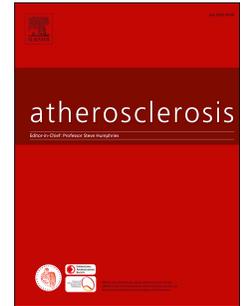


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Cholesterol efflux capacity of large, small and total HDL particles is unaltered by atorvastatin in patients with type 2 diabetes

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2 **Cholesterol efflux capacity of large, small and total HDL particles is unaltered by atorvastatin in**
3 **patients with type 2 diabetes**

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Abstract

33 *Background and aims:* Research on the biologic activities of HDL, such as cholesterol
34 efflux capacity and HDL composition, have allowed the understanding of the effect of
35 interventions directed to improve cardiovascular risk. Previously, statin therapy has shown
36 conflicting results in its effects on cholesterol efflux capacity of HDL, the underlying
37 mechanisms are unclear but studies with positive effect are associated with an increase of
38 HDL-cholesterol levels. We investigated if 10 weeks with atorvastatin therapy changes
39 HDL efflux capacity and the chemical composition of its subpopulations.

40 *Methods:* In a before-after design basis, HDL-cholesterol levels, chemical composition and
41 cholesterol efflux capacity from HDL subpopulations isolated by isopycnic
42 ultracentrifugation were assessed in plasma samples from 60 patients with type 2 diabetes
43 mellito (T2DM) at baseline and after 10 weeks of treatment with 20 mg of atorvastatin.
44 Cholesterol efflux was measured from human THP-1 cells using large, light HDL2b and
45 small, dense 3c subpopulations as well as total HDL as acceptors. Changes of cholesterol
46 efflux and chemical composition of HDL after treatment were analyzed. Correlations
47 among variables potentially involved in cholesterol efflux were evaluated.

48 *Results:* A significant decrease of 4% in HDL-cholesterol levels was observed from 47 (42-
49 54) to 45 (39-56) mg/dL, $p= 0.02$. Cholesterol efflux from total-HDL and HDL2b and 3c
50 subfractions was maintained unchanged after treatment. The total mass of HDL remained
51 unaffected, except for HDL3a subpopulation accounted for by a significant increase in total
52 protein content. No significant correlations for variables previously known to be associated
53 with cholesterol efflux were found in our study.

54 *Conclusions:* Short therapy of 10 weeks with 20 mg of atorvastatin does not modify the
55 cholesterol efflux capacity neither the total mass of HDL2b, HDL3c and total HDL. The

56 discrepancy with previous reports may be due to selective effects among different classes
57 of statins or differences in the approaches to measure cellular cholesterol efflux.

58

59 Key words: Cholesterol efflux, HDL subpopulations, Type 2 diabetes, Atorvastatin, HDL
60 chemical composition

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61 Introduction

62 Anti-atherogenic activities and chemical composition of high-density lipoprotein (HDL) are
63 altered in patients with Type 2 diabetes (T2DM)[1][2][3][4][5][6][7][8][9][10]. Diabetic
64 dyslipidemia is characterized by high levels of triglyceride-rich lipoproteins, low
65 concentrations of HDL-cholesterol (HDL-C) and elevated proportions of small, dense LDL
66 particles (sdLDL). Although the pathophysiology of diabetic dyslipidemia is not clearly
67 elucidated yet, insulin-resistant state and elevated activities of key proteins regulating HDL
68 and LDL metabolism, and notably cholesteryl ester transfer protein (CETP) and hepatic
69 lipase, play an important role[11][12][13].

70 Among their anti-atherogenic effects, HDL particles are able to stimulate cholesterol efflux
71 from peripheral cells, the first step of the reverse cholesterol transport pathway. HDL
72 capacity to act as an acceptor for free cholesterol depends on the integrity of specific
73 cellular cholesterol export pathways which involve ATP-binding cassette transporter
74 (ABC) A1, ABCG1, scavenger receptor class B type I (SR-BI) and passive
75 diffusion[14][15][16][17][18][19]. Currently, several assays have been described to
76 determine the cholesterol efflux capacity, most of them using cellular lines from murine
77 and human macrophages, tritiated cholesterol and Apo-B depleted-serum, HDL or ApoA-1
78 as cholesterol acceptors[20][21]. The recognition of different HDL subpopulations has
79 allowed the identification of particles with diverse degrees of efficacy, potentially
80 accounted by a particular distribution of their chemical component and biologic activity
81 [22][23][24][25][26]. Interestingly, liquid chromatography-mass spectrometry (LC-MS)
82 analysis has documented profound alterations in the HDL lipidome in patients with T2DM,
83 suggesting that the structure of HDL is under continuing remodeling in this disease and that

