

Supplementary Table 1. Chromosomal position, classification, and allele frequencies of sequence variants in *ANGPTL3*, *ANGPTL4*, *ANGPTL5*, and *ANGPTL6*

Chr	Position	nt change	Classification	MAF (%)		
				Whites	African-American	Hispanic
ANGPTL3 (Chr 1)						
62836324		in/del	intron	0.00	0.03	0.00
62836326		T/C	intron	0.00	0.03	0.00
62836893		C/T	intron	0.00	0.03	0.00
62836933		C/T	intron	0.00	0.03	0.00
62836939		T/A	intron	0.00	0.03	0.00
62839261		C/T	intron	0.00	0.03	0.00
62839321		G/T	intron	0.00	0.03	0.00
62839972		C/G	intron	0.00	0.03	0.00
62839988		T/G	intron	0.00	0.03	0.00
62840240		in/del	intron	0.00	0.03	0.00
62840261		G/A	intron	0.00	0.03	0.00
62842840		A/G	intron	0.00	0.03	0.00
62842849		A/G	intron	0.00	0.03	0.00
62843118		C/T	Noncoding	0.00	0.03	0.00
62843148		G/T	Noncoding	0.00	0.03	0.00
62843178		in/del	Noncoding	0.00	0.03	0.00
62843217		T/A	Noncoding	0.00	0.03	0.00
62836380		in/del	intron	0.05	0.00	0.00
62836856		C/T	intron	0.05	0.03	0.00
62836918		G/A	intron	0.05	0.00	0.00
62837198		G/A	intron	0.05	0.00	0.00
62840417		G/A	intron	0.05	0.00	0.00
62840452		C/T	intron	0.05	0.00	0.00
62836381		T/A	intron	0.00	0.05	0.00
62839541		T/A	intron	0.00	0.05	0.00
62842495		G/T	intron	0.00	0.05	0.00
62843151		T/C	Noncoding	0.00	0.05	0.00
62837154		A/C	intron	0.00	0.00	0.08
rs34927999		A/G	intron	0.00	0.05	0.08
62835790		G/C	Noncoding	0.00	0.00	0.08
62835817		A/C	Noncoding	0.05	0.03	0.08
62835752		T/C	Noncoding	0.00	0.14	0.00
62839481		C/T	intron	0.10	0.16	0.08
62836366		A/G	intron	0.05	0.00	0.17
62840163		T/C	intron	0.00	0.27	0.00
62842510		A/G	intron	0.00	0.30	0.00
62840278		T/C	intron	0.05	0.33	0.17
62843192		T/C	Noncoding	0.00	0.00	0.33
62836938		in/del	intron	0.00	0.49	0.00
rs10449755		A/C	Noncoding	0.00	0.81	0.25
62839551		C/T	intron	0.00	1.76	0.33
62840198		T/G	intron	0.05	1.84	0.25
62839294		in/del	intron	2.39	0.42	1.53

