

Exposure of human fallopian tube epithelium to elevated testosterone results in alteration of cilia gene expression and beating

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STUDY QUESTION: How does exposure to a testosterone rich environment affect the function and gene expression of human fallopian tube epithelium (hFTE)?

SUMMARY ANSWER: Elevated testosterone level alters several gene transcripts that regulate cilia expression and negatively impacts the rate of cilia beating.

WHAT IS KNOWN ALREADY: The presence of estrogen in the follicular phase of the menstrual cycle increases the human fallopian tube ciliary beating frequency. The luteal phase, triggered by ovulation and increasing progesterone, is marked by a decrease in ciliary beating. Women with polycystic ovarian syndrome (PCOS) may have twice the serum level of testosterone than ovulatory women. To date, the effect of elevated androgens on the function of the human fallopian tube is not well-understood. We chose to examine the impact of elevated testosterone on hFTE.

STUDY DESIGN, SIZE, DURATION: A prospective basic science study of human fallopian tube specimens from reproductive-aged women undergoing benign gynecologic surgery was performed. Fallopian tube removal at a large US academic center was collected and provided to us to continue with epithelium isolation and culturing. A total of 12 patients were analyzed in the study.

PARTICIPANTS/MATERIALS, SETTING, METHODS: Fallopian tube epithelium was isolated and exposed to two different conditions: normal with low testosterone concentration of 0.8 nM and PCOS-like, with high testosterone concentration of 2 nM. The study was conducted in both static and dynamic conditions in microfluidic devices for a total of 14 days, after which the tissue was collected for processing including RNA extraction, quantitative PCR and immunohistochemistry. After the first 7 days of each experiment, a sample of tissue from each condition was imaged to quantify cilia beating frequency.

MAIN RESULTS AND THE ROLE OF CHANCE: hFTE exposed to the 2 nM testosterone displayed slower cilia beating, inhibited estrogen signaling and decreased expression of the ciliary marker FOXJ1 when compared to stimulation with 0.8 nM testosterone.

LARGE SCALE DATA: N/A.

LIMITATIONS, REASONS FOR CAUTION: The *in vivo* response to elevated testosterone may differ from *in vitro* studies. RNA amount was limited from tissue cultured in the microfluidic devices as compared to static culture.

WIDER IMPLICATIONS OF THE FINDINGS: Understanding elevated testosterone in tubal function may explain an additional contribution to subfertility in women with PCOS and other hyper-androgen disorders, aside from oligo-ovulation. Furthermore, this adds to the body of literature of fallopian tube function using a microfluidic device.

STUDY FUNDING/COMPETING INTEREST(S): NIH grants: UH3 ES029073 and R01 CA240301. There are no competing interests.

Key words: fallopian tube / hyperandrogenism / estrogen / polycystic ovarian syndrome / microfluidic / cilia / FOXJ1

Introduction

The fallopian tubes are the site of fertilization but are also necessary for transport of the sperm, egg, and developing embryo, as well as tubal secretions which influence embryo development (Bui *et al.*, 2000; Hunter *et al.*, 2005; Lyons *et al.*, 2006a,b). The mechanisms involved in tubal transport include muscular contractions, flow of tubal secretions and the beating of cilia, which are hair-like projections extending into the tubal lumen (Jansen, 1984). It has been suggested that ciliary motion may play a critical role in fallopian tube transport (Lyons *et al.*, 2006a,b). With widespread availability of IVF, the physiology of the human fallopian tube has become increasingly overlooked. However, for mammalian *in vivo* fertilization (IVF), the fallopian tube is essential for reproduction (Lyons *et al.*, 2006a,b). When ciliary motion, and thus tubal function, is disrupted through trauma to the tubal epithelium by sexually transmitted infections, pelvic inflammatory disease or environmental exposures such as cigarette use, there is a significant negative impact on fertility (McGee *et al.*, 1981; Cooper *et al.*, 1986; Cooper *et al.*, 1990; McGee *et al.*, 1999; Bouyer *et al.*, 2003). In addition, cigarette smoke inhibits aromatase activity, suggesting that it may play a role in reproduction through increased testosterone levels (Biegon *et al.*, 2012).

The tubal epithelium responds to hormonal changes during the menstrual cycle (Novak, 1928; Critoph and Dennis, 1977a). Ovarian hormones regulate tubal epithelial structure and the expression of cilia (Lyons *et al.*, 2006a,b). Estrogen stimulates epithelial cell hypertrophy, secretion and ciliogenesis, while high levels of progesterone, as seen in the luteal phase, are associated with epithelial atrophy and deciliation (Verhage *et al.*, 1979). Animal studies found an increase in cilia beating frequency (CBF) following copulation, presumed to be due to high estrogen levels and the need for gamete transport (Borell *et al.*, 1957). This finding was corroborated in human fallopian tube specimens with an increase in CBF after ovulation, when estrogen predominates, and decreased CBF later in the progesterone dominant luteal phase (Critoph and Dennis, 1977b).

Ciliary beating frequency in response to estrogen and progesterone was studied in an *ex vivo* model using human fallopian tube epithelium (hFTE) specimens and a stepwise exogenous hormone preparation (Zhu *et al.*, 2016). A static culture model was able to maintain viability of hFTE for the duration of the 14 days experiment. After 7 days of hormone exposure, the CBF was significantly increased with estrogen exposure in a dose–response pattern, with a 25% increase in CBF in the tissue exposed to higher estrogen and in the absence of progesterone. However, when the hFTE was exposed to estrogen combined with progesterone, the CBF decreased by 30%. One effect not considered in this study was the influence of androgens, including testosterone, on CBF and overall tubal function.

In ovarian steroidogenesis, estrogen is made in the granulosa cells by the aromatization of precursor androgens, which originate from ovarian theca cells (Stocco, 2008). Ovarian androgens are continuously produced throughout the menstrual cycle (Lebbe and Woodruff, 2013). Testosterone peaks a couple of days after estrogen (Salonia *et al.*, 2008) suggesting that in physiological conditions it does not interfere with estrogen action. Ovulation is a sentinel event changing the hormonal environment of the human reproductive tract. However, in anovulatory or oligo-ovulatory women, the lack of cyclical progesterone production, and thus prolonged estrogen and testosterone exposure, can have various clinical consequences including subfertility and endometrial hyperplasia, which can result in endometrial cancer (Kim *et al.*, 2013). Clinically significant disruptions in ovulation most often occur in women with polycystic ovarian syndrome (PCOS) and during the perimenopausal transition.