84 HDL lipidomics can equally contribute to identify biomarkers of normal and deficient HDL
85 functionality[7][27][28]. In the last years, focus has been directed to therapies to improve
86 HDL-C and more recently HDL function. Therapies that increase HDL-C has failed to
87 show a diminution in cardiovascular risk and interventional studies directed to demonstrate
88 an improvement in HDL functions have produced conflicting
89 results[8][29][30][31][32][33][34] [35][36][37]. Recent studies revealed that cholesterol
90 efflux capacity was inversely associated with the incidence of cardiovascular events in a
91 population-base cohort study [38]. This report highlights the relevance of studying HDL
92 function in high-risk populations.

93 Current guidelines recommend early use of lipid-lowering drugs, particularly statins, in a
94 large proportion of patients with T2DM as a key approach to reduce cardiovascular
95 morbidity[39][40][41]. Despite that, cardiovascular mortality remains the leading cause of
96 death in T2DM, even in cases treated in accordance with current recommendations. Statins
97 are 3-hydroxi-3-methylglutaryl coenzyme A (HMG-CoA) reductase inhibitors. Early
98 blocking of this microsomal enzyme reduces hepatic cholesterol synthesis and promotes a
99 faster clearance of circulating cholesterol. In numerous large-scale studies statins have
100 proven to reduce the risk of acute cardiac events and death[42][43]. Previous reports, which
101 included T2DM patients as a target population, brought about contradictory results in terms
102 of the capacity of statins to improve anti-atherogenic activities of HDL, such a cholesterol
103 efflux, anti-inflammatory activity and anti-oxidative capacity[44][45][46][18] Moreover,
104 recent kinetic studies have reported that most of the effects of statins are molecule-specific
105 rather class-related and that statins have differential effects in HDL cholesterol
106 concentration and HDL functions[47][48]. Most of the studies of the effect of statins on

107 HDL function have been conducted in patients with preexisting cardiovascular disease
108 (CAD). Except for one small study on HDL cholesterol efflux after treatment with
109 simvastatin, T2DM patients have not been a matter of a properly controlled study.

110 We conducted a single group, before-after study to evaluate the effect on HDL function and
111 composition of a 10-week treatment with atorvastatin of T2DM patients. Five
112 subpopulations of HDL were isolated from plasma and their chemical composition and
113 cholesterol efflux capacity from human macrophagic THP-1 cells were evaluated

114 **Materials and methods**

115 *Patients and study design*

116 The study sample was composed by T2DM patients treated at the Lipid Clinic of the
117 Instituto Nacional de Ciencias Médicas y Nutrición (INCMNSZ) in Mexico. Clinical phase
118 was carried out at INCMNSZ, where the protocol was approved by the Ethics Institutional
119 Committee in accordance with the Helsinki Declaration. Further analyses were performed
120 at the INSERM Research Unit 1166 at the Hospital La Pitié - Salpêtrière in France.

121 Males or postmenopausal women aged 20 to 65 years, free of major diabetes-related
122 chronic complications and displaying HbA1c levels <8% were invited to participate.
123 Patients had to be out of statin therapy for at least 24 weeks to be eligible for the study.
124 Patients with previous diagnosis of elevated blood pressure had to be under a good control
125 using anti-hypertensive drugs (blood pressure below 130/80 mmHg). Smoking patients
126 were excluded. Patients with positive history of cardiovascular disease (CVD) or severe
127 hyperlipidemia (defined as plasma total cholesterol > 300 mg/dL or triglycerides > 400
128 mg/dL) as well as secondary causes of dyslipidemia or conditions altering lipid profile (i.e.

129 liver diseases, infection with human immunodeficiency virus, rheumatologic diseases, and
130 treatment with drugs that affect plasma lipid profile) were excluded. All patients signed an
131 informed consent.

132 The sample size for the study was calculated assuming one group (one-sample comparison
133 of mean) and 90% power to demonstrate a difference of 10% in the cholesterol efflux
134 capacity of HDL as a result of the treatment. According to these criteria, 33 patients were
135 required.

136 The study included four visits (at baseline and after 2, 4 and 10 weeks of the treatment).
137 Three weeks before the baseline visit, an isocaloric dietary plan was prescribed involving
138 50%, 20% and 30% of energy uptake from carbohydrates, proteins and fats, respectively.

