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**Evidences that Progesterone Receptor B  
decreases Estrogen Receptor  $\alpha$  gene  
expression through its interaction to a  
half-PRE site at Estrogen Receptor  $\alpha$   
gene promoter.**

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## SUMMARY

Regulated expression of Progesterone Receptor-A (PR-A) and Progesterone Receptor-B (PR-B) is critical for appropriate mammary gland responsiveness to Progesterone (Pg) which has a broad impact on target cells. There is considerable biological evidence for cross talk between the progesterone and estrogen hormone receptor signalling pathways even though the molecular mechanisms remains to be further elucidated.

In the present study we demonstrate that PR mediates the down-regulation of the E2 induced cell proliferation due to Pg cotreatment in MCF-7 breast cancer cells. In order to assess whether increased PR-B/PR-A ratio could affect response to E2 through alteration of ER $\alpha$  expression levels in MCF-7 cells, we transiently over-expressed PR-B isoform. We show that PR-B isoform over-expression, robustly inhibits ER $\alpha$  expression in terms of mRNA and protein content, causing concomitantly the repression of the estrogen regulated genes cyclin D1 and pS2, addressing a specific effect of PR/PR-B on ER $\alpha$  gene transcription. For instance ER $\alpha$  gene promoter activity was drastically inhibited by PR-B over-expression. ER $\alpha$  gene promoter analysis reveals a responsive region containing a half-PRE site located at -1757 bp to -1752 bp as putative effector of PR action. In presence of mutation of the half-PRE, the down-regulatory effect induced by PR/PR-B over-expression was abrogated. Moreover chromatin immunoprecipitation

analyses, reveal an increase of PR bound to the ER $\alpha$  regulatory region and recruitment of a corepressor complex containing NCoR but not SMRT and DAX1, concomitantly with hypoacetylation of the chromatin and displacement of RNA polymerase II. Furthermore PR-B ablation studies using small interfering RNA demonstrate the specific recruitment of PR-B isoform on ER $\alpha$  promoter.

Our study provide evidences for a mechanism by which over-expressed PR-B is able to actively repress ER $\alpha$  gene expression in vivo. The antiestrogenic effect of over-expressed PR-B also provides potential therapeutic strategy for breast cancer treatment.

## **INTRODUCTION**

The sex steroid hormones, Estrogen (E2) and Progesterone (Pg), play an important role in normal mammary gland development, and it is believed that breast cancer progression is influenced by them and their receptors (1, 2). The level of these steroid hormones receptors is a strong prognostic factor for patients with breast cancer and has been used in clinical management as an indicator of endocrine responsiveness (3, 4).

Approximately 75% of primary breast cancers express ER, and more than half of these cancers also express PR.(5) Both ER and PR are prognostic factors, although both are weak and lose their prognostic value after long-term followup.(6) PR is an estrogen-regulated gene, and its synthesis in normal and cancer cells requires estrogen and ER. Therefore, it is not surprising that ER positive/PR-positive tumors are more common than ER-positive/PR-negative tumors. The etiology of ER-positive/PR-negative tumors is currently unclear. Some studies have shown that ER and PR status can change over the natural history of the disease or during treatment.(7) For instance, sequential breast cancer biopsies have shown that ER levels are reduced slightly with intervening endocrine therapy, although complete loss is uncommon. In contrast, PR levels decrease more dramatically during tamoxifen therapy, with up to half of tumors completely losing PR expression when resistance develops.(8) These ER-

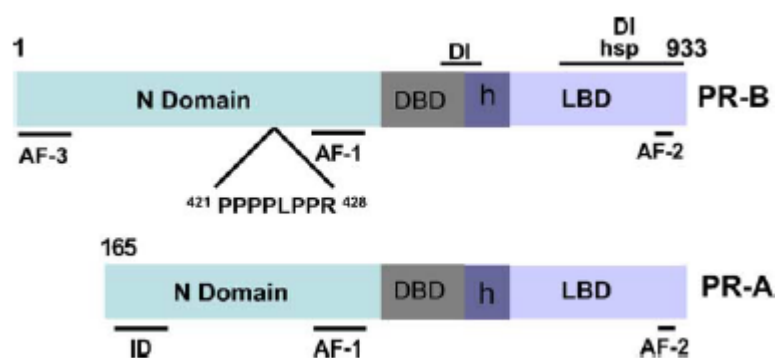
positive/PR-negative metastatic tumors then display a much more aggressive course after loss of PR compared with tumors retaining PR, and patients then have a worse overall survival, indicating a change in tumor cell-regulatory mechanisms.(9,10) Whether and how the loss of PR affects the poor clinical course of these tumors is at present unclear.

Although Estrogen Receptor (ER) is an accepted predictor of response to endocrine therapy the role of Progesterone Receptor (PR) has been more controversial. Recent studies published on the largest retrospective analysis of early breast cancer treated with tamoxifen found that patients with ER+/PR+ tumors benefited much more from adjuvant tamoxifen therapy than those with ER+/PR- tumors (11,12). Importantly, multivariate analyses, including lymph node involvement, tumor size, and age, show that PR status was independently associated with disease-free and overall survival.

Progesterone Receptors belong to the subfamily of classical nuclear steroid receptors; human PR proteins exist as two isoforms, termed PR-A and PR-B.

Both PR-A and PR-B contain three functional domains including the N-terminus, a centrally located DNA binding domain (DBD), and C-terminal ligand binding domain (LBD). Three-dimensional structures of isolated DBD and LBDs have revealed common motifs for these regions. By comparison, little is known about the structure of the N-terminal domain. This is the least conserved region among family members with re-spect to both length and amino acid sequence. The N-domain is functionally important, as it is required for full transcriptional activity of

steroid hormone receptors and for many cell- and target gene-specific responses. Other functional and structural determinants have been identified within these broader three domains. In addition to binding steroid hormone, the LBD contains determinants for dimerization (DI) in the absence of DNA, binding of heat shock proteins (hsps) and for nuclear localization sequence (NLS). The DBD contains a second NLS and dimerization domain that is dependent on DNA binding. Steroid receptors contain at least two transcription activation domains (AFs), AF-1 in the N-terminal domain and highly conserved AF-2 in the C-terminal LBD. These are autonomous transferable domains required for the DNA bound receptor to transmit a transcriptional activation response and they function as specific binding sites for coactivators. AF-2 located in the LBD is hormone-dependent and becomes activated as a result of the steroid hormone inducing a conformational change that creates a hydrophobic binding pocket for members of the p160 family of steroid receptor coactivators (SRCs). The coactivators that bind to and mediate the activity of AF-1 are yet not well defined.





The human PR is expressed as two isoforms from a single gene by alternate promoter usage. PR-A (94 kDa) differs from PR-B (120 kDa,) by lacking 164 amino acids (aa) at the N-terminus . Although the two forms of PR have similar steroid hormone and DNA binding activities, they have distinct transcriptional activities. PR-B in general, is a much stronger activator than PR-A. However, PR-A can be a strong activator under specific cell and target gene contexts. The stronger activation potential of PR-B is due in part to the existence of a third activation domain (AF-3) within the first N-terminal 164 aa that is unique to PR-B. Under certain cell and target promoter contexts PR-A is inactive as a transcription factor and can function as a ligand-dependent transdominant repressor of other steroid receptors including PR-B and the estrogen receptor (ER). PR-A can act in this repressor mode in response to the binding of either progestin agonists or antagonists. An inhibitory domain (ID) responsible for this transrepressor function has been mapped to the first 140 N-terminal (aa 165–305) amino acids of PR-A. Since the sequence within ID is present in both PR isoforms but is only active in the context of PR-A suggests the PR-B specific N-terminal segment plays a role in suppressing the ID domain.

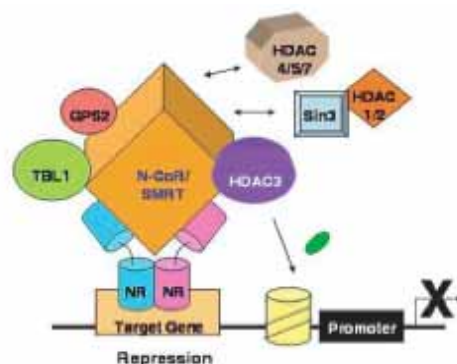
Despite structural similarities, PR-A and PR-B regulate different subsets of genes and although PR-B is transcriptionally more active, there are genes, known to be involved in breast cancer progression, that are uniquely regulated by PR-A (13,14,15).

In vivo the two PR isoforms are usually co-expressed at similar levels in normal cells, yet their ratio varies dramatically in different tissues, physiological states and disease (16,17). With regard to the mammary gland, in transgenic mice, 3:1 over-expression of PR-A over PR-B results in extensive epithelial cell hyperplasia, excessive ductal branching, and disorganized basement membrane, all features associated with neoplasia. In contrast, over-expression of PR-B leads to premature ductal growth arrest and inadequate lobulo-alveolar differentiation (18,19). Moreover the loss of coordinated PR-A and PR-B expression is an early event in carcinogenesis and is evident in pre-malignant lesions (20). A significant proportion of carcinomas expresses a predominance of PR-A isoform and this is associated with poor clinical outcomes: in endometrial cancer, PR-A isoform predominance is significantly more common in tumors of higher grade, indicating an association between PR-A isoform predominance and poor prognosis (21,22).

Although ER and PR are members of different steroid hormone receptor sub-families and recognize distinct hormone response elements, there is considerable biological evidence for cross-talk between the estrogen and progestin hormone receptor signaling pathways. In many cases, progestins suppress the stimulatory effects of estrogens in target cells, for example estrogen increases the expression of both *c-fos* and PR mRNA in uterine cells, and progestins block this effect (23,24) which appear to be mediated via PR, but it is unclear if ER or some other component of the estrogen-ER signaling pathway is the target for repression. Also liganded PR can suppress estradiol-stimulated ER activity, with the

magnitude of repression dependent on the PR isoform, progestin ligand, promoter, and cell type (25). The exact molecular mechanisms regulating Estrogen Receptor  $\alpha$  (ER $\alpha$ ) expression in breast tumors are unclear, but studies suggest that they are partly at the level of transcription (26).

In the present study we examined whether increased PR-B/PR-A ratio could affect response to estradiol in ER $\alpha$  positive breast cancer cells, MCF-7. We demonstrate that PR-B over-expression down regulates ER $\alpha$  mRNA, protein content and gene promoter activity. The investigation of the ER $\alpha$  gene promoter, site directed mutagenesis of half-PRE region in Estrogen Receptor  $\alpha$  promoter, and chromatin immunoprecipitation provide evidence for an active mechanism by which the PR-B is able to negatively regulate expression of Estrogen Receptor  $\alpha$  gene by recruitment to the Estrogen Receptor  $\alpha$  promoter of a corepressor complex containing NCoR but not SMRT and DAX1, causing hypoacetylation of the chromatin and displacement of RNA polymerase II. Furthermore PR-B ablation studies using small interfering RNA demonstrated the critical involvement of PR-B isoform in the recruitment on Estrogen Receptor  $\alpha$  promoter in MCF-7 PR-B overexpressing cells.





## **MATERIAL AND METHODS**

### ***Materials***

Dulbecco's modified Eagle's medium (DMEM/F-12), L-glutamine, Eagle's non-essential amino acids, penicillin, streptomycin, calf serum, bovine serum albumin, and phosphate-buffered saline were purchased from Eurobio (Les Ullis Cedex, France). Triazol reagent was obtained from Invitrogen (Carlsbad, CA), and FuGENE 6 was from Roche Applied Science (Indianapolis, Indiana). Taq DNA polymerase, 100-bp DNA ladder, dual luciferase kit, thymidine kinase and *Renilla* luciferase plasmid were provided by Promega (Madison, Wisconsin). Aprotinin, leupeptin, phenylmethylsulfonyl fluoride (PMFS), and sodium orthovanadate were purchased from Sigma (Milan, Italy)

Antobodies used in this study anti-ER $\alpha$ , anti-cyclin D1, anti-GAPDH, anti-Pol II, anti-NCoR, anti SMRT, anti DAX-1 were purchased from Santa Cruz Biotechnology (Santa Cruz, CA); anti-acetyl histone4-K16, from Upstate (Lake Placid, NY); anti PR, from NeoMarkers (Labvision, Fremont, CA). Salmon sperm DNA was from Santa Cruz Biotechnology (Santa Cruz, CA). Biotinylated horse anti-goat IgG and ABC complex/horseradish peroxidase were provided by Vector Laboratories (Burlingame, CA). Chromogen, 3-diaminobenzidine tetrachloride dihydrate, was purchased from Bio-Optica, MI. ECL System was

purchased from Amersham Pharmacia (Buckinghamshire, UK). VCX500 ultrasonic processor was provided by Sonics (Newtown, CT).

**Plasmid** Firefly luciferase reporter plasmid XETL is a construct containing an estrogen-responsive element from the *Xenopus* vitellogenin promoter (27). The wild type human ER $\alpha$  expression vector (HEGO) consists of the full-length ER $\alpha$  cDNA fused with the SV40 early promoter and expressed in the pSG5 vector (28). The full-length PR-B expression plasmid consists of the full-length PR-B cDNA fused with the SV40 early promoter and expressed in the pSG5 vector (gift from Dr. D. Picard University of Genève, Switzerland). The PR DNA-binding mutant (PR-B DBD mutant C587A) expression plasmids was previously described by Faivre et al (29) (gift from Dr. C. Lange University of Minnesota Cancer Center, Minneapolis-USA). The full-length PR-A expression plasmid was previously described by Kastner *et al.* (30). The deletion fragments of the ER $\alpha$  gene promoter are described by L.A. deGraffenried (31) (gifts from Dr. SAW Fuqua, Baylor College of Medicine, Houston, Texas-USA). The Renilla luciferase expression vector pRL-TK (Promega, Milan, Italy) was used as a transfection standard.

**Cell Culture** Wild type human breast cancer MCF-7 cells were gifts from Dr. B. Van der Burg (Utrecht, The Netherlands). Human uterine cervix adenocarcinoma (HeLa) cells were obtained from the ATCC (Manassas, VA). MCF-7 and HeLa cells were maintained in DMEM/F-12 medium containing 5% fetal calf serum (5% FCS), 1% L-glutamine, 1% Eagle's nonessential amino acids, and 1 mg/ml

penicillin/streptomycin in a 5% CO<sub>2</sub> humidified atmosphere. Cells were cultured in phenol red-free DMEM, 0.5% bovine serum albumin (0,5% BSA), and 2 mM L-glutamine (serum-free medium, SFM), for 48 h before each experiment. Hormone stimulation was performed in DMEM containing 5% charcoal-treated fetal calf serum to reduce the endogenous steroid concentration. (32)

***Proliferation assays*** For quantitative proliferation assays, 10,000 cells were seeded in 24-well plates in regular growth medium. Cells were washed once they had attached and then incubated in medium containing 5 % charcoal-treated FCS with the indicated treatments; medium was renewed every 2 days (with treatments) and cells were trypsinized and counted in a hemocytometer on day 6.

***Reverse Transcription-PCR Assay*** Total cellular RNA was extracted from MCF-7 cells using Triazol reagent as suggested by the manufacturer. 2 µg of total RNA were reverse-transcribed using 200 IU of Moloney murine leukemia virus-reverse transcriptase in a reaction volume of 20 µl (0.4 µg of oligo(dT), 0.5 mM deoxy-NTP, and 24 IU of RNasin) for 30 min at 37 °C, followed by heat denaturation for 5 min at 95 °C. 2 µl of cDNA were then amplified with a pair of 5' and 3' 50-pmol primers for either ER α or pS2 or acidic ribosomal phosphoprotein P0 (36B4) by using 2 units of Taq DNA polymerase. The primers used are as follows: ERα forward, 5'-GGAGACATGAGAGCTGCCA-3', and reverse, 5'-CCAGCAGCATGTCGACGATC-3', to amplify a 438-bp fragment; pS2 forward, 5'-TTCTATCCTAATACCATCGACG-3', and reverse, 5'-

TTTGAGTAGTCAAAGTCAGAGC-3' to amplify a 210-bp fragment; cyclin D1 5'-TCTAAGATGAAGGAGACCATC-3', and reverse, 5'-GCGGTAGTAGGACAGGAAGTTGTT-3', to amplify a 400 bp fragment; PR-B forward, 5'-TAGTGAGGGGGCAGTGGAAC-3', and reverse, 5'-AGGAGGGGGTTTCGGGAATA-3' to amplify a 442 bp fragment; 36B4, forward, 5'-CTCAACATCTCCCCCTTCTC-3', and reverse, 5'-CAAATCCCATATCCTCGTCC-3', to amplify a 408-bp fragment. PCR amplification was performed in 10 mM Tris-HCl, pH 9.0, containing 0.1% Triton X-100, 50 mM KCl, 1.5 mM MgCl<sub>2</sub>, and 0.25 mM each dNTP. Equal amounts of PCR product were electrophoresed on a 1% agarose gels and visualized by ethidium bromide staining. To check out for the presence of DNA contamination, a reverse transcription-PCR was performed without Moloney murine leukemia virus-reverse transcriptase (negative control).

**Western blotting** Cells were grown in 10-cm dishes, exposed to ligands, and then lysed in 500  $\mu$ L of 50 mmol/L HEPES (pH 7.5), 150 mmol/L NaCl, 1.5 mmol/L MgCl<sub>2</sub>, 1 mmol/L EGTA, 10% glycerol, 1% Triton X-100, 1% SDS, a mixture of protease inhibitors containing 1 mmol/L aprotinin, 20 mmol/L phenylmethylsulfonyl fluoride, and 200 mmol/L sodium orthovanadate. Protein concentration was determined using Bradford reagent according to the manufacturer's recommendations (Sigma-Aldrich). Equal amounts of whole protein extract were resolved on a 10% SDS-polyacrylamide gel, transferred to a nitrocellulose membrane (Amersham Biosciences, Milan, Italy), probed overnight



at 4°C with the antibody as indicated in the figure legends, followed by anti-goat or anti-mouse horseradish peroxidase-conjugated antibody and then revealed using the enhanced chemiluminescence system (Amersham Biosciences).

***Transfections and luciferase assays*** Cells ( $1 \times 10^5$ ) were plated into 24-well dishes with 500  $\mu$ L of regular growth medium per well the day before transfection. The medium was replaced with that lacking serum on the day of transfection, which was done using Fugene 6 reagent as recommended by the manufacturer (Roche Diagnostics, Milan, Italy) with a mixture containing 0.5  $\mu$ g of reporter plasmid, alone or in combination with ER $\alpha$  and/or PR-B or mDBD PR or PRA expression plasmids as indicated in the figure legends, and 5 ng of pRL-TK. Upon transfection, the serum-free medium containing the indicated treatment was renewed and then cells were treated for 24 h. Luciferase activity was measured with the Dual Luciferase kit (Promega) according to the manufacturer's recommendations. Firefly luciferase values were normalized to the internal transfection control provided by the *Renilla* luciferase activity.

For whole cell extracts and chromatin immunoprecipitation (ChIP) assay, cells shifted to serum-free medium for 24 h were plated on a 10-cm dish and transfected after 24 h using the FuGENE 6 reagent, with an appropriate amount of the various plasmids as indicated in the figure legends. Cells were treated as indicated in the figure legends or left untreated in DMEM containing 5% charcoal-treated fetal calf serum for 48 h.

Empty vector was used to ensure that DNA concentrations were constant in each transfection.

***Lipid-Mediated Transfection of siRNA Duplexes*** RNA oligonucleotides directed against PR-B or NCoR, were purchased from Invitrogen (Carlsbad, CA), and transfection was performed as described previously (33,34). Briefly, MCF-7 cells were plated in medium lacking antibiotics 2 days before transfection so that they would be 30–50% confluent at the time of transfection. On the day of transfection (d 0), siRNA duplexes (200 nM) and Lipofectamine (8  $\mu$ l per well; Invitrogen) were diluted in Optimem (Invitrogen) according to the manufacturer's instructions. The medium was changed 24 h later (d 1). On day 5, cells were placed in serum-free medium and treated as indicated in the figure legends.

***Site-directed mutagenesis*** Was performed on the D fragment of the ER $\alpha$  promoter using the QuikChange mutagenesis kit (Stratagene) following the manufacturer's instructions. The mutagenesis sense and antisense primers were designed using the Stratagene web site.

The sequences for the primers were: 5'-AGCAGGGAGATGAGGATTGCTGGaagTCCATGGGGGTATGT-3' and 5'-ACATACCCCATGGACTtcAGCAATCCTCATCTCCCTGCT-3'. The plasmids were then sequenced to confirm the mutation of the desired site.

**Chromatin immunoprecipitation (ChIP) Assay** We followed the ChIP methodology described by Shang *et al.* (35) with minor modifications. MCF-7 cells were grown in 100-mm plates. 90% confluent cultures were shifted to serum-free medium for 24 h and then treated left untreated as indicated in the figure legends. Following treatment, the cells were washed twice with phosphate-buffered saline and cross-linked with 1% formaldehyde at 37 °C for 10 min. Next the cells were washed twice with phosphate-buffered saline at 4 °C, collected, and resuspended in 200 µl of Lysis buffer (1% SDS, 10 mM EDTA, 50 mM Tris-Cl, pH 8.1) and left on ice for 10 min. Then the cells were sonicated four times for 10 s at 30% of maximal power and collected by centrifugation at 4 °C for 10 min at 14,000 rpm. Supernatants were collected and diluted in 1.3 ml of IP buffer (0.01% SDS, 1.1% Triton X-100, 1.2 mM EDTA, 16.7 mM Tris-Cl, pH 8.1, 16.7 mM NaCl) followed by immunoclearing with 80 µl of sonicated salmon sperm DNA/protein A-agarose for 1 h at 4 °C. The precleared chromatin was immunoprecipitated for 12 h either with PR, RNA Pol II, acetyl histone4-K16, NCoR, SMRT and DAX-1 antibodies or with normal goat IgG as the negative control. After that, 60 µl of salmon sperm DNA/protein A-agarose was added, and precipitation was continued for 2 h at 4°C. After pelleting, precipitates were washed sequentially for 5 min with the following buffers: Wash A (0.1% SDS, 1% Triton X-100, 2 mM EDTA, 20 mM Tris-Cl, pH 8.1, 150 mM NaCl), Wash B (0.1% SDS, 1% Triton X-100, 2 mM EDTA, 20 mM Tris-Cl, pH 8.1, 500 mM NaCl), and Wash C (0.25 M LiCl, 1% Nonidet P-40, 1% sodium deoxycholate, 1

mM EDTA, 10 mM Tris-Cl, pH 8.1), and then twice with TE buffer (10 mM Tris, 1 mM EDTA). The immune complexes were eluted with elution buffer (1% SDS, 0.1 M NaHCO<sub>3</sub>). The eluates were reverse cross-linked by heating at 65 °C for 12 h and digested with 0.5 mg/ml proteinase K at 45 °C for 1 h. DNA was obtained by phenol and phenol/chloroform extractions. 2 µl of 10 mg/ml yeast tRNA were added to each sample, and DNA was precipitated with ethanol for 12 h at -20 °C and resuspended in 20 µl of TE buffer. 5 µl of each sample were used for PCR with the following ER α promoter primers: forward 5'-ACGTTCTTGATCCAGCAGGGTA-3' and reverse 5'-ACCTGCCAAATTATATGCAAATGGCAG-3' containing the half-PRE site and forward 5'-GTGGCCATTGTTGACCTACAG-3' and reverse 5'-CTGTAGGTCAACAATGGCCAC-3' upstream the half-PRE site. The amplification products, obtained in 30 cycles, were analyzed in a 2% agarose gel and visualized by ethidium bromide staining. The negative control was provided by PCR amplification without DNA sample.

