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3 **Impact of concomitant Lp(a) level on LDL-cholesterol or apolipoproteinB-related risk**
4 **for incident coronary heart disease**

5 *(Arnold et al. Lp(a) and LDL-C or apoB-related CHD risk)*

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104 **Abstract**

105 **Background:** Conventional LDL-cholesterol (LDL-C) quantification includes cholesterol
106 attributable to lipoprotein (a) (Lp(a)-C) due to their overlapping densities.

107 **Objectives:** To compare the association between LDL-C and LDL-C corrected for Lp(a)-C
108 ($LDL_{Lp(a)corr}$) with incident coronary heart disease (CHD) in the general population and to
109 investigate whether concomitant Lp(a) values influence the association of LDL-C or apolipo-
110 protein B (apoB) with coronary events.

111 **Methods:** Among 68,748 CHD-free subjects at baseline $LDL_{Lp(a)corr}$ was calculated as “LDL-
112 C—Lp(a)-C”, where Lp(a)-C was 30% or 17.3% of total Lp(a) mass. Fine and Gray compet-
113 ing risk-adjusted models were applied for the association between the outcome incident CHD
114 and 1) LDL-C and $LDL_{Lp(a)corr}$ in the total sample; 2) LDL-C and apoB after stratification by
115 Lp(a) mass (\geq / $<$ 90th percentile (pctl.)).

116 **Results:** Similar risk estimates for incident CHD were found for LDL-C and $LDL_{Lp(a)corr30}$
117 or $LDL_{Lp(a)corr17.3}$ (sub-distribution Hazard Ratios (sHRs) with 95% CI) were 2.73 (2.34-
118 3.20) vs 2.51 (2.15-2.93) vs 2.64 (2.26- 3.10), respectively (top vs bottom fifth; fully-adjusted
119 models). Categorization by Lp(a) mass resulted in higher sHRs for uncorrected LDL-C and
120 incident CHD at $Lp(a)\geq 90^{th}$ pctl. (4.38 (2.08-9.22)) vs 2.60 (2.21-3.07) at $Lp(a)<90^{th}$ pctl. (top
121 vs bottom fifth; $p_{interaction}$ 0.39). In contrast, apoB risk estimates were lower in subjects with
122 higher Lp(a) mass (2.43 (1.34-4.40)) than in $Lp(a)<90^{th}$ pctl. (3.34 (2.78-4.01) ($p_{interaction}$ 0.49).

123 **Conclusion:** Correction of LDL-C for its Lp(a)-C content provided no meaningful infor-
124 mation on CHD-risk estimation at the population level. However, simple categorization of
125 Lp(a) mass (\geq / $<$ 90thpctl.) influenced the association between LDL-C or apoB with future
126 CHD mostly at higher Lp(a) levels.

127 **Key Words:** Lipoprotein (a), low-density lipoprotein, apolipoprotein B, coronary heart dis-
128 ease, general population

129 **Condensed Abstract:**

130 Correction of LDL-Cholesterol (LDL-C) for concomitant Lp(a) cholesterol represents a mat-
131 ter of debate. Among 68,748 subjects, who developed a CHD event over a median follow-up
132 of 9.72 years, similar association with future outcomes was found between uncorrected LDL-
133 C and $LDL_{Lp(a)corr}$, corrected for Lp(a) cholesterol (assumed as 30% or 17.3% of total Lp(a)
134 mass). In contrast, a simple categorization by Lp(a) values (\geq / $<$ 90thpercentiles) modified the
135 association between LDL-C or apolipoprotein B with future CHD mainly at higher Lp(a)
136 levels. Thus, an assessment of the conventional lipid profile without taking into account ac-
137 companying Lp(a) values might provide incomplete information on CHD risk.

138
139 **Abbreviations list:**

140 Lp(a) = lipoprotein (a)
141 LDL-C = low-density lipoprotein cholesterol
142 apoB = apolipoprotein B
143 CHD = coronary heart disease
144 ASCVD = atherosclerotic cardiovascular disease
145 FU = follow-up
146 IQR = interquartile range
147 sHRs = sub-distribution Hazard Ratios
148 CI = confidence interval
149 BMI = body mass index

150 **Introduction**

151 Low density lipoprotein (LDL)-targeted therapy has become a cornerstone in the management
152 of atherosclerotic cardiovascular disease (ASCVD) (1). Over the past decade, however,
153 emerging evidence has suggested that conventional assays for LDL-Cholesterol (LDL-C)
154 measurement quantify cholesterol attributable to a composite of atherogenic lipoproteins.
155 “LDL-C” not only measures cholesterol bound to LDL, but also cholesterol bound to inter-
156 mediate-density lipoprotein (IDL-C) and lipoprotein(a)- (Lp(a)-C) due to the overlapping
157 densities of these lipoproteins (1-2). This methodological limitation might have a significant
158 clinical impact, particularly in the setting of high Lp(a) (3). Whilst a fasting status results in
159 almost negligible contribution of IDL-C to measured LDL-C, an elevated Lp(a)-C could, by
160 contrast, account for a substantial proportion of conventionally measured LDL-C and in this
161 scenario, the real cholesterol content of LDL would be much lower than previously appreciat-
162 ed.

163 This assumption has led to the introduction of so called “corrected LDL-C”
164 ($LDL_{Lp(a)corr}$), i.e. a LDL-C without taking into account its Lp(a)-C content. Commonly,
165 Lp(a)-C has been calculated as 30% of Lp(a) mass, a correction factor derived from early
166 studies (3-5). However, one recent study in a small population measuring Lp(a)-C directly
167 showed a much higher variability of Lp(a)-C related to Lp(a) mass, ranging from 6 to 57%
168 (6).

