Supplementary Table 1. Baseline characteristics and intermediate phenotypes in those men who went on to develop CHD and those men who remained disease free

	Number of measures	**No CHD N=2751**	**With CHD N=301**	P- Value	OR per tertile
Age (years)		56.0 (3.5)	56.5 (3.6)	0.020	
Smoking (% Current)	6	27.6%	37.5%	<0.001	1.57
BMI (kg/m^2)†	6	26.4 (3.55)	26.9 (3.38)	0.006	1.22
SBP (mmHg)	6	137.6 (19.0)	142.6 (19.7)	<0.001	1.27
DBP (mmHg)	6	84.2 (11.2)	87.0 (11.4)	<0.001	1.30
Cholesterol (mmol/l)	6	5.70 (1.01)	6.05 (1.02)	<0.001	1.43
HDL (mmol/l)	1	1.72 (0.59)	1.60 (0.54)	0.002	0.77
LDL (mmol/l)	1	3.06 (1.01)	3.36 (0.96)	<0.001	1.39
Triglyceride (mmol/l)	6	2.05 (1.26)	2.35 (1.34)	<0.001	1.39
ApoB (g/l)	1	0.89 (0.26)	0.95 (0.23)	<0.001	1.53
ApoAI (g/l)	1	1.64 (0.32)	1.61 (0.29)	0.151	0.89
Homocysteine µmol/l	1	12.75 (5.06)	13.50 (5.65)	0.977	1.13
Folate µmol/l	1	233.6 (64.6)	242.0 (70.1)	0.155	1.18
Lp-PLA2 nmol/min/ml	1	49.4 (16.0)	50.6 (16.6)	0.200	1.15
CRP mg/l	1	5.57 (7.06)	7.79 (11.52)	<0.001	1.35
FVIIc g/l (FVII antigen/FVIIc/FVIIa)	6	109.2 (28.9)	108.2 (29.7)	0.572	0.99
Fibrinogen g/l	6	275.4 (55.7)	286.0 (49.8)	<0.001	1.37

Supplementary Table 2 Genic scan: genes showing association with blood phenotypes

Phenotype	Previous GWAS hit (alternate phenotype previously showing association)	rs	*gene*	p-value	FDR adj pvalue	Common vs Rare HMZ*	% R2	MAF
Triglycerides		rs301	*LPL*	1.59E-06	0.01	-0.314	1.12	24.0
	(1) (LDL)	rs6589566	*APOA5-A4-C3-A1*	1.98E-05	0.01	0.633	0.91	7.0
		E40K(2)	*ANGPTL4*	2.23E-05	0.01	-0.420	0.69	2.1
		rs264	*LPL*	2.46E-05	0.01	-0.217	0.89	15.1
		rs662799	*APOA5*	3.15E-05	0.01	0.817	0.81	6.1
		rs13702	*LPL*	6.20E-05	0.01	-0.247	0.81	28.4
		rs3135506	*APOA5*	1.50E-04	0.02	1.076	0.71	5.7
		rs3779788	*LPL*	2.26E-04	0.03	-0.184	0.70	15.3
		rs5072	*APOA5-A4-C3-A1*	2.78E-04	0.03	0.527	0.69	7.3
		rs579163	*APOA5-A4-C3-A1*	2.81E-04	0.03	0.258	0.69	32.6
	(3)	rs328	*LPL*	3.68E-04	0.03	-0.189	0.66	10.9
		rs5128	*APOA5-A4-C3-A1*	6.05E-04	0.04	0.444	0.58	8.0
		rs1131012	*PECAM1*	7.41E-04	0.05	-0.040	0.53	48.6
		rs4075015	*IL6R*	1.04E-03	0.06	0.220	0.58	41.5
		rs12285095	*APOA5-A4-C3-A1*	1.09E-03	0.06	0.880	0.57	5.9
		rs4803455	*TGFB1*	1.25E-03	0.06	-0.022	0.56	49.9
		rs7252574 (2)	*ANGPTL4*	1.26E-03	0.06	-0.008	0.50	49.5
		rs780094	*GCKR*	1.61E-03	0.07	0.213	0.48	37.8
		rs268	*LPL*	1.81E-03	0.08	-0.964	0.53	2.1
		rs4520	*APOA5-A4-C3-A1*	3.07E-03	0.11	0.298	0.45	21.3
		rs4508523	*LDLR*	3.15E-03	0.11	0.298	0.48	12.5
		rs5142	*APOA5-A4-C3-A1*	3.66E-03	0.12	0.211	0.47	8.0
		rs3781600	*LRP5*	4.07E-03	0.13	0.858	0.46	8.4
		rs3289	*LPL*	4.79E-03	0.14	1.267	0.45	2.7
		rs3211719	*F7*	5.26E-03	0.15	0.269	0.44	24.6
		rs316	*LPL*	5.42E-03	0.15	-0.463	0.44	11.9
		rs7255436 (2)	*ANGPTL4*	5.77E-03	0.16	-0.019	0.38	49.0
		rs7746553	*C2*	6.21E-03	0.16	0.282	0.43	16.4
		rs689	*INS*	6.39E-03	0.16	0.235	0.43	27.6
		rs2479408	*PCSK9*	6.74E-03	0.16	0.215	0.42	19.9
		rs11216157	*APOA5-A4-C3-A1*	7.08E-03	0.17	0.542	0.42	11.8
		rs499718	*PCSK9*	7.53E-03	0.17	0.315	0.41	18.1
		rs5128	*APOA5-A4-C3-A1*	7.77E-03	0.17	0.173	0.38	36.5
		rs4417316	*APOA5-A4-C3-A1*	8.93E-03	0.19	-0.085	0.40	10.0
		rs397211	*ILRN1*	9.08E-03	0.19	0.178	0.39	30.5
		rs676210	*APOB*	1.02E-02	0.20	-0.213	0.38	21.5
		rs11465699	*IL18RAP*	1.02E-02	0.20	-0.447	0.38	4.4
Cholesterol		rs429358/ rs7412	*APOE-C1-C2-C4*	1.09E-19	0.01	0.124	3.28	33.7
	(3)	rs6511720	*LDLR*	1.11E-07	0.01	-0.462	1.34	12.3
	(3,4,5), (6) (AD), (7) (AD)	rs4420638	*APOE-C1-C2-C4*	2.50E-06	0.01	0.405	1.08	19.1
		rs6413504	*LDLR*	5.54E-06	0.01	0.293	1.01	49.0
		rs2738446	*LDLR*	1.62E-05	0.01	0.273	0.92	45.3
		rs2228671	*LDLR*	3.11E-05	0.01	-0.419	0.87	13.1
		rs585967	*APOB*	1.60E-04	0.02	-0.391	0.73	14.8
		rs12713843	*APOB*	1.69E-04	0.02	0.799	0.59	49.5
		rs952275	*APOB*	4.02E-04	0.04	-0.222	0.66	49.2
		rs12714264	*APOB*	5.07E-04	0.04	-0.322	0.64	13.5
		rs11591147	*PCSK9*	1.04E-03	0.08	-2.741	0.56	0.9