### ***Statistical Analysis***

Each datum point represents the mean SE of three different experiments. Data were analyzed by ANOVA test using the STATPAC computer program.



## RESULTS

### **Progesterone decreases estradiol induced cell proliferation in MCF-7 breast cancer cells.**

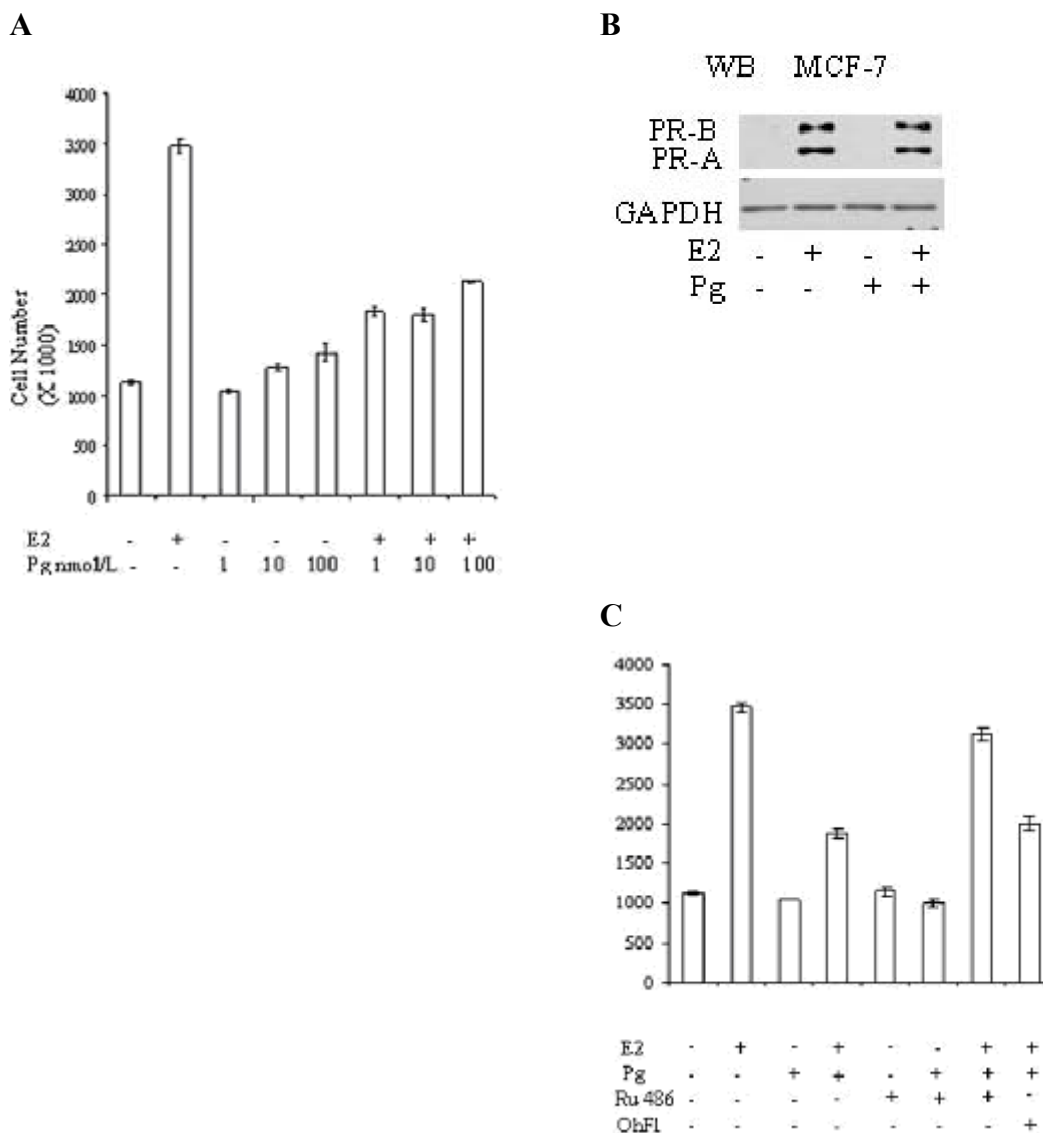
Reports about the effects of progestins on cell proliferation are contradictory and fuel the debate around the actions on breast tissue.

In order to understand the eventual ability of progestins to interfere with cell proliferation stimulated by estradiol (E2), we evaluated the effects of different concentrations of progesterone (Pg 1 nmol/L, Pg 10 nmol/L , Pg 100 nmol/L ) on the proliferation of MCF-7 cells which express low levels of Progesterone Receptor (PR) inducible by estradiol treatment.

The results obtained revealed that after 6 days, as expected 10 nmol/L E2 treatment significantly increased the cell number, Pg alone at all tested concentrations had no significant effect compared with the control experimental conditions, however 10 nmol/L Pg caused the maximal inhibitory effect (48 %) on the E2 induced cell proliferation (Fig. 1A). These data evidenced that the action of Pg in inhibiting cell proliferation is likely to be due on prior exposure to E2 inducing PR-A and PR-B expression as evidenced by western blotting analysis (Fig. 1B) RU 486 is an antagonist of progestins action in human and binds with high affinity to the corresponding receptors (36). In our experimental conditions 1  $\mu$ mol/L RU 486 did not affect the growth of the MCF-7 cells under control treatment conditions (Fig. 1C).

However, in combination with 10 nmol/L E2 and 10 nmol/L Pg significantly reduced the inhibitory effect due to Pg treatment on cell proliferation, suggesting that these inhibitory action is mediated by PR.

It is also interesting to note that 1  $\mu$ mol/L OH-FI, an androgen receptor (AR) antagonist, did not modify the inhibitory action of Pg on E2 stimulated proliferation, indicating that AR is not involved.



**Figure 1. Progesterone decreases the estradiol induced cell proliferation in MCF-7 breast cancer cells.** A, MCF-7 cells were treated with vehicle (–) or Estradiol 10 nmol/L (E2) and/or increasing amount of Progesterone (1 nmol/L, 10 nmol/L, 100 nmol/L) [Pg] in medium containing 5% charcoal-stripped FBS (medium was refreshed and treatments were renewed every 2 d) and then counted on day 6. B, Immunoblot analysis of PR. MCF-7 cells were treated with vehicle (–) or Estradiol 10 nmol/L (E2) and/or Progesterone 10 nmol/L (Pg) for 48 hours. GAPDH levels were used as a loading control. C, Cells cultured in the experimental conditions described in A, were also treated with vehicle (–) or Estradiol 10 nmol/L (E2) and/or Progesterone 10 nmol/L (Pg) in combination with RU 486 1  $\mu$ mol/L or OHF1 1  $\mu$ mol/L and counted on day 6. Columns, mean of three independent experiments done in triplicate; bars, SD.



**Progesterone Receptor B over-expression represses the E2/ ER $\alpha$  transcriptional activity.**

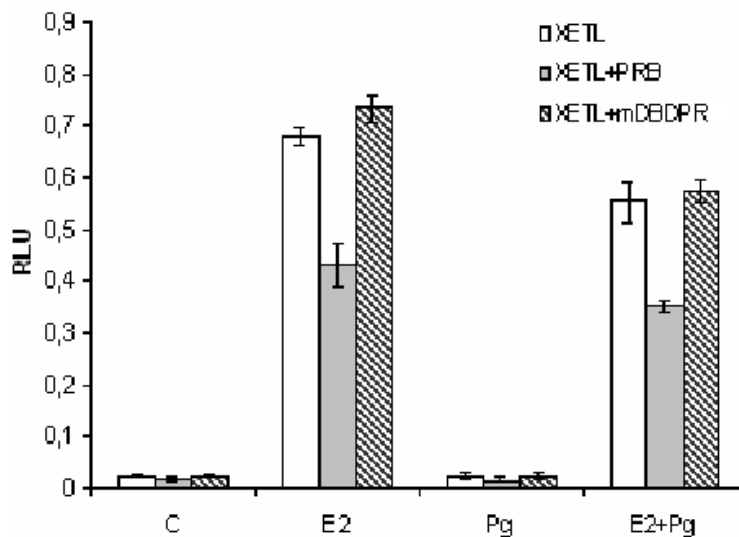
The PR system is characterized by several complexities; one is that PR is an estrogen receptor dependent gene product (3) and in MCF-7 cells the action of Pg requires priming treatment of E2 inducing PR-A and PR-B isoforms expression.

To study whether changes in relative expression levels of PR-B and A isoforms we analyze the specific effects of PR-B over-expression on genomic activity induced by estradiol treatment. To this aim a luciferase reporter plasmids containing a consensus estrogen-responsive element (ERE) sequence was transiently co-transfected into MCF-7, in presence or absence of expression plasmids encoding the full length PR-B. Cells were treated with 10 nmol/L E2 and/or 10 nmol/L Pg, as indicated in Fig. 2A for 18 h. Treatment with E2 alone resulted, as expected, a substantial increase in luciferase activity and interestingly PR-B transient over-expression repressed this effect as much as 36%; moreover when the cells were co-treated with E2 and Pg the inhibitory effect was of 49%. Conversely, over-expression of a PR mutant with a disrupted DBD (Cys587 to Ala) cannot mediate transcriptional responses and did not affect luciferase activity in the same experimental conditions.

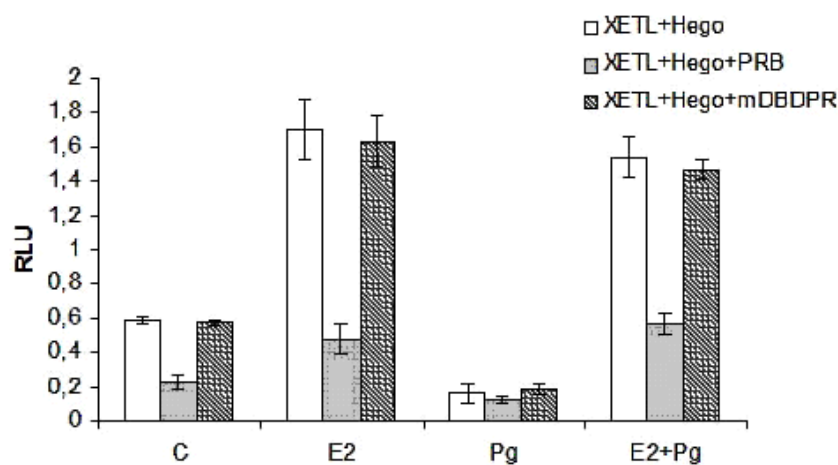
To confirm these findings, the same experiments were carried out in HeLa cells expressing ectopic ER $\alpha$ . As shown in Fig. 2B, PR-B co-expression repressed the 10 nmol/L E2 induced luciferase activity by 75% and 73 % when cells were co-treated

with E2 and 10 nmol/L Pg; over-expression of a PR mutant with a disrupted DBD had no inhibitory effects.

A



B



**Figure 2. Progesterone Receptor B over-expression represses the E2/ ER $\alpha$  transcriptional activity.** ERE luciferase reporter assay. MCF-7 cells (A) and HeLa cells expressing ectopic ER $\alpha$  (B), were transiently co-transfected with XETL in the presence or absence of full-length PR B expression plasmid or mDBD PR expression plasmid. After transfection cells were treated with vehicle (–) or Estradiol 10 nmol/L (E2) and/or Progesterone 10 nmol/L (Pg), harvested after 18 h, then luciferase activities were determined. *Columns*, mean of luciferase activities observed in three independent experiments. *bars*, SD.

### **Progesterone Receptor B over-expression down-regulates ER $\alpha$ protein levels and mRNA**

After demonstrating the inhibitory effects of PR-B over-expression on the estrogen receptor induced signal we examined whether this action might be due to a decrease in the expression levels of ER $\alpha$  protein.

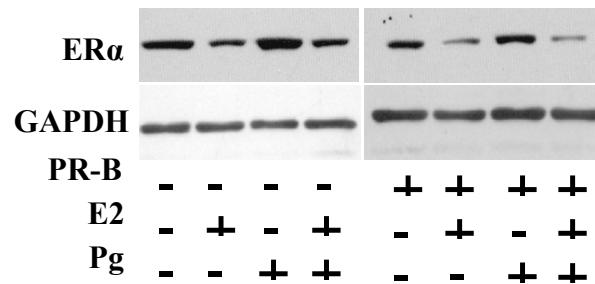
Immunoblot analysis of lysates from MCF-7 cells treated with 10 nmol/L E2 and/or 10 nmol/L Pg for 48 h, shown that as expected E2 treatment repressed the ER  $\alpha$  expression levels while Pg alone had no effect (Fig 3A).

Moreover, in MCF-7 cells over-expressing PR-B, a marked repression of ER $\alpha$  content was observed in control experimental conditions in absence of treatment, as well as after ligands exposure compared with the examined conditions in absence of exogenous PR-B. These effects were not observed in cells overexpressing PR-A.

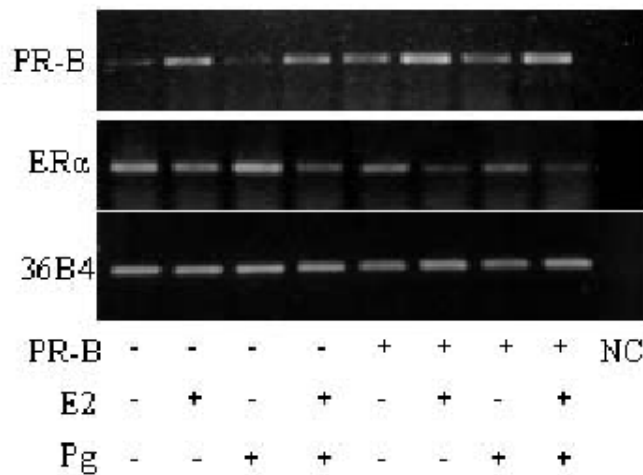
Previous studies provide strong evidences that ER $\alpha$  expression in breast cancer is regulated at both the transcriptional as well as the translational level. (38)

To further investigate the molecular basis of the PR-B mediated regulation of ER $\alpha$  we examined the effect of PR-B over-expression on ER $\alpha$  mRNA levels in MCF-7 cells treated for 48 h with 10 nmol/L E2 and/or 10 nmol/L Pg. As shown in Fig. 3B, ER $\alpha$  mRNA was reduced by E2 treatment compared to control, moreover the over-expression of PR-B further decreased ER $\alpha$  mRNA compared with the examined conditions in absence of exogenous PR-B.

A



B



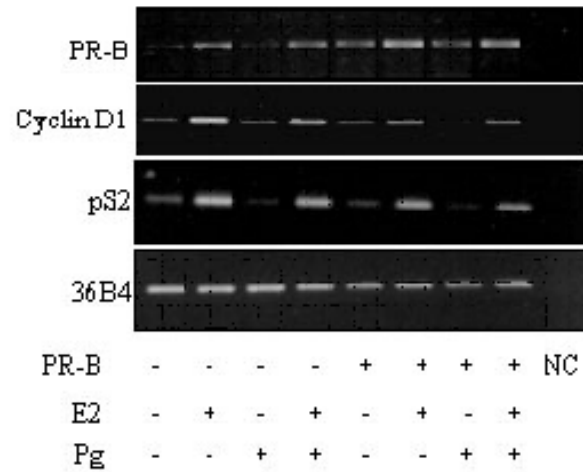
**Figure 3. Progesterone Receptor over-expression decreases the ER  $\alpha$  protein and mRNA levels.** (A) Immunoblot analysis of ER $\alpha$ . MCF-7 cells and MCF-7 transient over-expressing PR-B were treated with vehicle (-) or Estradiol 10 nmol/L (E2) and/or Progesterone 10 nmol/L (Pg) for 48 hours. GAPDH levels were used as a loading control. (B) Semiquantitative RT-PCR. mRNA expression of ER $\alpha$  in MCF-7 cells and MCF-7 cells transient over-expressing PR-B. The expression of ER $\alpha$  was evaluated in cells treated with vehicle (-) or Estradiol 10 nmol/L (E2) and/or Progesterone 10 nmol/L (Pg) for 48 h; the housekeeping gene 36B4 was determined as a control.

**PR-B overexpression decreases the levels of ER $\alpha$ -regulated genes in MCF-7 cells.**

The effects of E2 hormone are mediated by ER through its gene regulation activities. To determine whether Pg could inhibit the effect of E2 by modifying the transcriptional activity of ER $\alpha$ , the mRNA of well known estrogen target genes, cyclin D1 and pS2, were studied. (Fig. 3C) As expected after 48 h treatment, 10 nmol/L E2 markedly increased the mRNA levels of the estrogen-regulated cyclin D1 and pS2 in MCF-7 cells. However, the induction of cyclin D1 levels was impaired by 20% after concomitant 10 nmol/L E2 and 10 nmol/L Pg treatment; similarly, E2 induced pS2 expression was impaired by 15%. These data evidenced that the inhibitory action of Pg in MCF-7 cells is concomitant to high PR expression levels due to prior exposure to E2.

To analyze whether changes in relative expression levels of PR B and A isoforms causing the above described down-regulatory effect on ER $\alpha$  protein and mRNA levels, could affect the expression of estrogen regulated genes, plasmids encoding full length PR-B were transiently transfected into MCF-7 cells. After 48 h of ligand treatment, cells were harvested and subjected to semiquantitative RT-PCR experiments comparing mRNA levels encoding cyclin D1, and pS2 after standardization with a housekeeping gene encoding the ribosomal protein 36B4. As depicted in Fig. 3C, PR-B over-expression, represses both cyclin D1 and pS2 mRNA levels.

C



**Figure 3. PR-B overexpression decreases the levels of ER $\alpha$ -regulated genes in MCF-7 cells.** (C) mRNA expression of IRS-1 Cyclin D1 and pS2 and PR-B in MCF-7 cells and MCF-7 transient overexpressing PR-B. Cells were treated with vehicle (–) or Estradiol 10 nmol/L (E2) and/or Progesterone 10 nmol/L (Pg) for 48 hours. The housekeeping gene 36B4 was determined as a control.

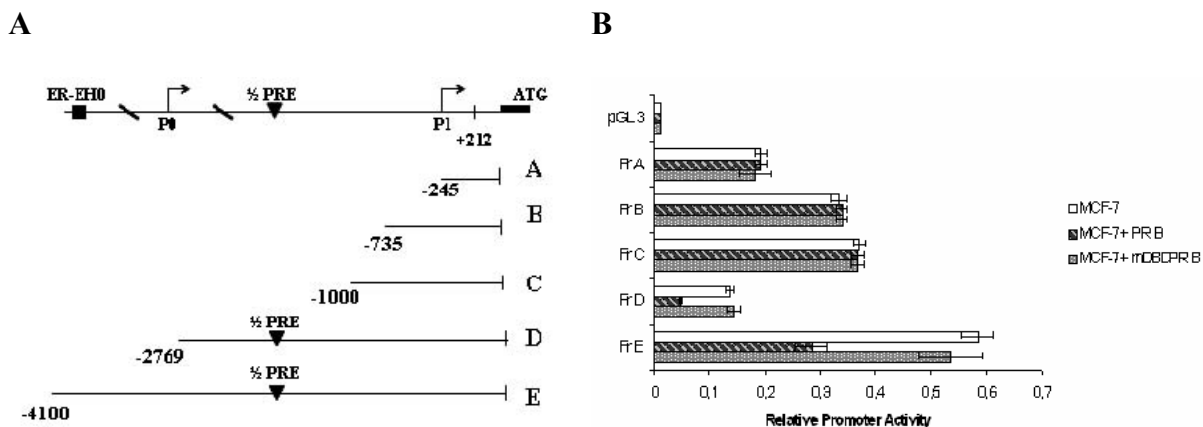
**The PR-B overexpression mediates the down-regulation of ER  $\alpha$  promoter activity by the region between  $-2769$  bp to  $-1000$  bp and mutation of the half-PRE site in ER $\alpha$  promoter abrogates the PR-B induced inhibitory effect.**

To analyze whether PR-B could interfere with ER $\alpha$  gene transcription the promoter of ER $\alpha$  was investigated first with a bioinformatics approach using the NCBI Genome data base ([www.ncbi.nlm.nih.gov](http://www.ncbi.nlm.nih.gov)). The region examined in this study covered from  $-4100$  bp to  $+212$  bp and in this portion one half-PRE ( $-1757$  bp to  $-1752$  bp) was identified as putative effector of PR action.

