# 1 Elevation of anti-müllerian hormone (AMH) in PCOS women in ART:

- 2 Effect of Insulin
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#### 18 Abstract

- 19 Objective: To measure blood and follicular AMH levels in PCOS woman in ART and
- 20 examine direct action of insulin on AMH expression in human granulosa cells
- 21 **Design:** Prospective clinical and experimental study.
- 22 **Setting:** University Hospital-based laboratory.
- Patients: Women with (n=86) and without (n=172) PCOS in ART.
- 24 Intervention(s): Blood, follicular fluid and luteinized granulosa cells were collected from
- 25 PCOS or non-PCOS women in ART.
- 26 Main Outcome Measure(s): Hormone levels in blood and fluid and gene expression in
- 27 granulosa cells were measured.
- 28 Result(s): Serum levels of AMH were elevated and inversely correlated with embryo
- 29 cleavage rate in PCOS women in ART. Significant higher levels of AMH were also found in
- 30 small and large follicles collected from PCOS women as compared to those in non-PCOS
- 31 women. Luteinized granulosa cells from PCOS women showed higher expression of AMH
- 32 and its receptor AMHR2. Direct effect of insulin in increasing the expression of AMH in the
- isolated luteinized granulosa cells was observed with the PCOS granulosa cells responding to
- 34 a high dose of insulin. Co-treatment of AMH attenuated insulin-induced aromatase
- 35 expression in the luteinized granulosa cells.
- 36 Conclusion(s): These results suggest insulin may contribute to AMH elevation in PCOS,
- 37 while AMH counteracts with insulin-promoted aromatase expression in granulosa cells.
- 38 Key Words: Polycystic Ovarian Syndrome (PCOS), Assisted reproductive technology
- 39 (ART), Anti-Müllerian hormone (AMH), Insulin.

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43 AMH elevation and insulin in PCOS under ART

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# 45 Abstract capsule

- 46 AMH is elevated in blood and follicular fluid in PCOS in ART. Insulin increases AMH
- 47 expression, while AMH counteracts with insulin-promoted aromatase expression in human
- 48 luteinized granulosa cells granulosa cells.

### Introduction

Polycystic ovary syndrome (PCOS), characterized by ovarian follicular arrest and hormonal disturbance, is a leading cause for female infertility affecting 5-10% women of reproductive age (1, 2). A variety of hormones are dysregulated in PCOS including excessive androgens (3), hypersecreted luteinizing hormone (LH) (4), reduced follicle-stimulating hormone (FSH) (4) as well as disturbed metabolic hormones (e.g. insulin and glucagon) (5). In particular, insulin resistance accompanied with compensatory hyperinsulinemia is commonly found in PCOS women (up to 75% of lean and 95% of overweight, respectively) (6, 7).

Anti-Müllerian hormone (AMH), a key regulator for sex differentiation during embryonic development (8), is recently recognized to be elevated in PCOS contributing to the ovarian dysfunction (9, 10). AMH is a homodimer glycoprotein belonging to the transforming growth factor  $\beta$  (TGF- $\beta$ ) family (11, 12). In addition to its role in inhibiting embryonic development of Müllerian duct and thus leading sex differentiation into males (13), AMH is also known to be involved in folliculogenesis (14). In women, AMH is produced exclusively by granulosa cells (15), starting at the primary follicle stage with its highest expression detected in pre-antral and small antral follicles, which declines afterwards till absent in large (> 8-10 mm diameter) follicles (16). Studies have shown that AMH inhibits FSH-stimulated expression of aromatase, a key enzyme for ovarian steroidogenesis (17), as well as the growth of pre-antral follicles(18). Therefore, the expression pattern of AMH during folliculogenesis is believed to preserve most antral follicles from all being matured and ovulated, which is evident by the accelerated folliculogenesis leading to early depletion of ovarian follicles observed in AMH-null mice (19). Abnormally elevated AMH in PCOS,

however, is believed to be detrimental to FSH-stimulated growth of selectable follicles leading to anovulation (20).

While its contribution to PCOS pathogenesis is generally accepted, the cause of AMH elevation in PCOS remains unknown. It was thought that the increased number of small antral follicles in PCOS might produce an excessive AMH environment, although later studies have shown that AMH production is increased in individual follicles or granulosa cells from PCOS women (21-23) with the underlying mechanism largely unexplored. Interestingly, hyperinsulinemia is proposed a reason for AMH elevation in PCOS, as AMH levels were found to be particularly higher in PCOS women with insulin resistance (24, 25) and long-term metformin treatment was reported to correct AMH levels in PCOS (26). However, contradictory findings were also reported (27, 28) and the direct effect of insulin on AMH production by granulosa cells is not clearly demonstrated.

PCOS women often seek in vitro fertilization (IVF)-based assisted reproductive technology treatment (ART). In ART, the AMH level has been associated with ovarian responses (29), implantation, pregnancy and other qualitative ART outcomes (30, 31), although controversial results are also reported (32, 33)(34, 35).

We undertook the present study to better characterize the association between AMH level with embryo quality as well as ART outcomes in PCOS. We also examined possible action of insulin on AMH expression in human granulosa cells, while comparing the cells from PCOS and non-PCOS women.

### **Materials and Methods**

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### Human subjects and clinical data collection

Total 258 women undergoing in vitro fertilization (IVF)-based assisted reproductive treatment (ART) at Shanghai First Maternity and Infant Hospital from June 2014 to June 2017 were recruited for the present study. The Rotterdam criteria (36): I, oligomenorrhea (cycle length > 35 days) or amenorrhea (cycle length > three months or no rise of progesterone before menstruation); II, hyperandrogenism as defined by hirsutism, and/or serum testosterone level > 0.7 ng/ml and/or androstenedione > 2.2 ng/ml; and III, the presence of more than 12 follicles of 2-9 mm in diameter in each ovary and/or unilateral ovarian volume >10ml under B-ultrasound, was used for diagnosis of PCOS. 86 women showed the presence of Rotterdam criteria- I and III and were diagnosed of PCOS. Androstenedione was not measured in them. 172 women who met the following inclusion criteria: i) age between 20 and 40, ii) both ovaries were present with no morphological abnormalities and adequately visualized in transvaginal ultrasound scans, iii) regular menstrual cycle, iv) no current or past diseases affecting ovaries or gonadotrophin or using sex-steroids for at least 3 months prior to inclusion, were grouped as non-PCOS for the present study. The cause of infertility in non-PCOS group was due to either tubal obstruct or male factor. The study was approved by the Ethics Committee of Tongji University and all the subjects gave informed consent. All the procedures were performed in accordance with the relevant guidelines and regulations. The subjects' clinical data including age and BMI were documented. Patients with oligo- or amenorrhea were pre-treated with progesterone to induce regular cycle before the blood test and ART treatment. A blood test was performed on the 3<sup>rd</sup>-5<sup>th</sup> day of the menstrual cycle before the treatment to determine basal levels of hormones.

