

# Advanced glycation end products alter steroidogenic gene expression by granulosa cells: an effect partially reversible by vitamin D

Z. Merhi <sup>1,\*</sup>, E. Buyuk<sup>2</sup>, and M.J. Cipolla<sup>3</sup>

<sup>1</sup>Department of Obstetrics and Gynecology, Division of Reproductive Biology, New York University School of Medicine, 4 Columbus Circle, Fourth Floor, New York, NY 10019, USA <sup>2</sup>Department of Obstetrics & Gynecology and Womens' Health, Division of Reproductive Endocrinology and Infertility, Montefiore's Institute for Reproductive Medicine and Health, Albert Einstein College of Medicine, 1300 Morris Park Avenue, Bronx, NY 10461, USA <sup>3</sup>Department of Neurological Sciences and Obstetrics, Gynecology and Reproductive Sciences, University of Vermont College of Medicine, 111 Colchester Ave, Burlington, VT 05401, USA

\*Correspondence address. NYU School of Medicine, 4 Columbus Circle, Fourth Floor, New York, NY 10019, USA. Tel: +1-21-22-63-2823; Fax: +1-21-22-63-0290; E-mail: zom00@hotmail.com  [orcid.org/0000-0002-7952-1428](https://orcid.org/0000-0002-7952-1428)

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**STUDY QUESTION:** Does vitamin D attenuate the adverse effects of advanced glycation end products (AGEs) on steroidogenesis by human granulosa cells (GCs)?

**SUMMARY ANSWER:** AGEs alter the expression of genes important in steroidogenesis while 1,25-dihydroxyvitamin D<sub>3</sub> (vit D<sub>3</sub>) *in vitro* attenuates some of the actions of AGEs on steroidogenic gene expression, possibly by downregulating the expression of the pro-inflammatory cell membrane receptor for AGEs (RAGE).

**WHAT IS KNOWN ALREADY:** Vitamin D attenuates the pro-inflammatory effects of AGEs in non-ovarian tissues.

**STUDY DESIGN, SIZE, DURATION:** Women who were undergoing IVF were enrolled. Follicular fluid samples ( $n = 71$ ) were collected and cumulus GCs ( $n = 12$ ) were treated in culture.

**PARTICIPANTS/MATERIALS, SETTING, METHODS:** Follicular fluid levels of the anti-inflammatory soluble RAGE (sRAGE), AGEs and 25-hydroxyvitamin D (25-OHD) were quantified for possible correlations. GCs of each participant were split equally and treated with either media alone (control) or with human glycated albumin (HGA as a precursor for AGEs) with or without vit D<sub>3</sub> after which RT-PCR and immunofluorescence were performed and cell culture media estradiol (E<sub>2</sub>) levels were compared.

**MAIN RESULTS AND THE ROLE OF CHANCE:** In follicular fluid, sRAGE levels were positively correlated with 25-OHD levels. HGA treatment (i) increased CYP11A1 (by 48%), 3 $\beta$ -HSD (by 38%), StAR (by 42%), CYP17A1 (by 30%) and LHR (by 37%) mRNA expression levels ( $P < 0.05$  for all) but did not alter CYP19A1 or FSHR mRNA expression levels; and (ii) increased E<sub>2</sub> release in cell culture media ( $P = 0.02$ ). Vit D<sub>3</sub> treatment (i) downregulated RAGE mRNA expression by 33% and RAGE protein levels by 44% ( $P < 0.05$ ); (ii) inhibited the HGA-induced increase in CYP11A1, StAR, CYP17A1 and LHR mRNA levels, but not the increase in 3 $\beta$ -HSD mRNA levels; and (iii) did not inhibit the HGA-induced E<sub>2</sub> release in cell culture media.

**LIMITATIONS REASONS FOR CAUTION:** This study used luteinized GCs that were collected from women who received gonadotropins thus the results obtained may not fully extrapolate to non-luteinized GCs *in vivo*.

**WIDER IMPLICATIONS OF THE FINDINGS:** This study suggests that there is a relationship between AGEs and their receptors (RAGE and sRAGE) with vitamin D. Understanding the interaction between AGEs and vitamin D in ovarian physiology could lead to a more targeted therapy for the treatment of ovarian dysfunction.

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**Key words:** advanced glycation end products / vitamin D / RAGE / sRAGE / granulosa / follicular fluid / steroidogenesis

## Introduction

Advanced glycation end products (AGEs) are physiologically formed by a non-enzymatic modification of proteins, lipids and nucleic acids by glucose (Monnier and Sell, 2006; Inagi, 2011; Piperi et al., 2012). AGEs constitute a heterogeneous group of compounds of more than 20 members (Monnier and Sell, 2006; Ramasamy et al., 2012). Pentosidine and *N*-carboxymethyl-lysine (CML) are well characterized AGEs and have been used as markers of AGE accumulation in various tissues (Diamanti-Kandarakis et al., 2007a; Merhi, 2014). Diabetes, insulin resistance, aging and oxidative stress accelerate the generation, or decrease the renal clearance, of AGEs (Bohlender et al., 2005; Unbarri et al., 2007; Unoki and Yamagishi, 2008; Tatone and Amicarelli, 2013; Yan et al., 2007). Additionally, AGEs play a role in the pathogenesis of cardiovascular disease (Yan et al., 2009) and are elevated in obesity and polycystic ovary syndrome (PCOS) (Diamanti-Kandarakis et al., 2008, 2005, 2007b). In addition to endogenous AGEs, serum and ovarian levels of AGEs depend on exogenous sources such as unhealthy diet and smoking (Cerami et al., 1997; Goldberg et al., 2004). Contemporary methods of cooking such as precooked fast-food meals and foods high in protein and fat such as fried eggs, meat and cheese dramatically increase serum concentrations of AGEs (Goldberg et al., 2004; Tantalaki et al., 2014), potentially leading to systemic inflammation. In animals, a high-AGE diet can initiate and provoke insulin resistance, an elevation in serum testosterone levels, an increase in ovarian weight and over-expression of the pro-inflammatory cell membrane receptor for AGEs, RAGE, in the ovaries (Diamanti-Kandarakis et al., 2007a).