rs12565895	C/T	Noncoding	0.19	7.81	2.53
rs34483103	in/del	Noncoding	21.97	19.04	21.72
62836000	G/C	Nonsynonymous	0.00	0.03	0.00
62836013	A/C	Nonsynonymous	0.00	0.03	0.00
62836072	in/del	Nonsynonymous	0.00	0.03	0.00
62836097	A/G	Nonsynonymous	0.00	0.03	0.00
62836315	C/T	Nonsynonymous	0.00	0.03	0.00
62836977	A/G	Nonsynonymous	0.00	0.03	0.00
62840570	C/T	Nonsynonymous	0.00	0.03	0.00
62840582	T/C	Nonsynonymous	0.00	0.03	0.00
62842264	T/C	Nonsynonymous	0.00	0.03	0.00
62842408	A/G	Nonsynonymous	0.00	0.03	0.00
62842485	G/C	Nonsynonymous	0.00	0.03	0.00
62843027	in/del	Nonsynonymous	0.00	0.03	0.00
62836188	in/del	Nonsynonymous	0.05	0.03	0.00
62836217	A/C	Nonsynonymous	0.05	0.00	0.00
62836264	in/del	Nonsynonymous	0.05	0.03	0.00
62840571	G/A	Nonsynonymous	0.05	0.00	0.00
62840595	A/G	Nonsynonymous	0.05	0.00	0.00
62842294	T/G	Nonsynonymous	0.05	0.00	0.00
62842419	G/A	Nonsynonymous	0.05	0.00	0.00
62842443	A/T	Nonsynonymous	0.05	0.00	0.00
62842463	in/del	Nonsynonymous	0.05	0.00	0.00
62842969	C/G	Nonsynonymous	0.05	0.00	0.00
62842976	G/T	Nonsynonymous	0.05	0.00	0.00
62835901	T/C	Nonsynonymous	0.00	0.05	0.00
62842417	C/T	Nonsynonymous	0.00	0.05	0.00
62843050	C/A	Nonsynonymous	0.00	0.05	0.00
62835946	G/A	Nonsynonymous	0.00	0.00	0.08
62836117	G/A	Nonsynonymous	0.00	0.00	0.08
62842291	G/A	Nonsynonymous	0.00	0.00	0.08
62842943	A/G	Nonsynonymous	0.00	0.11	0.00
62837034	in/del	Nonsynonymous	0.00	0.14	0.00
62836053	C/G	Nonsynonymous	0.24	0.08	0.17
62836041	T/G	Nonsynonymous	0.00	0.00	0.25
62836204	C/T	Nonsynonymous	1.00	0.08	0.00
62840082	T/C	Nonsynonymous	0.10	4.16	0.33
62835870	T/C	Synonymous	0.00	0.03	0.00
62839425	T/C	Synonymous	0.00	0.03	0.00
62843007	T/C	Synonymous	0.00	0.03	0.00
62837024	T/C	Synonymous	0.05	0.05	0.12
62836197	T/C	Synonymous	0.00	0.11	0.17
62842418	G/A	Synonymous	0.00	0.33	0.00
rs12563308	T/C	Synonymous	0.24	8.09	2.74

ANGPTL4 (Chr 19)

8335155	C>A	intron	0.0000	0.0432	0.0000
8335162	T>A	intron	0.0000	0.0428	0.0000
8335185	G>T	intron	0.2660	8.1240	0.5495
8335189	C>T	intron	0.0000	0.1255	0.0000
8335207	T>T	Nonsynonymous	0.0000	0.0400	0.0000

