

ROLE OF DECODING OF AMH IN THE DIAGNOSTIC ADVANCEMENTS IN PCOS

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DOI: [10.5281/zenodo.12669954](https://doi.org/10.5281/zenodo.12669954)

Abstract

AMH has emerged as a pivotal biomarker for understanding the pathophysiology, diagnosis, and fertility implications of PCOS. This review delves into AMH's multidimensional involvement in PCOS, with an emphasis on pathogenic insights and diagnostic improvements. AMH levels in PCOS patients are significantly higher than in healthy controls, indicating the syndrome's etiology. According to studies, AMH is intricately linked to the hyperandrogenic and insulin-resistant milieu that characterizes PCOS, offering light on the underlying mechanisms that contribute to the disorder's development. Furthermore, AMH plays an important role in the diagnosis of PCOS. Its capacity to reflect ovarian follicular activity and reserve makes it useful for evaluating ovarian dysfunction and reproductive potential in PCOS patients. Diagnostic breakthroughs have used AMH measures to improve the accuracy of PCOS diagnosis, particularly in difficult instances where traditional approaches may fail. This review also looks at how AMH is progressing in therapy options for PCOS. It examines how AMH levels may influence treatment outcomes, such as ovulation induction and weight management, and provides insights into individualized therapy techniques.

Keywords: AMH, PCOS, Ovarian Reserve, Fertility, Diagnostic, Receptors.

INTRODUCTION

Polycystic ovarian syndrome (PCOS) is an elaborate endocrine condition that affects a large proportion of women globally, with estimates indicating a frequency of up to 10% among reproductive-age women [1]. Its distinguishing characteristics include hyperandrogenism, ovulatory failure, and polycystic ovarian morphology [2]. Despite being a prevalent disorder, the actual etiology of PCOS remains unknown, and its care provides therapeutic hurdles due to its variability and diverse clinical symptoms [3]. Over the past several years, there appears to be an increasing interest in investigating potential treatment strategies for PCOS that go beyond traditional symptomatic care [4]. Anti-Müllerian hormone (AMH), a glycoprotein in the TGF- β family, could be a potential target [5]. AMH, which has previously been recognized for its significance in male sexual differentiation during embryogenesis, has recently received attention for its role in ovarian folliculogenesis and probable contribution to PCOS pathogenesis [6].

This introduction tries to explain the rationale for addressing AMH in PCOS therapies, beginning with a summary of current PCOS pathophysiology, and then moving on to examining AMH's involvement in ovarian function and its consequences in the setting of PCOS. Furthermore, it will discuss the limitations of existing treatment modalities

for PCOS and highlight the potential benefits of targeting AMH as a more targeted and effective approach to managing this complex syndrome.

The basics of AMH expression and action

AMH gene

Anti-Müllerian hormone (AMH) (Fig. 1) is a glycoprotein peptide weighing approximately 140 kDa. It is categorized within the subfamily of growth factors called transforming growth factor beta (TGF- β) [7]. AMH gene lies on the short arm of chromosome 19, between the regions p13.2 and p13.3. It is separated into five exons, comprising 275 base pairs (bp) [8].

The AMH gene is transcribed through a 180-bp region that is contiguous to the protein Sap62. It has three transcription binding sites: one is a 20-bp conserved motif that binds the orphan nuclear receptor SF-1, and the other is 50 bp upstream from the SF-1 binding site and stimulates the binding of SOX9, a high-mobility group protein. The final binding site, located downstream of the SF-1 binding site, binds to GATA-4 from the GATA transcription factor family. When SF-1 binds to the promoter, transcription upregulates. Furthermore, the interaction between SOX9, WT1, and GATA-4 modulates this process [9]. The product of the AMH gene is a precursor known as proAMH, consisting of 560 amino acids [10]. Following the removal of the 24 amino acid fragment, a molecule is glycosylated, resulting in the formation of two identical subunits having a molecular weight of 70 kDa connected by sulphide bridges. The N-terminal domain known as the "pro-region" (115 kDa AMHN) and the C-terminal domain, or "mature region" (25 kDa AMHC), are formed as a result of proAMH proteolysis [10]. The N-terminal segment is necessary for the protein to be active, which makes it unlikely that the other proteins in the TGF- β complex would be affected. The C-terminal segment is required for the biological functions of the protein and its binding to receptors [11].