Advanced glycation results in irreversible cross-linking of proteins, causing loss of protein structure and function (Monnier and Sell, 2006). Once formed, the protein cross-links damage cellular structures through a number of mechanisms (Inagi, 2011; Piperi et al., 2012). One mechanism of damage to cellular structures is the formation of cross-links between key molecules in the basement membrane of the extracellular matrix, e.g. collagen. Another mechanism involves the interaction of AGEs with RAGE, thus inducing intracellular inflammation and apoptosis. The circulating soluble RAGE (sRAGE) is a truncated form of RAGE and is secreted extracellularly and can be detected in the blood and the follicular fluid (Bonetti et al., 2013; Fujii and Nakayama, 2010; Malickova et al., 2010; Merhi et al., 2014b). sRAGE confers an anti-inflammatory role (Basta, 2008) by binding circulating AGEs, thus preventing the adverse intracellular events of the pro-inflammatory AGE-RAGE interaction (Merhi, 2014). Contrary to RAGE, sRAGE is often considered the good receptor (Basta et al., 2006) and circulating sRAGE levels have been used as a predictive biomarker of the development and progression of several diseases such as obesity, insulin resistance, PCOS, diabetes, cardiovascular disease and endothelial dysfunction (Pertynska-Marczewska and Merhi, 2015). As for ovarian physiology, follicular fluid sRAGE levels positively correlate with markers of ovarian reserve (Merhi et al., 2014b) and follicular fluid levels of sRAGE have been shown to be significantly lower in women with PCOS compared to healthy women (Garg et al., 2017).

Vitamin D (vit D) deficiency remains common worldwide (Holick et al., 2011). In the USA, 20–90% of reproductive-aged women are vit D deficient (Holick et al., 2011). Vit D deficiency is associated with obstetrical and reproductive complications including recurrent pregnancy loss, small for gestational age babies, abnormal puberty and infertility (Dicken et al., 2012; Ozkan et al., 2010; Twig et al., 2012). Emerging

data have shown that vit D plays a role in female reproductive physiology (Anifandis et al., 2010; Ozkan et al., 2010; Pal et al., 2016). Vit D binds to vit D receptor (VDR), which is a transcription factor that belongs to the steroid and nuclear hormone receptor superfamily. VDR is expressed in ovarian tissue including granulosa cells (GCs) (Parikh et al., 2010), indicating that VDR plays a role in ovarian function, and it has been shown that vit D affects genes involved in ovarian steroidogenesis and folliculogenesis in human GCs (Merhi et al., 2014a). Several recent data have shown a relationship between vit D and the AGE-RAGE/sRAGE axis (Guo et al., 2016; Irani et al., 2014; Lee et al., 2014). Vit D is considered to have an anti-inflammatory effect and has been shown to exert a protective effect against the inflammatory actions of AGEs; for example, vit D3 and vit D2 down-regulate RAGE expression and attenuate the deposition of AGEs in the blood-brain barrier (Guo et al., 2016), the endothelium (Dreyer et al., 2014) and the heart (Lee et al., 2014). Additionally, there is a significant positive correlation between follicular fluid sRAGE and follicular fluid 25-hydroxyvitamin D (25-OHD) (Garg et al., 2017) and administration of 1,25-dihydroxyvitamin D3 (vit D3) to vit D-deficient women with PCOS causes an increase in serum sRAGE levels (Irani et al., 2014).

Data on the relationship between vit D and AGEs and their receptors are scarce. Since *in-vitro* and animal studies have shown that vit D application could lower the toxic effects of AGEs (Salum et al., 2013; Talmor et al., 2008), we were interested in assessing the relationship between follicular fluid sRAGE and follicular fluid 25-OHD in a relatively large sample of women without PCOS. We also wanted to study the role of AGEs *in vitro*, in the absence or presence of vit D3, on GC function, which represents a long-standing effective tool in characterizing ovarian physiology in humans (Catteau-Jonard et al., 2008; Cloix et al., 2014; Merhi et al., 2014a, 2014b). We hypothesized that there is a positive correlation between follicular fluid sRAGE and follicular fluid 25-OHD and that AGEs affect GC steroidogenic function while vit D3 treatment attenuates the AGE-induced GC alterations.

## Materials and Methods

### Subjects

Among 76 women who were offered participation, 71 infertile women aged between 26 and 42 years old undergoing fresh IVF cycles agreed to participate in the study and were prospectively enrolled. Inclusion criteria included women with normal ovarian reserve defined as Day 3 FSH <10 mIU/mL and Day 3 estradiol (E2) <294 pmol/L. Women with PCOS as defined by the Rotterdam criteria (Franks, 2006) were excluded from the study. All patients gave informed consent and the study was approved by the Institutional Review Board (IRB# M13-062 and i6902 from University of Vermont and NYU School of Medicine; respectively).

Participants underwent ovarian stimulation with a combination of gonadotropins (Follistim, Merck, Whitehouse Station, NJ; Gonal-F, EMD-Serono, Rockland, MA; Menopur and Bravelle, Ferring, Parsippany, NJ) using either a long agonist (Lupron, AbbVie, North Chicago, IL) or antagonist (Ganirelix acetate, Merck, Whitehouse Station, NJ, or Cetorelix acetate, EMD-Serono, Rockland, MA) protocol. When two or more follicles reached a diameter of  $\geq 17$  mm, hCG (ovidrel, EMD-Serono, Rockland, MA; or Novarel, Ferring, Parsippany, NJ) was administered for oocyte maturation, followed by transvaginal ultrasound-guided oocyte retrieval 34–36 h later. All the participants were fasting from the night before the day of the oocyte retrieval.