139 Patients arrived to the clinical center after a 10-12 hour fasting period and were instructed
140 to take Atorvastatin 20 mg/day (one pill every night). Adherence to the therapy was
141 measured on the following visits using three-day food records and pill counts.
142 Anthropometric data were collected by a nutritionist. Body mass index (BMI) was
143 calculated as weight (kg) divided by height (meters) squared. Glucose-lowering therapies
144 and antihypertensive drugs remained constant during the study.

145 Laboratory measures

146 Blood samples were collected on every visit into EDTA-containing tubes as well as into
147 tubes without anticoagulant. EDTA plasma and serum were separated using low-speed
148 centrifugation, aliquoted and frozen at -70°C .

149 Lipid profile, levels of transaminases, insulin, apolipoproteins B and A-I, and core
150 laboratory clinical chemistry were determined on every visit. HbA1c concentration was
151 added to the evaluation at the first and the last visits. Plasma lipids and clinical chemistry
152 parameters were measured using commercially available kits (Synchron CX5-delta®,
153 Beckman Co®). Insulin was measured using an immunoenzymatic assay (Abbott®).
154 Concentrations of apolipoprotein B and A-I were evaluated using immunonephelometric
155 methods (Beckman®). Levels of glycated hemoglobin A1c were measured by HPLC
156 (BioRad®). Non-HDL cholesterol was calculated by subtracting HDL-C from total
157 cholesterol. LDL-C was calculated by the Friedewald formula in subjects with triglycerides
158 levels below 250 mg/dL.

159 Five subpopulations of HDL, specifically HDL2b, 2a, 3a, 3b and 3c, were isolated by
160 single-spin isopycnic density gradient ultracentrifugation using a potassium bromide
161 density gradient ultracentrifugal procedure for the isolation of the major lipoprotein classes
162 from human serum.[49]. After isolation, HDL subpopulations were extensively dialyzed for
163 48 hours and stored at 4°C for not longer than 5 days[50]. Total HDL from each subject
164 was prepared by mixing all five individual HDL subfractions at their equivalent plasma
165 concentrations.

166 Major chemical components of HDL (phospholipid, free cholesterol, total cholesterol,
167 triglyceride and total protein) were measured in all subpopulations. Phospholipid, free
168 cholesterol, total cholesterol and triglyceride were quantified using DiaSys® reagents.
169 Esterified cholesterol was calculated as a difference between total and free cholesterol
170 multiplied by 1.67 [49]. Total protein was determined using the BCA method (BioRad®).
171 Total mass of each HDL subfraction was calculated as a sum of phospholipid, free

172 cholesterol, esterified cholesterol, triglyceride and total protein concentrations. Total HDL
173 mass was calculated as the sum of total masses of five individual HDL subfractions.

174 The capacity to efflux cellular cholesterol was evaluated in HDL2b and 3c subpopulations
175 representative of large, light HDL2 and small, dense HDL3, respectively, as well as in total
176 HDL. Human monocyte-derived THP-1 cells were placed at 1×10^6 cells/well in 24-well
177 plates containing RPMI 1640, and PMA was added to differentiate the cells into
178 macrophages [27]. After 48 hour incubation under CO_2 , acetylated LDL was added to the
179 wells at $50 \mu\text{g/mL}$ followed by the addition of $12.5 \mu\text{L}$ of [^3H]-cholesterol ($1 \mu\text{L}$ of [^3H]-
180 cholesterol/mL RPMI). After 48 hours, cells were washed with PBS twice and serum-free
181 RPMI was added followed by HDL in PBS to a final concentration of $15 \mu\text{g}$
182 phospholipid/mL and final volume of $300 \mu\text{L}$. Cholesterol efflux capacity of HDL particles
183 was measured on the basis of their PL concentrations because PL was shown to represent
184 the key component determining cholesterol efflux capacity of HDL [51]. After 4 hours, the
185 supernatant was removed and radioactivity within the medium was determined by liquid
186 scintillation counting. The cells were lysed, $500 \mu\text{L}$ of the isopropanolol/hexane mixture
187 was added to each well to extract lipids, and cholesterol content was measured using
188 cholesterol reagent (DiaSys[®]); [^3H]-cholesterol was determined by scintillation counting
189 following addition of scintillation liquid. The percentage of cholesterol efflux was
190 calculated as $(\text{medium cpm}) / (\text{medium cpm} + \text{cell cpm}) \times 100\%$. Specific cholesterol
191 efflux was determined by subtracting non-specific cholesterol efflux occurring in the
192 absence of HDL. All measurements were done in triplicates.

