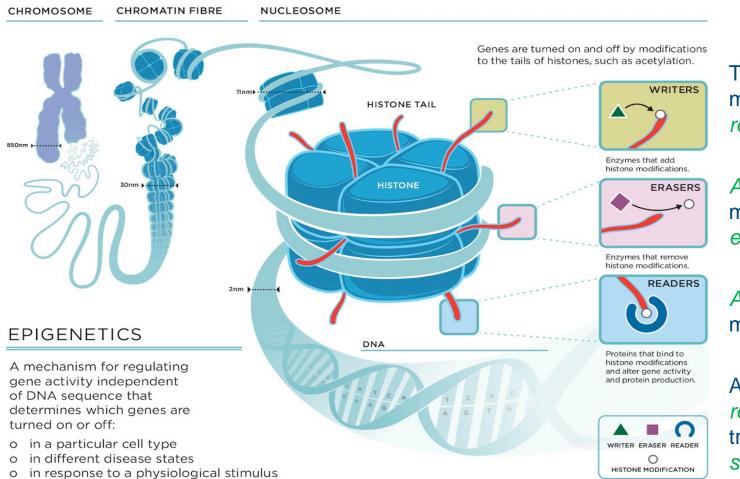
BET protein inhibitor apabetalone suppresses inflammatory hyper-activation of monocytes from patients with cardiovascular disease and type 2 diabetes

Sylwia Wasiak, Ph.D. Resverlogix Corp. AHA 2020



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The *epigenetic code* refers to secondary modifications to chromatin components that *regulate transcriptional activity*

Addition, removal or recognition of these modifications is done by proteins called *writers*, *erasers and readers*

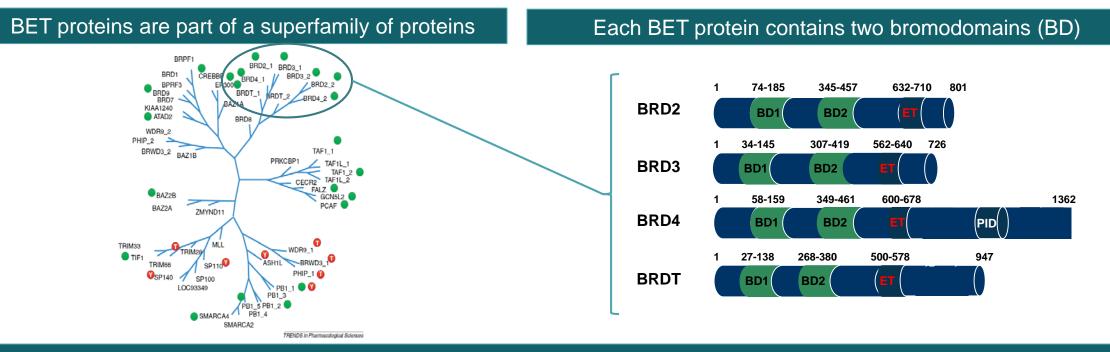
Acetylation of histone lysine residues by writers marks active regions of chromatin

Acetylated lysines on histones are recognized by readers called BET proteins that recruit transcriptional regulatory factors to activate or suppress genes

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Apabetalone (RVX-208) is a Small Molecule Inhibitor that Competitively Inhibits BET Bromodomains

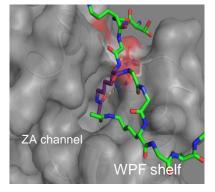


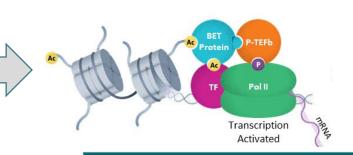


Bromodomains bind acetylated histones and transcription factors to regulate gene transcription

X-ray crystallography

Acetylated lysine (color) bound to bromodomain (grey)