		rs10199768	*APOB*	1.43E-03	0.10	0.213	0.55	44.0
		rs7259004	*APOE-C1-C2-C4*	2.61E-03	0.16	-0.269	0.50	10.8
		rs11602256	*LRP5*	3.35E-03	0.18	0.165	0.48	32.6
		rs1367117	*APOB*	3.47E-03	0.18	0.216	0.47	32.8
		rs934197	*APOB*	3.60E-03	0.18	0.216	0.47	32.7
LDL-cholesterol		rs429358/ rs7412	*APOE-C1-C2-C4*	3.49E-17	0.01	0.116	3.34	33.2
	(3), 4)	rs6511720	*LDLR*	5.57E-07	0.01	-0.612	1.39	12.2
		rs585967	*APOB*	1.51E-06	0.01	-0.564	1.29	14.6
		rs2228671	*LDLR*	6.36E-06	0.01	-0.588	1.15	13.0
		rs12714264	*APOB*	6.79E-06	0.01	-0.499	1.15	13.4
	(3)-4), (6) (AD), (7) (AD)	rs4420638	*APOE-C1-C2-C4*	3.75E-05	0.01	0.469	0.98	18.8
		rs952275	*APOB*	5.37E-05	0.01	-0.271	0.95	49.2
		rs12691202	*APOB*	4.42E-04	0.04	-0.730	0.74	3.8
		rs11591147	*PCSK9*	4.49E-04	0.04	-2.389	0.73	0.9
		rs729287	*ALX4*	9.20E-04	0.08	-0.297	0.68	29.6
		rs3740878	*EXT2*	1.11E-03	0.08	-0.291	0.66	29.9
		rs8110695	*LDLR*	1.75E-03	0.12	-0.187	0.61	21.6
		rs6413504	*LDLR*	2.10E-03	0.13	0.213	0.60	49.1
		rs10199768	*APOB*	2.81E-03	0.15	0.215	0.57	44.3
		rs7259004	*APOE-C1-C2-C4*	3.05E-03	0.15	-0.394	0.56	10.9
		rs1051931	*PLA2G7*	3.09E-03	0.15	0.298	0.56	19.1
		rs423490	*C3*	3.38E-03	0.15	-0.186	0.55	24.0
		rs1051931	*PLA2G7*	3.41E-03	0.15	0.239	0.49	19.0
		rs344548	*C3*	3.45E-03	0.15	-0.298	0.55	16.8
		rs10811661	*CDKN2A/2B*	4.34E-03	0.17	-0.385	0.47	15.8
		rs8112351	*C3*	4.80E-03	0.18	-0.188	0.52	23.9
		rs3814309	*GSTM3*	5.52E-03	0.20	-0.063	0.50	32.0
		rs1367117	*APOB*	5.81E-03	0.20	0.190	0.50	33.0
HDL-cholesterol		rs708272	*CETP*	8.88E-05	0.07	0.272	0.85	44.0
		rs1729410	*APOA5-A4-C3-A1*	4.11E-04	0.13	0.221	0.76	43.2
		rs1800588	*LIPC*	6.95E-04	0.13	0.313	0.71	20.9
		rs264	*LPL*	8.25E-04	0.13	0.214	0.69	15.1
		rs328	*LPL*	1.05E-03	0.14	0.459	0.60	10.3
		rs4769060	*ALOX5AP*	1.92E-03	0.19	0.217	0.61	45.1
ApoAI	(3) (HDL)	rs1800588	*LIPC*	4.94E-05	0.03	0.329	0.96	20.9
		rs708272	*CETP*	6.90E-05	0.03	0.273	0.87	43.9
		rs3885907	*ALOX5AP*	6.06E-04	0.15	0.240	0.71	45.7
		rs1729410	*APOA5-A4-C3-A1*	7.45E-04	0.15	0.214	0.69	43.2
		alu1(8)	*IGF2*	9.53E-04	0.15	0.231	0.74	47.3
		rs344550	*C3*	1.46E-03	0.20	0.033	0.63	34.8
		rs5167	*APOE-C1-C2-C4*	1.76E-03	0.20	0.206	0.62	34.4
		rs10162089	*ALOX5AP*	1.96E-03	0.20	0.220	0.60	47.9
	(3)(HDL, TG)	rs328	*LPL*	2.22E-03	0.20	0.574	0.53	10.3
APOB		rs429358/ rs7412	*APOE-C1-C2-C4*	1.27E-16	0.01	0.068	3.16	33.2
	(3) (LDL)	rs11591147	*PCSK9*	4.41E-05	0.02	-3.130	0.95	0.9
		rs7259004	*APOE-C1-C2-C4*	1.95E-04	0.06	-0.093	0.82	10.9
		rs585967	*APOB*	8.31E-04	0.14	-0.333	0.68	14.6
		rs952275	*APOB*	9.07E-04	0.14	-0.202	0.68	49.2
		rs1537236	*GSTM4*	1.07E-03	0.14	0.060	0.66	48.8
	(3) (LDL); (5)(LDL)	rs6511720	*LDLR*	1.25E-03	0.14	-0.199	0.64	12.2
		rs1121923	*LPL*	1.71E-03	0.15	-1.984	0.61	3.0
		rs729287	*ALX4*	1.75E-03	0.15	-0.295	0.61	29.6
		rs3740878	*EXT2*	1.87E-03	0.15	-0.289	0.61	29.9
		rs2738446	*LDLR*	2.28E-03	0.16	0.145	0.59	45.6
		rs2228671	*LDLR*	2.43E-03	0.16	-0.387	0.58	13.0