To evaluate whether this region was involved in Progesterone Receptor mediated down-regulation of ER $\alpha$  expression, five overlapping ER $\alpha$  promoter deletion constructs,  $-245$  bp to  $+212$  bp (fragment A),  $-735$  bp to  $+212$  bp (fragment B),  $-1000$  bp to  $+212$  bp (fragment C),  $-2769$  bp to  $+212$  bp (fragment D), and  $-4100$  bp to  $+212$  bp (fragment E), all relative to the first transcriptional start site, sub-cloned into the luciferase reporter vector pGL3-basic, were analysed as depicted in Fig. 4 A. The five fragments were transiently transfected into the ER positive MCF-7 cells. Data are shown as relative promoter activity in luciferase units. Consistent with previous ER $\alpha$  promoter studies of L.A. de Graffenried et al. 2002, fragment A exhibited high levels of activity compared to vector alone (Fig. 4B), the activity of fragments B and C was slightly increased relative to A and the activity of fragment D significantly decreased relative to the activity seen with A, the activity of fragment E increased. In the same experimental conditions PR-B over-expression had no effect on



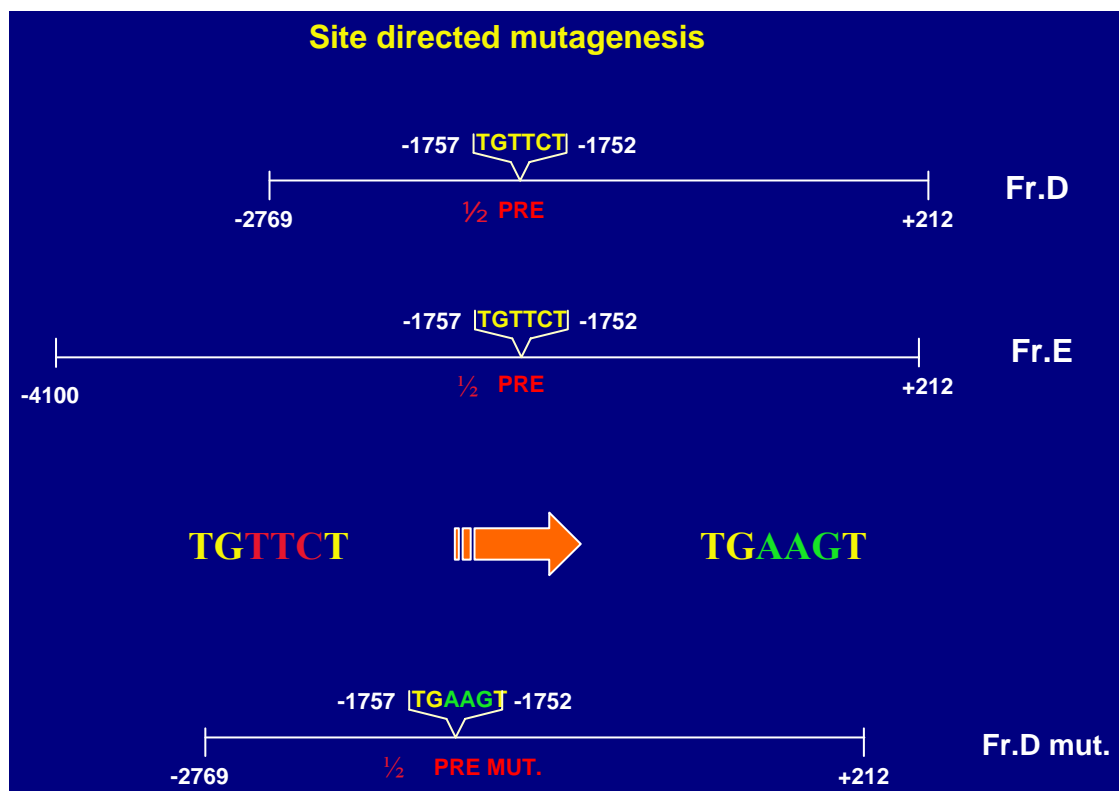
the promoter activity of A, B and C lacking the half-PRE site; in contrast was able to significantly and reproducibly reduce by 64 % the activity of D and by 50 % the activity of E tested, indicating that the region between -2769 bp to -1000 bp appeared to be responsible for the PR-B mediated down-regulation. These inhibitory effects were not observed co-transfecting an expression plasmid encoding PR mutant with a disrupted DBD (Cys587 to Ala) into MCF-7 cells.



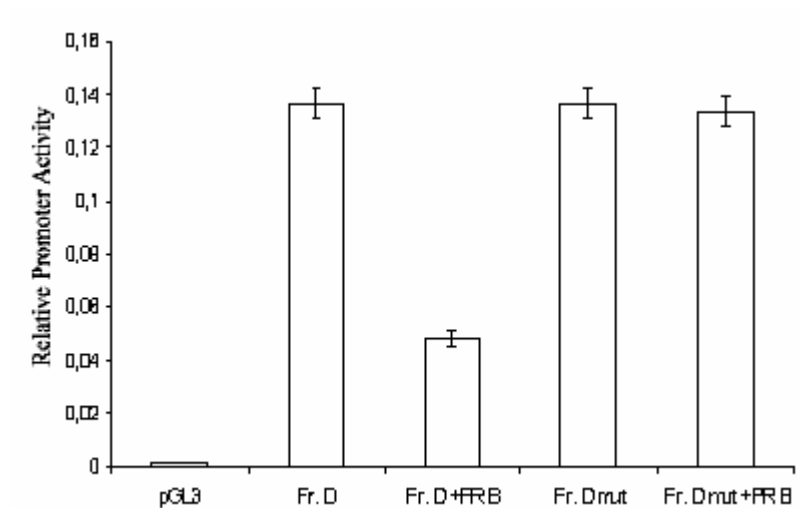
**Figure 4. ER  $\alpha$  promoter activity is down-regulated by PR-B over-expression and mutation of the half-PRE abrogates this effect** *A*, Schematic representation of deletion fragments of the ER $\alpha$  gene promoter. Fragment coordinates are expressed relative to the primary transcription start site *B*, Promoter activity of the ER  $\alpha$  5'-flanking region. Constructs described in *A* were transiently co-transfected in MCF-7 cells in presence or absence of full-length PR-B expression plasmid or mDBD PR expression plasmid. After 24 h, cells were harvested, then luciferase activities were determined. Columns, mean of luciferase activities observed in three independent experiments. *bars*, SD.

### Site directed mutagenesis reveals a role for the half-PRE in the fragment D of the ER $\alpha$ promoter

To evaluate the role of the half-PRE present in the fragment D of ER  $\alpha$  gene promoter (-1757 bp to -1752 bp), we used site-directed mutagenesis to alter this site. We changed 3 of the half-PRE to ensure that the altered binding site would not be recognizable by the PR. Transient transfections were performed in MCF-7 cells with multiple independent clones containing the desired mutation. We found that all of the half-PRE mutated clones (D mut) were unaffected by PR-B over-expression (Fig. 4 C). These results indicate that the half-PRE element in the estrogen receptor  $\alpha$  promoter is necessary for repression due to PR-B over-expression.



C



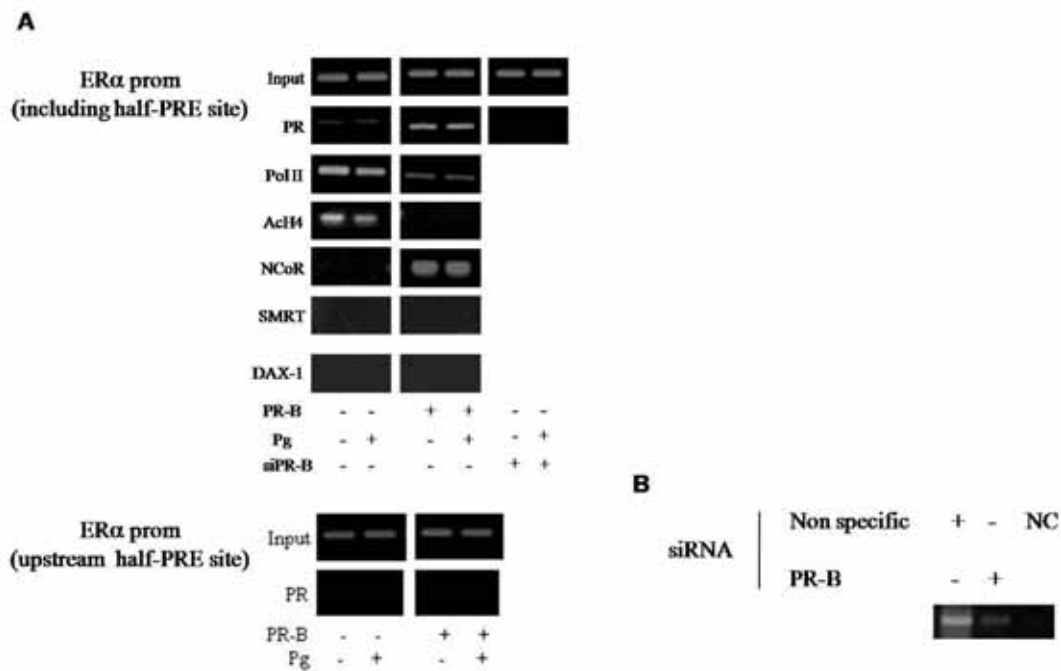
**Figure 4. Site directed mutagenesis reveals a role for the half-PRE in the fragment D of the ER  $\alpha$  promoter** C The half-PRE site in the Fr D promoter construct was mutated using site-directed mutagenesis. The Fr D and Fr D mut promoter constructs were co-transfected into MCF-7 cells, and promoter activity was assessed in the absence or presence of full-length PR-B expression plasmid. Luciferase assay was performed after 24 h. *Columns*, mean of luciferase activities observed in three independent experiments. *bars*, SD.

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### **A corepressor complex containing N-CoR bound PR-B is recruited at the ER $\alpha$ promoter “in Vivo”**

To examine PR recruitment to the ER  $\alpha$  gene promoter *in vivo*, we used ChIP assays. MCF-7 cells overexpressing or not PR-B were treated for 1 h with either control vehicle or 10 nmol/L Pg, after which chromatin was cross-linked with formaldehyde, and protein-DNA complexes were immunoprecipitated with antibodies against PR, RNA polymerase II, acetyl histone H4-K16, NCoR, DAX-1 or SMRT. The PCR primers used encompass the half-PRE site of ER $\alpha$  promoter. Results obtained in MCF-7 PR-B overexpressing cells and depicted in Fig. 5A, show an increased recruitment of PR to the ER $\alpha$  promoter encompassing the half-PRE site in a ligand independent manner, but not to the unrelated region located upstream the half-PRE site. Moreover Histone H4 became deacetylated and RNA polymerase II was released from the ER  $\alpha$  promoter upon PR-B overexpression, indicating that consequently to the PR interaction the chromatin in this region becomes a less permissive environment for transcription. To examine whether corepressor could be recruited by PR bound ER $\alpha$  promoter we tested NCoR, SMRT and DAX-1, previously described as PR corepressors (39) by ChIP assay. Of note NCoR showed hormone independent recruitment to the ER $\alpha$  promoter in MCF7 cells overexpressing PR-B, highlighting a possible new role of this corepressor in the down regulation of ER $\alpha$  expression. SMRT and DAX 1 were not detected in the examined experimental conditions.

To determine which PR isoform mediates the interaction with ER  $\alpha$  promoter, MCF-7 cells were transfected with PR-B siRNAs to achieve knockdown of PR-B proteins (Fig. 5 B). Remarkably the PR recruitment on half-PRE site of ER  $\alpha$  promoter was strongly reduced by PR-B specific siRNA (Fig.5 A) indicating the specific recruitment of PR-B isoform on ER $\alpha$  promoter .



**Figure 5. PR is recruited to the ER $\alpha$  promoter, RNA polymerase II is dismissed.** *A* ChIP was performed on the PR-responsive region of the ER $\alpha$  promoter using various antibodies as indicated. A nonspecific region upstream of the half-ERE site in the ER $\alpha$  promoter was used as negative control. The cells were treated for 45 min with vehicle (–) or Progesterone 10 nmol/L (Pg). Similar results were obtained in multiple independent experiments. *B* Knockdown of PR-B. MCF-7 cells were transfected with siRNA targeted against PR-B. The medium was changed 6 h later and cells were treated with vehicle (–) or Progesterone 10 nmol/L (Pg) for 48 hrs. RNA was isolated and the expression of PR-B was analyzed by semi-quantitative RT-PCR.

## **DISCUSSION**

Approximately 75% of primary breast cancers express ER and more than half of these cancers coexpress PR. ER and PR are considered prognostic factors for early breast cancer therapeutic management although both are weak and lose their prognostic value after long-term follow-up. PR exists as two isoforms, called PR-A and PR-B, which have distinct roles in regulating the effects of progesterone and there is increasing evidence that enhanced PR-A/PR-B ratio is coincident with a major breast cancer growth and progression.(40)

It is well documented that in hormone dependent breast cancer cells, progesterone may inhibit the induction of classically estrogen regulated genes even though the molecular mechanisms underlying these effects remain to be fully elucidated.(41, 42)

In the present study we demonstrated that PR-B isoform overexpression represses Estrogen Receptor  $\alpha$  expression in MCF-7 breast cancer cells.

We first demonstrated that Progesterone Receptor mediates the down-regulatory effect induced by progesterone treatment on estradiol stimulated cell proliferation, since RU486 prevents this inhibitory effect. In addition we show that the overexpression of PR-B produces an inhibition of estrogen induced ERE reporter gene activity due to the down-regulatory effect of PR-B on ER $\alpha$  expression. The same findings were reproduced in Hela cells expressing ectopic ER $\alpha$  and PR-B.



The latter event may explain the reduced expression of classically ER $\alpha$  regulated genes as pS2 and Cyclin D1 upon PR-B overexpression.

In order to clarify the molecular mechanisms through which PR-B may interfere with ER $\alpha$  gene transcription we explored the ER $\alpha$  promoter sequence. We identify the existence of a PRE element as a putative effector of the inhibitory action of PR-B on ER $\alpha$  expression. Utilizing five deleted constructs we observed that the down-regulatory effect induced by PR-B overexpression on ER $\alpha$  promoter activity was still present but was no longer noticeable in the deleted construct lacking the half PRE site that we considered as putative effector site of PR-B inhibitory action. For instance when we performed site directed mutagenesis in the above mentioned region of ER $\alpha$  promoter, the effects due to PR-B overexpression on ER $\alpha$  promoter activity were completely abrogated. The inhibitory action of PR-B isoform over-expression on ER $\alpha$  promoter transcriptional machinery address most likely the capability of PR-B to recruit in these circumstances some corepressors interfering with ER $\alpha$  gene transcription.

To determine which PR isoform could mediate the interaction of specific coregulators with ER $\alpha$  promoter, siRNA directed against PR-B alone was used. Our ChIP assay results demonstrate the key role of PR-B isoform in the interaction with half-PRE site on ER  $\alpha$  gene promoter.

Although there are far fewer nuclear receptor corepressors, these molecules serve important roles in negatively regulating receptor-dependent gene expression. It was reported that unliganded nuclear receptors as TR and RXR can repress basal

transcription in the absence of their cognate ligands, and this function is mediated, at least in part, by two nuclear proteins, silencing mediator of retinoic acid and thyroid hormone receptor (SMRT) and nuclear receptor corepressor (NCoR). (43) The latter has been recently documented to be also recruited by both estrogen and progesterone receptors in the presence of ligand antagonist. Here we demonstrated that PR-B bound to ER $\alpha$  promoter, in presence or absence of its own natural ligand, is able to recruit a repressor of cell transcriptional machinery as NCoR. Recent data regarding PR transcriptional regulation of different gene promoter mostly address its capability to recruit in those circumstances NCoR. We tested if the latter event may also occur in our experimental conditions during the binding of PR-B isoform to the ER $\alpha$  promoter. Our ChIP experiments proved that among different corepressor molecules able to interact with PR, NCoR is the only present on PR-B/ DNA complex. These results could support a model in which elevated expression levels of PR-B increase the interaction of the receptor with NCoR on the half-PRE site of ER $\alpha$  promoter, an event incompatible with PR-coactivator interactions.

In conclusion we highlight the importance of regulated expression of PR-A and PR-B for cellular properties in breast cancer cells and we have provided a new insight into the molecular mechanism through which PR-B mediates the antiestrogenic effect in breast cancer cells. We think that the inhibition of Estrogen Receptor  $\alpha$  by PR-B is a critical regulatory pathway in ER-positive cells, and we can speculate that deregulation of this repression in breast cancer may

have dramatic effects such as the promotion of transformation and metastasis. Our combined results suggest that this antiestrogenic mechanism is of therapeutic relevance for breast cancer treatment.

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# Human sperm express a functional androgen receptor: effects on PI3K/AKT pathway

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**BACKGROUND:** Results from mice lacking the androgen receptor (AR) showed that it is critical for the proper development and function of the testes. The aim of this study was to investigate whether a functional AR is present in human sperm. **METHODS:** The expression of AR and its effects on sperm were evaluated by RT-PCR, Western Blot, Immunocytochemistry, PI3Kinase and DNA laddering assays. **RESULTS:** We showed in human sperm that AR is located at the head region. Dihydrotestosterone (DHT), in a dose-dependent manner, leads to the rapid phosphorylation of the AR on tyrosine, serine and threonine residues and this effect was reduced by the AR antagonist hydroxyflutamide (OH-Flut). The effects of AR were evaluated on the phosphoinositide-3 kinase/protein kinase B (PI3K/AKT) pathway. Specifically, 0.1 and 1 nM DHT stimulated PI3K activity, whereas 10 nM DHT decreased PI3K activity and levels of p-AKT S473 and p-AKT T308, p-BCL2, and enhanced phosphatase and tensin homologue (PTEN) phosphorylation. In addition, 10 nM DHT was able to induce the cleavage of caspases 8, 9 and 3 and cause DNA laddering, and these effects were reversed either by casodex or OHFlut. By using wortmannin, a specific PI3K inhibitor, the cleavage of caspase 3 was reproduced, confirming that in sperm the PI3K/AKT pathway is involved in caspase activation. **CONCLUSIONS:** Human sperm express a functional AR that have the ability to modulate the PI3K/AKT pathway, on the basis of androgen concentration.

*Keywords:* androgen receptor; androgens; human sperm; male reproduction; PI3K/AKT

## Introduction

A functional androgen receptor (AR) is required for male embryonic sexual differentiation, pubertal development and regulation of spermatogenesis in mammals. The role of AR during spermatogenesis has been the subject of intense interest for many years (Collins *et al.*, 2003). Several findings have shown that AR function is required for the completion of meiosis and the transition of spermatocytes to haploid round spermatids (De Gendt *et al.*, 2004). Studies of androgen withdrawal and disruption of AR activity, either by surgical, chemical or genetic means, have demonstrated that spermatogenesis rarely proceeds beyond meiosis. In all of these model systems, very few round and even fewer elongated spermatids are observed, as clearly demonstrated in a previous study (Yeh *et al.*, 2002). However, the mechanisms by which androgens regulate male fertility are not fully understood and the sites of androgen action within the male reproductive system are not yet resolved.

Whereas few studies have raised the intriguing possibility that some germ cells may exhibit immunoreactive AR (Kimura *et al.*,

1993; Vornberger *et al.*, 1994), other reports point to Sertoli cells or Leydig cells or peritubular/myoid cells as the exclusive androgen target cells in the testis (Ruizeveld de Winter *et al.*, 1991; Iwamura *et al.*, 1994; Goyal *et al.*, 1996; Suarez-Quian *et al.*, 1999). Recently, the presence of the AR in human sperm was demonstrated by western blot and by immunofluorescence assay (Solakidi *et al.*, 2005).

It is generally accepted that androgens bind to intracellular ARs resulting in mRNA and protein synthesis (McPhaul and Young, 2001). Nevertheless, rapid responses to androgens have been observed in different tissues, that cannot be explained by involvement of mRNA and protein synthesis (Peterziel *et al.*, 1999; Castoria *et al.*, 2004). These rapid, nongenomic effects are also seen for other steroid hormones (Cato *et al.*, 2002) and their importance as a complementary route for cell regulation has recently become evident. Different nuclear receptors (Calogero *et al.*, 2000; Aquila *et al.*, 2004) have been found to be present in human spermatozoa, regulating cellular processes through nongenomic mechanisms. This may

represent an exclusive modality of action in spermatozoa since they are apparently transcriptionally inactive cells.

In addition to stimulating cell growth, androgens and/or AR play important roles in the promotion of cell apoptosis (Heisler *et al.*, 1997; Olsen *et al.*, 1998; Shetty *et al.*, 2002; King *et al.*, 2006). The term apoptosis defines programmed cell death, which is executed by the activation of caspases, a family of cytoplasmic cysteine proteases (Cohen, 1997) through two major pathways: the intrinsic and the extrinsic. The intrinsic pathway involves the cell sensing stress that triggers mitochondria-dependent processes, resulting in cytochrome *c* release and activation of caspase 9 (Olson and Kornbluth, 2001). The extrinsic pathway involves the final cleavage of caspase 8 (Schulze-Osthoff *et al.*, 1998). Both caspases 8 and 9 can be directly regulated through protein phosphorylation from protein kinase B (AKT) (Cardone *et al.*, 1998; Shim *et al.*, 2004). The phosphoinositide-3 kinase (PI3K) signalling pathway is an important intracellular mediator of cell survival and antiapoptotic signals (Parsons, 2004). PI3K activation leads to production of 3'-phosphoinositide second messengers, such as phosphatidylinositol 3,4,5-trisphosphate, which activate a variety of downstream cell survival signals. Accumulation of phosphatidylinositol 3,4,5-trisphosphate at the membrane recruits a number of signalling proteins containing pleckstrin homology domains, including AKT. On recruitment, AKT becomes phosphorylated and activated and exerts its antiapoptotic activity through inactivation of proapoptotic proteins. In addition, the PI3K pathway has also been shown to be negatively regulated by phosphatase and tensin homologue (PTEN), a lipid phosphatase that cleaves the D3 phosphate of the second messenger phosphatidylinositol 3,4,5-trisphosphate (Maehama and Dixon, 1998; Wu *et al.*, 1998). Recently in fibroblasts, it has been demonstrated that AR mediates androgen nongenomic function, and that androgen activates PI3K/AKT through the formation of a triple complex between AR, the regulatory subunit p85 of PI3K (PIK3R1) and SRC tyrosine kinase. Indeed, this interaction is dependent on androgen concentration, and a particularly high androgen concentration will dissociate the AR-SRC tyrosine kinase-PI3K complex (Castoria *et al.*, 2003).

The functional impact of programmed cell death in human sperm is poorly understood (Sakkas *et al.*, 2003). Up to now it has been unclear whether apoptosis in ejaculated spermatozoa takes place in a similar way as in somatic cells or whether spermatozoa, which are thought to have a transcriptionally inactive nucleus, undergo abortive forms of this process (Sakkas *et al.*, 2003). However, sperm constitutively express proteins required to execute apoptosis. Active caspases have been observed predominantly in the postacrosomal region (caspases 8, 1 and 3) and caspase 9 has been particularly localized in the midpiece, associated with mitochondria (Paasch *et al.*, 2004). Moreover, a wide spectrum of cell cytoskeletal proteins and membrane components are also targets of caspase 3 (Paasch *et al.*, 2004), and the proper regulation of the caspase cascade plays an important role both in sperm differentiation and testicular maturity (Said *et al.*, 2004). In addition, caspases have been implicated in the pathogenesis of multiple andrological pathologies such as impaired spermatogenesis, decreased sperm motility, increased levels of

sperm DNA fragmentation, testicular torsion, varicocele and immunological infertility (Said *et al.*, 2004). Further studies are needed to evaluate the full significance of caspases activation in spermatozoa. A direct link between AR and sperm survival has not been investigated previously.