193 *Statistical analysis*

194 Statistical analysis was performed using STATA 13 software package (StataCorp LP,
195 USA). Graphs were plotted using GraphPad Prism 5 software (2007 GraphPad Software
196 Inc)

197 The demographic, anthropometric and biochemical continuous variables are presented as
198 mean \pm SD or median and interquartile range (25-75) after testing distributions for
199 normality, whereas categorical variables are presented as percentages. To calculate
200 significance of the treatment effects, paired t-test or sign-rank Wilcoxon-test for continuous
201 variables with normal or non-normal distribution were applied, respectively. McNemar X^2
202 test for categorical variables was used.

203 Cholesterol efflux and chemical components of HDL subpopulations are presented as
204 median and IQ range (10-90). To calculate significance of the effects of atorvastatin on
205 these variables, Wilcoxon signed-rank test was employed.

206 Significance level of $p \leq 0.05$ was considered significant, except for multiple comparisons
207 when the level of $p \leq 0.05/5 = 0.01$ was used according to Bonferroni's adjustment, in order
208 to account for multiple comparisons across five HDL subpopulations.

209 According to the response to the treatment, two subgroups were built selecting patients who
210 achieved 35% of reduction in apoB levels or not.

211 We estimated the level of sdLDL using the triglyceride/HDL-cholesterol ratio. Patients
212 with the ratio above 4 were assumed to display elevated levels of sdLDL. We generated
213 two subgroups according this criterion[52].

214 Finally, we did Spearman correlation r^2 tests between clinical and biochemical variables
215 with cholesterol efflux to identified determinants of cholesterol efflux capacity of HDL.
216 Regression models were done as another approach to identified determinants of cholesterol
217 efflux.

218

219 **Results**

220 Seventy patients were recruited for the study. Ten patients were excluded for different
221 reasons and 60 patients completed the protocol (*Figure 1*).

222 The population predominately consisted of postmenopausal women (62%), with a mean age
223 for all patients of 58 ± 10 years (*Table 1*). Most (95%) of the patients were treated with
224 metformin as a single therapy. Thirty percent of patients used metformin in combination
225 with insulin or other glucose-lowering drugs, except thiazolidinediones. Thirty four percent
226 were previously diagnosed with elevated blood pressure; all of them were well controlled
227 (SBP, 127 ± 16 mmHg and DBP, 78 ± 10 mmHg). Former smoking was reported by 33% of
228 subjects and the time between smoking withdrawal and the start of the study was at least
229 one year. Patients maintained an isocaloric diet during the study (*Table 2*).

230 No significant effect of atorvastatin treatment on clinical parameters was observed (*Table*
231 *3*). The treatment resulted in significant changes in all lipid-related parameters assessed
232 (*Table 4*). The treatment reduced plasma concentrations of total cholesterol (by -32%),
233 LDL-C (by -50%), triglycerides (by -19%), and apoB (by -34%); all changes were
234 significant ($p<0.001$). LDL-C or apoB levels did not change after the treatment in 6% of
235 patients. There was a -4% ($p=0.02$) decrease in HDL-C levels. Finally, S-creatinine and

236 aspartate aminotransferase levels were decreased (by 5 and 4 %, $p=0.005$ and 0.02 ,
237 respectively) by the treatment (*Table 4*).

238 Total mass of HDL subpopulations remained unchanged, except in HDL3a, with an
239 increase from 65.7 to 69.2 mg/dL ($p=0.01$) after treatment. By contrast, atorvastatin
240 significantly modified the concentrations of several components of the HDL subpopulations.
241 Thus, free cholesterol in the small, dense HDL3c subpopulation was reduced from 0.29
242 (0.22-0.39) to 0.20 (0.15-0.28) mg/dL ($p=0.0001$). Triglycerides in the HDL3c
243 subpopulation decreased from 0.49 (0.38-0.80) to 0.34 (0.20-0.56) mg/dL ($p=0.0001$).
244 Total protein increased in the HDL3a subpopulation from 29.5 (27.2-35.1) to 32.3 (29.0-
245 38.5) mg/dL, $p=0.0008$. (Figure 2-A, B and E, respectively). These changes conducted to
246 some significant modifications in the proportional distributions of chemical components.
247 The free cholesterol proportion in the HDL3c subpopulation decreased from 1.97 (1.4-2.4)
248 to 1.5 (1.0-1.8)% ($p\leq 0.0001$). Triglycerides in the HDL3c subpopulation decreased from
249 3.2 (2.3-4.8) to 2.4 (1.5 - 3.2)% ($p<0.001$). Phospholipids increased in the HDL2b
250 subpopulation from 30.1 (27.3-33.3) to 31.3 (28.6-33.9)% ($p=0.01$) and decreased in
251 HDL2a from 31.14 (29.3-33.3) to 30.8 (28.5-32.3)%, HDL3a from 28.2 (26.6-29.9) to 28.1
252 (26.3-29.2)% and HDL3b from 23.6 (21.5-25.7) to 22.8 (20.8-24.9)%, ($p= 0.001$, 0.01 and
253 0.003 , respectively). Total protein increased in HDL3a from 46.7 (43.6-49.4) to 47.8 (44.8-
254 49.2)%, HDL3b from 53. (50.5-58.5) to 54.8 (52.5-58.9)% and HDL3c from 63.2 (58.8-
255 68.8) to 67.4 (63.1-70.2)%, ($p=0.001$, 0.01 and <0.001 , respectively) (Figure 2-G, H, J, K).