## Follicular fluid collection for measurement of sRAGE, AGEs (pentosidine and CML), 25-OHD, vit D binding protein, total and free testosterone, sex hormone binding globulin, insulin and glucose

Follicle size was estimated immediately at the time of oocyte retrieval by ultrasound. Follicular fluid from the first large (>14 mm) aspirated follicle was used for protein measurement in order to avoid blood contamination. Follicular fluid was then centrifuged at 5000g for 5 min at 20°C to pellet the cells and debris, and the supernatant was stored at -80°C for protein analysis.

sRAGE was measured by ELISA using reagents from R&D Systems (Human RAGE Immunoassay kit; Minneapolis, MN). The assay sensitivity was 0.1 nmol/L and the interassay coefficient of variation (CV) was 8.3, 8.2 and 6.6% at concentrations of 14.8, 41.2 and 82.1 nmol/L, respectively. The two AGEs, CML and pentosidine, were quantified. CML was quantified by a competitive enzyme immunoassay using reagents (CML Competitive ELISA Kit) from Cell Biolabs, Inc. (San Diego, CA). The assay sensitivity was 10.7 μmol/L and the interassay CV was <10%. Pentosidine was measured by ELISA using reagents from Antibody Research (St. Charles, MO). The assay sensitivity was 2 nmol/L and the interassay CV was <10%.

25-OHD was measured by radioimmunoassay (RIA) using reagents (25-OHD <sup>125</sup>I RIA Kit) from DiaSorin (Stillwater, MN). The RIA of 25-OHD involved a preceding extraction step with acetonitrile. The assay sensitivity was 3.7 nmol/L and the interassay CV was 9.4, 8.2 and 9.1% at concentrations of 21.5, 56.7 and 82.4 nmol/L, respectively. Vit D binding protein (VDBP) was quantified by ELISA using reagents (Human Vit D BP Immunoassay kit) from R&D Systems (Minneapolis, MN). The assay sensitivity was 12.3 nmol/L, and the interassay CV was 5.1, 6.0 and 7.4% at concentrations of 999.4, 1983.8 and 3098.6 nmol/L, respectively.

Testosterone was measured by RIA with preceding organic solvent extraction and Celite column partition chromatography. It was eluted off the column with 40% toluene in isooctane. The assay sensitivity was 0.05 nmol/L and the interassay CV was 8, 12 and 12% at 0.45, 1 and 3.3 nmol/L, respectively. Free and bioavailable (non-SHBG-bound) testosterone were calculated using the measured total testosterone levels and sex hormone binding globulin (SHBG) concentrations as well as an average assumed concentration for albumin (Sodergard et al., 1982; Vermeulen et al., 1999)—this method has been shown to have high validity (Rinaldi et al., 2002). SHBG was measured by direct immunoassay on the Immulite analyzer (Siemens Healthcare Diagnostics, Deerfield, TX). The assay sensitivity was 1 nmol/L and the interassay CV was 9.1% at 69 nmol/L.

Insulin was measured by a solid-phase chemiluminescent immunometric assay using the Immulite 2000 analyzer (Siemens Healthcare Diagnostics, Deerfield, IL). The assay sensitivity was 2 μU/mL. The interassay CV was 4.2 and 2.9% at 10.0 and 47.8 μU/mL, respectively. Glucose was measured by a standard procedure using the Vitros Chemistry System.

## Cumulus GC collection and culture for RNA extraction, reverse transcription and RT-PCR for steroid gene expression

Cumulus GCs of 4 out of the 71 participants were collected. After identification of the cumulus–oocyte complex in the aspirate, cumulus GCs were mechanically collected by cutting the cumulus layer from each oocyte. Cumulus GCs collected from each participant were pooled and then suspended in phosphate buffered saline (PBS), centrifuged for 5 min at 325g and re-suspended in fresh PBS at 4°C, as we previously described (Merhi et al., 2014a). Cells were briefly treated with 0.614 U/mL hyaluronidase

(Sigma, MO), followed by 2× wash with PBS and centrifugation. The pellet of GCs of each participant was washed with fresh medium (DMEM-F12 and 1% fetal bovine serum, FBS). Since naturally occurring glycosylated albumin presents a sensitive glycemic indicator (Stensen et al., 2014), commercially available human glycosylated albumin (HGA) was used as a precursor for AGEs (Diamanti-Kandarakis et al., 2016) throughout the study.

GCs of each participant were split into three equal groups and then cultured in 24 well-culture plates (pretreated with poly-L-lysine for 5 min). All cells were plated 2–4 h after the participants underwent oocyte collection. From each participant, one-third of GCs was treated with media alone (control), another third of GCs was treated with HGA (0.4 mg/mL, Sigma Aldrich St. Louis, MO), and the last third of cells was treated with HGA (0.4 mg/mL) and vit D3 (100 nM, Sigma Aldrich St. Louis, MO) together. The treatment was for 48 h after which RT-PCR was performed. Since vit D3 was prepared with ethanol, ethanol was used as a vehicle control thus it was added to the all control groups of cells in all experiments. The doses of vit D3 (Merhi et al., 2014a) and HGA (Diamanti-Kandarakis et al., 2016; Stensen et al., 2014; Xu et al., 2014) used were chosen based on previous studies related to their effects on human GC function. Plates were incubated under a humidified atmosphere of 95% air and 5% CO<sub>2</sub> at 37°C.

