

IMPROVING THE PREDICTION OF OVARIAN RESERVE AND RESPONSE IN ART USING BIOMARKERS

Christos Venetis

MD, FRANZCOG, MSc (Res), MSc (HCM), PhD

University of New South Wales

Sydney, Australia

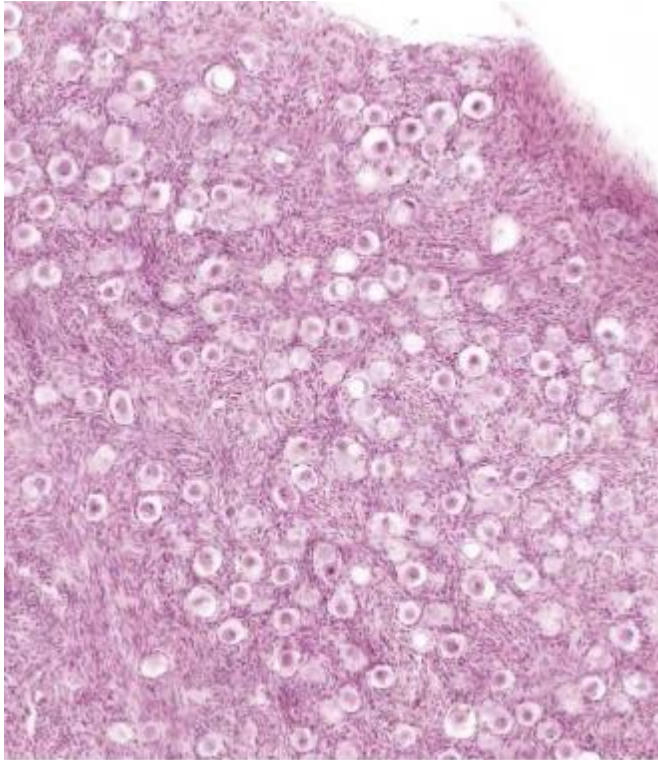
c.venetis@unsw.edu.au



Disclosures

- Consultancy fees and Honoraria received for speaker engagements:
Merck; Merck, Sharp & Dohme

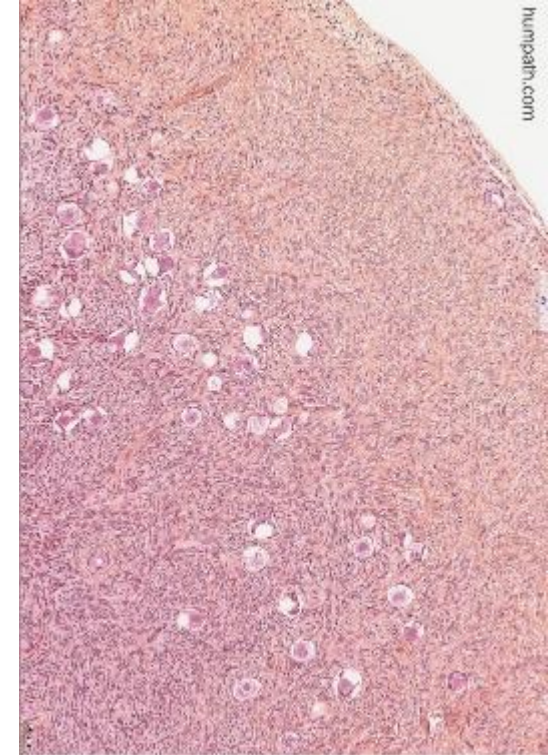
Ovarian reserve in humans (1)



Infant



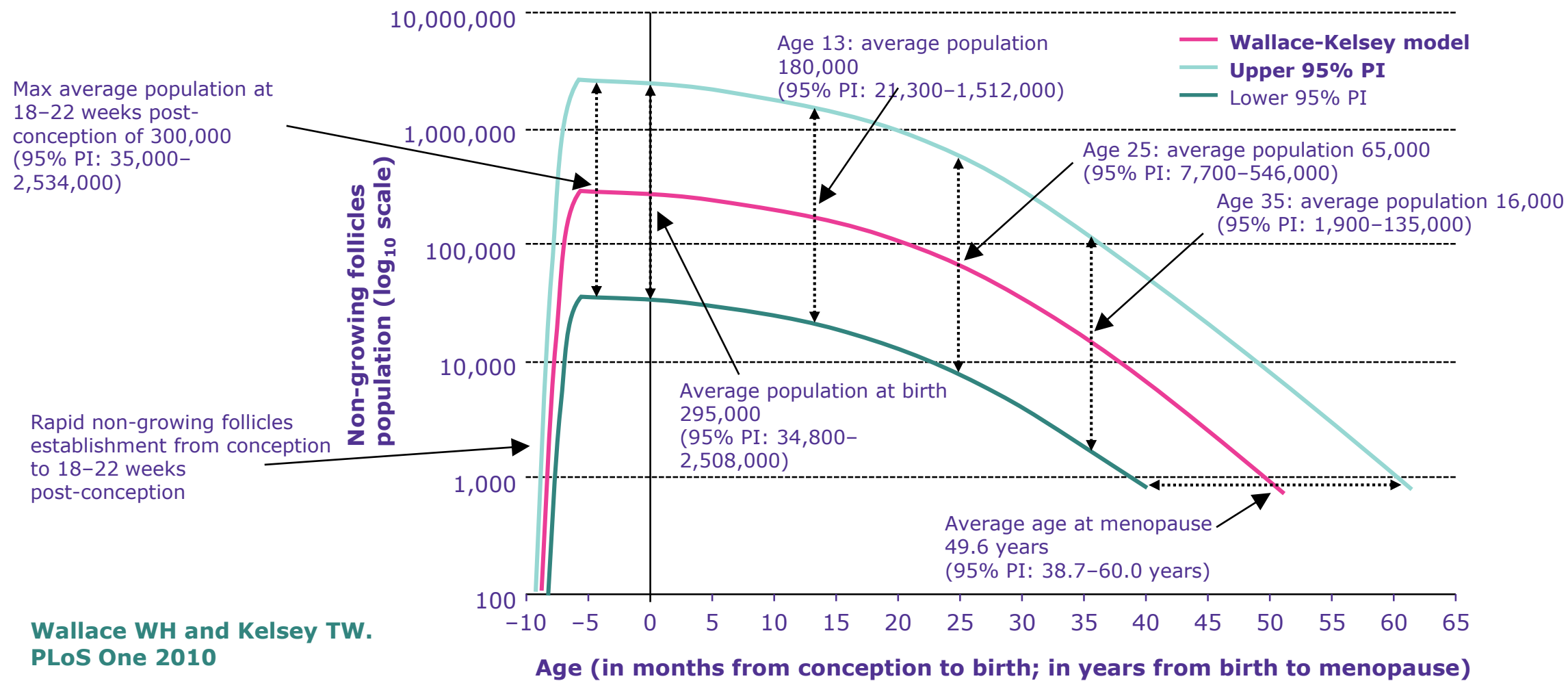
Reproductive age



Menopause

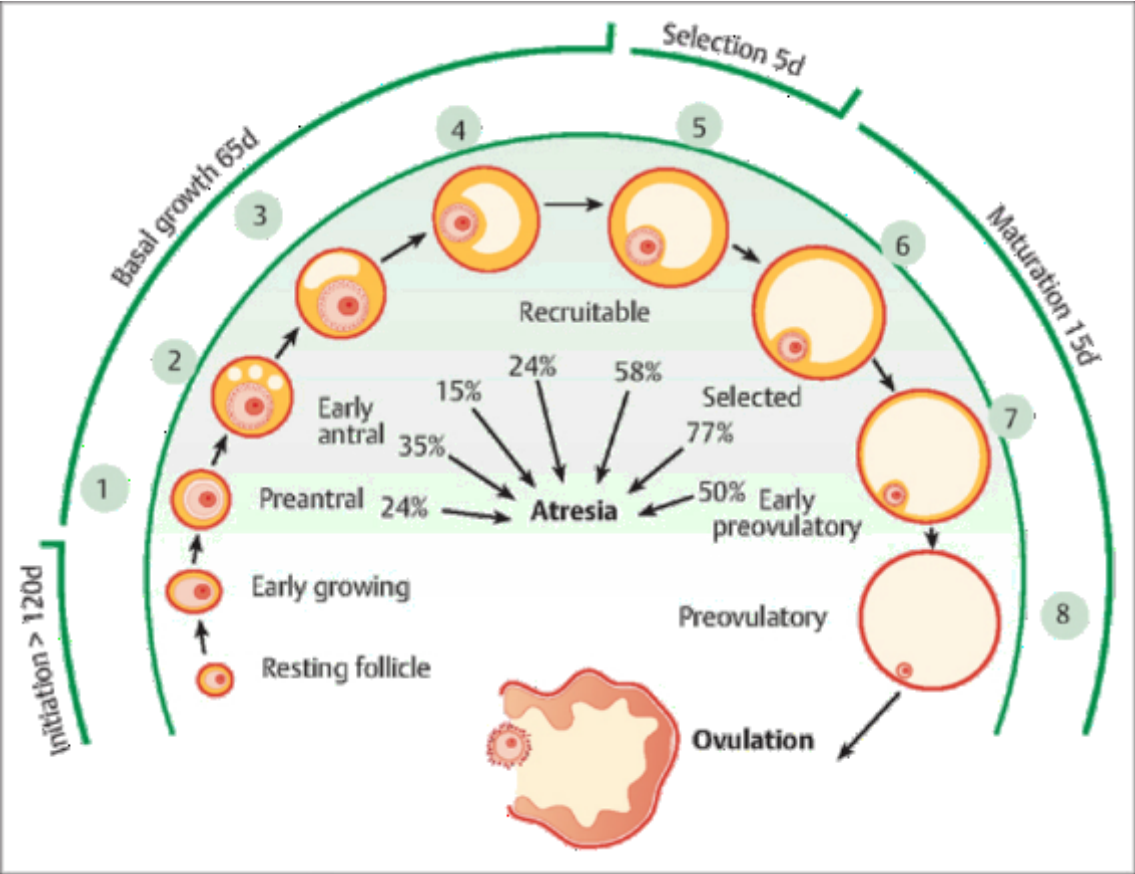
Dramatic loss of primordial follicles with advancing age

Ovarian reserve in humans (2)