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#### Ovarian hyperstimulation

All the subjects were given diphereline, (Ipsen Pharma Biotech, France, 1.25 mg) a gonadotropin-releasing hormone agonist, at their mid-luteal phase of the menstrual cycle, to suppress pituitary secretion of gonadotropin hormones and prevent premature ovulation. After pituitary suppression was achieved, which was determined by  $E_2 \leq 50 \text{pg/ml}$ , endometrial thickness ≤6mm and no follicles >10 mm under transvaginal ultrasound, subjects were daily injected with Gonal-F, a recombinant human FSH (rhFSH, Merck-Serono, Switzerland), starting from day 5 of a normal menstrual cycle (initial day). Serum sex hormone levels were monitored and transvaginal ultrasound (5-MHz, Model SSD-620, Aloka Co. Ltd, Tokyo Japan) was performed to evaluate the development of ovarian follicles and adjust the dose of rhFSH. When the lead follicle achieved 18 mm in diameter, the lead two were 17mm or the lead three were 16mm, patients were subcutaneously injected with recombinant human chorionic gonadotropin (rhCG, 250µg, Ovidrel, MERCK SERONO, Switzerland) to trigger oocyte maturation (hCG day). 36 hours afterwards, oocytes were retrieved by transvaginal ultrasound-guided approach. Fertilization of the retrieved oocytes was done either by IVF or intracytoplasmic sperm injection (ICSI), according to the sperm quality. Successful fertilization and cleavage were determined by reported ART standards (37). Quality of embryo were evaluated and graded into I to V: grade I – embryos with equal blastomeres and less than 5% cytoplasm fragmentation, grade II - embryos with equal blastomeres and 5-25% cytoplasm fragmentation, grade III - embryos with unequal blastomeres and less than 5% cytoplasm fragmentation, grade IV -embryos with unequal blastomeres and 5-25% cytoplasm fragmentation, grade V – embryos with a few blastomeres and with a strong or complete cytoplasm fragmentation. Embryos of grade I and II were considered as good, grade III as intermediate, grade IV and V as poor quality embryos. 3-5 days after cleavage was observed, up to 2 embryos were transferred in each patient per cycle. Grade I and II embryos were transferred as priority. Grade III embryos were transferred in case no grade I-II embryo was available. In rare cases, grade IV-V embryos were cultured till blastocyst-stage before the transfer. Patients were intramuscularly injected with progesterone (60 mg/day, Tongyong Pharmaceutical Co., Shanghai, China) from oocyte retrieval day till 14 days after embryo transfer. Clinical pregnancy was defined as the identification of a gestational sac with fetal heart activity on ultrasound examination 4 to 5 weeks after embryo transfer. Implantation rate was calculated as the number of gestational sacs (observed at week 4 post-transfer) divided by the number of embryos transferred in each patient. Other downstream parameters - pregnancy, miscarriage, ectopic and live birth rates were calculated with the total number of patients in each group as the denominator.

#### Follicular fluid and granulosa cell collection

The ovarian follicles were collected after hCG injection from the patients as we previously reported (38). Follicles were classified into two groups, small (<10mm diameter) and large (>18mm), according to their size measured by ultrasonography. Under the guidence of ultrasonography, a sterile needle, connected to a vaculum device, was transvaginally inserted. With gental suction, each follicle was individually collected through the needle to a collection vial. Large follicles were collected before the collection system was washed sufficiently and small follicles were collected afterward. For each individually collected follicle, the oocyte was carefully dissected out under a dissecting microscope, the remaining granulosa cells with fluid were transferred into sterile tubes (Axygen Scientific, Union City, CA, USA) for centrifugation at 400g for 10 min. Afterwards, the supernatant was used as

follicular fluid. Cell pellets were washed with PBS, subsequently a red blood cell lysis buffer and further centrifuged at 400 g for 5 min. For ELISA measurements of the follicular fluid, a single large follicle with sufficient amount of fluid was used for one measurement, whereas follicular fluid gathered from 3-4 small follicles from a patient was used for one measurement. The collected granulosa cells and follicular fluid were stored at -80 °C until mRNA analysis or re-suspended for culture.

### Hormonal immunoassays

Blood was collected from subjects on the 3<sup>rd</sup>-5<sup>th</sup> day of the menstrual cycle prior to the treatment (Basal), the day when FSH treatment started (Initial day) and the day when hCG was given (hCG day) to determine testosterone, FSH, LH or E<sub>2</sub> levels by an automated chemiluminescence immunoassay analyzer (Siemens, Tarrytown,NY, USA). On hCG injection day and before the injection, blood was collected to examine serum AMH levels by an ELISA kit, Anti-Mullerian Hormone ELISA (Biomatik), according to the supplier's instructions. For follicular fluid samples which are of small volume and have lower AMH levels, we used a more sensitive and specific chemiluminescence kit (Roche, Basel, Switzerland) according to manufacturer's instruction. Insulin in follicular fluid was measured using a specific ELISA-kit (Mai Bio, Shanghai, China) according to manufacturer's instruction. The plates were read by an ELISA reader (Thermo Fisher Scientific, Inc.).

