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# Transcriptional regulation of Endothelin-1 expression by Advanced Glycation End-products in human aortic endothelium is mediated via NF-kappaB and AP-1

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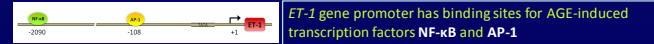


### Introduction:

- Advanced Glycation End-products (AGEs) are produced by the non-enzymatic glycation of proteins, lipids and nucleic acids, resulting in an overload of highly reactive molecules of endogenous or exogenous (dietary) origin.
- Increased AGE levels in circulation and concomitant elevated tissue deposition have been associated with diabetic complications, atheromatosis, ageing and more recently with polycystic ovary syndrome pathogenesis.
- Interaction of AGEs with their receptor RAGE (Receptor for AGEs) activates intracellular signaling pathways which induce targeted gene expression in endothelium including upregulation of cell adhesion molecules and endothelin-1 (ET-1), implicated in vascular injury and endothelial dysfunction.

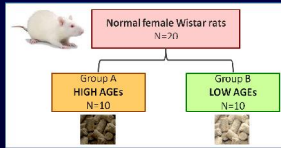
### Aim:

To explore the molecular mechanism of AGE-induced regulation of ET-1 gene/protein expression in human endothelial cells and investigate its functional relevance in normal rat vascular endothelium.



### Methods & Materials:

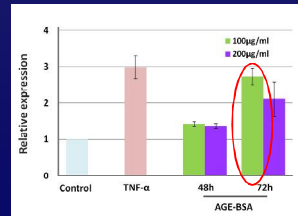
- Cell Cultures: Human Aortic Endothelial Cells (HAECs)
  - Phase contrast image HAECs
- Treatment with AGE-Bovine Serum Albumin (AGE-BSA)
  - Concentrations: 100, 200 µg/ml
  - Time points: 24, 48, 72h
- Quantitative real-time Polymerase Chain Reaction (real-time qPCR): ET-1 mRNA expression
- Flow cytometry analysis: RAGE expression
- Western Blot analysis: activated/phosphorylated ERK1/2 (p-ERK1/2) expression
- Electrophoretic-Mobility Shift Assay (EMSA): NF-kB and AP-1 binding to ET-1 gene promoter
- Immunohistochemistry: AGEs, RAGE, ET-1 expression in normal aortic endothelium of rats fed with low- or high-AGE content diet.



HIGH AGEs diet was prepared through exposure to 191° C for 30 min

### Results:

#### I. ET-1 gene expression analysis (mRNA levels) in HAECs after treatment with AGEs



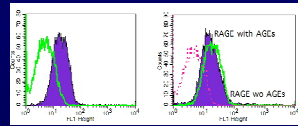
total RNA extraction  
reverse transcription (RT-PCR)  
real time qPCR

Forward Primer: CCAGGAGGCTCCAGAAACAG  
Reverse Primer: -GATGTCCAGGTGGCAGAAATG

TNF-α: positive control

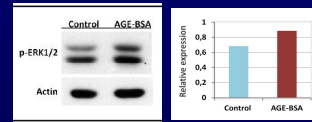
Treatment of HAECs with AGE-BSA induced ET-1 transcription in a time- and dose- dependent manner.

#### II. Flow cytometric analysis of RAGE expression in HAECs



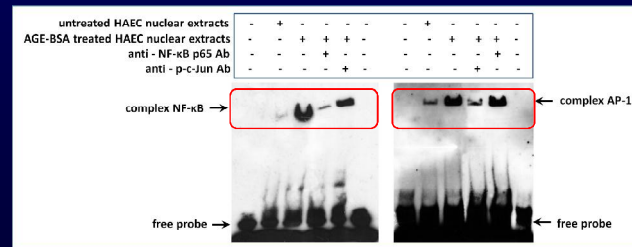
RAGE expression in HAECs remained unchanged after AGE-BSA treatment (72h, 100µg/ml).

#### III. Western blot for p-ERK1/2 in untreated (Control) and AGE-treated HAEC (AGE-BSA) protein extracts



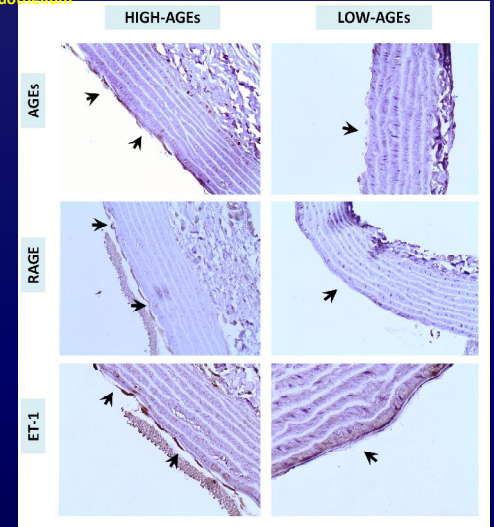
Induction of p-ERK1/2 expression after AGE-BSA administration (72h, 100µg/ml).

#### IV. Analysis of ET-1 gene promoter binding capacity of transcription factors NF-kB and AP-1 (EMSA)



Treatment of HAECs with AGE-BSA (72h, 100µg/ml) induced NF-kB and AP-1 activation and binding to ET-1 gene promoter.

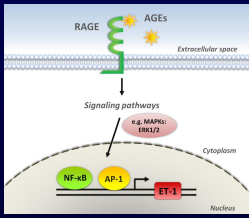
#### V. Immunohistochemical investigation of AGEs, RAGE and ET-1 in normal rat aortic endothelium



Increased expression and co-localization of AGEs, RAGE and ET-1 were observed in the aortic endothelium of normal rats fed with high-AGE diet compared with controls.

### Conclusion

AGE-RAGE signaling induces ET-1 protein expression in endothelium through regulation of ET-1 gene promoter by the transcription factors, NF-kB and AP-1 constituting a molecular mechanism that potentially contributes to the characteristic endothelial dysfunction of obesity, diabetic microvascular complications, atherogenesis and polycystic ovary syndrome.



### References:

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