169 Lp(a) measurement has recently gained increasing attention (7-8) and the current
170 ESC/EAS guidelines on dyslipidemia management recommend the measurement of Lp(a) at
171 least once in a person’s lifetime (1). Nonetheless, Lp(a) testing in the real world still remains
172 low (9-10), despite its prominent role in atherogenesis and the potentially meaningful contri-
173 bution of Lp(a)-C content to overall measured LDL-C. Although the relevance of this meth-
174 odological limitation is not completely appreciated, it might be important at least in two clini-

175 cal situations. First, extremely high Lp(a) levels might “create” low- or non-responders to
176 statin therapy, since Lp(a)-C would probably reflect the significant proportion of overall
177 measured LDL-C, which can not be lowered by statins. Second, it could also have a relevant
178 impact on the diagnosis of familial hypercholesterolemia (FH), where high Lp(a) might mim-
179 ic the classical monogenic form of FH simply by the contribution of Lp(a)-C to measured
180 LDL-C (11-12). A hitherto unanswered question is whether the methodological limitation of
181 LDL-C measurement could also be clinically important within the general population, where
182 the vast majority of subjects have only moderately elevated Lp(a) levels.

183 In the present analysis we therefore aimed to compare the association between LDL-C
184 and $LDL_{Lp(a)corr}$ and incident coronary heart disease (CHD) in the general population. Fur-
185 thermore, to circumvent the potential inaccuracies of the conventionally used methods to cor-
186 rect LDL-C for Lp(a)-C, we investigated whether the association between uncorrected LDL-C
187 and future CHD events might be affected particularly by concomitant Lp(a) levels and wheth-
188 er a similar pattern of association might be observed for other lipid parameters, such as
189 apolipoprotein B (apoB).

190 **Material and Methods:**

191 Study Design, Study Population and Outcome

192 The design and rationale of the BiomarCaRE (Biomarker for Cardiovascular Risk assessment
193 across Europe; <http://www.biomarcare.eu>) consortium have been published elsewhere (13).
194 Briefly, based on the Monitoring of Trends and Determinants in Cardiovascular Diseases
195 (MONICA) Risk Genetics Archiving and Monograph (MORGAM) Project, BiomarCaRE
196 represents a EU-funded initiative, which harmonized data from population-based cohorts
197 across Europe.

198 All participating cohorts obtained approval by the responsible local ethical review
199 boards. Participation was voluntary and written informed consent was obtained from each
200 subject upon entry into the study. This study was performed according to the principles of
201 Good Clinical Practice and the Declaration of Helsinki.

202 For the present analysis, data from 10 cohorts were used, resulting in a total of 93,313
203 individuals. Detailed cohort descriptions, including enrollment and follow-up procedures are
204 provided elsewhere (13). Two cohorts were excluded from the analysis due to missing data on
205 Lp(a) (PRIME/Belfast) or an analytical issue in Lp(a) determination (DAN-MONICA, due to
206 significant variations in Lp(a) levels between three surveys, compared to the remaining Bi-
207 omarCARE cohorts). After further exclusion of subjects with missing information on CHD
208 and Lp(a), as well as those with prevalent CHD at baseline, the final study sample comprised
209 68,748 CHD-free subjects (Northern Sweden (n=8,774), FINRISK (n=6,048), SHHEC (Scot-
210 tish Heart Health Extended Cohort) (n=12,585), MONICA/KORA Augsburg (Cooperative
211 Health Research in the Region of Augsburg), (n=7,405), MATISS (Malattie ATerosclerotiche
212 Istituto Superiore di Sanità) cohort (n=3,081), MONICA Brianza (n=4,303), Moli-Sani
213 (n=21,640), MONICA Catalonia (n=4,912)). The study population was further stratified ac-

214 cording to accompanying Lp(a) values with the 90th percentile chosen as the cut-off (<90th
215 pctl.: n=61,861; ≥90th pctl.: n=6,887). A flowchart of the study is presented in Figure 1.

216 All study participants were followed-up prospectively for an overall range of 2.5-25
217 years for incident CHD events, defined as fatal or non-fatal (definite or possible) myocardial
218 infarction (MI), coronary death, unstable angina pectoris, cardiac revascularization, and un-
219 classifiable death (i.e. death with insufficient evidence of coronary origin and no competing
220 cause). Most centers adjudicated the events using MONICA diagnostic criteria. The MOR-
221 GAM manual provides further information on endpoint classifications (14).

222 Data collection and risk factor definition

223 For detailed information on data collection and risk factor definition please see the **online**
224 **data supplement.**

225 Laboratory measurements

226 Baseline Lp(a) mass was measured from stored blood samples in the BiomarCaRE central
227 laboratory in either Mainz (until 2011) or Hamburg, (since 2011) Germany, using a fully au-
228 tomated, particle-enhanced turbidimetric immunoassay (Biokit Quantia Lp(a)-Test; Abbott
229 Diagnostics, USA) (15). LDL-C levels were calculated using the Friedewald formula without
230 any additional hypertriglyceridemia-related adjustments. ApoB was measured using Immuno-
231 turbidimetric assay (Abbott, Architect c8000). The remaining lipid parameters (total choles-
232 terol, HDL-C or triglycerides) were measured locally at each participating center by routine
233 methods or in the BiomarCaRE central laboratory.

234 The cohort-specific intra- and interassay coefficients of variation for Lp(a), LDL-C
235 and apoB are provided in the supplemental Table 1.

236 Statistical analysis

237 Baseline characteristics of the study population are reported in a descriptive way and shown
238 as frequencies (percentage) for binary variables and as medians with their lower and upper
239 quartile (interquartile range, IQR) for continuous variables.

240 Median follow-up (FU) times were estimated by the Kaplan-Meier potential follow-up
241 estimator (16). All lipoproteins were categorized into fifths (F) using cohort-specific quintiles.
242 The mean cut-point values for LDL-C were 2.72 mmol/L, 3.26 mmol/L, 3.78 mmol/L and
243 4.42 mmol/L, whilst for apoB they were 0.80 g/L, 0.95 g/L, 1.09 g/L and 1.27 g/L.

