

SUPPLEMENTARY INFORMATION

C2238/ α ANP modulates Apolipoprotein E through Egr-1/miR199a in vascular smooth muscle cells in vitro

Rosita Stanzione^{1*}, Sebastiano Sciarretta^{1,2*}, Simona Marchitti¹, Franca Bianchi¹, Sara Di Castro¹,
Stefania Scarpino³, Maria Cotugno¹, Giacomo Frati^{1,2}, Massimo Volpe^{1,3}, Speranza Rubattu^{1,3}

¹IRCCS Neuromed, Pozzilli; ²Department of Medical-Surgical Sciences and Biotechnologies, Sapienza University of Rome, Latina; ³Department of Clinical and Molecular Medicine, School of Medicine and Psychology, Sapienza University of Rome, Ospedale S. Andrea, Rome; Italy

*these authors contributed equally to this work

Running title: C2238/ α ANP and Apolipoprotein E

Correspondence to:

Speranza Rubattu, MD.
Clinical and Molecular Medicine Department,
School of Medicine and Psychology
Sapienza University, S.Andrea Hospital, Rome;
IRCCS Neuromed, Pozzilli (Is), Italy
e-mail: rubattu.speranza@neuromed.it
Tel. 0039 06 33775979; Fax 0039 06 33775061

Supplementary Figure legends

Supplementary Figure 1

Role of oxidative stress on Egr-1 stimulation by CC2238/ α ANP

Cells exposed to CC2238/ α ANP showed significant increase of Egr-1 expression levels, which was abolished in the presence of apocynin. Number of independent experiments=3.

CC2238/ α ANP = variant α ANP; Egr-1= early growth response protein-1; CTR= control

Supplementary Figure 2

Modulation of proteins involved in apoptosis, necrosis and inflammation by CC2238/ α ANP in HUVMSCs

Panels A-F. Representative western blots of cleaved-caspase-3, CREG, JNK, p38MAPK, Nf-kBp65, Smad4 under CC2238/ α ANP both in the presence and in the absence of NPR-C. Corresponding densitometric analysis is shown below each blot.

Data are expressed as mean \pm SD. Number of independent experiments=6.

** $p < 0.0001$ for CC2238/ α ANP vs all other points.

CC2238/ α ANP = variant α ANP; siRNA1=NPRC gene silencer 1; siRNA2=NPRC gene silencer 2; CREG=cellular repressor of E1A-stimulated gene; JNK=c-Jun N-terminal kinase; CTR=control.

Supplementary Figure 3

Modulation of proteins involved in apoptosis, necrosis and inflammation by CC2238/ α ANP in CAMSCs

Panels A-F. Representative western blots of cleaved-caspase-3, CREG, JNK, MAPKp38, Nf-kB, Smad4 under CC2238/ α ANP both in the presence and in the absence of NPR-C. Corresponding densitometric analysis is shown below each blot.

Data are expressed as mean \pm SD. Number of independent experiments=6.

** p<0.0001 for CC2238/ α ANP vs all other points.

CC2238/ α ANP = variant α ANP; siRNA1=NPRC gene silencer 1; siRNA2=NPRC gene silencer 2;

CREG=cellular repressor of E1A-stimulated gene; JNK=c-Jun N-terminal kinase; CTR= control.

Supplementary Figure 4

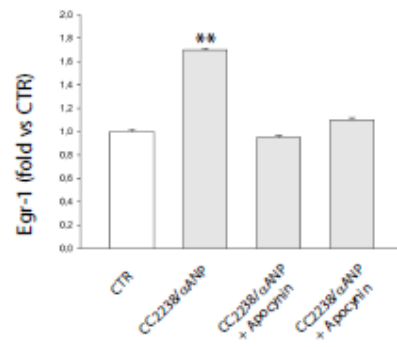
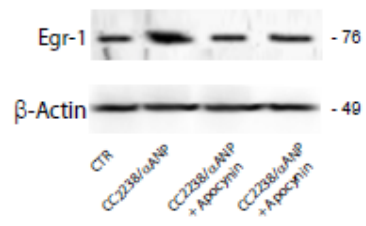
Results of ApoE gene sequencing in both HUVSMCs (panel A) and CASMCs (panel B).

