

Laura Tsujikawa¹, Li Fu¹, Shovon Das¹, Brooke D. Rakai¹, Chris D. Sarsons¹, Chris Halliday¹, Stephanie C. Stotz¹, Emily Daze¹, Sylwia Wasiak¹, Deborah Studer², Sam Dorosz², Kristina D. Rinker², Michael Sweeney³, Jan O. Johansson³, Norman C. Wong¹ and Ewelina Kulikowski¹
¹Resverlogix Corp, Calgary, AB, Canada ² Department of Chemical and Petroleum Engineering, University of Calgary, Calgary, AB, Canada ³ Resverlogix Inc., San Francisco, CA, USA

Abstract

Background: Apabetalone (RVX-208) is a small molecule bromodomain & extraterminal (BET) protein inhibitor, targeting the second bromodomain (BD2) of BET proteins. In cardiovascular disease (CVD) patients enrolled in phase 2 trials apabetalone treatment reduced the relative risk of a CV event by 44% (Nicholls 2017). Elevated cytokines, such as TNF α , promote vascular inflammation (VI) and monocyte adhesion in CVD and Diabetes Mellitus, driving atherosclerosis. Here we test the impact of apabetalone on cell types that contribute to atherosclerosis.

Methods: Human endothelial cells (HUVECs) and THP-1 monocytes were stimulated with TNF α and treated with apabetalone or MZ-1 PROTAC. mRNA (qPCR, Nanostring) and protein levels (FACS, western blot) were compared. HUVEC-THP-1 adhesion assays assessed the functional consequences of TNF α stimulation and apabetalone treatment. The phase 2 ASSURE CVD patient plasma proteome (SOMAscan[®]) was analyzed using Ingenuity[®] Pathway Analysis (IPA[®]) to predict canonical and upstream regulator pathways impacted by apabetalone.

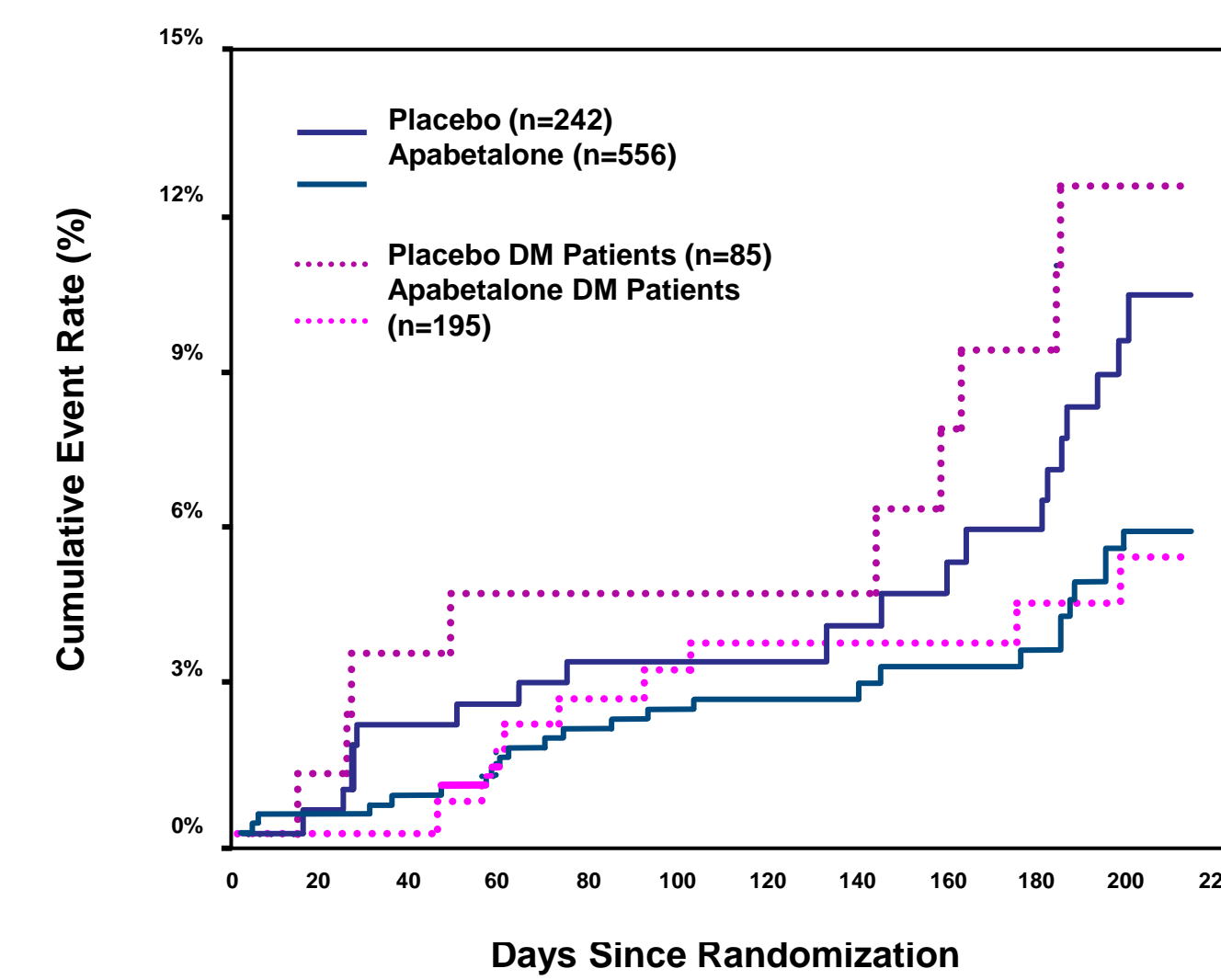
Results: Apabetalone repressed transcription of inflammatory and adhesion genes in TNF α -stimulated HUVEC and THP-1 cells. Corresponding HUVEC protein abundance was also reduced. MZ-1 BET protein degradation blocked TNF α responses, indicating BET-dependency. Functionally, apabetalone suppressed monocyte THP-1 cell adhesion to inflamed endothelial cells. CVD patient plasma proteome analysis revealed that apabetalone reduced key players in adhesion (VCAM-1, ICAM-1) and plaque stability (MMP-3, MMP12). IPA[®] analysis of the clinical proteome data predicted that apabetalone inhibits pro-atherogenic mediators and inflammatory pathways.