8335219	C>T	Nonsynonymous	0.0000	0.0403	0.0000
8335323	G>A	Nonsynonymous	1.4650	0.3182	0.9474
8335328	G>C	Nonsynonymous	0.0000	0.0000	2.2105
8335349	C>T	Synonymous	0.0000	0.0400	0.0000
8335361	C>A	Synonymous	0.0000	0.0796	0.0000
8335406	C>G	Nonsynonymous	0.0635	0.0000	0.0000
8335420	G>T	Nonsynonymous	0.0000	0.1195	0.0000
8335434	G>A	Nonsynonymous	0.0000	0.0400	0.0000
8336793	G>A	intron	0.0631	0.0000	0.0000
8336802	A>T	intron	0.0698	0.0472	0.0000
8336810	C>T	intron	0.1397	0.1873	0.3371
8336811	C>T	intron	0.0000	0.0400	0.0000
8336819	G>T	intron	0.0000	0.0466	0.0000
8336918	G>A	Synonymous	0.0000	0.7504	0.2096
8336996	C>T	intron	0.0000	0.7504	0.2096
8337000	T>C	intron	0.3788	0.0395	0.1048
8337027	G>A	intron	0.0631	1.0285	0.1048
8337074	G>C	intron	0.0000	0.0400	0.0000
8337082	in/del	intron	0.0631	3.7915	0.1048
8337155	G>A	Nonsynonymous	0.0640	0.0000	0.0000
8337176	C>T	Nonsynonymous	0.0000	0.0404	0.0000
8337204	G>A	intron	0.0638	0.0000	0.0000
8337248	C>T	intron	0.0000	0.0400	0.0000
8337249	G>A	intron	0.0631	0.0000	0.0000
8337250	C>A	intron	0.0639	0.0000	0.0000
8340030	G>A	intron	0.0639	0.0000	0.0000
8340051	C>T	intron	0.0000	0.5556	0.0000
8340055	C>T	intron	0.0000	0.5733	0.0000
8340057	in/del	intron	0.1299	0.0820	0.2151
8340062	G>A	intron	0.0649	5.5419	0.3226
8340097	C>T	intron	0.0000	0.3419	0.0000
8340100	C>T	intron	0.0670	3.8462	0.1106
8340123	G>C	Nonsynonymous	0.2005	0.6399	0.6652
8340141	G>A	Nonsynonymous	0.0676	0.0000	0.0000
8340185	G>A	Synonymous	0.1362	8.9255	0.4454
8340204	A>T	Nonsynonymous	0.0649	0.0000	0.0000
8340226	G>A	intron	0.0000	0.2042	0.0000
8340229	in/del	intron	0.0671	0.0000	0.0000
8340249	C>T	intron	0.0000	0.2024	0.0000
8340295	in/del	intron	0.0000	0.1221	0.0000
8340299	in/del	intron	0.0000	0.0000	0.1087
8341802	C>T	intron	0.0649	0.0000	0.0000
8341829	G>A	intron	0.0000	0.0000	0.1050
8341879	in/del	intron	0.0000	0.0398	0.0000
8341894	G>A	intron	0.0000	1.0350	0.0000
8341945	G>C	Nonsynonymous	0.0635	0.0000	0.0000
8341966	C>T	Nonsynonymous	0.0000	0.0000	0.1050
8341987	T>G	Nonsynonymous	0.0649	0.0000	0.0000
8342012	in/del	Nonsynonymous	0.0635	0.0000	0.0000
8342029	C>A	Nonsynonymous	0.0635	0.0000	0.0000
8342126	C>T	Synonymous	0.0000	0.0789	0.0000

8342164	C>T	Nonsynonymous	29.9492	22.8175	35.9705
8342165	G>A	Synonymous	0.0000	0.0794	0.0000
8342200	G>A	Nonsynonymous	0.0636	5.7143	0.3165
8342216	C>T	Synonymous	0.0000	0.0397	0.0000
8342238	G>A	Nonsynonymous	0.0637	0.0000	0.0000
8342244	C>A	Nonsynonymous	0.0000	0.0397	0.0000
8342254	A>T	Nonsynonymous	0.0637	0.0000	0.0000
8342271	in/del	Nonsynonymous	0.0639	0.0000	0.0000
8342286	C>T	Nonsynonymous	0.0635	0.0000	0.0000
8342288	C>T	Synonymous	0.1904	11.6878	0.4228
8342289	G>A	Nonsynonymous	0.0635	0.0000	0.1057
8342321	A>T	Synonymous	0.0000	0.0396	0.0000
8342373	C>T	Nonsynonymous	0.1904	0.0000	0.1053
8342381	C>A	Nonsynonymous	0.0000	0.0396	0.0000
8342438	A>G	intron	0.2548	0.0000	0.3165
8344596	G>T	Nonsynonymous	0.0000	0.0402	0.0000
8344629	C>T	Synonymous	0.1926	8.6747	0.4292
8344630	G>C	Nonsynonymous	0.0642	0.0000	0.0000
8344630	G>A	Nonsynonymous	0.1284	0.0000	0.0000
8344638	C>T	Synonymous	0.0643	0.0000	0.0000
8344644	C>T	Synonymous	0.0000	0.0800	0.0000
8344661	G>A	Nonsynonymous	0.0000	0.0404	0.0000
8344699	C>T	Nonsynonymous	0.0000	0.0401	0.0000
8344716	G>A	Synonymous	15.0763	2.3600	5.9574
8344771	C>T	intron	0.0000	3.5515	0.3165
8344772	G>A	intron	0.0000	0.3990	0.0000
8344778	G>A	intron	0.1904	12.2408	0.4219
8344782	G>T	intron	0.0000	0.0399	0.0000
8344800	C>T	intron	0.0000	0.0800	0.0000
8344808	G>T	intron	0.0000	0.5591	0.1053
8344840	G>A	intron	0.0634	0.0000	0.0000
8344885	T>C	intron	0.0000	0.0447	0.0000
8344924	T>C	intron	0.0000	0.0453	0.0000
8344936	G>C	intron	0.0000	0.0642	0.0000
8344943	T>C	intron	0.0000	0.0400	0.0000
8344965	C>T	intron	0.0000	1.4091	0.0000