244 To assess the association between lipoproteins and future CHD events, Fine and Gray
245 models accounting for competing risk of death from a non-CHD cause, stratified by sex and
246 study cohort were calculated using individual level data from the available cohorts. Lp(a) and
247 apoB were cubic-root transformed prior to the analysis. The data are presented as sub-
248 distribution Hazard Ratios (sHRs) with their 95% confidence interval (95% CI).

249 As a first step, we performed an analysis for the association of LDL-C, apoB and
250 Lp(a) (separately, as sole biomarkers) with future CHD events at different levels of adjust-
251 ment. Model 1 included age and fasting status; model 2 was additionally adjusted for systolic
252 blood pressure, antihypertensive drugs, diabetes mellitus, body mass index (BMI), daily
253 smoking, family history of CHD, average daily alcohol consumption, highest level of educa-
254 tion and lipid-lowering medication. Further, we repeated the analysis for LDL-C after correct-
255 ing it for its Lp(a)-C content. Two different corrections were applied, which estimated Lp(a)-
256 C content as 30% ($LDL_{Lp(a)corr\ 30}$) (4) or 17.3% ($LDL_{Lp(a)corr\ 17.3}$) (6) of total Lp(a) mass. Thus,
257 $LDL_{Lp(a)corr\ 30}$ was calculated as $LDL-C - (Lp(a)*0.30)$ and $LDL_{Lp(a)corr\ 17.3}$ was calculated as
258 $LDL-C - (Lp(a)*0.173)$. The Spearman correlation coefficients were calculated for LDL-C
259 and the two corrected versions

260 Finally, we looked into the relationship between LDL-C or apoB and incident CHD
261 according to concomitant Lp(a) level and therefore stratified the entire population into low-to-
262 moderate and high Lp(a) values using cohort-specific Lp(a) cut-offs (<90th pctl.
263 (n=61,861 versus ≥90th pctl. (n=6,887)). The mean 90th pctl. is about 43.5 mg/dL. Same levels
264 of adjustment were used for the regression models. Importantly, the same categorization of
265 LDL-C or apoB into fifths using cohort-specific quintiles were applied. Terms for the interac-
266 tion of continuous Lp(a) as well as fifths with LDL-C and apoB were added to the models.

267 Due to the exploratory nature of this analysis, a significance threshold was not defined
268 for p-values. R version 4.2.1 software (R Foundation for Statistical Computing, Vienna, Aus-
269 tria) was used to perform all statistical analyses.

270 **Results:**

271 For the present analysis, data from 68,748 subjects from eight prospective population-based
272 cohorts across Europe participating in the BiomarCaRE project, were used. All included par-
273 ticipants were free of CHD at the time of enrollment.

274 Table 1 describes the baseline demographic, clinical and biochemical characteristics of
275 the entire population, as well as after stratification of the study sample according to Lp(a)
276 mass with a 90th percentile used as a cut-off. The median Lp(a) was found to be 9.3 (IQR 4.2-
277 20.4) mg/dL in the total study group; 8.0 (IQR 3.8-15.2) mg/dL in subject with Lp(a) values
278 below 90th percentile and 60.5 (IQR 52.1-70.3) mg/dl in those with a Lp(a) mass $\geq 90^{\text{th}}$ pctl. of
279 Lp(a) distribution. The prevalence of most cardiovascular risk factors was comparable be-
280 tween subject with low versus high Lp(a) levels, the only exception being a family history of
281 CHD, which was more frequent in individuals with high Lp(a) mass (19.7% vs 16.9%). Medi-
282 an values for LDL-C and non HDL-C were slightly higher in subjects with high Lp(a) mass
283 (3.4 (IQR 2.8-4.2) vs 3.8 (IQR 3.2-4.5) mmol/L for LDL-C and 4.2 (IQR 3.4-5.0) vs 4.5 (IQR
284 3.8-5.3) mmol/L for non-HDL-C, respectively). In contrast, the concentration of apoB was
285 almost identical in the two subgroups, being 1.0 (IQR 0.8-1.2) in those with low vs 1.1 (IQR
286 0.9-1.2) g/L in those with high Lp(a) level. For the baseline characteristics of each individual
287 cohort please see Supplementary material online (Supplemental Table 2).

288 During a median FU of 9.72 years (95% CI 9.64-9.79 years) 3,536 of subjects who
289 were free of CHD at baseline developed an event, defined as fatal or non-fatal MI, coronary
290 death, unstable angina pectoris, coronary revascularization, or unclassifiable death.

291 As a first step, we looked into the association between LDL-C, apoB and Lp(a) and
292 incident CHD in the entire population (Table 2). Comparing the fifths of lipoprotein distribu-
293 tions , we found that all studied lipoproteins were associated with future CHD events after

294 multivariable adjustment for cardiovascular risk factors with sHRs of 2.73 (95% CI 2.34-
295 3.20) for LDL-C, 3.33 (95% CI 2.79-3.97) for apoB and 1.49 (95% CI 1.31-1.69) for Lp(a)
296 (exemplarily for top versus bottom fifth (used as reference), fully adjusted models, all
297 $p < 0.001$). Interestingly, for LDL-C and apoB, the associations with outcome were already
298 evident from the second fifths onwards, whereas there was no evidence of association be-
299 tween Lp(a) and incident CHD within F2 and F3 of the Lp(a) distribution (Table 2).