256 By contrast, no effect of atorvastatin on cholesterol efflux capacity of either large, light
257 HDL2b, small, dense HDL3c or total HDL was observed (*Figure 3*). No correlations were
258 found between HDL2b, HDL3c and total HDL cholesterol efflux and clinical and

259 biochemical parameters after adjusting for glycemic control, statin response, sLDL, Apo B
260 and triglyceride levels. As mentioned in Materials and methods, we used phospholipid
261 concentrations to calculate the cholesterol efflux experiments, but we also recalculated it
262 using total proteins, no changes in results were found (HDL 3c subfraction $p= 0.07$ and 2b
263 subfraction $p=0.39$ and total HDL $p= 0.20$). We recalculated the cholesterol efflux taking
264 baseline correlations of subpopulation components and cholesterol efflux showed a positive
265 relationship between total protein content and cholesterol efflux capacity of HDL2b,
266 Spearman Rho 0.31 ($p=0.03$). Regression models with cholesterol efflux capacity of HDL
267 as dependent variable did not reveal significant determinants of the response to treatment.
268 We did not find any correlations with changes in ApoB and composition or function of
269 HDL.

270 Discussion

271 Our study did not find any differences in cellular cholesterol efflux towards large HDL2b
272 and small HDL3c subpopulations, or towards total HDL after a short course of treatment
273 with moderate doses of atorvastatin in patients with well controlled type 2 diabetes.
274 Although simvastatin and fibrates have earlier been tested for their capacity to modify
275 cholesterol efflux properties of HDL in patients with T2DM, this is the first study to
276 evaluate the effect of atorvastatin on this functional metric in patients with T2DM.
277 Furthermore, we report, for the first time, the effect of atorvastatin treatment on cholesterol
278 efflux capacity of HDL subpopulations.

279 Previous studies showed a significant improvement on cholesterol efflux in HDL from
280 THP-1 cells, after treatment with simvastatin and pitavastatin in men with type 2 diabetes

281 and in dyslipidemic patients, respectively. Both studies demonstrated a significant increase
282 in HDL-cholesterol concentrations and used ApoB-depleted serum as cholesterol acceptor.
283 As a consequence, the increase in HDL levels potentially explained the improvement in
284 cholesterol efflux capacity[44][45]. By contrast, we used HDL isolated from plasma by
285 ultracentrifugation rather apoB-depleted serum. Our technique adjusts for alterations in the
286 HDL concentration and phospholipid content, the latter providing a major contribution to
287 the efflux capacity [27]. The methodological differences in cholesterol efflux experiments
288 and the use of other type of statin can explain contradictions between our results and those
289 reported earlier [44][45].

290 Indeed, the different type and doses of statins can have differential effects on HDL function
291 and lipid metabolism[47]. There are two previous studies of atorvastatin on HDL function,
292 although not in patients with type 2 diabetes. One of them found that proteome of small
293 HDL3 of patients with cardiovascular arterial disease (CAD) was altered by the treatment
294 with atorvastatin after one year[53]. Rader et al. studied patients with preexisting CAD, and
295 in accordance with our findings, did not find modification of cholesterol efflux from J774
296 cells to apoB-depleted serum after 16 weeks of treatment with low (10mg) or high (80mg)
297 doses of atorvastatin, even after a modest increase in HDL-cholesterol concentrations[18].
298 These contradictory findings between studies on proteomics of HDL and cholesterol efflux
299 capacity after treatment with atorvastatin highlight the possibility that atorvastatin displays
300 a null effect on cholesterol efflux capacity but positive effects on other biological activities
301 where proteins play a pivotal role, for example its anti-oxidative capacity[46]. New studies
302 are necessary to elucidate this theory.