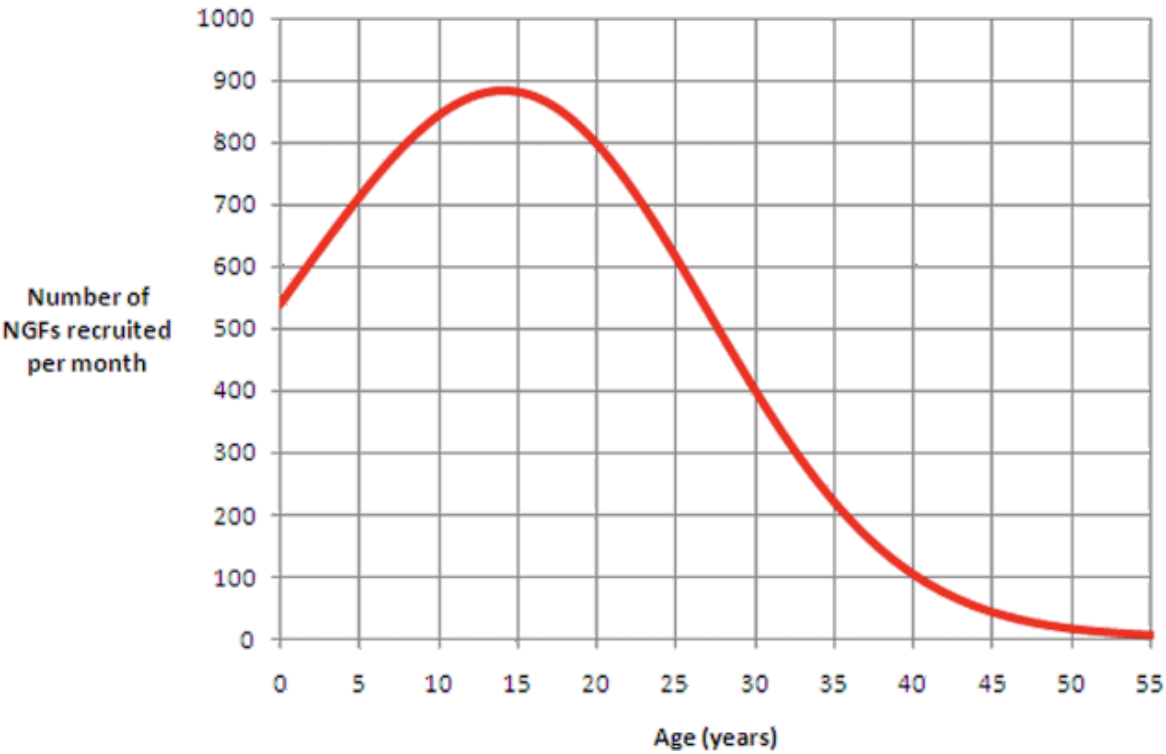


Wallace WH and Kelsey TW.
PLoS One 2010

Folliculogenesis/oogenesis in humans



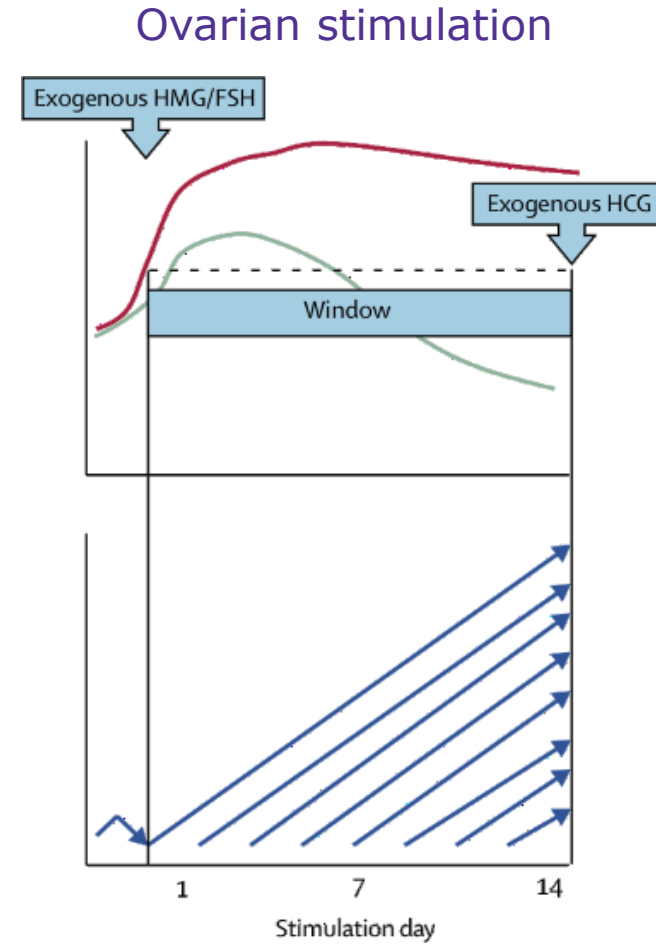
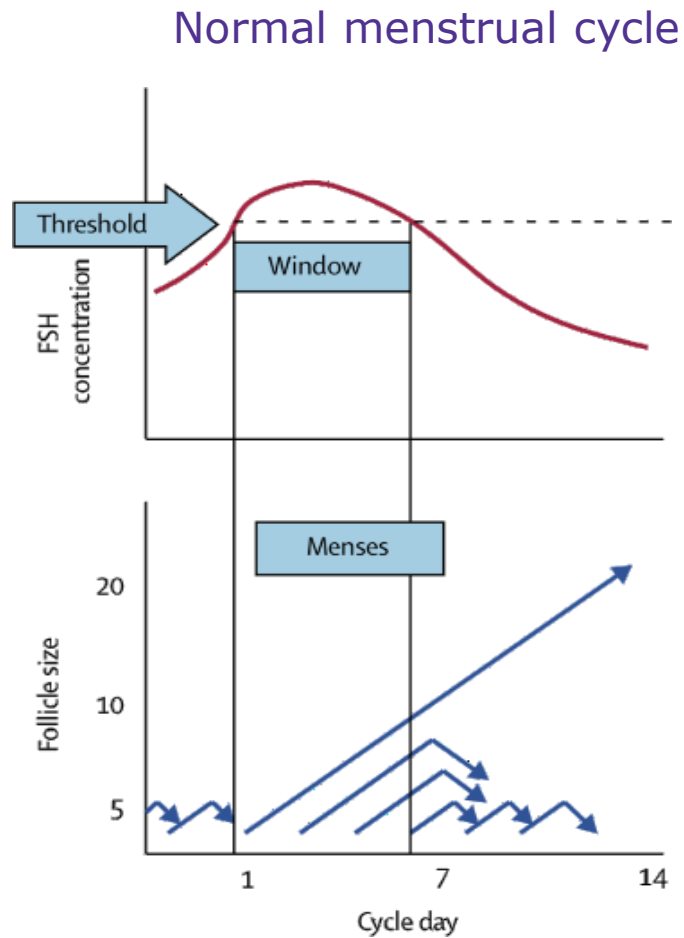
NGF recruitment - average age at menopause



Wallace WH and Kelsey TW. PLoS One 2010

Ovarian stimulation

Follicular rescue

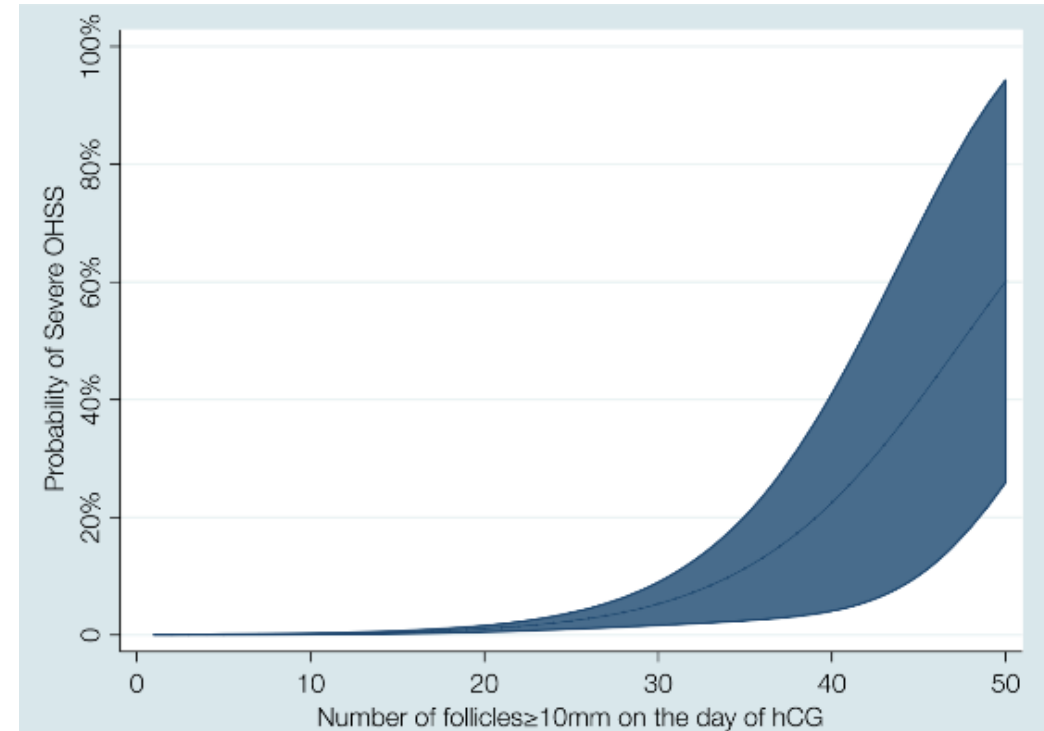


Fauser BC et al. Lancet 2005

Why is prediction of ovarian response important? (1)

Poor ovarian response
is associated with
disappointingly low pregnancy
rates and is one of the most
challenging patient groups
to manage

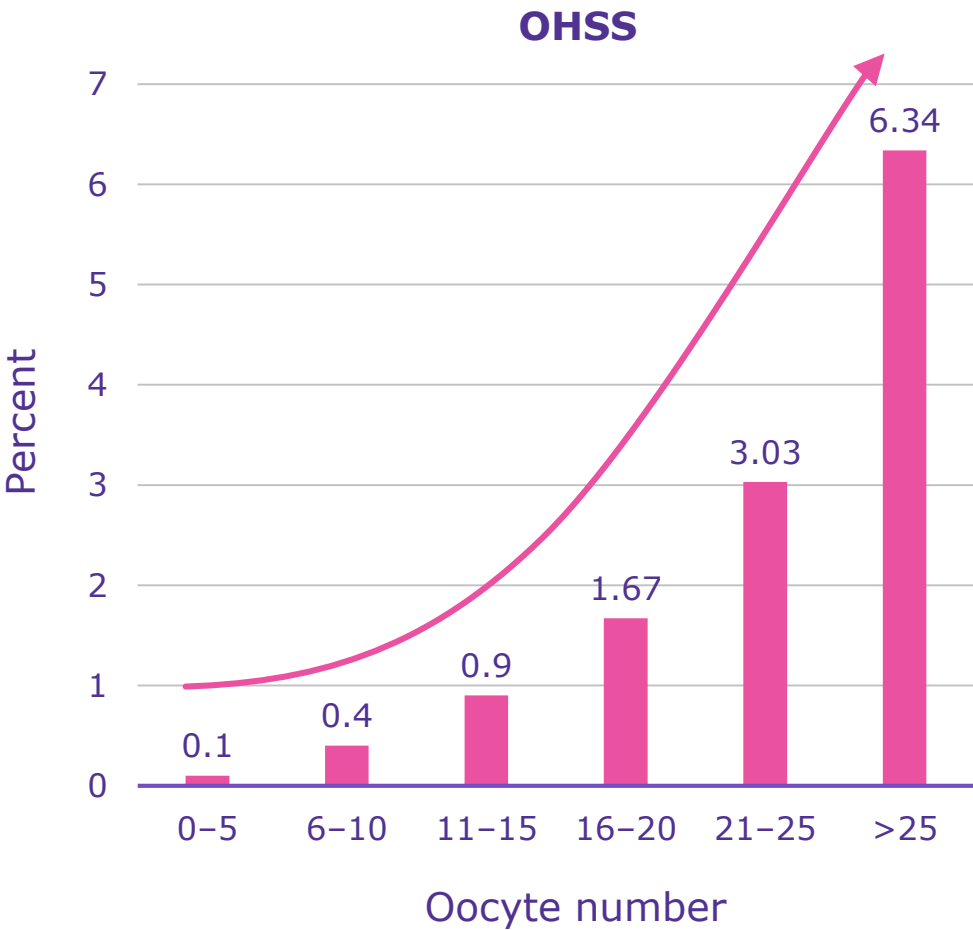
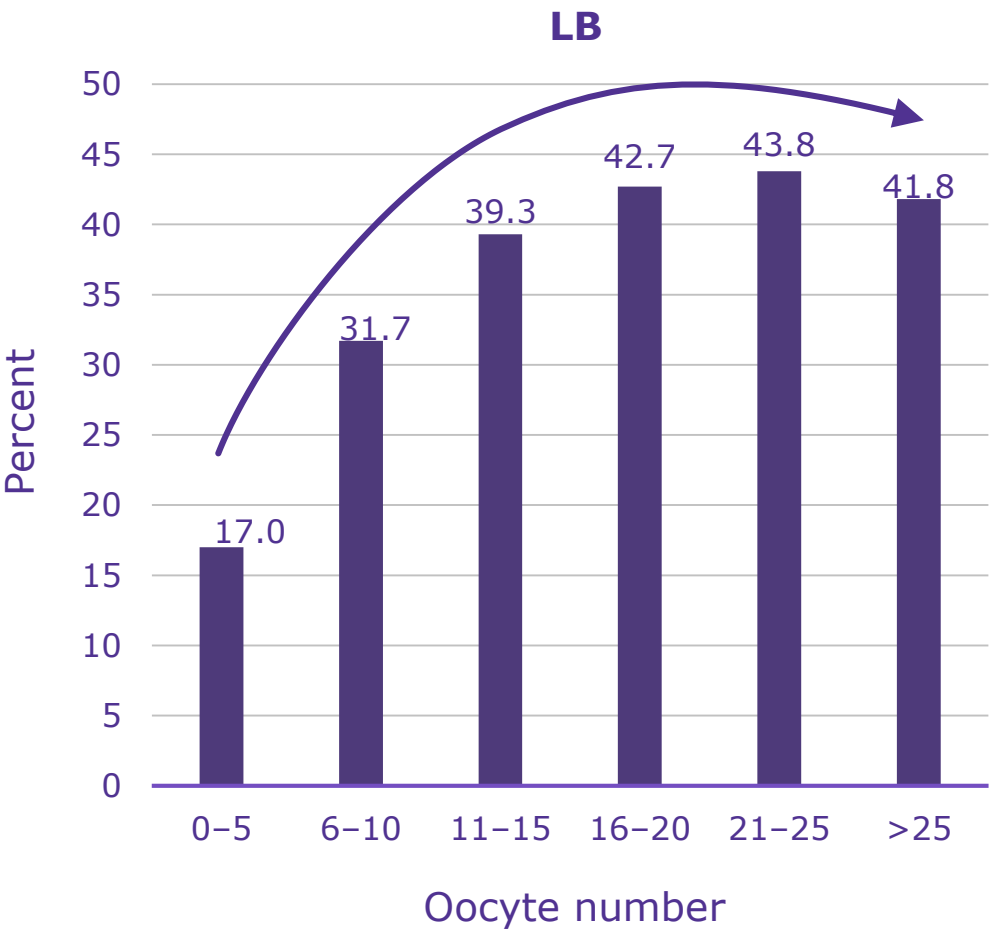
Venetis C, et al. Ann NY Acad Sci 2010



