

559 **Supplementary Materials**

<b>Supplementary Table 1</b>	<b>Adjusted rate ratios (RR) and 95% confidence intervals (CI) for risk of ischaemic stroke (IS; N = 5475) and intracerebral haemorrhage (ICH; N = 4776) by concentrations of major blood lipids in observational analyses in CKB, with and without adjustment for body mass index (BMI)</b>
<b>Supplementary Table 2</b>	<b>Characteristics of the 46 LDL-C associated genetic variants obtained from the Global Lipids Genetics Consortium</b>
<b>Supplementary Table 3</b>	<b>Summary-level Mendelian randomisation sensitivity analyses of associations of genetically-instrumented LDL-C with risk of ischaemic stroke (N = 5567) and intracerebral haemorrhage (N = 4911) in CKB</b>
<b>Supplementary Table 4</b>	<b>Age-specific rates of ischaemic stroke, intracerebral haemorrhage, and major coronary events in the CKB prospective study, by different levels of background vascular risk</b>

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**Supplementary Table 1: Adjusted rate ratios (RR) for risk of ischaemic stroke (IS; N = 5475) and intracerebral haemorrhage (ICH; N = 4776) by concentrations of major blood lipids in observational analyses in CKB, with and without adjustment for body mass index (BMI)<sup>a</sup>**

Major blood lipids (SD)	RR (95% CI) per SD higher				P for heterogeneity between IS and ICH	
	Ischaemic stroke (N = 5475)		Intracerebral haemorrhage (N = 4776)			
	Without BMI	With BMI	Without BMI	With BMI		
<b>LDL-C</b> (0.68 mmol/L)	1.12 (1.07-1.16)	1.10 (1.06-1.15)	0.90 (0.86-0.95)	0.91 (0.87-0.96)	P=4.2x10 <sup>-11</sup>	
<b>HDL-C</b> (0.29 mmol/L)	0.93 (0.89-0.97)	0.94 (0.90-0.99)	1.00 (0.96-1.05)	0.98 (0.94-1.02)	P=0.01	
<b>Triglycerides<sup>b</sup></b> 0.58	1.05 (1.00-1.10)	1.03 (0.98-1.08)	0.87 (0.82-0.91)	0.88 (0.84-0.93)	P=1.3x10 <sup>-8</sup>	

SD=Standard deviation; CI=Confidence intervals.

<sup>a</sup> Cox regression was used to estimate the rate ratios (95% CI), as shown in column 2, 3, 4, and 5, per SD unit higher concentrations of measured blood lipids. The analyses were stratified by age-at-risk (5-year), sex, and study area, and adjusted for education, smoking, alcohol consumption, physical activity, diabetes, and baseline systolic blood pressure, with and without adjustment for BMI. Chi-square test was used for heterogeneity. All P-values (two-sided) were uncorrected for multiple testing.

<sup>b</sup> Values for triglycerides are natural logarithm of triglyceride measurements.

**Supplementary Table 2. Characteristics of the 46 LDL-C associated genetic variants obtained from the Global Lipids Genetics Consortium**

