

# VDR GENE EXPRESSION AND GLOBAL DNA METHYLATION ARE ASSOCIATED WITH INSULIN RESISTANCE IN ESRD PATIENTS ON CHRONIC HAEMODIALYSIS

D. Kirmizis<sup>1</sup>, F. Chatzopoulou<sup>2</sup>, D. Chatzidimitriou<sup>2</sup>, G. Tzimagiorgis<sup>3</sup>, A. Papagianni<sup>1</sup>, G. Efstratiadis<sup>1</sup>

<sup>1</sup>Department of Nephrology, Hippokration General Hospital, Aristotle University, Thessaloniki, Greece;

<sup>2</sup>Laboratory of Microbiology, Aristotle University, Thessaloniki, Greece;

<sup>3</sup>Laboratory of Biochemistry, Aristotle University, Thessaloniki, Greece

## OBJECTIVES

Evidence suggest a probable role of genetic inactivation of the vitamin D receptor (VDR) gene, and possibly of epigenetic gene modifications, in insulin resistance in uraemia.

The aim of the present study was to investigate the potential genetic and epigenetic associations of insulin resistance, with vitamin D deficiency, VDR gene expression, and DNA methylation, in nondiabetic end-stage renal disease (ESRD) patients on chronic haemodialysis (HD).

## METHODS

- Cross-sectional study: 50 non-diabetic ESRD Caucasian patients stabilized on HD (>6 months) (URR  $\geq 0.65$ ; eKt/V  $\geq 1.2$ )
- Exclusion criteria: diabetes mellitus, pre-diabetes, liver disease, active infection, malnutrition, autoimmune diseases, malignancies, folate supplementation, antibiotics, corticosteroids or cytotoxic drugs.
- Blood samples taken from a peripheral vein under fasting conditions. Biochemistry, plasma 25(OH)D, fasting glucose, insulin, total homocysteine (Hcy), intact parathormone (iPTH), and serum high-sensitivity C-reactive protein (hsCRP) levels were measured by routine techniques.
- Insulin resistance was assessed with the use of the Homeostasis Model Assessment (HOMAIR=[fasting serum insulin ( $\mu$ U/ml)  $\times$  fasting plasma glucose (mmol/L)]/22.5).
- VDR gene expression (VDRG) (real-time PCR): relative quantification of leukocyte VDR mRNA, reported as  $2^{-\Delta\Delta Ct}$ , was performed with the use of calibration curves using GAPDH mRNA as an endogenous control.
- Global DNA methylation (GMeth) status (colorimetry): the final concentration was then calculated by reference to standard curves performed with the corresponding recombinant molecule.