Increased number of follicles (>10 mm) on
the day of triggering final oocyte maturation
= increased risk of severe OHSS

Tarlatzi TB, et al. JARG 2017

Why is prediction of ovarian response important (2)



Note: percent LB and OHSS are per retrieved oocyte numbers per IVF cycle among SART members from 2008 to 2010.

LB, live birth.

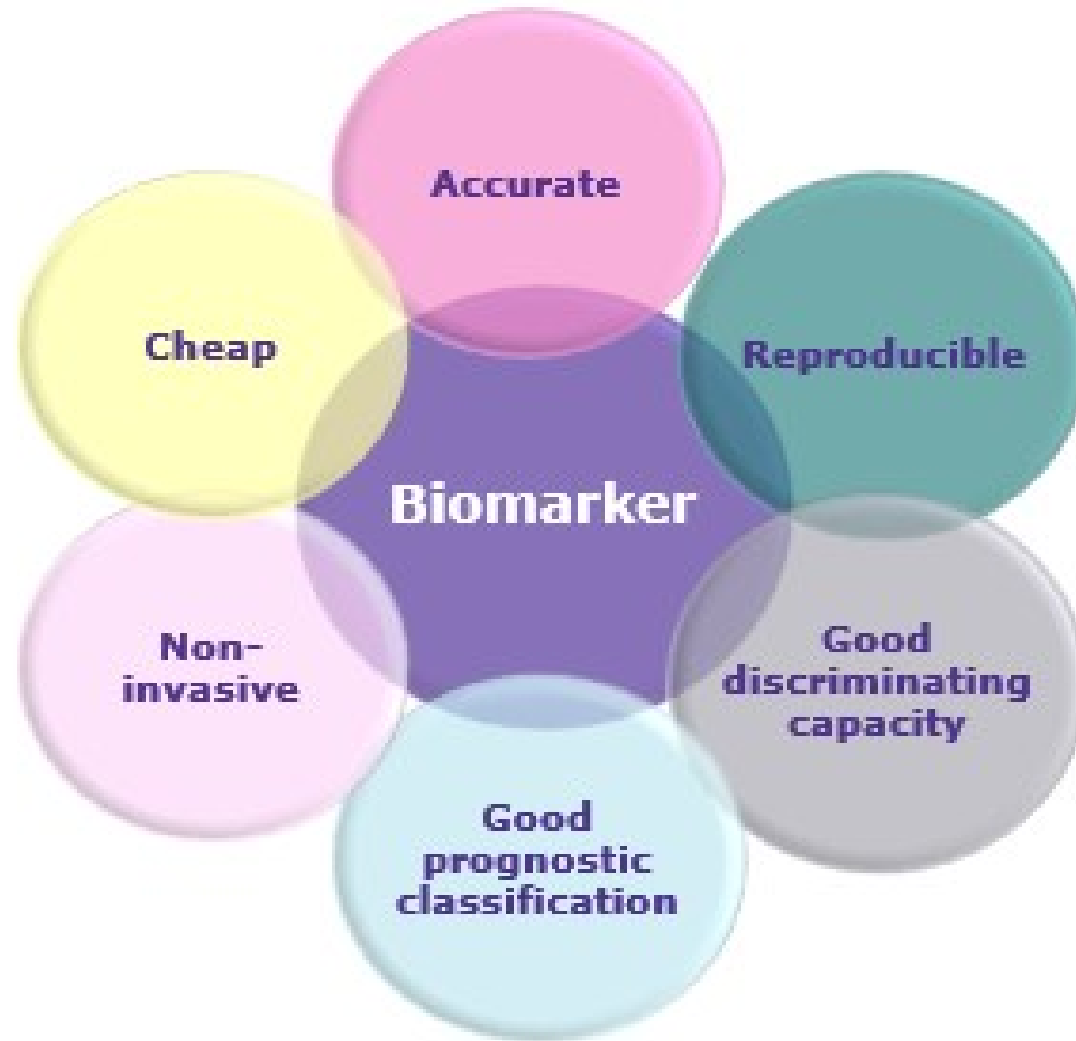
Steward RG et al. Fertil Steril. 2014;101:967-73.

Prediction of ovarian reserve and response...



...increases the efficiency and safety
of ovarian stimulation

What makes a good predictor/biomarker?



Do we have reliable predictors/biomarkers of ovarian reserve and response?

- **Limited value**
 - Basal FSH or LH
 - Basal estradiol
 - Inhibin B
 - Ovarian volume
 - Ovarian vascular flow
- **Most promising**
 - Female age
 - AFC
 - AMH

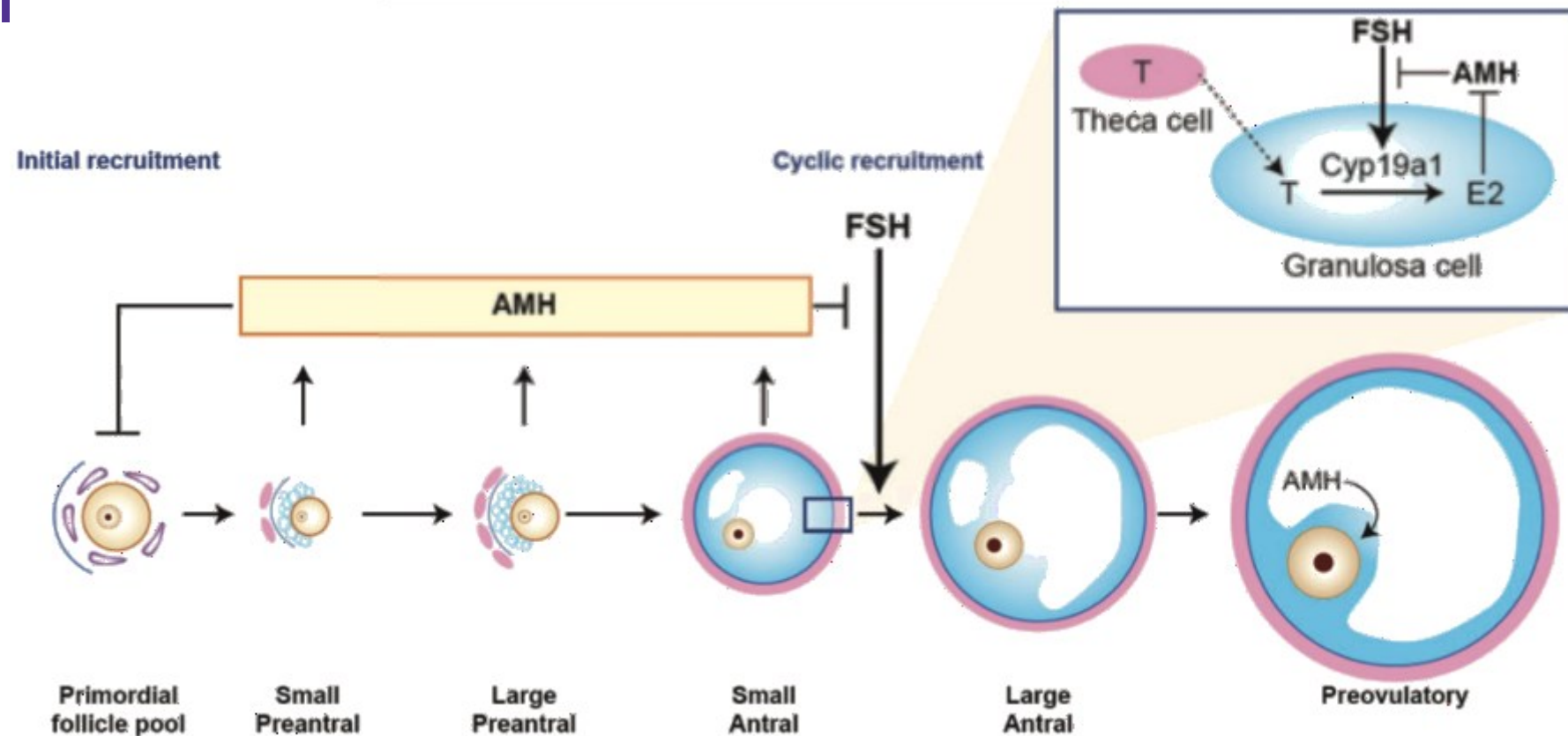


The era of AMH



Welcome to the new era

What is AMH?



Dimeric glycoprotein
Member of the TGF- β superfamily

How is AMH measured? (1)

1990's: First report of measurement of serum AMH with ELISAs



Early 2000's: The first ultrasensitive AMH assay
(IOT assay/Beckman Coulter)



2006: An additional AMH ELISA assay using different epitopes was developed (DSL) –
different result to IOT



2010: **AMH Gen II assay** developed by Beckman Coulter using
the DSL antibodies – calibrated to IOT assay

How is AMH measured? (2)

Table I Anti-Müllerian hormone (AMH) assays characteristics.