RNA was isolated using Trizol reagent (Invitrogen, Carlsbad, CA) and chloroform extraction according to the manufacturer's instructions. RNA quality was assessed by a Nanodrop Spectrophotometer and Agilent Bioanalyzer (Santa Clara, CA). Only samples with a minimum concentration of 10 ng/μL and with an optical density (OD) 260/280 ratio of 1.8–2.0 were used for evaluation of RAGE, CYP19A1 (aromatase), 17α-hydroxylase/17,20 lyase (CYP17A1), P450 side-chain cleavage enzyme (CYP11A1), steroidogenic acute regulatory protein (StAR), 3 beta-hydroxysteroid dehydrogenase (3β-HSD), FSH receptor (FSHR) and LH receptor (LHR) mRNA expression levels. Although CYP17A1 is usually suppressed in non-luteinized GCs (Patel et al., 2009), luteinized GCs do express CYP17A1 (Moran et al., 2000). RT-PCR kinetics was achieved by using the SYBR Green I chemistry as we described elsewhere (Merhi et al., 2013). The primers used (Table I) were synthesized by Fisher (Pittsburg, PA). Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) primers were used as a loading control and the levels of mRNA for each gene relative to GAPDH was calculated using the 2<sup>-ΔΔCT</sup> method (Livak and Schmittgen, 2001).

## Immunofluorescence for RAGE in luteinized GCs

Cumulus GCs obtained from 4 out of the 71 participants were used for immunofluorescence. GCs of each participant were split into two equal groups and then cultured on eight-well culture slides (Permanox® slide, 0.8 cm<sup>2</sup>/well, sterile, 96/cs Nunc® Lab-Tek® Chamber Slide™ system; Sigma, MO). From each participant, one half of GCs were treated with media alone (control) and the other half of GCs were treated with vit D3 (100 nM) for 24 h. GCs were fixed in 3.7% formaldehyde and permeabilized in 0.2% Triton X-100 for 10 min at room temperature, and nonspecific binding was inhibited by blocking with 1% bovine serum albumin (BSA) in PBS for 1 h. Fixed GCs were incubated with primary antibody for RAGE (1:100 dilution; Cell Signaling Technology Inc., Danvers, MA) overnight at 4°C, washed and then incubated in secondary antibody (1:200 dilution; Alexa Fluor 488 goat anti-rabbit IgG; Invitrogen) for 1–2 h at room temperature. Nuclear staining was performed using DAPI mounting media (Vector Labs, Inc., Burlingame CA) and observed on a Zeiss 510 META Laser Scanning Confocal microscope. Cellular images were captured using the Plane-NEOFLUAR 25× Immersion objective lens. Using MetaMorph software analysis, a threshold of 62 (0–255 scale of intensity) was used to exclude any background produced by DAPI. RAGE intensity on individual cells was performed by subtracting the nuclear staining from the cell surface staining.

## Measurement of E2 concentration in cell culture media

Cumulus GCs obtained from 4 out of the 71 participants were used for measuring E2. GCs of each participant were split into three equal groups and then cultured in 24 well-culture plates (pretreated with poly-L-lysine for 5 min) in media for 24 h after which the culture media was collected for E2 concentrations (baseline levels). Thus baseline E2 levels reflect 24-h accumulation of E2 in the culture media. From each participant, one-third of GCs were treated with media alone (control), another third of GCs were treated with HGA (0.4 mg/mL), and the last third of GCs were treated with HGA (0.4 mg/mL) and vit D3 (100 nM) together for another 48 h after which culture media was collected for E2 concentrations (48 h levels). Cell culture media E2 levels were quantified by RIA. Prior to the RIA, E2 was extracted with ethylacetate:hexane (3:2) and was then separated by Celite column partition chromatography, using ethylene glycol as the stationary phase. E2 was eluted with 40% ethylacetate in isooctane. The assay sensitivity was 7.3 pmol/L and the interassay CV ranged from 9 to 14%.

## Statistics

In order to determine the normality of the data, sktest in STATA 10.0 program that assesses both the skewness and kurtosis of the data simultaneously was used. Comparisons were performed using paired, unpaired *t*-test or ANOVA with a post hoc Bonferroni test for multiple comparisons if the data were normally distributed and Wilcoxon matched-pairs signed rank test, Mann–Whitney *U* test or Kruskal–Wallis test if the data were not normally distributed. Demographic and clinical data of the 71 participants were expressed as mean  $\pm$  SD. Pearson correlation analyses between follicular fluid sRAGE versus 25-OHD and between follicular fluid sRAGE versus other relevant markers (VDBP, pentosidine, CML, total and free testosterone, SHBG, insulin and glucose) were performed.

For cell culture experiments, comparisons were performed using multi-level (hierarchical) analysis to account for cells in different conditions being from the same participant. This approach adjusts standard errors for clustering of cells within participant so the type I error rate was not inflated.

**Table I Primers used for RT-PCR in the study.**

Gene	Sequence primers (5'–3')
CYP19A1	Forward: GAC GGA AGG TCC TGT GCT C Reverse: GGG GGC AAT TTA GAG TCC ACA
CYP17A1	Forward: TGGGCACCAAGACTACAGTG Reverse: CAGAGTCAGCGAAGCGGATA
CYP11A1	Forward: TGGGTCGCCTATCACCAGTA Reverse: TGCAGGACACTGACGAAGTC
StAR	Forward: AGGACGAAGAACCACCCTTG Reverse: CATCACAGCCTGTTGCCTCA
3 $\beta$ -HSD	Forward: GCTTCTGGGTCAGAGGATCG Reverse: CTGGCAGGCTCTTTTCAGGA
FSHR	Forward: CCA GAA CCT TCC CAA CCT TCA Reverse: TTT CAA AGC TCA GCC CCA CG
LHR	Forward: GTTGACTTACCCAGCCACT AGTCCCAGCCACTCAGTTCA
RAGE	Forward: AGCCTCCCCCTCAAATCCACT Reverse: CAGCCCAGACCCATCCACAG
GAPDH	Forward: ACC CAC TCC TCC ACC TTT GA Reverse: TGT TGC TGT AGC CAA ATT CGT T

Condition was the fixed effect in the model and participant was the random effect. For RT-PCR and immunofluorescence, a sample size of  $n = 4$  has been shown to be adequate to produce statistical difference with 80% power and two-tailed  $\alpha$  error of 0.05 (Merhi *et al.*, 2015). RT-PCR results were expressed as relative number of copies  $\pm$  SEM in cell culture experiments with controls being set at 1. Immunofluorescence was determined by taking the mean densitometry values of each treated cell group. All statistical procedures, except testing for normality of the data, were run on GraphPad Prism 7.  $P \leq 0.05$  was considered statistically significant.