## RESULTS

Parameter	1 <sup>st</sup> tertile (n=17) ( $\leq 2.31$ )	2 <sup>nd</sup> tertile (n=18) (2.32-4.57)	3 <sup>rd</sup> tertile (n=15) ( $> 4.57$ )
Age, years	68 (42-87)	65 (47-85)	74 (43-86)
Gender (male), n(%)	12 (71)	10 (44)*	8 (53)*
HD duration, months	71 (12-291)	45 (12-108)	58 (12-96)
Dialysis method: ⊙ HD, n (%) ⊙ Online HDF, n(%)	10 (77) 3 (23)	8 (57) 6 (43)	8 (67) 4 (33)
BMI, kg/m <sup>2</sup>	22.6 $\pm$ 1.9	24.9 $\pm$ 2.8	25.6 $\pm$ 2.3*
Kt/V	1.18 $\pm$ 0.3	1.19 $\pm$ 0.44	1.21 $\pm$ 0.38
Mean BP (mmHg)	104 $\pm$ 7	106 $\pm$ 9	105 $\pm$ 4
CVD, n(%)	8 (62)	8 (57)	6 (50)
Art. hypertension, n(%)	14 (82)	16 (89)	14 (93)
Hb, g/dL	10.8 $\pm$ 0.2	11.3 $\pm$ 0.4	11.1 $\pm$ 0.2
WBC, $\times 10^9$ /mL	8.6 $\pm$ 0.4	8.2 $\pm$ 1.0	7.8 $\pm$ 0.5
Glucose, mg/dL	85 $\pm$ 12	104 $\pm$ 11	112 $\pm$ 30
Insuline, $\mu$ U/mL	9.1 (3.0-38.3)	13.7 (3.4-33.5)	16.7 (3.6-53.6)*
HOMAIR	2.21 (0.58-13.32)	3.48 (0.65-13.80)	4.93 (0.86-18.27)*
hsCRP, mg/L	0.80 (0.22-2.11)	1.79 (1.1-3.9)*	3.26 (2.1-5.9)*
Albumin, g/dL	3.5 $\pm$ 0.4	3.4 $\pm$ 0.3	3.3 $\pm$ 0.2
Urea, mg/dL	146 $\pm$ 39	148 $\pm$ 37	141 $\pm$ 42
Alkaline phosphatase, U/L	101 $\pm$ 54	122 $\pm$ 98	188 $\pm$ 122*
25(OH)D, pg/mL	10.2 $\pm$ 6.2	9.1 $\pm$ 2.6	6.7 $\pm$ 1.8*
iPTH, pg/mL	222 $\pm$ 183	369 $\pm$ 255	651 $\pm$ 460*
Hcy, $\mu$ mol/L	25.5 $\pm$ 11.9	21.7 $\pm$ 10.8	16.3 $\pm$ 6.7*
GMeth (%)	3.66 $\pm$ 1.03	4.28 $\pm$ 0.95	4.68 $\pm$ 0.84*
VDRG	0.84 $\pm$ 0.65	0.66 $\pm$ 0.79	0.16 $\pm$ 0.38*

Results are expressed as mean  $\pm$ SD, median (range) or patient number (%), as appropriate. \*P<0.05 compared to the respective 1st tertile