# Human luteinized granulosa cell culture

The collected granulosa cells were cultured in DMEM/F12 (Gibco, Rockville, MD, USA) with 10% fetal bovine serum (Gibco), 100 U/ml penicillin (Life Technologies), 100 ug/ml streptomycin sulfate (Life Technologies), and 1×GlutaMAX (Life Technologies) at 37

°C in CO<sub>2</sub> incubator. Cells were deprived of serum for 24 hours before stimulated with 5, 10, 100ng/ml of insulin (Sigma; bovine pancreatic insulin) or GSK1904529A (5μM, Selleck), an insulin receptor (INSR) inhibitor.

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#### RNA extraction, reverse transcription and quantitative real-time PCR

Total RNAs from cells were extracted using the Trizol reagent (Life Technologies, Inc.) following the manufacturer's protocol. The concentration of RNA was measured using the NanoDrop 2000cUV-Vis spectrophotometer (Thermo Fisher Scientific, Inc.).1 µg RNA was transcribed to cDNA using the PrimeScript RT reagent Kit (Takara, Shiga, Japan). Briefly, a master mix containing 4 µL 5×PrimeScript Buffer, 1 µL Rime Script RT Enzyme Mix I, 1 µL Oligo dT Primer (50µM), 1 µL Random 6 mers (100µM) and RNase Free water was prepared for each 20 µL reaction, which followed by 37 °C for 30 min, at 85 °C for 5 s and cooling at 4 °C. Gene expression levels were evaluated by the manufacturer's protocol on the StepOnePlus™ Real-Time PCR System (Thermo Fisher Scientific, Inc.) using the SYBR reagent (Tiangen, China). The primers were from Sangon Biotech (Shanghai, China) and the sequences are listed in Table 1. The general thermal profile of the reaction was for all the genes tested 10s at 95°C to activate the enzyme followed by 40 cycles of 95°C for 5s and 60°C for 34s for each cycle. The dissociation stage was initiated at 95°C for 15s, followed by one cycle at 60°C for 60s and 95°C for 15s. Each reaction was assayed in triplicate and three separate experiments were performed on different cultures. A mean value (generated from the triplicate data of the target genes and reference gene of each sample) was used for the determination of mRNA levels by the comparative Ct (2- $\Delta\Delta$ Ct) method with  $\beta$ -actin as the housekeeping gene,  $\Delta\Delta Ct = [Ct \text{ of target (sample A)} - Ct \text{ of reference (sample A)}] - [Ct \text{ of }]$ target (sample B) – Ct of reference (sample B)]. This equation can be used to compare the

gene expression in two different samples (sample A and sample B); each sample is related to a housekeeping gene ( $\beta$ -actin).

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#### Western blot

Cells were washed with PBS and lysed in RIPA buffer (Beyotime, Shanghai, China) containing protease inhibitor cocktail and PMSF (Sigma-Aldrich) on ice. The extract was centrifuged at 15000g for 30 min at 4°C to remove cellular debris and the protein concentration was quantified using the BCA assay (Thermo Fisher Scientific, Inc.). The same amount of protein (20µg) of each sample was separated by SDS-PAGE (Beyotime, Shanghai, China) with 5% stacking gel and 10% separating gel at 120V for 2 hours and then transferred onto the polyvinyl difluoride membranes (PVDF) at 250 mA for 1 hour. Membrane nonspecific binding was blocked with 5% nonfat milk in TBST (0.01M Tris-HCl, 0.15M NaCl, 0.1%Tween 20, PH7.4, Beyotime), containing 0.05% Tween-20 for 1 hour at room temperature and then incubated with rabbit antihuman aromatase (Cell Signal Technology, USA, 1:1000) and rabbit antihuman GAPDH (Abmart, Shanghai, China, 1:1000) diluted in TBST with 5% milk overnight at 4°C. Next, the membranes were washed three times with TBST and then incubated for 1 h with the horseradish peroxidase –conjugated secondary antibody (Abmart, 1:2000). Immunoreactive bands were detected using an enhanced chemiluminescent substrate (Millipore, Watford, UK). The intensities of the bands were quantified using Image- J software.

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### Statistical analysis

Data were mean  $\pm$  SEM of at least three independent experiments. Comparisons of two

independent groups were using the Student's t-test. One-way ANOVA followed by Tukey's post hoc tests was used to compare more than two groups. The correlation analysis was done by the Spearman's coefficient. *P*<0.05 was considered statistically significant. All statistic procedures were run on SPSS 20.0 (IBM, Armonk, NY, USA). Since there were differences in age and BMI as well as in the duration of infertility between PCOS and non-PCOS in Cohorts I and III, we corrected a priori for these differences by analysis of covariance and multiple liner regression.

### **Results**

#### Clinical characteristics of PCOS and non PCOS women in ART

Blood, follicular fluid and granulosa cells were collected from 3 cohorts of woman subjects respectively. Clinical characteristics are summarized in Supplementary Table 2. In general, the PCOS women, when examined, were about 2 years younger with a higher BMI in average, as compared to the non-PCOS women. The PCOS women in average were given a lower initial dose of rhFSH with longer treatment duration than the non-PCOS, although the total dose of rhFSH were comparable with no significant difference between the two groups.

Hormonal profiles of the two groups are shown in Supplementary Table 2. Prior to the treatment (basal), the PCOS group exhibited lower FSH and higher LH levels as compared to the non-PCOS group. The serum FSH, LH and E<sub>2</sub> in both PCOS and non-PCOS subjects were largely decreased to a low level before rhFSH treatment started (Initial day), indicating successful pituitary suppression. After rhFSH treatment and right before hCG injection (hCG day), the LH level remained low while E<sub>2</sub> level was highly elevated in both groups with no significant difference. Progesterone levels on hCG day were also found to be comparable

between PCOS and non-PCOS women.

Clinical outcomes are shown in Supplementary Table 2. More PCOS patients were considered to have risk of ovarian hyperstimulation syndrome (OHSS), and therefore chose to cancel embryo transfer within the same cycle and freeze the formed embryos for later treatment. Among the assessed parameters including fertilization, cleavage, embryo quality, implantation and pregnancy, PCOS and non-PCOS women exhibited similar ART outcomes.