Competitor assays	Beckman Coulter AMH Gen II (Kumar <i>et al.</i> , 2010; Beckman Coulter AMH Gen II ELISA package insert, 2013)	Beckman Coulter Immunotech AMH (Immunotech AMH (Beckman Coulter) package insert, 2012)	AnshLabs Ultrasensitive (AnshLabs UltraSensitive AMH/ MIS ELISA package insert, 2014)	AnshLabs picoAMH (AnshLabs picoAMH ELISA package insert, 2014)	Roche Assay Elecsys® AMH (Gassner and Jung, 2014)	Beckman Coulter Access 2 IA AMH (Beckman Coulter, 2014)
Assay type	Manual	Manual	Manual	Manual	Automated	Automated
Imprecision	<8%	<14%	<6%	<6%	1.8–2.0%	2.87–4.34%
Sample type	Serum, plasma	Serum, plasma	Serum, plasma	Serum, plasma	Serum, Li-heparin plasma	Serum, Li-heparin plasma
Minimum sample volume	20 µl	25 µl	50 µl	100 µl	50 µl	20 µl
Incubation time	<3 h	3 h	2.5 h	4.5 h	18 min	39 min
Limit of detection (LoD)	0.08 ng/ml	0.14 ng/ml	0.023 ng/ml	0.0012 ng/ml	0.01 ng/ml	≤0.02 ng/ml
Limit of Quantification (LoQ)	0.16 ng/ml	0.35 ng/ml (Decanter <i>et al.</i> , 2014)	0.06 ng/ml	0.0039 ng/ml	0.03 ng/ml	≤0.08 ng/ml
Measurement range	0.16–22.5 ng/ml	0.42–21.0 ng/ml	0.06–11.6 ng/ml	0.003–0.75 ng/ml	0.01–23.0 ng/ml	0.02–24.0 ng/ml

Results are not directly comparable and have to be interpreted in an assay specific manner

Issues with AMH measurement?

Assay specific

- Manual assays:
 - Substantial variability between labs/lack of standardization
 - Recent field notices regarding complement interference
 - Complement interference seems to be addressed with freezing storage
- Automated assays:
 - Better standardization
 - Recent tests seem to suggest that complement does not interfere with AMH measurement in automated protocols

Issues with AMH measurement?

Inter/intra-individual

Inter-individual factors



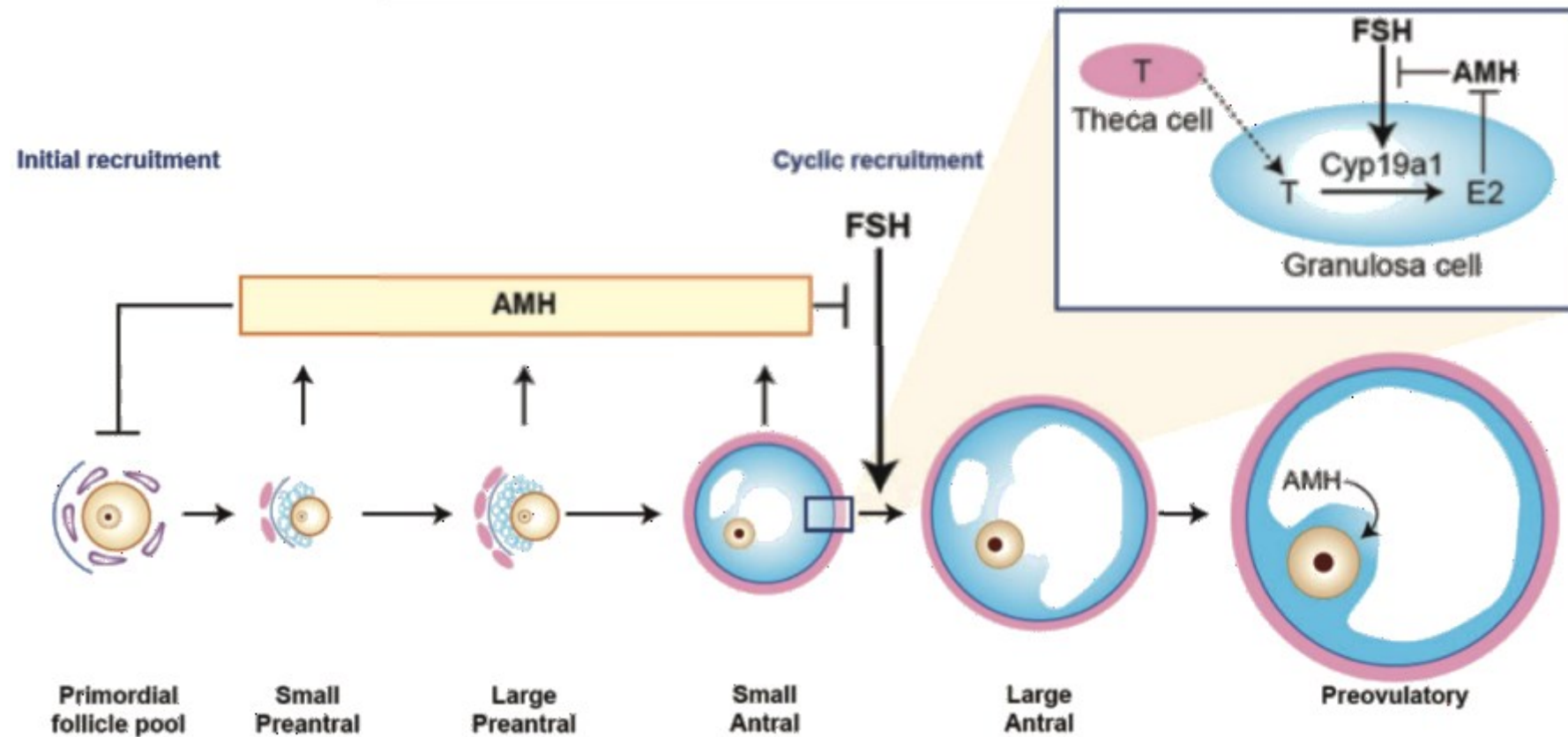
- Age
- Ovarian surgery/ Endometriosis/ Prior gonadotoxic chemotherapy
- Ethnicity (Chinese, Black African, Hispanic and South Asian < Caucasian)
- Smoking
- Pregnancy, GnRH analogues, combined hormonal contraceptives
- Chronic disease/ malignancies

Intra-individual factors



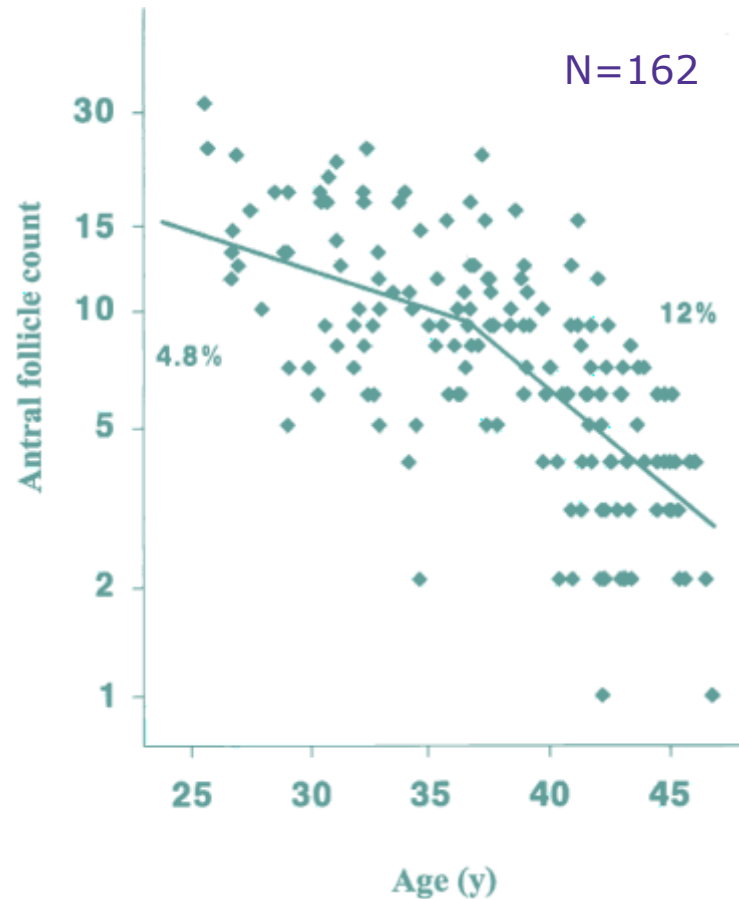
- Minimal cyclic variation during the menstrual cycle (late follicular phase)
- Low variability between menstrual cycles

What is AFC?

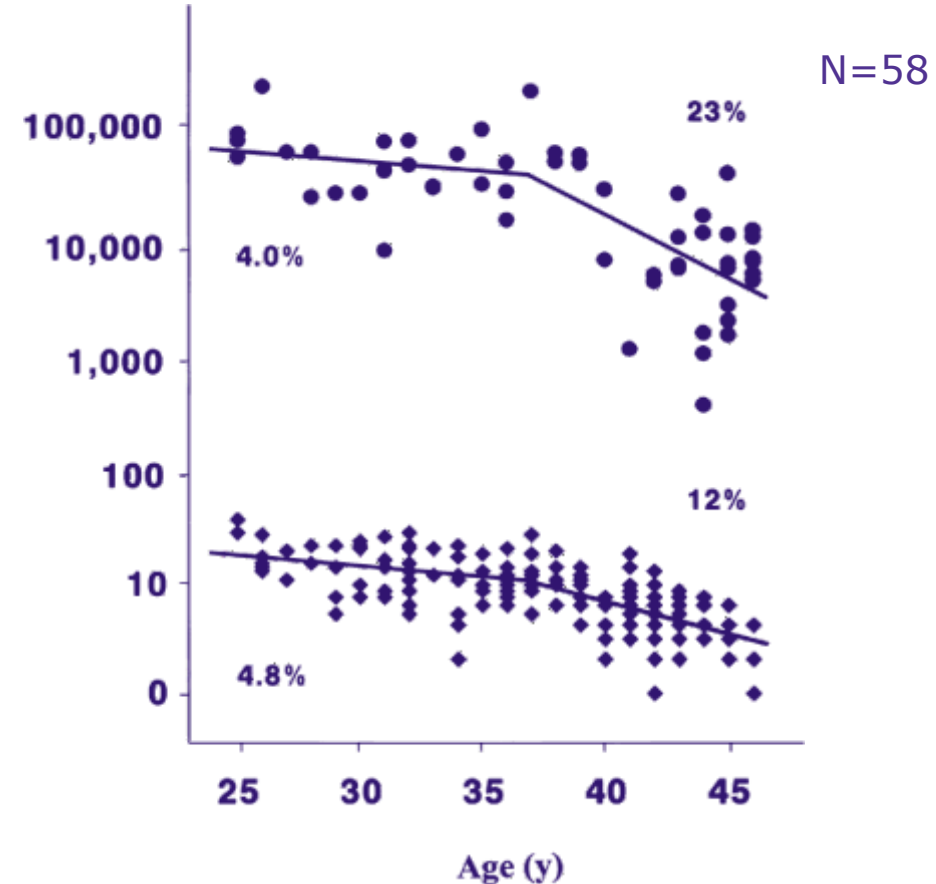


Follicles with an antrum, 2–10 mm in diameter

Is AFC associated with ovarian reserve?