Marker	Locus	Chr	Position <sup>a</sup>	Effect Allele	Other Allele	GLGC		CKB <sup>b</sup>	
						Beta	P-value	Beta	P-value
rs10903129	<i>MACO1</i>	1	25768937	G	A	0.033	4E-19	0.034	4E-03
rs1998013 <sup>c</sup>	<i>RNU6-830P</i>	1	55958030	C	T	0.381	4E-67	NA	NA
rs646776	<i>CELSR2</i>	1	109818530	T	C	0.161	1E-292	0.209	2E-20
rs267733	<i>ANXA9</i>	1	150958836	A	G	0.033	5E-10	-0.013	6E-01
rs2642438	<i>MARC1</i>	1	220970028	G	A	0.035	4E-17	0.061	3E-05
rs1367117	<i>APOB</i>	2	21263900	A	G	0.119	2E-196	0.102	2E-10
rs515135	<i>APOB</i>	2	21286057	C	T	0.139	1E-188	-0.011	6E-01
rs6544713	<i>ABCG8</i>	2	44073881	T	C	0.081	6E-85	-0.107	1E-01
rs2710642	<i>EHBP1</i>	2	63149557	A	G	0.024	3E-10	0.031	8E-03
rs2030746	<i>LOC105373585</i>	2	121309488	T	C	0.021	2E-08	0.006	6E-01
rs2287623	<i>ABCB11</i>	2	169830155	G	A	0.022	7E-09	0.031	1E-02
rs1250229	<i>LOC105373868</i>	2	216304384	C	T	0.024	8E-09	0.034	2E-01
rs11563251	<i>UGT1A</i>	2	234679384	T	C	0.035	2E-08	0.021	3E-01
rs7640978	<i>CMTM6</i>	3	32533010	C	T	0.039	1E-08	-0.037	1E-01
rs17345563	<i>DNAJC13</i>	3	132209203	A	G	0.036	3E-10	0.016	4E-01
rs7703051	<i>ANKRD31</i>	5	74625487	A	C	0.073	5E-85	0.081	3E-14
rs4530754	<i>CSNK1G3</i>	5	122855416	A	G	0.028	4E-14	-0.013	2E-01
rs6882076	<i>TIMD4</i>	5	156390297	C	T	0.046	5E-33	0.059	1E-06
rs1800562	<i>HFE</i>	6	26093141	G	A	0.062	2E-14	-0.012	9E-01
rs1564348	<i>SLC22A1</i>	6	160578860	C	T	0.048	3E-22	0.075	4E-01
rs12670798	<i>DNAH11</i>	7	21607352	C	T	0.034	7E-16	0.019	7E-02
rs4722551	<i>LOC105375199</i>	7	25991826	C	T	0.039	7E-16	0.031	3E-01
rs2073547	<i>NPC1L1</i>	7	44582331	G	A	0.049	5E-23	-0.003	8E-01
rs217386	<i>NPC1L1</i>	7	44600695	G	A	0.036	8E-22	0.013	7E-01
rs10102164	<i>TRMT112P7</i>	8	55421614	A	G	0.032	3E-12	0.044	7E-04
rs3780181	<i>VLDLR</i>	9	2640759	A	G	0.045	1E-09	0.063	3E-03
rs8176720	<i>ABO</i>	9	136132873	T	C	0.033	6E-18	0.034	2E-03
rs579459	<i>ABO</i>	9	136154168	C	T	0.067	3E-49	0.088	9E-12
rs174532	<i>MYRF</i>	11	61548874	A	G	0.035	5E-17	0.068	6E-01
rs1535	<i>FADS2</i>	11	61597972	A	G	0.053	3E-43	0.041	2E-04
rs11220462	<i>ST3GAL4</i>	11	126243952	A	G	0.059	3E-23	0.029	1E-02
rs1186380	<i>RPL12P33</i>	12	121376416	C	T	0.024	1E-08	0.015	2E-01
rs1169288	<i>HNF1A</i>	12	121416650	C	A	0.038	9E-21	0.018	1E-01
rs4942486	<i>BRCA2</i>	13	32953388	T	C	0.024	3E-11	0.041	2E-04
rs8017377	<i>NYNRIN</i>	14	24883887	A	G	0.031	3E-15	0.062	8E-03
rs2000999	<i>TXNL4B</i>	16	72108093	A	G	0.065	1E-45	0.033	6E-03
rs314253	<i>ASGR1</i>	17	7091650	T	C	0.024	2E-10	0.031	5E-03
rs4791641	<i>PFAS</i>	17	8161149	C	T	0.021	5E-08	-0.007	6E-01
rs6511720	<i>LDLR</i>	19	11202306	G	T	0.221	3E-289	0.184	3E-03
rs688	<i>LDLR</i>	19	11227602	T	C	0.054	9E-48	0.077	6E-08
rs6859	<i>NECTIN2</i>	19	45382034	A	G	0.084	1E-101	-0.041	5E-04
rs7254892	<i>NECTIN2</i>	19	45389596	G	A	0.485	8E-365	0.702	9E-246
rs492602	<i>FUT2</i>	19	49206417	G	A	0.029	3E-14	0.043	5E-01
rs364585	<i>LINC01722</i>	20	12962718	G	A	0.025	4E-11	-0.003	8E-01
rs2328223	<i>LOC107985440</i>	20	17845921	C	A	0.031	2E-09	0.021	1E-01
rs5763662	<i>MEMR3</i>	22	30378703	T	C	0.077	2E-10	0.022	2E-01

Chr.=Chromosome; Effect alleles are given as the LDL-C raising allele in the Global Lipids Genetics Consortium data<sup>49,50</sup>;

<sup>a</sup>Genomic coordinates were in Build hg19;

<sup>b</sup>The analyses were conducted in 17,567 CKB participants with available data. General linear regression was used to estimate SD differences in LDL-C (after rank-inverse-normal transformation) per 1 effect allele, adjusted for sex, age, age-squared, and case status. The analyses were performed for each study area, and the overall estimates were obtained by inverse-variance-weighted meta-analyses. All P-values (two-sided) were uncorrected for multiple testing.

<sup>c</sup>Monogenic in the present study.

