Supplemental Figure 1. Experimental design for in vivo vascular disease models. ApoE-KO mice were employed in studies designed to investigate: a) aortic vascular disease burden and aneurysm formation, and b) vascular remodeling in a vein graft model characterized by the development of neointimal hyperplasia.

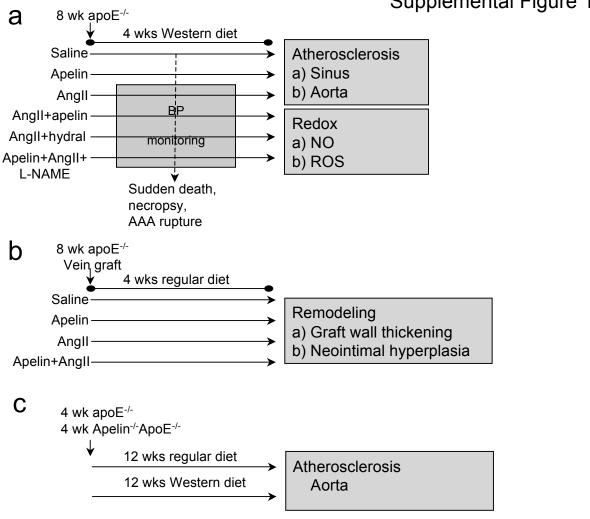
Supplemental Figure 2. Apelin inhibits native atherosclerosis in the ApoE null model. Apelin (2mg/kg/day) and AngII (1.4mg/kg/day) were administered by osmotic minipumps to ApoE-KO mice on a high fat diet. Relative area of vascular disease in the aortic root was significantly reduced in apelin treated mice (\*P<0.05 compared to saline). Black bar represents 200 $\mu$ m.

Supplemental Figure 3. Apelin suppresses disease-related  $O_2^{\bullet-}$  production in vein grafts. Cellular  $O_2^{\bullet-}$  production in vein grafts, measured by 2-hydroxyethidium (DHE) fluorescent nuclear staining in ApoE-KO mice was assessed in the endothelial and neointimal layers of the vessel wall. There was no difference in  $O_2^{\bullet-}$  production in the endothelium among the different conditions.  $O_2^{\bullet-}$  production was attenuated in the neointima in apelin treated mice even when coadministered with AngII (54%, \*P<0.05, apelin vs. saline; and 48%, †P<0.05, AngII+apelin vs. AngII alone). Vein graft vessel wall exhibits green autofluorescence. White arrows denote DHE staining, white arrowheads denote neointima, white bar represents 5  $\mu$ m.

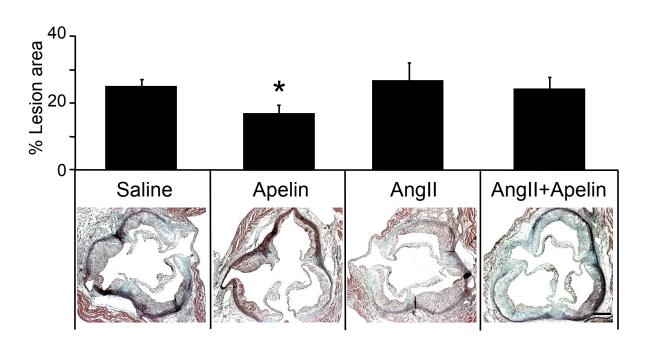
Supplemental Figure 4. Apelin does not compete with Angl I for binding to the primary Angl I receptor AT1R. a) FAM-labeled

- AngII (10 nM) binds to cells transfected with AT1R (blue curve) but not to cells with control vector transfection (red curve).
- **b)** Unlabeled AngII (100 nM) inhibits binding of FAM-AngII (10 nM) to AT1R (blue curve), compared to FAM-AngII (10 nM) in the absence of unlabeled AngII (red curve).
- **c)** Unlabeled apelin (100 nM) does not compete with FAM-AngII for binding to AT1R (blue curve) and shows no difference to FAM-AngII (10 nM) binding in the absence of apelin (red curve). The x-axis represents fluorescence intensity.

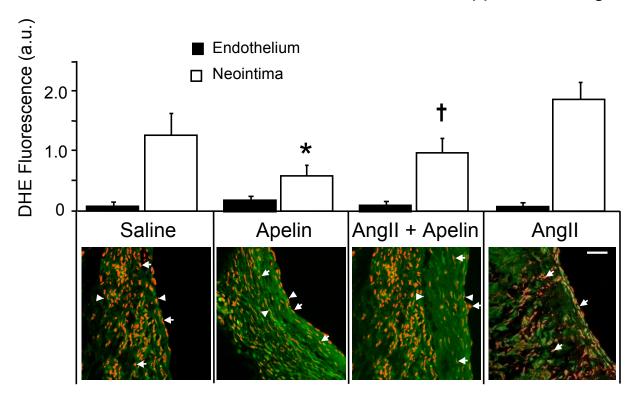
### Supplemental Figure 1



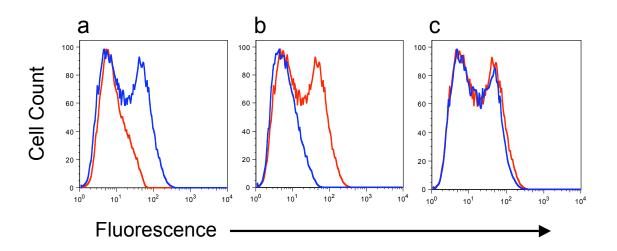
#### Supplemental Figure 2



## Supplemental Figure 3



## Supplemental Figure 4



	Saline	Apelin	Ang+apelin	Angli
Total Cholesterol	1201 <u>+</u> 167	1095 <u>+</u> 93	1236 <u>+</u> 146	1111 <u>+</u> 121
HDL-Cholesterol	105 <u>+</u> 12	86 <u>+</u> 8	99 <u>+</u> 9	108 <u>+</u> 11
Triglycerides	223 <u>+</u> 112	211 <u>+</u> 102	191 <u>+</u> 97	209 <u>+</u> 126
Body Weights	27 <u>+</u> 2.4	24 <u>+</u> 3.0	23 <u>+</u> 4.6	27 <u>+</u> 2.9

# Supplemental Table 1. - Lipid Profiles and Body Weights of Experimental Animals.

Serum lipid values are expressed in mg/dl, and body weights in grams (statistical analysis with ANOVA).