## AMH is elevated and negatively correlated with oocyte/embryo quality in PCOS women

We collected blood samples on hCG-injection day from 31 PCOS and 59 non-PCOS women (Cohort I, Supplementary Table 2) to determine AMH blood levels. Results showed that the averaged blood level of AMH in PCOS women (39.77  $\pm$  1.84) was significantly higher than that in non-PCOS women (28.26  $\pm$  1.00) (Fig.1A). To explore possible effect of AMH on embryo quality, correlation analysis was performed between the AMH blood level and the number of oocytes retrieved (Fig.1B), fertilization rate (Fig.1C), cleavage rate (Fig.1D), number of good quality embryo (Fig.1E) and implantation rate (Fig.1F) in PCOS and non-PCOS subjects, respectively. Results showed significantly negative correlation of the AMH blood levels with the cleavage rate (Fig.1D) and number of good quality embryos (Fig.1E) in PCOS women. In non-PCOS women, only a weak inverse correlation of AMH with number of good quality embryos was found.

### AMH is elevated in follicular fluid and granulosa cells in PCOS women

Since the follicular environment is particularly important to oocyte development and maturation, we further examined the AMH level in the follicular fluid in another cohort of

PCOS (n=19) and non-PCOS women (n=36) (Cohort II, Supplementary Table 2). Both small (<10mm) and large (>18mm) follicles were achieved under ultrasonography during oocyte pick-up and the follicular fluid was collected. In both PCOS and non-PCOS groups, the small follicles contain higher concentrations of AMH as compared to the large follicles, consistent with a role of AMH during folliculogenesis (Fig.2A). Importantly, PCOS follicles exhibited higher AMH levels (large: 6.15 ± 0.98 ng/ml, small: 16.51 ± 1.48 ng/ml), as compared to non-PCOS group (large: 2.29 ± 0.32 ng/ml, small: 6.29 ± 1.00 ng/ml) in large and small follicles respectively (Fig.2A). Since AMH is known to be exclusively produced by granulosa cells in women (8), we tested whether the elevated AMH in follicular fluid would be due to upregulation of AMH expression in PCOS granulosa cells. Indeed, in collected luteinized granulosa cells from large follicles, significantly higher mRNA level of AMH was observed in PCOS women as compared to non-PCOS group (Fig.2B). Moreover, the AMH receptor, AMHR2, was also found to be increased at the mRNA level in PCOS granulosa cells, as compared to non-PCOS (Fig.2C).

#### Insulin induces upregulation of AMH in PCOS granulosa cells

We next explored possible reason for AMH upregulation in granulosa cells in PCOS. Hyperinsulinemia is proposed to be responsible for AMH elevation in PCOS (25). We also detected the presence of insulin in both PCOS and non-PCOS follicular fluid, with a relatively higher level detected in the PCOS group (Fig.3A). We next examined the direct effect of insulin on AMH expression in culture human granulosa cells. Luteinized granulosa cells from 24 non-PCOS and 6 PCOS women (Cohort III, Supplementary Table 2) were isolated and cultured as we previously reported (38). The expression of insulin receptor showed no significant difference between PCOS and non PCOS granulosa cells (Fig. 3B).

After 24-hour insulin treatment, the mRNA levels of AMH in granulosa cells were significantly increased by insulin in a dose-dependent manner in both non-PCOS and PCOS (Fig.3C) groups. At 100ng/ml, insulin no longer increased AMH expression in non-PCOS cells, suggesting desensitization to a high dose of insulin in these cells (Fig. 3C). Whereas, PCOS cells continued to respond to insulin at 100 ng/ml resulting in further increases in AMH mRNA levels as compared to those at 10 ng/ml, suggesting their loss of desensitization to high dose insulin (Fig.3C). In addition, pretreatment with an antagonist of insulin receptor, GSK1904529A (5μM), for 8 hours, blocked the insulin-induced AMH upregulation in non-PCOS granulosa cells (Fig.3E). No significant changes in AMHR2 mRNA were observed after the insulin treatment in either non-PCOS or PCOS cells (Fig.3D).

#### AMH inhibits insulin-stimulated aromatase expression in granulosa cells

Granulosa cells function is essential to oocyte development and maturation. Given the presence of both insulin and AMH in follicular fluid and the direct effect of insulin in increasing AMH expression as suggested above, we wondered whether the expression of genes essential to in granulosa cell function would be subject to regulation by either the follicular insulin or AMH. Aromatase (CYP19), CYP17, FSH receptor (FSHR), androgen receptor (AR), vascular endothelial growth factor receptor (FLK), insulin-like growth factor 1 receptor (IGF1R) and LH receptor (LHR) were examined in luteinized granulosa cells from non-PCOS women after treatment with insulin (10 ng/ml), AMH (20 ng/ml) or their combination for 24 hours. Results showed that insulin (10 ng/ml) alone significantly increased mRNA levels of CYP19 (Fig.4A), FSHR (Fig.4B), AR (Fig.4C) and CYP17 (Fig.4D), as compared to controls cells. AMH (20 ng/ml) alone did not alter any of these genes (Fig.4). Co-treatment of AMH, however, substantially attenuated the effect of insulin

(10ng/ml) in increasing CYP19 mRNA expression (Fig.4A). Slight effect of AMH co-treatment in inhibiting insulin-induced FSHR expression was also observed (Fig.4B). No obvious changes in FLK, IGF1R and LHR by either insulin or AMH were observed in these cells (Fig.4E-G). To confirm the inhibitory effect of AMH on insulin-induced CYP19, we also examined the protein level of CYP19, which consistently showed that as compared to cells treated with insulin (10ng/ml) alone, cells treated with combination of AMH (20 ng/ml) and insulin (10ng/ml) exhibited significantly lower CYP19 protein levels (Fig.4H). Antagonizing insulin receptor by GSK1904529A (5μM Fig. 4I) also inhibited insulin-induced upregulation of CYP19 in these cells.

### **Discussion**

Collectively, the present study has indicated a higher blood AMH level in PCOS than non-PCOS women during ART treatment. The blood AMH levels are in negative correlations with the cleavage rate and number of good quality embryos in PCOS women. AMH is also elevated in collected follicular fluid and upregulated in isolated luteinized granulosa cells from PCOS women. Direct effect of insulin in increasing the expression of AMH in the isolated granulosa cells is observed with the PCOS cells responding to a high dose of insulin. The presence of AMH exerts an inhibitory effect on insulin-induced aromatase expression in isolated luteinized granulosa cells.