PCOS is the most common endocrinopathy of reproductive-aged women with an incidence of up to one in eight women of reproductive age. Despite being a common diagnosis worldwide, there is no universally accepted definition of the condition. In fact, several criteria are used to determine the diagnosis based on the presence of hyperandrogenism, oligoamenorrhea or amenorrhea, and polycystic morphology of the ovaries on ultrasound (Rotterdam ESHRE/ASRM-Sponsored PCOS Consensus Workshop Group, 2004). Among these, hyperandrogenism is the most consistently agreed upon criterion between expert panels. Testosterone is the most frequently tested androgen in the clinical diagnosis of PCOS. Although there is no cutoff for the definition of hyperandrogenemia, women with PCOS have been noted to have two times the level of testosterone than ovulatory women (Winters *et al.*, 2000). The paracrine and endocrine actions of ovarian testosterone on the human fallopian tube are of interest as there is evidence that ovarian hormones are very highly concentrated in the ovarian follicular fluid, which is found in the fallopian tubes after ovulation (Emori and Drapkin, 2014). Furthermore, the event of ovulation is a risk factor for high grade serous ovarian cancer that originates in the fallopian tube (Fathalla, 2013). There is emerging evidence of elevated circulating testosterone increasing the risk of epithelial ovarian cancers (Ose *et al.*, 2017). Exploring the sequelae of elevated testosterone exposure on fallopian tube health and function may be important to understanding the lifetime health risks for women living with PCOS and/or hyperandrogenism.

This study sought to investigate the impact of androgens on hFTE using *ex vivo* models that were previously validated in a microfluidic device (Zhu *et al.*, 2016; Xiao *et al.*, 2017). Our objective was to evaluate the impact of elevated testosterone on hFTE function and gene expression in static and dynamic conditions used for primary human explant culture. Given the importance of tubal cilia to reproduction, we choose to examine cilia beating and cilia related genes as a marker of overall tubal health.

Materials and methods

Chemical and reagents

Seventeen beta-estradiol (E2) was purchased from innovative research of America (NE-121). Testosterone (T1500), rhFSH (F4021) and fetuin (F3385) were purchased from Sigma (St. Louis, USA). F12 was purchased from Gibco (1967615). Alpha minimum essential medium (MEM) was purchased from Corning (10-022-CV). ITS (5 µg/ml insulin, 5 µg/ml transferrin and 5 ng/ml selenium) was purchased from Sigma (11074547001).

Human fallopian tissue

Human fallopian tube specimens were obtained from hysterectomies and salpingectomies done for benign indications in pre-menopausal women, aged 35–50, at Northwestern University Prentice Women's Hospital (Chicago, IL, USA) from 1 October 2018 to 30 November 2019. Written informed consent was obtained from each subject prior to surgery. Each portion of fallopian tube received was cut lengthwise and the inner epithelium and underlying stroma were removed with blunt dissection using forceps. The tissue was dissected in MEM and Nutrient Mixture F-12 medium (50:50) with 10% fetal bovine serum within a 24-h period from arrival. The tissue was then transferred to MEM with 0.3% bovine serum albumin, 0.5 mg/ml fetuin, 1% penicillin/streptomycin and ITS medium. The epithelium was then isolated from the underlying stroma, again with blunt dissection using forceps, and fashioned into 2 × 2 mm sections for culture. Tissues from the same patient were divided into normal and PCOS-like conditions and cultured on 0.4 µm pore Millicell inserts (Millipore, MA, USA). After 24 h, the static condition experiments continued in static culture in 12-well plates while microfluidic experiments were transferred to the microfluidic devices for dynamic culture. Both static and microfluidic conditions were maintained in incubators at 37°C and 5% CO₂ for 14 days in one of the two experimental conditions. The physiologic media condition, referred to as 'normal' contained estrogen (E₂) 0.1 nM and testosterone 0.8 nM on Days 0–6 then E₂ 1 nM and testosterone 0.8 nM Days 7–13 and finally E₂ 1 nM and testosterone 1.25 nM for 1 day prior to the end of the experiment on Day 14. The hyperandrogenic media conditions referred to as 'PCOS' contained E₂ 0.1 nM and testosterone 2 nM on Days 0–6 then E₂ 1 nM and testosterone 2 nM Days 7–14 until the experiment ended on Day 14. In this model, recombinant follicle-stimulating hormone (rFSH) was present in the media from Days 0 to 14, to replicate the follicular phase of the menstrual cycle. After 14 days of culture, the tissue was either frozen for RNA extraction or fixed in 4% paraformaldehyde for immunohistochemistry.

Ethical committee approval

Written informed consent was obtained from each subject and experiments, procedures and methods were performed in accordance with the IRB-approved guidelines and regulations (IRB# STU00205203).

RNA sequencing

RNA was isolated from cells using Qiagen RNeasy Mini kit (#74104) as per manufacturer's protocol. RNA libraries (three technical

replicates) were created. RNA quality determination, mRNA enrichment, library construction, sequencing and transcriptome statistical analysis were performed at the Genomics Core Facility at Northwestern University as previously described (Colina et al., 2019).

Microfluidic systems

Prior to all microfluidic experiments, all systems were sterilized and subsequently assembled under sterile conditions. The single microfluidic platform (Solo-MFP) system operation was previously described (Xiao et al., 2017) (Fig. 1). All fallopian tube epithelium isolated as above was cultured on 0.4 µm pore Millicell inserts and the microfluidic experiments were placed into the Solo-MFP tissue culture modules containing 200 µl of the corresponding culture media, either normal or PCOS. An additional 2 ml of media was placed in each donor module, for circulation throughout the Solo-MFP system. Acceptor modules were sampled daily, and the collected media stored at –20°C prior to analysis. The flow rate was between 35 and 50 µl/h. The donor modules were refilled daily.

Immunohistochemistry

After 14 days, cultured tissue was fixed for 24 h in 10% paraformaldehyde. Tissues were then dehydrated in increasing concentrations of 50–70% ethanol. Dehydrated tissues were then embedded in paraffin using an automated tissue processor. Later 5 µm serial sections were created for hematoxylin and eosin (H&E) and immunohistochemistry staining as previously described (Russo et al., 2018). Antibodies used are described in Supplementary Table S1. In brief, tissues were incubated with primary antibodies overnight at 4°C and secondary for 1 h at room temperature.