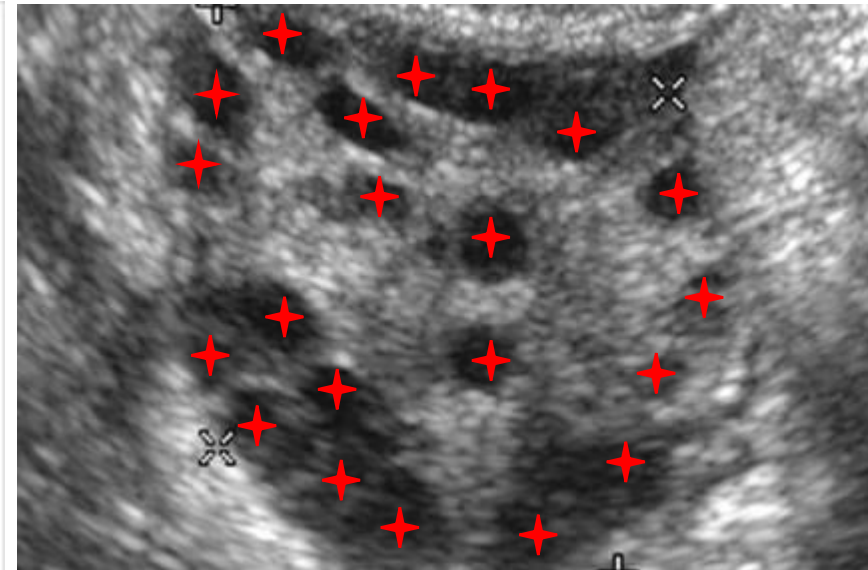
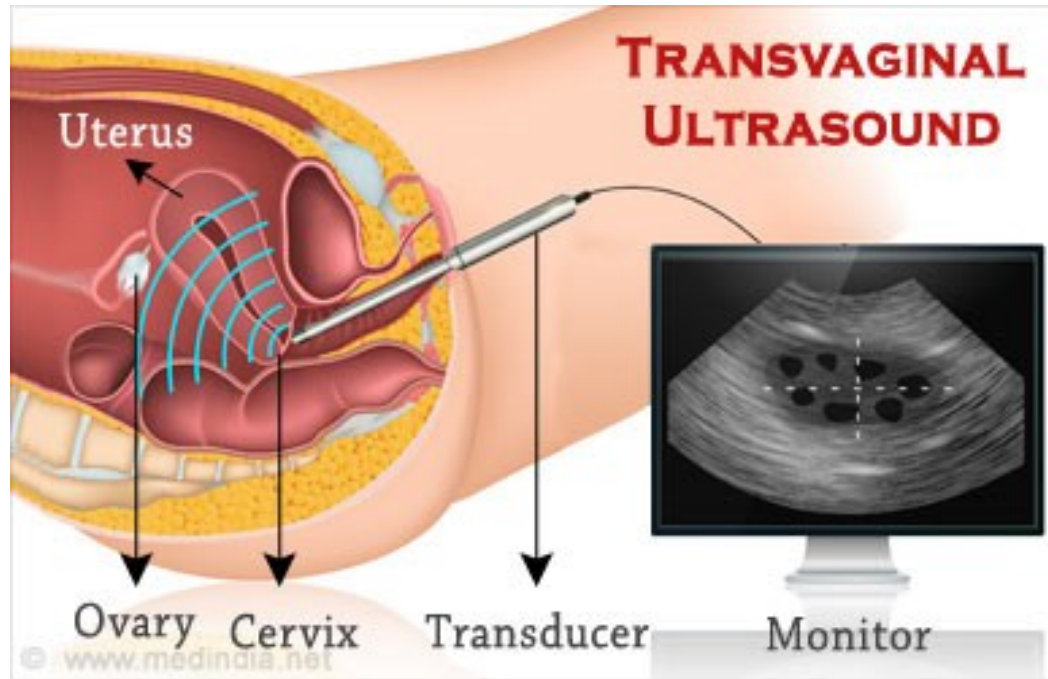


AFC declines with age



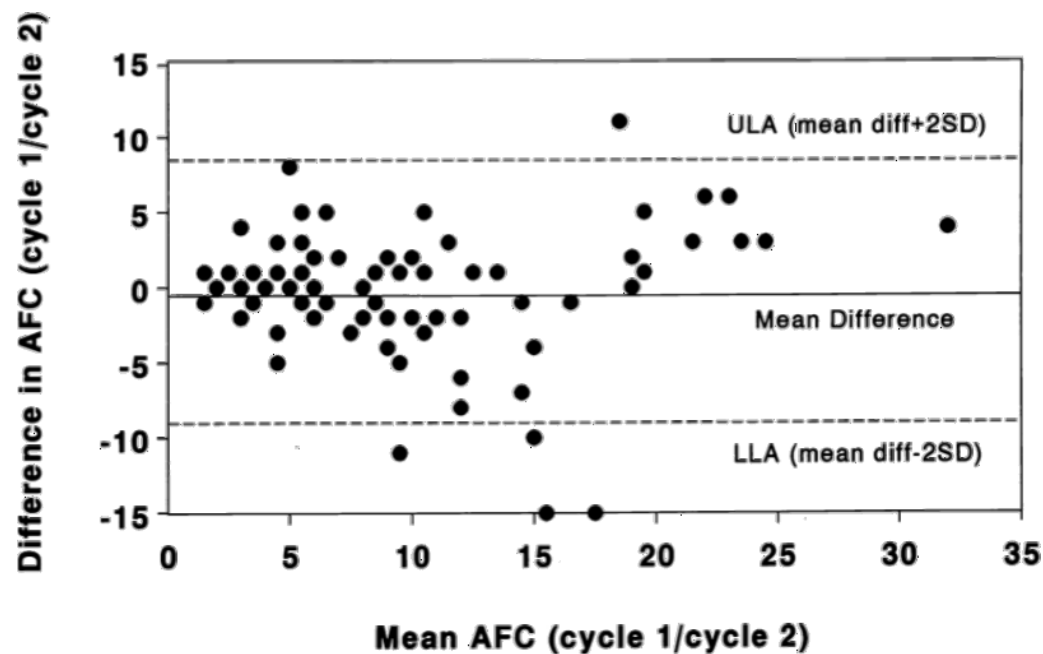
Similar pattern of loss of primordial follicles and AFCs with age

How is AFC measured?

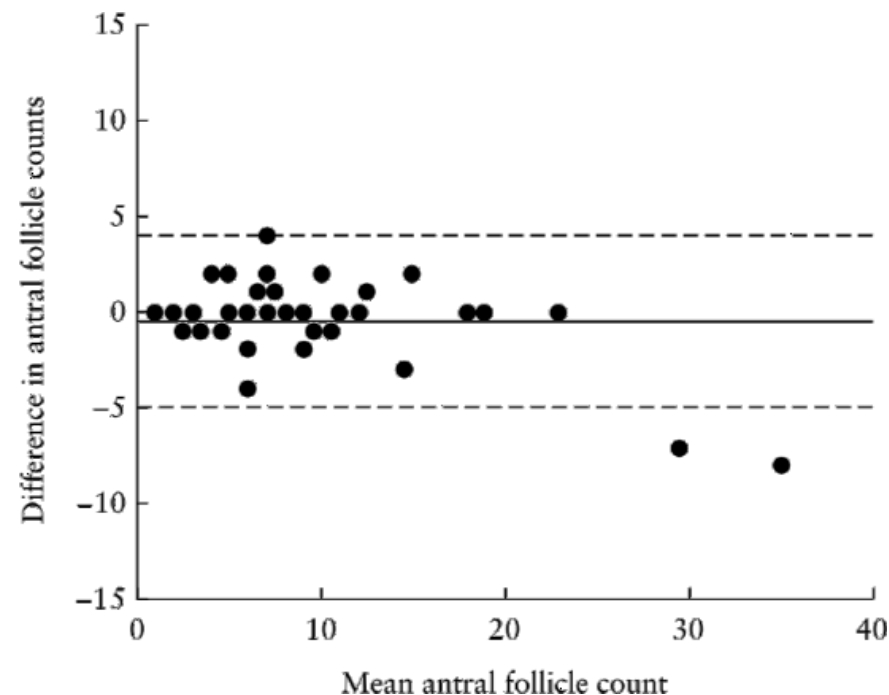


21 AFC

AFC measurement: Inter-cycle & inter-observer variability



Moderate inter-cycle agreement
at the entire range of AFC



Moderate inter-observer variability
increasing with higher AFCs

AFC measurement: Pros and cons

Pros

- Allows the assessment of recruitable follicles for the specific cycle
- Allows the assessment of the potential of each ovary
- Provides information regarding the sizes of follicles
- Can provide information regarding other ovarian pathology (cysts etc.)

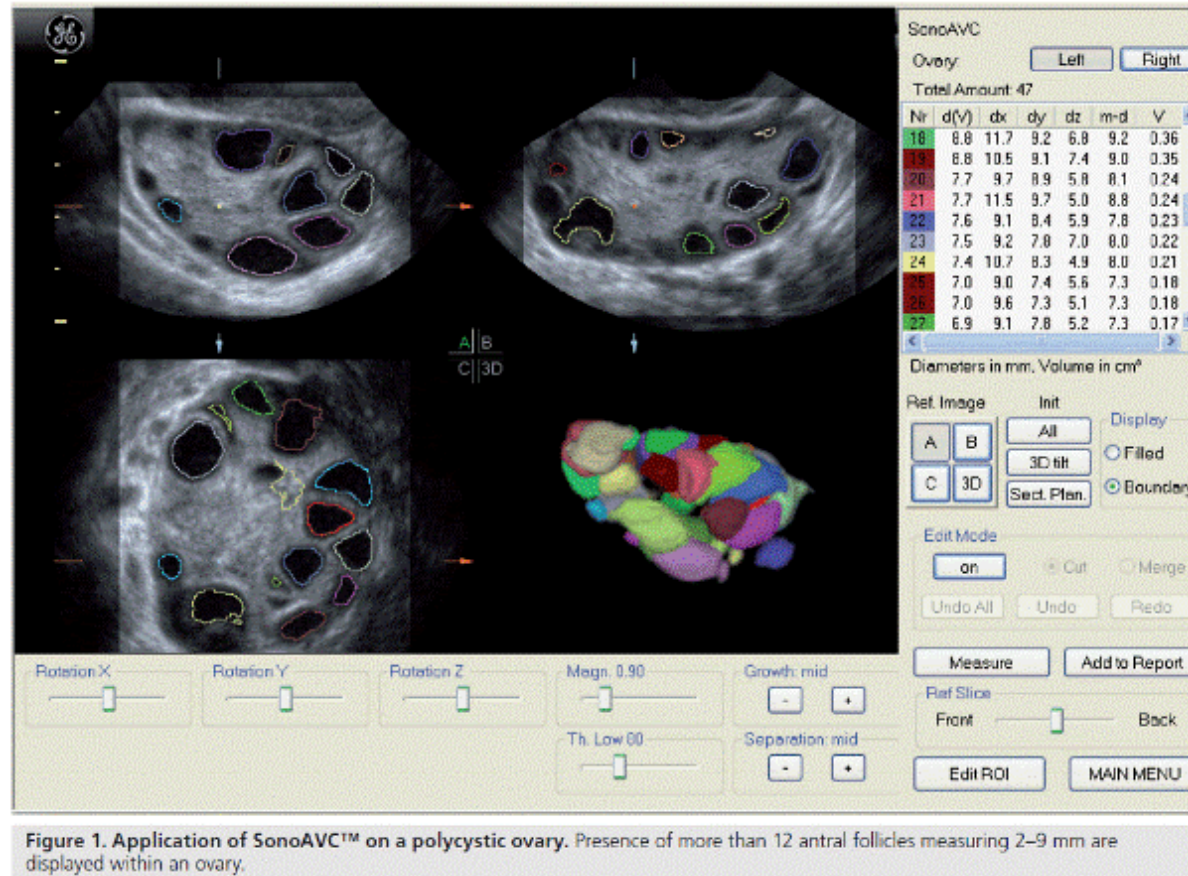
Cons

- Invasive/ TVUS
- Requires expensive equipment
- Performed at a specific time of the cycle
- Inter-operator variability/lack of standardization
- Inter-cycle variability
- May include atretic follicles

Standardizing AFC measurement

- Patients with regular menstrual cycles with no coexisting ovarian pathologies
- Day 2–4 of cycle
- Include all follicles 2–10mm in diameter
- Have appropriately trained personnel
- Use TVUS with a probe of a minimum frequency of 7 MHz
- Systematically sweep each ovary
- Measure the largest follicle in two dimensions
- If that follicle is $\leq 10\text{mm}$ then count all round or oval transonic structures within the ovarian margins – combine the number of follicles in each ovary
- If the largest follicle is $> 10\text{mm}$ measure the largest follicles until a follicle with a mean diameter of $\leq 10\text{mm}$ is identified – measure all follicles and subtract the ones $> 10\text{mm}$

Automated AFC measurement?



