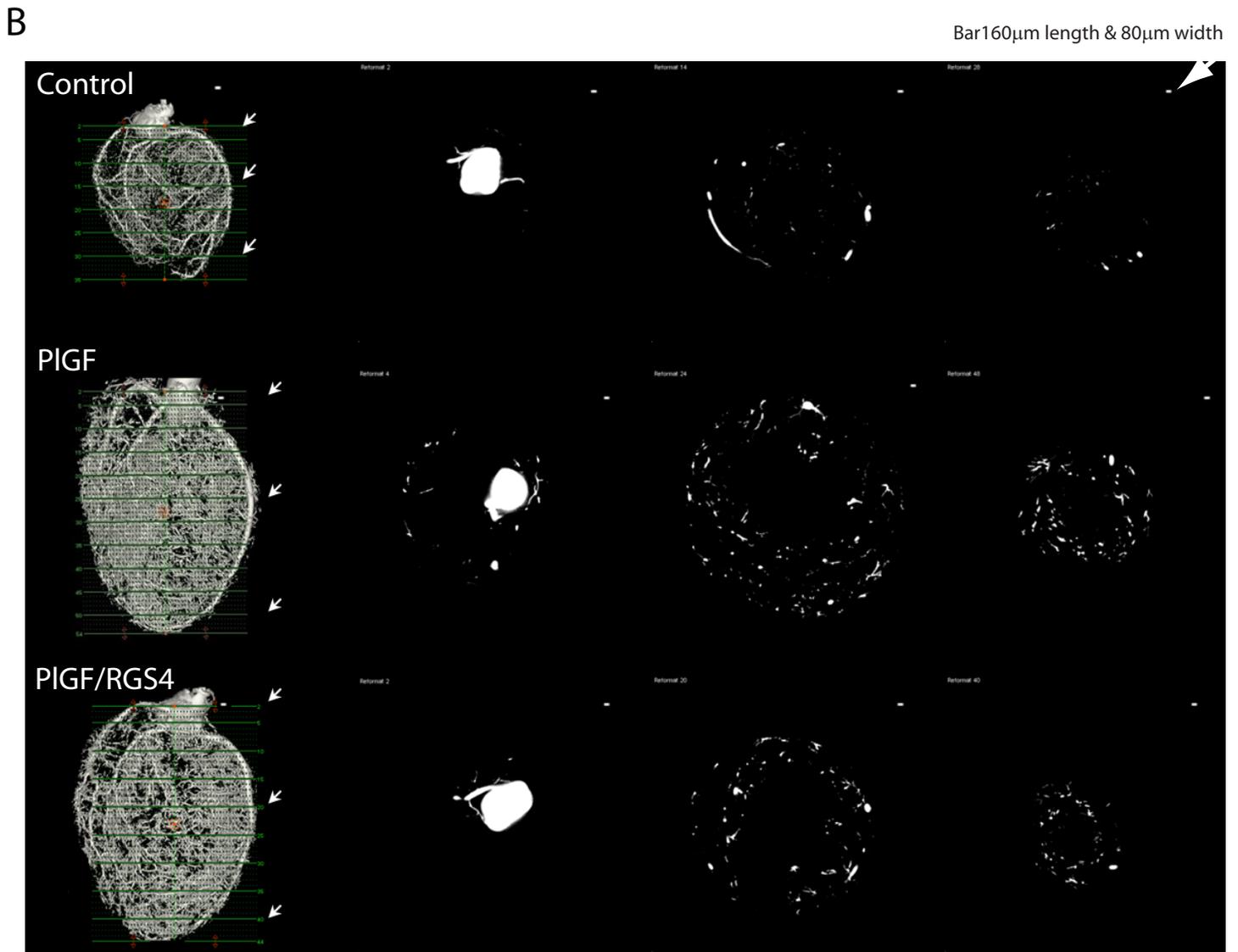
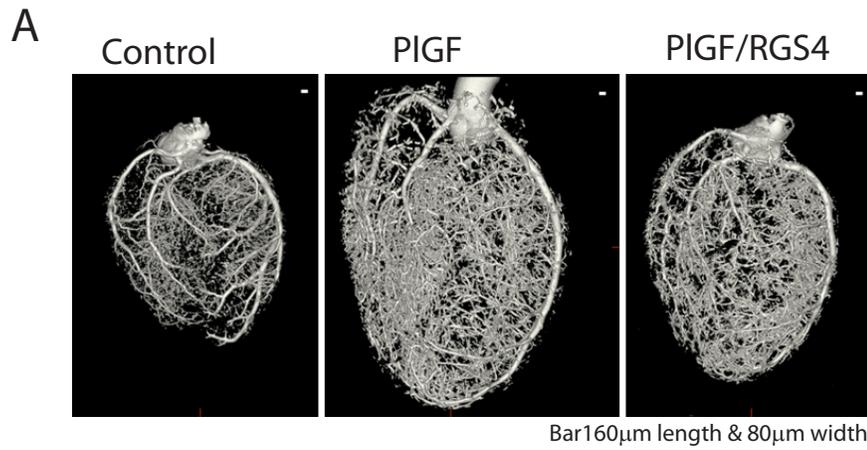
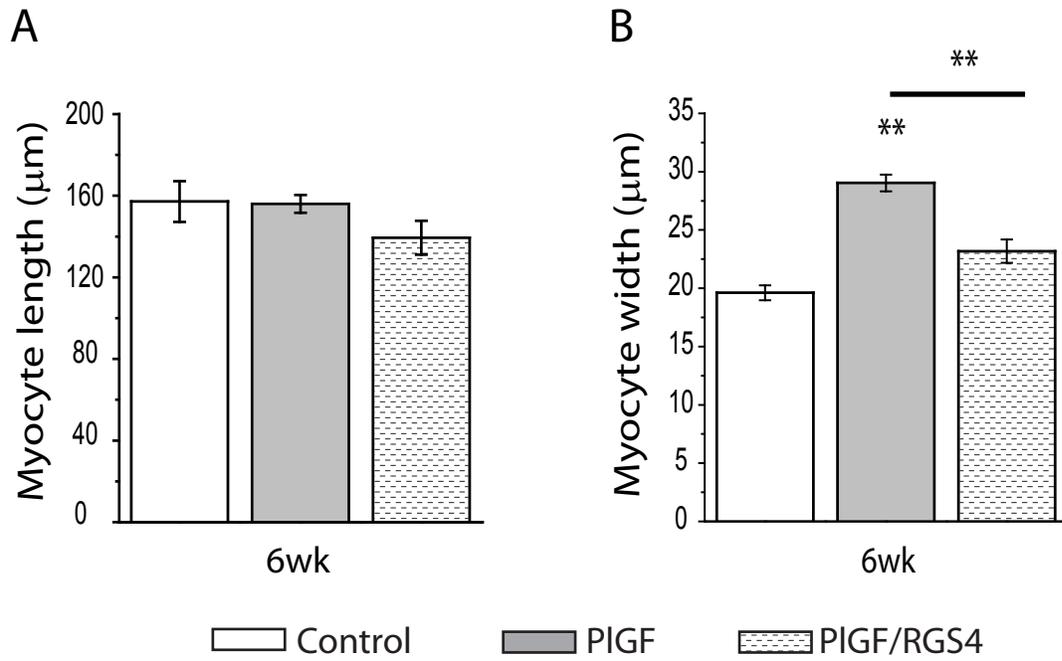