HUVSMCs express both ApoE2 (Cys112/Cys158) and ApoE3 (Cys112/Arg158) isoforms.

CASMCs express both ApoE3 (Cys112/Arg158) and ApoE4 (Arg112/Arg158) isoforms.

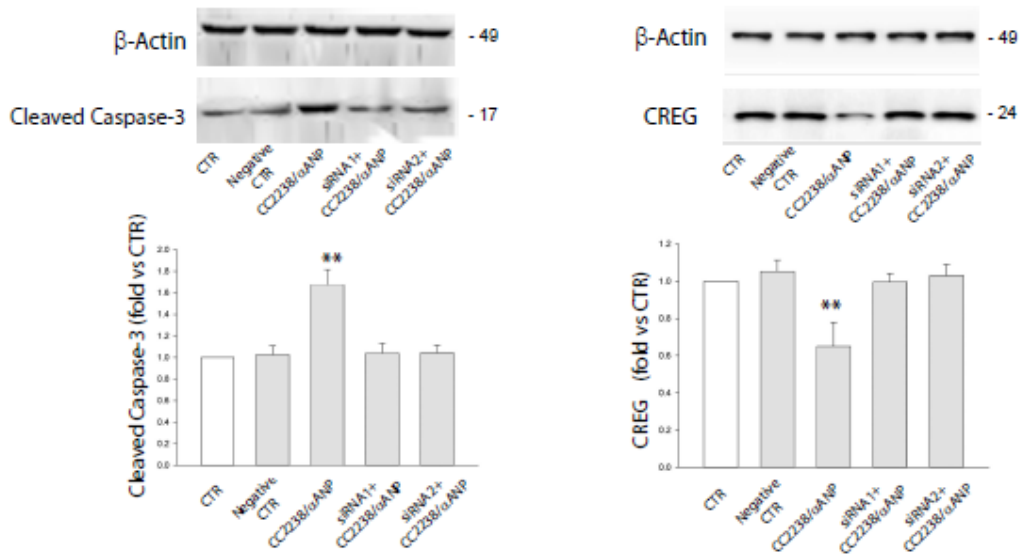
Cys=cysteine; Arg=arginine

Supplementary Figure 1

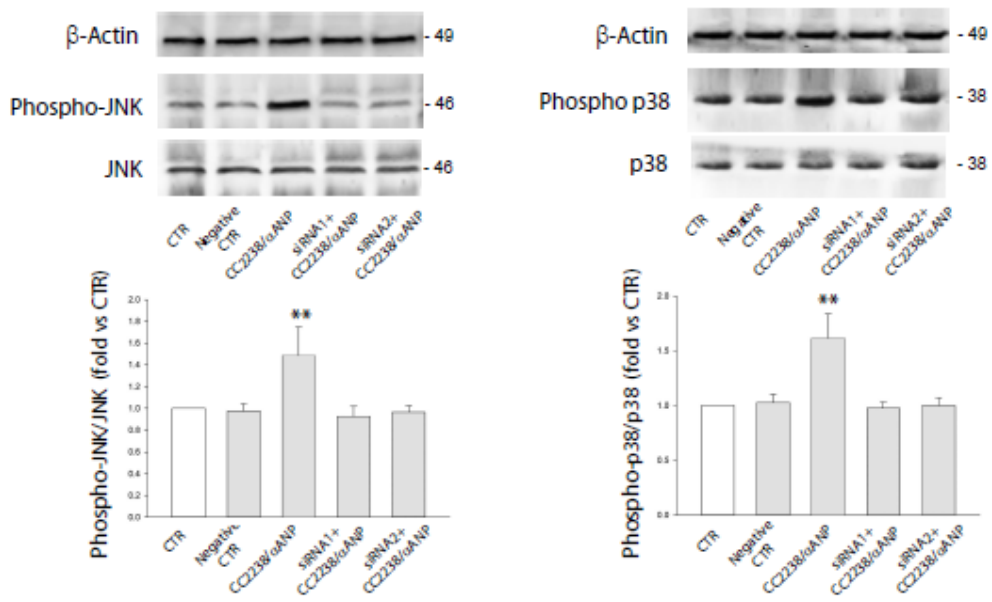


Supplementary Figure 2

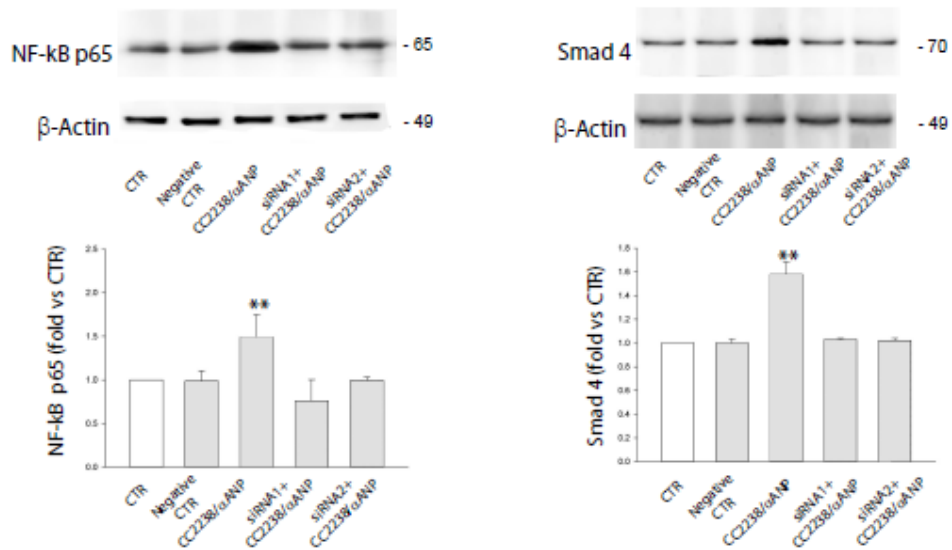