Conclusion: Apabetalone attenuates VI through the epigenetic regulation of inflammatory and adhesion gene transcription. Downregulation of VI by apabetalone may contribute to the reduction in CVD events observed in phase 2 studies. The ability of apabetalone to prevent major adverse cardiac events in post-acute coronary syndrome patients with type 2 diabetes mellitus (DM) and low HDL-C is being assessed in phase 3 trial (BETonMACE).

All information in this presentation is copyrighted and no reproduction is permitted.

Results

Apabetalone lowers MACE, phase 2b analysis



Apabetalone counters pro-inflammatory gene expression

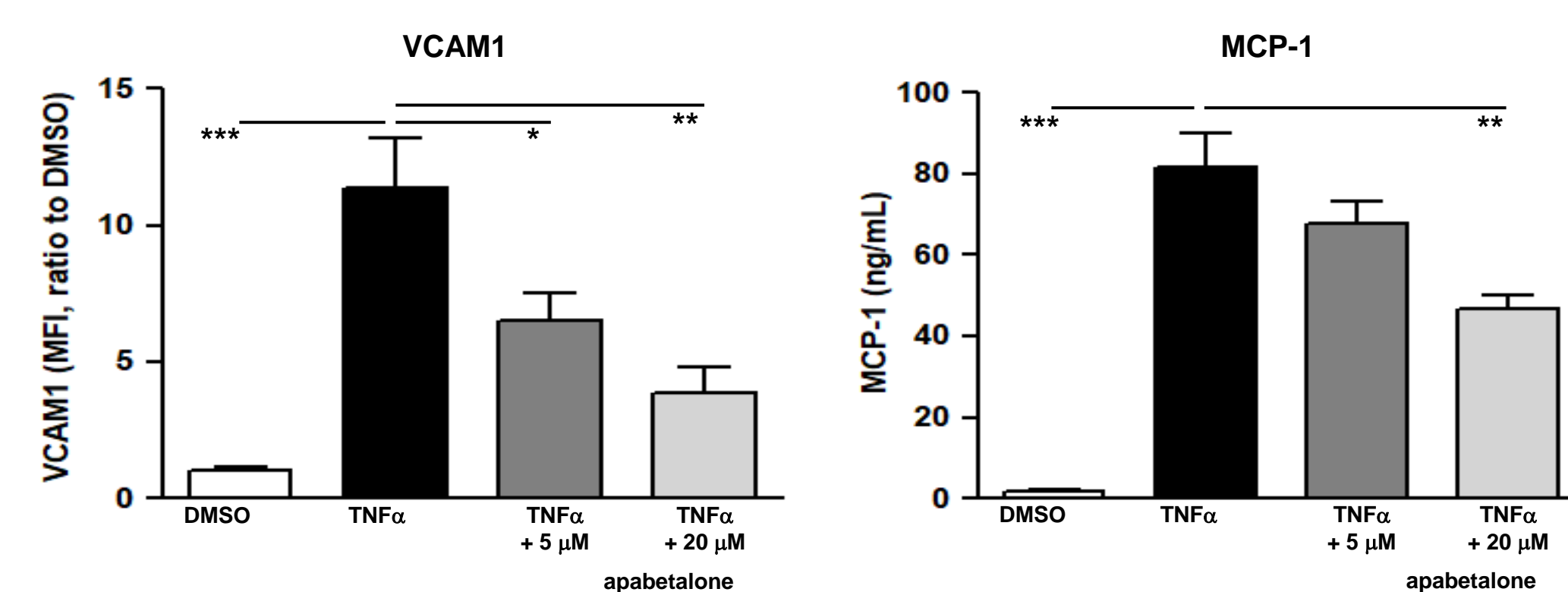
mRNA transcripts of key HUVEC and THP-1 cytokine, chemokine, TLR Signaling, and adhesion molecules are induced by TNF α /stimulants (4h co-treatment), and reduced by apabetalone (1h pre-treatment+ 4hr co-treatment).

| Simulation: | | TNF α | | IL-1 β | | LPS | | | | |
|--------------------|--------------|----------------|-------------|----------------|-------------|----------------|-------------|-----|-----|-----|
| Gene | Control | Fold induction | % reduction | Fold induction | % reduction | Fold induction | % reduction | | | |
| Cytokines | COX2 | 4 | NS | -86 | 19 | -46 | -85 | 1 | -42 | -83 |
| | CSF2 | 945 | -82 | -98 | 8096 | -59 | -91 | 9 | -64 | -85 |
| | IL-1 β | 1685 | -90 | -99 | ND | ND | ND | ND | ND | ND |
| | IL-6 | 9 | -51 | -91 | 191 | -54 | -84 | 1.6 | -67 | -69 |
| | IL-8 | 26 | ND | -48 | ND | ND | ND | ND | ND | ND |
| Chemokine | OPG | 43 | -95 | -99 | 142 | -96 | -99 | 1.4 | -71 | -84 |
| | MCP-1 | 4 | -21 | -71 | 44 | -35 | -62 | 4 | -50 | -82 |
| TLR signaling | MYD88 | 1 | NS | -56 | 1 | -30 | -66 | 1.6 | -44 | -38 |
| | CD44 | 2 | NS | -34 | 3 | NS | NS | 1 | -33 | -34 |
| Adhesion molecules | SELE | 1164 | NS | -54 | 368 | -17 | -40 | 11 | -51 | -76 |
| | VCAM1 | 196 | -59 | -83 | 96 | -72 | -91 | 6 | -73 | -96 |

| THP-1 | | TNF α | | |
|--------------------|--------------|----------------|-------------|-----|
| Gene | Control | Fold induction | % reduction | |
| Cytokines | IL-1 β | 3.5 | -75 | -84 |
| | TNF α | 3.8 | NS | -54 |
| Chemokines | CCR1 | 1.4 | -51 | -85 |
| | CCR2 | 0.5 | -50 | -92 |
| TLR signaling | MCP-1 | 3.7 | -77 | -91 |
| | MYD88 | 2.6 | -39 | -71 |
| Adhesion molecules | TLR4 | 0.7 | NS | -51 |
| | CD44 | 1.8 | -26 | -39 |
| | VLA-4 | 0.9 | -35 | -61 |