Imaging and quantification of CBF

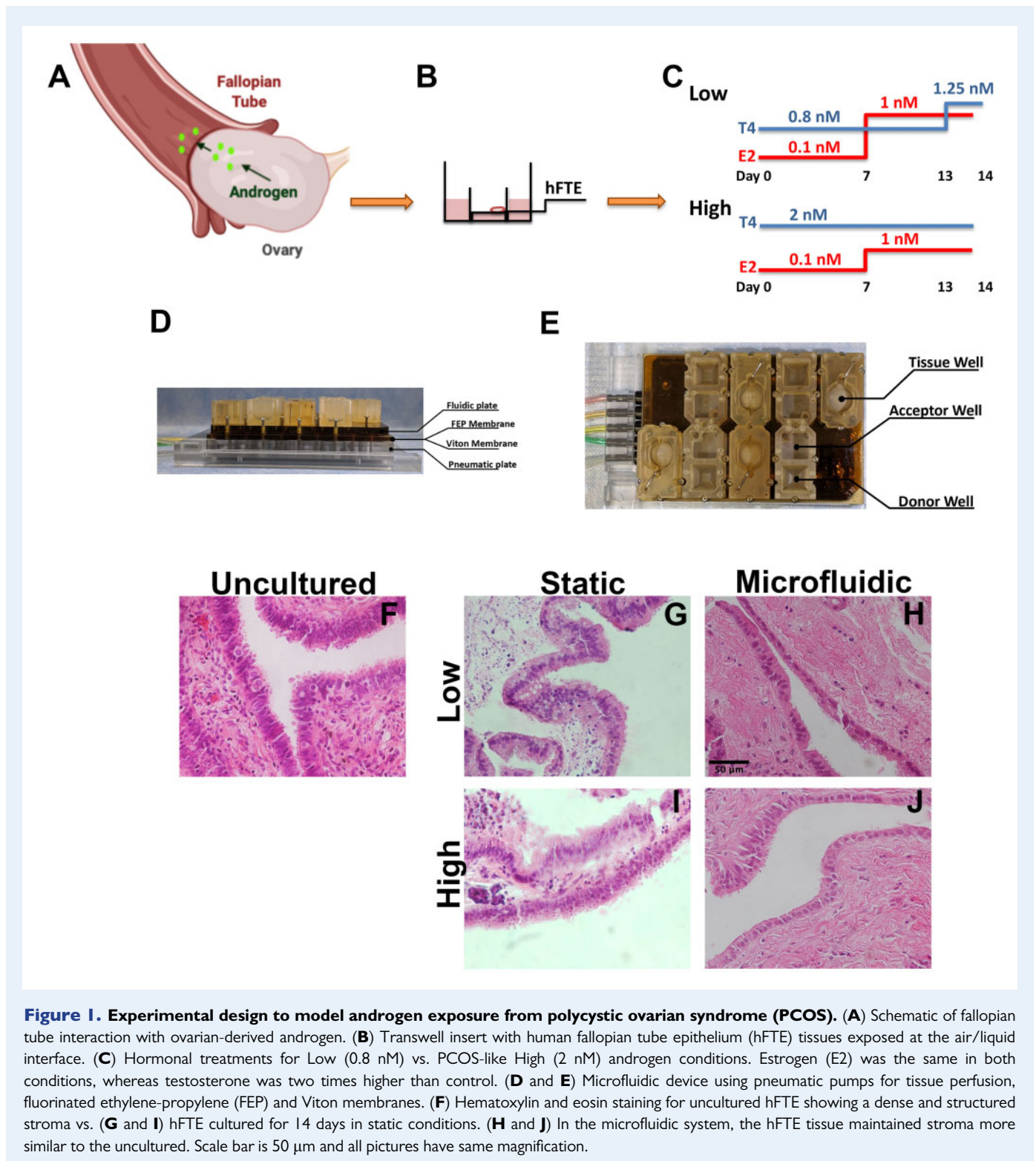
After 7 days of culture, the hFTE was imaged using a Nikon CFI SR HP Plan Apo Lambda S 100xSil Objective 1.35NA on a Nikon Ti2 microscope stand and captured with a Photometrics Prime 95B 25 mm camera with 5 ms frame rate. Five movies in different areas of the sample were taken. The movies were analyzed within ImageJ Fiji software by creating kymograph images to determine the CBF in Hertz (Hz) (Zhu et al., 2016).

Immunoassay analysis

Vascular endothelial growth factor (VEGF) and insulin-like growth factor (IGF) concentrations in media collected from both the static and microfluidic experiments were measured using commercial ELISA kits (RayBiotech, Peachtree Corners, GA, USA) according to the manufacturer's protocols. A 100 µl of conditioned media from hFTE from three subjects at Day 14 of treatment, were loaded on ELISA plates in duplicate.

Quantitative PCR

Tissues were homogenized in lysis buffer plus beta-mercaptoethanol (RLT buffer) using zirconia beads (Invitrogen, Waltham, MA, USA) and the Mini-BeadBeater cell/tissue disrupter (BioSpec, Bartlesville, OK, USA). Tissues were centrifuged and supernatant processed for RNA extraction as described by RNeasy kit manufacturer's instructions



(Qiagen, Germantown, MD, USA). iScriptTM cDNA synthesis kit (BioRad, Hercules, CA, USA) and SYBR green (Roche, Madison, WI, USA) were used according to manufacturer's instructions. All quantitative PCR (qPCR) runs were performed on the BioRad machine. Primers used are listed in [Supplementary Table SII](#).

Statistical analysis

All data are represented as mean \pm SE. Statistical analysis was carried out using Prism software (GraphPad, La Jolla, CA, USA). All conditions were tested in three replicates in at least triplicate experiments. Statistical significance was determined by Student's *t*-test, or one-way

ANOVA with Dunnett's *post hoc* test or two-way ANOVA. * $P \leq 0.05$ was considered significant.

