorase as a marker for NOS was found in the isthmus at oestrus (Gawronska *et al.* 2000). High expression was noticed during the luteal phase (Gawronska *et al.* 2000). The presence of NADPH diaphorase activity in the porcine oviduct has been demonstrated during the oestrous cycle. Bryant *et al.* (1995) showed reduced NO activity during late pro-oestrus by measuring the conversion of L-arginine to L-citrulline in the rat oviduct, and eNOS in the

Figure 2 Messenger RNA expression (real-time RT-PCR) of (A) eNOS and (B) iNOS in bovine oviduct cells during the oestrous cycle in either ampulla (■) or isthmus (□). eNOS mRNA was significantly upregulated in the ampulla between oestrus (day 0) and pro-oestrus (day 18). There was a remarkable downregulation of iNOS mRNA in the isthmus compared with the ampulla at oestrus (day 0) and dioestrus (day 18). Data are presented as means of mRNA/total RNA $\pm$ S.E.M. normalised by 18S ($n=6$). *Significant differences between ampulla and isthmus ($P<0.05$). Different superscript letters indicate significant differences between days of the oestrous cycle ($P<0.05$).

rat oviduct was most prevalent at pro-oestrus and oestrus (Chatterjee *et al.* 1996). The latter result may point to a direct oestradiol effect on eNOS in rats, but this was not confirmed in the present study for the cow. Instead, the rising levels of eNOS mRNA between oestrus (day 0), dioestrus (day 3.5 and 12) and pro-estrus (day 18) indicate the dependency of eNOS on progesterone. This assumption is supported by the stimulation of NOS mRNA expression by progesterone in cultured BOEC as well as the eNOS protein staining *in vivo* mainly at pro-oestrus.

Current data on the localisation of NOS in the oviduct are controversial. NADPH diaphorase was demonstrated mainly in the oviduct epithelium of man, rat and pig (Bryant *et al.* 1995, Ekerhovd *et al.* 1999, Gawronska *et al.* 2000), and eNOS protein was consistently present in oviduct epithelial cells of different species (Bryant *et al.* 1995, Rosselli *et al.* 1996, Ekerhovd *et al.* 1997, Gawronska *et al.* 2000). In the present study, we clearly

demonstrate the presence of both eNOS and iNOS in BOEC. Previously, iNOS protein was observed in the epithelium of the human and the rat oviduct (Bryant *et al.* 1995, Ekerhovd *et al.* 1997), but not in the pig (Gawronska *et al.* 2000). These peculiarities may indicate species differences. In the porcine oviduct, eNOS was found to be the predominant isoform (Gawronska *et al.* 2000). The present results reveal that the absolute expression of iNOS is much higher than eNOS, in addition to a more pronounced regulation during the oestrous cycle. Since the latter isoform produces lower quantities of NO (Ekerhovd *et al.* 1999), eNOS in the bovine oviduct might be the isoenzyme responsible for the rather constitutive presence of NO.

NADPH diaphorase as well as eNOS was demonstrated in the myosalpinx of the rat and pig (Bryant *et al.* 1995), and iNOS was also found in smooth muscle of the human oviduct (Ekerhovd *et al.* 1997). The relaxing effect of NO

Figure 3 Immunohistochemical localisation of eNOS (A and B) and iNOS (C–F) in bovine oviducts. Black arrowheads (A–F) point at specific immunopositive secretory epithelial cells with supranuclear staining mainly toward the lumen. The staining appeared mostly at the apical, and not in the basal, parts of the luminal branching folds (E). Arrows (A–F) point at immunopositive cells for eNOS and iNOS in the lamina muscularis with pronounced nuclear staining. *In situ* hybridisation of iNOS (G and H) in bovine oviducts (brown staining) clearly showed positive iNOS mRNA transcripts specifically in epithelial cells of the ampulla (black spearheads). The black bar indicates 50 µm.

Figure 4 Messenger RNA expression (real-time RT-PCR) of (A) eNOS and (B) iNOS transcripts in a bovine oviduct epithelial cell suspension culture after stimulation with progesterone or oestradiol-17 β . Transcripts of both eNOS and iNOS were stimulated by progesterone, but not by oestradiol-17 β . Data are presented as means of mRNA/total RNA $\pm$ S.E.M. normalised by 18S ($n=4$). *Significant differences between control and stimulation ($P<0.05$).