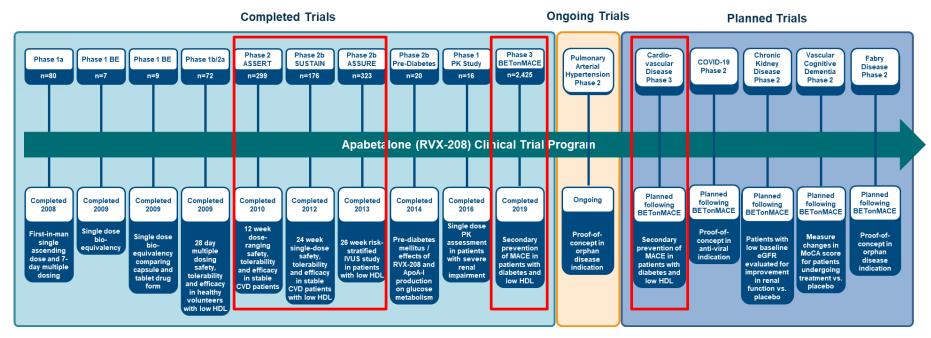


Apabetalone disrupts BD-chromatin binding

Apabetalone in Human Clinical Trials



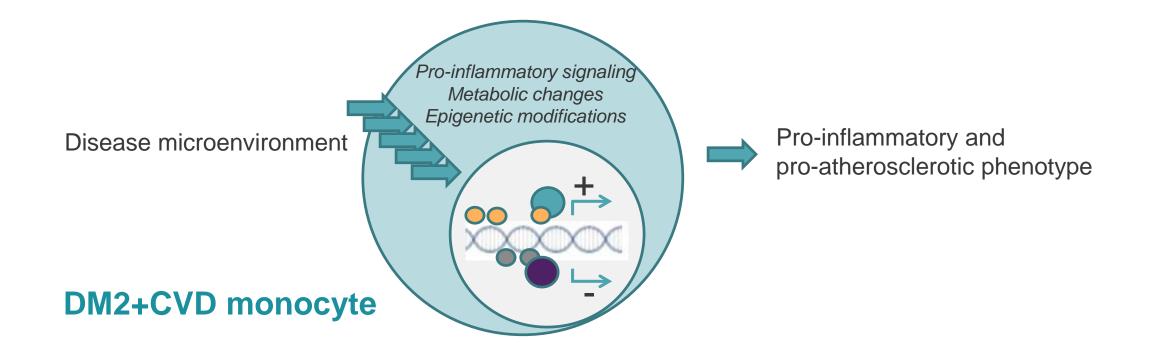
- Apabetalone/RVX-208/RVX000222 (2-(4-(2-hydroxyethoxy)-3,5-dimethylphenyl)-5,7-dimethoxyquinazolin-4(3H)-one) was discovered in 2006.
- Tested in multiple phase 2 trials in CVD patients (endpoints: HDL, ApoA-I elevation)
- Phase 3 cardiovascular event-driven trial BETonMACE
 - Design: Multi-centre, double-blind, randomized, parallel group, placebo-controlled
 - Patients: 2400+ high risk type 2 diabetes with CAD, up to 104 weeks of dosing
 - Results: Apabetalone treatment showed a favorable trend on all cardiac endpoints and reached nominal statistical significance for CHF
 - On February 3, 2020, the FDA granted *Breakthrough Therapy Designation* to apabetalone in combination with top standard of care, including high-intensity statins, for the secondary prevention of MACE in patients with T2DM and recent ACS.
 - A follow-up phase 3 trial BETonMACE2 is currently being planned.



Lowering Monocyte Inflammation In Patients With Type 2 Diabetes and CVD with Apabetalone