300 Next, we investigated whether correction of LDL-C for its Lp(a)-C content might af-
301 fect its relationship with outcome. In order to account for the variability of Lp(a)-C in relation
302 to Lp(a) mass we used two different estimations, one calculating Lp(a)-C as 30%, the other as
303 17.3 % of total Lp(a) mass. Interestingly, the Spearman correlation between the original LDL-
304 C and the two corrected LDL-C were found to be 0.99 for the correlation between LDL-C and
305 $LDL_{Lp(a)corr\ 30}$ and 1.0 for the correlation between LDL-C and $LDL_{Lp(a)corr\ 17.3}$. The results of
306 Fine and Gray competing risk-adjusted models revealed that these corrections did not affect
307 the relationship between corrected LDL-C and incident CHD meaningfully, demonstrating
308 very comparable sHRs to those seen for uncorrected LDL-C (sHR of 2.51 (95% CI 2.15-2.93)
309 for $LDL_{Lp(a)corr\ 30}$ and 2.64 (95% CI 2.26- 3.10) for $LDL_{Lp(a)corr\ 17.3}$ versus 2.73 (95% CI 2.34-
310 3.20) for uncorrected LDL-C (top versus bottom fifth, fully-adjusted models; both $p < 0.001$)
311 (Table 3 and Table 2).

312 Considering that the estimation of a “true” LDL-C by the above mentioned equations
313 has limited applicability in routine practice, we sought for a more practical solution to assess
314 the association between LDL-C and future events depending on its Lp(a)-related cholesterol
315 content and stratified our study population according to Lp(a) into low-moderate ($< 90^{th}$ pctl.)
316 and high values ($\geq 90^{th}$ pctl.) (Table 4). In subjects with low-moderate Lp(a) mass, the corre-
317 sponding sHR for the association between LDL-C and incident CHD were almost identical to

318 those obtained within the total study sample (i.e. without categorization for Lp(a) mass) (sHR
319 2.60 (95% CI 2.21-3.07); top versus bottom fifth of LDL-C distribution; fully-adjusted mod-
320 el). In contrast, the risk estimates were higher in those with high Lp(a) with a sHR of 4.38
321 (95% CI 2.08-9.22) (top versus bottom fifth of LDL-C distribution; fully-adjusted model)
322 ($p_{\text{interaction}}$ 0.39).

323 To investigate whether apoB would demonstrate a similar pattern of association with
324 outcome like LDL-C taking into account Lp(a) values, we repeated the analysis using apoB as
325 the independent variable (Table 5).Corresponding sHRs for apoB obtained within the total
326 study sample (i.e. without categorization for Lp(a) mass) were very similar to the risk esti-
327 mates found in subjects with low-moderate Lp(a) (<90th pctl.), for whom the sHR was 3.34
328 (95% CI 2.78-4.01) for apoB (top versus bottom fifth, fully-adjusted model). Surprisingly, the
329 association between apoB and future CHD was weaker in subjects with Lp(a) mass $\geq 90^{\text{th}}$ pctl.
330 with a sHR of 2.43 (95% CI 1.34-4.40) ($p_{\text{interaction}}$ 0.49). No association with incident CHD
331 was found for the second and third fifth of the apoB distribution (F2 and F3) in the group with
332 high Lp(a) values. In contrast, corresponding sHRs in those without stratification for Lp(a)
333 mass or in those with low-moderate Lp(a) values were markedly higher, varying from 1.5 to
334 1.9 for F2 and F3 respectively (Table 2 and Table 5).

335 **Discussion:**

336 To the best of our knowledge, the present analysis represents the largest study so far investi-
337 gating the impact of concomitant Lp(a) level on LDL-C or apoB-related risk of incident CHD
338 in the general population. Our study demonstrates that correction of LDL-C for its Lp(a)-C
339 content (assumed either as 30% or as 17.3% of total Lp(a) mass) did not substantially change
340 LDL-C-associated CHD-risk estimation at the population level. Echoing this, similar LDL-C-
341 associated risk for future CHD was observed in subjects with low/moderately increased Lp(a)
342 mass (i.e. being under 90th pctl. of Lp(a) distribution), which represents the vast majority of
343 studied individuals. The situation, however, became more complex when Lp(a) values were
344 high (e.g. in this instance exceeding the 90th cohort-specific pctl.) since LDL-C-related CHD
345 risk estimates were higher in subjects with higher Lp(a) mass. In the case of apoB, opposite
346 patterns of association with incident CHD were observed with slightly lower sHR for apoB-
347 associated risk found in those with high Lp(a) mass compared to subjects with lower Lp(a)
348 mass.

349 Correction of LDL-C for the cholesterol content of Lp(a)

350 Despite several attempts to understand the clinical relevance of Lp(a)-C content of routinely
351 measured LDL-C, this critical issue has not been fully resolved (17). Although the current
352 ESC/EAS Lp(a) statement indicates that correcting of LDL-C for its Lp(a)-C content is not
353 precise enough and should not be applied routinely (7), it is still broadly used in the research
354 community. The findings of the present analysis showed that the two most commonly applied
355 corrections of LDL-C for its Lp(a)-C content to date (calculated as 30% or 17.3% of total
356 Lp(a) mass) do not provide any additional meaningful information on top of LDL-C-related
357 CHD risk prediction in subjects from the general population, where Lp(a) on an absolute scale
358 is little to only moderately increased in the vast majority of subjects. Interestingly, our data