## Results

### Correlation between follicular fluid sRAGE versus 25-OHD and other relevant markers

Table II shows the demographics and clinical characteristics of all the participants including follicular fluid levels of sRAGE, 25-OHD, VDBP, pentosidine, CML, total and free testosterone, SHBG, insulin and glucose. As seen in Table III, follicular fluid sRAGE levels positively correlated with 25-OHD, pentosidine, CML and SHBG. There was a negative correlation between follicular fluid sRAGE levels and follicular fluid insulin and glucose levels. No correlation was found between follicular fluid sRAGE levels and follicular fluid VDBP, total testosterone or free testosterone levels.

### Vit D3 treatment *in vitro* downregulates RAGE mRNA and protein expression levels in human luteinized GCs

In order to evaluate whether vit D3 alters RAGE expression in human GCs, we treated human GCs with vit D3 after which RT-PCR for RAGE mRNA and immunofluorescence for RAGE protein were

**Table II Demographics and clinical characteristics of all the 71 participants.**

	Mean $\pm$ SD
Age (years)	33.4 $\pm$ 2.7
BMI (kg/m <sup>2</sup> )	25.8 $\pm$ 2.8
Serum Day 3 FSH (mIU/mL)	6.9 $\pm$ 2.3
Serum Day 3 E2 (pmol/L)	171.1 $\pm$ 40.8
Antral follicle count by ultrasound	15.9 $\pm$ 4.9
Number of days of stimulation	9.8 $\pm$ 3.3
Total gonadotropin IUs per cycle	3481.5 $\pm$ 956.5
Serum peak E2 level (pmol/L) on day of hCG	6689.2 $\pm$ 2892.3
Number of oocytes retrieved	10.3 $\pm$ 4.9
Follicular fluid sRAGE (nmol/L)	155.7 $\pm$ 62.3
Follicular fluid 25-hydroxyvitamin D (nmol/L)	95.1 $\pm$ 26
Follicular fluid VDBP (nmol/L)	4553.4 $\pm$ 1481.3
Follicular fluid pentosidine (nmol/L)	20.3 $\pm$ 18.6
Follicular fluid CML ( $\mu$ mol/L)	0.34 $\pm$ 0.1
Follicular fluid total testosterone (nmol/L)	9.7 $\pm$ 18.9
Follicular fluid free testosterone (nmol/L)	1.2 $\pm$ 2.5
Follicular fluid SHBG (nmol/L)	1297.9 $\pm$ 466.3
Follicular fluid insulin ( $\mu$ IU/mL)	5.9 $\pm$ 6.9
Follicular fluid glucose (mmol/L)	2.92 $\pm$ 0.82

E2, estradiol; sRAGE, soluble receptor for advanced glycation end products; CML, N-(carboxymethyl) lysine; SHBG, sex hormone binding globulin.

**Table III** Correlation between follicular fluid sRAGE versus 25-hydroxyvitamin D (25-OHD) and other relevant markers.

n = 71 participants	Variable	R-value	P-value
sRAGE			
	25-OHD	0.27	0.02
	VDBP	0.02	0.8
	Pentosidine	0.24	0.04
	CML	0.32	0.006
	SHBG	0.37	0.001
	Insulin	-0.28	0.02
	Glucose	-0.39	0.0007
	Total testosterone	0.0002	0.9
	Free testosterone	-0.04	0.7

sRAGE, soluble receptor for advanced glycation end products; VDBP, vitamin D-binding protein; CML, N-(carboxymethyl) lysine; SHBG, sex hormone binding globulin.

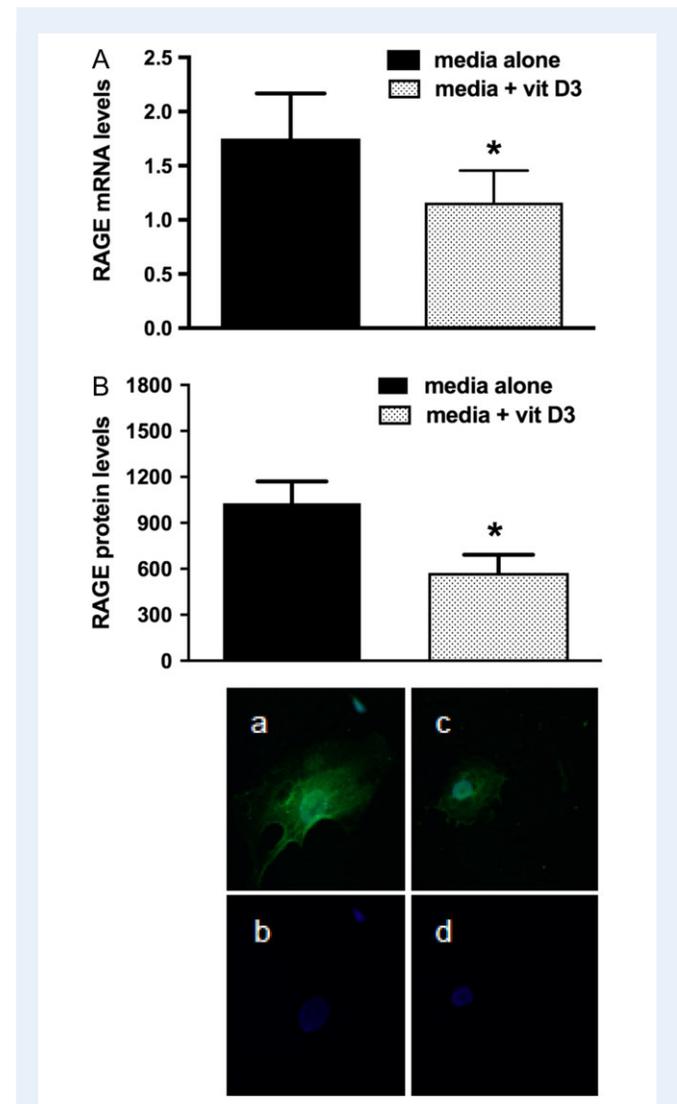
performed. Compared to GCs treated with media alone (control), the addition of vit D3 significantly suppressed RAGE mRNA (by 33%) and protein (by 44%) levels ( $P < 0.05$ ) (Fig. 1) indicating that vit D3 might attenuate the effects of AGEs by suppressing RAGE expression.