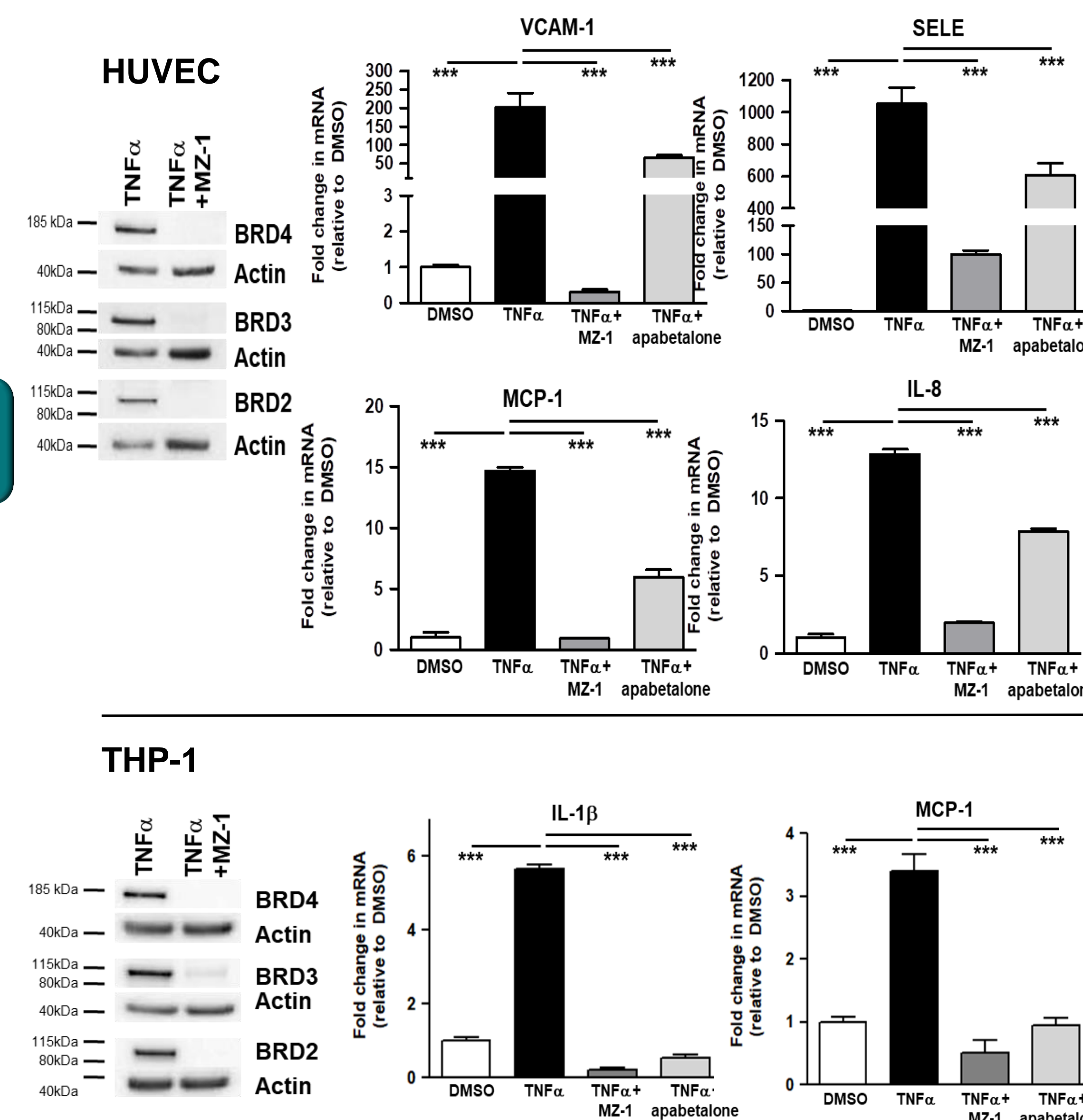
Apabetalone suppresses protein expression of VCAM1 and MCP-1 in endothelial cells.

