

# Apabetalone (RVX-208) inhibits key drivers of vascular inflammation, calcification, and plaque vulnerability through a BET-dependent epigenetic mechanism

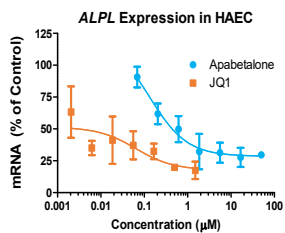
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Disclosure: Authors were employed by Resverlogix & held stock options with the exception of K. Rinker, et al. and D. Studer.

## ABSTRACT

Apabetalone (RVX-208) is an orally available small molecule bromodomain & extraterminal (BET) protein inhibitor that targets the second bromodomain (BD2) of BET proteins. Apabetalone returns dysregulated BET-dependent transcription toward normal physiological levels. In phase 2 trials, apabetalone treatment reduced the incidence of major adverse cardiac events by 44% in CVD patients and by 57% in diabetic CVD patients. Previous studies have highlighted apabetalone's positive impact on vascular calcification (VC) and inflammation (VI) marker expression in vitro, as well as its ability to lower serum alkaline phosphatase (ALP) levels, and improve atherosclerotic plaque stability parameters in treated patients. In CVD, elevated inflammatory mediators and cell surface adhesion molecules drive VI, resulting in leukocyte adhesion, infiltration, uptake of oxLDL, and ultimately plaque formation. Here we show in vitro that THP-1 monocyte adhesion to human aortic endothelial cells (HAECs) increases with TNF $\alpha$  stimulation and is attenuated by apabetalone treatment, with fewer monocytes attaching to HAECs under flow conditions. This functional outcome is attributed to apabetalone's reduction of key endothelial adhesion genes, VCAM-1 (50%, p=0.0001) and SELE (37%, p=9 x 10<sup>-5</sup>). Apabetalone also prevents TNF $\alpha$  induction of endothelial recruitment genes (MCP-1; 75%, p=0.0002) and genes involved in plaque rupture (IL8; 24%, p=2x10<sup>-5</sup>). Basal HAEC ALP expression, a potential contributor to endothelial dysfunction and VC, also decreases with apabetalone treatment (70%, p=0.005). Induction of VI genes by TNF $\alpha$  is BET-dependent as degradation of BET proteins by MZ-1 prevents an increase in transcripts in response to TNF $\alpha$  treatment. Ingenuity<sup>®</sup> Pathway Analysis (IPA<sup>®</sup>), GSEA, and GO analysis of HAEC gene expression data predicts apabetalone inhibition of pro-atherogenic pathways, gene sets, and upstream regulators induced by TNF $\alpha$ . These include cytokine and chemokine, Toll-Like Receptor (TLR), NF $\kappa$ B, Interferon and TNF $\alpha$  signaling. In addition, IPA<sup>®</sup> disease and biological function analysis predicts inhibition of immune cell activation and recruitment by apabetalone. Plasma proteomics (SOMAscan<sup>®</sup>) and IPA<sup>®</sup> analysis from apabetalone-treated CVD patients in ASSERT and ASSURE phase 2 trials indicate that apabetalone inhibits pro-atherogenic upstream regulators (IL-6 and IFN $\gamma$ ), canonical pathways, and diseases and functions. Serum ALP also decreases dose dependently with apabetalone treatment (ASSERT). Epigenetic inhibition of VI and VC driven atherogenesis likely contributes to the reduction in MACE observed in phase 2 apabetalone treated patients. The ongoing phase 3 post-acute coronary syndrome (ACS) clinical trial in T2DM patients, BETonMACE, is currently testing this

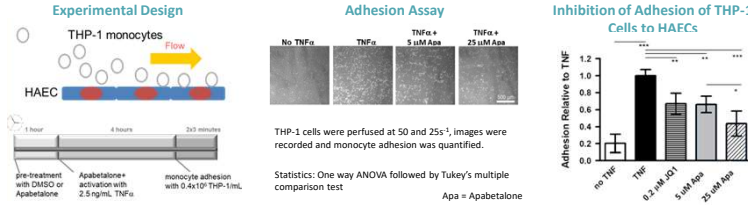
## Apabetalone inhibits alkaline phosphatase expression in HAEC



