

Table S1. Summary of outcome assessments comparing circulating biomarker concentration between groups in unadjusted, and adjusted LME analyses.

	Unadjusted analyses			Adjusted analyes*						
Parameter assessed	β (std error) at 3 Weeks	β (std error) at 24 weeks	p-value	β (std error) at 3 Weeks	β (std error) at 24 weeks	p-value				
Quality assessment										
Triglycerides+	0.277 (0.050)	0.203 (0.050)	< 0.001	0.278 (0.050)	0.204 (0.050)	< 0.001				
eGFR	12.596 (4.369)	11.727 (4.361)	0.006	12.589 (4.395)	11.722 (4.386)	0.006				
Homocysteine†	-0.243 (0.041)	-0.269 (0.041)	< 0.001	0.243 (0.041)	-0.268 (0.041)	<0.001 <sup>‡</sup>				
Primary outcome measures										
OPN+	0.047 (0.064)	-0.022 (0.064)	0.554	0.048 (0.064)	-0.021 (0.064)	0.551				
Kallistatin	0.490 (0.419)	0.324 (0.416)	0.491	0.488 (0.419)	0.322 (0.416)	0.494				
Secondary outcome measures										
MMP9†	0.082 (0.072)	0.085 (0.072)	0.406	0.082 (0.072)	0.085 (0.072)	0.405				
Resistin†	-0.033 (0.054)	-0.028 (0.054)	0.805	-0.032 (0.054)	-0.028 (0.054)	0.809				
OPG	0.020 (0.031)	0.066 (0.031)	0.095	0.019 (0.031)	0.066 (0.031)	0.095				
Plasma D-dimer† <sup>§</sup>	0.070 (0.074)	-0.007 (0.074)	0.528	0.070 (0.074)	-0.007 (0.074)	0.524				
CRP†§	0.296 (0.166)	0.119 (0.165)	0.202	0.304 (0.166)	0.122 (0.165)	0.186				
Total cholesterol†§	0.059 (0.023)	-0.000 (0.023)	0.011‡	0.060 (0.023)	0.000 (0.023)	0.014 <sup>‡</sup>				
HDL-C+	-0.009 (0.029)	0.021 (0.029)	0.576	-0.009 (0.029)	0.021 (0.029)	0.579				
LDL-C†§	0.034 (0.037)	-0.046 (0.038)	0.110	0.034 (0.037)	-0.046 (0.038)	0.108				
VLDL-C†	0.270 (0.052)	0.166 (0.053)	<0.001	0.271 (0.052)	0.166 (0.053)	<0.001				

Beta coefficients refer to the interaction of time and treatment allocation (using patients allocated fenofibrate as a reference) for that time point.

P-values relate to the interaction between time and treatment allocation over the whole 24 weeks.\* Model adjusted to include history of CHD and prescription for statins, antiplatelet medications and anticoagulants. † Data transformed using natural log to conform to LME assumptions. †

P-values from sub-set analysis excluding outliers. § Data normalized to baseline levels to account for initial differences between groups.

Table S2. Summary of models assessing differences in AAA growth between treatment groups using unadjusted and adjusted LME analyses.

	Unadjusted a	nalyses	Adjusted analyses*						
AAA measurement	β (std error) at 24 weeks	p-value	β (std error) at 24 weeks	p-value					
Observer 1									
Inner-wall to inner wall	0.435 (0.398)	0.277	0.432 (0.398)	0.279					
Leading edge to leading edge	0.324 (0.327)	0.322	0.323 (0.327)	0.325					
Outer wall to outer wall	0.228 (0.313)	0.467	0.227 (0.313)	0.470					
Observer 2									
Inner-wall to inner wall	0.078 (0.359)	0.828	0.076 (0.359)	0.832					
Leading edge to leading edge	0.115 (0.342)	0.738	0.113 (0.342)	0.742					
Outer wall to outer wall	-0.029 (0.327)	0.929	-0.031 (0.327)	0.925					

Beta coefficients refer to the interaction of time and treatment allocation (using patients allocated fenofibrate as a reference) for that time point.

P-values relate to the interaction between time and treatment allocation over the whole 24 weeks.\* Model adjusted to include history of CHD and prescription for statins, antiplatelet medications and anticoagulants.