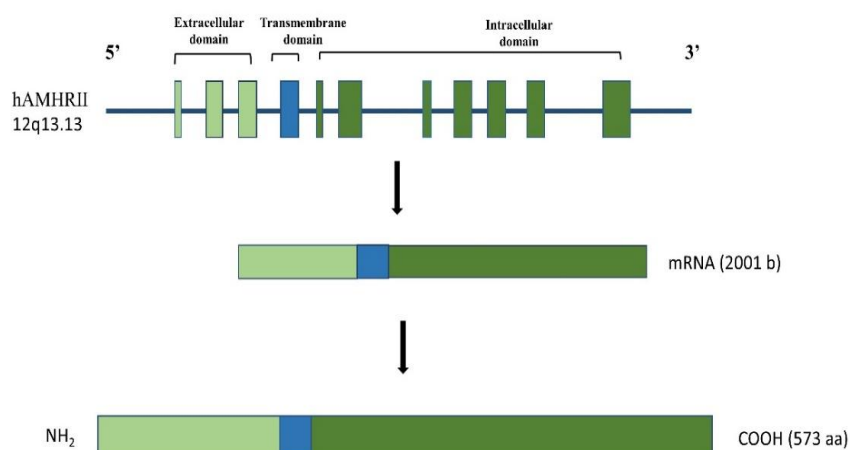


Fig 1: The human AMH receptor (AMHR II) gene, mRNA, and protein. The human AMHR II gene localized in chromosome 12, 12q13.13, is transcribed into 2001 base (b) mRNA, which is then translated into a 573 amino acid (aa) protein

AMH Receptors

The AMH signaling pathway is regulated by two heteromeric serine/threonine kinase transmembrane receptors, categorized into two types: type I and type II [12].

Anti-Müllerian hormone receptor I (AMHRI) is phosphorylated when free AMH binds to AMHRII. This phosphorylation activates cytoplasmic Smad proteins, which start downstream signaling. More specifically, AMH initiates Smad1 protein signal transduction but not Smad2 protein [12].

The Smad proteins translocate into the nucleus after becoming phosphorylated, where they can either stimulate or inhibit the transcription of particular genes [9].

The only receptor available for the AMH hormone is the AMH type II receptor. It expresses itself in the ovary soon after birth and lasts the entirety of an individual's life [13]. The ductal epithelium of the mammary gland, the prostate, the endometrial, the Sertoli and Leydig cells in the testes, and the theca and granulosa cells in the ovaries are all known to contain the AMHRII receptor [14–17].

Numerous cancer cell lines, including those from the ovarian, cervical, endometrial, and breast epithelium, have also been shown to express the receptor AMHRII [8,18].

In recent years, it has also been demonstrated that organs other than the gonads, including the motoneurons [19], neurons [20], hypothalamus [21], and gonadotropic cells of the pituitary gland [22], also express AMH, as shown in Table 1.

However, since serum AMH's expression profile corresponds to that of the gonads, this expression has little effect on circulating levels. The fact that AMHRII is more widely expressed emphasizes the importance of the effects of AMH shown in tissues that express AMHRII.

Table 1: AMH and AMHR2 distribution in both sexes

	Organs	Men	Women	Both	Reference
AMH Expression	Ovaries		****		23
	Testes	****			24
	Nervous system			*	25,26
	Pituitary gland			*	27
	Hypothalamus			*	28
AMHR2 Expression	Testes	****			24
	Ovaries		****		23
	Nervous system			*	25,26
	Pituitary gland			*	27
	Hypothalamus			*	28
	Uterus		**		23
	Placenta		*		23
	Breasts		*		29
	Pancreas			*	30
	Lungs			*	31
	Adrenals			**	30
	Prostate	*			32

The number of * represents the level of expression of AMH and AMHR2 in the different tissues.

Expression of AMH

AMH expression begins in primary follicles in the ovaries of every species that has been studied thus far. It peaks in pre-antral and small antral follicles, declines in large antral follicles, and eventually becomes almost undetectable—except in cumulus cells [23].

AMH secretion in the gonads corresponds to its detection in serum, even though AMH is expressed in endometrial and endometriotic cells [33].