In the present study, we have demonstrated the presence of a functional AR in sperm. It emerges from our data that low androgen concentrations stimulate PI3K activity, which is inhibited at higher levels. Additionally, in the latter circumstance increases in PTEN phosphorylation and cleavages of caspases 8, 9 and 3 were evident.

## Materials and Methods

### Chemicals

PMN Cell Isolation Medium was from BIOSPA (Milan, Italy). Total RNA Isolation System kit, enzymes, buffers, nucleotides 100 bp ladder used for RT-PCR were purchased from Promega (Milan, Italy). Moloney Murine Leukemia Virus (M-MLV) was from Gibco BRL Life Technologies Italia (Milan, Italy). Oligonucleotide primers and TA Cloning kit were made by Invitrogen (Milan, Italy). Gel band purification kit was from Amersham Pharmacia Biotech (Buckinghamshire, UK). DMEM-F12 medium, BSA protein standard, laemmli sample buffer, prestained molecular weight markers, percoll (colloidal PVP coated silica for cell separation), sodium bicarbonate, sodium lactate, sodium pyruvate, dimethyl sulfoxide, anti-rabbit IgG fluorescein isothiocyanate (FITC) conjugated, Earle's balanced salt solution, Hoechst 33 342, steroids and all other chemicals were purchased from Sigma Chemical (Milan, Italy). RPMI 1640 medium was from Life Technologies, Inc. (Gaithersburg, MD) and DMEM (PRF-SFM) was from Eurobio (Milan, Italy). Acrylamide bisacrylamide was from Labtek Eurobio (Milan, Italy). Triton X-100 and Eosin Y was from Farmitalia Carlo Erba (Milan, Italy). ECL Plus western blotting detection system, Hybond<sup>TM</sup> ECL<sup>TM</sup>, [ $\gamma$ -<sup>32</sup>P]ATP and HEPES sodium salt were purchased from Amersham Pharmacia Biotech (Buckinghamshire, UK). Goat polyclonal actin antibody (1–19), monoclonal mouse anti-AR (AR 441) and anti-PIK3R1 antibodies, monoclonal anti-p-tyrosine (PY99), normal mouse serum, peroxidase-coupled anti-rabbit and anti-goat and protein A/G-agarose plus were from Santa Cruz Biotechnology (Heidelberg, Germany). Monoclonal mouse anti-p-SRC tyrosine kinase was from Oncogene (Milan, Italy). Polyclonal rabbit anti-p-serine, anti-p-threonine, anti-p-AKT1/AKT2/AKT3 S473, anti-p-AKT1/AKT2/AKT3 T308, anti-p-BCL2, anti-p-PTEN, anti-caspases (8, 9 and 3) antibodies were from Cell Signaling (Milan, Italy). PY20 (Transduction Laboratories, Lexington, UK), anti-phospho-serine and anti-phospho-threonine Abs were from Zymed Laboratories (San Francisco, CA). Casodex (Cax) was from Astra Zeneca (Milan, Italy) and hydroxy-flutamide (OH-Flut) was from Schering (Milan, Italy). The specific caspases inhibitor Z-VAD-FMK (ZVF) was from R&D Systems (Milan, Italy). PCR 2.1 vector was from Promega and the sequencing was by MWG AG Biotech (Ebersberg, Germany).

### Semen samples and spermatozoa preparations

Semen specimens from normozoospermic men were obtained after three days of sexual abstinence. The samples were ejaculated into sterile containers and left for at least 30 min in order to completely liquefy before being processed. Sperm from ejaculates with normal parameters of semen volume, sperm count, motility, vitality and morphology, according to the WHO Laboratory Manual (World Health Organization, 1999), were included in this study. In each experiment, three normal samples were pooled. Spermatozoa preparation was

performed as previously described (Aquila *et al.*, 2002). An independent observer, who observed several fields for each slide, inspected the cells. Percoll-purified sperm were washed with un-supplemented Earle's medium and were incubated in the same medium (uncapacitating medium) for 30 min at 37°C and 5% CO<sub>2</sub>, without (control) or with treatments (experimental). Some samples were incubated in capacitating medium (CAP) (Earle's balanced salt solution medium supplemented with 600 mg BSA /100 ml and 200 mg sodium bicarbonate/100 ml). When the cells were treated with the inhibitors Cax, OH-Flut and ZVF, a pretreatment of 15 min was performed. The study was approved by the local medical Ethical Committees and all participants gave their informed consent.

#### LNCaP cells culture

LNCaP, human prostate adenocarcinoma cells, were grown in RPMI 1640 medium supplemented by 5% heat inactivated fetal bovine serum and Penicillin–Streptomycin 1%. Cultures were maintained at 37°C, 5% CO<sub>2</sub> and 100% humidity. In the experiments, steroids and growth factors were withdrawn from cells, and they were grown in phenol red-free DMEM containing 0.5% BSA and 2 mM L-glutamine for 24 h. LNCaP were treated for 30 min at 37°C and 5% CO<sub>2</sub>, without (control) or with the indicated treatments (experimental).

#### RNA isolation and RT–PCR

Total RNA was isolated from human ejaculated spermatozoa and purified as previously described (Aquila *et al.*, 2002). Contamination by leucocytes and germ cells in the sperm preparations was assessed by amplifying *PTPRC* and *KIT* transcripts, respectively. The applied PCR primers and the expected lengths of the resulting PCR products are shown in Table 1. *AR* primers were chosen to amplify the region of the DNA binding domain plus the hinge region of the receptor. PCR was carried for 40 cycles using the following parameters: 95°C/1 min, 55°C/1 min, 72°C/2 min for *AR*; 95°C/1 min, 52°C/1 min, 72°C/2 min for *KIT*; 95°C/1 min, 55°C/1 min, 72°C/2 min for *PTPRC*. For all PCR amplifications, negative (reverse transcription-PCR performed without M-MLV reverse transcriptase) and positive controls (LNCaP for *AR*, human testis for *KIT* and human leucocytes for *PTPRC*) were included.

**Table 1:** Oligonucleotide sequences used for RT–PCR

Gene	Sequence (5'–3')	Size of PCR product (bp)
<i>AR</i>	5'-TGCCCATGACTATTACTTTCC-3' 5'-TGTCCAGCACACTACACC-3'	400
<i>KIT</i>	5'-AGTACATGGACATGAAACCTGG-3' 5'-GATTCTGCTCAGACATCGTCG-3'	780
<i>PTPRC</i>	5'-CAATAGCTACTACTCCATCTAAGCCA-3' 5'-ATGTCTTATCAGGAGCAGTACATG-3'	230

#### Gel extraction and DNA sequence analysis

The *AR* RT–PCR product was extracted from the agarose gel by using a gel band purification kit, and the purified DNAs were subcloned into PCR 2.1 vector and then sequenced.

#### Western blot analysis of sperm proteins

Sperm samples washed twice with Earle's balanced salt solution (uncapacitating medium), were incubated for 30 min without or with the treatments indicated in the figures. During western blot analysis, sperm samples were processed as previously described (Aquila

*et al.*, 2002). The negative control was performed using a sperm lysate that was immunodepleted of AR (i.e. preincubation of lysates with anti-AR antibody for 1 h at room temperature and immunoprecipitated with protein A/G-agarose) (Aquila *et al.*, 2004). As internal controls, all membranes were subsequently stripped (glycine 0.2 M, pH 2.6 for 30 min at room temperature) of the first antibody and re-probed with anti-actin antibody. As a positive control, LNCaP (prostate cancer cell line) was used. The intensity of bands representing relevant proteins was measured by Scion Image laser densitometry scanning program.

#### Immunofluorescence assay

Sperm cells, were rinsed three times with 0.5 mM Tris–HCl buffer (pH 7.5) and were fixed using absolute methanol for 7 min at –20°C. AR staining was carried out, after blocking with normal human serum (10%), using the monoclonal anti-human AR (1 µg/ml) as primary antibody and an anti-mouse IgG FITC conjugated (4 µg/ml) as secondary antibody. To stain DNA in living cells, Hoechst 33 342 (Hoechst) was added at a final concentration of 10 µg/ml. The specificity of AR was tested by using normal mouse serum instead of the primary antibody; sperm cells incubated without the primary antibody were also used as negative controls. The cellular localization of AR and Hoechst was studied with a Bio-Rad MRC 1024 confocal microscope connected to a Zeiss Axiovert 135 M inverted microscope with 600×. The fluorophores were imaged separately to ensure no excitation/emission wavelength overlap, and a minimum of 200 spermatozoa per slide were scored.

#### Immunoprecipitation of sperm proteins and LNCaP cells proteins

Spermatozoa were washed in Earle's balanced salt solution and centrifuged at 800g for 20 min. Sperm resuspended in the same uncapacitating medium and LNCaP cells were incubated without (control, UC) or in the presence of dihydrotestosterone (DHT) at increasing concentrations (0.1, 1, 10 and 100 nM) for 30 min. Other samples were pre-treated for 15 min with 10 µM OH-Flut. In order to evaluate the rapid effect of DHT on AR, spermatozoa were incubated in the un-supplemented Earle's medium at 37°C and 5% CO<sub>2</sub> at different times (5 and 30 min and 1 h). To avoid non-specific binding, sperm lysates were incubated for 2 h with protein A/G-agarose beads at 4°C and centrifuged at 12 000g for 5 min. The supernatants (each containing 600 µg total protein) were then incubated overnight with 10 µl anti-AR and 500 µl HNTG (IP) buffer (50 mM HEPES, pH 7.4; 50 mM NaCl; 0.1% Triton X-100; 10% glycerol; 1 mM phenylmethylsulfonylfluoride; 10 µg/ml leupeptin; 10 µg/ml aprotinin and 2 µg/ml pepstatin). Immune complexes were recovered by incubation with protein A/G-agarose. The beads containing bound proteins were washed three times by centrifugation in immunoprecipitation buffer, then denatured by boiling in Laemmli sample buffer and analysed by western blot to identify the coprecipitating effector proteins. Immunoprecipitation using normal mouse serum was used as negative control. Membranes were stripped of bound antibodies by incubation in glycine (0.2 M, pH 2.6) for 30 min at room temperature. Before re-probing with the different indicated antibodies, stripped membranes were washed extensively in Tris buffered saline with Tween 20 (TBS-T) and placed in blocking buffer (TBS-T containing 5% milk) overnight.

#### Evaluation of sperm viability

Viability was assessed by using Eosin Y method. Spermatozoa were washed in uncapacitating medium and centrifuged at 800g for 20 min. To test androgen effects on sperm viability, spermatozoa

were incubated in unsupplemented Earle's medium at 37°C and 5% CO<sub>2</sub> without (control, UC) or in the presence of DHT at increasing concentrations (0.1, 10 and 100 nM) or 10 nM testosterone (T) for 2 h. In a different set of experiments, sperm were incubated in unsupplemented Earle's medium at 37°C and 5% CO<sub>2</sub> without (UC) or in the presence of 10 nM DHT or T at different times (0, 10 and 30 min, 2, 6 and 24 h). Some samples were pretreated for 15 min with 10 µM OH-Flut. 10 µl of Eosin Y [0.5% in phosphate-buffered saline (PBS)] were mixed with an equal volume of sperm sample on a microscope slide. The stained dead cells and live cells that excluded the dye, were scored among a total of 200 cells and by an independent observer. Further, viability was evaluated before and after pooling the samples.

### PI3K activity

PI3K activity was performed as previously described (Aquila *et al.*, 2004). The negative control was performed using a sperm lysate, where p110 catalyzing subunit of PI3K was previously removed by preincubation with the respective antibody (1 h at room temperature) and subsequently immunoprecipitated with protein A/G-agarose. The PI3K1 was precipitated from 500 µg of sperm lysates. The immunoprecipitates were washed once with cold PBS, twice with 0.5 M LiCl, 0.1 M Tris (pH 7.4) and finally with 10 mM Tris, 100 mM NaCl and 1 mM EDTA. The presence of PI3K activity in immunoprecipitates was determined by incubating the beads with reaction buffer containing 10 mM HEPES (pH 7.4), 10 mM MgCl<sub>2</sub>, 50 µM ATP, 20 µCi [ $\gamma$ -<sup>32</sup>P] ATP and 10 µg L- $\alpha$ -phosphatidylinositol-4,5-bis phosphate (PI-4,5-P<sub>2</sub>) for 20 min at 37°C. Phospholipids were extracted with 200 µl CHCl<sub>3</sub>/methanol. The labelled products of the kinase reaction, the PI phosphates, in the lower chloroform phase were spotted onto *trans*-1,2-diaminocyclohexane-*N,N,N',N'*-tetraacetic acid-treated silica gel 60 thin-layer chromatography plates state running solvent used for TLC. Radioactive spots were visualized by autoradiography.

### DNA laddering

DNA laddering was determined by gel electrophoresis. Spermatozoa were washed in Earle's balanced salt solution and centrifuged at 800g for 20 min, then were resuspended in the same uncapacitating medium and in different tubes containing no androgens (control, UC), T or DHT or estrogen or progesterone or wortmannin at the indicated concentrations for 30 min. Some samples were resuspended in CAP. Some samples were pretreated for 15 min with 10 µM Cax or 10 µM OH-Flut or ZVF alone or each combined with 10 nM DHT. After incubation cells were pelleted at 800g for 10 min. The samples were resuspended in 0.5 ml of extraction buffer (50 mM Tris-HCl [pH 8], 10 mM EDTA, 0.5% sodium dodecyl sulphate (SDS)] for 20 min in rotation at 4°C. DNA was extracted with phenol/chloroform for three times and once with chloroform. The aqueous phase was used to precipitate acids nucleic with 0.1 volumes of 3 M sodium acetate and 2.5 volumes cold EtOH overnight at -20°C. The DNA pellet was resuspended in 15 µl of H<sub>2</sub>O treated with RNase A for 30 min at 37°C. The absorbance of the DNA solution at 260 and 280 nm was determined by spectrophotometry. The extracted DNA (2 µg/lane) was subjected to electrophoresis on 1.5% agarose gels. The gels were stained with ethidium bromide and then photographed.

### Statistical analysis

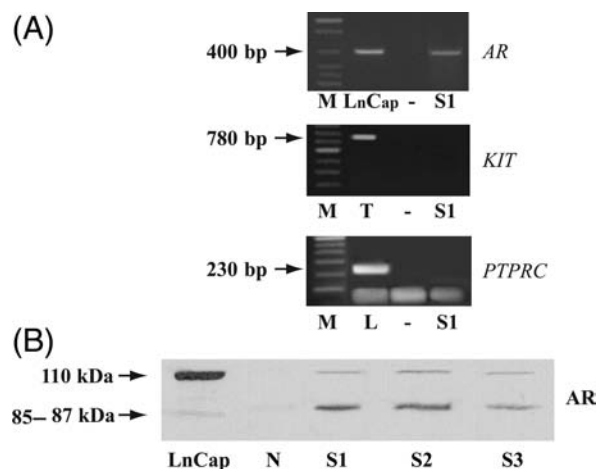
The experiments for RT-PCR, immunofluorescence and immunoprecipitation assays were repeated on at least four independent occasions, and western blot analysis was performed in at

least six independent experiments, PI3K activity and DNA laddering assay were performed in at least four independent experiments. The data obtained from viability (six replicate experiments using duplicate determinations) were presented as the mean  $\pm$  SEM. Statistical analysis was performed using analysis of variance (ANOVA) followed by Newman-Keuls testing to determine differences in means.  $P < 0.05$  was considered as statistically significant.

## Results

### AR mRNA and protein were detected in human sperm

To determine whether mRNA for AR is present in human ejaculated spermatozoa, RNA isolated from percoll-purified sperm samples from normal men was subjected to reverse PCR. The nucleotide sequence of AR was deduced from the cDNA sequence of the human AR gene and our primers amplified a region from 1648 to 2055 bp corresponding to the DNA binding domain plus the hinge region of the AR. RT-PCR amplification of AR in human sperm revealed the expected PCR product size of 400 bp. This product was sequenced and found identical to the classical human AR. No detectable levels of mRNA coding either *PTPRC*, a specific marker of leucocytes, or *KIT*, a specific marker of germ cells, were found in the same semen samples (Fig. 1A), thus ruling out any potential contamination.

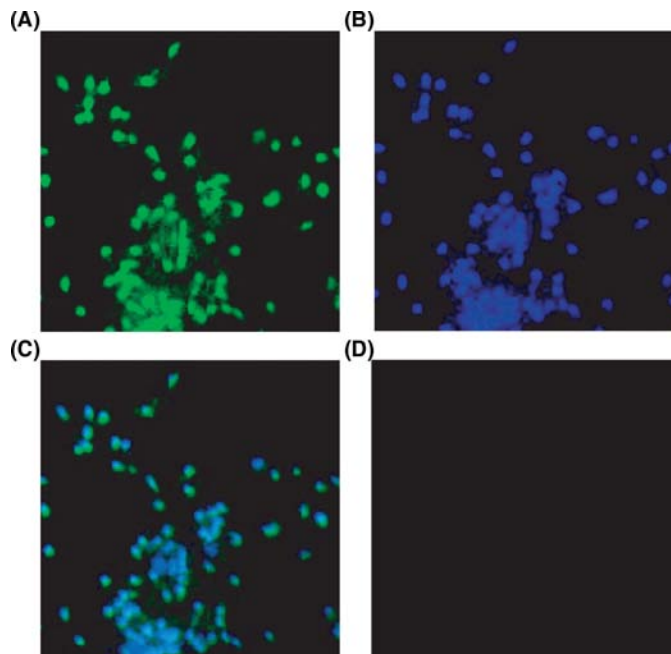


**Figure 1:** AR expression in human ejaculated spermatozoa (A) reverse transcription-PCR analysis of human AR gene, *KIT* and *PTPRC* in percollated human ejaculated spermatozoa (S1), negative control (no M-MLV reverse transcriptase added) (-), positive control (LnCap, prostate cancer cell; T, human testis and L, human leucocytes), marker (M). Arrows indicate the expected size of the PCR products; (B) western blot of AR protein by using a monoclonal antibody raised against the epitope 299–316 of the AR from human origin: extracts of percollated sperm, were subjected to electrophoresis on 10% SDS-polyacrylamide gels, blotted onto nitrocellulose membranes and probed with the above mentioned antibody. Expression of the receptors in three samples of ejaculated spermatozoa from normal men (S1, S2 and S3). LNCap cells were used as positive control. N, negative control performed as described in *Materials and Methods*. The experiments were repeated at least four times and the autoradiographs of the figure show the results of one representative experiment

The presence of AR protein in human ejaculated spermatozoa was investigated by western blot using a monoclonal antibody raised against the epitope mapping at the 299–316 aa in the N-terminus of AR from human origin (Fig. 1B). The antibody revealed the presence in sperm of two protein bands with molecular weights of 110 and 85–87 kDa, the latter expressed to a greater extent.

### Immunolocalization of AR in human sperm

Using an immunofluorescence technique, we identified a positive signal for AR in human spermatozoa (Fig. 2A). No immunoreaction was detected either by replacing the anti-AR antibody by normal mouse serum (Fig. 2D) or when the primary antibody was omitted (data not shown), demonstrating the immunostaining specificity. AR immunoreactivity was specifically compartmentalized at the sperm head (Fig. 2A), where the DNA is packaged, as it can be seen in Fig. 2B in which the DNA is stained by Hoechst. Fig. 2C shows the merged images of Fig. 2A and B.



**Figure 2:** Immunolocalization of AR in human ejaculated spermatozoa

Spermatozoa were extensively washed and incubated in the unsupplemented Earle's medium for 30 min at 37°C and 5% CO<sub>2</sub>. Spermatozoa were then fixed and analyzed by immunostaining as detailed in *Materials and Methods*. (A) AR localization in sperm; (B) staining with Hoechst of spermatozoa nuclei; (C) overlapping images of A and B; (D) sperm cells incubated replacing the anti-AR antibody by normal mouse IgG were utilized as negative control. The pictures shown are representative examples of experiments that were performed at least four times with reproducible results

### AR is phosphorylated in human sperm

It was reported that the function of AR is strongly correlated with the phosphorylation status (Wang *et al.*, 1999), which is rapidly enhanced upon androgen exposure when it is able to activate signal transduction pathways. AR immunoprecipitates were blotted with three different antibodies: anti-p-tyrosine, anti-p-threonine and anti-p-serine. As shown in Fig. 3A, two

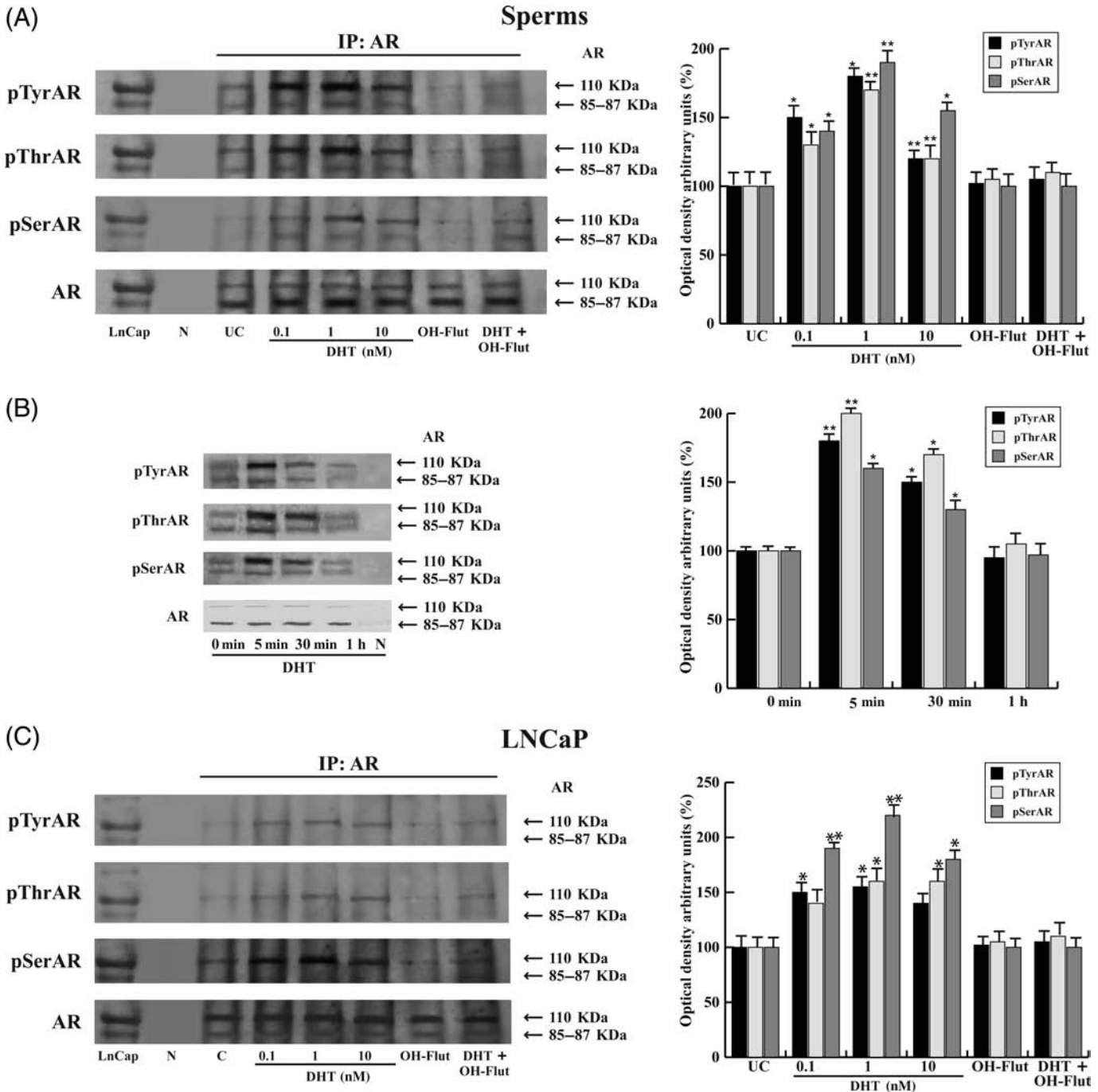
major AR antibody reactive proteins corresponding to the 85–87 and 110 kDa were observed. To determine if the changes in phosphorylation status of AR under androgen treatments may occur in ejaculated sperm, these were exposed for 30 min to varying concentrations of DHT (0.1–10 nM). We observed that the AR phosphorylation was enhanced in a dose related manner (Fig. 3A) and was significantly reduced by OH-Flut, an AR antagonist. To investigate if the enhanced phosphorylation status may represent an early event, we performed a time course study revealing that AR phosphorylations occurred rapidly as they were observed from 0 to 15 min and then dropped significantly after 1 h (Fig. 3B). Moreover, all three phospho-antibodies demonstrated a prevalence for phosphorylation of the 110 kDa isoform. Furthermore, we repeated the experiments with the LNCaP cells to see whether they show similar results. As evidenced in Fig. 3C, in LNCaP cells, the major phosphorylation event appears to affect serine residues to a higher extent.

