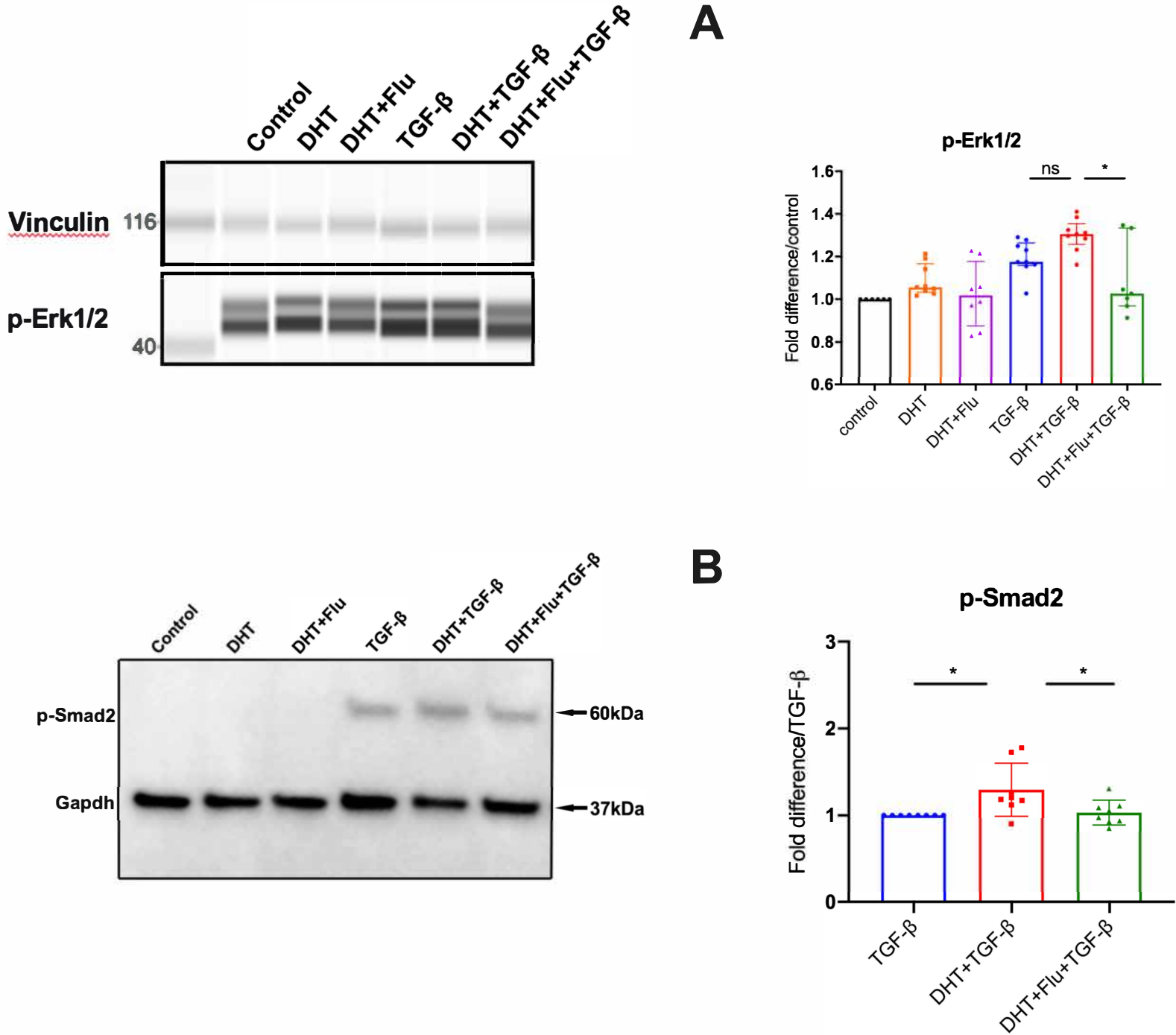


Supplemental Material

Figure S1. p-Erk1/2 and p-Smad2 signaling in female *Fbn1*^{c1039G/+} aortic root/ascending derived smooth muscle cells following hormone treatment.



A phosphorylated-Erk1/2 (p-Erk1/2) and **B** phosphorylated-Smad2 (p-Smad2) in aortic root/ascending (ASC) aorta derived SMC from *Fbn1*^{C1039G/+} (*Fbn1*) female mice treated with dihydrotestosterone 10nM (DHT), DHT + flutamide 1 μ M (Flu), TGF- β 5ng/mL, TGF- β + DHT, TGF- β + DHT + Flu or vehicle control (n=8-9 replicates for each group). Relative protein level quantified using vinculin or Gapdh as loading control as indicated. Values expressed as fold difference compared to control for p-Erk1/2 and fold difference compared to TGF- β for p-Smad2. Results presented as median +/-interquartile. Kruskal-Wallis test with Dunn's posttest was used for multiple comparison among the groups: control vs. DHT vs. DHT + Flu, TGF- β vs. TGF- β + DHT vs. TGF- β + DHT + Flu. * $p \leq 0.05$.