For instance, there is an undetectable level of AMH in the serum of women who have had their ovaries removed. Beginning in the 36th week of pregnancy, the ovarian granulosa cells (GC) of the pre-antral and antral follicles release AMH in females [34].

Serum AMH levels are two to four times higher in PCOS-affected women and in their daughters. The higher number of small antral follicles, which express AMH the most, as well as the overexpression of AMH by these GCs are the two causes of the raised serum AMH levels in women with PCOS [35].

Modified GC receptivity to numerous hormones dysregulated in PCOS might lead to the overexpression of AMH and AMHR2. Indeed, androgens increase in vitro AMH mRNA levels exclusively in GCs from PCOS patients which overexpress the androgen receptor [23].

Estradiol (E2), on the other hand, mitigates AMH expression in the GCs of normal women while not regulating the expression of this gene in the GCs of PCOS women [36]. LH and E2 reduce AMHR2 mRNA levels in GCs from normal women, however, they are not modulated by these hormones in GCs from anovulatory women with PCOS.

Anti-Müllerian Hormone (AMH) regulates the number and selection of developing follicles for ovulation [37]. It functions as a negative regulator in the early stages of follicular development [38]. Essentially, AMH suppresses follicle recruitment and growth by lowering the influence of growth factors and gonadotropins, particularly follicle-stimulating hormone (FSH) [39]. To put it simply, AMH regulates the reproductive cycle by limiting the number of follicles that mature and preventing excessive proliferation.

Immature Sertoli cells in the testes generate AMH during embryonic development. AMH secretion begins in the ninth post-conception week and lasts until 2 years postnatally. It decreases during puberty and becomes undetectable in adults owing to high testosterone levels [40,41]. Whenever a male fetus has low or no AMH levels, both male and female genitalia develop simultaneously.

In the absence of AMH, the Müllerian ducts form the fallopian tube, uterus, cervix, and upper one-third of the vagina. Changes in AMH levels or receptors are capable of impacting the development of the female reproductive system.

In women, production begins in the 36th week after conception. After a brief neonatal rise, AMH levels remain low until puberty. Adolescent girls experience escalating AMH serum levels that eventually plateau. AMH serum levels start to decline in the mid-20s and become undetectable many years before menopause [42,43].

AMH also alters follicular growth and dynamics. It serves as an antagonist of follicle recruitment, effectively halting the process of primordial follicles developing into

expanding follicles. This mechanism contributes to follicular pool balance and regulates the follicle maturation rate. AMH also modulates follicular sensitivity to follicle-stimulating hormone (FSH), an important hormone in ovarian function [44].

At different phases of development, AMH affects the hypothalamic-pituitary-gonadal (HPG) axis and is implicated in the formation of ovarian follicles [45].

The number of antral follicles in the initial follicular stage of the menstrual cycle strongly influences the number of ovarian follicles, which is reflected in the serum AMH concentration [45,46,47].

Consequently, low serum AMH levels can be attributed to a decreased antral follicle count. Furthermore, neither the menstrual cycle phase nor exogenous sex steroids affect the blood level of AMH. Consequently, any day of the cycle can be used to collect blood samples [48].

The fact that AMH is a validated biomarker of female reproductive potential should be underlined. It represents the quantity of primary follicles and how they react to exogenous gonadotrophins [45,47,49].

Mechanism of action (MOA) of AMH (Fig. 2)

AMH in humans is activated by cleaving its 140-kDa precursor at a certain location, which results in a 110-kDa N-domain and a 25-kDa C-terminal domain which carries the bioactive site [50]. Although the half-lives of the precursor and its noncovalent forms in female serum are similar, methodological differences have made it difficult to determine the exact half-life [51]. The cleavage most likely takes place in the producing cells and is mediated by enzymes PC5 [52,53,54]. While serum and follicular fluid include both precursor and noncovalent forms, serum contains more of the latter, signifying further cleavage or degradation from outside cells. But without the N-terminal domain, the instability of the C-terminal segment makes it unlikely to circulate on its own. More research is necessary to fully understand the cleavage dynamics and functional consequences in the ovary [55].