### Androgens effect on sperm viability

To evaluate sperm viability under androgen treatment, we performed different sets of experiments. Sperm were incubated in the presence of 10 nM T or 10 nM DHT at the indicated times (Fig. 4A). Other samples were incubated in uncaptivating medium for 2 h in the absence or presence of different T or DHT concentrations (0.1–100 nM). As shown in Fig. 4B, the majority of cells remained viable in the control at 2 h. Cell viability significantly decreased with 10 and 100 nM T or DHT. Interestingly, the effect of androgen was reversed by using OH-Flut, addressing an AR mediated effect. It should be mentioned that the 100 nM androgen concentration is much higher than that commonly found circulating *in vivo* in man, while about 3 nM is detected in the seminal plasma of normal subjects (Luboshitzky *et al.*, 2002).

### Androgen action on PI3K activity, p-AKT, p-BCL2 and p-PTEN is mediated by AR

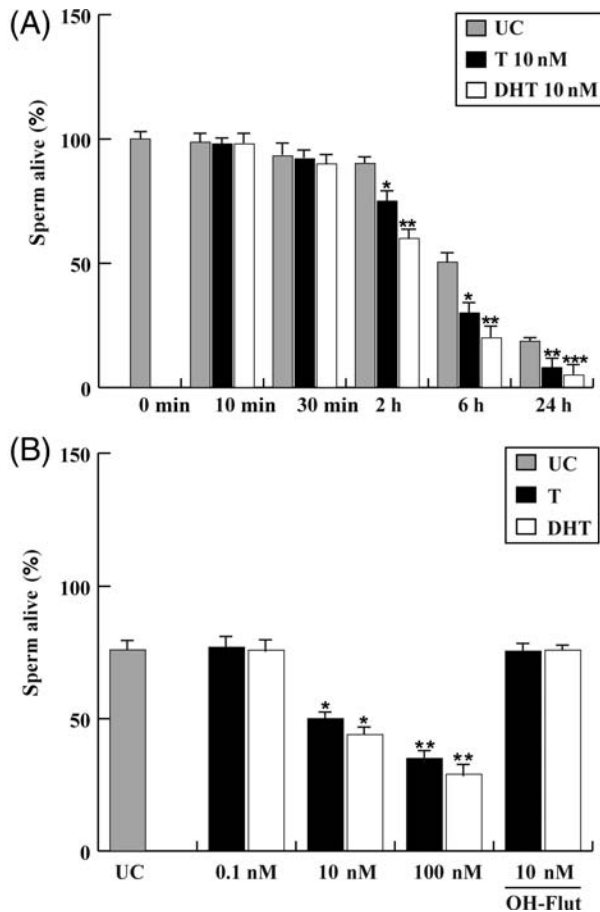
As shown in Fig. 5A low androgen concentration (0.1 and 1 nM) induced PI3K activity, while it was reduced by using 10 and 100 nM DHT. Both 10 nM T and to a greater extent 10 nM DHT treatments decreased PI3K activity (Fig. 5A). The 10 nM DHT effect was reversed in the presence of OH-Flut. Concomitantly, we observed a reduction on the levels of the downstream p-AKT S473 and p-AKT T308 (Fig. 5B and C) as well as p-BCL2 (Fig. 5D), a known antiapoptotic protein (Ito *et al.*, 1997). Specifically, DHT but not T had a significant inhibitory effect on p-AKT S473 and p-AKT T308 levels. Further, 10 nM of T or DHT significantly increased the phosphorylation of PTEN, a specific inhibitor of PI3K (Fig. 5E). All the above mentioned effects were reversed by Cax and OH-Flut, indicating that the effects of androgens are mediated by the classic AR in sperm. Recently, it was found that estradiol (E<sub>2</sub>) enhances sperm survival signalling (Aquila *et al.*, 2004). Therefore, we aimed to evaluate whether, in sperm, a functional interaction exists between androgen and estrogen on PI3K activity. In sperm samples incubated with 100 nM E<sub>2</sub> combined with increasing DHT



**Figure 3:** AR is phosphorylated in human sperm

AR phosphorylation was determined by immunoprecipitation using an AR specific antibody. The immunoprecipitates were blotted with three different antibody: anti-p-tyrosine (pTyrAR), anti-p-threonine (pThrAR) and anti-p-serine (pSerAR). (A) sperm were incubated without (control, UC) or in the presence of DHT at increasing concentrations (0.1, 1 and 10 nM) for 30 min. Some samples were pretreated for 15 min with 10  $\mu$ M OH-Flut. Sperm lysates (600  $\mu$ g) were immunoprecipitated using anti-AR and then blotted with specific antibodies raised to anti-p-tyrosine, anti-p-serine, anti-p-threonine, anti-AR. Immunoprecipitation by using normal mouse serum was used as negative control (N). The autoradiographs presented are representative examples of experiments that were performed at least four times with repetitive results. Molecular weight markers are indicated on the right of the blot. The histograms indicated on the right of each blot are the quantitative representation after densitometry of data (mean  $\pm$  SD) of four independent experiments. \* $P$  < 0.05, \*\* $P$  < 0.01 DHT-treated versus untreated cells. (B) time course of sperm incubated without (control, UC) or in the presence of 10 nM DHT. The autoradiographs presented are representative examples of experiments that were performed at least four times with repetitive results. Molecular weight markers are indicated on the right of the blot. The histograms indicated on the right of each blot are the quantitative representation after densitometry of data (mean  $\pm$  SD) of four independent experiments. \* $P$  < 0.05, \*\* $P$  < 0.01 DHT-treated versus untreated cells. (C) LNCaP cells were incubated without (control, C) or in the presence of DHT at increasing concentrations (0.1, 1 and 10 nM) for 30 min. Some samples were pretreated for 15 min with 10  $\mu$ M OH-Flut. The autoradiographs presented are representative examples of experiments that were performed at least four times with repetitive results. Molecular weight markers are indicated on the right of the blot. The histograms indicated on the right of each blot are the quantitative representation after densitometry of data (mean  $\pm$  SD) of four independent experiments. \* $P$  < 0.05, \*\* $P$  < 0.01 DHT-treated versus untreated cells





**Figure 4:** Effect of androgen on sperm viability. Viability was assessed by using Eosin Y as described in *Materials and Methods*. (A) time course of sperm incubated in the unsupplemented Earle's medium at 37°C and 5% CO<sub>2</sub> without (UC) or in the presence of 10 nM T or 10 nM DHT. (B) sperm were incubated without (UC) or in the presence of T or DHT at increasing concentrations (0.1, 10 and 100 nM) for 2 h. Some samples were pretreated for 15 min with 10 μM OH-Flut and then treated with 10 nM T or DHT. All experiments were repeated at least six independent times with duplicate samples, and the values represent the mean ± SEM. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.005 versus control

concentrations, the E<sub>2</sub>-induced PI3K activity progressively decreased (Fig. 5F).

#### **Androgens induce AR, PIK3R1 and phospho-src tyrosine kinase coimmunoprecipitation in human sperm**

It was reported that a triple complex between AR, PIK3R1 and SRC tyrosine kinase is required for androgen-stimulated PI3K/AKT activation (Castoria *et al.*, 2003; Sun *et al.*, 2003), therefore we investigated whether it also occurs in sperm. At the 0.1 nM DHT concentration, phospho-SRC tyrosine kinase coimmunoprecipitated with the two proteins immunodetected by the C-19 anti-AR antibody, migrating at 110 and 85–87 kDa. Remarkably, no association of phospho-SRC tyrosine kinase with AR occurred at the 100 nM DHT concentration. Fig. 6 shows immunocomplexes blotted with anti-AR (Fig. 6A) or phospho-SRC tyrosine kinase (Fig. 6B) or anti-PIK3R1 (Fig. 6C) antibodies. The possibility that androgen treatment could modify the AR level was excluded since in

the immunoprecipitated proteins the same amount of AR was detected.

Since the coimmunoprecipitation of phospho-SRC tyrosine kinase and PI3-kinase with AR decreased as androgens concentration increased, we may suppose that this is the mechanism through which high DHT concentration reduced PI3k activity.

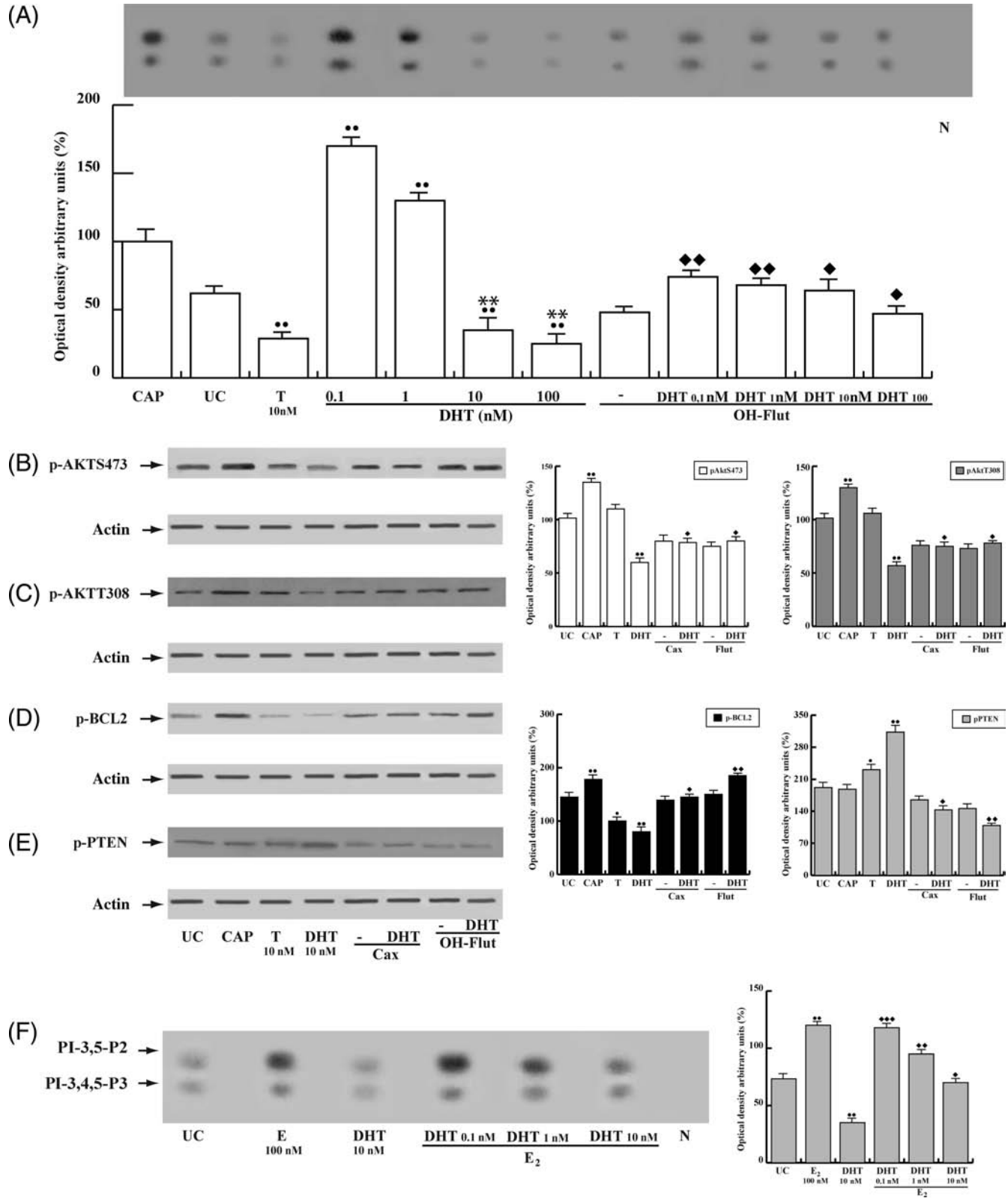
#### **Androgens effects on caspases are mediated by AR**

On the basis of the abovementioned results we sought to evaluate androgen action on the caspases family (Paasch *et al.*, 2004), since these proteins are involved in cell death. Particularly, caspase 3 which is the main effector of both caspases 8 and 9, executes the final disassembly of the cell by cleaving a variety of cell structure proteins and generating DNA strand breaks. Our study revealed activation of the caspases 8, 9 and 3 upon 30 min of 10 nM T or 10 nM DHT treatments. The DHT effect was reversed by both AR antagonists, 10 μM Cax or 10 μM OH-Flut (Fig. 7A). Notably, the effect on caspases was specific for androgen as it was not observed with E<sub>2</sub> or progesterone treatments. Particularly, progesterone treatment was performed because of the similarity in structure between progesterone receptor and AR. The cleavage of caspase 3 was increased by androgens in a dose-dependent manner (Fig. 7C). In the presence of wortmannin, a specific inhibitor of PI3K activity, the cleavage of caspase 3 was also observed, addressing a regulatory role of PI3K in caspase activation in sperm. Furthermore, in order to demonstrate a specific effect on caspase activation, an additional control experiment was included showing that activation of caspases by androgens can be inhibited with a specific caspases inhibitor such as ZVK. All these data were confirmed by DNA laddering assay (Fig. 7B and D).

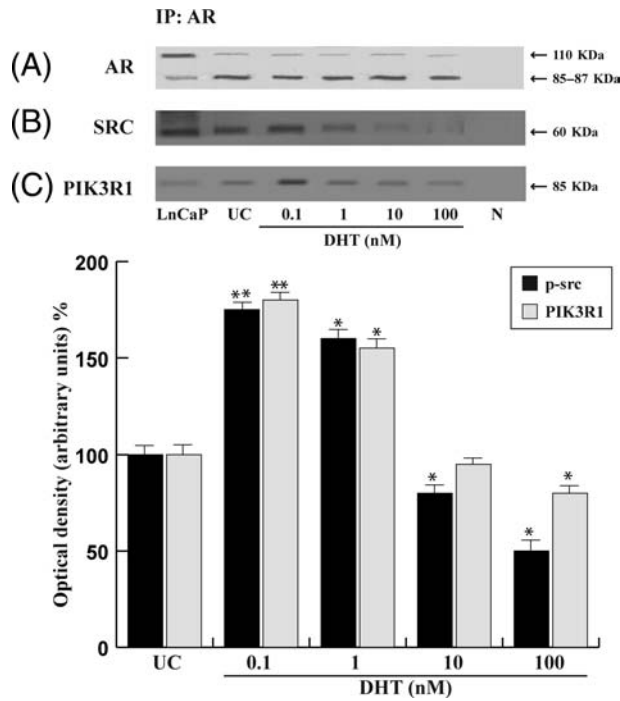
#### **Discussion**

Androgens and AR have been shown to play critical roles in testis function (Collins *et al.*, 2003). AR has been detected in Sertoli, Leydig, peritubular myoid and spermatid cells (round and elongated) (Kimura *et al.*, 1993; Vornberger *et al.*, 1994; Suarez-Quian *et al.*, 1999). The currently prevailing view is that sperm does not contain AR and this stems from previous studies reporting that no AR immunostaining of germ cells was observed both in rat and in human testis (Suarez-Quian *et al.*, 1999). However, several studies reported that in spermatozoa, the binding capacity of androgens was greater than that of estrogens or progesterone (Hyne and Boettcher, 1977; Cheng *et al.*, 1981), and recently AR was shown to be present in sperm by western blot and immunofluorescence assays (Solakidi *et al.*, 2005).

In this study, we have demonstrated the presence of AR in human sperm at different levels: mRNA expression, protein expression and immunolocalization. By RT-PCR, we amplified a gene region corresponding to the DNA binding domain plus the hinge region of the human AR. This product was sequenced and found to be identical to the classical human AR. As it concerns the presence of mRNAs in mammalian ejaculated spermatozoa, originally it was hypothesized that these transcripts were carried over from earlier stages of



**Figure 5:** Androgens action on PI3K activity, p-AKT, p-BCL2 and p-PTEN is mediated by AR. Washed pooled sperm from normal samples were incubated in the unsupplemented Earle's medium at 37°C and 5% CO<sub>2</sub>, in the absence (UC) or in the presence of 10 nM T or in the presence of DHT at increasing concentrations (0.1, 1, 10 and 100 nM) for 30 min. 500 µg of sperm lysates were used for PI3K activity in sperm incubated at the indicated DHT concentrations in the absence or in the presence of 10 µM OH-Flut (A). The autoradiograph presented is representative example of experiments that were performed at least four times with repetitive results. The histograms indicated on the bottom of the figure are the quantitative representation after densitometry of data (mean ± SD) of four independent experiments. \*\**P* < 0.01 T- and DHT-treated versus untreated cells, \*\**P* < 0.01 10 and 100 nM DHT versus 0.1 and 1 nM DHT, ♦♦*P* < 0.01 and ♦*P* < 0.05 10 µM OH-Flut plus DHT-treated versus DHT-treated cells. 50 µg of sperm lysates were used for western blot analysis of p-AKT S473 (B) and



**Figure 6:** AR, PIK3R1 and phospho-SRC tyrosine kinase coimmunoprecipitate in human sperm

Washed spermatozoa from normal samples were incubated in the unsupplemented Earle's medium for 30 min at 37°C and 5% CO<sub>2</sub>, without (UC) or in the presence of DHT at increasing concentrations (0.1, 1, 10 and 100 nM). 600 µg of sperm lysates were immunoprecipitated using anti-AR antibody and then blotted with specific antibodies raised to AR (A), p-SRC tyrosine kinase (B) and PIK3R1 (C). LnCaP lysates were used as positive control (lane 1); Immunoprecipitation by using normal mouse serum was used as negative control (N). The autoradiographs presented are representative examples of experiments that were performed at least four times with repetitive results. Molecular weight markers are indicated on the left of the blot. \**P* < 0.05, \*\**P* < 0.01 DHT-treated versus untreated cells

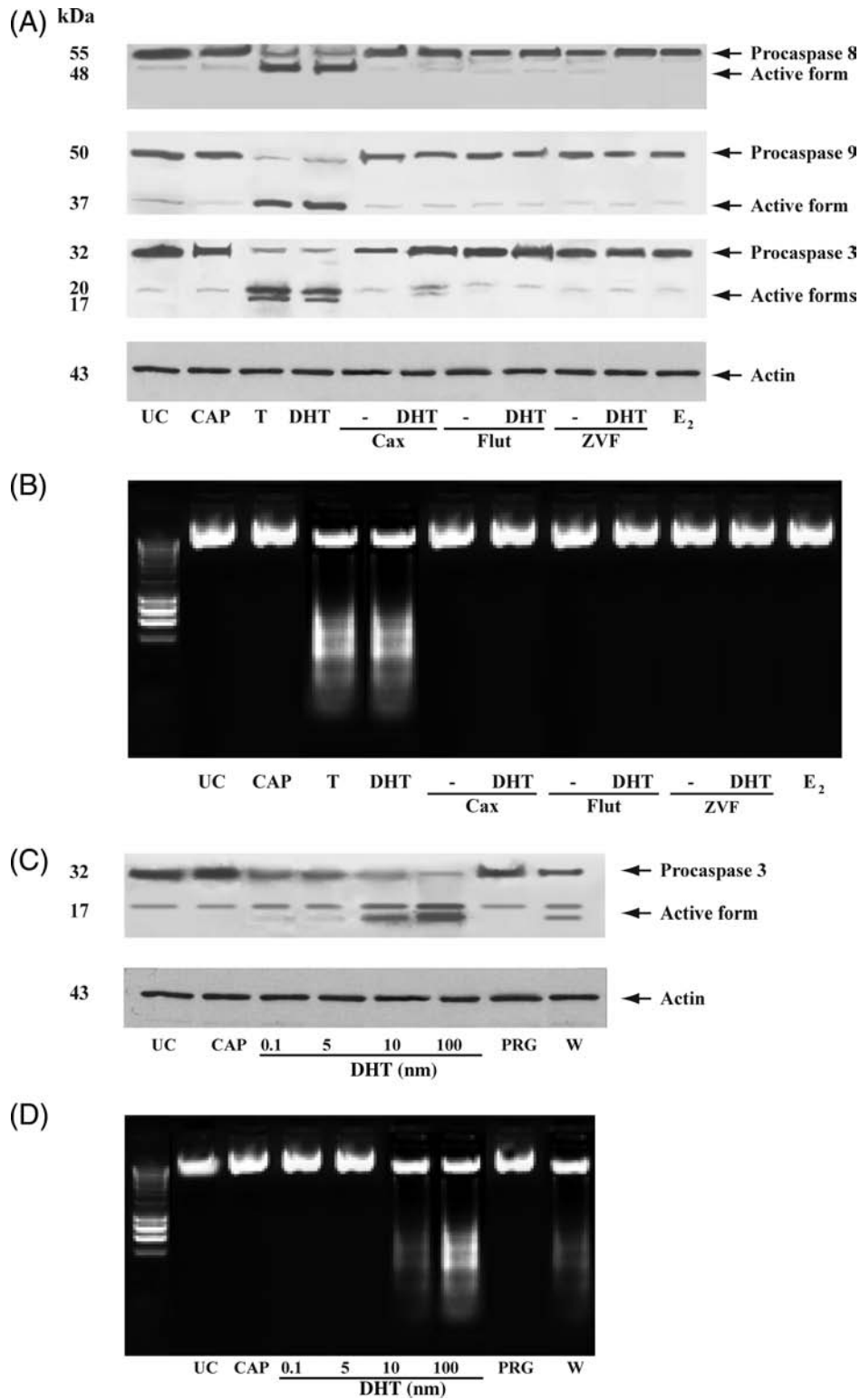
spermatogenesis, however, new reports re-evaluate the significance of mRNA in these cells (Andò and Aquila, 2005; Miller *et al.*, 2005) and the issue is currently under investigation.