Consistent with others (10, 39), the present study demonstrates that AMH is abnormally upregulated in PCOS. First, protein levels of AMH are increased in the blood and follicular fluid of either small (presumably pre-antral/antral) or large (matured) follicles. Second, we were only able to collect analysable granulosa cells from large but not small follicles, which are usually believed luteinized cells and supposed to have reduced AMH expression.

Nevertheless, in these cells, significantly increased mRNA expression of both AMH and its receptor AMHR-II is observed in PCOS compared to non-PCOS. It should be also noted that the blood AMH level was examined on hCG day, when pituitary suppression and exogenous FSH was already applied and other hormones became comparable between PCOS and non-PCOS. AMH is elevated in PCOS even after hormonal correction in ART, which in together suggests intrinsic changes in PCOS granulosa cells resulting in abnormally upregulated AMH and its downstream signaling.

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It has been controversial whether AMH can affect outcomes of ART (40-44). The present study has found negative correlations of AMH blood level with the cleavage rate and number of good quality embryos in PCOS women. Therefore, a negative impact on embryo quality, particularly at early stages of embryo development, by the elevated AMH particularly in PCOS is suggested. However, since only cleaved or good quality embryos are selected to proceed with embryo transfer and implantation in ART, the later stages and ultimate ART outcomes seem not influenced by the elevated AMH in PCOS. Interestingly, a most recent study has shown that prenatal AMH exposure may account for the pathogenesis of PCOS in adulthood (45), suggesting the influence of AMH on embryo development in PCOS too. Of note, it was previously reported in a general population that no relationship between AMH and embryo quality was found (46). Others showed that low maternal AMH level could be a predicting marker for fetal aneuploidy (47), suggesting that a low AMH level is not good to embryos either. In the present study, although a high AHM level correlates with poor embryo quality in PCOS women, the maternal AMH levels in non-PCOS women have no or weak correlation with embryo quality parameters. It seems consistent that AMH level can hardly predict embryo quality within a general population. Nevertheless, detailed mechanisms underlying the effect of AMH on embryo quality in PCOS await further investigation.

An association between the environmental insulin and granulosa cell production of AMH is suggested. A direct effect of exogenous insulin, mediated by insulin receptor, in increasing AMH expression is observed in both PCOS and non-PCOS luteinized granulosa cells. Interestingly, luteinized granulosa cells isolated from PCOS women seem to be quite sensitive to insulin, rather insulin-resistant, in terms of AMH expression (Fig.3C). PCOS cells respond to insulin even at a high dose that already desensitizes non-PCOS cells (Fig.3C). Given that the expression level of the insulin receptor in PCOS cells showed no difference with that of non-PCOS cells (Fig.3B), the difference observed might due to functional change of the insulin receptor or alterations in its regulatory pathways, which is possibly a cause for the observed elevation of AMH in PCOS. However, the detailed mechanisms underlying the discrepancy in responses to insulin between PCOS and non-PCOS granulosa cells await further investigation. Of note, it was suggested that the overexpression of AMH and AMHR-II in PCOS women could be due to increased LH levels, especially in PCOS women with oligo/anovulation (48). Since insulin and LH are believed to have synergy effects (49), hyperinsulinemia may enhance the effect LH in increasing AMH expression in PCOS women.

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It should be noted that insulin is known for long to increase aromatase activity in granulosa cells from either normal and PCOS women (50, 51). Similar effect of insulin in increasing aromatase expression is also observed presently. Whereas, aromatase is known to be dysfunctional or downregulated (38) contributing to a hyperandrogenic environment in PCOS. In addition to previously reported effect of AMH in reducing FSH-induced aromatase expression (52), the present study has shown the capacity of AMH in counteracting with insulin's action on aromatase expression. Together with the finding of insulin's role in

402	promoting AMH production, it may provide an explanation to the reduced aromatase activity
403	despite a high insulin environment in PCOS.
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416	Competing financial interests
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418	
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### 571 Figure legends

### Fig.1. Blood AMH is elevated in PCOS women

- 573 (A) ELISA measurement of AMH levels in the blood collected on hCG injection day (before
- the injection) from PCOS and non-PCOS women. n is shown in each column. \*\*\*P < 0.001,
- 575 t-test. B-F) Correlation analysis of the blood AMH level and oocytes retrieved (B),
- 576 fertilization(C), cleavage (D), good quality embryo(E) and implantation (F) in PCOS and
- non-PCOS women. \*P < 0.05, \*\*P < 0.01, Spearman's coefficient correlation test.

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### Fig.2. AMH is elevated in follicular fluid and granulosa cells in PCOS women

- 580 (A) ELISA measurement of AMH levels in follicular fluid from non-PCOS and PCOS
- women. Data are mean  $\pm$  SEM, n is indicated in each column. \*\*\* P<0.001, t-test. **B-C**)
- Quantitative PCR (qPCR) analysis of AMH(B) and AMHR-II(C) mRNA levels in luteinized
- 583 granulosa cells freshly isolated from non-PCOS and PCOS women. β-actin was used as the
- internal control for relative mRNA level measurement by qPCR. Data are mean  $\pm$  SEM. n is
- 585 indicated in each column, \**P*<0.05, \*\**P*<0.01, t-test.