AMH mostly interacts with AMHR2 via its C-terminal domain, most likely as a noncovalent complex. This notion is corroborated by the fact that AMH's N-terminal segment increases the C-terminal domain's activity and that, in lab settings, the noncovalent complex and the pure C-terminal fragment have similar affinities for AMHR2 [55].

While the noncovalent complex from follicular fluid efficiently binds to AMHR2 in vitro, the serum-derived complex fails to do so, implying that serum-related variables may impair AMH bioactivity. Laboratory investigations have demonstrated that when AMH and AMHR2 bind, the N-terminal domain of the noncovalent complex is cleaved, allowing for AMH signaling, with the C-terminal version being more potent than AMH noncovalent complex in stimulating the phosphorylation of SMAD 1, 5, or 8 [54,56].

The Q496 residue in the C-terminal domain is critical for promoting interactions with AMH type I receptors. Mutations at this location cause chronic Müllerian duct syndrome in men, making the AMH physiologically inactive. Further study on AMHR2 residues and their involvement in AMH signaling, together with experimental data indicating early monomer binding before dimerization, highlights the intricacy of AMH-mediated pathways [57].

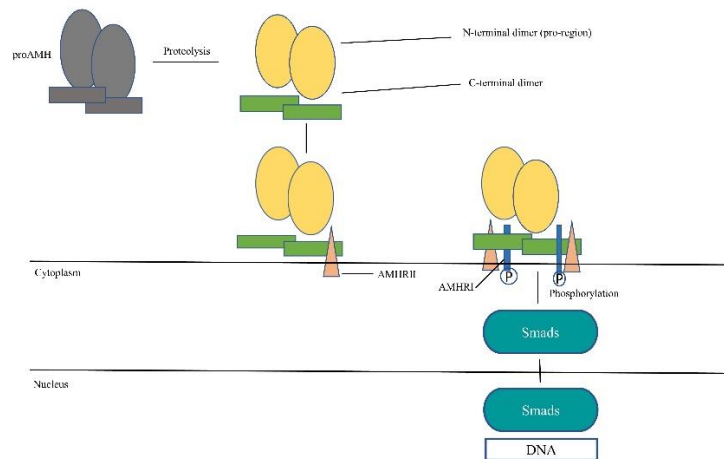


Fig 2: The mechanism of action of AMH action via receptor. Proteolysis of proAMH leads to a conformational change in the C-terminal domain that allows binding AMHR2. AMHR2 induces binding to the type I receptor, which is then phosphorylated by a kinase receptor type II.

AMH variability in PCOS

PCOS is a prevalent endocrine condition that impacts around 5 to 10 percent of women in their reproductive years worldwide [58]. According to the Rotterdam criteria, the diagnosis needs to include at least two of the following: polycystic ovarian morphology visible on ultrasound, evidence of elevated androgens either in biological tests or clinical signs like acne and hirsutism (excessive hair growth), and ovulation irregularities [59].

Four phenotypes can be defined by these criteria, the most severe of which combines all three features. Disruptions in the GnRH/LH pulsatility are also linked to PCOS, and these can result in high LH (luteinizing hormone) levels and frequently a raised LH/FSH (follicle-stimulating hormone) ratio [60]. PCOS-afflicted women frequently have metabolic difficulties such as weight gain, resistance to insulin, elevated levels of lipids, and type 2 diabetes.

Given that 20% to 40% of individuals exhibit familial clustering of PCOS symptoms, there is evidence of a genetic component [61]. Developmental variables may also be involved since, by the theory of the Developmental Origins of Health and Disease, abnormalities during the foetal and prepubertal stages might have long-term effects on reproduction and metabolism [62,63].

Levels of AMH are 2- to 4-fold greater in follicular fluid and serum of PCOS patients [64]. This rise is explained by the overexpression of AMH and its receptor (AMHR2) by granulosa cells (GCs), which may have been influenced by hormonal dysregulation, in addition to the number of tiny antral follicles releasing AMH [65,66].

According to research, hyperandrogenism, a defining symptom of PCOS, causes granulosa cells (GCs) to express more AMH in afflicted women. Although factors such as FSH, estradiol, and follicle proliferation under androgen influence can complicate this association, research consistently reveals a positive relationship between androgens and AMH levels for PCOS patients [67,68].