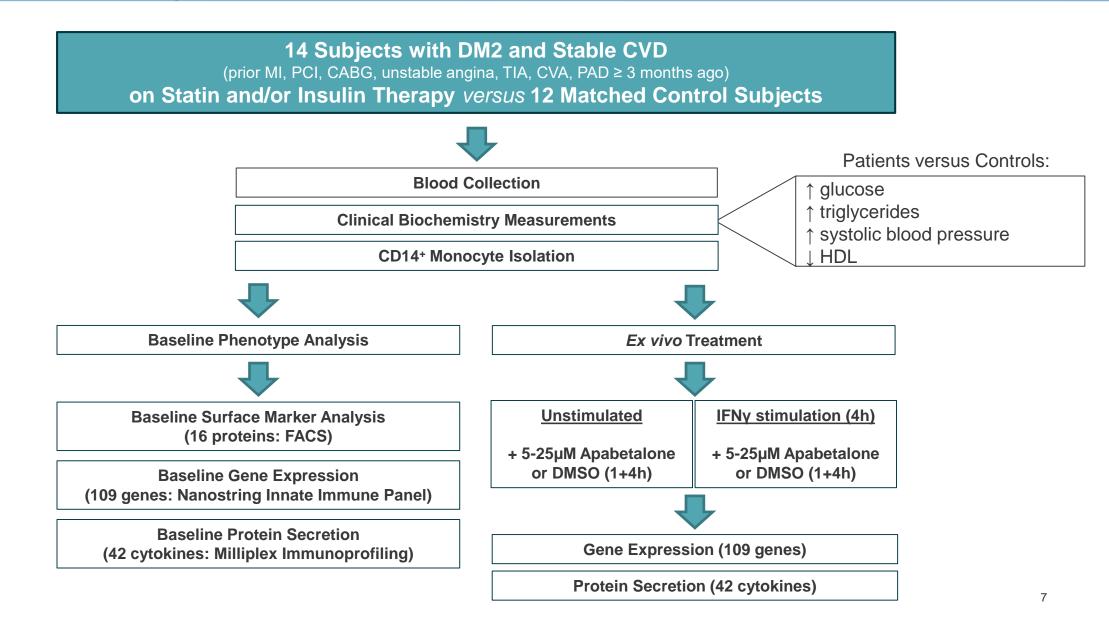


- In diabetes and CVD, microenvironmental factors can trigger pro-inflammatory signaling in monocytes that leads to cytokine production and vascular wall invasion which, in turn, can promote atherosclerosis.
- This "hyper-activation" is partially ascribed to epigenetic reprogramming.
- Hypothesis: Epigenetic modulators such as <u>apabetalone</u> would "correct" the pro-inflammatory hyper-activation of circulating monocytes.



Lowering Monocyte Inflammation In Patients With Type 2 Diabetes And CVD with Apabetalone



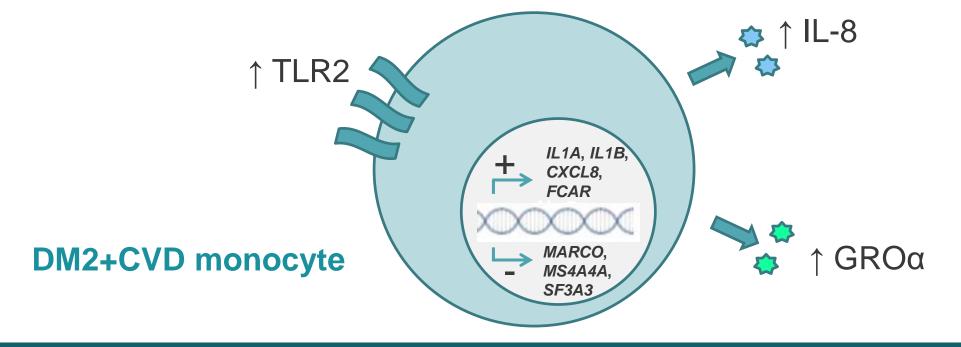


Comparison of Baseline Characteristics of Non-Stimulated DM2+CVD vs. CONTROL monocytes

- Surface marker expression: Pro-inflammatory pattern recognition receptor TLR2 was expressed at higher levels on the surface of DM2+CVD monocytes vs. controls
- Gene expression: <u>Pro-inflammatory</u> genes *IL1A, IL1B, CXCL8 (IL8), FCAR* were upregulated in DM2+CVD monocytes vs. controls, whereas genes associated with an <u>anti-inflammatory</u> phenotype *MARCO, MS4A4A* and *SF3A3* were downregulated

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• **Protein secretion**: Cytokines **IL-8 and GRO**α were <u>secreted at higher levels</u> in DM2+CVD monocytes vs. controls



Monocytes from DM2+CVD patients on SoC therapy have higher expression of pro-inflammatory genes and proteins at baseline indicating *in vivo* pro-inflammatory hyper-activation

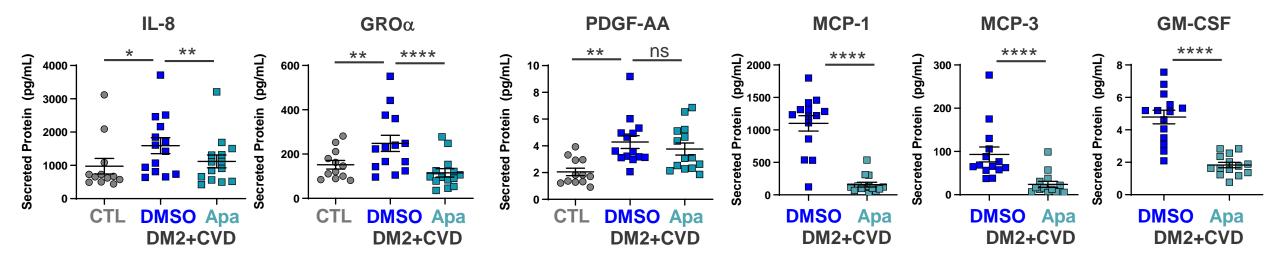
Apabetalone Attenuates Baseline Pro-inflammatory "Hyper-Activation" in Monocytes from DM2+CVD Patients on SoC Therapy