303 Importantly, our study included only patients with type 2 diabetes with moderately low
304 concentrations of HDL cholesterol at baseline; this alteration is typical and part of the
305 phenotype called “diabetic dyslipidemia”. Our results showed that after treatment, HDL-
306 cholesterol levels decreased even more. We do not believe that the significant diminution of
307 HDL concentration accounted for the lack of the effect of atorvastatin on cholesterol efflux
308 capacity because, as mention earlier, our experiments adjusted for changes in the
309 phospholipid content of HDL. Typically, statins display moderately positive effects on
310 HDL-C levels; however, the effects of statins seem to be altered in patients with T2DM. In
311 2004, CARDS, a multicenter randomized placebo-controlled trial, showed a 9% reduction
312 of HDL-C levels in patients with T2DM after 4 years of treatment with 10 mg of
313 atorvastatin [54]. Chang (2013) reported a high prevalence of diminution of HDL-C (-3%)
314 after one year of atorvastatin treatment in patients with T2DM [55]. Other clinical trials of
315 statins provided similar findings. The mechanism of this response is not fully elucidated but
316 can involve some enzymes and transfer protein, including lipoprotein lipase, hepatic lipase,
317 and phospholipid transfer protein, involved in HDL metabolism and remodeling. Indeed,
318 the function of these proteins has been reported to be impaired in an insulin-resistant
319 milieu[2]. Additionally, it is known that the liver may represent the major source of
320 cholesterol that circulates as HDL-C; prolonging the inhibition of HMG-CoA reductase by
321 statins may, therefore, result in depletion of hepatic cholesterol, leading to decreased
322 production of HDL-C[56].

323 Our studies have limitations. As we evaluated a short course of treatment with moderate
324 doses of atorvastatin, we cannot rule out that higher doses or longer treatment modify the
325 cholesterol efflux capacity of HDL. On the other hand, the strengths of the present study

326 include a higher number of patients in comparison with previous studies of statin effects on
327 cholesterol efflux capacity of HDL [44] as well as homogeneity of patients. As to our
328 knowledge, there is no other study with a similar design and technique to evaluate the
329 cholesterol efflux capacity of HDL under statin treatment. The lack of effect of atorvastatin
330 on cholesterol efflux is consistent with the residual risk observed in these patients even
331 after they reach lipid goals with statins. Our results thereby suggest that some biological
332 activities of HDL can be independent of the statins effect on ApoB or cholesterol levels.

333 In conclusion, our study showed that 10 weeks of treatment with a moderate 20 mg dose of
334 atorvastatin does not modify the cholesterol efflux capacity of HDL particles in patients
335 with well controlled type 2 diabetes. Due to the nature of our study, we cannot, however,
336 translate these results into the commonly prescribed long-term statin therapy neither to
337 other statin types.

338

339 **Conflict of interest**

340 The authors declared they do not have anything to disclose regarding conflict of interest
341 with respect to this manuscript.

342

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348

349 **Author contributions**

350 All authors contributed equally in the design, development, results and discussion of the
351 research.

352

353

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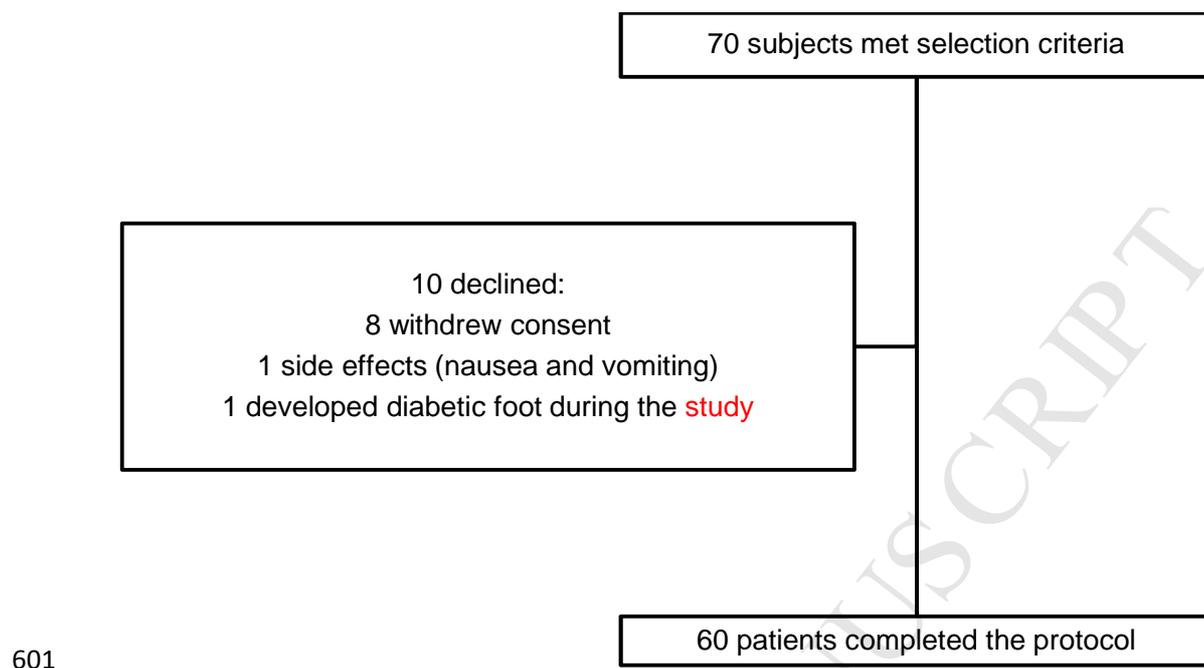
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599 Tables and figures