**Supplementary Table 3: Summary-level Mendelian randomisation sensitivity analyses of associations of genetically-instrumented LDL-C with risk of ischaemic stroke (N = 5567) and intracerebral haemorrhage (N = 4911) in CKB<sup>a</sup>**

Methods	Ischaemic stroke (N = 5567)					Intracerebral haemorrhage (N = 4911)				
	Beta	SE	LCI	UCI	P-value	Beta	SE	LCI	UCI	P-value
Main analyses:	0.282	0.117	0.054	0.511	0.015	-0.122	0.109	-0.337	0.092	0.264
Sensitivity analyses:										
Weighted median	0.126	0.122	-0.114	0.366	0.304	-0.072	0.114	-0.296	0.152	0.530
Inverse-variance weighted method	0.210	0.114	-0.014	0.433	0.066	-0.099	0.100	-0.295	0.097	0.322
MR-Egger (Causal estimates)	0.101	0.131	-0.156	0.359	0.440	-0.056	0.118	-0.287	0.174	0.631
MR-Egger (Intercept)	0.011	0.007	-0.002	0.024	0.112	-0.004	0.006	-0.016	0.008	0.484

SE=standard error; LCI=lower confidence interval; UCI=upper confidence interval.

<sup>a</sup> The Betas and SEs were estimated using published methods with R package<sup>51</sup>, per 1 mmol/L higher concentrations of genetically-instrumented LDL-C. All P-values (two-sided) were uncorrected for multiple testing.

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**Supplementary Table 4: Age-specific rates of ischaemic stroke, intracerebral haemorrhage, and major coronary events in the CKB prospective study, by different levels of background vascular risk**

Age at risk	Ischemic stroke				Intracerebral hemorrhage				Major coronary events			
	Person-years	No. of events	Rates (% p.a.)	Events avoided	Person-years	No. of events	Rates (% p.a.)	Excess events	Person-years	No. of events	Rates (% p.a.)	Events avoided
<b>Low-risk (n=336,696)</b>												
<50	1,258,546	1,356	0.11	11 (1)	1,261,482	324	0.03	-2 (1)	1,262,025	209	0.02	2 (0)
50-60	979,911	4,041	0.41	41 (4)	990,273	538	0.05	-5 (2)	990,637	507	0.05	6 (1)
60-70	558,263	4,636	0.83	83 (9)	571,871	739	0.13	-11 (5)	572,227	753	0.13	17 (1)
≥70	226,053	4,470	1.98	198 (20)	240,434	929	0.39	-33 (14)	240,596	1,149	0.48	60 (4)
Overall	3,022,772	14,503	0.48	48 (5)	3,064,060	2,530	0.08	-7 (3)	3,065,485	2,618	0.09	11 (1)
<b>Medium-risk (n=153,066)</b>												
<50	228,332	960	0.42	42 (4)	229,893	459	0.2	-17 (7)	230,589	158	0.07	9 (1)
50-60	408,045	3,788	0.93	93 (9)	416,966	1,238	0.3	-25 (11)	419,006	513	0.12	15 (1)
60-70	404,001	6,427	1.59	159 (16)	422,308	1,860	0.44	-37 (16)	424,513	1,090	0.26	32 (2)
≥70	258,092	7,191	2.79	279 (28)	281,546	2,183	0.78	-66 (28)	282,573	2,212	0.78	98 (6)
Overall	1,298,469	18,366	1.41	141 (15)	1,350,713	5,740	0.42	-36 (15)	1,356,680	3,973	0.29	37 (2)
<b>High-risk (n=23,129)</b>												
<50	9,394	140	1.49	149 (15)	9,666	58	0.6	-51 (21)	9,720	41	0.42	53 (3)
50-60	37,871	1,025	2.71	271 (28)	40,387	262	0.65	-55 (23)	40,739	173	0.42	53 (3)
60-70	64,497	2,486	3.85	385 (39)	72,127	534	0.74	-63 (26)	72,534	500	0.69	86 (5)
≥70	57,237	3,129	5.47	547 (56)	69,696	604	0.87	-74 (31)	69,428	1,029	1.48	185 (11)
Overall	169,000	6780	4.01	401 (41)	191,876	1,458	0.76	-65 (27)	192,421	1,743	0.91	113 (7)
<b>Total</b>	<b>4,490,241</b>	<b>39,649</b>	<b>0.88</b>	<b>88 (9)</b>	<b>4,606,649</b>	<b>9,728</b>	<b>0.21</b>	<b>-18 (8)</b>	<b>4,614,586</b>	<b>8,334</b>	<b>0.18</b>	<b>23 (1)</b>

Low-risk: no measured hypertension, or prior history of cardiovascular disease (coronary heart disease, stroke or transient ischemic attack); Medium-risk: with measured hypertension, but with no prior history of cardiovascular disease; High-risk: with prior history of cardiovascular disease. Participants were considered to have measured hypertension if they had measured systolic blood pressure of at least 140 mm Hg or a measured diastolic blood pressure of at least 90 mm Hg or were receiving treatment for hypertension. The latter was defined as those who reported a diagnosis of hypertension by a physician and use of anti-hypertensives at recruitment.