ANGPTL5 (Chr 11)

101281894	in/del	intron	0.00	0.03	0.00
101280686	C/A	intron	0.00	0.03	0.00
101281614	G/C	intron	0.00	0.03	0.00
101276365	C/A	intron	0.00	0.03	0.00
101278738	G/T	intron	0.00	0.03	0.00
101283858	T/C	intron	0.00	0.03	0.00
101281603	T/G	intron	0.00	0.03	0.00
101267641	C/T	intron	0.00	0.03	0.00
101278751	T/C	intron	0.00	0.03	0.00
101267033	G/C	non-coding	0.00	0.03	0.00
101278786	G/C	intron	0.00	0.03	0.00
101270720	A/G	intron	0.00	0.03	0.00
101283220	T/C	intron	0.00	0.03	0.00

101283951	G/A	intron	0.00	0.03	0.00
101283040	in/del	intron	0.03	0.03	0.00
101276518	T/C	intron	0.00	0.03	0.00
101270819	C/T	intron	0.05	0.00	0.00
101281829	G/A	intron	0.05	0.00	0.00
101281844	A/T	intron	0.05	0.00	0.00
101278553	T/C	intron	0.05	0.00	0.00
101281594	T/C	intron	0.05	0.00	0.00
101267080	A/G	non-coding	0.05	0.00	0.00
101284008	A/G	intron	0.06	0.00	0.00
101281643	T/C	intron	0.00	0.06	0.00
101280691	C/T	intron	0.00	0.06	0.00
101283025	A/T	intron	0.08	0.00	0.00
101278395	A/G	intron	0.00	0.09	0.00
101280906	A/T	intron	0.00	0.00	0.09
101283739	C/T	intron	0.00	0.00	0.09
101283647	A/G	intron	0.00	0.06	0.09
101267555	C/G	intron	0.00	0.00	0.09
101267574	C/A	intron	0.00	0.00	0.09
101278751	in/del	intron	0.00	0.03	0.09
101270705	C/T	intron	0.00	0.00	0.09
101283214	in/del	intron	0.00	0.00	0.09
101283285	A/G	intron	0.00	0.00	0.10
101270753	A/G	intron	0.00	0.14	0.00
101278833	G/A	intron	0.00	0.17	0.00
101281562	G/T	intron	0.00	0.00	0.18
101280915	G/A	intron	0.00	0.20	0.00
101270797	A/T	intron	0.00	0.00	0.26
101271057	in/del	intron	0.00	0.00	0.26
101266976	G/C	non-coding	0.30	0.00	0.10
101278769	in/del	intron	0.00	0.00	0.31
101271010	A/G	intron	0.00	0.43	0.00
101283241	G/T	intron	0.05	0.58	0.09
101280698	C/T	intron	0.05	0.64	0.00
101266954	C/T	non-coding	0.65	0.00	0.30
101281629	T/C	intron	0.00	1.26	0.26
101280687	A/C	intron	0.00	1.46	0.26
101278494	C/T	intron	0.05	1.55	0.27
101278717	T/A	intron	0.00	1.83	0.09
101278803	C/T	intron	0.00	2.04	0.27
101280883	T/C	intron	1.77	0.39	2.11
101281847	T/C	intron	0.05	4.02	0.26
101270715	C/T	intron	0.00	4.09	1.11
101280934	G/C	intron	3.43	4.50	2.60
101276359	T/A	intron	4.24	2.18	8.68
101270814	G/T	intron	4.55	6.82	11.11
101283920	G/A	intron	26.46	15.54	21.40
101267688	in/del	intron	26.51	21.17	21.08
101280873	T/G	intron	27.01	16.38	21.55
101276591	C/T	intron	19.74	20.83	29.24
101283756	T/C	Nonsynonymous	0.00	0.03	0.00