### In-vitro effect of AGEs (HGA) with or without vit D3 on steroid gene expression in human luteinized GCs

HGA, a precursor for AGEs, binds and activates the pro-inflammatory cell membrane receptor RAGE. Compared to GCs cultured in media alone (control), HGA treatment significantly increased CYP11A1 (by 48%), 3 $\beta$ -HSD (by 38%), StAR (by 42%), CYP17A1 (by 30%) and LHR (by 37%) mRNA expression levels ( $P < 0.05$ ) but did not alter CYP19A1 or FSHR mRNA levels (Fig. 2). The addition of vit D3 to HGA inhibited the HGA-induced increase in CYP11A1, StAR, CYP17A1 and LHR mRNA levels, but not 3 $\beta$ -HSD mRNA levels, which remained 48% higher than controls ( $P < 0.05$ ). Although HGA alone did not increase CYP19A1 mRNA expression, the addition of vit D3 to HGA significantly increased CYP19A1 mRNA expression as compared to controls ( $P < 0.05$ ).

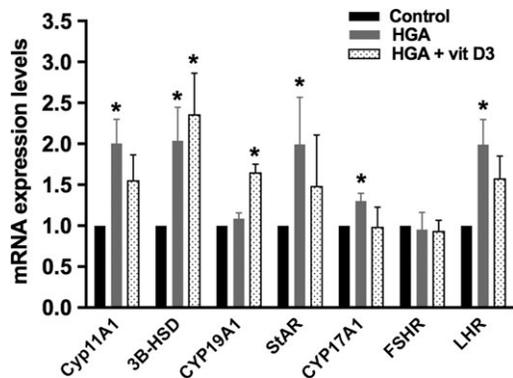
### In-vitro effect of AGEs (HGA) with or without vit D3 on E2 release by human luteinized GCs

To further study the effect of HGA in the presence of absence of vit D3 on E2 release by GCs, we treated GCs with media alone (control) or with HGA with or without vit D3. It has been shown that E2 production of cultured human luteinized GCs gradually decreases during culture in media alone (Foldesi et al., 1998). Similarly in our results as seen in Fig. 3 in control cells, E2 in cell culture media of control GCs decreased from  $356 \pm 99$  pmol/L at baseline to  $224 \pm 55$  pmol/L following 48 h of culture ( $P = 0.03$ ). Baseline E2 concentration levels in culture media of GCs treated with media alone, HGA alone or HGA + vit D3 were identical ( $P = 0.42$ ) (Fig. 3). GCs treated for 48 h with HGA alone or



**Figure 1** Real-time PCR and immunofluorescence for RAGE in cumulus granulosa cells treated with media alone or media with 1,25 dihydroxyvitamin D3 (vit D3). Pooled cumulus granulosa cells of women undergoing oocyte retrieval following ovarian stimulation for IVF were mechanically collected. Cells were treated with media alone or media + vit D3 (100 nM) for 24 h after which (A) RT-PCR for RAGE mRNA ( $n = 4$ ) and (B) immunofluorescence for RAGE protein ( $n = 4$ ) were performed. The addition of vit D3 significantly suppressed RAGE mRNA and protein levels. For immunofluorescence, the positive signal is seen in green and the blue signal represents DAPI; (a, b) are for cells treated with media alone (control) and (c, d) are for cells treated with media + vit D3. \* $P < 0.05$  for media + vit D3 versus media alone.

HGA + vit D3 had significantly higher E2 concentration levels compared to GCs treated with media alone for 48 h ( $P = 0.02$ ). In GCs treated with HGA alone, E2 in cell culture media remained stable between baseline ( $378 \pm 114$  pmol/L) and 48 h following culture ( $385 \pm 110$  pmol/L,  $P = 0.84$ ). Similarly in GCs treated with HGA + vit D3, E2 in cell culture media remained stable between baseline ( $330 \pm 84$  pmol/L) and 48 h following culture ( $382 \pm 106$  pmol/L,  $P = 0.22$ ).

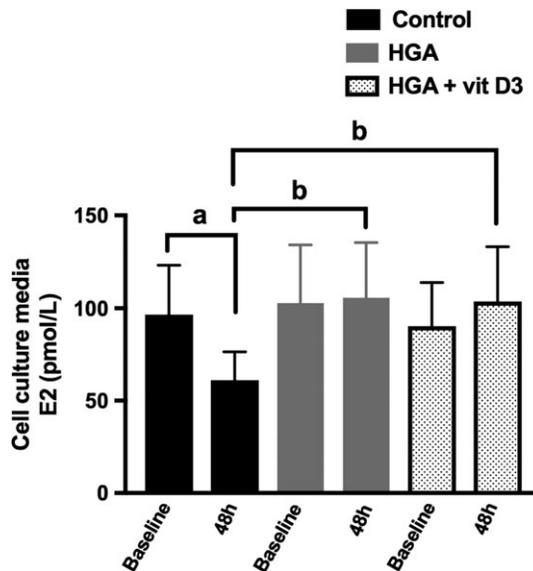


**Figure 2** Real-time PCR for steroid gene expression following treatment of cumulus granulosa cells with human glycated albumin (HGA), as a precursor for AGEs, with or without 1,25 dihydroxyvitamin D3 (vit D3). Pooled cumulus granulosa cells of women ( $n = 4$ ) undergoing oocyte retrieval following ovarian stimulation for IVF were mechanically collected by cutting the cumulus layer from each oocyte. Cells were treated with HGA (0.4 mg/mL) with or without vit D3 (100 nM) for 24 h after which real-time PCR was performed. HGA significantly increased CYP11A1 (side chain cleavage), 3 $\beta$ -HSD, StAR, CYP17A1 and LHR mRNA expression levels but did not alter CYP19A1 or FSHR mRNA levels. The addition of vit D3 to HGA inhibited the HGA-induced increase in CYP11A1, StAR, CYP17A1 and LHR but not increase in 3- $\beta$ HSD mRNA levels. The combination of HGA + vit D3 significantly increased CYP19A1 mRNA expression levels. Controls cells were set at 1. \* $P < 0.05$  for HGA versus control, and  $^aP < 0.05$  for HGA + vit D3 versus control.