Lp-PLA2		rs429358/ rs7412	*APOE-C1-C2-C4*	2.55E-14	0.01	0.210	3.00	35.0
	(4)(LDL); (9)(LDL); (5)(LDL) (6) (AD) (7) (AD)	rs4420638	*APOE-C1-C2-C4*	1.41E-06	0.01	0.490	1.48	19.5
		rs3918232	*NOS3*	4.32E-04	0.12	-1.031	0.68	49.7
BMI		6815A>T (8)	*IGF2*	1.59E-05	0.02	-0.439	0.93	23.5
		rs680	*IGF2*	4.81E-04	0.19	-0.302	0.60	26.9
Fibrinogen		rs4508864	*FGA-B-G*	1.31E-07	0.01	0.425	1.32	18.9
		rs1800790	*FGB*	2.74E-07	0.01	0.368	1.18	19.3
		rs4220	*FGA-B-G*	9.66E-07	0.01	0.347	1.16	17.1
		rs2070016	*FGA-B-G*	6.50E-06	0.01	0.377	1.00	14.8
		rs1685354	*UCP3*	8.50E-04	0.14	0.311	0.59	25.2
		rs1537236	*GSTM4*	1.61E-03	0.20	0.199	0.54	48.9
		rs618923	*ApoA5-A4-C3-A1*	1.75E-03	0.20	-0.064	0.53	23.9
FVII		rs6046	*F7*	9.20E-140	0.01	-2.154	23.59	10.3
		insdel(10)	*F7*	1.89E-126	0.01	-2.065	19.84	10.6
		rs555212	*F7*	1.76E-34	0.01	0.814	6.33	23.8
		-670A>C (10)	*F7*	1.39E-31	0.01	0.806	6.06	23.2
		-402G>C (10)	*F7*	3.57E-22	0.01	0.580	3.96	25.1
		rs867186	*PROCR*	5.81E-13	0.01	0.825	2.25	9.1
		rs11576175	*CTSS*	2.39E-04	0.03	0.609	0.70	8.3
		rs780094	*GCKR*	2.69E-04	0.03	0.234	0.61	37.8
		rs4676411	*CAPN10*	1.84E-03	0.14	0.129	0.53	37.9
		rs11602906	*UCP2*	1.95E-03	0.14	0.200	0.52	7.5
		rs5167	*APOE-C1-C2-C4*	2.26E-03	0.15	0.048	0.52	34.3
		rs6413504	*LDLR*	2.63E-03	0.16	0.193	0.50	48.9
		rs3211719	*F7*	2.90E-03	0.17	0.281	0.51	24.6
		rs11536857	*TLR4*	3.12E-03	0.17	0.698	0.48	7.8
CRP		rs429358/ rs7412	*APOE-C1-C2-C4*	1.28E-10	0.01	-0.320	1.90	33.9
	(4)(LDL); (3)(LDL); (5)(LDL); (6) (AD), (7)(AD)	rs4420638	*APOE-C1-C2-C4*	2.41E-07	0.01	-0.401	1.39	19.4
		rs3093077	*CRP*	1.03E-04	0.03	-0.245	0.75	33.7
		rs1130864	*CRP*	2.19E-04	0.05	0.229	0.98	32.0
		rs3091244	*CRP*	2.70E-04	0.05	0.188	0.67	5.4

*Standardised difference

** Standardised difference common HMZ vs HTZ

Supplementary Table 3 Categories and candidate genes and the number of SNPs analysed in each, included in the 768 Illumina bundle

Inflammation genes		***Lipid genes***		***ROS genes***		***Clotting genes***		***T2D***		***Complement genes***		***Genes of Interest***		***CNV tSNPs***
IFNG	10	*APOA2*	11	*CAT*	13	*F2*	12	*CALPAIN10*	14	*BF*	13	*GSK3*	4	59
IL10	7	*APOB*	39	*GSTM1*	5	*F7*	7			*C2*	8	*G6PD*	2	
IL18RAP	17	*APOE-C1-C2-C4*	13	*GSTM3*	7	*FIBA-FIBB-FIBG*	16	*ALX4*	1	*C3*	33	*ZNF202*	11	
IL1A	3	*LPL*	27	*GSTM5*	9		35	*CAMTA1*	1	*CFD*	7	*ALOX5AP*	26	
IL1B	6	*LIPE*	6	*HMOX1*	7			*CXCR4???*	1	*CFH*	18	*NDST1*	9	
IL6	10	*LRP1*	21	*HSPD1*	3			*CYBA*	1		79	*MMP3*	6	
IL6R	14	*SREBP1*	4	*NOS3*	14			*EXT2*	1			*CTSS*	3	
IL6ST	10	*A5-A4-C3-A1*	35	*SELS*	9			*HHEX*	2			*TIMP1*	4	
ILRN1	12	*LDLR*	22	*SOD2*	14			*KCNJ11*	1				65	
LTA	11	*LRP5*	27	*UCP2*	8			*KCTD12*	1					
MYD88	4	*PCSK9*	21	*UCP3*	6			*MMP26*	1					
PLA2G7	13		226	*PRDX5*	3			*NEUROG3*	1					
PLA2GX	1			*GSTM4*	1			*PSMA6*	1					
TGFB1	8				99			*SLC30A8*	1					
TGFB3	9							rs1804451	1					
TLR2	13							rs1256517	1					
TLR4	20								29					
TNF	2													
	170													

Supplementary Table 4 Additional candidate genes analysed in NPHSII

Gene	No of SNPs	Gene	No of SNPs	Gene	No of SNPs	Gene	No of SNPs	Gene
ACE	1	*F2*	1	*LTBP4*	1		*TPO*	2
ADH1C	1	*F5*	1	*LTBP4*	7		*UCP2*	1
ADIPOQ	4	*F7*	4	*MMP3*	1		*UCP3*	1
ADRB1	1	*FGA-B-G*	4	*MTHFR*	1		*VEGF*	1
ADRB3	1	*FTO*	1	*MTP*	3		*WNT5*	1
AGT	8	*GAL*	1	*MTR*	1		*ZNF202*	1
ANGPTL3	2	*GCH1*	1	*NOS3*	4			
ANGPTL4	7	*GCKR*	1	*PCSK9*	3			
APOA1-C3-A4-A5	11	*GSTM1*	1	*PECAM1*	3			
APOB	7	*GSTT1*	1	*PLA2G5*	2			
APOE	1	*HABP2*	1	*PLA2G7*	1			
ARH	1	*IGF12*	1	*PLG*	1			
AT1	1	*IGF2*	10	*PON1*	2			
AT2	2	*IL18*	8	*PON2*	1			
BK1	2	*IL6*	3	*PON3*	1			
BK2	2	*INS*	1	*PPARA*	3			
CBS	2	*INSIG1*	1	*PPARD*	1			
CDKN2A	2	*LCAT*	1	*PPARG*	3			
CETP	1	*LDLR*	2	*PROCR*	1			
CNR1	1	*LIPC*	1	*SERPINA1*	3			
CRP	4	*LIPE*	2	*SERPINE1*	1			
CTGF	1	*LPL*	5	*TCF7L2*	2			
CYP11P2	1	*LTBP1*	1	*THBD*	2			
F12	1	*LTBP3*	3	*TNFA*	1			