Results

A PCOS mimetic protocol for human fallopian tube in a microfluidic system

An *in vitro* model for studying the role of fallopian tube epithelium in PCOS was created to simulate a key feature of the disease, the presence of increased androgen, with a concentration of testosterone approximately two times higher (2 nM) than what is detected in control (0.8 nM). The hFTE was isolated from tissue collected from women having surgery for benign gynecological conditions and cultured in the normal (Low) or PCOS-like (High) conditions for 14 days (Fig. 1C). The same culture medium and concentration of hormones were used for tissues in static versus microfluidic conditions. In both cases, the tissues were added to transwells (Fig. 1A and B) inserted into 12-well plates (static) or into microfluidics wells (Fig. 1D and E). In the microfluidic system, the tissues maintained its structure with a more compact stroma that resembled uncultured tissue as shown through H&E staining (Fig. 1F–J).

Hormonal responses for fallopian tube epithelium in static or dynamic cultures

Previously, hFTE was cultured in the microfluidic device in the presence of hormones to model the normal reproductive cycle. VEGF has been shown to be induced in the presence of the estrogen-dominated phases (Zhu et al., 2016) and to play a role in embryo implantation and is expressed and secreted by hFTE (Zarezade et al., 2015). In order to compare the role of elevated testosterone levels, the amount of VEGF-A secretion was quantified. Reduced VEGF-A secretion was found in PCOS-like (High) vs. normal (Low) in the static condition, but not in microfluidic conditions potentially due to the effect of increased flow (Fig. 2A). In the microfluidic system, secretions are distributed by the constant flow of the pumps and this results in the generation of gradients that might result in increased dilution, limiting detection. IGF has been shown to be secreted by fallopian tube and to play a role in embryo pre-implantation (Winger et al., 1997) and development (Kirby et al., 1996). IGF-1 secretion was not changed (Fig. 2B). The reduced VEGF-A secretion suggests an anti-estrogenic effect of testosterone. In order to confirm the impact on estrogen receptor signaling, a well-known estrogen-dependent target, OVGPI (estrogen target oviduct-specific glycoprotein 1), was measured and the protein expression was downregulated in the presence of testosterone in static and fluidic conditions (Fig. 2C–F). OVGPI downregulation was also confirmed by qPCR in the static condition where estrogen receptor and the estrogen-dependent target, OVGPI, were both downregulated in the presence of testosterone (Fig. 2G).

Androgen and estrogen receptor localization were analyzed using immunohistochemistry in normal (Low) and PCOS-like (High) conditions. In both Low static and Low fluidic culture conditions, the androgen receptor was localized more in the cytoplasm, while in the PCOS-like High conditions, there was more receptor localized in the nucleus (Figs 3A and B and 3E and F). In contrast, estrogen receptor was

localized less in the nucleus in High condition treated hFTE as compared to Low (Figs 3C and D and 3G–F). The receptor localization further suggests that the increased testosterone concentration was activating androgen receptor and thereby its nuclear accumulation, while inhibiting estrogen receptor activity and causing it to reside in the cytosol.

Androgen stimulation alters cilia genes profile

Fallopian tube epithelium expresses androgen receptors (Home et al., 2009); however, very few analyses have been reported on the signaling pathways of androgens in the fallopian tube, particularly using primary human tissue. To gain a better insight into the androgen-regulated pathways, RNA-sequencing (RNAseq) was performed from RNA extracted from tissues treated with High vs. Low testosterone concentrations. Unbiased two-way hierarchical clustering analysis revealed transcripts were most similar between individual patients (Supplementary Fig. S1). The RNAseq identified 556 transcripts differentially and significantly altered upon exposure to high testosterone concentrations. GSEA (gene set enrichment analysis) and Cytoscape clustering revealed treatment with high testosterone-regulated pathways associated with ciliary function and assembly (Supplementary Figs S2 and S3) with *FOXJ1* and *LRRC6* appearing in the GSEA leading edge analysis. Based on the data analysis, we generated a heat map of cilia transcripts regulated upon high testosterone stimulation in hFTE that were involved in cilia function and cilia machinery, including *FOXJ1*, *LRRC6* and *KIF5C* (Fig. 4A). Other members of the LRRC dynein family were also regulated. Dyneins are involved in the retrograde intraciliary transport and are balanced with kinesins that regulate anterograde transport (Fig. 4B). A member of the kinesins *KIF5C* showed up in our top 20 upregulated genes (Fig. 4A). *FOXJ1*, *LRRC6* and *KIF5C* were therefore further validated in static conditions using qPCR (Fig. 4C).

Androgen stimulation alters the frequency of cilia beating

In order to further validate the RNAseq, we analyzed the expression and localization of cilia proteins in hFTE tissues by immunohistochemistry. In particular acetylated tubulin was chosen to stain cilia to confirm the cilia structure. FOXJ1, which is a master regulator of cilia function, was also selected as this transcription factor regulates expression of important proteins that constitute cilia. In both static (Fig. 5A and B) and fluidic cultures (Fig. 5E–H), cilia structures were present based on acetylated tubulin staining in Low and High conditions. However, FOXJ1 expression and localization in the nucleus, which indicates its activity, was reduced with increased testosterone in High conditions (Fig. 5C and D) and (Fig. 5G and H). This suggests that cilia function may be altered by elevated testosterone. After 7 days of culture, we imaged the hFTE using a Nikon Ti2 microscope and found that elevated testosterone exposure reduces CBF (Fig. 6A and B).

Discussion

This study investigated the impact of androgens on *ex vivo* models of primary hFTE. We chose to focus on testosterone as this is most