## Discussion

This study evaluated the correlation between follicular fluid sRAGE, a decoy receptor that captures the circulating AGEs thus preventing the activation of the pro-inflammatory RAGE signaling pathway, and follicular fluid 25-OHD in a relatively large sample of women without PCOS. Our results showed that there is a positive correlation between follicular fluid sRAGE and 25-OHD. We also used human luteinized GCs as a model to assess whether treatment with AGEs alters genes involved in steroid production and to assess whether vit D3 application attenuates the effect of AGEs. The results demonstrated that HGA (AGEs) significantly increased CYP11A1, 3 $\beta$ -HSD, StAR, CYP17A1 and LHR mRNA levels but did not change CYP19A1 or FSHR mRNA levels. The addition of vit D3 inhibited most of the HGA-induced changes in these genes. Additionally, HGA (AGEs), in the presence or absence of vit D3, significantly increased E2 release by GCs in culture media. We also found that vit D3 downregulated RAGE mRNA and protein expression, which could constitute a mechanism by which vit D3 inhibited the HGA-induced changes in steroid gene expression.

The functions of sRAGE in the follicular fluid are still unclear, but it may impact the activities of the AGE-RAGE system in ovarian follicles, and it has been shown to correlate with markers of ovarian reserve, such as follicular fluid anti-Mullerian hormone and number of oocytes retrieved following oocyte retrieval for IVF (Merhi *et al.*, 2014b). Similar to what has been reported in the follicular fluid of women with PCOS (Garg *et al.*, 2017), we found, in women without PCOS, that there is a



**Figure 3** Effect of human glycated albumin (HGA), as a precursor for AGEs, with or without 1,25 dihydroxyvitamin D3 (vit D3) on estradiol (E2) release by cumulus granulosa cells. Pooled cumulus granulosa cells of women ( $n = 4$ ) undergoing oocyte retrieval following ovarian stimulation for IVF were mechanically collected by cutting the cumulus layer from each oocyte. Cells were treated with HGA (0.4 mg/mL) with or without vit D3 (100 nM) for 24 h after which cell culture media was collected for E2 radioimmunoassay. Between baseline and following 48 h of culture, E2 release significantly decreased in control cells ( $P = 0.03$ ) but remained unchanged in cells treated with HGA alone or with HGA + vit D3 ( $P = 0.84$ ). Following 48 h of culture, E2 in cell culture media was significantly higher with HGA alone or with HGA + vit D3 compared to control cells. <sup>a</sup> $P < 0.05$  for baseline versus 48 h levels in the control cells, and <sup>b</sup> $P < 0.05$  for HGA versus control at 48 h of culture and <sup>b</sup> $P < 0.05$  for HGA + vit D3 versus control at 48 h of culture.

weak positive correlation ( $R$ -value was only 0.27) between follicular fluid sRAGE and 25-OHD. Additionally, Sung *et al.* (2013) reported a positive correlation between serum sRAGE and serum 1,25 dihydroxyvitamin D3 in hemodialysis patients who received vit D3 supplementation. Interestingly, vit D3 replacement to vit D-deficient women with PCOS increases serum sRAGE levels, potentially suggesting a direct effect of vit D3 on sRAGE production (Irani *et al.*, 2014). Both sRAGE and vit D have anti-inflammatory actions, which could explain this positive relationship between the two molecules. Our findings also showed that follicular fluid sRAGE levels correlated positively with follicular fluid levels of the two AGEs: pentosidine and CML. Similarly, Willemssen *et al.* (2012) showed that serum pentosidine and CML levels were positively correlated with serum sRAGE levels in patients hospitalized for heart failure. Similarly, Kerkeni *et al.* (2012) observed that sRAGE serum levels were positively correlated with serum AGEs in patients with diabetic retinopathy. An explanatory hypothesis for this correlation is that as the pro-inflammatory AGEs increase, there is a counter-effect of anti-inflammatory response as reflected by more sRAGE production, i.e. increased AGEs stimulate sRAGE expression as a feedback mechanism. Finally, serum sRAGE is related to the insulin resistance state; e.g. it

has been shown that there is an inverse relationship between serum sRAGE and serum glucose, insulin and HOMA-IR levels (Basta et al., 2006). Our data showed similar findings in the follicular fluid where sRAGE was negatively correlated with follicular fluid insulin and glucose and positively correlated with SHBG. The relationship between sRAGE, glucose and insulin in the follicular fluid environment might be important in the dynamics of the follicle development and ultimately in oocyte maturation. More studies pertaining to this topic are needed.