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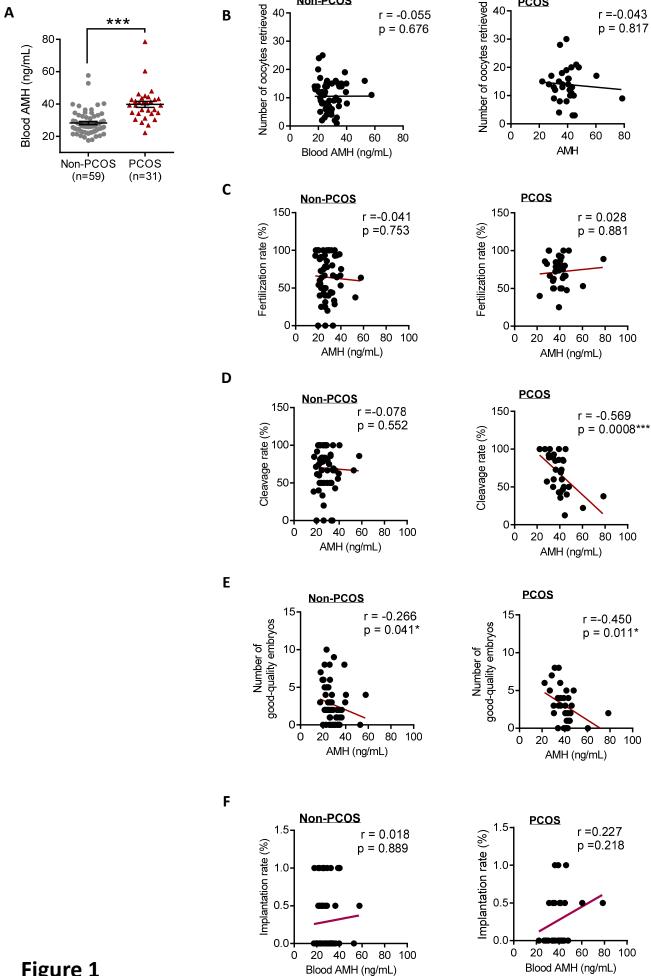
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#### Fig.3. Effect of insulin on AMH expression in human luteinized granulosa cells.

- 588 (A) ELISA measurement of insulin levels in follicular fluid from non-PCOS and PCOS
- women. Data are mean  $\pm$  SEM, n is indicated in each column. t-test. (B) qPCR analysis of
- 590 mRNA levels of insulin receptor (INSR) in cultured luteinized granulosa cells isolated from
- 591 non-PCOS and PCOS women. (C-D) qPCR analysis of mRNA levels of AMH (C) and
- 592 AMHR2 (D) in cultured luteinized granulosa cells isolated from non-PCOS and PCOS
- women, after incubation with insulin (0-100 ng/ml) for 24 hours. Cells isolated from 14

non-PCOS women and 6 PCOS women were pooled together for the experiments. n =3 (number of independent experiments). \*P<0.05, \*\*\*P<0.001. One-way ANOVA with post-tests. **(E)** qPCR analysis of mRNA levels of AMH in non-PCOS granulosa cells pretreated with or without GSK1904529A (5 $\mu$ M, an insulin receptor antagonist) for 8 hours before incubated with or without insulin (10 ng/ml) for 24 hours. Cells isolated from 10 women were pooled together for the experiments. Data are mean  $\pm$  SEM, n = 3 (number of independent experiments). \*P<0.05 one-way ANOVA.

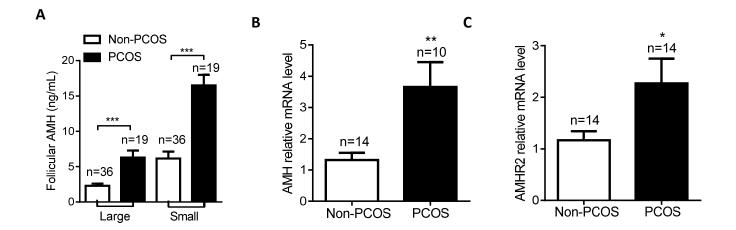
Fig.4. Effect of insulin and AMH on genes expression in human luteinized granulosa cells. (A-I) qPCR analysis of mRNA levels of CYP19 (A), FSHR (B), AR (C), CYP17 (D), FLK (E), IGF1R (F) and LHR (G) in luteinized granulosa cells from non-PCOS women, treated with or without insulin (10 ng/ml) and AMH (20 ng/ml) for 24 hours. β-actin was used as the internal control for relative mRNA level measurement by qPCR. Cells isolated from 9 women were pooled together for the experiments. Data are mean  $\pm$  SEM, n = 3 (number of independent experiments). \*P<0.05 and \*\*\*P<0.001, one-way ANOVA. (H) Western blotting for CYP19 in luteinized granulosa cells isolated from non-PCOS women, treated with or without insulin (10ng/ml) or AMH (20ng/ml). β-actin was used as a loading control for western blot. (I) Western blotting for CYP19 in luteinized granulosa cells isolated from non-PCOS women, treated with or without GSK1904529A (5μM, an insulin receptor antagonist) for 8 hours before incubated with or without insulin (10 ng/ml) for 24 hours. Cells isolated from 10 women were pooled together for the experiments. Data are mean  $\pm$  SEM, n = 3 (number of independent experiments). \*P<0.05, one-way ANOVA.