5 α -dihydrotestosterone (5 α -DHT), a non-aromatizable androgen, specifically enhances AMH mRNA levels in GCs from PCOS patients with elevated androgen

receptors (AR) [69]. This proposes a direct androgen-induced mechanism responsible for AMH overexpression in PCOS.

These findings emphasize the importance of hyperandrogenism in the dysregulation of AMH in PCOS, as well as the intricate interplay between androgens, follicular growth, and AMH production. Understanding these molecular processes is critical for deciphering PCOS pathogenesis and creating targeted treatments.

Although research on the AMH-insulin resistance link in PCOS is inconsistent, studies such as Liu et al. show that insulin can dose-dependently boost AMH mRNA expression in GCs from both PCOS and normal women [36]. Furthermore, enhanced AMH cleavage in association with metabolic parameters shows that metabolic variables may worsen AMH dysregulation in PCOS, especially given these women's heightened vulnerability to metabolic diseases.

The modulation of AMHR2 expression in PCOS has received little attention, although it is emerging as an important factor. In vitro studies indicate that the increased AMHR2 expression in PCOS granulosa cells (GCs) may be related to the lack of the inhibitory effects of estrogen (E2) and luteinizing hormone (LH) seen in normal women [23].

Recent suspicions have emerged about the role of AMH overexpression in PCOS pathogenesis. Certain single nucleotide polymorphisms (SNPs) in the AMH and AMHR2 genes, such as AMH Ile49Ser and AMHR2-482A>G, have been associated with PCOS and demonstrate decreased bioactivity. Furthermore, heterozygous mutations near AMH and AMHR2, which are seen in certain PCOS-affected individuals with low serum AMH levels, reinforce this association [70-73].

While these findings may explain enhanced theca cell testosterone synthesis and primordial follicle recruitment, such effects could also be caused by increased androgen and LH expression, respectively. However, a reduction in AMH/AMHR2 system activity alone does not fully account for the range of reproductive problems reported in PCOS [74].

These findings emphasize the intricate interplay of genetic, hormonal, and regulatory variables that contribute to PCOS pathogenesis. To precisely determine how AMHR2 dysregulation contributes to the variety of symptoms associated with this condition, further research is required.

AMH Measurement

AMH has been measured using a variety of commercial enzyme-linked immunosorbent test (ELISA) kits, each with its differences in antibody pairings, standard curve ranges, and limits of detection [75] (Table 2). The development of AMH tests is an important step forward in clinical epidemiology. Nevertheless, despite these advancements, there are still no well-recognized, standardized assay techniques or materials for determining serum AMH concentrations. Furthermore, the lack of conversion techniques to evaluate the comparability of AMH assays makes it difficult to interpret results from various research studies. For any clinical application of AMH, these exact definitions are necessary. With a lower detection threshold, the Ansh Laboratories picoAMH test is particularly notable for being the most sensitive assay for identifying low amounts of AMH [76].

Table 2: Measurements of different AMH assays

AMH Assay Year	Manufacture Company	Limit	Standard Curve Range	Description
IOT, 1999	Immunotech	0.05 ng/mL	0.1-24.5 ng/mL	A monoclonal antibody pair was directed, one directed at the pro region and the other at the mature region.
DSL, 2003	Diagnostic systems laboratories	0.006 ng/mL	0.05-15 ng/mL	Both monoclonal antibodies were directed at the mature region
GEN II Generation, 2010	Beckman coulter	0.16-22.5 ng/mL	0.08 ng/mL	The DSL antibodies were used in the assay, which was standardized to the IOT assay
Ultrasensitive, 2012	Ansh labs	0.083-14.2 ng/mL	0.023 ng/mL	Monoclonal antibody pair directed against specific linear epitopes in the stable pro region and mature region of the associated form of human recombinant AMH
PicoAMH, 2013	Ansh labs	0.001-0.746 ng/mL	0.001 ng/mL	

The following are the limitations of AMH:

- Women's AMH levels fall at different rates with age. It is still necessary to develop an international standard for age-specific AMH diagnostic thresholds for assessing functional ovarian reserves or estimating menopausal age [77].
- There are a few endogenous and exogenous factors that may affect serum AMH levels, making it more difficult to accurately interpret AMH results in a clinical environment [78].
- AMH levels shouldn't be used as a fertility test because the predictive value of AMH for a successful clinical pregnancy (in both natural and assisted reproduction) is not very high [79].
- Due to intra-assay/interassay variations, the manual enzyme-linked immunosorbent assay (ELISA) for AMH evaluation has limitations and requires careful sample preparation and preservation. On the other hand, automated AMH assay technologies provide increased sensitivity, improved precision, quicker turnaround times, and national accessibility. Because of their better performance and useful benefits, automated platforms ought to be used as the industry standard for calculating AMH [77].