4 hour TNF α stimulation induced HUVEC VCAM1 surface expression (FACS) and MCP-1 secretion (BD[™] cytometric bead array). Co-treatment with apabetalone reduced VCAM1 and MCP-1 protein levels. Statistical analysis: 1-way ANOVA, Dunnett's Multiple Comparison Test



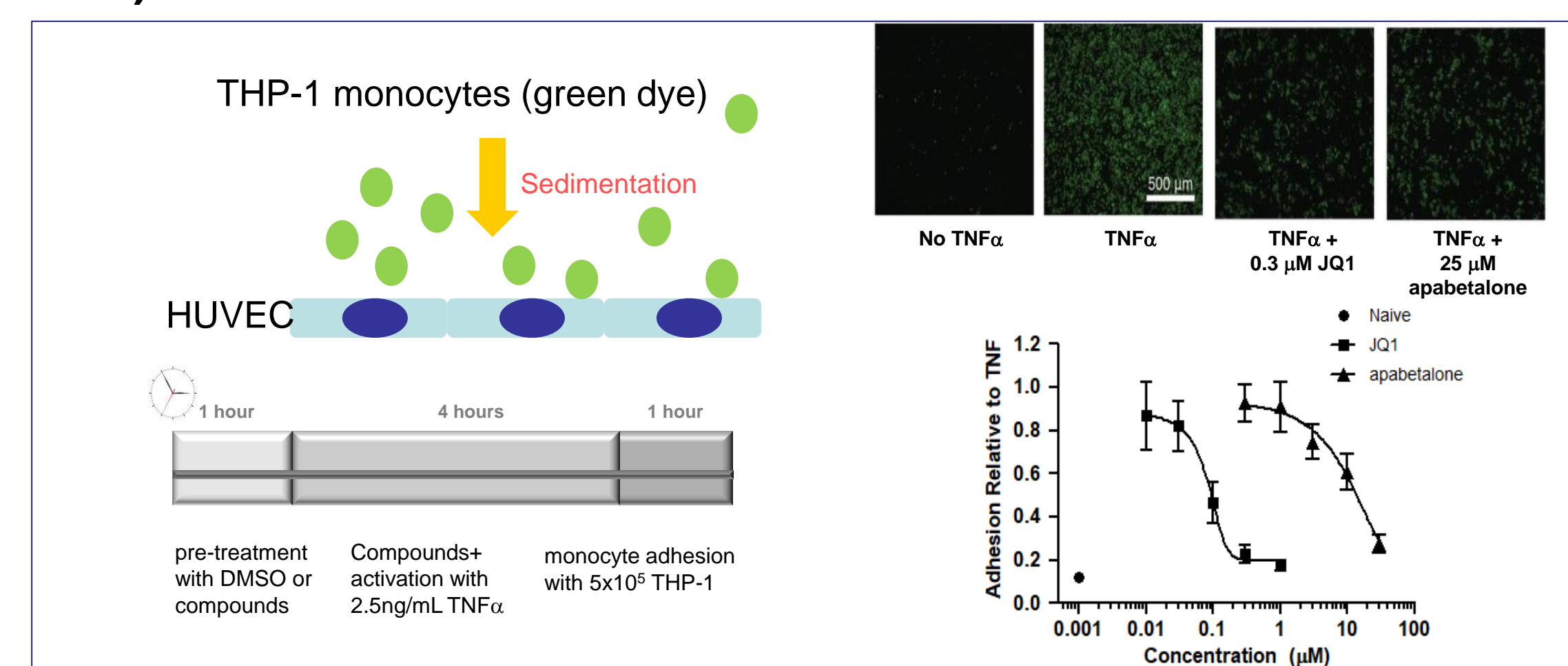
Pro-inflammatory gene expression is BET-dependent

MZ-1 and apabetalone (4hr pre-treatment) prevent TNF α (2hr co-treatment) induction of key pro-inflammatory markers through the degradation or inhibition of BET proteins