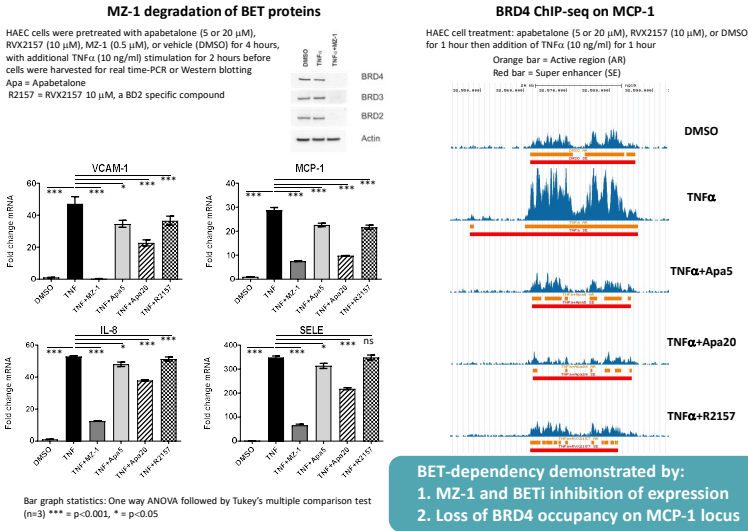
HAECs were treated with BET inhibitors for 4 hours, followed by analysis of alkaline phosphatase expression (gene symbol *ALPL*) by real-time PCR. JQ1: a pan selective BET inhibitor with a chemical scaffold different from apabetalone, used as a comparator molecule.

**Conclusion:** Apabetalone maximally reduced *ALPL* mRNA by 70% (p<0.001, Student's t-test versus cells treated with solvent only)

## Apabetalone suppresses monocyte adhesion to endothelial cells



## Apabetalone inhibits pro-atherogenic gene expression in endothelial cells in a BET-dependent manner by reducing BRD4 occupancy at gene loci



**BET-dependency demonstrated by:**  
 1. MZ-1 and BETi inhibition of expression  
 2. Loss of BRD4 occupancy on MCP-1 locus

## Bioinformatics of HAEC RNA-seq data: IPA<sup>®</sup>

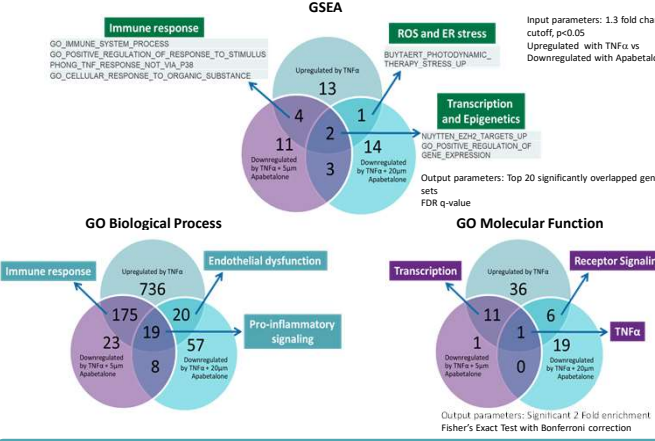
Upstream Regulators	TNF $\alpha$	TNF $\alpha$ +Apa5	TNF $\alpha$ +Apa20	TNF $\alpha$ +R2157
TNF signaling	13.6	-6.3	-4.4	-3.8
Tnf (family)	8.7	-1.7	-1.6	-1.8
IL6 signaling	11.9	-1.9	-2.4	-1.8
TLR signaling	6.1	-3.2	-2.8	-1.4
TLR8	6.6	-2.4	-2.9	-3.1
NF $\kappa$ B signaling	10.9	-6.3	-3.6	-3.9
NF $\kappa$ B (complex)	7.3	-1.9	-1.1	-2.1
RELA	7.1	-2.2	-1.4	-1.9
IL1 signaling	10.2	-4.4	-3.8	-3.8
IL1B	8.8	-4.7	-3.5	-4.2
Interferon signaling	6.6	-4.4	-3.4	-4.8
IFN7	6.8	-4.9	-2.8	-3.3
STAT1	6.9	-4.9	-2.8	-3.3
Transcription	6.8	-3.7	-6.9	-1.4
CSF2	6.8	-2.8	-1.9	-1.4
OSM	6.6	-1.4	-3.2	-4.9
RANK	6.8	-2.6	-2.6	-1.5
TGFB1	6.6	-2.1	-0.9	-1.8
IFN	7.1	-3.2	-0.9	-1.6
IFN (family)	7.8	-2.9	-2.7	-2.1
Msi (family)	6.6	-2.3	-2.9	-1.6