Role of AMH as a marker of ovarian reserve

The quantity and quality of eggs, or oocytes, in the ovaries are referred to as ovarian reserve. An ovary that has lost ovarian reserve as a result of age-related factors is considered senile. One important indicator for determining ovarian reserve is the quantity of primordial follicles, or early-stage eggs [80]. It can be difficult to quantify this count precisely, though Research indicates that there may be a relationship between the number of follicles that enter the pool of growing follicles and the length of the primordial follicular pool [81-83]. The only source of AMH is these developing follicles. Consequently, the amount of AMH in the blood can serve as a proxy for the primordial follicle pool's size, which in turn represents ovarian reserve. Numerous research investigations have noted this association [84-87]. An initial quantity of

oocytes, typically ranging from one to two million, is present in females at birth [88]. AMH inhibits Leydig cell differentiation and promotes follicular maturation, but it is not crucial to ovarian differentiation [89]. By attracting primordial follicles during folliculogenesis (Fig. 3), it plays a crucial role in controlling follicular maturation. The amount of ovarian reserve and a woman's potential for ovulation are directly correlated with the levels of AMH found in serum [90].

An additional differentiation can be pointed out between a functional ovarian reserve (FOR) and an ovarian reserve. There is no correlation between the blood concentration of AMH and the number of primordial follicles in young women since AMH is expressed by a certain population of developing follicles [91]. The pool of follicles, with a diameter of 2 to 5 mm, among which one follicle is chosen by FSH and will ovulate, is known as the FOR [90]. FOR release AMH during the follicular phase. The quantity of developing follicles drawn from the primordial pool is indicated by the blood's level of AMH. Since there isn't a direct blood marker for ovarian reserve, this functions as an indirect indicator. Serum AMH levels essentially indicate the activity of developing follicles and offer important insights into ovarian reserve [90]. Serum levels of AMH gradually rise to a peak and plateau at age 25; the serum level begins to decline then. Individuals with elevated AMH levels who do not have PCOS are typically quite fertile [92]. It is crucial to remember that serum levels of AMH are influenced by various factors, including vitamin D and hormonal contraception. Women using hormonal contraception had reduced AMH serum levels, which range from 14 to 55% [91]. levels of AMH are affected during pregnancy. The second and third trimesters are marked by a significant fall in AMH levels, which suggests that ovarian activity is lower during this time. On the other hand, compared to preconception levels, the AMH concentration is rather steady during the first trimester and recovers to normal following delivery [93]. Recent research suggests a link between higher serum levels of vitamin D and higher amounts of AMH in the bloodstream. Vitamin D and AMH levels might change periodically throughout the year [92]. This link is due to the occurrence of a vitamin D-responsive element inside the AMH gene promoter, which allows vitamin D levels to impact AMH concentrations [94].

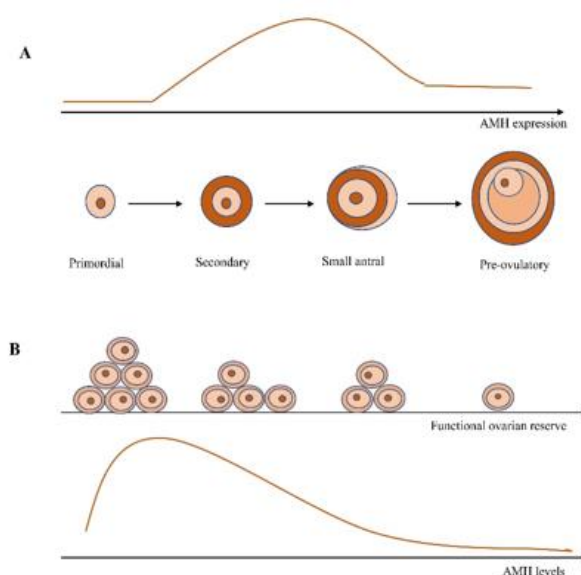


Fig 3: Anti-müllerian hormone (AMH) expression and concentration about folliculogenesis and ovarian reserve