359 are contrary to the previously published individual-patient-data meta-analysis from 5 land-
360 mark statin trials including 18,043 patients, which demonstrated that corrected LDL-C did not
361 predict future CVD events in contrast to “uncorrected” LDL-C (HR for incident cardiovascu-
362 lar disease 1.07 (95% CI 0.93–1.22), $p=0.36$) versus HR 1.17 (95% CI 1.05–1.31), $p=0.005$,
363 respectively; top versus bottom quartile) (18). In contrast, within the present analysis almost
364 identical risk estimates between uncorrected and corrected LDL-C for incident CHD were
365 found. Interestingly, uncorrected LDL-C was highly correlated with $LDL-C_{Lp(a)corr}$ 30 with a
366 pooled correlation coefficient of 0.96 (95% CI, 0.94–0.97) (18), which was very similar to the
367 correlations observed within the present analysis. It should be noted here that our analysis
368 cannot be directly compared to the study by Willeit et al. due to differences in the study de-
369 sign, population studied and concomitant medications used. Furthermore, although correction
370 of LDL-C for its Lp(a)-C content did not markedly change the LDL-C-associated risk in the
371 present analysis, lack of such differences at a population level does not establish clinical
372 equivalence at an individual level. Whenever subjects with low/moderate Lp(a) mass are
373 pooled with those with high Lp(a) mass, corrected LDL-C risk in the whole population might
374 remain largely unchanged due to a simple predominance of subjects with only low/moderately
375 increased Lp(a) values. Our results thus argue for a clear separation of subjects with high ver-
376 sus low/moderate Lp(a) values and that assessment of the conventional lipid parameters, such
377 as LDL-C or apoB without taking into account accompanying Lp(a) values would provide
378 only incomplete information on CHD risk estimation. While in subjects with low to moderate-
379 ly increased Lp(a) values, the contribution of Lp(a)-C to “overall” LDL-C might not be sub-
380 stantial with only negligible impact on the associated risk estimation, in those with high Lp(a)
381 levels it might impact risk prediction.

382 Lp(a) and LDL-C or apoB-related risk for incident CHD

383 The present analysis indirectly touches upon the debate related to the superior accuracy of
384 apoB compared to LDL-C as a marker of cardiovascular risk (19). Multiple lines of epidemio-
385 logical evidence including Mendelian randomization studies (19-21) suggest apoB as a domi-
386 nant marker of ASCVD risk when compared to LDL-C. Our data in the general population, as
387 well as in a subgroup of subjects with lower Lp(a) values (< 90th pctl. of its distribution) also
388 revealed slightly higher sHRs for apoB and incident CHD than for LDL-C, with 3.33 (95% CI
389 2.79-3.97) vs 2.73 (95% CI 2.34-3.20) respectively (exemplarily for the entire population,
390 both for top versus bottom fifths, fully adjusted models). More surprising were our findings
391 on the association between apoB and incident CHD in subjects with high Lp(a) levels, in
392 whom the risk estimates for the top vs bottom fifths of apoB distribution were lower than
393 those in subjects with lower Lp(a) values or from the entire population (sHR 2.43 (95 % CI
394 1.34-4.40) versus 3.34 (95% CI 2.78-4.01) or 3.33 (95% CI 2.79-3.97), respectively). Fur-
395 thermore, at high Lp(a) values no association with CHD risk was observed for the second and
396 third fifths of the apoB distribution, in contrast to the associations observed in those with low-
397 er Lp(a) levels or the entire population. Interestingly, apoB concentrations in both Lp(a)
398 groups were almost identical (median 1.0 (0.8-1.2) versus 1.1 (0.9-1.2) g/L)). It has to be not-
399 ed here, that these findings should be interpreted with caution because of a small number of
400 events and some overlap between point estimates and their 95% CIs. Nonetheless, our results
401 are rather hypothesis generating and encouraging further research. A mechanistic explanation
402 for the diminished or even lost association between apoB and future CHD events is unclear
403 and needs to be investigated in more details. It is well established that native/unmodified
404 apoB-100 does not seem to be atherogenic, whereas the risk of ASCVD seems to be mostly
405 driven by oxidized phospholipids (oxPLs) on apoB particles (22). On the other hand, it is well
406 known that Lp(a) carries the largest fraction of oxPLs among apoB-containing lipoproteins in
407 the circulation, where they are bound covalently to the KIV10 of its apo(a) fragment (23).

408 Interestingly, experimental studies have demonstrated differences in oxPL content among
409 individuals with different plasma Lp(a) levels, where oxPL contents seem to be low in those
410 with low Lp(a) and increase proportionally with increasing plasma Lp(a) levels (23-24).
411 Moreover, in plasma from individuals with elevated plasma Lp(a) levels nearly all oxPLs
412 were found in density fractions containing apo(a) (24). Thus, one might suggest that in set-
413 tings of chronically elevated Lp(a) levels, oxPL might be preferentially transferred from non-
414 Lp(a) apoB-100 particles to Lp(a). Generally, such kind of transfer/shift is possible, as shown
415 by Tsimikas et al. (25) in the post percutaneous coronary intervention (PCI) setting, where
416 only 50% of the oxPLs were associated with Lp(a) directly after PCI and almost all oxPL
417 were localized on Lp(a) 6 hours later. However, whether such shift or transfer of oxPLs might
418 be responsible for the diminished predictivity of the “total” apoB is highly speculative. No
419 similar pattern was observed for LDL-C, where we found that the association between LDL-C
420 and outcome was higher at high Lp(a) values, than in subjects with low/moderate Lp(a) mass
421 (HR 4.38 (95% CI 2.08-9.22) versus 2.60 (95% CI 2.21-3.07), respectively (top versus bottom
422 fifth of LDL-C distribution; fully-adjusted models)). One potential explanation for such dis-
423 cordant findings might be related to the methodological issues of LDL-C and apoB quantifi-
424 cation (26). Despite the biological linkage between these measures, it is well established that
425 LDL-C does not represent an accurate equivalent of apoB due to the highly variable choles-
426 terol content within each particle, with either cholesterol-depleted or cholesterol-enriched
427 LDL (27). In contrast, apoB concentration reflects the total number of apoB particles in plas-
428 ma. Taking into account these considerations, it is not surprising that the association between
429 LDL-C and incident CHD was found to be stronger at high Lp(a) values than in subjects with
430 low to moderate Lp(a) increase, reflecting a significant contribution of Lp(a) cholesterol mass
431 to a given mass of overall LDL-C due to their overlapping densities (2). Interestingly, the
432 question which rather should be raised here is whether a molar measurement of Lp(a), which

433 yields Lp(a) particle number rather than Lp(a) mass (28), would also enhance LDL-C predic-
434 tive potential for future CHD or are we dealing with the effects that are attributable to the
435 Lp(a) mass assay only. Although very intriguing, our results need to be replicated within larg-
436 er populations, since we cannot exclude that lack of statistical power due to the low number
437 of participants within the reference group among subjects with high Lp(a) values might have
438 led to an overestimation of LDL-C associated CHD risk. On the other hand, the discordant
439 results for the association with outcome revealed for LDL-C and apoB would rather speak
440 against this assumption.