101267495	G/C	Nonsynonymous	0.00	0.03	0.00
101267290	C/T	Nonsynonymous	0.00	0.03	0.00
101270843	A/G	Nonsynonymous	0.00	0.03	0.00
101283102	in/del	Nonsynonymous	0.00	0.03	0.00
101270909	A/G	Nonsynonymous	0.05	0.00	0.00
101270970	A/C	Nonsynonymous	0.05	0.03	0.00
101267302	G/A	Nonsynonymous	0.05	0.00	0.00
101270979	C/G	Nonsynonymous	0.00	0.06	0.00
101270994	in/del	Nonsynonymous	0.00	0.06	0.00
101270838	T/G	Nonsynonymous	0.00	0.06	0.00
101276377	A/G	Nonsynonymous	0.06	0.00	0.00
101267510	A/T	Nonsynonymous	0.00	0.06	0.00
101280821	C/T	Nonsynonymous	0.00	0.00	0.09
101267405	G/A	Nonsynonymous	0.10	0.00	0.00
101276449	T/C	Nonsynonymous	0.00	0.00	0.10
101278579	A/G	Nonsynonymous	0.00	0.11	0.09
101267401	C/T	Nonsynonymous	0.05	0.20	0.09
101281752	A/G	Nonsynonymous	0.00	0.00	0.26
101270862	C/T	Nonsynonymous	0.00	0.28	0.00
101276457	T/G	Nonsynonymous	0.36	0.27	0.41
101267480	A/G	Nonsynonymous	0.00	0.43	0.00
101267229	in/del	Nonsynonymous	0.05	0.68	0.00
101281737	G/A	Nonsynonymous	0.10	0.08	0.86
101267462	T/C	Nonsynonymous	0.00	1.07	0.09
101270864	T/C	Nonsynonymous	2.93	3.30	1.65
101280843	C/T	Synonymous	0.00	0.03	0.00
101270908	C/G	Synonymous	0.05	0.00	0.00
101278652	A/C	Synonymous	0.05	0.03	0.00
101267328	C/T	Synonymous	0.05	0.00	0.00
101283797	A/T	Synonymous	0.00	0.00	0.09
101278601	C/T	Synonymous	0.00	0.00	0.09
101267379	C/T	Synonymous	0.00	0.32	0.09
101267394	A/G	Synonymous	0.00	0.57	0.00
101267223	T/G	Synonymous	2.25	0.34	1.58

ANGPTL6 (Chr 19)

10065351	G/C	intron	0.00	0.03	0.00
10064135	C/T	non-coding	0.00	0.03	0.00
10066634	G/A	intron	0.00	0.03	0.00
10066268	A/C	intron	0.00	0.03	0.00
10065574	C/T	intron	0.06	0.00	0.00
10065618	C/G	intron	0.06	0.00	0.00
10064105	C/T	non-coding	0.06	0.00	0.00
10064474	A/G	intron	0.06	0.00	0.00
10064172	A/G	non-coding	0.00	0.06	0.00
10066257	G/A	intron	0.00	0.07	0.00
10064252	C/T	non-coding	0.00	0.00	0.09
10065562	C/G	intron	0.00	0.03	0.09
10065024	G/A	intron	0.00	0.00	0.09
10066320	T/G	intron	0.06	0.00	0.10
10066356	A/C	intron	0.00	0.00	0.10

