Epigenetic BET Reader Inhibitor Apabetalone (RVX-208) Counters Proinflammatory Aortic Gene Expression In a Diet-Induced **Obesity Mouse Model**

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BACKGROUND

Obesity is associated with insulin resistance, dyslipidemia, high blood pressure, chronic inflammation and it increases the risk of type 2 diabetes (DM2) and cardiovascular disease (CVD) in patients. Bromodomain and extraterminal (BET) proteins enhance the expression of proinflammatory genes in DM2 and CVD models by recruiting the transcription regulation factors. Apabetalone is a small molecule that inhibits BET interactions with chromatin, leading to anti-inflammatory and anti-atherogenic effects.

OBJECTIVE: Here we evaluate the effects of apabetalone on vascular inflammation in a mouse model of diet induced obesity.

METHODS

From 8 to 30 weeks of age, C57 BL/6J mice were fed a low-cholesterol high-fat (HFD) or low-fat diet (LFD). Serum glucose, GTT and ITT were measured every 6 weeks. Mice received apabetalone at 150 mg/kg b.i.d between 14 and 30 weeks. Gene expression was analyzed post necropsy in aortic tissue by PCR and nCounter® Inflammation Panel.

RESULTS

Metabolic dysregulation in the DIO model: As previously published, HFD induced weight gain, visceral obesity, high fasting blood glucose, glucose intolerance and insulin resistance compared to LFD controls (not shown), confirming metabolic dysregulation. Apabetalone treatment did not impact metabolic parameters measured.

Gene expression profiling in the aorta: Analysis showed an upregulation of 27 inflammatory genes in HFD-fed mice as compared to LFD, including transcription factors *Rela* (22%), *Hif1a* (25%) and *Tcf4* (44%) (p<0.05) which are associated with DM2 risk. Differentially expressed genes map to cytokine signaling, cytoskeletal remodeling, coagulation and complement pathways.

Effects of apabetalone treatment: Apabetalone reduced the expression of proinflammatory transcription factors *Rela* and *Nfkb* as well as the expression of potent chemokines Ccl2 and Ccl8, and the chemokine receptor Ccr2. Genes associated with vascular inflammation were also reduced by apabetalone, including leukocyte receptors Itgam (Cd11b) and Ccr2, as well as endothelial receptors Sele and Icam1.

CONCLUSION

HFD induces vascular inflammation in mice. Apabetalone treatment diminishes this proinflammatory phenotype, providing mechanistic insight into how apabetalone reduces CVD risk in DM2 patients in clinical trials.

8 weeks 14 weeks Vehicle Low-fat (LFD) or High-fat (HFD) **APABETALONE** 150 mg/kg b.i.d. Diet Gene expression (mouse aorta) HFD (n=7) LFD (n=6) Gene CCIS TRE7 TRADD PTGFRI C10A GAPDH C3AB1 CCL2CE TLRS TREL TSID PIPK1 CCP2TLRA N FRB 1 MF F 2D TCF4 CCL11 CXCR4 SMAD7 GRB2 TL18 CSF3 CEBFASLG MAPK1 CCL7NOD 2 STAT1CRP PTGS1 DAXX TLR2 C1SCD163 OASL CD 40 CD86 CCL24 I.TATGFB1PRKCA TREM2 MAEG TT.1 Δ (대민)

Gene Set Enrichment Analysis (GSEA) Red: Upregulated Blue: Downregulated

APABETALONE INHIBITS EPIGENETIC ACTIVITY OF BET PROTEINS FIG 1: THE DIET INDUCED OBESITY (DIO) MOUSE MODELS HUMAN CHARACTERISTICS OF OBESITY, PRE-DIABETES AND DYSLIPIDEMIA Apabetalone is a clinical-stage small molecule that showed a trending reduction in CV events in high-risk CVD and DM2 patients. It binds **Pharmacological intervention with APABETALONE** competitively to bromodomains in BET histone acetylation "readers", (clinical-stage BET inhibitor) causing their release from chromatin, and downregulation of BETsensitive genes with proinflammatory and pro-atherosclerotic activities. 20-21 weeks 30 weeks Aorta dissection Tests: BET: bromodomain and extraterminal proteins; ac: acetylated Fasting glucose Gene expression with lysine residue on DNA-associated proteins; BD: bromodomain; **DIO Characteristics:** TF: transcription factor Glucose tolerance Nanostring Inflammation Increased adiposity

Insulin resistance

Panel (254 genes)

Glucose intolerance Insulin resistance

FIG 2: HFD INDUCES PROINFLAMMATORY AORTIC GENE EXPRESSION

HFD enhances the expression of pro-inflammatory gene sets in the mouse aorta

GENE SETS (GSEA) Upregulated in HFD-fed mice	NES	Nominal p-value	FDR q-value
HALLMARK_INTERFERON_GAMMA_RESPONSE	1.43	0.019	0.364
HALLMARK_INTERFERON_ALPHA_RESPONSE	1.41	0.038	0.210
HALLMARK_COMPLEMENT	1.39	0.042	0.167
HALLMARK_ALLOGRAFT_REJECTION	1.38	0.033	0.135
HALLMARK_INFLAMMATORY_RESPONSE	1.34	0.117	0.170
HALLMARK_APOPTOSIS	1.34	0.089	0.142
HALLMARK_PI3K_AKT_MTOR_SIGNALING	1.28	0.200	0.182
HALLMARK IL6 JAK STAT3 SIGNALING	1.25	0.160	0.189

HFD enhances the expression of proinflammatory genes in the mouse aorta



HFD enhances the expression of key proinflammatory transcription factors











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LFD HFD HFD APS





DISCLOSURE INFORMATION

LED HED AD