Canonical Pathways	TNF $\alpha$	TNF $\alpha$ +Apa5	TNF $\alpha$ +Apa20	TNF $\alpha$ +R2157
Cardiac Hypertrophy Signaling (Enhanced)	4.2	-2.6	-3.9	-4.2
Dendritic Cell Maturation	4.0	-3.6	-2.3	-2.6
HMG1 Signaling	3.9	-1.4	-2.3	-1.3
Production of Nitric Oxide and ROS in Macrophages	3.8	-3.6	-2.5	-2.0
IL-8 Signaling	3.7	-1.7	-2.8	-2.7
TREM1 Signaling	3.6	-1.9	-3.0	-1.7
IL-6 Signaling	3.6	-3.3	-2.8	-1.1
Type I Diabetes Mellitus Signaling	3.3	-3.8	-1.6	-1.9
INDS Signaling	3.0	-4.1	-2.2	-1.0
Role of Pattern Recognition Receptors	2.9	-1.9	-4.4	-2.0
NF $\kappa$ B Activation by Viruses	2.7	-2.1	-3.9	-2.8
Type II Diabetes Mellitus Signaling	2.6	-2.4	-2.9	-1.6
STAT3 Pathway	2.6	-2.5	-2.8	-2.8
Acute Phase Response Signaling	2.6	-2.4	-1.9	-4.2
RANK Signaling in Osteoclasts	2.4	-2.0	-3.3	-1.7
Thrombin Signaling	1.9	-2.1	-1.6	-1.6

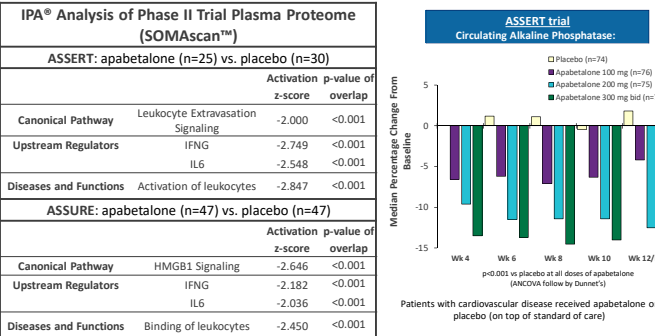
Apabetalone and R2157 inhibit TNF $\alpha$  activated pro-atherogenic upstream regulators, pathways, and diseases and functions

## Bioinformatics of HAEC RNA-seq data: Gene Set Enrichment Analysis (GSEA) and Gene Ontology (GO) Analysis



Apabetalone opposes TNF $\alpha$  activated gene sets: Pro-inflammatory, immune response, endothelial dysfunction

## Apabetalone inhibits pro-atherogenic drivers and processes and downregulates ALP in CVD patients



Plasma proteins cutoff = 10% Δ (vs. placebo, p<0.05). IPA<sup>®</sup> z-score s-2 predicts inhibition based on target gene modulation. p-value = Fisher's Exact Test.

## SUMMARY

1. Apabetalone inhibits the expression of pro-atherogenic genes in a BET-dependent manner resulting in the inactivation of pathologic pathways and regulators.
2. BET-dependent downregulation of vascular inflammation, cell adhesion, and ALP by apabetalone may contribute to the reduced incidence of MACE (phase 2), a hypothesis currently being tested in the phase 3 cardiovascular outcomes trial, BETonMACE.