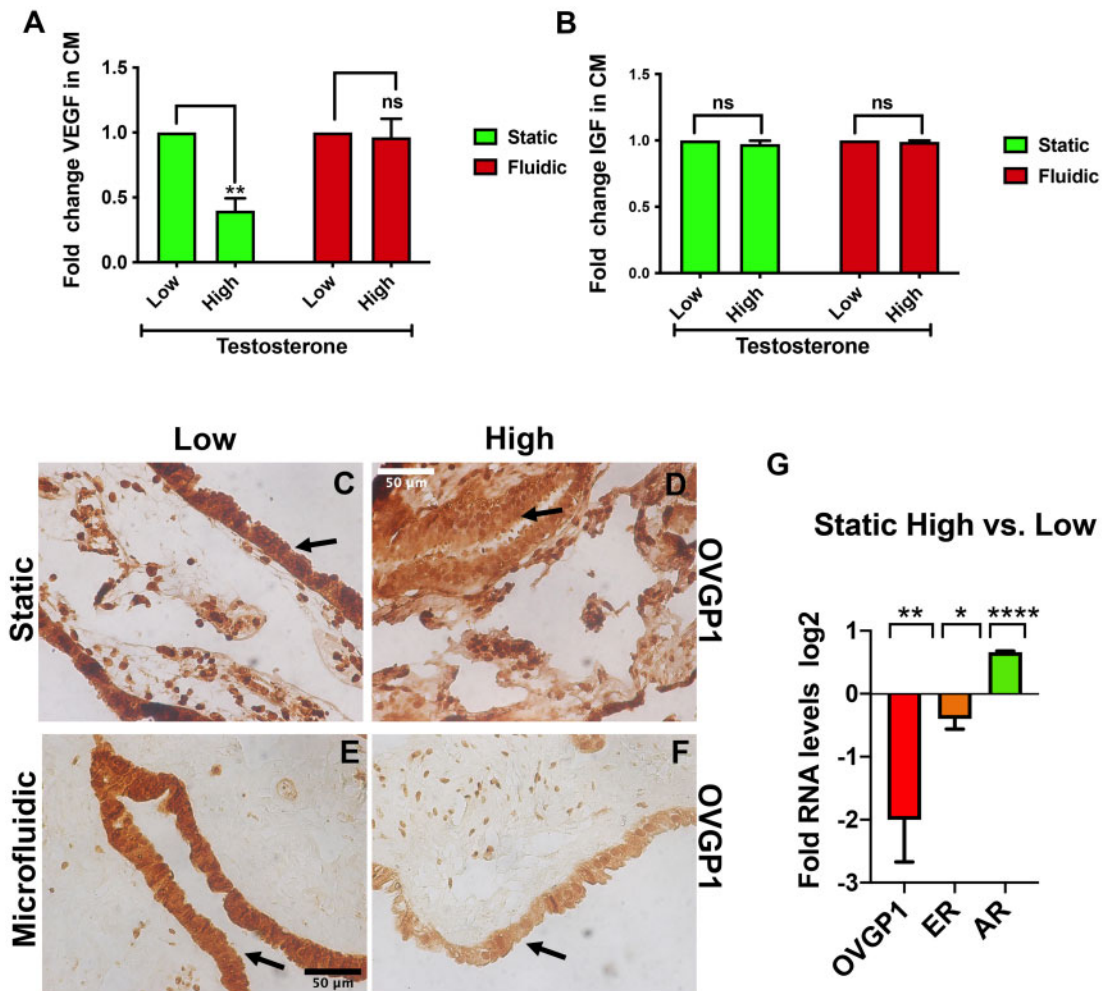


Figure 2. Androgen-induced response on human fallopian tube static vs. dynamic conditions. (A) ELISA for vascular endothelial growth factor-A (VEGF-A) in conditioned media from hFTE treated with low testosterone vs. high, show a significant decrease in VEGF-A after 14 days. (B) Insulin-like growth factor-I (IGF-I) secretion was unchanged in the presence of high testosterone. (C and D) Decreased oviduct-specific glycoprotein-I (OVGP1) protein, in static culture, after exposure to high testosterone is compared to low testosterone using immunohistochemistry (IHC). Scale bar is 50 μ m and all pictures have same magnification. (E and F) Decreased OVGP1 protein in the presence of high testosterone as compared to low using IHC in the microfluidic. (G) Reduction of *OVGP1* expression confirmed by quantitative PCR (qPCR) in static tissues treated for 14 days. Experiments include four different patients and graph are represented as mean \pm SEM. * $P \leq 0.05$; ** $P \leq 0.01$; **** $P \leq 0.0001$

clinically relevant and is documented as the predominantly elevated androgen in women with PCOS (Stocco, 2008). Excessive testosterone stimulation of hFTE altered the genomic profile, specifically of genes related to ciliary assembly and function. In addition, higher testosterone concentration altered CBF (Cilia Beating Frequency) and hormonal signaling. These results suggest that the hyperandrogenic environment, found in PCOS patients, may impact human fallopian tube function in ways that can modulate reproduction, as compared to a physiologic ovulatory environment.

Herein, we addressed the role of androgen on hFTE in static and dynamic culture and showed that human fallopian tube maintains a healthier and compact structure, when cultivated in microfluidic conditions (Fig. 1), consistent with other studies showing that dynamic condition preserves fallopian and ovarian health (Zhu et al., 2016; Xiao

et al., 2017). Our model is based on isolation of the epithelium monolayer that allows a reduced thickness of the tissue that is critical for high-resolution microscopy of CBF in addition to tissue exposure to nutrients and hormone stimulation.

In vivo study of fallopian tube epithelial function is contraindicated in humans as the potential for trauma to the fallopian tubes can predispose to scarring, infection, ectopic pregnancy and infertility (Lyons et al., 2006a,b). Therefore, building upon previous work using *ex vivo* cultures of hFTE, we designed experiments using exogenous hormones and investigated their role on hFTE explants. We also chose to use exogenous hormones as opposed to co-culturing with murine ovaries as a way to test our hypothesis and eliminate the impact of other ovarian derived hormones or peptides which could affect tubal function. Limitations of this study include limited sample size and consistency in