Supplementary Table 5a List of Genes and SNPs with rs numbers included in the 768 Illumina bead Array

rs	*Gene*	*rs*	*gene*	*rs*	*gene*
rs1256517	*LOC646279*	rs3766404	*CFH*	rs17547610	*LRP1*
rs1800764	*ACE*	rs380390	*CFH*	rs1800127	*LRP1*
rs4459609	*ACE*	rs402032	*CFH*	rs1800159	*LRP1*
rs2055049	*AGTR1*	rs419137	*CFH*	rs1800168	*LRP1*
rs2276736	*AGTR1*	rs460184	*CFH*	rs2229278	*LRP1*
rs388915	*AGTR1*	rs515299	*CFH*	rs4759044	*LRP1*
rs5182	*AGTR1*	rs534399	*CFH*	rs715948	*LRP1*
rs5186	*AGTR1*	rs7524776	*CFH*	rs7304504	*LRP1*
rs931490	*AGTR1*	rs1136774	*CTSS*	rs7306742	*LRP1*
rs10162089	*ALOX5AP*	rs11576175	*CTSS*	rs7398375	*LRP1*
rs10507391	*ALOX5AP*	rs7511673	*CTSS*	rs7485650	*LRP1*
rs11147439	*ALOX5AP*	rs932206	*CXCR4*	rs7956957	*LRP1*
rs12019512	*ALOX5AP*	rs2254073	*CYBA*	rs7968719	*LRP1*
rs12429692	*ALOX5AP*	rs3740878	*EXT2*	rs922963	*LRP1*
rs17222814	*ALOX5AP*	rs1799963	*F2*	rs11574422	*LRP5*
rs17238738	*ALOX5AP*	rs2070852	*F2*	rs11602256	*LRP5*
rs17245120	*ALOX5AP*	rs3136430	*F2*	rs11826287	*LRP5*
rs17245317	*ALOX5AP*	rs3136431	*F2*	rs12417792	*LRP5*
rs3885907	*ALOX5AP*	rs3136440	*F2*	rs2242340	*LRP5*
rs3935644	*ALOX5AP*	rs3136456	*F2*	rs2508836	*LRP5*
rs4073261	*ALOX5AP*	rs3136457	*F2*	rs312009	*LRP5*
rs4075131	*ALOX5AP*	rs3136520	*F2*	rs312014	*LRP5*
rs4076128	*ALOX5AP*	rs3136532	*F2*	rs312016	*LRP5*
rs4254165	*ALOX5AP*	rs4752930	*F2*	rs312786	*LRP5*
rs4503649	*ALOX5AP*	rs5896	*F2*	rs312788	*LRP5*
rs4597169	*ALOX5AP*	rs5898	*F2*	rs314756	*LRP5*
rs4769055	*ALOX5AP*	rs1475931	*F7*	rs3736228	*LRP5*
rs4769060	*ALOX5AP*	rs3093248	*F7*	rs3781596	*LRP5*
rs4769873	*ALOX5AP*	rs3211719	*F7*	rs3781600	*LRP5*
rs4769875	*ALOX5AP*	rs510317	*F7*	rs491347	*LRP5*
rs9315051	*ALOX5AP*	rs555212	*F7*	rs4930573	*LRP5*
rs9508835	*ALOX5AP*	rs574229	*F7*	rs4988300	*LRP5*
rs9578196	*ALOX5AP*	rs6046	*F7*	rs4988327	*LRP5*
rs9579648	*ALOX5AP*	rs1118823	*FGA-B-G*	rs4988331	*LRP5*
rs9671124	*ALOX5AP*	rs1800788	*FGA-B-G*	rs587397	*LRP5*
rs729287	*ALX4*	rs1800792	*FGA-B-G*	rs606989	*LRP5*
rs11590967	*APOA2*	rs2066860	*FGA-B-G*	rs608343	*LRP5*
rs12721035	*APOA2*	rs2066870	*FGA-B-G*	rs624947	*LRP5*
rs1803091	*APOA2*	rs2070016	*FGA-B-G*	rs638051	*LRP5*

rs2070902	*APOA2*	rs2070018	*FGA-B-G*	rs676318	*LRP5*
rs3557	*APOA2*	rs2070022	*FGA-B-G*	rs7116604	*LRP5*
rs5082	*APOA2*	rs2070025	*FGA-B-G*	rs1041981	*LTA*
rs5085	*APOA2*	rs4220	*FGA-B-G*	rs1799964	*LTA*
rs3813627	*APOA2*	rs4463047	*FGA-B-G*	rs1800629	*LTA*
rs4233368	*APOA2*	rs4508864	*FGA-B-G*	rs1800683	*LTA*
rs4489574	*APOA2*	rs6054	*FGA-B-G*	rs2009658	*LTA*
rs7528588	*APOA2*	rs7659024	*FGA-B-G*	rs2071590	*LTA*
rs11216157	*ApoA5-A4-C3-A1*	rs7659613	*FGA-B-G*	rs2229094	*LTA*
rs11216159	*ApoA5-A4-C3-A1*	rs7673587	*FGA-B-G*	rs2844484	*LTA*
rs12284864	*ApoA5-A4-C3-A1*	rs1050828	*G6PD*	rs3093662	*LTA*
rs12285095	*ApoA5-A4-C3-A1*	rs2230037	*G6PD*	rs3093668	*LTA*
rs12365440	*ApoA5-A4-C3-A1*	rs2302485	*GSK3*	rs915654	*LTA*
rs12365462	*ApoA5-A4-C3-A1*	rs3745233	*GSK3*	rs2499953	*MMP26*
rs1263173	*ApoA5-A4-C3-A1*	rs3810149	*GSK3*	rs3025090	*MMP3*
rs12718464	*ApoA5-A4-C3-A1*	rs708598	*GSK3*	rs476762	*MMP3*
rs1729410	*ApoA5-A4-C3-A1*	rs12068997	*GSTM1*	rs522616	*MMP3*
rs2071521	*ApoA5-A4-C3-A1*	rs12097277	*GSTM1*	rs527832	*MMP3*
rs2071523	*ApoA5-A4-C3-A1*	rs12562055	*GSTM1*	rs650108	*MMP3*
rs2075291	*ApoA5-A4-C3-A1*	rs4147565	*GSTM1*	rs679620	*MMP3*
rs2075294	*ApoA5-A4-C3-A1*	rs4147567	*GSTM1*	rs6796045	*MYD88*
rs2098453	*ApoA5-A4-C3-A1*	rs1055259	*GSTM3*	rs6853	*MYD88*
rs2216311	*ApoA5-A4-C3-A1*	rs1332018	*GSTM3*	rs7625290	*MYD88*
rs2727789	*ApoA5-A4-C3-A1*	rs1537234	*GSTM3*	rs7744	*MYD88*
rs2849174	*ApoA5-A4-C3-A1*	rs1803687	*GSTM3*	rs10074650	*NDST1*
rs33989105	*ApoA5-A4-C3-A1*	rs2234696	*GSTM3*	rs11959543	*NDST1*
rs35625356	*ApoA5-A4-C3-A1*	rs3814309	*GSTM3*	rs12516924	*NDST1*
rs35631472	*ApoA5-A4-C3-A1*	rs4970737	*GSTM3*	rs2273232	*NDST1*
rs35833621	*ApoA5-A4-C3-A1*	rs1537236	*GSTM4*	rs2273233	*NDST1*
rs4417316	*ApoA5-A4-C3-A1*	rs10776699	*GSTM5*	rs2545342	*NDST1*
rs5072	*ApoA5-A4-C3-A1*	rs11101992	*GSTM5*	rs2748216	*NDST1*