441 Limitation and Strengths of the Study

442 There are limitations of our study, which merit consideration. Since we used a mass-based
443 assay for Lp(a) measurement, we could not apply the molar-based correction recently pro-
444 posed by Rosenson and Marcovina (29), and could not calculate a proportion of total apoB
445 attributable to Lp(a) (8). The studied biomarkers were measured only once and therefore the
446 results cannot account for a regression dilution bias. The present data cannot be extrapolated
447 to other ethnic populations or age groups, since only middle aged Caucasians were included
448 in this analysis. Finally, CHD assessment at baseline mainly relied on medical reviews or to a
449 much lesser extent was self-reported, which may have led to some misclassification, but we
450 expect this to be non-differential across Lp(a) levels.

451 The current study has also several strengths. It represents the largest population-based
452 analysis so far investigating the role of Lp(a) as a possible modifier of LDL-C or apoB-related
453 CHD risk. Centralized measurements of biomarkers by the same assays minimized analytical
454 imprecision in measurements between individual BiomarCaRE cohorts. Moreover, possible
455 Lp(a)-increasing effects of statins are negligible within the present analysis, since only 2.8%
456 of the entire study population were reported to be on lipid-lowering drugs at baseline and

457 most BiomarCaRE cohorts were recruited in the late 1980s-early 1990s, when statins were not
458 broadly used. Finally, largely standardized baseline measurements and careful harmonization
459 of the data from eight European general population-based studies lead to comparable and reli-
460 able data on risk factors and endpoint validation.

461 **Conclusions**

462 Within the present analysis, correction of LDL-C for its Lp(a)-C content by established esti-
463 mations did not provide any meaningful information on LDL-C related CHD-risk in the gen-
464 eral population. In contrast, a simple categorization of Lp(a) mass into high ($\geq 90^{\text{th}}$ pctl.) vs
465 low/moderate ($< 90^{\text{th}}$ pctl.) values demonstrated that concomitant Lp(a) might impact the lipo-
466 protein-related risk for future CHD events mostly at higher Lp(a) levels. Thus, an assessment
467 of the conventional lipid profile without taking into account accompanying Lp(a) values
468 would provide incomplete information on CHD risk estimation, especially in subjects with
469 high Lp(a) values.

470

471 **CLINICAL PERSPECTIVES**

472 **COMPETENCY IN PATIENT CARE:** Since concomitant increased Lp(a) mass might im-
473 pact the association between conventional lipoproteins and future CHD, Lp(a) should be tak-
474 en into account for more comprehensive assessment of LDL-C- or apo(B)-related risk for fu-
475 ture coronary events. Ideally, Lp(a) should be included in the conventional lipid panel at first
476 presentation of the patient.

477 **TRANSLATIONAL OUTLOOK:** More accurate estimation the cholesterol content of Lp(a)
478 is urgently needed to understand the tight interplay between Lp(a), LDL-C and apoB.

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566 **Figure legends**

567 **Figure 1: Flow chart of the study.**

568 MONICA = Monitoring of Trends and Determinants in Cardiovascular Diseases; KORA =
569 Cooperative Health Research in the Region of Augsburg; MATISS = Malattie ATeroscle-
570 rotiche Istituto Superiore di Sanità; SHHEC = Scottish Heart Health Extended Cohort; CHD =
571 coronary heart disease; Lp(a) = lipoprotein (a); pctl. = percentiles.

572 **Central Illustration**

573 CHD = coronary heart disease; Lp(a) = lipoprotein (a); pctl. = percentiles; FU = follow up;
574 yrs. = years; sHR = sub-distribution Hazard Ratio; CI = confidence interval; LDL-C = low
575 density lipoprotein cholesterol; $LDL_{Lp(a)corr}$ = LDL corrected for Lp(a) cholesterol; apoB =
576 apolipoprotein B.

	All	Lp(a) <90 th pctl.	Lp(a) ≥90 th pctl.
N	68,748	61,861	6,887
Examination age (years)	49.7 (41.0- 58.7)	49.6 (40.9- 58.7)	50.5 (41.7- 59.2)
Male, n (%)	33,270 (48.4)	30,218 (48.8)	3,052 (44.3)
Systolic BP (mmHg)	131.0 (119.0-145.5)	131.0 (119.0-145.5)	131.0 (119.0-146.0)
BMI (kg/m ²)	26.3 (23.6-29.4)	26.3 (23.6-29.4)	26.2 (23.7-29.1)
Hypertension, n (%)	27,726 (40.4)	24,873 (40.2)	2,853 (41.5)
Daily smoker, n (%)	17,628 (25.7)	15,938 (25.8)	1,690 (24.6)
Diabetes, n (%)	2,628 (3.8)	2,400 (3.9)	228 (3.3)
Family history of CHD, n (%)	9,071 (17.2)	8,030 (16.9)	1,041 (19.7)
Daily alcohol (g)	5.0 (0-22.0)	5.0 (0-22.0)	5.0 (0-20.0)
Lipid-lowering drugs, n (%)	1,715 (2.8)	1,475 (2.7)	240 (3.9)
Antihypertensive drugs, n (%)	9,561 (14.0)	8,553 (13.9)	1,008 (14.7)
Aspirin intake, n (%)	1,257 (2.8)	1,129 (2.8)	128 (2.8)
Non-HDL-C (mmol/L)	4.2 (3.5-5.0)	4.2 (3.4-5.0)	4.5 (3.8-5.3)
LDL-C (mmol/L)	3.5 (2.9-4.2)	3.4 (2.8-4.2)	3.8 (3.2-4.5)
LDL <i>Lp(a) corr 17.3</i> (mmol/L)	3.4 (2.8-4.1)	3.4 (2.8-4.1)	3.5 (2.9-4.2)
LDL <i>Lp(a) corr 30</i> (mmol/L)	3.4 (2.7-4.0)	3.4 (2.7-4.1)	3.3 (2.7-4.0)
Lp(a) (mg/dL)	9.3 (4.2-20.4)	8.0 (3.8-15.2)	60.5 (52.1-70.3)
Apolipoprotein B (g/L)	1.0 (0.8-1.2)	1.0 (0.8-1.2)	1.1 (0.9-1.2)

Table 1. Baseline demographic and clinical characteristics of the study participants.