AMH and cancer gynecological tumors and cancer

One common type of gynecological cancer is ovarian cancer, which comes in three primary categories: germ cell tumors, stromal tumors, and epithelial tumors. Of these, 82% of all ovarian malignancies are epithelial ovarian tumors, which are the most common type. In contrast, germ cell tumors account for 3% of ovarian cancers and are further divided into juvenile and adult types [95]. It is interesting to note that AMH can affect the development of breast, endometrial, and prostate cancers in addition to its main association with germ cell malignancies [15,32,96]. Considering its stability throughout the menstrual cycle, AMH is a reliable tumor marker for both primary and recurrent GCT incidence, in addition to inhibin B [97].

Although serum AMH levels are higher in these individuals than in those without tumours, it can only be used effectively in patients under the age of 65 [98]. Regardless of their illness, elderly women's serum AMH levels are too low to assess. The higher level of AMH in tumors is attributed to bigger granulosa cells that release more AMH into the circulation [97]. The same positive connection between high circulating AMH levels and an increased risk of breast cancer has been discovered in breast cancer patients [99].

While most ovarian cancers emerging from the Müllerian tract are thought to grow via the fimbriated end of the fallopian tube, there are distinct tumor types that arise from the secondary Müllerian system [100]. The involvement of AMH in the regression of Müllerian ducts during male gender development in embryos has prompted scientists to investigate its potential as a therapy for epithelial ovarian cancer [101]. Recombinant AMH has been shown in studies to successfully limit the growth of several ovarian cancer cell lines, including OVCAR 8 and IGROV 1. These findings point to a possible option for using AMH in targeted therapy for epithelial ovarian cancer [102]. The levels of AMH differ between healthy individuals and cancer patients, which has consequences for ovarian reserve. For example, young breast cancer patients (28–44 years old) had AMH levels comparable to healthy women of a comparable age (30–44 years old), implying that breast cancer has no substantial impact on ovarian reserve [103].

However, those suffering from Hodgkin lymphoma had lower AMH levels and a reduced ovarian reserve when compared to healthy women [104]. Notably, serum AMH levels can indicate post-cancer treatment ovarian function and prospective recovery; greater AMH levels correspond to faster restoration of ovarian function and fertility. As a result, testing AMH in patients' serum is regarded as a more robust and reliable approach to determining ovarian reserve than measuring FSH or inhibin B levels [105].

Artificial Reproductive Technology

Exogenous FSH treatment affects the hormonal modulation of ovarian function in women with reproductive challenges. Predicting an ovarian response before stimulation allows for personalized counseling and optimal gonadotrophin dosage for each patient [106]. Assisted reproductive technology (ART) results are strongly reliant on the ovarian response to stimulation, which reflects ovarian reserve in terms of egg number and quality. Because of its higher sensitivity than procedures such as day-three FSH, AMH measurement has become routine practice in ART centres [107]. A low response to controlled ovarian hyperstimulation (COH) is commonly characterized as extracting five or fewer oocytes or canceling the cycle, although the optimal range

for oocyte retrieval is between 10 and 12 [108]. To optimize stimulation, national recommendations advocate adjusting gonadotropin doses depending on individual ovarian reserve markers, such as specific AMH levels [12].

Endometriosis is prevalent amongst infertile women, with studies indicating that 20-50% of those facing infertility have endometriosis [109]. Endometriosis and infertility are associated via complicated pathways, with reduced ovarian reserve being indicated as a contributing factor. Chronic inflammation, increased oxidative stress, dysregulated cell cycles, and endometriosis-related poor angiogenesis all contribute to this impairment [110]. Kitajima et al. found reduced AMH levels in peritoneal fluid among women with endometriosis compared to a control group without the disorder. This study emphasizes the impact of endometriosis on ovarian function and the potential role of AMH as a diagnostic marker in measuring ovarian reserve in women with endometriosis-related infertility [111]. Endometriosis is usually treated with an amalgam of surgery and medications, such as pain relievers and hormone therapy [112]. However, these treatments frequently fail to cure the disease and are linked with high rates of clinical recurrence. While surgical intervention is successful at lowering symptoms, it can also reduce ovarian reserve and AMH levels.