rs5101	*ApoA5-A4-C3-A1*	rs11807	*GSTM5*	rs3095902	*NDST1*
rs5142	*ApoA5-A4-C3-A1*	rs12743015	*GSTM5*	rs3797621	*NDST1*
rs579163	*ApoA5-A4-C3-A1*	rs366631	*GSTM5*	rs10823406	*NGN3*
rs595049	*ApoA5-A4-C3-A1*	rs3768490	*GSTM5*	rs10277237	*NOS3*
rs601634	*ApoA5-A4-C3-A1*	rs4970772	*GSTM5*	rs1799983	*NOS3*
rs603446	*ApoA5-A4-C3-A1*	rs886177	*GSTM5*	rs1800779	*NOS3*
rs618354	*ApoA5-A4-C3-A1*	rs929166	*GSTM5*	rs1800783	*NOS3*
rs618923	*ApoA5-A4-C3-A1*	rs1111875	*HHEX*	rs2070744	*NOS3*
rs6589566	*ApoA5-A4-C3-A1*	rs7923837	*HHEX*	rs2373929	*NOS3*
rs6589567	*ApoA5-A4-C3-A1*	rs11912889	*HMOX1*	rs3918166	*NOS3*
rs6589568	*ApoA5-A4-C3-A1*	rs2071746	*HMOX1*	rs3918186	*NOS3*
rs9804646	*ApoA5-A4-C3-A1*	rs2285112	*HMOX1*	rs3918188	*NOS3*
rs10199768	*APOB*	rs5755713	*HMOX1*	rs3918201	*NOS3*
rs1042031	*APOB*	rs5755720	*HMOX1*	rs3918227	*NOS3*
rs11126598	*APOB*	rs743811	*HMOX1*	rs3918232	*NOS3*
rs11676704	*APOB*	rs9282702	*HMOX1*	rs3918234	*NOS3*
rs12691202	*APOB*	rs2340690	*HSPD1*	rs743507	*NOS3*
rs12713450	*APOB*	rs6434930	*HSPD1*	rs10465832	*PCSK9*
rs12713540	*APOB*	rs788016	*HSPD1*	rs10888896	*PCSK9*
rs12713554	*APOB*	rs10878763	*IFNG*	rs11206514	*PCSK9*
rs12713559	*APOB*	rs11177074	*IFNG*	rs11583680	*PCSK9*
rs12713675	*APOB*	rs2069709	*IFNG*	rs13312	*PCSK9*
rs12713843	*APOB*	rs2069713	*IFNG*	rs17111495	*PCSK9*
rs12714097	*APOB*	rs2069716	*IFNG*	rs2182833	*PCSK9*
rs12714214	*APOB*	rs2069718	*IFNG*	rs2479406	*PCSK9*
rs12714225	*APOB*	rs2069720	*IFNG*	rs2479408	*PCSK9*
rs12714264	*APOB*	rs2069727	*IFNG*	rs2479409	*PCSK9*
rs12720791	*APOB*	rs2069728	*IFNG*	rs2495481	*PCSK9*
rs12720801	*APOB*	rs3181033	*IFNG*	rs4927193	*PCSK9*
rs12720843	*APOB*	rs1800871	*IL10*	rs499718	*PCSK9*
rs12720848	*APOB*	rs1800890	*IL10*	rs505151	*PCSK9*
rs12720854	*APOB*	rs1878672	*IL10*	rs529787	*PCSK9*
rs13306187	*APOB*	rs3021094	*IL10*	rs533375	*PCSK9*
rs1367117	*APOB*	rs3024505	*IL10*	rs557435	*PCSK9*
rs1713223	*APOB*	rs3024509	*IL10*	rs562556	*PCSK9*
rs17398765	*APOB*	rs6673928	*IL10*	rs568052	*PCSK9*
rs1801700	*APOB*	rs11465699	*IL18RAP*	rs572512	*PCSK9*
rs1801701	*APOB*	rs11465716	*IL18RAP*	rs7552841	*PCSK9*