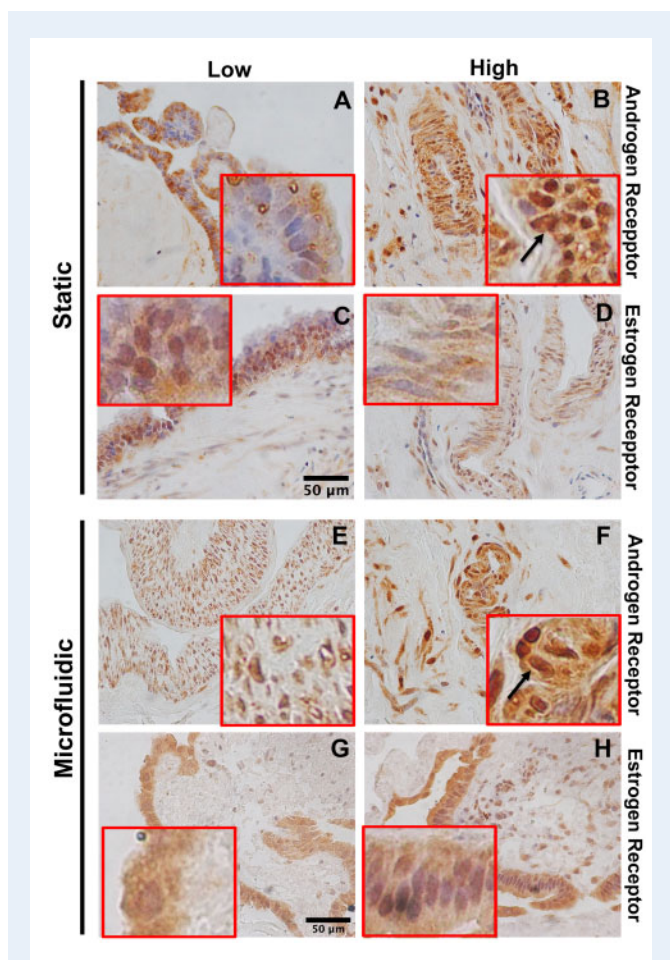


Figure 3. Estrogen and androgen receptor expression in responses to hFTE static vs. dynamic conditions in response to high androgen. IHC for androgen receptor (A and B) and estrogen receptor (C and D) in static conditions. IHC for androgen receptor (E and F) and estrogen receptor (G and H) in microfluidic conditions (E–H). Scale bar 50 μm . Magnification is the same for all pictures. Red squares show a 4 \times magnification of the image to visualize receptor localization.

terms of age, diagnosis and potential contraception or treatment of benign gynecologic conditions, as well as the intrinsic variability between human specimens. Future work could examine the relevance of our findings using either a mouse or rat model of PCOS, both of which have been developed using exposure to either dehydroepiandrosterone or testosterone by repeated injection or ingestion that gives rise to ovaries with some biological characteristics of human disease (Paixao et al., 2017).

Androgen exposure in the fallopian tube epithelium reduced estrogen receptor as well as the estrogen target OVGPI, which is produced by fallopian secretory cells. OVGPI facilitates sperm capacitation and motility for fertilization (Erickson-Lawrence et al., 1989; Verhage et al., 1997) suggesting a potential role for testosterone on sperm function through reduction of OVGPI. VEGF-A, which increases during periovulatory stages (Lam et al., 2003a,b), was reduced with high

concentration androgen treatment, consistent with less nuclear estrogen receptor and therefore less estrogen receptor action.

RNA-sequencing analysis of hFTE after high testosterone exposure revealed repression of *FOXJ1*, a master regulator of ciliogenesis, that modifies the expression of other cilia-related genes (Yu et al., 2008). *FOXJ1* affects the assembly of the cilium by anterograde transport (Johnson and Rosenbaum, 1992) and cilia turnover via retrograde transport (Rosenbaum and Witman, 2002; Scholey, 2003; Ishikawa and Marshall, 2011). Duloherly et al. (2019) characterized the alterations to the fallopian tube after surgical removal in female-to-male transsexuals who had undergone testosterone therapy for 1–3 years prior to surgery, as compared to normally cycling controls and found luminal accumulations of thick mucus-like secretions and cellular debris, which caused ciliary clumping and potential luminal blockage. The abundance of viscous secretions in the lumen may be a mechanism by which hyperandrogenism impairs fertility and decreases ciliary beating frequency by ciliary entrapment. However, this was not tested, and mucus accumulation may be actually a consequence of cilia impairment. This study corroborates our own as both studies show the immediate and long-term impact of testosterone on fallopian tube epithelium to be disruptive to normal function.

Tubal transport may play the dominant role in transport of gametes and contributes to early embryo development (Lyons et al., 2006a,b). Cilia beating is an integral component of the reproductive function of fallopian tubes, but tubal contraction and secretions are also required (Jansen, 1984). Nevertheless, experiments that blocked tubal contractility still allowed for transport of ovum suggesting that the cilia alone can facilitate this task (Halbert et al., 1976; Halbert et al., 1989). Genetic conditions, such as primary ciliary dyskinesia, impair ciliary motion and affected women with varying degrees of infertility based on the specific genes involved and thus the degree of ciliary motility (Halbert et al., 1997; Raidt et al., 2015). In women with primary ciliary dyskinesia, who present with subfertility (Afzelius et al., 1978; Pedersen, 1983; Lurie et al., 1989), rare cases of successful pregnancy may be explained by the presence of additional tubal factors in the absence of normal ciliary function, which allow for gamete and embryo transport (McComb et al., 1986; Halbert et al., 1997). In a study of 36 women with primary ciliary dyskinesia, only 61.1% were found to be infertile. However, in the infertile patients, mutations in *LRRC50* and *LRRC6* that encode for cilia machinery have been found (Vanaken et al., 2017) suggesting that the gene profile of ciliary components is a critical contribution to infertility in addition to cilia beating. Furthermore, common sexually transmitted reproductive pathogens cause deciliation and acute and chronic inflammation in the fallopian tube (McGee et al., 1981; Cooper et al., 1986). CBF is significantly lower in the surviving cilia of fallopian tubes showing evidence of edema, dilation or distal obstruction as a result of salpingitis (Leng et al., 1998). This deciliation appears to be irreversible which significantly increases infertility and ectopic pregnancy risk (Cooper et al., 1990).

There is a paucity of published data on the risk of ectopic pregnancy in women with PCOS. One retrospective cohort study found a higher ectopic pregnancy risk in women with PCOS who underwent controlled ovarian hyperstimulation (COH) with subsequent embryo transfer compared to women without PCOS (Wang et al., 2013). Higher estrogen levels in women without PCOS also conferred a higher ectopic pregnancy risk than women with lower estrogen levels,