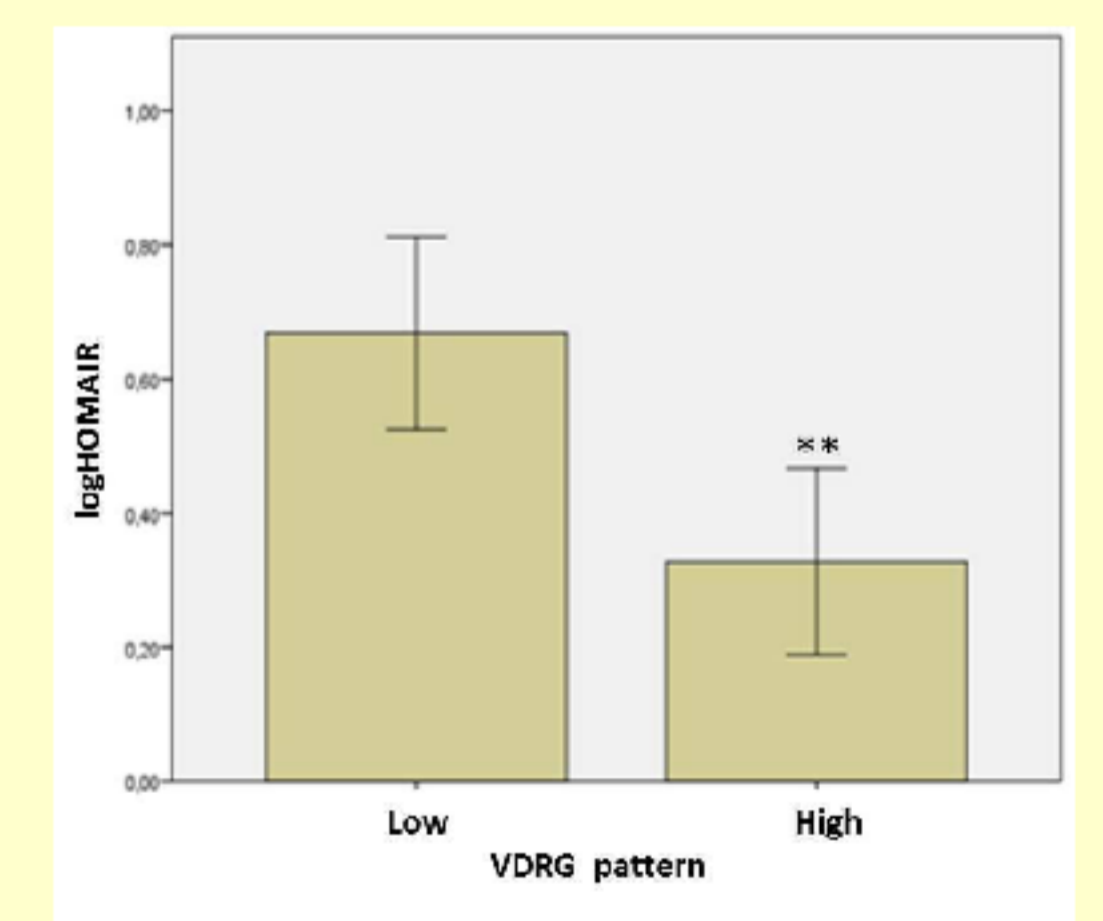
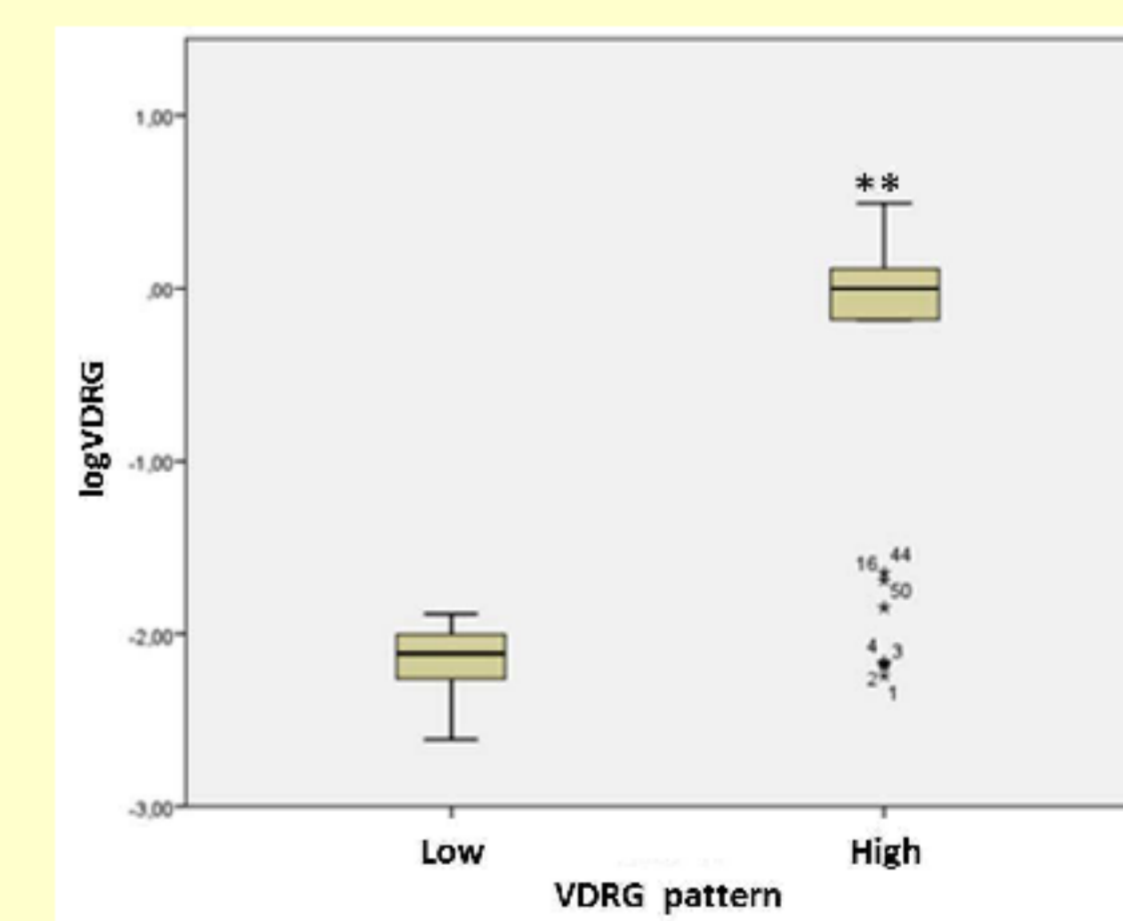
Parameter	Low VDRG (n=20)	High VDRG (n=30)
Age, years	69 (47-90)	67 (42-87)
Gender (male), n(%)	14 (70)	18 (60)*
HD duration, months	49 (6-108)	64 (6-291)*
BMI, kg/m <sup>2</sup>	25.6 $\pm$ 2.7	23.3 $\pm$ 2.2*
Glucose, mg/dL	115 $\pm$ 24	95 $\pm$ 18
Insulin, $\mu$ U/mL	19.3 $\pm$ 10.7	11.9 $\pm$ 10.4**
HOMAIR	4.22 (0.62-16.31)	2.29 (0.58-18.27)**
hsCRP, mg/L	0.82 (0.11-5.5)	0.7 (0.1-6.2)
Albumin, g/dL	3.4 $\pm$ 0.5	3.3 $\pm$ 0.3
Urea, mg/dL	134 $\pm$ 35	155 $\pm$ 36
Alkaline phosphatase, U/L	140 $\pm$ 95	121 $\pm$ 94
25(OH)D, pg/mL	9.8 $\pm$ 4.6	10.1 $\pm$ 5.3
iPTH, pg/mL	663 $\pm$ 436	295 $\pm$ 225*
Hcy, $\mu$ mol/L	19.1 $\pm$ 7.3	26.8 $\pm$ 12.8*
GMeth (%)	4.26 $\pm$ 0.87	4.24 $\pm$ 2.0
VDRG	0.0078 $\pm$ 0.0031	1.1 $\pm$ 0.92**