Data are presented as median with their interquartile range for continuous variables. Categorical variables are reported as frequency and percentage.

Lp(a) = lipoprotein (a); pctl = percentile; BP = blood pressure; BMI = body mass index; CHD = coronary heart disease; HDL = high density lipoprotein; C = cholesterol; LDL = low density lipoprotein. Cohort-specific Lp(a) cut-off (90th pctl.) = 43.53 mg/dL.

Table 2: Association between circulating lipoproteins and risk of incident CHD

Lipoprotein (a)					LDL-Cholesterol				Apolipoprotein B			
sHR (95% CI)	p-value	N _{overall}	N _{events}		sHR (95% CI)	p-value	N _{overall}	N _{events}	sHR (95% CI)	p-value	N _{overall}	N _{events}
Model 1												
F 1	REF.	-	12,528	554	REF.	-	12,022	276	REF.	-	12,577	203
F 2	1.00 (0.89-1.13)	0.97	12,130	523	1.24 (1.07-1.45)	0.0051	11,969	391	1.58 (1.34-1.87)	<0.001	12,620	393
F 3	1.03 (0.92-1.16)	0.58	12,136	545	1.53 (1.33-1.77)	<0.001	12,006	523	2.04 (1.74-2.40)	<0.001	11,783	552
F 4	1.24 (1.11-1.38)	<0.001	12,168	680	1.84 (1.60-2.11)	<0.001	11,876	664	2.53 (2.16-2.95)	<0.001	12,282	758
F 5	1.42 (1.28-1.59)	<0.001	12,231	725	2.72 (2.38-3.11)	<0.001	11,823	988	3.76 (3.23-4.37)	<0.001	11,841	1,119
Model 2												
F 1	REF.	-	7,787	413	REF.	-	7,338	201	REF.	-	7,763	149
F 2	0.98 (0.85-1.13)	0.79	7,590	379	1.30 (1.08-1.55)	0.0045	7,353	285	1.56 (1.28-1.89)	<0.001	7,805	290
F 3	1.10 (0.96-1.26)	0.16	7,525	419	1.58 (1.33-1.87)	<0.001	7,430	382	1.94 (1.61-2.34)	<0.001	7,205	407
F 4	1.26 (1.10-1.43)	<0.001	7,467	499	1.92 (1.63-2.26)	<0.001	7,418	495	2.31 (1.93-2.77)	<0.001	7,706	564
F 5	1.49 (1.31-1.69)	<0.001	7,551	540	2.73 (2.34-3.20)	<0.001	7,230	730	3.33 (2.79-3.97)	<0.001	7,388	838

Fine and Gray competing risk-adjusted models stratified by study cohort were calculated and the data are presented as sub-distribution Hazard ratios (sHRs) with their 95% confidence interval (95% CI). Biomarkers were transformed for the analysis (Lp(a) and apolipoprotein B: cubic-root transformed). CHD = coronary heart disease; LDL = low density lipoprotein; REF = reference; F = fifth.

The mean cut-point values for Lp(a) were 3.46 md/dL, 6.66 mg /dL, 11.27 mg/dL and 24.85 mg/dL; for LDL-C were 2.72 mmol/L, 3.26 mmol/L, 3.78 mmol/L and 4.42 mmol/L; for apoB were 0.80 g/L, 0.95 g/L, 1.09 g/L and 1.27 g/L.

All models were stratified by sex and study cohort.

Model 1: Adjusted for examination age and fasting status

Model 2: Additionally adjusted for systolic blood pressure, antihypertensive medication, diabetes mellitus, BMI, daily smoker, family history of CHD, average daily alcohol consumption, highest level of education, lipid-lowering medication

Table 3: Association between LDL-Lp(a) corrected and risk of incident CHD

	LDL_{Lp(a)corr 30}				LDL_{Lp(a) corr 17.3}			
	sHR (95% CI)	p-value	N _{overall}	N _{events}	sHR (95% CI)	p-value	N _{overall}	N _{events}
Model 1								
F 1	REF.		11,945	287	REF.		11,949	277
F 2	1.15 (0.99-1.34)	0.070	11,969	383	1.22 (1.04- 1.42)	0.012	11,963	391
F 3	1.47 (1.27-1.69)	<0.001	11,943	528	1.50 (1.30- 1.73)	<0.001	11,943	522
F 4	1.73 (1.51-1.99)	<0.001	11,916	664	1.84 (1.60- 2.11)	<0.001	11,919	676
F 5	2.52 (2.21-2.87)	<0.001	11,922	980	2.61 (2.28- 2.98)	<0.001	11,921	976
Model 2								
F 1	REF.		7,276	208	REF.		7,270	198
F 2	1.18 (0.99-1.41)	0.068	7,364	276	1.27 (1.06- 1.52)	0.0094	7,361	284
F 3	1.51 (1.28-1.79)	<0.001	7,396	387	1.56 (1.32- 1.85)	<0.001	7,412	385
F 4	1.82 (1.55-2.14)	<0.001	7,428	503	1.93 (1.63- 2.27)	<0.001	7,419	504
F 5	2.51 (2.15-2.93)	<0.001	7,304	719	2.64 (2.26- 3.10)	<0.001	7,306	722

Fine and Gray competing risk-adjusted models stratified by study cohort were calculated and the data are presented as sub-distribution Hazard ratios (sHRs) with their 95% confidence interval (95% CI). Biomarkers were transformed for the analysis (Lp(a): cubic-root transformed). CHD = coronary heart disease; LDL = low density lipoprotein; REF = reference; F = fifth. LDL_{Lp(a)corr 30} calculated as LDL-(Lp(a)*0.30); LDL_{Lp(a)corr 17,3} calculated as LDL-(Lp(a)*0.173).