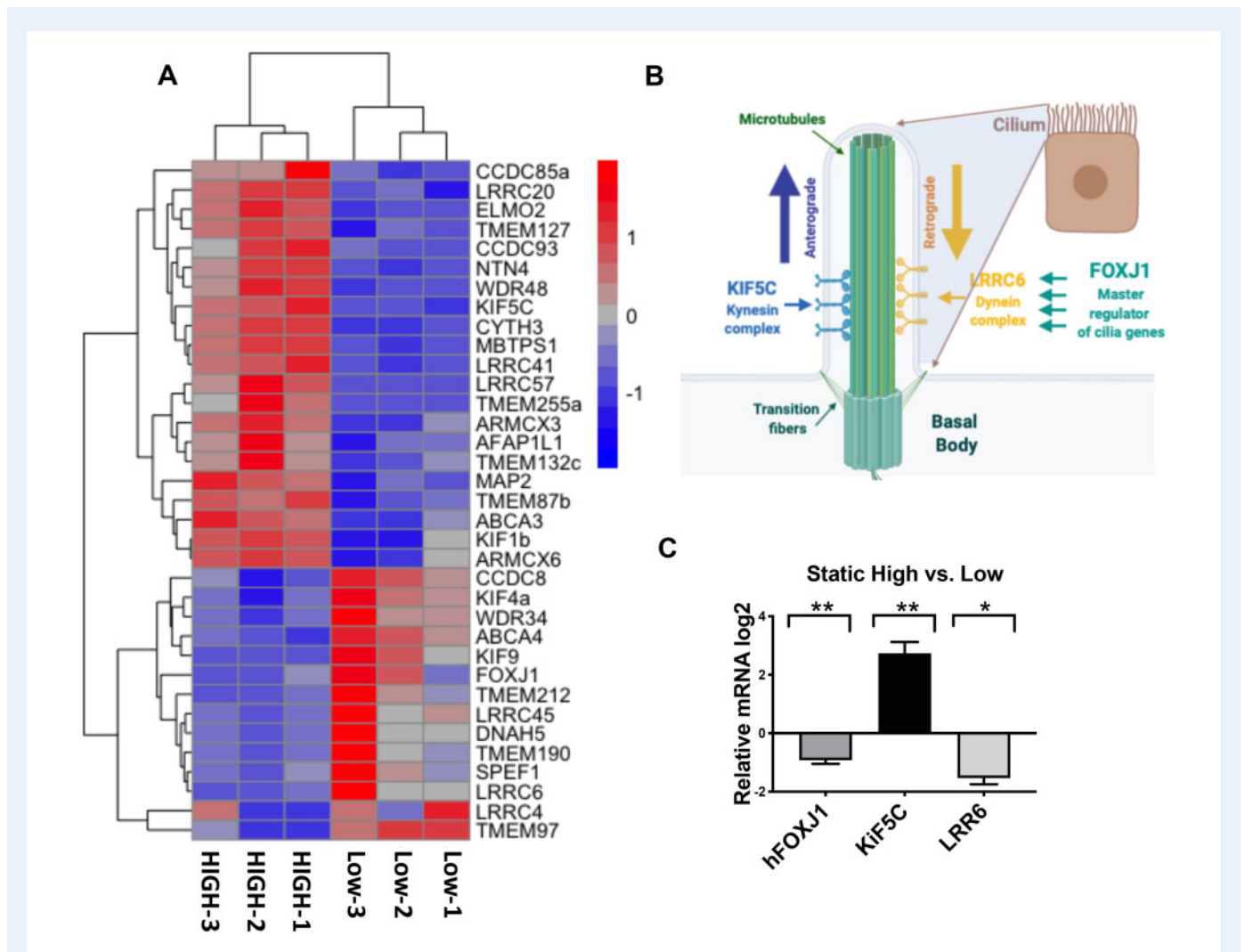


Figure 4. Exposure to PCOS-like (high) androgen concentrations alters expression of transcripts that encode for cilia in hFTE.

(A) Heatmap of cilia mRNA expression in PCOS-like (High) vs. control (Low). These data were generated using hFTE from three subjects treated with control or PCOS-like androgen levels in static culture for 14 days. Only significant genes with false discovery rate (FDR) adjusted P -value < 0.05 were included. (B) Schematic showing the role of *FOXJ1* as master regulator of cilia genes and of *LRRC6* and *KIF5C* regulation of intra-cilia transport that is critical for proper cilia function. (C) qPCR validation of *FOXJ1*, *KIF5C*, *LRRC6* mRNA expression from three independent patient samples. $*P \leq 0.05$; $**P \leq 0.01$.

although women with PCOS in both the high and low estrogen groups had a higher risk of ectopic than women without PCOS. This risk was ameliorated when frozen embryo transfers were performed, where serum hormone levels are more similar to natural conception cycles than COH. Testosterone levels were not included in their analyses. Women with PCOS tend to have higher estrogen levels during COH and are at higher risk of ovarian hyperstimulation syndrome (Luke *et al.*, 2010). The derangement of serum ovarian hormones in COH may also include elevated testosterone, which increases with estrogen levels prior to ovulation. Interestingly, in our study, higher androgen levels resulted in reduced estrogen receptor nuclear localization and OVGPI expression suggesting that despite higher steroid levels of estrogen, the fallopian tube epithelium of PCOS women may respond to hormones differently due to higher androgen exposure. Given the effect of testosterone on tubal cilia in our *ex vivo* models, we are left to

question the clinical reproductive significance and whether women with hyperandrogenism have an increased risk of ectopic pregnancy or other fallopian tube dysfunction. Unfortunately, no specimens included in this study were obtained from women with a clinical diagnosis of PCOS at the time of surgery. This could be viewed as a limitation but due to the heterogeneity of PCOS phenotypes and differences in diagnostic criteria, using PCOS tubal specimens may have added to the patient variability. Further women of reproductive age with PCOS are not frequent among surgical cases for bilateral salpingectomy for sterilization or hysterectomy for benign indications. Potential PCOS specimens may be captured by including specimens from surgery for ectopic pregnancy, hydrosalpinges or endometrial cancer, but these fallopian tubes may also confer a high degree of tubal damage. We sought to reduce selection bias by limiting our specimens to healthy tubal specimens. If the epithelium appeared damaged or scant in