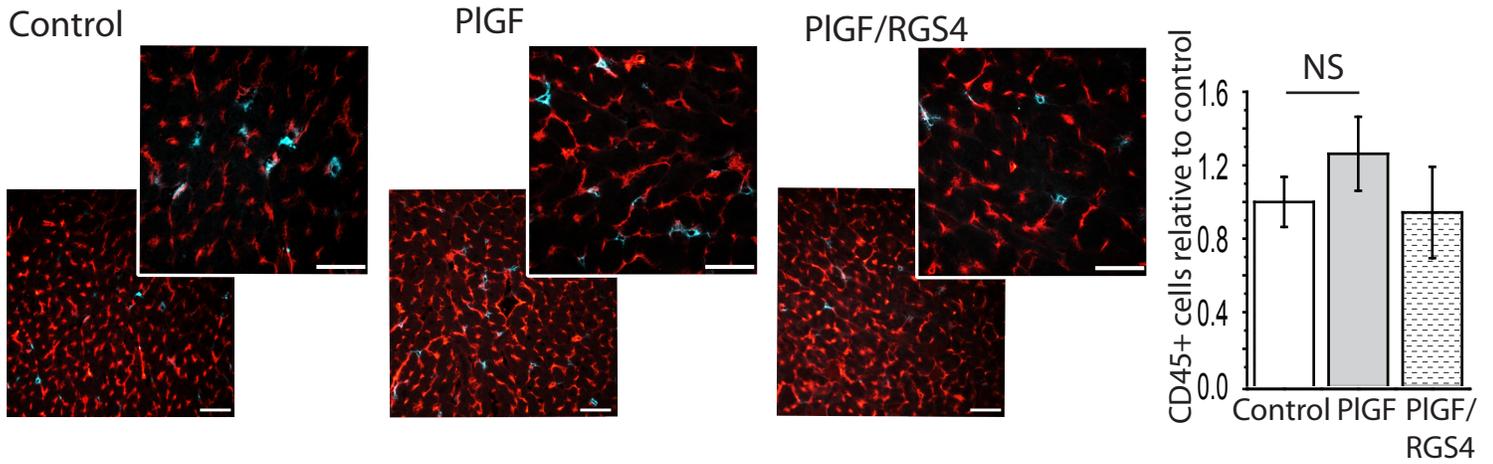
**Supplementary Figure 1.** PIGF and PIGF/RGS4 inducible mouse models. (A) Design of PIGF/LacZ and PIGF/RGS4 inducible bidirectional tetracycline responsive (TRE-promoter) constructs. (B) TRE-PIGF/LacZ and TRE-PIGF/RGS4 induced expression in MEF/3T3 Tet-OFF cells in the absence of doxycycline (western blot analysis for PIGF and RGS4 expression). (C) PIGF and PIGF/RGS4 transgene level in PIGF and PIGF/RGS4 mice, respectively, as determined by real-time PCR based genotyping (Transnetyx, Inc.); n=50 mice/group. Note similar level of transgene relative to the housekeeping gene indicating a relatively equal transgene copy number for both conditional mouse models. (D) Representative example of PIGF induced mice: upper panel before induction (Tet-ON); lower panel after 6 weeks induction (Tet-OFF). Note homogenous expression of LacZ transgene as demonstrated by X-Gal staining (blue) of the entire heart in Tet-OFF and no staining in Tet-ON. (E) Increased PIGF mRNA expression at 3 and 6 weeks in PIGF and PIGF/RGS4 mice compared with control mice, as determined by RT-PCR. (F) Increased PIGF expression at 6 weeks in PIGF and PIGF/RGS4 mice (western blot analysis). (G) Increased RGS4 transgene mRNA expression in PIGF/RGS4 mice at 3 and 6 weeks compared with controls, as determined by qPCR; n=6 mice/group. (H) Heart weight/Body weight ratio and (I) LV mass/Body weight ratio in PIGF mice (littermates of PIGF mice induced by doxycycline withdrawal) continuously fed with doxycycline including the 6 weeks stimulation period. n=4 mice/group. NS - not significant; \* $p$ <0.05, \*\* $p$ <0.01 vs. control.