10064466	T/C	intron	0.00	0.13	0.00
10064984	G/A	intron	0.00	0.13	0.00
10065338	G/A	intron	0.17	0.00	0.00
10065339	G/A	intron	0.17	0.00	0.00
10065575	C/G	intron	0.06	0.20	4.73
10066618	C/T	intron	0.06	0.03	5.02
10066329	C/T	intron	0.36	0.57	5.15
10064171	T/C	non-coding	9.39	3.90	3.86
10067864	G/C	Nonsynonymous	0.00	0.02	0.00
10067836	C/T	Nonsynonymous	0.00	0.03	0.00
10064362	G/A	Nonsynonymous	0.00	0.03	0.00
10065052	G/A	Nonsynonymous	0.00	0.03	0.00
10068107	A/C	Nonsynonymous	0.00	0.04	0.00
10067774	C/T	Nonsynonymous	0.00	0.04	0.00
10064432	G/C	Nonsynonymous	0.06	0.03	0.00
10065478	A/G	Nonsynonymous	0.06	0.00	0.00
10065481	G/A	Nonsynonymous	0.06	0.00	0.00
10065243	A/G	Nonsynonymous	0.06	0.00	0.00
10067973	G/A	Nonsynonymous	0.06	0.00	0.00
10067997	G/C	Nonsynonymous	0.06	0.03	0.00
10067858	C/G	Nonsynonymous	0.00	0.06	0.00
10068100	C/G	Nonsynonymous	0.06	0.00	0.00
10067763	G/T	Nonsynonymous	0.07	0.00	0.00
10064372	G/A	Nonsynonymous	0.00	0.00	0.09
10067908	G/A	Nonsynonymous	0.00	0.00	0.09
10064324	C/T	Nonsynonymous	0.00	0.00	0.09
10064357	C/T	Nonsynonymous	0.00	0.00	0.09
10064294	G/A	Nonsynonymous	0.00	0.03	0.09
10065461	T/G	Nonsynonymous	0.00	0.00	0.09
10065491	C/T	Nonsynonymous	0.00	0.00	0.09
10065244	T/C	Nonsynonymous	0.00	0.00	0.09
10068066	C/G	Nonsynonymous	0.00	0.00	0.10
10068068	A/G	Nonsynonymous	0.00	0.00	0.10
10068069	C/T	Nonsynonymous	0.00	0.00	0.10
10068070	C/T	Nonsynonymous	0.00	0.00	0.10
10068073	C/G	Nonsynonymous	0.00	0.00	0.10
10068024	C/G	Nonsynonymous	0.00	0.11	0.00
10064434	G/A	Nonsynonymous	0.11	0.00	0.00
10065469	C/G	Nonsynonymous	0.11	0.00	0.00
10065229	C/T	Nonsynonymous	0.17	0.00	0.00
10067848	A/T	Nonsynonymous	0.33	0.03	0.36
10067988	in/del	Nonsynonymous	0.00	0.00	0.40
10065175	C/T	Nonsynonymous	1.90	0.32	0.74
10067953	T/C	Nonsynonymous	2.21	0.32	0.36
10065275	C/T	Synonymous	0.00	0.03	0.00
10065534	G/A	Synonymous	0.00	0.03	0.00
10066579	G/A	Synonymous	0.00	0.03	0.00
10068096	A/G	Synonymous	0.00	0.04	0.00
10067826	G/A	Synonymous	0.06	0.00	0.00
10067805	C/T	Synonymous	0.06	0.00	0.09
10065494	C/T	Synonymous	0.06	0.10	0.00

10068072	C/A	Synonymous	0.00	0.00	0.10
10065143	C/T	Synonymous	1.05	0.00	0.37
10068048	G/A	Synonymous	0.06	0.53	4.33
10065230	C/T	Synonymous	0.06	0.03	4.83

Abbreviations: in, insertion; del, deletion

Supplemental Table 2. Properties of sequence variations identified in *ANGPTL3*, *ANGPTL4*, *ANGPTL5*, and *ANGPTL6*