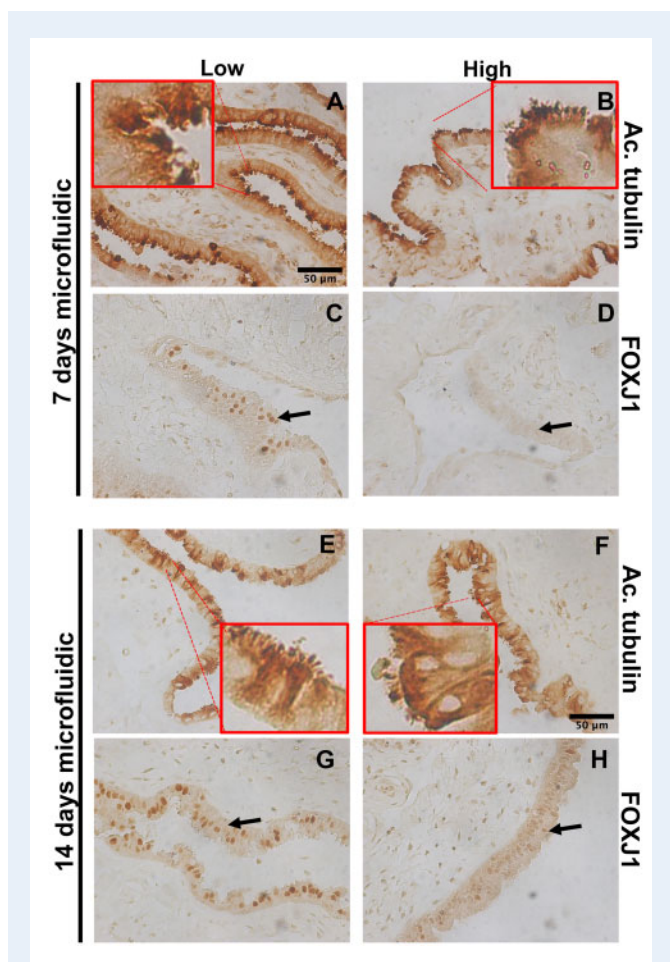


Figure 5. High androgen alters protein expression of FOXJ1, a regulator of cilia expression, in fluidic cultures.

(A and B) IHC of acetylated tubulin (Ac. Tubulin), which marks cilia, in low vs. high testosterone-treated hFTE after 7 days treatment and 14 days. (C and D) IHC of FOXJ1 shows reduced expression and nuclear localization in high testosterone-treated hFTE after 7 days treatment. (E and F) IHC of acetylated tubulin (Ac. Tubulin), in low vs. high testosterone-treated hFTE after 14 days of treatment. (G and H) IHC of FOXJ1 shows reduced expression and nuclear localization in high testosterone-treated hFTE after 14 days treatment. Scale bar is 50 μm and magnification is the same for all pictures. Red squares show a 4 \times magnification of the image to visualize receptor localization.

amount on initial dissection, that tissue was not used for experiment and was banked for potential future research. In addition, another interesting question that we are currently addressing is the role of anti-androgenic drugs (bicalutamide) and environmental endocrine disrupting factors (phthalates) on human fallopian tube health and cilia function. Furthermore, the uncovered role of testosterone on cilia assembly and function may suggest an interesting implication on the effect on androgen on lung function. Sex disparity has been demonstrated in lung diseases, but the role of testosterone has been shown to be controversial. In a mouse model, testosterone exacerbated pulmonary fibrosis (Voltz et al., 2008).

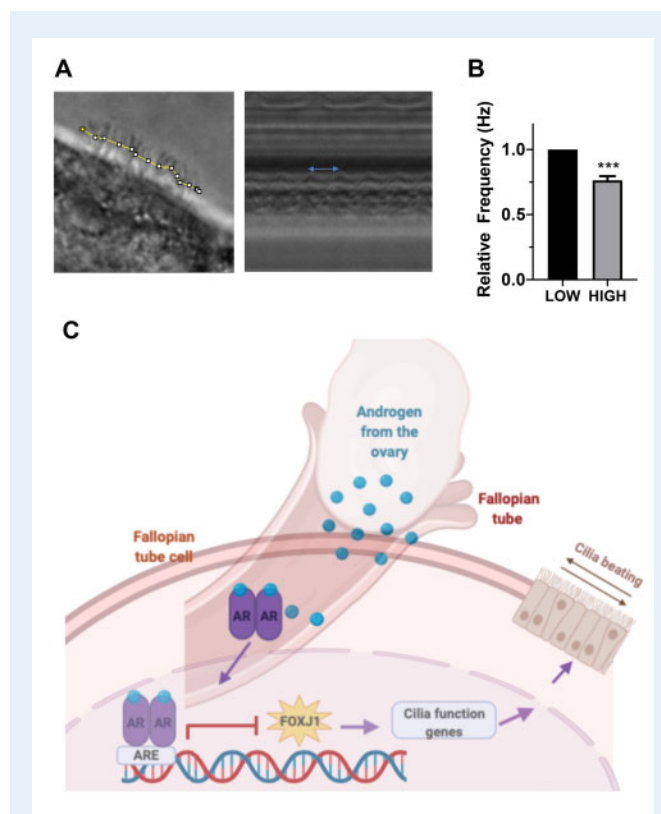


Figure 6. High androgen reduces cilia beating frequency.

(A and B) Tubal cilia were imaged using a spinning disc Nikon Ti2 microscope which showed that high androgen exposure significantly reduced cilia beating. In the left panel of 'A', the yellow line indicates a section of moving cilia selected for quantification. The blue arrow in the right hand panel shows the distance between two waveforms. Experiments shown in 'B' include four unique subjects. The graphs are represented as mean \pm SEM. (C) Schematic representing the androgen signaling pathway with ovarian over-production of androgen in PCOS patients and the stimulation of receptors on the hFTE. Activation of the receptor caused translocation to the nucleus to inhibit transcription of cilia genes, FOXJ1 (as well as LRR6 and KIF5C) that are critical for proper cilia assembly and movement.*** $P < 0.001$.

In summary, this study demonstrated that elevated testosterone exposure impaired tubal ciliary function and gene expression. This direct effect on ciliary function, hormone profile and gene expression were observed at a relatively low dose of testosterone and short duration of time. The *in vitro* models sustained tissue integrity and viability for 14 days of culture on both static and microfluidics systems, again adding to the literature the validity of using microfluidic systems to understand nuances of human reproduction and factors that contribute to impaired fertility. This study adds to our understanding of the mechanisms of subfertility and reproductive health risks in women living with hyperandrogenic disorders, including PCOS.

Supplementary data

Supplementary data are available at *Human Reproduction* online.

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Authors' roles

T.J.-B. contributed on experimental execution, paper figures and writing. J.C. contributed on paper figures. B.C.I. and J.C. provided the microfluidic devices. M.U., J.J.K. and W.K.T. are part of the Northwestern Reproductive Science Center and collaborated on the experimental design, subject recruitment and provision of human fallopian tube specimens. J.E.B. supervised the project. A.R. generated the project idea and contributed on experimental design, paper figures and writing.

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Conflict of interest

The authors declare no conflict of interest.

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