Laparoscopic surgery has been regarded as the primary treatment for endometriosis-related infertility, with assisted reproductive technology (ART) as a backup option [109]. A combination of laparoscopic surgery and ART has resulted in higher pregnancy rates in endometriosis-related infertility cases. However, such procedures have the potential to produce iatrogenic damage, including ovarian reserve loss and scar formation [109], particularly following ovarian surgery, which specifically lowers AMH levels and antral follicle count (AFC), contributing to diminished fertility [113]. To address the reduction in fertility associated with endometriosis treatment, women with endometriosis can get oocyte cryopreservation as a preoperative fertility preservation therapy. This method seeks to reduce the effect of surgical operations on ovarian function and fertility [109].

Menopause

AMH has emerged as a useful marker for predicting menopause age, providing important information about women's reproductive health and possible fertility loss [114]. As women age, their ovarian reserve steadily declines, resulting in menopause—a natural physiological shift that signals the end of reproductive potential [115]. Given that genetic variables have been demonstrated to influence menopause, a woman's mother's age at the onset of menopause may be a good indicator of her own. It is known that genetic variations account for up to 50% of the variation in menopausal age [116]. Ovarian follicles produce AMH, which indicates the number of remaining primordial follicles in the ovaries and so serves as an indirect indicator of ovarian reserve [117]. Lower levels of AMH are often associated with decreasing ovarian function and imminent menopause. Dolleman et al. found from population-based cohort studies that the AMH predicts menopause onset age more accurately than the mother's age hereditary factor [118]. The accuracy of the AMH in predicting the onset of menopause over estradiol, FSH, or inhibin B has also been demonstrated by another research. From the moment of a female's birth, the AMH progressively rises and peaks at approximately 25 years of age. After that, AMH levels in post-menopausal women start to progressively decline until they are almost undetectable [119]. On the other hand, some argue that AMH may not be a reliable indicator of total

ovarian age. This opinion is supported by De Kat et al., who note that although AMH and other ovarian reserve indicators are connected to menopause age, they might not accurately depict the precise course of an individual's ovarian aging cycle [120].

While AMH was a strong predictor of time till menopause (TTM) and time until early menopause, it was discovered in a prospective cohort research by Depmann et al. that AMH had low predictive power for the age at which menopause begins. This suggests that although AMH can be a helpful predictive model for younger women who want to know if they might go through menopause early, its accuracy in identifying the precise age at which menopause begins may have some mistakes [121]. Another important factor to note is that as women age, AMH's predictive capacity declines. This suggests that, while AMH may be more trustworthy in predicting menopause timing in younger women, its accuracy declines as women approach menopause. These findings underscore the complexities of utilizing AMH alone to predict menopausal age, implying that other factors, such as genetic predispositions and lifestyle effects, also play important roles in determining the time of menopause onset. Thus, while AMH is still a useful tool for monitoring ovarian reserve and possible fertility reduction, its capacity to predict the exact age of menopause onset should be regarded with caution, particularly in older women or those with unique health profiles.

CONCLUSION

Serum anti-Müllerian hormone (AMH) levels are significantly higher among individuals with polycystic ovary syndrome (PCOS) than in their healthy counterparts, underscoring the significance of AMH as a PCOS diagnostic tool. It is especially beneficial in situations where evaluating the ovaries is difficult, including in obese, nulliparous, and individuals with limited echogenicity during ultrasound examinations, where reliable determination of the antral follicle count (AFC) is challenging.

Although AMH helps identify PCOS and provides insight into its pathophysiology and phenotypic variations, determining an exact AMH concentration threshold is a difficult task. Nevertheless, AMH has the potential to develop into a reliable diagnostic tool for PCOS with improved assay standardization. AMH is linked to the suppression of follicular growth and ovulation in the context of PCOS pathophysiology, which may add to treatment complications. Further in-depth investigations are imperative to elucidate AMH's intricate role in PCOS etiology, paving the way for targeted therapeutic interventions.

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