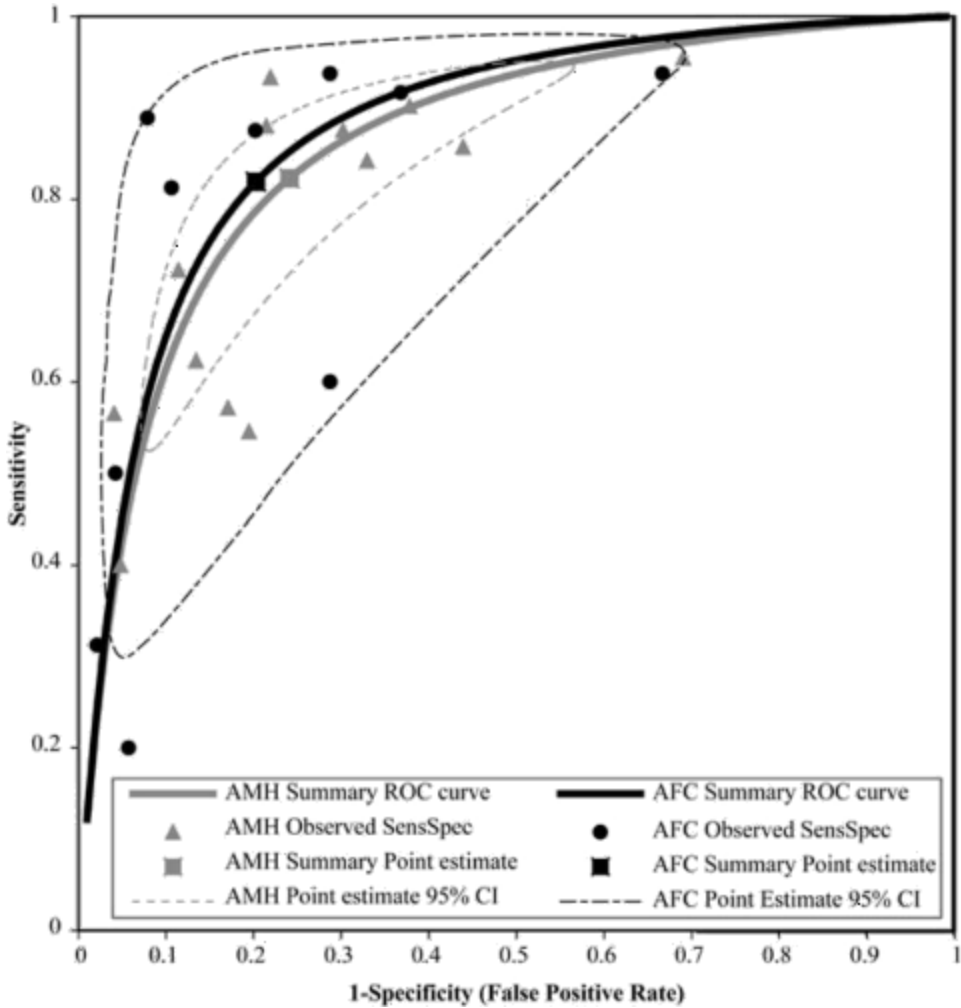
Accurate and reproducible automated system that determines the absolute AFC measurement and volume

AFC vs AMH: clinical data



Which one is best at predicting ovarian response after ovarian stimulation?

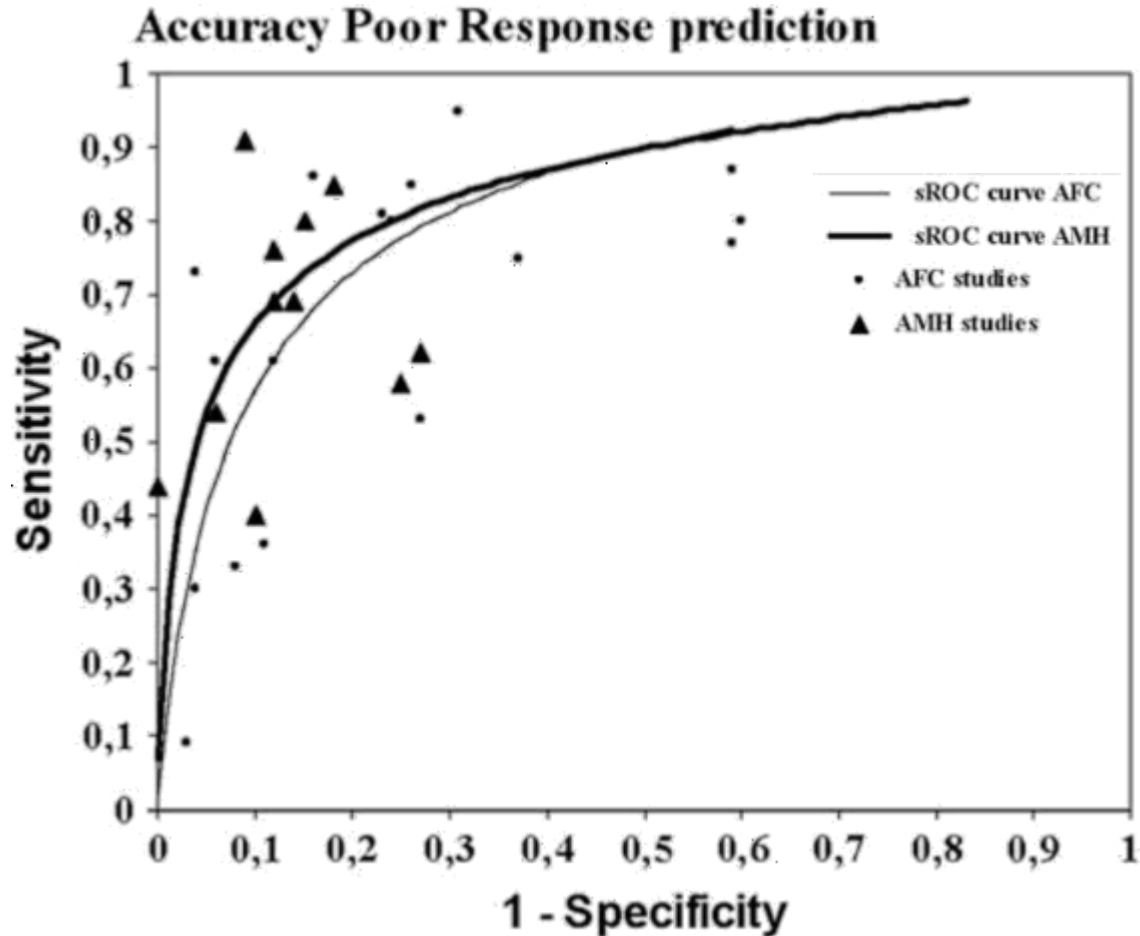
AFC vs AMH: hyper-response



Only AMH 6 studies AMH + AFC 3 studies AFC only 2 studies		
Predictor	Sensitivity	Specificity
AMH	82%	76%
AFC	82%	80%

"Both AMH and AFC are accurate predictors of excessive response to ovarian hyperstimulation. Moreover, both tests appear to have clinical value"