rs1801702	*APOB*	rs11465724	*IL18RAP*	rs9804032	*PCSK9*
rs1801703	*APOB*	rs11465730	*IL18RAP*	rs1051931	*PLA2G7*
rs2163204	*APOB*	rs11465732	*IL18RAP*	rs10948300	*PLA2G7*
rs2854725	*APOB*	rs11695455	*IL18RAP*	rs12195701	*PLA2G7*
rs520354	*APOB*	rs11886793	*IL18RAP*	rs12528807	*PLA2G7*
rs533617	*APOB*	rs17027173	*IL18RAP*	rs1421368	*PLA2G7*
rs550619	*APOB*	rs17821875	*IL18RAP*	rs1421378	*PLA2G7*
rs584542	*APOB*	rs2075186	*IL18RAP*	rs16874962	*PLA2G7*
rs585967	*APOB*	rs2293225	*IL18RAP*	rs17288905	*PLA2G7*
rs6752026	*APOB*	rs3771150	*IL18RAP*	rs1805018	*PLA2G7*
rs676210	*APOB*	rs6543139	*IL18RAP*	rs2216465	*PLA2G7*
rs679899	*APOB*	rs6741230	*IL18RAP*	rs4498351	*PLA2G7*
rs934197	*APOB*	rs7559479	*IL18RAP*	rs6929105	*PLA2G7*
rs952275	*APOB*	rs7559845	*IL18RAP*	rs974670	*PLA2G7*
rs10413089	*APOE-C1-C2-C4*	rs917998	*IL18RAP*	rs584367	*PLA2GX*
rs12691089	*APOE-C1-C2-C4*	rs17561	*IL1A*	rs11600990	*PRDX5*
rs405509	*APOE-C1-C2-C4*	rs2071373	*IL1A*	rs4930702	*PRDX5*
rs429358	*APOE-C1-C2-C4*	rs2856838	*IL1A*	rs9787810	*PRDX5*
rs439401	*APOE-C1-C2-C4*	rs1143630	*IL1B*	rs1048990	*PSMA6*
rs4420638	*APOE-C1-C2-C4*	rs1143634	*IL1B*	rs12438654	*SELS*
rs4803770	*APOE-C1-C2-C4*	rs1143643	*IL1B*	rs2101171	*SELS*
rs5112	*APOE-C1-C2-C4*	rs16944	*IL1B*	rs2291250	*SELS*
rs5126	*APOE-C1-C2-C4*	rs2853550	*IL1B*	rs2306534	*SELS*
rs5127	*APOE-C1-C2-C4*	rs4848306	*IL1B*	rs4965372	*SELS*
rs5167	*APOE-C1-C2-C4*	rs10242595	*IL6*	rs4965815	*SELS*
rs7259004	*APOE-C1-C2-C4*	rs10499563	*IL6*	rs7178239	*SELS*
rs7259679	*APOE-C1-C2-C4*	rs11766273	*IL6*	rs8036413	*SELS*
rs1048709	*BF*	rs12700386	*IL6*	rs9874	*SELS*
rs1270942	*BF*	rs13306435	*IL6*	rs13266634	*SLC30A8*
rs2072633	*BF*	rs1800795	*IL6*	rs1804451	*SOD2*
rs4151651	*BF*	rs1800796	*IL6*	rs2758329	*SOD2*
rs4151657	*BF*	rs2069830	*IL6*	rs2758331	*SOD2*
rs4151659	*BF*	rs2069840	*IL6*	rs2842980	*SOD2*
rs4151660	*BF*	rs2069861	*IL6*	rs2855115	*SOD2*
rs4151664	*BF*	rs11584622	*IL6R*	rs4342445	*SOD2*
rs4151667	*BF*	rs1386821	*IL6R*	rs4987023	*SOD2*
rs537160	*BF*	rs1552481	*IL6R*	rs5746101	*SOD2*
rs641153	*BF*	rs28730736	*IL6R*	rs5746128	*SOD2*
rs9332730	*BF*	rs4075015	*IL6R*	rs5746129	*SOD2*
rs9501161	*BF*	rs4240872	*IL6R*	rs5746133	*SOD2*
rs2734335	*C2*	rs4329505	*IL6R*	rs5746136	*SOD2*
rs3020644	*C2*	rs4601580	*IL6R*	rs5746151	*SOD2*
rs4151648	*C2*	rs4845615	*IL6R*	rs7855	*SOD2*
rs497309	*C2*	rs4845617	*IL6R*	rs9456442	*SOD2*
rs547154	*C2*	rs4845618	*IL6R*	rs11656665	*SREBP1*
rs7746553	*C2*	rs6684439	*IL6R*	rs2228461	*SREBP1*

rs9267677	*C2*	rs7526293	*IL6R*	rs2229590	*SREBP1*
rs9332739	*C2*	rs8192284	*IL6R*	rs4925118	*SREBP1*
rs1047286	*C3*	rs10076283	*IL6ST*	rs10417924	*TGFB1*
rs11569422	*C3*	rs10180	*IL6ST*	rs12980942	*TGFB1*
rs11569429	*C3*	rs1063560	*IL6ST*	rs1800469	*TGFB1*
rs11569450	*C3*	rs10940495	*IL6ST*	rs1800471	*TGFB1*
rs11569523	*C3*	rs11574780	*IL6ST*	rs1800472	*TGFB1*
rs11569534	*C3*	rs1900173	*IL6ST*	rs4803455	*TGFB1*
rs11569538	*C3*	rs2228043	*IL6ST*	rs8110090	*TGFB1*
rs11569541	*C3*	rs2228046	*IL6ST*	rs8179181	*TGFB1*
rs11672613	*C3*	rs3729960	*IL6ST*	rs11466414	*TGFB3*
rs1389623	*C3*	rs6870870	*IL6ST*	rs2268625	*TGFB3*
rs163913	*C3*	rs17486819	*ILRN1*	rs2268626	*TGFB3*
rs2230199	*C3*	rs2637988	*ILRN1*	rs3917148	*TGFB3*
rs2230203	*C3*	rs3087263	*ILRN1*	rs3917158	*TGFB3*
rs2230204	*C3*	rs315920	*ILRN1*	rs3917200	*TGFB3*
rs2230205	*C3*	rs315936	*ILRN1*	rs3917210	*TGFB3*
rs2230210	*C3*	rs315946	*ILRN1*	rs4252315	*TGFB3*
rs2241393	*C3*	rs315952	*ILRN1*	rs7156293	*TGFB3*
rs2241394	*C3*	rs380092	*ILRN1*	rs13440825	*TIMP1*
rs2250656	*C3*	rs397211	*ILRN1*	rs2070584	*TIMP1*
rs2277984	*C3*	rs4251961	*ILRN1*	rs2077407	*TIMP1*
rs237554	*C3*	rs4252019	*ILRN1*	rs6520278	*TIMP1*
rs2642196	*C3*	rs4252041	*ILRN1*	rs11938228	*TLR2*
rs344540	*C3*	rs5219	*KCNJ11*	rs1339	*TLR2*
rs344548	*C3*	rs2876711	*KCTD12*	rs1898830	*TLR2*
rs344550	*C3*	rs10409044	*LDLR*	rs3804099	*TLR2*
rs344555	*C3*	rs11669576	*LDLR*	rs3804100	*TLR2*
rs3745568	*C3*	rs12608650	*LDLR*	rs4696480	*TLR2*
rs379527	*C3*	rs13306505	*LDLR*	rs4696483	*TLR2*
rs423490	*C3*	rs13306510	*LDLR*	rs5743699	*TLR2*
rs432001	*C3*	rs1799898	*LDLR*	rs5743703	*TLR2*
rs432823	*C3*	rs2228671	*LDLR*	rs5743704	*TLR2*
rs7257062	*C3*	rs2569540	*LDLR*	rs5743708	*TLR2*
rs8112351	*C3*	rs2738446	*LDLR*	rs720633	*TLR2*
rs1193179	*CAMTA1*	rs2738459	*LDLR*	rs7656411	*TLR2*
rs10933621	*CAPN10*	rs2738460	*LDLR*	rs893629	*TLR2*
rs1133353	*CAPN10*	rs2738464	*LDLR*	rs10759930	*TLR4*
rs11680323	*CAPN10*	rs4508523	*LDLR*	rs10818073	*TLR4*
rs2953147	*CAPN10*	rs5927	*LDLR*	rs11536857	*TLR4*
rs2953149	*CAPN10*	rs5928	*LDLR*	rs11536869	*TLR4*
rs2953161	*CAPN10*	rs5930	*LDLR*	rs11536889	*TLR4*
rs2975766	*CAPN10*	rs5931	*LDLR*	rs11536897	*TLR4*
rs3749164	*CAPN10*	rs6413504	*LDLR*	rs1554973	*TLR4*
rs3792267	*CAPN10*	rs6511720	*LDLR*	rs16906079	*TLR4*
rs3792270	*CAPN10*	rs7508676	*LDLR*	rs1927906	*TLR4*