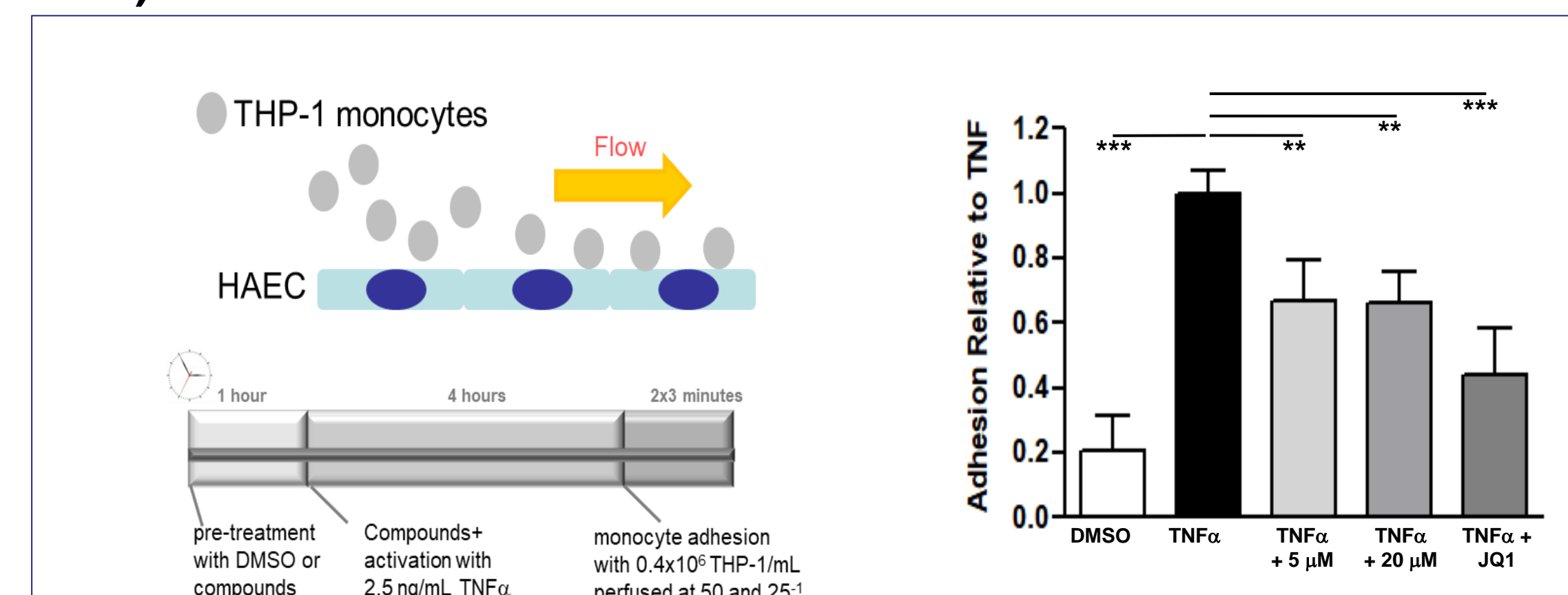


Apabetalone suppressed monocyte THP-1 cell adhesion to inflamed endothelial cells under both static and flow conditions.

1.) Adhesion of THP-1 to HUVEC cells under static conditions



2.) Adhesion of THP-1 to HAEC cells under flow conditions



Apabetalone suppresses pro-atherogenic and plaque rupture mediators in CVD patient plasma

| SOMAscan [®] CVD Patient Plasma Protein Apabetalone (n=47) vs. Placebo (n=47) | | | |
|--|---------------|----------------------|---------|
| Target Gene | Molecule type | % change vs. placebo | p-value |
| VCAM-1 | Adhesion | -12.2 | 0.005 |
| ICAM-1 | Adhesion | -10.9 | 0.006 |
| MMP-3 | Enzyme | -26.8 | 0.005 |
| MMP-12 | Enzyme | -24.6 | 0.003 |

Ingenuity Pathway Analysis (IPA[®]) of SOMAscan[®] CVD Patient Plasma Protein Apabetalone vs. Placebo