To date, multiple isoforms of the AR have been described and among them two proteins are well characterized: AR-B and AR-A (Wilson and McPhaul, 1996). They are believed to be derived from the same gene and differ only in the NH<sub>2</sub>-terminal transactivation domain (Wilson and McPhaul, 1996). Our antibody against an epitope (aminoacids 299–311), i.e. common to both the AR-A and AR-B isoforms, detected two protein bands: one of 85–87 kDa and another one approximately of 110 kDa. Both, the AR-B and AR-A isoforms, are expressed in a variety of fetal and adult (male and

female) human tissues and especially in reproductive tissues (Wilson and McPhaul, 1994). The B form migrates with an apparent mass of 110 kDa and constitutes >80% of the immunoreactive receptor in most cell types. The A form of the AR migrates with an apparent mass of 87 kDa. It was identified as an NH<sub>2</sub>-terminally (from 1 to 187 aa) truncated protein of AR-B and it was first described in human genital skin fibroblasts. The detection of two distinct forms of the AR raised a number of issues. AR-A is expressed at low levels in many androgen-responsive tissues; however, it appears to have functions similar to those of the full-length AR-B isoform. Functional activities of cDNAs containing the two isoforms were assessed using cotransfection assays that employed two models of androgen-responsive genes (MMTV-luciferase and PRE2-tk-luciferase) in response to mibolerone, a potent androgen agonist, in three different cell lines (Gao and McPhaul, 1998). These studies demonstrated subtle differences in the activities of the A and B isoforms, which depended on the promoter and cell context. Additional studies failed to reveal any major differences in the responses of the AR-A and AR-B isoforms to a variety of androgen agonists and antagonists, suggesting that the previously reported functional defect of the AR-A is due principally to its level of expression. When assays of AR function are performed under conditions in which levels of expression of the two isoforms are equivalent, the AR-A and AR-B possess similar functional activities (Gao and McPhaul, 1998). The ratio of AR-B:AR-A may vary among tissues and at different stages of development. However, it is unknown whether these isoforms have divergent biologic signal transduction capacities in humans, therefore we cannot predict what is the physiological correlate of a low AR-B:AR-A ratio as observed in sperm. By immunohistochemical assays, we have demonstrated that AR protein is detectable in the sperm head. Solakidi *et al.* (2005) reported AR prevalently localized in the midpiece region and the labeling pattern was similar to that of ERα. The apparent discrepancy between the latter finding and ours may be due to the different methods used to process samples.

An increasing body of evidence suggests that androgens and other steroid hormones can exert rapid, nongenomic effects (Peterziel *et al.*, 1999; Cato *et al.*, 2002). Different nuclear receptors such as progesterone receptor (Calogero *et al.*, 2000), estrogen receptors α and estrogen receptor β (Aquila *et al.*, 2004) were found to be present in human ejaculated spermatozoa, regulating cellular processes through nongenomic mechanisms. All these findings strengthen the importance of the nuclear receptors in nongenomic signalling (Cato *et al.*, 2002) which may represent their exclusive modality of action

p-AKT T308 (C), p-BCL2 (D) and p-PTEN (E). The autoradiographs presented are representative examples of experiments that were performed at least six times with repetitive results. The histograms indicated on the right of each blot are the quantitative representation after densitometry of data (mean ± SD) of six independent experiments. \*\**P* < 0.01 capacitated (CAP) or DHT-treated versus untreated cells, ♦*P* < 0.05 T-treated versus untreated cells, ♦♦*P* < 0.05 10 and \*\**P* < 0.01 µM OH-Flut plus DHT or Cax plus DHT-treated versus DHT-treated cells. (E) p13K activity of sperm incubated with E<sub>2</sub> and/or DHT at the indicated increasing concentrations. The negative controls were performed using a sperm lysate, where p110 catalyzing subunit of PI3K was previously removed by preincubation with the respective antibody (1 h at room temperature) and subsequently immunoprecipitated with protein A/G-agarose (N). The autoradiographs presented are representative examples of experiments that were performed at least four times with repetitive results. The histograms indicated on the bottom of the figure are the quantitative representation after densitometry of data (mean ± SD) of four independent experiments. \*\**P* < 0.01 DHT- and E<sub>2</sub>-treated versus untreated cells, ♦*P* < 0.05 10, ♦♦*P* < 0.01 and \*\*\**P* < 0.001 E<sub>2</sub> plus DHT-treated versus DHT-treated cells



**Figure 7:** Androgens effects on caspases are mediated by AR

(A) washed pooled sperm from normal samples were incubated in the unsupplemented Earle's medium at 37°C and 5% CO<sub>2</sub> (UC) in the presence of 10 nM T or 10 nM DHT or 100 nM E for 30 min. Some samples were washed with the unsupplemented Earle's medium and incubated in capacitating medium (CAP). Some samples were treated with Cax or Flut or ZVF each alone or combined with 10 nM DHT. The sperm were lysed and subjected to western blot analysis. 70 µg of sperm lysates were used for western blot analysis of caspase 8, caspase 9 and caspase 3. (B) DNA laddering was performed in sperm treated as indicated. (C) effect of increasing DHT concentrations (0.1–100 nM), 100 nM PRG and 10 µM wortmanninn (W) on caspase 3 cleavage. The experiments were repeated at least six times and the autoradiographs of the figure show the results of one representative experiment. (D) DNA laddering was performed in sperm treated as indicated

in spermatozoa since they are apparently transcriptionally inactive cells. Here, we have demonstrated that in human ejaculated sperm, short exposure to androgens produces an increase in AR phosphorylation in a dose-dependent manner, while the antagonist OH-Flut significantly reduces this effect. Furthermore, we observed the most prominent phosphorylation on the 110 kDa band which is the less expressed isoform in sperm. It is known that the function of nuclear receptors is strongly correlated with their phosphorylation status rather than the level of total receptor proteins. The 110 kDa isoform exhibits a major length of the N-terminal domain which is an important effector of the cell signalling (Wilson and McPhaul, 1994, 1996). This may explain why the phosphorylated status of the 110 kDa appears much more pronounced than the smaller isoform. From these findings it emerges that in sperm the 110 kDa is the most involved isoform in mediating AR trafficking signals. Furthermore, in sperm, the phosphorylation of the less expressed isoform appears to occur, upon androgen binding, in tyrosine, threonine and to a greater extent with respect to the basal values, on serine aminoacidic residues. Therefore, we repeated the experiments with the LNCaP cells to see whether they show a similar result. In LNCaP cells, the major phosphorylation event appears to affect serine residues. In any case, we should take into account that even in uncipated sperm, the autocrine effect of a pool of cytokines, insulin-like growth factors, as we previously demonstrated (Aquila *et al.*, 2005a,b) may *per se* influence the phosphorylation status of AR in addition to that determined by its natural ligand.

On the basis of our data androgens are able to modulate sperm survival depending on their concentration. To investigate the molecular mechanism involved in these effects we evaluated their action on the PI3K/AKT pathway, since it represents the main cell survival pathway and it was identified in sperm (Aquila *et al.*, 2004). The 0.1 and 1 nM androgens induced PI3K activity, which was reduced by higher concentrations (10 and 100 nM). The 10 nM DHT was able to reduce the PI3K downstream signalling, while phosphorylation of PTEN, a proapoptotic marker which inhibits the PI3K pathway, was enhanced. To gain further insight into the mechanism involved in the PI3K/AKT modulation by AR, we investigated the association between AR/PIK3R1/p-SRC tyrosine kinase since it was reported depending on androgen concentration in somatic cells (Castoria *et al.*, 2003; Sun *et al.*, 2003). In our study, high androgen concentrations (10 and 100 nM) produce a detachment of SRC tyrosine kinase from the PIK3R1/AR complex, confirming that the triple complex is needed for the PI3K pathway activation. Furthermore, wortmannin, a specific PI3K/AKT inhibitor, induced caspase 3 cleavage in sperm, showing that the PI3K/AKT pathway is involved in the modulation of the caspases activity. The sperm death under high androgens (10 nM T, 10 and 100 nM DHT) was confirmed both by DNA laddering and cleavage of caspases 8, 9 and 3. In addition, increasing androgen concentrations were able to counteract the E<sub>2</sub>-induced PI3K activity previously documented (Aquila *et al.*, 2004).

It is well established that in men intratesticular T levels are ~800 nM (Coviello *et al.*, 2004), whereas they are ranging from 16 to 20 nM in serum (Luboshitzky *et al.*, 2002; Coviello

*et al.*, 2004). The androgenic milieu in seminal plasma is dependent on circulating androgen levels and no longer intratesticular levels (Kuwahara, 1976; Andò *et al.*, 1983). The biologically active amount of T, represented by its free fraction, in the genital tract is mostly converted in DHT by 5 alpha-reductase which is particularly expressed in the epididymis and in the adnexal glands (Steers, 2001). A careful evaluation of the total androgenic milieu in seminal plasma, prevalently represented by the two most important androgens T and DHT, reveals the presence of about 1 nM of T and 2 nM of DHT and their ratio is about T/DHT 0.61 (Andò *et al.*, 1983). Therefore, the seminal androgenic milieu, prevalently represented by the total molar concentration of T plus DHT corresponds to ~3 nM. In our study, the effects induced by 10 nM DHT were opposite to those induced by the lower doses and the same opposite pattern of androgen effects on PI3K pathway was previously documented in other cell type (Castoria *et al.*, 2003).

In conclusion, the importance of androgen in the completion of male gamete maturation during the epididymal transit has been proved by the presence of AR in epididymal tissue (Zhou *et al.*, 2002). Now on the basis of our results we may speculate how the importance of androgen in the sperm maturative process goes beyond the length of their life in seminiferous tubules, and continues when they became transcriptionally silent, through the AR nongenomic signalling. Even though to date, we cannot establish the physiopathologic correlates of these findings, we observed that an excess of androgens in the local hormonal milieu inhibits PI3K activity and negatively interferes with sperm survival. Further work will be required to more fully elucidate the role that AR plays in this aspect of male fertility.

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# Reproduction

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## Leptin and leptin receptor in pig spermatozoa: evidence of their involvement in sperm capacitation and survival.

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1 **Title page**

2

3 **Leptin and leptin receptor in pig spermatozoa: evidence of their involvement**

4 **in sperm capacitation and survival.**

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## Abstract

Much effort has recently focused on understanding the role of leptin, the obese gene product secreted by adipocytes, in regulating growth and reproduction in rodents, humans and domestic animals. The present study was designed to investigate the expression of leptin and its receptor in pig spermatozoa. Successful western blot evidenced a 16 kDa band for leptin and six isoforms ranging from 120 to 40 kDa for the leptin receptor, suggesting a possible biological effects in these cells. Intriguingly, leptin and leptin receptor were both located at the acrosomal level, addressing an action in the events that render the sperm able to fertilize the oocyte. In fact, both capacitation indexes and acrosin activity were enhanced upon 10 nM leptin and these effects were reduced by the anti-leptin receptor antibody. Afterwards, we explored the main transduction pathways regulated by the hormone. Our results showed that in pig sperm leptin can trigger the signal transducer and activator of transcription 3, a classical component of cytokine signal transduction pathways, which expression was never reported in male gamete and that we found constitutively activated. Besides, leptin was able to induce the activation of Phosphatidylinositol Phosphate Kinase 3 and Mitogen-Activated Protein Kinase pathways as well as of Bcl-2, a known antiapoptotic protein. Altogether, these data also address a role for leptin and its receptor on pig sperm survival. The presence of leptin and its receptor in pig sperm suggests that they, through an autocrine short loop, may induce signal transduction and molecular changes associated to sperm capacitation and survival.

## INTRODUCTION

1  
2 Leptin (Lep) is known to be the hormonal link between energy stores and several vital  
3 functions, including food intake and energy homeostasis. It is a 16-kDa adipocyte-  
4 derived hormone that suppresses food intake, stimulates energy expenditure, increases  
5 metabolic rate, and ultimately causes loss of body fat. (Boulounie *et al.* 1998; Kamohara  
6 *et al.* 1997; Rossetti *et al.* 1997). Much attention has focused on the effects of Lep as a  
7 central satiety agent and its action was reported to be involved in the regulation of  
8 fatness via feed intake. Although originally postulated to act largely via the central  
9 nervous system, several recent studies have demonstrated that Lep exerts a wide  
10 repertoire of peripheral effects through direct actions on target tissues. These direct  
11 effects include stimulation of fatty acid oxidation in adipocytes (Muller *et al.* 1997;  
12 Boulounie *et al.* 1998; Lord *et al.* 1998; Sierra-Honigmann *et al.* 1998; Zhao *et al.* 1998  
13 ) and increased glucose uptake in skeletal muscle and brown adipose tissue in vivo  
14 (Kamohara *et al.* 1997; Yaspelkis *et al.* 1999) and in a myotube cell line in vitro (Berti  
15 *et al.* 1997 ).

16 The pig is emerging rapidly as an important biomedical research model, and whereas  
17 genetic influences may impact the degree of similarity between pig and human systems,  
18 the regulation of some endocrine and metabolic processes in the pig may be more  
19 similar to humans than are rodents (Tsiaoussis *et al.* 2001; Vilei *et al.* 2001). Lep plays a  
20 key role in the regulation of energy homeostasis and is also linked to mechanisms  
21 controlling reproductive processes. With respect to reproductive function, the  
22 importance of Lep has grown rapidly in the past few years in fact, numerous lines of  
23 evidence point to the participation of this peptide in the regulation of neuroendocrine  
24 axis and thereby, among others , in reproductive processes of different species  
25 (Campfield *et al.* 1996; Foster *et al.* 1999; Ahima *et al.* 2000; Amstalden *et al.* 2000),

1 including pigs (Barb *et al.* 1998; 1999). However, data concerning the involvement of  
2 Lep in controlling reproductive functions at the level of hypothalamus and pituitary in  
3 the pig are very limited. The relationship between nutrition and reproduction in swine  
4 has been extensively reviewed (Zurek *et al.* 1995; Quesnel & Prunier 1995). Nutrition  
5 may influence reproductive performance by a number of mechanisms, including central  
6 effects on gonadotropin secretion (Booth *et al.* 1994) and local effects on ovarian  
7 function (Cosgrove *et al.* 1992; Booth *et al.* 1996). Consistent with the data reported by  
8 Zak (Zak *et al.* 1997), the different feeding regimens caused differential sow body  
9 weight changes. In the pig nutritional signals such as Lep are detected by the central  
10 nervous system (CNS) and translated by the neuroendocrine system into signals, which  
11 regulate appetite, hypothalamic gonadotropin-releasing hormone (GnRH) release and  
12 subsequent luteinizing hormone (LH) secretion. Changes in body weight or nutritional  
13 status are characterized by altered adipocyte function, a reduction in adipose tissue  
14 leptin expression, serum leptin concentrations and a concurrent decrease in LH  
15 secretion.

16 In both male and female mice it has been shown that leptin have direct effects on  
17 fertility since the hormone is able to reverse the infertility of *ob/ob* mice, lacking of the  
18 leptin gene (Cunningham *et al.* 1999). The biological actions of Lep on body weight  
19 homeostasis, neuroendocrine function and fertility are carried out through interactions  
20 with its specific obese receptor (Ob-R), in target tissues. The pattern of leptin receptor  
21 expression in the testis is specie-specific. Several isoforms of Ob-R exist that differ  
22 mainly in the length of the cytoplasmic domain (Takaya *et al.* 1996; Dieterichet *et al.*  
23 1996; Lee *et al.* 1996; Lollmann *et al.* 1997; Murakami *et al.* 1997; Bjorbaek *et al.*  
24 1998; Yamanita *et al.* 1998). The Ob-R, a single membrane-spanning glycoprotein,  
25 belongs to the class I cytokine receptor superfamily, sharing sequence homologies for  
26 interaction with Janus kinase as well as signal transducers and activators of transcription

1 (STATs) (Tartaglia *et al.* 1997). To date, the human Ob-R is identified as a full-length  
2 Ob-Rb form (Tartaglia *et al.* 1995) and as several short form receptors, generated by  
3 alternative splicing, OB-ra, OB-rc, OB-rd, OB-rf and OB-re (Bennet *et al.* 1996; Cioffi  
4 *et al.* 1996, Tartaglia *et al.* 1997). The hypothalamus is supposed to be the only tissue  
5 with predominantly expressed full-length Ob-R. However, an increasing number of  
6 peripheral tissues are described to express the long form as well as several isoforms  
7 (Cioffi *et al.* 1996; Tartaglia *et al.* 1997; Glasow *et al.* 1998; Breidert *et al.* 1999),  
8 including liver, heart, kidneys, lungs, small intestine, pituitary cells, testes, ovaries,  
9 spleen, pancreas, adrenal gland and adipose tissue (Margetic *et al.* 2002). Furthermore  
10 studies on different OB-r splice variants indicated that many tissues may contain a  
11 heterologous mix of OB-r subtypes. In a variety of porcine tissues Ob-R mRNA (long  
12 form) has been detected, including adipose tissue (Lin *et al.* 2000). Using *in situ*  
13 hybridisation techniques, other groups have shown that leptin receptor mRNA is  
14 expressed in Sertoli cells of adult rats (Tena-Sempere *et al.* 2001).

15 Rodent testis revealed the Ob-R expression but with a differential cellular site, in fact,  
16 the immunoreactivity was confined in Leydig cells of rat but in germ cells of mouse (El-  
17 Hefnawy *et al.* 2000, Caprio *et al.* 2003). This discrepancy may be due to interspecie  
18 difference. Lep and Ob-R were found to be present in the human seminiferous tubules  
19 (Soyupek *et al.* 2005), in human seminal plasma (Jope *et al.* 2003), in human sperm  
20 (Aquila *et al.* 2005) and recently Ob-R was also identified in boar sperm (De Ambrogi  
21 *et al.* 2007). In the present study we investigated Lep and Ob-R expression in pig sperm,  
22 evaluating their potential role on sperm capacitation and survival by the assessment of  
23 the main molecular markers involved in these processes.

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## Materials and Methods

### *Chemicals*

BSA protein standard, Laemmli sample buffer, prestained molecular weight marker, Percoll (colloidal PVP coated silica for cell separation), sodium bicarbonate, dimethyl sulfoxide, Earle's balanced salt solution, triethanolamine buffer, MgCl<sub>2</sub>, propidium iodide, LY294002 and all other chemicals were purchased from Sigma Chemical (Milan, Italy). Recombinant porcine leptin was purchased from Protein Laboratories Rehovot (Rehovot, Israel), acrylamide bisacrylamide was from Labtek Eurobio (Milan, Italy). Triton X-100, ECL Plus Western blotting detection system, Hybond™ ECL™, were purchased from Amersham Pharmacia Biotech (Buckinghamshire, UK). Antibodies (Abs) polyclonal rabbit anti-leptin (A-20), rabbit anti-Ob-R(H-300), rabbit anti-p-Akt1/Akt2/Akt3 (Ser473), rabbit anti-p-Akt1/Akt2/Akt3 (Thr308), rabbit anti-phosphotyrosine (PY99), mouse anti-p-signal transducer and activator of transcription-3 p-STAT3 (B-7), peroxidase-coupled anti-rabbit, and FITC/ Texas Red conjugated anti-rabbit IgG were from Santa Cruz Biotechnology (Heidelberg, Germany). Rabbit anti-p-Bcl-2 and p-ERK1/2 Abs were from Cell Signalling (Milan, Italy). Cholesterol-oxidase (CHOD)-peroxidase (POD) enzymatic colorimetric kit was from Inter-Medical (Biogemina Italia Srl, Catania, Italy).

### *Animals and semen samples*

The investigation has been conducted on semen from 7 fertile male pigs (*Sus scrofa domestica*, Large White) kept at " Swine Artificial Insemination Centre " (Rende, Cosenza, Italy). The animals were 24 to 30 month-old and their weights were from 280 to 320 kg. Individual fresh ejaculates were collected by the gloved hand method and filtered immediately by Universal Semen bags (Minitub, Tiefenbech, Germany). Semen was transported within half an hour to the laboratory, it was diluted 1:10 with TBS

1 buffer and centrifuged on a discontinuous Percoll density gradient (72 % / 90 %) to  
2 remove bacteria and debris (Kuster *et al.* 2004).

### 3 ***Evaluation of sperm viability***

4 Viability of pig sperm was assessed using the DNA-specific fluorochrome propidium  
5 iodide (PI). Sperm suspension ( $1 \times 10^6$  mL) was exposed to PI (12  $\mu\text{mol/L}$ ) for 5 min at  
6 room temperature. Then spermatozoa were fixed by adding 1 mL of 12.5% (w/v)  
7 paraformaldehyde in 0.5 mol Tris /l (pH 7.4) and the slides were immediately examined  
8 under an epifluorescence microscope (Olympus BX41) observing a minimum of 200  
9 spermatozoa  $\times$  slide (100 $\times$  objective).

