

Wasiak et al. Table S1 (Supporting Figure 1)

Cytokine-induced gene expression in hCMEC/D3 cells is reduced by BET protein inhibition after 24h of treatment.

Target Name		TNF $\alpha$ + IFN $\gamma$	TNF $\alpha$ + IFN $\gamma$ + Apabetalone	
Protein	Gene	Fold Induction*	% Inhibition†	p-value‡
MCP-3	<i>CCL7</i>	149	96	<0.0001
Fractalkine	<i>CX3CL1</i>	964	89	<0.0001
MCP-1	<i>CCL2</i>	15	66	<0.0001
RANTES	<i>CCL5</i>	593	83	<0.0001
IL-6	<i>IL6</i>	66	81	<0.0001
IL-8	<i>CXCL8</i>	32	60	<0.0001
IP-10	<i>CXCL10</i>	14073	58	<0.0001
GM-CSF	<i>CSF2</i>	17	89	<0.0001

\*mRNA fold induction in response to 24h cytokine treatment (100ng/mL) was calculated relative to cytokine-naïve cells treated with vehicle for the same amount of time (0.05% DMSO).

†Gene expression inhibition was calculated relative to the induced state in cells co-treated with cytokines and 25 $\mu$ M apabetalone.

‡Statistical significance was calculated with one-way ANOVA with Dunnett's correction. ns, non-significant, ns: non-significant, n=3.

Wasiak et al. Table S2 (Supporting Table 3)

Pharmacokinetic distribution of apabetalone in mouse plasma and brain tissue.

Tissue*	Mean Concentration ( $\mu\text{M}$ )
	Apabetalone
Brain	2.4
Plasma	32
Brain to Plasma Ratio	0.07

\*Plasma and saline-perfused brains were collected from naïve mice at 3h post single oral dose of 150mg/kg apabetalone, n=3.