All models were stratified by sex and study cohort.

Model 1: Adjusted for examination age and fasting status

Model 2: Additionally adjusted for systolic blood pressure, antihypertensive medication, diabetes mellitus, BMI, daily smoker, family history of CHD, average daily alcohol consumption, highest level of education, lipid-lowering medication

Table 4. Association between LDL-C and risk of incident CHD, according to Lp(a) mass

	Lp(a) < 90 th percentile				Lp(a) ≥ 90 th percentile			
	sHR (95% CI)	p-value	N _{overall}	N _{events}	sHR (95% CI)	p-value	N _{overall}	N _{events}
Model 1								
LDL F 1	REF.		11,392	259	REF.		630	17
F 2	1.23 (1.05-1.45)	0.0095	10,997	353	1.18 (0.68-2.07)	0.56	972	38
F 3	1.56 (1.34-1.81)	<0.001	10,770	467	1.12 (0.66-1.90)	0.67	1,236	56
F 4	1.82 (1.57-2.11)	<0.001	10,399	571	1.60 (0.97-2.64)	0.065	1,477	93
F 5	2.69 (2.34-3.09)	<0.001	10,091	828	2.31 (1.43-3.75)	<0.001	1,732	160
							P interaction 0.62	
Model 2								
LDL F 1	REF.		6,959	194	REF.		379	7
F 2	1.23 (1.03-1.48)	0.025	6,756	256	2.53 (1.12-5.68)	0.025	597	29
F 3	1.55 (1.30-1.85)	<0.001	6,670	341	2.19 (1.00-4.81)	0.051	760	41
F 4	1.84 (1.55-2.18)	<0.001	6,492	428	3.14 (1.47-6.73)	0.0033	926	67
F 5	2.60 (2.21-3.07)	<0.001	6,152	605	4.38 (2.08-9.22)	<0.001	1,078	125
							P interaction 0.39	

Fine and Gray competing risk-adjusted models stratified by study cohort were calculated and the data are presented as sub-distribution Hazard ratios (sHRs) with their 95% confidence interval (95% CI). Biomarkers were transformed for the analysis (Lp(a): cubic-root transformed). LDL = low density lipoprotein; CHD = coronary heart disease; Lp(a) = lipoprotein (a); REF = reference; F = fifth. Cohort-specific Lp(a) cut-off (90th pctl.) = 43.53 mg/dL. The mean cut-point values for for LDL-C were 2.72 mmol/L, 3.26 mmol/L, 3.78 mmol/L and 4.42 mmol/L

All models were stratified by sex and study cohort.

Model 1: Adjusted for examination age and fasting status

Model 2: Additionally adjusted for systolic blood pressure, antihypertensive medication, diabetes mellitus, BMI, daily smoker, family history of CHD, average daily alcohol consumption, highest level of education, lipid-lowering medication

Table 5. Association between apolipoprotein B and risk of incident CHD, according to Lp(a) mass

		Lp(a) < 90 th percentile				Lp(a) ≥ 90 th percentile			
		sHR (95% CI)	p-value	N _{overall}	N _{events}	sHR (95% CI)	p-value	N _{overall}	N _{events}
Model 1									
apoB	F 1	REF.	-	11,733	180	REF.	-	844	23
	F 2	1.66 (1.39-1.99)	<0.001	11,446	355	0.84 (0.51-1.39)	0.50	1,174	38
	F 3	2.19 (1.85-2.60)	<0.001	10,523	496	0.90 (0.56-1.44)	0.66	1,260	56
	F 4	2.64 (2.24-3.11)	<0.001	10,847	657	1.35 (0.87-2.10)	0.18	1,435	101
	F 5	3.93 (3.35-4.61)	<0.001	10,409	959	1.97 (1.29-3.00)	0.0018	1,432	160
								P interaction <0.01	
Modell 2									
apoB	F 1	REF.		7,251	138	REF.		512	11
	F 2	1.57 (1.28-1.93)	<0.001	7,081	264	1.22 (0.62-2.42)	0.57	724	26
	F 3	1.99 (1.64-2.42)	<0.001	6,437	366	1.26 (0.66-2.41)	0.49	768	41
	F 4	2.28 (1.88-2.75)	<0.001	6,802	484	1.96 (1.06-3.60)	0.031	904	80
	F 5	3.34 (2.78-4.01)	<0.001	6,472	714	2.43 (1.34-4.40)	0.0035	916	124
								P interaction 0.49	

Fine and Gray competing risk-adjusted models stratified by study cohort were calculated and the data are presented as sub-distribution Hazard ratios (sHRs) with their 95% confidence interval (95% CI). Biomarkers were transformed for the analysis (Lp(a): cubic-root transformed). CHD = coronary heart disease; Lp(a) = lipoprotein (a), apoB = apolipoprotein B; REF = reference; F = fifth. Cohort-specific Lp(a) cut-off (90th pctl.) = 43.53 mg/dL. The mean cut-point values for apolipoprotein B were 0.80 g/L, 0.95 g/L, 1.09 g/L and 1.27 g/L.

All models were stratified by sex and study cohort.

Model 1: Adjusted for examination age and fasting status

Model 2: Additionally adjusted for systolic blood pressure, antihypertensive medication, diabetes mellitus, BMI, daily smoker, family history of CHD, average daily alcohol consumption, highest level of education, lipid-lowering medication