### 10 ***Sample treatments***

11 Percoll-purified sperm were washed with unsupplemented Earle's medium  
12 (uncapacitating medium) and were incubated for 30 minutes (min) at 39 °C and 5 %  
13 CO<sub>2</sub>, without (control) or with increasing concentration of Lep (10 and 60 nM). These  
14 doses were chosen on the basis of physiological concentrations to reproduce the  
15 environment of the sperm journey. In fact, Lep concentration in porcine semen ranged  
16 from 9.2 - 13.38 ng/ml (considering a mean of 10 ng/ml we obtain 62 nM) (Lackey *et al.*  
17 2002) while the value for Lep in follicular fluid was about 1.9 ng/ml (12 nM)  
18 (Gregoraszczyk *et al.* 2004). Some cells were also pre-treated (15 min) with the  
19 inhibitor, LY 294002, or with the anti-Ob-R Ab (autocrine blockage). No adverse  
20 effects among the different treatments have been observed on pig sperm viability.

21 Then, samples were centrifuged (3000 g  $\times$  5min) using the upper phase for further  
22 determinations and sperm pellet for western blot analysis.

### 23 ***Western blot (WB) analysis of sperm proteins***

24 Western blot analysis was used to identify leptin and Ob-R in individual pig sperm.  
25 Sperm samples were washed twice with Earle's balanced salt solution (uncapacitating  
26 medium) and then centrifuged for 5 min at 5000 x g. The pellet was resuspended in lysis

1 buffer as previously described (Aquila *et al.* 2002). Equal amounts of proteins (80 µg)  
2 were boiled for 5 min, separated by 10 % polyacrylamide gel electrophoresis,  
3 transferred to nitrocellulose sheets and probed with an appropriate dilution of the  
4 (indicated) specific antibody (Ab). The bound of the secondary antibody was revealed  
5 with the ECL Plus WB detection system according to the manufacturer's instructions.  
6 The negative control was performed using a sperm lysate that was immunodepleted of  
7 leptin or Ob-R (i.e. lysates pre-incubated with anti-leptin Ab or anti-Ob-R Ab for 1 hour  
8 at room temperature and immunoprecipitated with Protein A/G-agarose).

9 To further validate the results for Ob-R, as negative control, non-immune rabbit serum,  
10 instead of the first antibody, was used at the same dilution ratio (1 : 1000).

11 Western blot was also performed to identify STAT3, Bcl-2, ERK 1/ 2, Akt1/ Akt2 in pig  
12 sperm extracts and to evaluate the cell signaling induced by Lep (p-Akt1/Akt2/Akt3, p-  
13 STAT-3, p-Bcl-2 and p-ERK1/2). MCF7 cell lines were used as positive controls while  
14 negative controls were performed as above indicated These last blots were stripped  
15 (glycine 0.2M ,pH2.6 for 30 min at room temperature) and riprobed with anti-βactin Ab  
16 as loading control.

17 The experiments were repeated at least four times for each sample.

### 18 ***Immunofluorescence assay***

19 Following Percoll separation, sperm cells were rinsed three times with 0.5mM Tris-HCl  
20 buffer, pH 7.5; then 10µl of concentrated cell suspension was added to 250µl drop of  
21 warm (37°C) TBS and allowed to settle onto slides in a humid chamber. The overlying  
22 solution was carefully pipetted off and replaced by absolute methanol for 7 minutes at –  
23 20 °C. After methanol removal, sperm cells were washed in TBS, containing 0.1%  
24 Triton X-100 and were treated for immunofluorescence

25 Leptin and Ob-R stainings were carried out, after blocking with normal goat serum (10  
26 %), using anti leptin(1:100) and anti-Ob-R (1:100) as primary antibodies, followed by

1 anti-rabbit IgG Texas Red conjugated/ FITC conjugated antibodies (1:200) respectively.  
2 Sperm cells incubated without the primary antibodies were utilized as negative controls.  
3 The slides were immediately examined under an epifluorescence microscope (Olympus  
4 BX41) observing a minimum of 200 spermatozoa × slide (100× objective).

#### 5 ***Measurement of cholesterol in the sperm culture medium***

6 Cholesterol was measured (in duplicate) in the incubation medium from pig  
7 spermatozoa by a CHOD - POD enzymatic colorimetric method according to  
8 manufacturer's instructions Sperm samples, washed twice with uncapacitating medium,  
9 were incubated in the same medium (control, NC) for 30 minutes (min) at 39 °C and 5  
10 % CO<sub>2</sub>. Other samples were incubated in the presence of 10 and 60 nM Lep  
11 concentrations . Some samples were incubated with anti-Ob-R combined with 10nM  
12 Lep. At the end of the sperm incubation, the culture media were recovered by  
13 centrifugation, lyophilized and subsequently dissolved in 1 ml of buffer reaction. The  
14 samples were incubated for 10 minutes at room temperature, then the cholesterol  
15 content was measured with the spectrophotometer at 505 nm. Cholesterol standard used  
16 was 200 mg/dl. The limit of sensitivity for the assay was 0.05 mg/dl. Inter- and intra-  
17 assay variations were 0,71 % and 0,57 % respectively.

#### 18 ***Acrosin activity assay***

19 Acrosin activity was assessed by the method of Glogowski (Glogowski *et al.* 1998). Briefly,  
20 sperm were washed in Earle's medium and centrifuged at 800 g for 20 min, then they were  
21 resuspended (final concentration of  $100 \times 10^3$  sperm/ml) in different tubes containing no  
22 treatment (control) or the indicated treatments (experimental). 1 ml of substrate-detergent  
23 mixture (23 mmol/l BAPNA in DMSO and 0.01% Triton X-100 in 0.055 mol/l NaCl, 0.055  
24 mol/l HEPES at pH 8.0 respectively) was added for 1 hour at room temperature. After  
25 incubation, 0.5 mol/l benzamidine was added (0.1 ml) to each of the tubes and then they were



1 centrifuged at 1000 g for 30 min. The supernatants were collected and the acrosin activity  
2 measured spectrophotometrically at 410 nm. In this assay, the total acrosin activity is defined  
3 as the amount of the active (non-zymogen) acrosin associated with sperm plus the amount of  
4 active acrosin that is obtained by pro-acrosin activable. The acrosin activity was expressed as  
5 mIU/10<sup>6</sup> sperms.

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For Review Only

## RESULTS

### ***Western Blot showed both Lep and Ob-R expression in pig sperm***

The presence of Lep protein in pig spermatozoa was investigated by WB using an antibody raised against the carboxyl-terminus of the human protein. One immunoreactive band was observed at 16 kDa in the lysates from pig sperm samples at the same mobility of the adipocyte extract used as positive control (Fig 1A).

Ob-R by WB, using the same antibody as used for immunohistochemistry (Fig. 1B) showed six different immunoreactive isoforms (120, 90, 80, 65, 60 and 40 kDa) (Fig 1B,b1). The bands were not detected by non immune rabbit serum indicating that these proteins are specific for Ob-R (Fig. 1B,b2).

### ***Immunolocalization of Lep and OB-R in pig sperm***

An intense red fluorescence localized Leptin in the acrosoma of pig sperm while the other cellular regions were unlabelled (Fig 2 A). In addition, a brilliant green light revealed that Ob-R was confined in the apical portion of sperm acrosoma ( 2 B). No fluorescent signal was obtained when primary antibodies (anti-Lep Ab or anti-Ob-R Ab) were omitted (inserts) thus confirming the specificity of the antibody binding.

### ***Lep influences pig sperm capacitation enhancing both cholesterol efflux and protein tyrosine phosphorylation***

One of the early events associated with capacitation of mammalian spermatozoa is the cholesterol efflux, after which protein phosphorylation begins. Our results showed a significant increase in cholesterol efflux (Fig 3 A). Protein tyrosine phosphorylation pattern of pig sperm, similar to that previously reported in porcine (Tardif *et al.* 2003), was also induced by leptin treatment (Fig 3 B). However, leptin effect on both processes

1 was not dose-dependent, because only 10 nM lep was able to sustain the increases.  
2 Further, anti-Ob-R Ab was able to abolish Lep effect.

### 3 ***Lep stimulates pig sperm acrosin activity***

4 Capacitation prepares the sperm for the acrosome reaction and acrosin is a sperm  
5 representative acrosomal enzyme. Acrosin activity showed that 10 nM Lep were able to  
6 stimulate the enzymatic activity, while no effect was observed with 60 nM Lep. Anti-Ob-R  
7 Ab combined with 10 nM Lep were able to abolish Lep effect (Fig 4).

### 8 ***STAT3 is activated by Lep/Ob-R in pig sperm***

9 The JAK/STAT pathway is one of the main signalling cascades activated by Lep  
10 (Thomas 2004). Our results showed in pig sperm the expression of the STAT3 (Fig 5  
11 A) which is a critical mediator of leptin action. Therefore, we aimed to investigate  
12 whether this important factor of Lep signal is activated in pig male gamete. As shown in  
13 Fig 5 B,C constitutively activated (phosphorylated) STAT3 is present in pig sperm and  
14 10 nM Lep treatment induced a slight but significant activation that was reversed by the  
15 anti-Ob-R Ab.

### 16 ***ERK 1/2 is activated by Lep/Ob-R in pig sperm***

17 Activation of MAPK has also been implicated as a signalling intermediate for Lep in  
18 various cell types (Takahashi *et al.* 1997; Tanabe *et al.* 1997). Besides, several lines of  
19 evidence indicate that components of the extracellular signal-regulated kinase (ERK)  
20 family of mitogen-activated protein kinases (MAPK) are present in spermatozoa and are  
21 involved in motility and capacitation (Naz *et al.* 1992; Luconi *et al.* 1998; de Lamirande  
22 & Gagnon 2002). Our results showed the expression of ERK 1/2 (Fig 6 A) and  
23 evidenced that Lep, through its receptor, is able to activate ERK 1/2 (Fig 6 B,C).

### 1 *Lep/Ob-R signalling modulates pig sperm survival*

2 Fig 7 A shows in pig sperm the expression of Bcl-2, a key protein in survival signaling.  
3 Furthermore, our data (Fig 7 B, C) revealed that leptin induced the Serine 70  
4 phosphorylation of Bcl-2, that is the physiologically relevant phosphorylation site,  
5 necessary for its full and potent antiapoptotic function. This effect was inhibited by the  
6 anti-Ob-R Ab.

7 PI3-K plays an important role in cell survival and metabolism in somatic cells as well as  
8 in sperm (Aquila *et al.* 2004; Aparicio *et al.* 2006). The PI3-K main downstream  
9 effector is the Akt, identified in human and boar spermatozoa. Fig 7 D shows the  
10 expression of Akt in pig sperm. Then we investigated the possible Akt phosphorylation  
11 by Lep evaluating the dual serine 473 and threonine 308 aminoacidic residues that give  
12 the full Akt activation. Our results showed that Akt is constitutively activated and that  
13 Lep induced an increase in the kinase phosphorylation, which was reduced by using anti-  
14 Ob-R Ab (Fig 7 E, F). To ensure that the increase in Akt activation is driven by PI3-K  
15 activation, the effect of a specific PI3-K inhibitor, 10 $\mu$ M LY294002 was assessed. The  
16 inhibitor abolished Lep-induced Akt S473 and Akt T308 phosphorylations (Fig 7 E, F).

### 17 *Statistical analysis*

18 Data, presented as mean  $\pm$  SEM, were evaluated by the one -way analysis of variance  
19 (ANOVA) The differences in mean values were calculated at a significance level of  
20  $P \leq 0.05$ .

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## DISCUSSION

1  
2 In recent years, it has been suggested that Lep is a metabolic signal to the reproductive  
3 system (Barash *et al.* 1996). To date, the mechanisms by which Lep regulate  
4 reproductive function remain to be determined. Lep and Ob-R were recently found in  
5 human sperm (Jope *et al.* 2003; Aquila *et al.* 2005) and the Ob-R in boar sperm (De  
6 Ambrogi *et al.* 2007). However, Lep and its signaling as well as its relationship with  
7 male gamete function across non-human species were not investigated. In the present  
8 study, we investigated, for the first time, the presence of Lep and Ob-R in pig sperm and  
9 evaluated their potential action on capacitation and sperm survival. The chief aim of our  
10 study was also to identify the potential signalling molecules set in motion by Lep in pig  
11 sperm.

12 First we evidenced the presence of Lep and OB-R by WB and immunofluorescence  
13 assay. Lep as protein was evidenced at the same size as it was observed in human sperm  
14 (Aquila *et al.* 2005), whereas different isoforms were obtained for the Ob-R. However,  
15 the pattern of leptin receptor expression is specie-specific and different Ob-R isoform  
16 sizes were reported in several tissues (Margetic *et al.* 2002). Interestingly, both Lep and  
17 Ob-R in pig sperm were found to be present exclusively at the acrosomal level. Since  
18 spermatozoa are highly polarized cells, they compartmentalize specific metabolic and  
19 signaling pathways to the regions where they are needed. On the basis of our data it is  
20 reasonably to hypothesize a role for Lep/Ob-R in the events that render the sperm able  
21 to fertilize the oocyte. As it concerns Ob-R, our data are similar to that of De Ambrogi (  
22 De Ambrogi *et al.* 2007), where the receptor was prevalently located on the acrosome of  
23 boar sperm, although they found weak immunofluorescence also in the subequatorial  
24 area and either on the whole tail or just on the midpiece. Whereas, in human sperm the  
25 receptor was visualized at the tail (Jope *et al.* 2003). These differences may be due to  
26 the sample management or they are specie-specific. Indeed, particularly intriguing is the

1 difference in the leptin expression site between human and pig, although the use of the  
2 same antibody. In fact, the hormone was specifically compartmentalized in the  
3 equatorial segment and the midpiece in human sperm (Aquila *et al.* 2005), whereas in  
4 pig on the acrosome. We can only hypothesize that leptin is differently involved in the  
5 regulation of sperm events in the two species.

6 In order to fertilize, the mammalian spermatozoa must undergo capacitation in the  
7 female tract. The mechanisms of capacitation remain poorly understood; although most  
8 studies have been conducted on mammals. During capacitation various biochemical and  
9 biophysical changes occur in the gamete and the molecular details of capacitation  
10 appear to vary somewhat among species. Up-stream of these events are subtle but all-  
11 determining changes to the sperm plasma membrane, among which may be mentioned  
12 the efflux of cholesterol (Visconti *et al.* 1999 a, b; Shadan *et al.* 2004). Cholesterol  
13 efflux from spermatozoa destabilizes lipid raft structures in the plasma membrane  
14 thereby initiating protein phosphorylation and acquisition of the capacitated state.  
15 Tyrosine phosphorylation of sperm proteins occurs during capacitation in mouse,  
16 human, bull, hamster (Visconti *et al.* 1995; Leclerc *et al.* 1996; Galatino-Homer *et al.*  
17 1997) and also in the pig ( Kalab *et al.* 1998; Flesch *et al.* 1999). It appears from our  
18 data that Lep through its receptor is able to affect both capacitation and acrosome  
19 reaction addressing their role in the pig sperm acquisition of fertilizing ability.

20 During the last few years, the study of the signaling events derived from Lep binding to  
21 its receptor has promoted a better understanding of the biochemical and molecular  
22 mechanisms of Lep function. The Ob-R is a member of the class I cytokine receptor  
23 superfamily that prevalently signals via association with non-receptor tyrosine kinases.  
24 Since Ob-R does not have intrinsic enzymatic activity, it activates associated JAK2,  
25 which auto-phosphorylates numerous tyrosine residues at the same time as it  
26 phosphorylates tyrosine residues on the functional Ob-R, upon ligand binding. The

1 phosphorylated intracellular domain then provides a binding site for STAT proteins.  
2 Although signals mediated directly by motifs on the Ob-R-associated JAK2 are poorly  
3 characterized, it is well known that Tyr<sub>985</sub> recruits the binding of the protein tyrosine  
4 phosphatase SHP-2 and the inhibitory suppressor of cytokine signaling-3, whereas  
5 Tyr<sub>1138</sub> specifically binds and activates the latent STAT3 (Munzberg & Myers 2005).  
6 The activation of STAT proteins in response to Lep was assayed in a variety of mouse  
7 tissues or cells known to express Ob-R. In our study, for the first time, we have  
8 demonstrated that STAT3 is expressed in the pig sperm, it is constitutively activated and  
9 its phosphorylation increased following Lep binding to its receptor. Here, we provide  
10 new information to indicate that Lep can stimulate the JAK-STAT pathway in pig sperm,  
11 resulting in a phosphorylation of STAT-3 depending on Lep concentration.

12 Since capacitation is a crucial step in the acquisition of sperm fertilizing ability, it is  
13 likely that it is controlled by redundant mechanisms and that cross talks between  
14 different pathways occur during this process (de Lamirande *et al.* 1997; Leclerc *et al.*  
15 1998). The Ras-Raf-MAPK (mitogen-activated protein kinase) pathway is also a  
16 downstream target of Ob-Rs (Takahashi *et al.* 1997; Tanabe *et al.* 1997) and several  
17 lines of evidence indicate that components of the extracellular signal-regulated kinase  
18 (ERK) family of mitogen-activated protein kinases (MAPK), as well as the key players  
19 in this cascade (Shc, Grb2, Ras, Raf-1, mitogen-activated protein kinase [MEK], ERK1,  
20 and ERK2), are present in spermatozoa and are involved in capacitation (Naz *et al.*  
21 1992; Luconi *et al.* 1998; de Lamirande & Gagnon 2002). Therefore, the activation of  
22 this pathway upon Lep was examined in pig sperm. Interestingly, we have shown that  
23 Lep positively modulates ERK1/2 through its receptor indicating that MAPK-dependent  
24 processes are involved in the hormone action.

25 Several pathways are also activated by JAKs including PI3K. Indeed, Lep signals via PI  
26 3-kinase in insulinoma cells (Harvey *et al.* 2000), muscle cells (Berti *et al.* 1997) and

1 hepatocytes (Zhao *et al.* 2000). Sperm capacitation is regulated by multiple pathways  
2 and among these signalling cascades the PI3K and its role in modulating sperm  
3 postejaculatory changes remain elusive. In somatic cells PI3K is implicated in many  
4 biological processes, including cell survival and chemotaxis, membrane ruffling and  
5 DNA synthesis, receptor internalization, and vesicular trafficking (Wymann & Pirola  
6 1998; Cantrell 2001). A role for PI3K has been also suggested in sperm functions such  
7 as survival and metabolism (Fisher *et al.* 1998; Luconi *et al.* 2001, Aquila *et al.* 2004,  
8 Aquila *et al.* 2005). In somatic cells, PI3K phosphorylates a large spectrum of protein  
9 substrates (Wymann & Pirola 1998). One of the main effectors of the PI3K, is the Akt  
10 (also named protein kinase B [PKB]), identified as a serine/threonine (Ser/Thr) protein  
11 kinase and it was also found in human ejaculated spermatozoa (Aquila *et al.* 2004).  
12 From our result it emerges that PI3K and Akt are implicated in mediating Lep signals in  
13 pig sperm, suggesting that this hormone may be involved in sperm survival.  
14 Concomitantly, Bcl-2 (Ito *et al.* 1997), a key protein in survival signaling, is enhanced  
15 upon Lep exposure and this effect was reduced by the anti-Ob-R Ab. Previous work  
16 have shown that Lep attenuates apoptosis of different cell types, such as osteoblasts,  
17 granulosa cells, and pancreatic islet cells (Almong *et al.* 2001; Okuya *et al.* 2001;  
18 Gordeladze *et al.* 2002). Moreover, very recently, Lep has been found to inhibit stress-  
19 induced apoptosis of T lymphocytes in vivo ( Fujita *et al.* 2002). Data presented here  
20 clearly demonstrate that Lep, inducing the phosphorylation of classical key survival  
21 proteins, such as p42/44 MAPK, PI3K, Akt and Bcl-2, allows pig sperm survival  
22 process.

23 It is important to point out that in all our experiments we obtained different responses by  
24 using low or high Lep doses, in a pattern in which only 10nM are stimulatory. The  
25 outcome of signalling activation can depend on differences in ligand concentration as it  
26 was demonstrated in human sperm (Marshall 1995; Aquila *et al.* 2005). Besides,



1 recently it was hypothesized that the net effect of Lep upon male reproductive function  
2 may depend on the circulating level of the molecule. Thus, predominant stimulatory  
3 effects are observed at Lep levels above a minimal threshold, in contrast, direct  
4 inhibitory actions may take place in the presence of a significantly elevated Lep  
5 concentration (Tena-Sempere & Barreiro 2002; Caprio *et al.* 2001).

6 It has been demonstrated that pig seminal plasma contains a significant amount of Lep,  
7 which decreases a lot in the follicular fluid (Lackey *et al.* 2002). As sperm leave seminal  
8 plasma during their transit in the female reproductive tract, they are exposed to  
9 decreased leptin concentrations. It may be speculated that the high Lep observed in  
10 seminal plasma may contribute to maintain sperm in a quiescent metabolic condition.  
11 Instead, the low Lep concentrations of the pig female reproductive tract secretions could  
12 reduce its inhibitory effect on sperm, thus rendering them suitable for capacitation and  
13 fertilizing ability acquisition.

14 Lep regulates food intake and energy expenditure via central and peripheral mechanisms.  
15 Fatness in pigs is of prime economic importance due to market incentives for production  
16 of lean pork and elevated fatness increasing the feed costs. Lep was identified as a  
17 metabolic signal affecting central regulation of reproduction in the pig and our data have  
18 shown that this hormone affects pig sperm acquisition of fertilizing ability. Therefore,  
19 on the basis of our results the manipulation of Lep concentration, suggesting a strategy  
20 for altering body composition, may affect pig reproduction.

## 21 22 23 **ACKNOWLEDGMENTS**

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### FIGURE LEGENDS

**Figure 1:** Western blot analysis of leptin and leptin receptor from pig sperm extracts

**A:** Immunoblot of leptin. *lane 1:* adipocyte cells used as positive controls(+), *lane 2:* negative control (-) *lanes P1, P2:* representative pig sperm samples **B:** Immunoblot of Ob-R from two representative pig sperm samples (P1, P2). **b1:** membrane was incubated with primary anti Ob-R Ab, **b2:** membrane was tested using the normal rabbit serum at the same dilution of the anti Ob-R Ab.