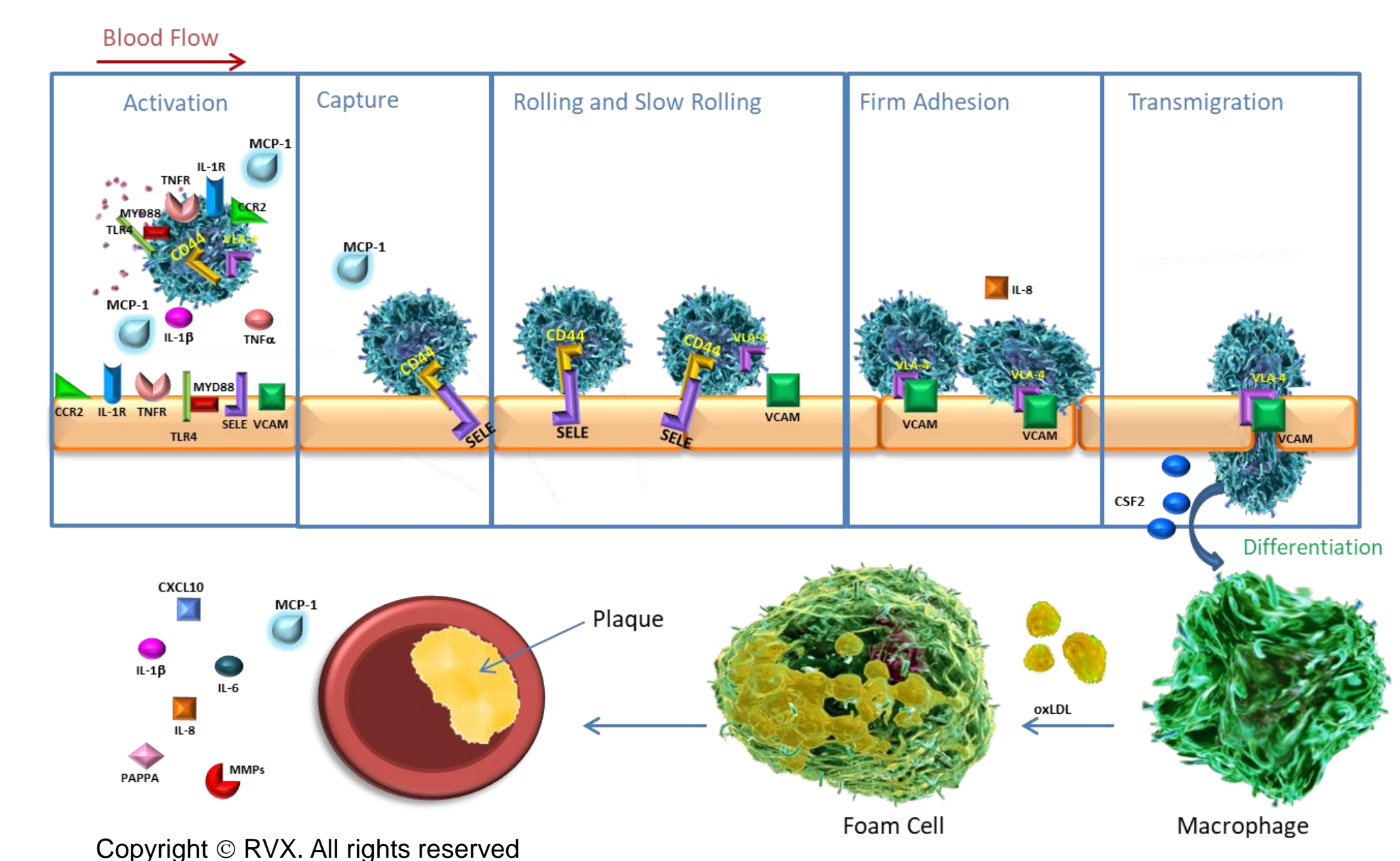
| Upstream Regulator | Molecule type | z-score/direction | p-value |
|--------------------|-------------------------|-------------------|---------|
| IL1B | cytokine | -2.8 | <0.001 |
| IL2 | cytokine | -2.7 | <0.001 |
| CSF2 | cytokine | -2.6 | <0.001 |
| IL6 | cytokine | -2.5 | <0.001 |
| lfn gamma | complex | -2.4 | <0.001 |
| NFKB1B | transcription regulator | -2.4 | <0.001 |
| TNF | cytokine | -2.1 | <0.001 |

Ingenuity Pathway Analysis (IPA[®]) of SOMAscan[®] CVD Patient Plasma Protein Apabetalone vs. Placebo

| Canonical Pathway | z-score | p-value | Contributors |
|-------------------|---------|---------|---|
| HMGB1 Signaling | -1.7 | <0.001 | TLR4, VCAM1, ICAM1, IL5, PTPN11, GRB2, OSM, IL17F, IL17B, TNFRSF11B |

Summary

Apabetalone downregulates pro-atherogenic gene expression in endothelial and monocytic cells.



BET-dependent downregulation of vascular inflammation and subsequent cell adhesion by apabetalone may contribute to the reduction in MACE, a hypothesis currently being tested in the phase 3 cardiovascular outcomes trial BETonMACE in patients with CVD, diabetes mellitus and low HDL-c.

[†]Disclosure: Resverlogix employees received salaries & stock options from RVX.