AFC vs AMH: poor response



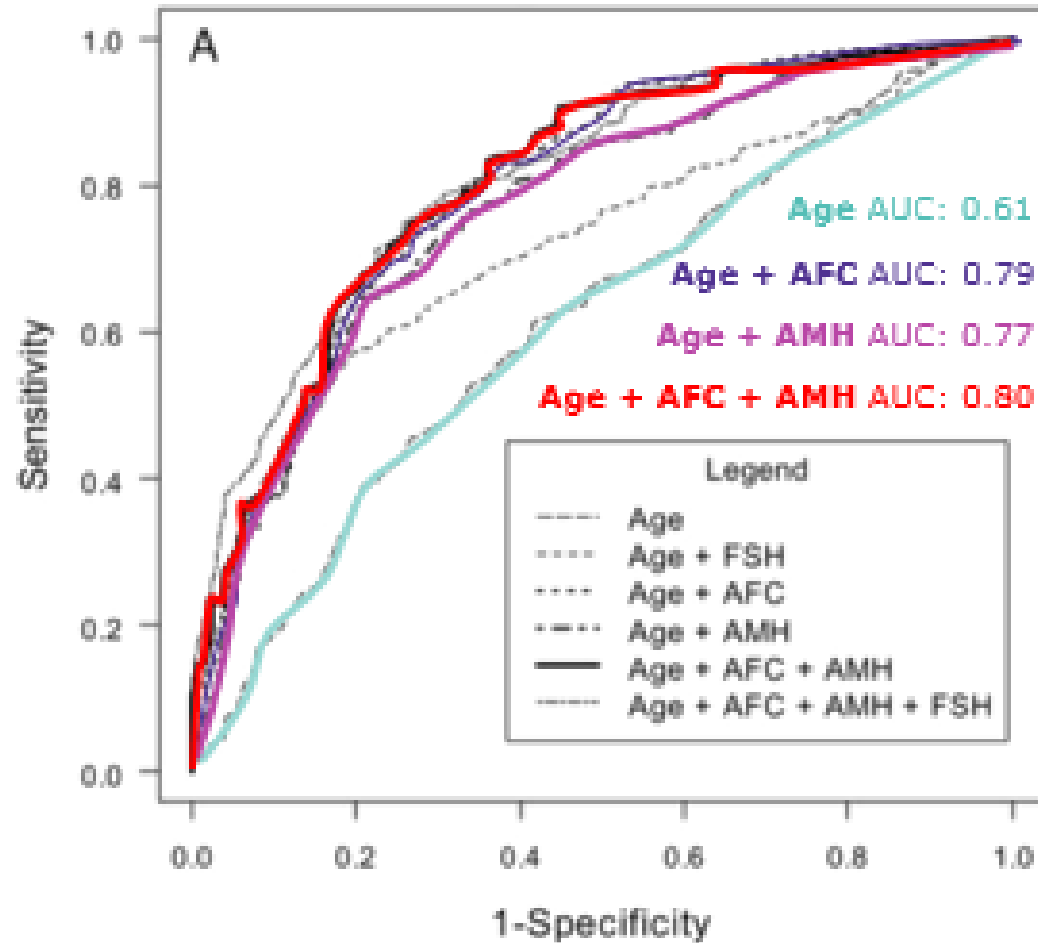
Only AMH
13 studies

AFC only
17 studies

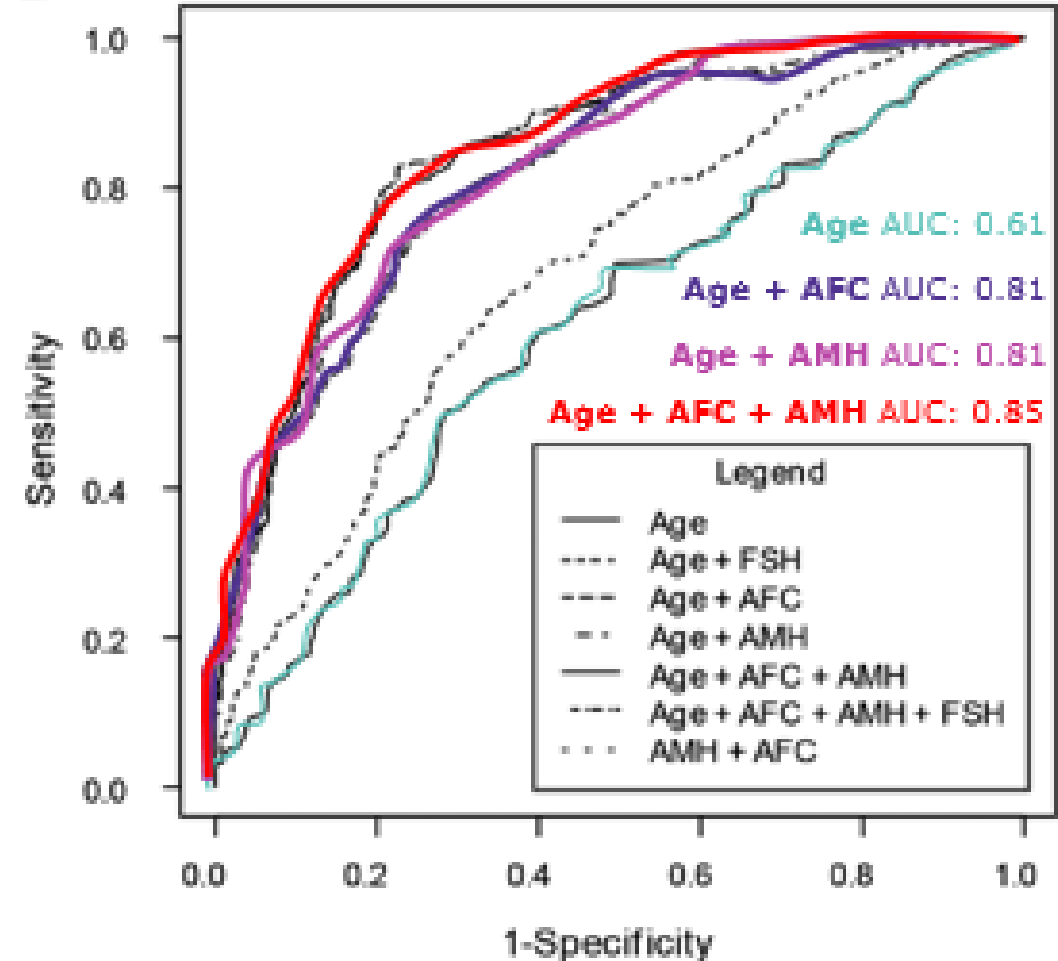
- *"Both AMH and AFC are accurate predictors of poor response to ovarian hyperstimulation"*
- *No significant difference seems to be present between the two tests*
- *Both tests appear to have clinical value*

What about combining ORTs? (1)

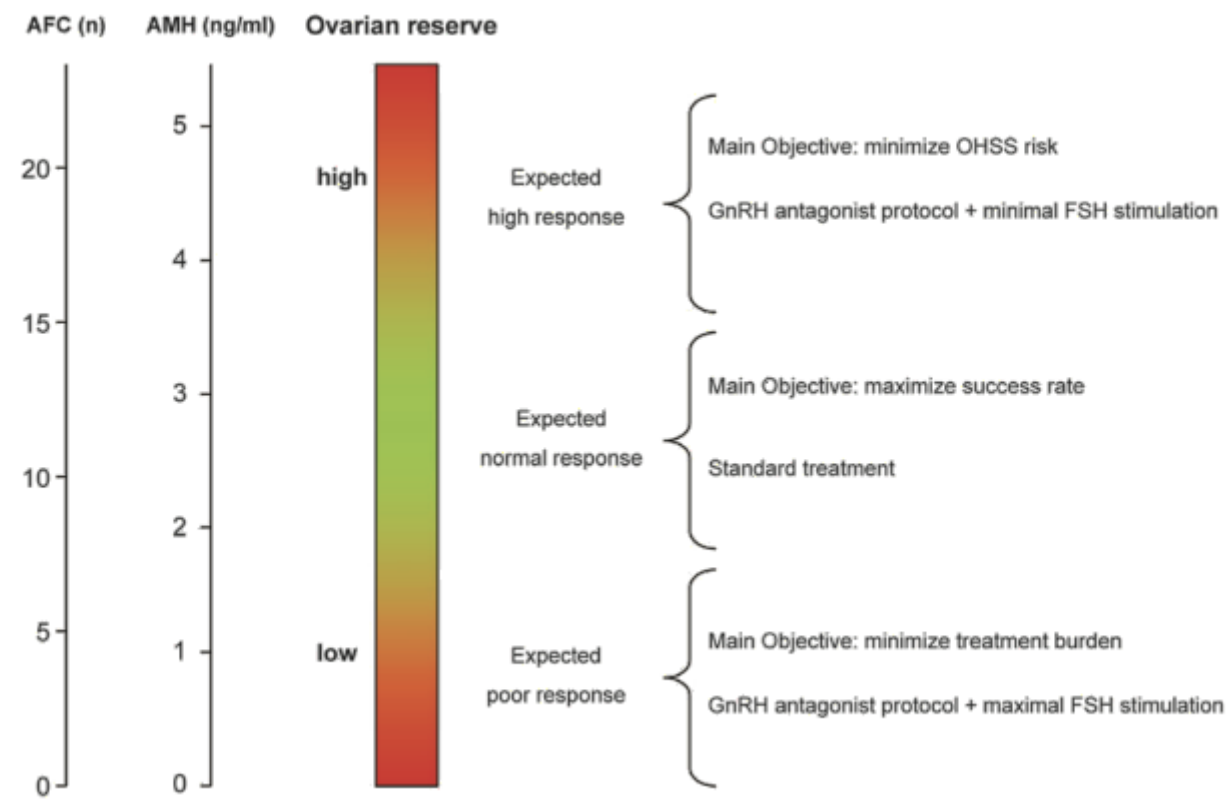
IPD Meta-analysis: 28 studies
Poor response



IPD Meta-analysis: 32 datasets
Hyper-response



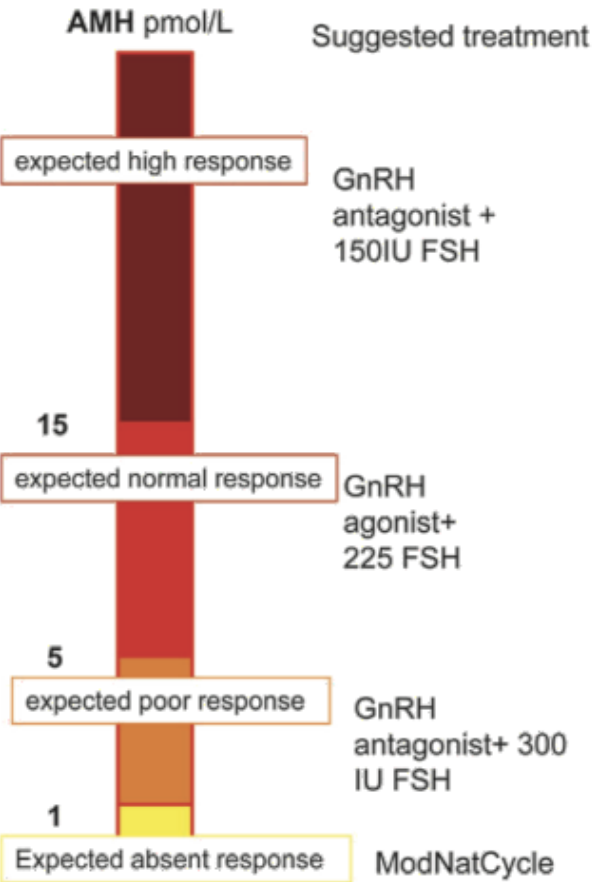
Purpose of ORTs



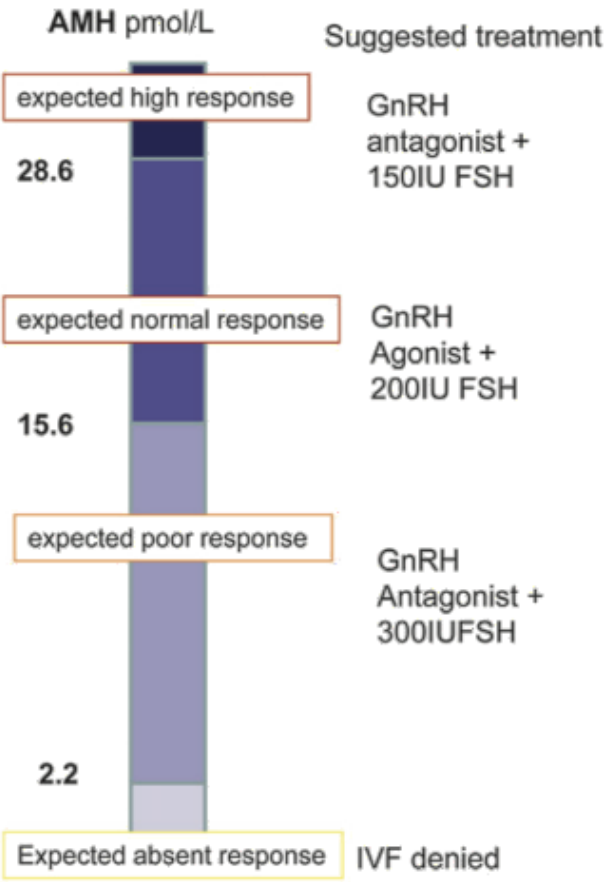
Nomograms

Note: ORT before the first IVF cycle would permit to categorize patients as expected poor-, normal- or hyper-responders. Since there is no evidence of superiority of one approach over another in the treatment of poor responders, the protocol associated with reduced discomfort and treatment burden should be preferred. In hyper-responder patients, one of the most important objectives of medical counselling is to prevent OHSS. Hence the first line protocol would be based on administration of low doses of FSH in a GnRH-antagonist-based scheme

Examples of ORT nomograms



Nelson SM et al. 2009



Yates AP et al. 2011

Yates AP, et al. Hum Reprod. 2011;26:2353–62;
Nelson SM, et al. Hum Reprod. 2009;24:867–75;

figure reproduced from La Marca A and Sunkara SK. Hum Reprod Update. 2014;20:124–40.

Prediction of response using ORT: RCTs? (1)

ORT algorithm vs. standard dose (150 IU)

Study	N	Population	Predictors	Aim	Stimulation protocol
Popovic-Todorovic et al., 2003	262	First IVF/ICSI, bFSH<12.5 IU/L, regular cycles, <39 yo	AFC, ovarian stromal blood flow, serum testosterone, smoking	5–14 oocytes	Long GnRH agonist
Olivennes et al., 2015	200	18–34 yo, Normoovulatory, BMI<30 kg/m ² , bFSH <12 IU/L	bFSH, age, BMI, AFC	COCs	Long GnRH agonist
Oudshoorn et al., 2017	1032*	First IVF/ICSI, AFC<11 or >15, regular cycles, no PCOS, <44 yo	AFC	CLBR	Long GnRH agonist or daily antagonist
Nyboe-Andersen et al., 2017	1329	First IVF/ICSI, bFSH: 1–15 IU/L, BMI: 17.5–32 kg/m ² , regular cycles, <40 yo	AMH (Roche)	OPR	GnRH antagonist fixed Day 6

*Data from two of the direct dose comparison studies conducted in tandem (Oudshoorn 2017 and Van Tilborg 2017) were included as one ORT-algorithm study (Oudshoorn 2017) in this analysis. bFSH, basal follicle-stimulating hormone; BMI, body mass index; CLBR, cumulative live birth rate; ICSI, intracytoplasmic sperm injection; OPR, ongoing pregnancy rate; PCOS, polycystic ovary syndrome; RCT, randomized controlled trial; yo, year's old.