rs3792272	*CAPN10*	rs8108932	*LDLR*	rs1927907	*TLR4*
rs3828345	*CAPN10*	rs8110695	*LDLR*	rs2149356	*TLR4*
rs4234123	*CAPN10*	rs10099160	*LPL*	rs2770144	*TLR4*
rs4676411	*CAPN10*	rs1121923	*LPL*	rs2770150	*TLR4*
rs10488736	*CAT*	rs13702	*LPL*	rs4987233	*TLR4*
rs11032699	*CAT*	rs1534649	*LPL*	rs5030717	*TLR4*
rs11032702	*CAT*	rs17410577	*LPL*	rs5030718	*TLR4*
rs11032703	*CAT*	rs1800590	*LPL*	rs5030719	*TLR4*
rs16925614	*CAT*	rs1801177	*LPL*	rs5031050	*TLR4*
rs208682	*CAT*	rs2197089	*LPL*	rs7044464	*TLR4*
rs2266630	*CAT*	rs248	*LPL*	rs7869402	*TLR4*
rs2284368	*CAT*	rs249	*LPL*	rs2507961	*TNF*
rs511895	*CAT*	rs253	*LPL*	rs4645843	*TNF*
rs554518	*CAT*	rs264	*LPL*	rs1050892	*UCP2*
rs564250	*CAT*	rs268	*LPL*	rs11602906	*UCP2*
rs566979	*CAT*	rs270	*LPL*	rs2306820	*UCP2*
rs7933285	*CAT*	rs281	*LPL*	rs2632725	*UCP2*
rs7947841	*CAT*	rs283	*LPL*	rs643064	*UCP2*
rs12985692	*CFD*	rs285	*LPL*	rs655717	*UCP2*
rs13090	*CFD*	rs298	*LPL*	rs659366	*UCP2*
rs1651895	*CFD*	rs301	*LPL*	rs660339	*UCP2*
rs1683564	*CFD*	rs316	*LPL*	rs1685354	*UCP3*
rs17684161	*CFD*	rs328	*LPL*	rs1800849	*UCP3*
rs2230216	*CFD*	rs3289	*LPL*	rs2229707	*UCP3*
rs3826945	*CFD*	rs3779788	*LPL*	rs2734827	*UCP3*
rs1061170	*CFH*	rs4921684	*LPL*	rs3781907	*UCP3*
rs1065489	*CFH*	rs5934	*LPL*	rs7926165	*UCP3*
rs10801560	*CFH*	rs7016529	*LPL*	rs1077911	*ZNF202*
rs10922096	*CFH*	rs9644636	*LPL*	rs10893079	*ZNF202*
rs1329421	*CFH*	rs10783814	*LRP1*	rs11219263	*ZNF202*
rs1329423	*CFH*	rs10876966	*LRP1*	rs11219270	*ZNF202*
rs17434860	*CFH*	rs11172113	*LRP1*	rs12794996	*ZNF202*
rs2019727	*CFH*	rs12369345	*LRP1*	rs1557429	*ZNF202*
rs3753394	*CFH*	rs12578897	*LRP1*	rs2282641	*ZNF202*
rs3753395	*CFH*	rs12814239	*LRP1*	rs2282646	*ZNF202*
		rs1466535	*LRP1*	rs3183878	*ZNF202*
				rs597530	*ZNF202*
				rs9326264	*ZNF202*

Supplementary Table 5b List of CNV SNPs with rs numbers included in the 768 Illumina bead Array

rs	*CNV (Redon et al, 2007)(11)*
rs10002047	4q26
rs10155079	4p15.1
rs1037388	1q31
rs10405246	19q13
rs1048374	11q12.1
rs10494095	1p13
rs10501812	11q21
rs10503141	18q22.1
rs10958546	8q11.23
rs1118333	11q11
rs11203531	8p22
rs11581756	1q21
rs11697985	20p13
rs11753874	6p25
rs12213776	6q14
rs12455301	18q22.1
rs13220198	6q14
rs1339867	1q41
rs1342114	9q31
rs136989	22q13
rs1424241	16q22.3
rs1553911	4p15.1
rs1559698	4q31
rs1779540	14q21

Supplementary Table 6. Comparison between statistically significant SNPs and independent genetic loci associated with each of the phenotypes tested. Arrangement into loci using LD clumping in PLINK with an R^2 of 0.01.

Trait	Significant SNPs	Independent SNPs or loci	no. genes
Triglycerides	37	20	16
Cholesterol	16	7	5
LDL-cholesterol	23	8	10
HDL-cholesterol	6	6	5
ApoAI	9	7	7
APOB	12	7	8
Lp-PLA2	3	2	2
BMI	2	1	1
Fibrinogen	7	3	4
FVII	14	9	9
CRP	5	2	2
Total	134	72	69

Supplementary Table 7. Additional genes, SNPs and their rs numbers genotyped in NPHSII