	All	<i>ANGPTL3</i>	<i>ANGPTL4</i>	<i>ANGPTL5</i>	<i>ANGPTL6</i>
Total	347	87	92	98	70
Singletons	200	58	49	51	42
MAF<0.01	294	79	74	80	61
MAF<0.050	326	84	83	92	67
NS Total	129	35	32	26	36
NS Singletons	84	27	20	12	25
NS MAF<0.01	120	33	28	25	34
NS MAF<0.050	126	34	30	26	36
S Total	40	7	13	9	11
S Singletons	21	3	6	6	6
S MAF<0.01	31	6	9	8	8
S MAF<0.050	35	6	9	9	11
NC Total	178	45	47	63	23
NC Singletons	103	28	30	34	11
NC MAF<0.01	144	40	37	48	19
NC MAF<0.050	164	43	44	57	20
Black only	140	46	39	35	20
White Only	67	14	23	14	16
Hispanic Only	45	7	5	16	17

Table entries are numbers of sequence variants identified.

Abbreviations: MAF, minor allele frequency; NS, nonsynonymous; S, synonymous; NC, noncoding

Supplementary Table 3. Clinical characterization of African-American in the DHS and ARIC study stratified by *ANGPTL3* genotype (M259T).

	DHS (n=1,521)				ARIC (n=3,682)			
	MM	MT	TT	P Value ^A	MM	MT	TT	P Value ^A
n	1380	138	3	-	3308	361	13	-
Male/Female	585/795	58/80	0/3	0.987	1242/2066	151/210	4/9	0.170
Age (years)	44.0±10	44.0±9	44.0±8	0.500	53.6±6	52.9±6	54.2±6	0.104
BMI (kg/m ²)	32.0±8	31.6±10	35.8±4	0.748	29.6±6	29.2±6	28.6±4	0.564
SBP (mm/Hg)	129±20	133±22	132±28	0.064	129±21	128±21	128.6±15	0.885
DBP (mm/Hg)	81±11	82±11	79±15	0.287	78±12	80±12	79±10	0.890
Glucose (mg/dL)	104±49	98±26	105±22	0.114	118±57	117±56	120±55	0.892
Insulin (mIU/L)	14.1(15)	11.4(14)	21,0 (4)	0.220	12.4(12)	11.4(11)	12.4(9)	0.544
Cholesterol (mg/dL)	178±41	173±32	154±10	0.210	215±45	206±39	219±32	0.006
Triglyceride (mg/dL)	84(61)	73(49)	64(47)	0.006	97(64)	88(54)	80(44)	0.014
LDL-C (mg/dL)	105±36	102±33	93±6	0.515	138.1±43	130±37	143±25	0.005
HDL-C (mg/dL)	52±15	55±17	52±7	0.259	55±18	56±18	56±15	0.414
HTGC (%)	3.3 (3.4)	3.2 (3.1)	4.0 (1.9)	0.615	-	-	-	-

^AValues are means ± standard deviations, excluding hepatic triglycerides content (HTGC) and plasma fasting insulin and triglyceride levels which are medians (interquartile ranges). P values were calculated using linear regression model including gender, age and diabetes mellitus status as covariates and after excluding individuals on lipid-lowering medications. BMI, body mass index; SBP, systolic blood pressure; DBP, diastolic blood pressure. HTGC was measured using proton magnetic resonance spectroscopy.

Supplementary Table 4: Oligonucleotide Primers used for Cloning

h-ANGPTL3F	CACCATGTTTACAATTAAGCTCCTTC
h-ANGPTL3R	TTCAAAGCTTTCTGAATCTGTTGGAT
h-ANGPTL4F	CACCATGAGCGGTGCTCCGACGGC
h-ANGPTL4R	GGAGGCTGCCTCTGCTGCCA
h-ANGPTL5F	ATGATGTCTCCATCCCAAGCCTCACTCTTA
h-ANGPTL5R	AGAAGAATGTACAATCCATATTTTAAATAA
h-ANGPTL6F	ATGGGGAAGCCCTGGCTGCGTGCG
h-ANGPTL6R	TCACAGCTTCAGGGGCCGAATGAGCAT