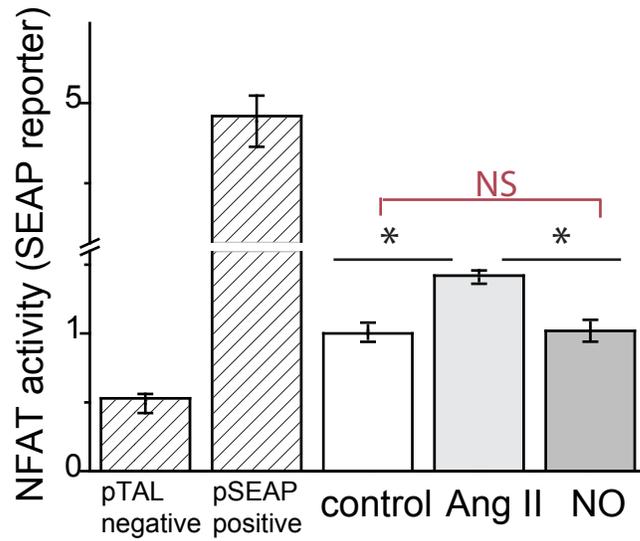
Supplementary Figure 2. Cardiac vasculature assessed by micro-CT. (A) Micro-CT angiograms after 6 weeks of PIGF or PIGF/RGS4 transgene expression compared with control. (B) Homogenous distribution of vascular structures in the myocardium in 2D microCT sections in PIGF and PIGF/RGS4 mice at 6 weeks. Representative images at three comparable levels.