Results are expressed as mean  $\pm$ SD, median (range) or patient number (%), as appropriate. \* P<0.05 compared to low VDRG; \*\* P<0.001 compared to low VDRG

Parameters	Low VDRG		High VDRG		
	r	P value	r	P value	
VDRG	-0.493	0.002	Hcy	-0.419	0.002
GMeth	0.513	0.003	VDRG	-0.398	0.003
iPTH	0.591	0.04	GMeth	0.38	0.05
Out of Model:			Out of Model:		
BMI	0.228	0.207	Urea	0.310	0.102
Hcy	-0.215	0.258	iPTH	-0.214	0.264
logCRP	-0.301	0.265	logCRP	-0.197	0.281
25(OH)D	0.17	0.276	25(OH)D	0.172	0.374
Urea	0.094	0.389	Calcitriol	0.144	0.422
Calcitriol	0.144	0.389	BMI	0.115	0.518
HD duration	-0.038	0.464	HD duration	-0.022	0.908

†Adjusted for age.

- HOMAIR tertiles differed significantly in hsCRP, fasting insulin, iPTH, GMeth values, Hcy, 25(OH)D, and VDRG.
- Two discrete patterns of VDRG, a high and a low expression pattern, were noticed. They differed significantly in iPTH (295 $\pm$ 225 vs 663 $\pm$ 436 pg/ml respectively, P=0.014), HOMAIR (3.14 $\pm$ 3.66 vs 5.88 $\pm$ 4.27 respectively, P<0.001) and Hcy values (26.75 $\pm$ 12.85 vs 19.07 $\pm$ 7.27  $\mu$ mol/L respectively, P=0.02).
- On multivariate analysis, VDRG and GMeth were strong independent correlates of HOMAIR in both VDRG groups, whereas iPTH and Hcy were correlates of HOMAIR in the low and the high VDRG groups, respectively.



## CONCLUSIONS

VDR gene expression and DNA methylation are independent correlates of insulin resistance in non-diabetic ESRD patients.

## REFERENCES:

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