600



602 *Figure 1.* Study population, screening and follow-up.

603 Table 1 Clinical characteristics of the study population

N	60
Women/men %	62/38
Age yr	58 ± 10
BMI kg/m ²	28.6 ± 3.7
SBP mmHg	127 ± 16
DBP mmHg	78 ± 10
Age at diagnosis yr	50 ± 10
Diabetes evolution years	8 (4-12)
Metformin treatment %	95
Insulin treatment %	30
Other anti-diabetic drugs ^a %	25
Cardiovascular disease history %	39
Smoking history ^b %	21

604 N= 60. Data are presented as mean ± SD or median and interquartile
 605 range (10-90) and as a percentage.

606 ^a Except thiazolidinediones

607 ^b At least one year after smoking withdrawal, Stata 13.

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610 Table 2 Energy intake through the study

	Before ^a treatment	After ^a treatment	<i>p</i> value
Energy intake kcal	1596 ± 439	1666 ± 438	0.21
Carbohydrates gr	206 ± 66	221 ± 62	0.10
Proteins gr	80 ± 19	83 ± 24	0.30
Fat gr	50 ± 21	50 ± 16	0.99
Fiber gr	27.4 ± 7.6	29.4 ± 8.3	0.09
Carbohydrates %	51 ± 6	53 ± 6	0.14
Proteins %	20 ± 2	19 ± 3	0.56
Fat %	28 ± 6	27 ± 5	0.23

611 ^aData are presented as mean ± SD

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618 Table 3.- Clinical parameters before and after the atorvastatin treatment of patients with
 619 T2DM

Characteristics	Before ^a	After ^a	<i>p</i> value
Weight kg (F/M)	66.2 (59.5-71.6)	65.7 (59.5-71.6)	0.27
	80.8 (74.5-86.7)	81.4 (73.8-86.0)	0.78
BMI Kg/m ² (F/M)	29.1 (26.1-31.1)	28.4 (25.9-30.7)	0.29
	28.4 (26.7-32.5)	29.1 (25.8-32.9)	0.70
SBP mmHg	127 ±16	121 ±18	0.06
DPB mmHg	78 ±10	76 ± 9	0.09
Waist cm (F/M)	94.5 (88.5-99)/	91.8 (88.5-99.8)	0.08
	99.5 (95-104.2)	97.5 (95.104.5)	0.06

620 ^a Data are presented as mean ± SD or median and interquartile range (10-90)

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630 Table 4.- Serum biochemistry before and after the atorvastatin treatment of patients with
 631 T2DM

	Before ^a	After ^a	Change %	<i>p</i> value
Total cholesterol mg/dL	184 ±32	127±27	-32	>0.001
LDL-cholesterol mg/dL	98 (82-117)	49 (39-62)	-50	>0.001
HDL-cholesterol mg/dL	47 (42-54)	45 (39-56)	-4	0.02
ApoB mg/dL	91 (82-120)	60 (49-76)	-34	>0.001
ApoA-I mg/dL	147 (129- 163)	143 (124-165)	-3	0.02
Triglycerides mg/dL	154 (113- 228)	125 (103-169)	-19	>0.001
Glucose mg/dL	125 (108- 145)	127 (105-161)	+1.6	0.08
Insulin Ui/L	11.9 (8.4- 18.1)	13.2 (9.8-19.3)	+11	0.16
HbA1c %	6.98 ±0.83	6.91 ±0.99	-2	0.89
Creatinine	0.72±0.16	0.69 ±0.16	-5	0.005
AST	24 (21-30)	23(21-28)	-4	0.02
ALT	23 (19-32)	25 (19-30)	+8	0.53
GGT	20 (15-27)	19 (15-26)	-5	0.11

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633 ^aData are presented as mean ± SD or median (interquartile range 10-90)

634 HbA1c, glycated hemoglobin ; apo B, apolipoprotein B; apo A-I, apolipoprotein A-I;
635 AST, aspartate transaminase; ALT, alanine transaminase; GGT, gamma glutamil
636 transaminase.