A



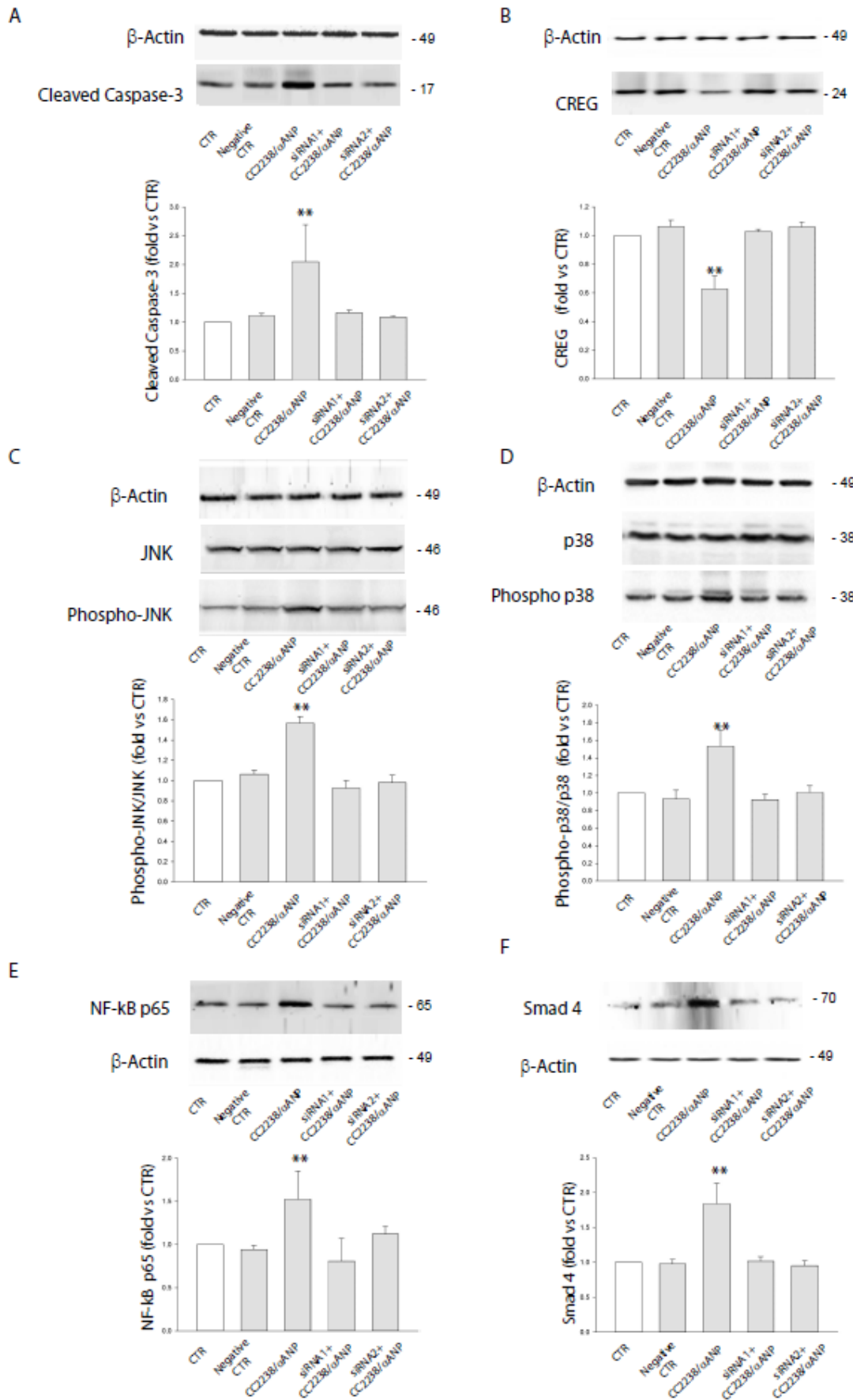
C



E

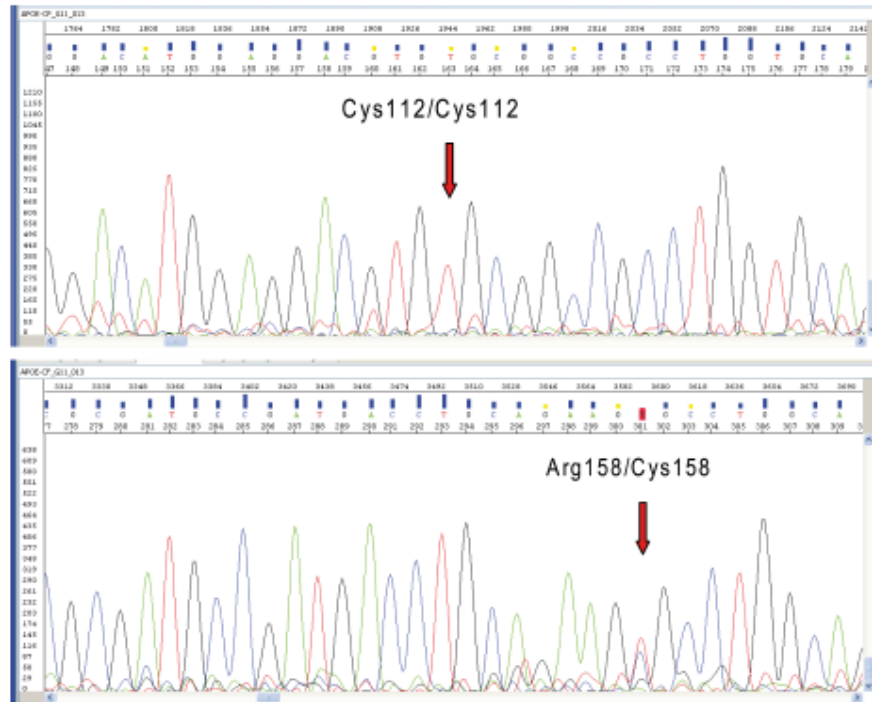


Supplementary Figure 3



Supplementary Figure 4

A



B