**Figure 2:** Representative immunofluorescence labelling of leptin and leptin receptor (Ob-R) in pig spermatozoa: **A** A red intense fluorescence localized leptin in the sperm acrosomal region. **B** A green brilliant light showed Ob-R in the apical acrosome.

*Inserts:* immunonegative controls. *Scale bars:* 5 $\mu$ m

1 **Figure 3:** Leptin affects cholesterol efflux and protein tyrosine phosphorylation of pig  
2 sperm. Spermatozoa were incubated in the absence (NC) or in the presence of leptin ( 10  
3 and 60 nM ) as well as in presence of anti Ob-R antibody + 10nM leptin. **A** Cholesterol  
4 in culture medium from pig sperm. Results are presented as mean  $\pm$  SEM and are given  
5 per  $10 \times 10^6$  spermatozoa. \*,  $P < 0.05$  vs.control **B** Western blot analysis of protein  
6 tyrosine phosphorylation from sperm lysates.

7  
8 **Figure 4:** Leptin affects acrosin activity of pig sperm. Spermatozoa were incubated in  
9 the absence (NC) or in the presence of leptin ( 10 and 60 nM ) as well as in presence of  
10 anti Ob-R antibody + 10nM leptin. Values are mean  $\pm$  SEM \*,  $P < 0.05$  vs.control.

11  
12 **Figure 5:** Western blot analysis of STAT 3 from one representative pig sperm lysate.  
13 **A:** Immunoblot of STAT3 in the pig sample (P). **B** p-STAT 3 band in sperm incubated  
14 in the absence (NC) or in the presence of leptin (10 and 60 nM ) as well as in presence  
15 of anti Ob-R antibody + 10nM leptin (Ab Ob-R+lep). MCF7 cell lines were used as  
16 positive controls (+). Negative control (-) was performed as previously described.  $\beta$ -  
17 actin served as loading control. **C** Band intensities were evaluated in term of arbitrary  
18 densitometric units. Values are as mean  $\pm$  SEM \*,  $P < 0.05$  vs.control

19  
20 **Figure 6:** Western blot analysis of ERK1/2 from one representative pig sperm lysate. **A**  
21 Immunoblot of ERK1/2 in the pig sample (P). **B** p-ERK1/2 band in spermatozoa  
22 incubated in the absence (NC) or in the presence of leptin (10 and 60 nM ) as well as in  
23 presence of anti Ob-R antibody + 10nM leptin (Ab Ob-R+lep). MCF7 cell lines were  
24 used as positive controls (+).Negative control(-) was performed as previously described.  
25  $\beta$ -actin served as loading control. **C** Band intensities were evaluated in term of arbitrary  
26 densitometric units. Values are as mean  $\pm$  SEM \*,  $P < 0.05$  vs.control



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**Figure 7:** Western blot analysis of Bcl-2 and Akt from one representative pig sperm lysate. **A, D** Immunoblots of Bcl-2 and Akt, in the pig sample (*P*). **B** p-Bcl-2 band when sperm cells were incubated in the absence (*NC*) or in the presence of leptin (*10 and 60 nM*) as well as in presence of anti Ob-R antibody + 10nM leptin (*Ab Ob-R+lep*). **E** p-AKTS (Akt ser) and p-AKTT (Akt thr) bands in sperm incubated in the absence (*NC*) or in the presence of leptin (*10 and 60 nM*) as well as in presence of anti Ob-R antibody + 10nM leptin (*Ab Ob-R+lep*) or LY 294002 (*LY*). **C, F:** Band intensities were evaluated in term of arbitrary densitometric units. Values are as mean  $\pm$  SEM \*,  $P < 0.05$  vs. control . (+) positive control. (-) negative control.  $\beta$ -actin served as loading control.

For Review Only

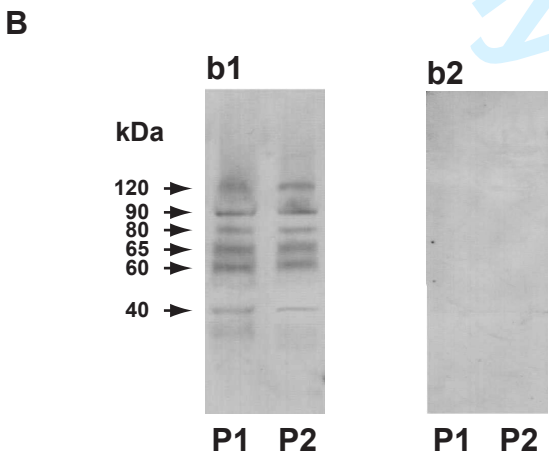
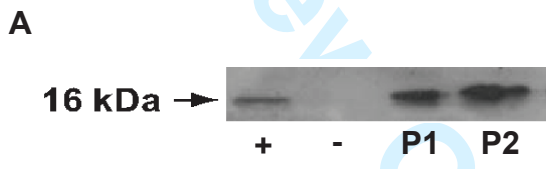
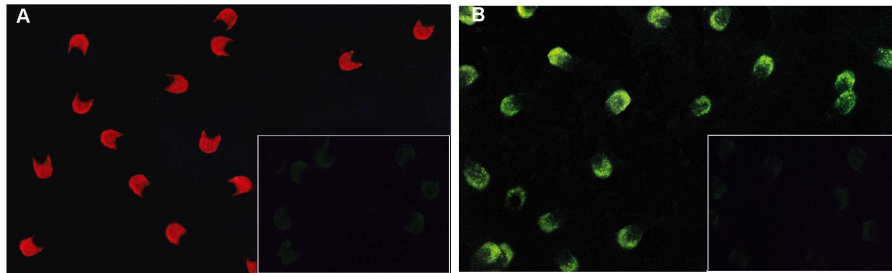
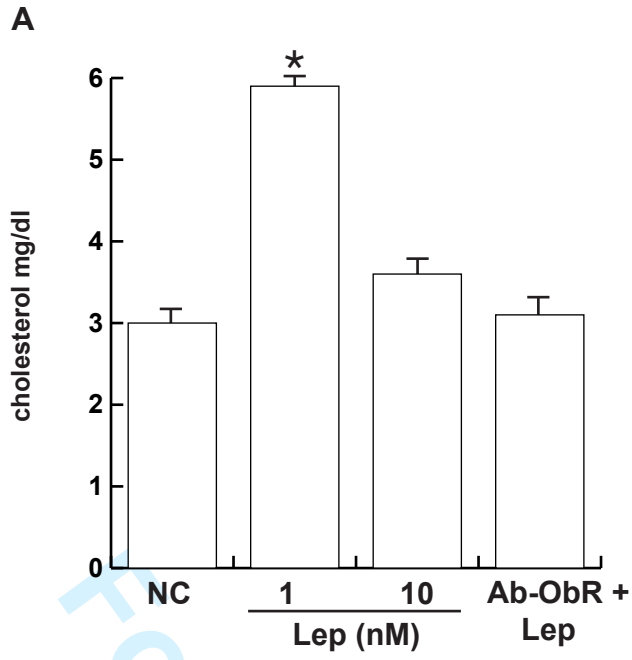


Fig. 1



**Figure 2: Representative immunofluorescence labelling of leptin and leptin receptor (Ob-R) in pig spermatozoa: A A red intense fluorescence localized leptin in the sperm acrosomal region. B A green brilliant light showed Ob-R in the apical acrosome. Inserts: immunonegative controls. Scale bars: 5  $\mu$  m  
180x180mm (300 x 300 DPI)**



**B**

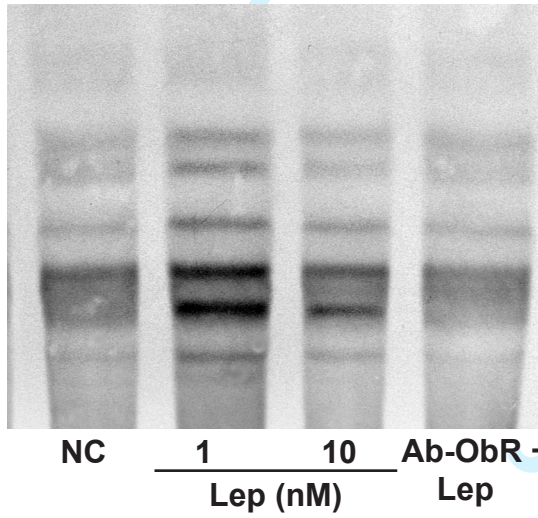


Fig. 3

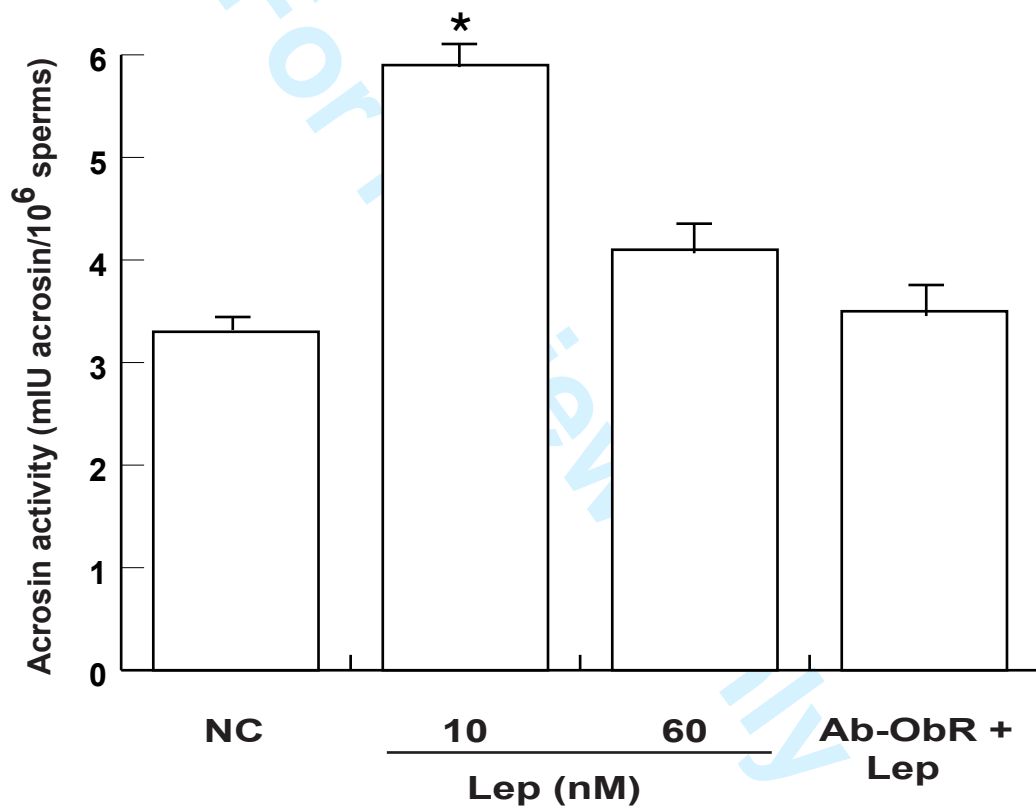


Fig. 4

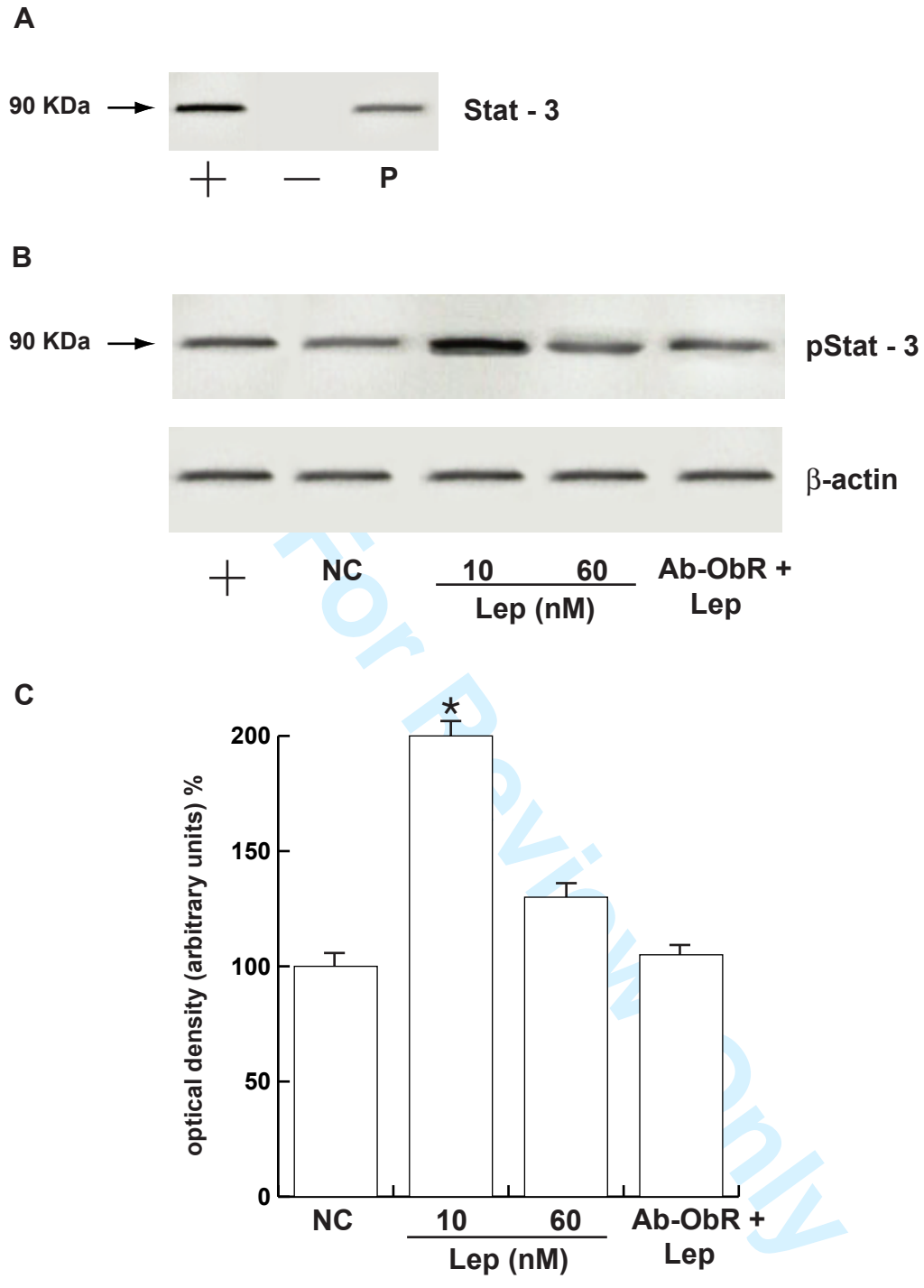
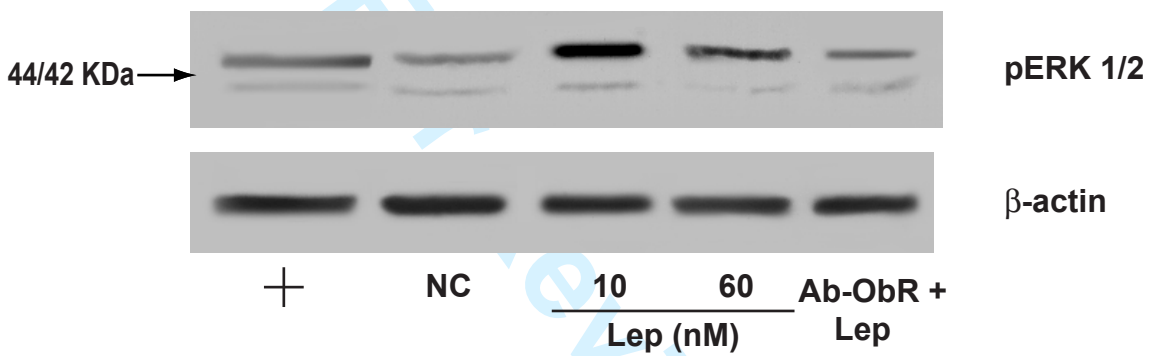


Fig. 5

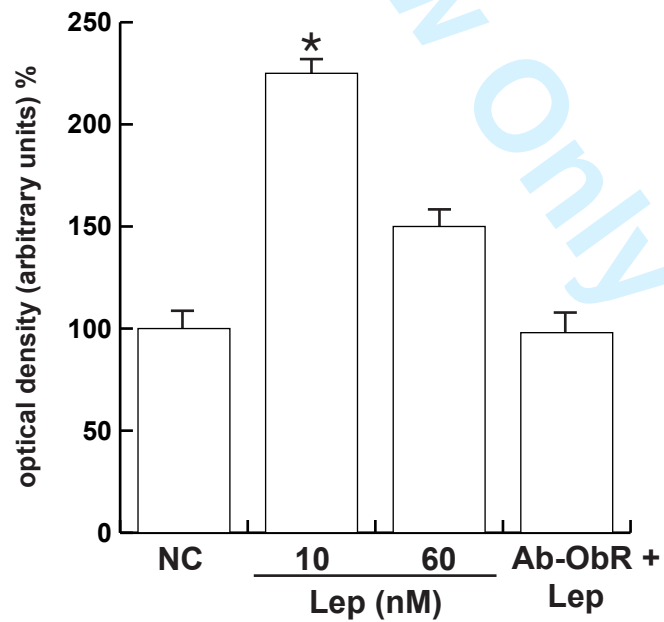
**A**



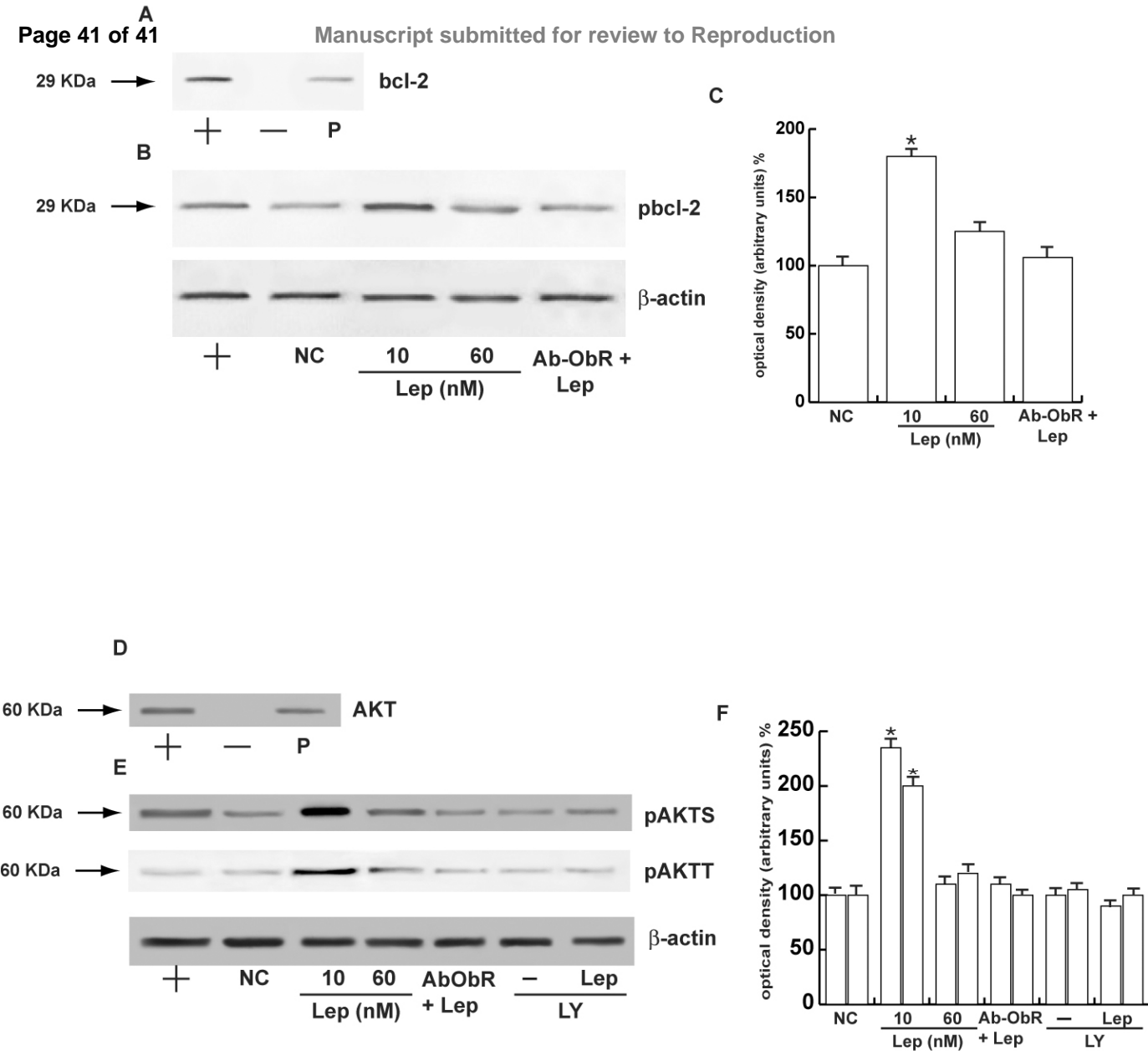
**B**



**C**



**Fig. 6**





## ATTI DI CONVEGNO

1. Francesca De Amicis, Silvia Zupo, Rocco Malivindi and Sebastiano Andò. "Evidences that progesterone receptor b decreases estrogen receptor gene expression through its interaction to a half-PRE site at estrogen receptor gene promoter." Atti del convegno "30th San Antonio Breast Cancer Symposium", San Antonio 13-16 December 2007.
2. De Amicis F. , Zupo S. , Malivindi R. , Ando' S. , "Il Recettore Progestinico B decrementa l'espressione del Recettore Estrogenico interagendo con un emisito-PRE presente sul promotore del Recettore Estrogenico ". Atti del convegno "X Congresso Nazionale AIBG", Torino (Italy), 2007.
3. Lanzino M. , De Amicis F. , Marsico S. , Zupo S. , Ando' S. , " ARA70 interagisce con il recettore estrogenico e modula il cross-talk funzionale tra recettore estrogenico e recettore androgenico in cellule MCF-7". Contributo a VIII Congresso AIBG, Sirolo (AN), 15-17 settembre 2005.
4. Lanzino M. , De Amicis F. , Zupo S. , Tirota E. , Ando' S. , " Endogenously expressed coactivator ARA70 interacts with ERalpha and modulates the functional interaction between ERalpha and AR in MCF-7 breast cancer cells". Atti del convegno "12th ICE", Lisbon, august 31-september, 2004.