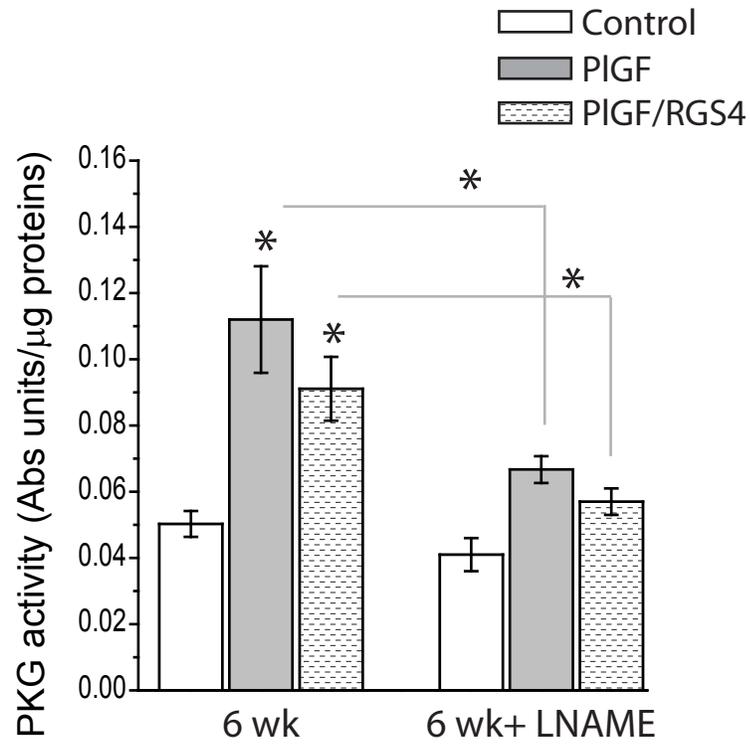
Supplementary Figure 3. Myocyte length (A) and width (B) characteristics in PIGF and PIGF/RGS4 mice at 6 weeks. Note the increase in myocyte thickness in PIGF mice as compared with PIGF/RGS4 or control mice. n=80-180 per group; \*\* $p < 0.01$  vs. control or as indicated.



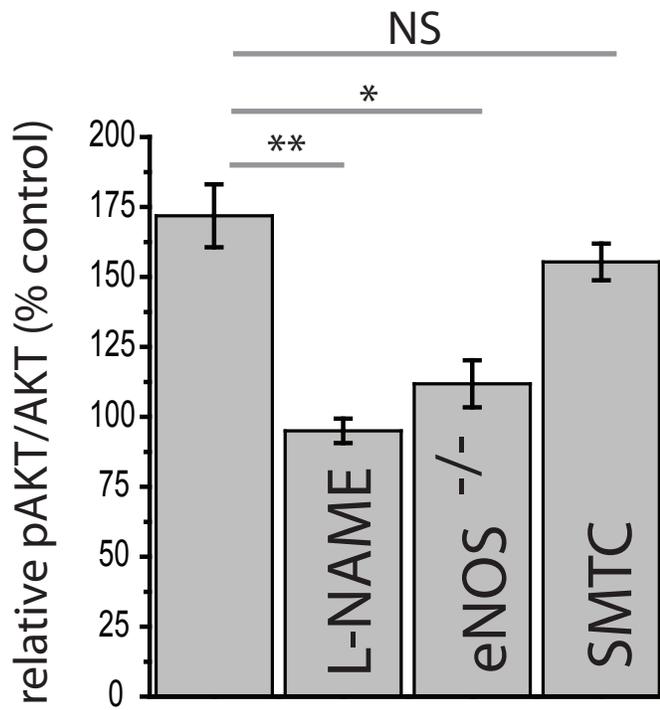
Supplementary Figure 4. CD45<sup>+</sup> cell recruitment in the heart after 6 weeks of transgene stimulation. Representative immunostaining of LV myocardium with specific antibodies against CD45 and CD31. Bar 50  $\mu$ m. Note no recruitment of CD45<sup>+</sup> cells in the heart in PIGF or PIGF/RGS4 mice at 6 weeks compared with control mice (histogram). NS- not significant.



Supplementary Figure 5. NFAT promoter activity determined with SEAP (secreted alkaline phosphatase) reporter in NO-treated versus AngII-treated or untreated cells, after 24 hr. \* $p < 0.05$ , vs. control or NO-treated cell. NS- not significant.



Supplementary Figure 6. PKG activity in PIGF and PIGF/RGS4 mice at 6 weeks with and without L-NAME treatment. n=6-14 per group; \* $p < 0.05$  vs. control or as indicated.



Supplementary Figure 7. The level of Akt phosphorylation at Ser 473 determined as pAkt /Akt ratio in western blots in PIGF mice: untreated, L-NAME-treated, eNOS<sup>-/-</sup> background and SMTC treated, relative to their appropriate controls. n=4-6 per group. \* p<0.05, \*\* p<0.01.

**Supplementary Table 1.** Heart rates for untreated, L-NAME-treated, SMTC-treated and eNOSKO mice. No significant differences were found between groups.

	<b>Heart Rate</b>	
Control	463.4 $\pm$ 18.04	n=24
PIGF	495.5 $\pm$ 13.25	n=24
L-NAME-treated control	493.7 $\pm$ 22.07	n=9
L-NAME-treated PIGF	500.9 $\pm$ 20.68	n=8
eNOSKO control	423.7 $\pm$ 22.75	n=7
PIGF-eNOSKO	421.8 $\pm$ 44.32	n=5
SMTC-treated control	449.2 $\pm$ 18.62	n=5
SMTC-treated PIGF	454.4 $\pm$ 41.02	n=7