Non-PCOS

**PCOS** 

Figure 1



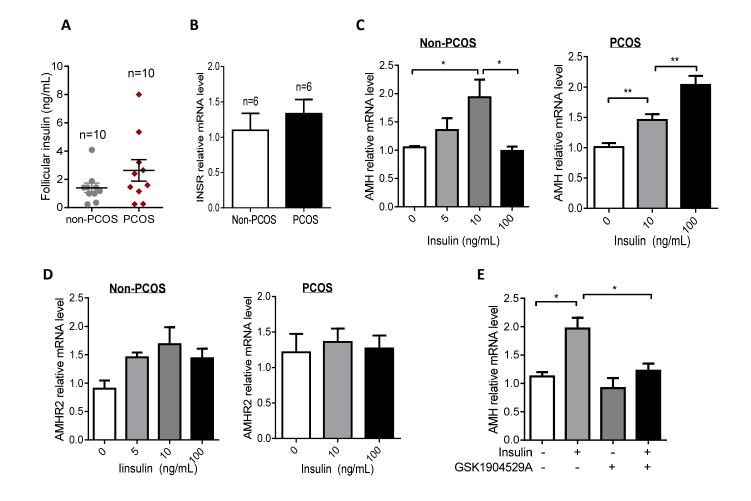


Figure 3

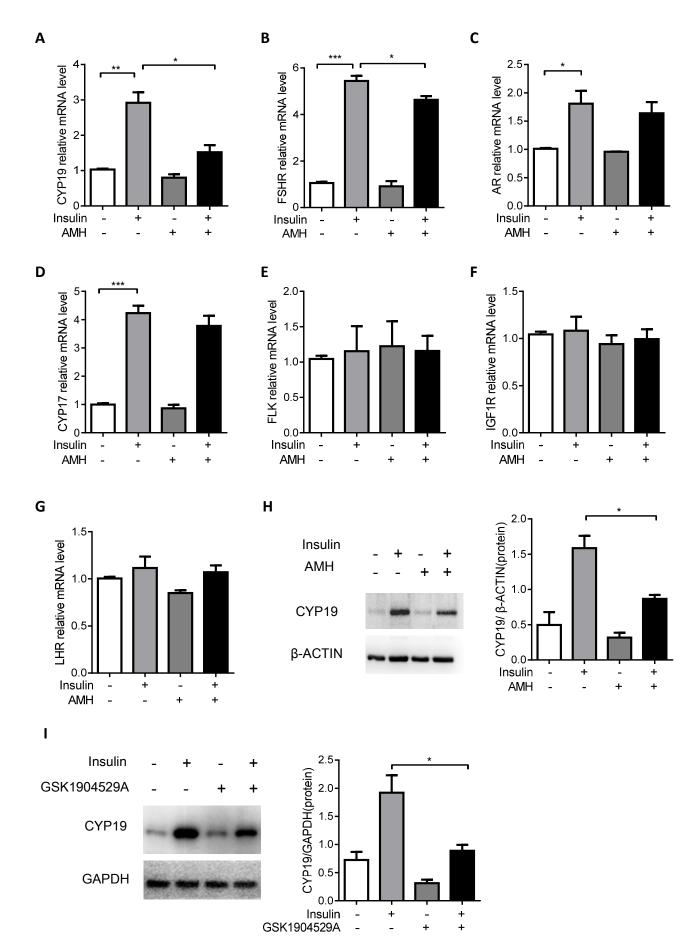


Figure 4

**Table.1 Sequences of primers** 

Gene	Forward sequence	Reverse sequence	Product size (bp)
AMH	TGGCCTCATCTTCCGAGAAG	GGAGCTGCTGCCATTGCT	100
AMHR-II	TTGGGGCTTTGGGCATTACTT	AGCTCTCCCAGTGTCTTTGTG	113
CYP19	TGGAAATGCTGAACCCGATAC	AATTCCCATGCAGTAGCCAGG	161
CYP17	GCTGCTTACCCTAGCTTATTTGT	ACCGAATAGATGGGGCCATATTT	174
FSHR	TCTGTCACTGCTCTAACAGGG	TGCACCTTTTTGGATGACTCG	131
AR	CCTGGCTTCCGCAACTTACAC	GGACTTGTGCATGCGGTACTCA	168
LHCGR	ATTTGTCAATCTCCTGGAGGC	CACTCAGTTCACTCTCAGCA	191
FLK	GGCCCAATAATCAGAGTGGCA	TGTCATTTCCGATCACTTTTGGA	105
INSR	AAAACGAGGCCCGAAGATTTC	GAGCCCATAGACCCGGAAG	90
β-actin	GCATGGGTCAGAAGGATTCCT	TCGTCCCAGTTGGTGACGAT	106