Steroidogenesis is crucial for the synchronization of follicle growth and oocyte development (Dumesic et al., 2015). Our current findings demonstrated that HGA, a precursor for AGEs, altered steroidogenic enzymes involved in E2 production and maintained a significantly higher level of E2 in the culture media compared to control cells, despite having no effect on aromatase (CYP19A1) mRNA levels. This was reflected by an upregulation in CYP11A1, 3 $\beta$ -HSD, StAR, CYP17A1 and LHR (Fig. 4). One explanation could be that AGEs, along with the upregulation of these genes, increase E2 'release' from GCs rather than just increasing production by induction of aromatase. Also there could have been a change in aromatase enzymatic activity/protein expression without a change in mRNA levels. Another hypothesis is that HGA could be blocking E2 metabolism rather than directly affecting directly E2. Vit D3 has been shown to downregulate RAGE expression in non-ovarian cells

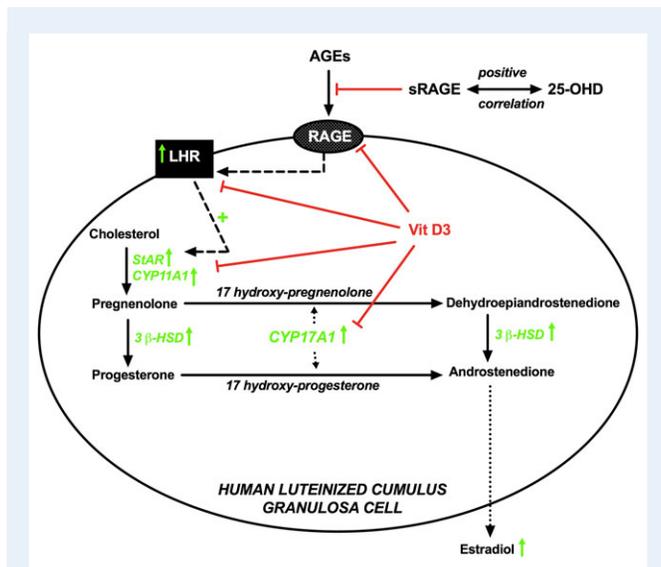
(Dreyer et al., 2014, Guo et al., 2016; Lee et al., 2014). In cell culture studies, calcitriol was shown to attenuate AGE-induced upregulation of RAGE mRNA and protein and counteracted the stimulating effect of AGEs on NF- $\kappa$ B pathway, which is a common pathway between vit D and AGEs (Talmor et al., 2008). In streptozotocin-induced diabetic rats, the increased expression of cardiac RAGE was attenuated by calcitriol (Lee et al., 2014). Also in an *in-vitro* blood-brain barrier model, RAGE expression was reduced following vit D3 application (Guo et al., 2016). Similarly, our current findings demonstrated that vit D3 suppressed RAGE mRNA and protein expression in human GCs, which could constitute a mechanism by which vit D3 inhibited the HGA-induced changes in steroid gene expression. The mechanisms of this relationship require further studies in order to better elucidate their role in ovarian function and female reproduction.

There are several limitations for this study. We did not have serum levels for the markers measured in the follicular fluid. Additionally, several types of ovarian cells play a role in steroidogenesis, such as mural GCs and theca cells. In this study, we focused on cumulus GCs that express characteristics distinct from mural GCs (Eppig et al., 1997, Vanderhyden and Tonary, 1995). Cumulus cells communicate with each other and with the oocyte through specialized gap junctions which allow metabolic exchange and transport of signaling molecules appropriate for follicular development (Dumesic et al., 2015, Simerman et al., 2015, Uyar et al., 2013). Thus, cumulus GCs, arguably, play a more important role in regulating oocyte maturation and in reflecting oocyte quality (Dumesic et al., 2015; Uyar et al., 2013). Another limitation is that our GCs were luteinized, as they were collected from women who were stimulated with gonadotropins. However, although these luteinized GCs may not be the best tool for studying steroidogenesis, we (Merhi et al., 2013) and others (Catteau-Jonard et al., 2008) have shown that luteinized GCs are responsive to several hormonal treatments *in vitro*. Another potential limitation is that we have pooled cumulus GCs of all small and large follicles retrieved during IVF from each participant. However, large follicles are more mature than smaller follicles, and thus AGEs and vit D3 *in vitro* could act differently on GCs obtained from large versus small follicles. Finally, the doses of HGA and vit D3 and the culture time used in the study were based on published studies but may not be ideal to reflect changes that occur physiologically.

In summary, understanding the role of the AGE-RAGE system in ovarian dysfunction, particularly for women who have elevated levels of these AGEs, such as obese or PCOS women or women who eat unhealthy diets, may help in our understanding of the mechanisms involved ovarian function. The results of this study provide a proof-of-principle for the potential for vit D supplementation in preventing some ovarian dysfunction and provide justification for the continued development of clinical trials of vit D as a cost-effective strategy to improving function of the ovaries. Additionally, the development of new therapeutic agents, such as AGE blockers, could represent a novel strategy to treat and/or prevent ovarian dysfunction.

## Authors' roles

Z.M. designed the study, performed the experiments and wrote major parts of the article. E.B. helped with the study design, performed some of the experiments and wrote parts of the article. M.J.C. supervised all of the experiments, provided feedback on the progress of the study and wrote parts of the article.



**Figure 4** Schematic diagram illustrating the relationship between vitamin D and AGEs in follicular fluid and human luteinized granulosa cells. A positive correlation exists between follicular fluid 25-hydroxyvitamin D (25-OHD) and follicular fluid sRAGE, which inhibits the circulating AGEs from activating their pro-inflammatory RAGE signaling pathway. AGEs affect steroidogenesis by upregulating luteinizing hormone receptor (LHR), which is known to induce steroidogenic acute regulatory protein (StAR) and P450 side-chain cleavage enzyme (CYP11A1). AGEs also upregulate 17 $\alpha$ -hydroxylase/17,20 lyase (CYP17A1) and 3 beta-hydroxysteroid dehydrogenase (3 $\beta$ -HSD) ultimately leading to more production and release of estradiol. Upward green arrows indicate upregulation by AGEs. 1,25-dihydroxyvitamin D3 (vit D3) downregulates RAGE expression and inhibits the effect of AGEs on LHR, StAR, CYP11A1 and CYP17A1, but not the effects on 3 $\beta$ -HSD.

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## Conflict of interest

All authors have nothing to disclose.

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