on smooth muscle, possibly controlled by progesterone (Chwalisz 1994), is well known, particularly for uterine quiescence during pregnancy (Yallampalli *et al.* 1993). The motility patterns of the oviduct show rising frequency and amplitude of motility around oestrus (Bennett *et al.* 1988). Therefore, several mediators of contraction could be involved in this regulation, influencing or orchestrating the NOS expression in the oviduct, namely, oestradiol, oxytocin, prostaglandin (PG) $F_{2\alpha}$, PGE_2 and endothelin-1 (Moore & Croxatto 1988, Gilbert *et al.* 1992, Salvemini *et al.* 1993, Rosselli *et al.* 1994, Perez *et al.* 1998). $PGF_{2\alpha}$ can induce NO production by NOS in rat oviduct cells (Perez *et al.* 1998). The highest concentrations of $PGF_{2\alpha}$ receptors are found around oestrus in the rat (Orlicky & Williams-Skipp 1993), and oestradiol is known to activate PG synthase. NO could therefore negatively modulate or antagonise the contractile response of $PGF_{2\alpha}$. Furthermore, NO participates in the release of PGE_2 (Salvemini *et al.* 1993). PGE_2 is increased at oestrus (Wijayagunawardane *et al.* 1998) and has been held to

cause relaxation of the oviduct in the presence of progesterone (Gawronska *et al.* 2000). Subsequently, PGE_2 , together with progesterone, could regulate NOS expression (Milano *et al.* 1995), an effect that would agree with the *in vitro* findings of the present study. Beside these findings, endothelin-1 (ET-1) is high in the ipsilateral oviduct during the follicular and postovulation stage (Wijayagunawardane *et al.* 1998). ET-1 has been shown to stimulate NO in BOEC via endothelin receptor beta (Rosselli *et al.* 1994). NO reduces the contractile effects of ET-1, hence the interplay of ET-1 and NO might contribute to the physiological relaxation of the oviduct.

Using L-NAME (*N*-nitro-L-arginine methyl ester), a well-known inhibitor of NOS, Perez *et al.* (2000) found evidence of increased tubal motility that resulted in accelerated ovum transport into the uterus. Moreover, oestradiol treatment caused increasing contraction frequency of the smooth muscle of the isthmus (Moore & Croxatto 1988). This could be mediated by oestrogen

receptor β , which is more abundant in the isthmus (Ulbrich *et al.* 2003). The endogenous local downregulation of iNOS in the isthmus could support similar effects. Therefore, our hypothesis is that the downregulation of iNOS at oestrus in the isthmus leads to an increase of oviduct motility by circular smooth muscle activity.

There is evidence that the accelerated movement of microspheres through the isthmus is due to peristaltic smooth muscle contractions, and to ciliary activity, and this is supported by the observation of a reduced number of ciliated cells in the isthmus (Perez *et al.* 1998). Moreover, the ciliated epithelial cells were mostly devoid of iNOS protein, whereas the lamina muscularis was clearly stained.

In secretory epithelial cells of the oviduct, the nucleus is shifted toward the apical side of the epithelium (unpublished data). We may deduce from this that both eNOS and iNOS proteins target mostly secretory epithelial cells, as has also been shown for eNOS in the human Fallopian tube (Ekerhovd *et al.* 1999). iNOS protein was found in secretory cells toward the oviduct lumen in the supranuclear region; therefore, it can be assumed that NO is released into the lumen. It might be of further interest to know whether the produced NO is also secreted toward the stroma and myosalpinx, or whether eNOS found in smooth muscle cells affects the contractibility of the oviduct. The contribution of NOS might be important for relaxation together with a local exertion during (pro-)oestrus, facilitating capture, retention and fertilisation of the released oocyte and the active transport of the conceptus (Chatterjee *et al.* 1996). Furthermore, through the ability of iNOS to produce cytotoxic levels of NO (Guo *et al.* 1995), the downregulation of iNOS in the isthmus at oestrus could be an implicit protective mechanism for advancing sperm and the developing embryo.

The present study does not take into account further regulation of NOS enzyme activity. In particular, the proposed differences in enzyme activity of both NOS isoforms (Presta *et al.* 1997) cannot be measured through expression analysis solely. Nonetheless, this investigation provides new data on a local NO-regulating mechanism in the bovine oviduct.

In summary, the present results provide evidence of the presence of NOS in the bovine oviduct. The region-specific modulated mRNA expression patterns of eNOS and iNOS during the oestrous cycle indicate a local regulatory system of NO in the bovine oviduct and suggest a different role of NO in the ampulla and the isthmus region. Although functional analyses still remain to be done, the conspicuous downregulation of iNOS at oestrus in the isthmus requires further research. NO might represent another important local factor regulating oviducal functions with possible impact on contractility response. The present findings underline the physiological influence of both NOS in supporting a successful fertilisation by regulating the oviduct environment.

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