	Cohort I (Blood)			Cohort II (Follicular fluid )		Cohort III (Granulosa cells)			Corhort I + II + III			
Variables	Non-PCOS (n=59)	PCOS (n=31)	P	Non-PCOS (n=36)	PCOS (n=19)	P	Non-PCOS (n=77)	PCOS (n=36)	P	Non-PCOS (n=172)	PCOS (n=86)	P
Basic characteristics	•			, ,			<u> </u>			,		
Age (years)	$31.9 \pm 0.6$	29.4 ± 0.7	**	31.8 ± 0.6	$30.1 \pm 0.9$		31.3 ± 0.5	29.5 ± 0.6	*	31.6 ± 0.3	29.6 ± 0.4	***
BMI (kg/m²)	$20.9 \pm 0.4$	22.4 ± 0.5		22.07 ± 0.5	$23.4 \pm 0.7$		20.5 ± 0.5	$23.5 \pm 0.8$	**	21.2 ± 0.2	22.6 ± 0.4	***
Infertility causes												
Primary infertility	28 (47.5%)	17 (54.8%)		26 (72.2%)	9 (47.4%)		41 (53.2%)	22 (61.1%)		95 (55.2%)	48 (55.8%)	
Secondary infertility	31 (52.5%)	14 (45.2%)		10 (27.8%)	10 (52.6%)		36 (46.8%)	14 (38.9%)		77 (44.8%)	38 (44.2%)	
Initial dose of rhFSH (IU)	173.3 ± 5.3	162.4 ± 7.4		170.8 ± 8.1	153.3 ± 6.5		178.2 ± 5.2	153.5 ± 8.3	*	170.4 ± 4.0	154.2 ± 5.9	*
Total dose of rhFSH (IU)	2065.1 ± 150.1	2062.8 ± 205.2		2208.3 ± 116.4	2154.6 ± 182.3		2139.3 ± 98.1	2038.1 ± 157.1		2076.7 ± 76.85	1998.6 ± 113.5	
Duration of rhFSH (day)	10.6 ± 0.6	11.9 ± 0.9		11.7 ± 0.4	$12.6 \pm 0.8$		11.1 ± 0.3	12.6 ± 0.5	**	10.9 ± 0.3	11.6 ± 0.5	
Hormonal test												
Basal												
Testosterone	0.4 ± 0.1	0.6 ± 0.1	**	0.5 ± 0.1	0.6 ± 0.01	*	0.4 ± 0.0	0.7 ± 0.0	***	0.4 ± 0.0	0.6 ± 0.0	***
(ng/ml)	0.4 ± 0.1	0.0 ± 0.1		0.5 ± 0.1	0.0 ± 0.01		0.4 ± 0.0	0.7 ± 0.0		0.4 ± 0.0	0.0 ± 0.0	
FSH (IU/I)	$6.9 \pm 0.3$	$6.45 \pm 0.4$		7.8 ± 0.3	$6.5 \pm 0.4$	*	7.4 ± 0.2	$6.5 \pm 0.3$	*	7.3 ± 0.1	$6.5 \pm 0.2$	**
LH (IU/I)	$4.2 \pm 0.5$	$7.3 \pm 07$	***	6.1 ± 0.8	$6.7 \pm 1.0$		4.7 ± 0.3	$7.6 \pm 0.5$	***	4.9 ± 0.3	$7.1 \pm 0.4$	***
E <sub>2</sub> (ng/ml)	$42.0 \pm 3.1$	$44.4 \pm 4.0$		48.0 ± 3.1	46.4 ± 4.2		46.8 ± 2.8	47.9 ± 4.2		45.4 ± 1.8	47.1 ± 2.6	
Initial day												
FSH (IU/I)	$2.6 \pm 0.6$	$4.0 \pm 0.9$		3.3 ± 0.5	$3.3 \pm 0.3$		3.5 ± 0.3	$2.6 \pm 0.6$		3.1 ± 0.3	$3.5 \pm 0.4$	
LH (IU/I)	$1.4 \pm 0.1$	$0.9 \pm 0.2$	*	2.2 ± 0.4	$1.7 \pm 0.3$		2.2 ± 0.2	$1.1 \pm 0.3$	**	1.9 ± 0.1	$1.2 \pm 0.2$	***
E <sub>2</sub> (ng/ml)	20.3 ± 1.7	20.3 ± 2.5		41.70 ± 20.4	27.1 ± 5.5		22.0 ± 1.6	$26.3 \pm 3.0$		21.6 ± 1.2	23.5 ± 1.9	
hCG day												
LH (IU/I)	$2.0 \pm 0.2$	$1.4 \pm 0.3$		1.8 ± 0.4	$1.4 \pm 0.4$		1.5 ± 0.3	$2.4 \pm 0.5$		1.8 ± 0.1	$1.4 \pm 0.2$	
E <sub>2</sub> (ng/ml)	2615.7 ±	3081.8 ±		2387.6 ±	2419.0 ±		2933.6 ±	2398.4 ±		2622.1 ±	2794.9 ±	
	223.0	311.8		264.5	403.6		190.9	300.9		133.0	195.8	
Progesterone (ng/ml)	$0.8 \pm 0.1$	0.8 ± 0.1		0.8 ± 0.1	1.0 ± 0.3		0.9 ± 0.1	0.7 ± 0.1	*	0.8 ± 0.0	0.8 ± 0.1	
Clinic outcomes												
Fertilization methods												
IVF	35 (59.3%)			25 (69.4%)	17 (89.5%)		50 (64.9%)	22 (61.1%)		110 (64.0%)		
ICSI	24 (40.7%)	, ,		11 (30.6%)	2 (10.5%)		27 (35.1%)	14 (38.9%)		62 (36.0%)	24 (27.9%)	
Oocyte number	$10.7 \pm 0.7$	13.6 ± 1.0	*	8.3 ± 0.8	9.6 ± 1.4		12.7 ± 0.8	10.4 ± 1.1		10.8 ± 0.5	12.4 ± 0.7	
Fertilization	$63.3 \pm 3.4$	73.6 ± 4.8		75.9 ± 3.7	66.4 ± 5.8		71.0 ± 2.4	70.6 ± 3.6		69.3 ± 1.8	70.1 ± 2.7	
Cleavage	69.2 ± 3.5	$65.0 \pm 4.9$		91.5 ± 2.8	91.8 ± 2.9		89.8 ± 2.4	85.7 ± 3.6		83.7 ± 1.9	79.2 ± 2.8	
Number of good-quality embryos	2.7 ± 0.3	$3.0 \pm 0.5$		2.2 ± 0.4	$2.6 \pm 0.6$		3.5 ± 0.4	$3.3 \pm 0.5$		2.8 ± 0.2	$3.4 \pm 0.3$	
Cancellation	12 (20.3%)	10 (32.3%)		9 (25.0%)	8 (42.1%)		21 (27.3%)	16 (44.4%)	*	42 (24.4%)	34 (39.5%)	*
OHSS risk	7 (11.9%)	8 (25.8%)		3 (8.3%)	6 (31.6%)	*	15 (19.5%)	12 (33.3 <del>%</del> )		25 (14.5%)	26 (30.2%)	*
No embryos formed	5 (8.5%)	2 (6.5%)		16.7 (6/36)	2 (10.5%)		6 (7.8%)	4 (11.1%)		17 (9.9%)	8 (9.3%)	
Implantation rate (%)	37.8 ± 6.1	36.9 ± 9.5		27.7 ± 7.3	13.6 ± 7.0		30.7 ± 4.7	19.1 ± 8.2		31.8 ± 3.4	31.0 ± 5.5	
Pregnancy	24 (40.7%)			11 (30.6%)	3 (15.8%)		30.7 ± 4.7	5 (13.9%)	*	65 (37.8%)	21 (24.4%)	
Miscarriage	2 (3.4%)	13 (41.9%)		0 (0%)	0 (0%)		2 (2.6%)	0 (0%)		4 (2.3%)	1 (1.2%)	
Ectopic pregnancy	2 (3.4%)	0 (0%)		0 (0%)	0 (0%)		0 (0%)	0 (0%)		2 (1.2%)	0 (0%)	
Live-birth		12 (38.7%)		11 (30.6%)	3 (15.8%)		28 (36.4%)	5 (13.9%)		59 (34.3%)	20 (23.3%)	
LIVE-DII (II	20 (33.3%)	12 (30.7%)		11 (30.0%)	3 (13.070)		20 (30.4%)	J (13.5/0)		33 (34.3%)	20 (23.3%)	

Values are Mean ± SEM or Number (Percentage).

Chi-square test was used for categorical variables.

Unpaired t-test was used to analyze Age and BMI before the rest data were analyzed by covariance analysis with Age and BMI adjustment.

P values are indicated (\*<0.05, \*\*<0.01, \*\*\*<0.001) or otherwise >0.05