- Gene expression (1+4h): Apabetalone downregulates genes overexpressed in DM2+CVD monocytes: *IL1A, CXCL8 (IL8), FCAR*
- Protein secretion (24h): Apabetalone downregulates secretion of cytokines and chemokines in DM2+CVD monocytes:

 \downarrow IL-8 and GROa (overexpressed in DM2+CVD monocytes)

 \downarrow MCP-1, MCP-3 and GM-CSF



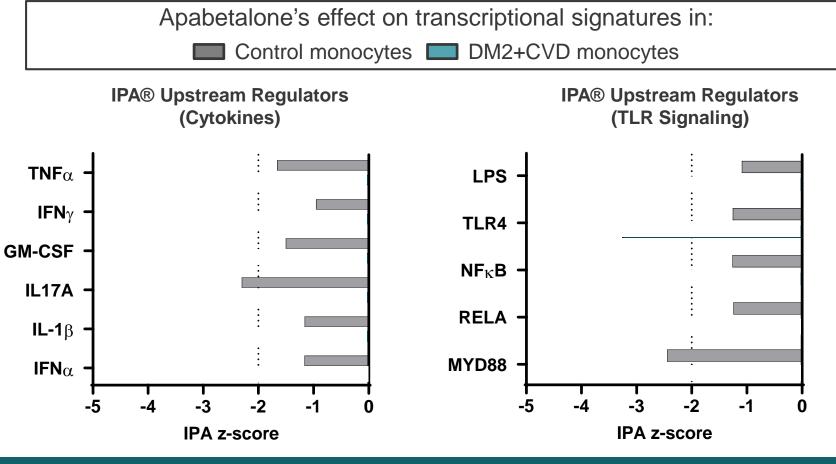
Monocytes from DM2+CVD patients have higher expression of pro-inflammatory genes and proteins <u>at baseline</u>. This "hyper-activation" is attenuated ex vivo by apabetalone treatment.

Apabetalone Downregulates Pro-Inflammatory Gene Signatures More Potently in DM2+CVD Patient Monocytes than in Control Monocytes



1. Nanostring Gene Expression Data: 25µM Apabetalone vs. DMSO

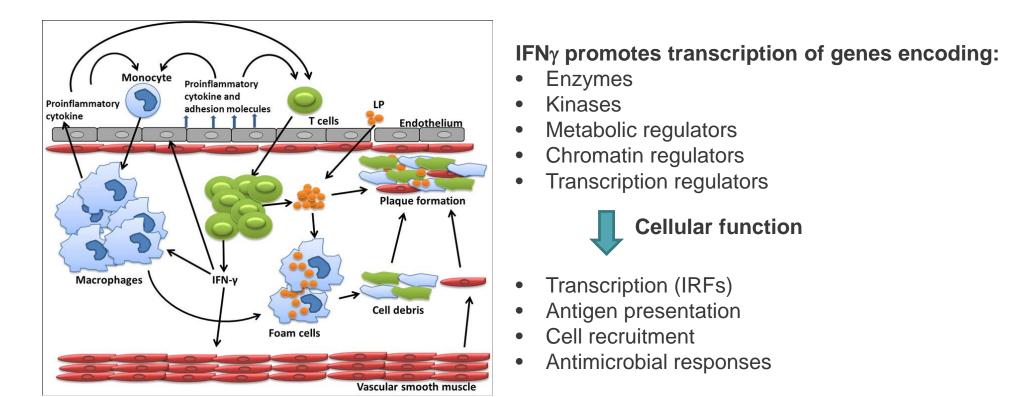
- DM2+CVD monocytes: 53 genes
- Control monocytes: 46 genes (out of 109; >20%Δ; adj. p<0.05)
- 2. Analysis of Transcriptional Gene Signatures using IPA software:
- Gene lists were uploaded into IPA software®
- z-scores determine the transcriptional impact of apabetalone on pro-inflammatory pathways



- Apabetalone treatment causes a more robust suppression of inflammatory gene signatures in DM+CVD monocytes as compared to controls (more negative z-scores).
- This data suggests that transcriptional activity of **BET proteins** is greater in DM2+CVD monocytes than in control monocytes.