Prediction of response using ORT: RCTs? (2)

ORT algorithm vs. standard dose (150 IU)

Outcome	RCTs/ N	Effect size (95% CI)	P-value	Interpretation
Live birth or ongoing pregnancy	4/2823	OR: 1.04 (0.88-1.23)	0.68	No association
Moderate or severe OHSS	4/2823	OR: 0.58 (0.34-1.00)	0.048	Negative association
Poor response	2/1294	OR:1.16 (0.90-1.50)	0.24	No evidence of an effect
Normal response	3/2623	OR: 1.22 (1.04-1.43)	0.013	Positive association
Hyper-response	2/1294	OR: 0.56 (0.42-0.76)	<0.001	Negative association
Cancellation due to poor response	4/2823	OR: 1.19 (0.89-1.60)	0.25	No evidence of an effect
Cancellation due to hyper response	4/2823	OR: 0.37 (0.24-0.57)	<0.001	Negative association
Total dose of FSH	3/1494	WMD: -157 (-215 to -98)	<0.001	Negative association

AMH vs. AFC based algorithm: RCT data

RCT: 368 patients

AMH VS. AFC

Starting dose of rFSH

AMH (ng/mL)

<0.7: 375 IU

0.7-2.1: 225 IU

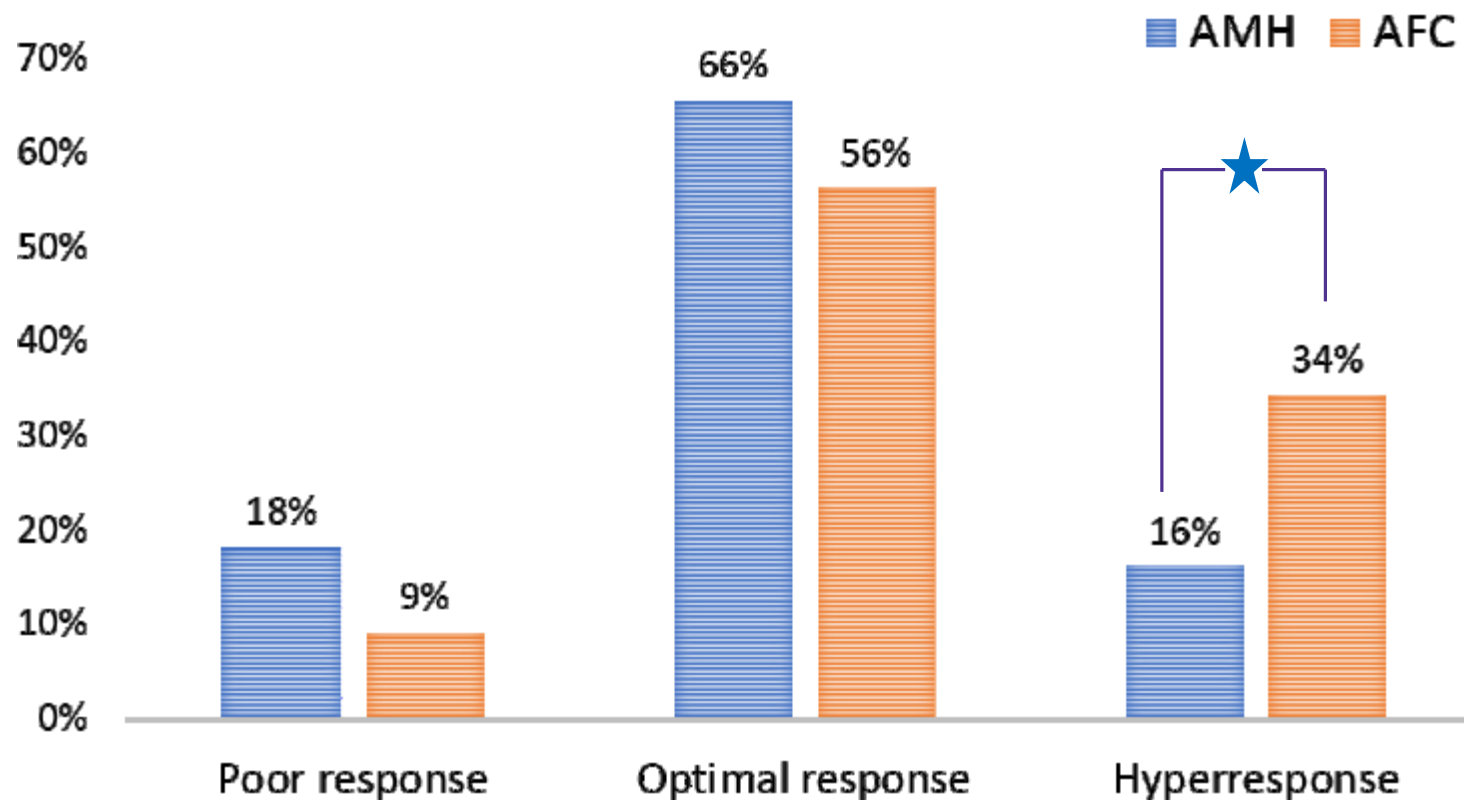
>2.1: 150 IU

AFC

<6: 375 IU

6-15: 225 IU

>15: 150 IU



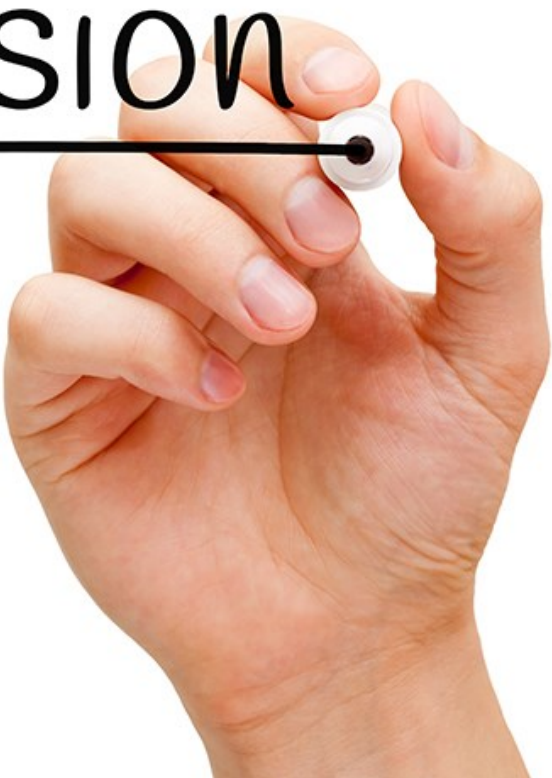
Prediction of response using ORT: genetic markers

- **FMR1 CGG repeat length:** Not associated with ovarian response or no added value in prediction of ovarian response
- **AMH and AMHR2 SNPs:** Conflicting evidence
- **Gonadotrophins and their receptors SNPs:** No strong association except for the p.Asn680Ser which modulates response both to endogenous and exogenous FSH



Morin et al., RBM Online. 2016;32:496–502; Banks et al. Fertil Steril. 2016;105(6):1537–46; Peluso et al. Cell Physiol Biochem. 2015;35:1401–12; Cerra et al. JARG. 2016;33:1085–91; Ricetti et al. Best Pract Res Clin Obstet Gynaecol. 2017;44:15–25.

Conclusion

A close-up of a right hand holding a white marker. The hand is positioned to draw a horizontal line under the word 'Conclusion', which is written in a black, cursive-style font. The line is already partially drawn and ends with a small black dot at the tip of the marker.

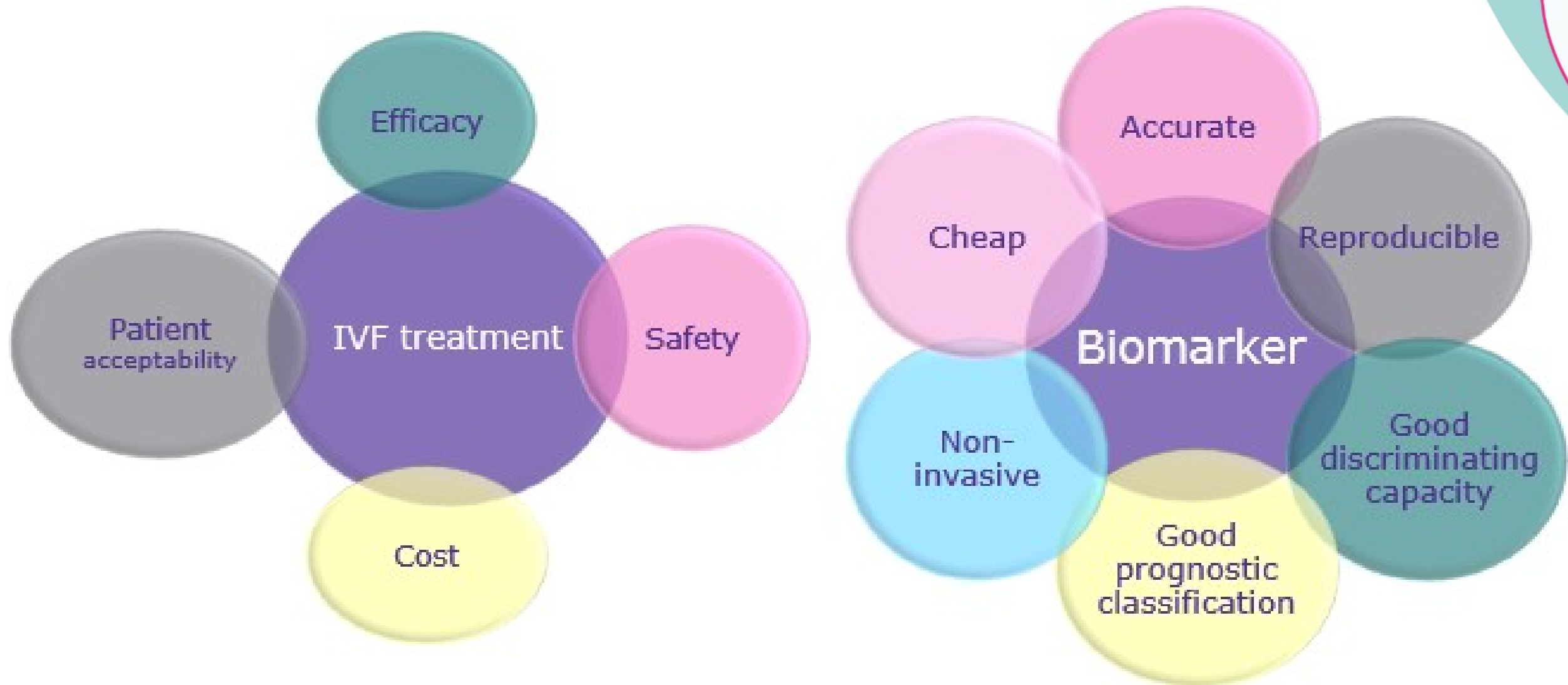
Conclusions (1)

- Out of the many biomarkers that have been evaluated, **female age, AFC and serum AMH** seem to be the strongest predictors of ovarian response
- Both AMH and AFC have pros and cons, and issues with reproducibility; these might be resolved by standardizing AFC measurement and by using automated AMH assays
- AMH and AFC seem to have a similar capacity to predict poor or excessive response and can add to female age in predicting poor or excessive ovarian response

Conclusions (2)

- Using an ORT based algorithm to determine the starting dose is superior to 150 IU of rFSH in terms of OHSS prevention, prediction of normal or high response and cycle cancellation due to hyper-response
- One randomized study suggests that a dosing algorithm based on AMH is superior to one based on AFC when it comes to preventing hyper-response
- Genetic markers of ovarian response are still scantily reported in the literature and in most cases lack validation

Improving ART outcome using biomarkers



IMPROVING THE PREDICTION OF OVARIAN RESERVE AND RESPONSE IN ART USING BIOMARKERS

Christos Venetis

MD, FRANZCOG, MSc (Res), MSc (HCM), PhD

University of New South Wales

Sydney, Australia

c.venetis@unsw.edu.au