Rs number /SNP	*Gene*	*Rs number /SNP*	*Gene*	*Rs number /SNP*	*Gene*
In10 in/del(11)	*ACE*	rs429358	*APOE*	c2207t *(12)*	*IGF2*
rs698	*ADH1C*	rs7412	*APOE*	rs2230949	*IGF2*
rs17300539	*ADIPOQ*	rs5186	*AT1*	c2722t *(12)*	*IGF2*
rs1501299	*ADIPOQ*	rs1675 (13)	*AT2*	c8173t *(12)*	*IGF2*
rs266729	*ADIPOQ*	3123 (13)	*AT2*	AluI*(12)*	*IGF2*
rs2241766	*ADIPOQ*	-699G>C(14)	*BDKRB1*	rs680	*IGF2*
rs1801253	*ADRB1*	Ex1del/ins 9bp(14)	*BDKRB2*	rs3741211	*IGF2*
rs4994	*ADRB3*	Ex8 del/ins 68bp(15)	*CBS*	rs360729	*IL18*
rs2067853	*AGT*	rs10757274	*CDKN2A*	rs549908	*IL18*
rs11122574	*AGT*	rs2383206	*CDKN2A*	rs1946519	*IL18*
rs11122580	*AGT*	rs708272	*CETP*	rs2043055	*IL18*
rs3789679	*AGT*	rs2229579	*CNR1*	rs3882891	*IL18*
rs11568030	*AGT*	rs1130864	*CRP*	rs3814721	*IL18*
rs6687360	*AGT*	rs1205	*CRP*	rs2298455	*IL18*
rs3789666	*AGT*	rs3093077	*CRP*	rs1541304	*IL18*
rs10864772	*AGT*	rs3091244	*CRP*	rs1800795	*IL6*
rs 2031373	*ANGPTL3*	rs6918698	*CTGF*	rs1800796	*IL6*
rs 12042319	*ANGPTL3*	rs1799998	*CYP11P2*	rs10499563	*IL6*
rs4076317	*ANGPTL4*	rs4314418	*F12*	rs689	*INS*
rs7255436	*ANGPTL4*	20210G-A(16)	*F2*	rs7566605	*INSIG1*
rs1044250	*ANGPTL4*	rs6025	*F5*	rs4986970	*LCAT*
rs11672433	*ANGPTL4*	-670A>C (17)	*F7*	rs11669576	*LDLR*
rs7252574	*ANGPTL4*	rs6046	*F7*	rs45508991	*LDLR*
rs1808536	*ANGPTL4*	-402G>A (17)	*F7*	-514C>T*(18)*	*LIPC*
g118a E40K(19)	*ANGTPL4*	In/del(17)	*F7*	-60C>G*(20)*	*LIPE*
rs670	*APOA1*	rs2070011	*FGA*	In4T>C*(20)*	*LIPE*
rs675	*APOA4*	rs1800790	*FGB*	rs1801177	*LPL*
rs5110	*APOA5*	rs1800791	*FGB/G*	rs1800590	*LPL*
rs3135506	*APOA5*	rs1421085	*FTO*	-95G>T *(21)*	*LPL*
rs662799	*APOA5*	rs2187331	*GAL*	rs268	*LPL*
rs1729408	*APOA5-A4*	rs780094	*GCKR*	Rs328	*LPL*
rs693	*APOB*	mu-1*(22)*	*GSTM1*	256G>C	*LTBP1*
rs1801701	*APOB*	theta-1*(22)*	*GSTT1*	rs947791	*LTBP3*
rs1042031	*APOB*	rs7080536	*HABP2*	rs4244811	*LTBP3*
Ins/del(23)	*APOB*	rs7873	*IGF12*	rs11601767	*LTBP3*
rs4520	*APOC3*	rs1003483	*IGF2*	rs2369006	*LTBP4*
3'utr g>(24)	*APOC3*	a6815t*(12)*	*IGF2*	rs393569	*LTBP4*
rs2854117	*APOC3*	rs3168310	*IGF2*	rs11083560	*LTBP4*

Supplementary Table 8. Power calculation showing the effect sizes that could be detected at 80% power for an α value of 0.05, for a range of minor allele frequencies.

MAF	Rare Hz	St Effect size
0.05	0.0025	1.702
0.1	0.01	0.8463
0.2	0.04	0.4467
0.3	0.09	0.3223
0.4	0.16	0.2653
0.5	0.25	0.2343

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Supplementary Figure 1 Correlation between continuous phenotypes in men from the Northwick Park Heart Study
Values in the cells are absolute p-values for the between-phenotype associations

	BMI	Systolic	Diastolic	Cholesterol	HDL	LDL	Triglycerides	APOB	APOA	Homocyst	Folate	LpPLA2	CRP	F7
Systolic	1.85E-21													
Diastolic	6.57E-33	6.39e-315												
Cholesterol	2.76E-06	4.58E-06	9.45E-05											
HDL	7.27E-17	8.50E-05	1.87E-12	5.23E-01										
LDL	3.97E-03	1.99E-01	5.03E-01	**1.00E-250**	6.15E-10									
Triglycerides	1.64E-75	2.75E-25	3.57E-29	2.85E-75	1.58E-123	3.94E-04								
APOB	8.69E-14	9.17E-04	1.94E-01	2.06E-245	7.28E-10	4.21E-166	5.26E-41							
APOA	1.68E-12	9.13E-01	8.05E-02	3.74E-07	1.27E-123	1.18E-01	2.67E-27	2.72E-37						
Homocyst	7.11E-02	6.34E-04	8.07E-02	4.75E-01	7.05E-01	5.98E-01	6.27E-01	2.49E-01	1.66E-01					
Folate	2.39E-05	2.99E-01	6.57E-01	1.63E-01	8.09E-01	8.84E-01	3.35E-05	7.24E-01	3.20E-01	1.12E-78				
LpPLA2	1.30E-04	3.61E-05	4.39E-05	6.29E-53	5.61E-24	1.71E-29	1.80E-29	2.34E-17	4.75E-04	1.15E-01	6.03E-01			
CRP	1.79E-40	1.08E-14	2.96E-08	1.30E-07	1.09E-20	7.68E-04	4.51E-34	8.93E-11	1.52E-16	5.54E-02	1.14E-01	4.30E-02		
F7	5.78E-06	1.09E-05	1.27E-08	1.32E-35	1.94E-01	3.05E-07	7.25E-49	6.40E-20	2.19E-01	1.26E-02	6.25E-01	7.79E-01	2.00E-11	
FIB	7.05E-05	1.91E-08	3.54E-03	1.95E-07	9.38E-17	1.13E-07	1.44E-04	2.01E-15	1.34E-15	1.98E-02	1.55E-01	3.92E-01	3.86E-126	1.04E-0

Supplementary Figure 2 Correlations between protein phenotypes linked to CHD in 2775 men from the NPHS-II Study. Values in cells indicate Pearson's correlation co-efficient r. * p<0.01, ** p<0.001.

	ApoB	ApoAI	LpPLA2	CRP	F7
ApoAI	-0.2433**				
LpPLA2	0.193**	-0.0802**			
CRP	0.1344**	-0.1707**	0.0433*		
F7	0.1755**	0.0238	0.0058	0.1277**	
FIB	0.1528**	-0.1538**	0.0178	0.434**	0.0942

Supplementary Figure 3: Power calculation plot showing the relationship between MAF and effect size