IFNγ Differentiates Monocytes into M1 "Classically Activated" Pro-Inflammatory Tissue Macrophages





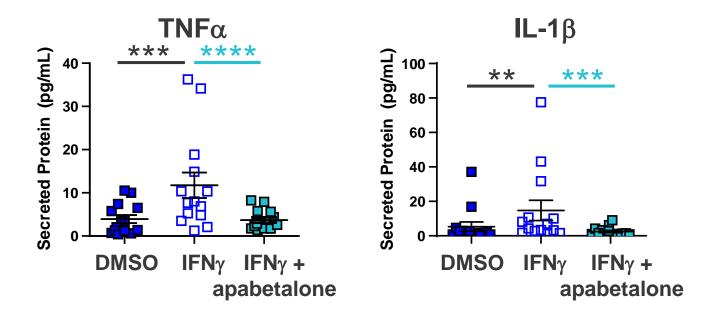
Lin et al., 2013, Adv. Biosci. Biotech.

IFN_γ treatment confers to monocytes a pro-inflammatory M1-like phenotype characterized by enhanced cytokine and chemokine production, phagocytosis, and intracellular killing of microbial pathogens.

Apabetalone Counters Pro-Inflammatory Hyper-Response to IFNγ in DM2+CVD Monocytes

Gene expression (1+4h): Apabetalone downregulates genes hyper-activated by IFNγ in DM2+CVD monocytes encoding cytokines *CCL7, CCL8, TNF* and NF-κB signaling proteins *RELA and MYD88*

Protein secretion (24h): Apabetalone suppressed IFNγ-induced **IL-1β and TNFα secretion** in DM2+CVD monocytes



Apabetalone reduces pro-atherogenic genes and proteins in IFNγ stimulated monocytes from DM2+CVD patients.

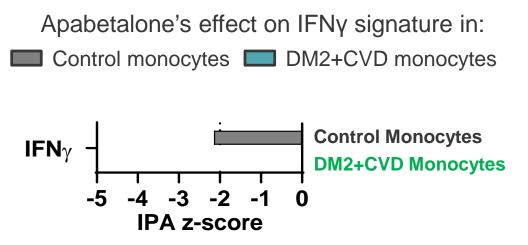
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Gene expression: Apabetalone robustly downregulates IFNγ targets in DM2+CVD monocytes

Gene	IFNγ 4h	IFNγ+Apa	Gene	IFNγ	IFNγ+Apa
Chemokines			Pattern Recognition Signaling		
CCL2	2.2	-91%	TLR8	6.8	-68%
CCL7	1.75	-90%	LY96	2.1	-67%
CXCL1	0.38	-88%	TLR1	1.5	-60%
CCL8	135	-85%	FPR2	1.67	-58%
CXCL9	297	-80%	TICAM2	2.7	-44%
CCR1	1.4	-79%	RELA	1.43	-42%
CXCL10	755	-61%	MYDD88	1.41	-40%
oxLDL Receptor			ROS Production		
MSR1	3.36	-79%	CYBB	1.77	-42%

Pathway analysis: IFNγ transcriptional signature is **preferentially inhibited** by apabetalone **in DM2+CVD monocytes**



Note: IPA z-score<-2 predicts pathway downregulation

Statistics: Two-Way RM ANOVA, Tukey's test. Significant p<0.05

Cytokine-stimulated DM2+CVD monocytes are more sensitive to BET inhibition by apabetalone than control monocytes, indicating that aberrant pro-inflammatory gene transcription is BET dependent in diseased cells.



- Monocytes from DM2+CVD patients exhibit pro-inflammatory hyper-activation <u>at baseline</u>.
- Monocytes from DM2+CVD patients are hyper-responsive to IFNγ upon ex vivo stimulation.
- This pro-inflammatory hyper-activation indicates that **diseased monocytes are "primed"** to **produce pro-inflammatory molecules** in patients which may contribute to disease progression.
- Apabetalone attenuates monocyte hyper-activation by downregulating key inflammatory genes and secreted cytokines in both non-stimulated and stimulated cells.
- Pro-inflammatory gene transcription is more sensitive to BET inhibitor treatment in monocytes from DM2+CVD patients than control monocytes, indicating that BET proteins are driving maladaptive gene expression in a diseased state.
- Findings support the development of apabetalone as a therapy for high risk CVD patients with epigenetic dysregulation of the innate immune response.

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