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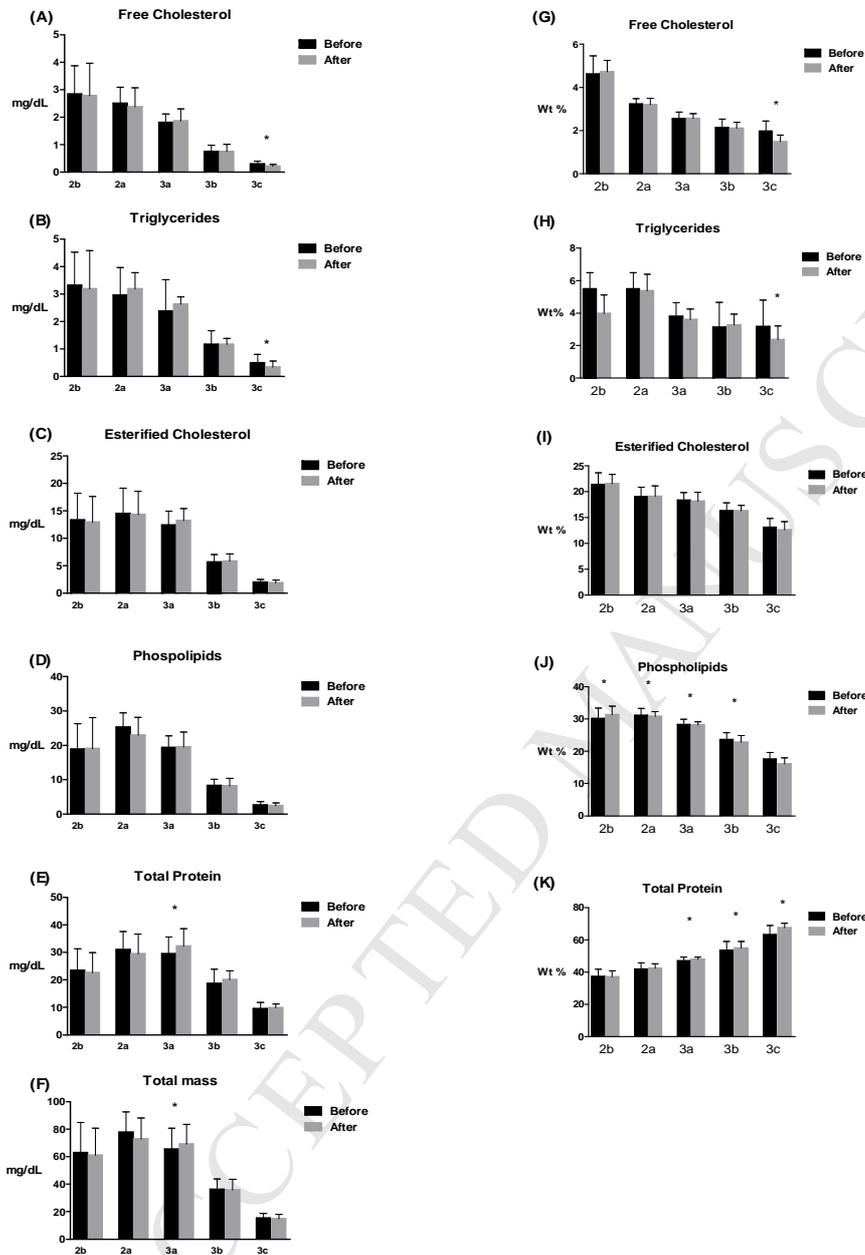
653 Table 5.- Cholesterol efflux capacity of HDL particles before and after the atorvastatin

654 treatment of patients with T2DM

Cholesterol efflux capacity	Before ^a	After ^a	<i>p</i> value
Total HDL %	3.78 (3-12-4.76)	3.59 (3.01-4.34)	0.32
HDL2b subpopulation %	2.86 (2.29-3.85)	2.93 (2.32-3.42)	0.84
HDL3c subpopulation %	6.89 (6.08-8.06)	6.66 (6.04-8.15)	0.88

655 ^aData are presented as median (interquartile range 10-90)

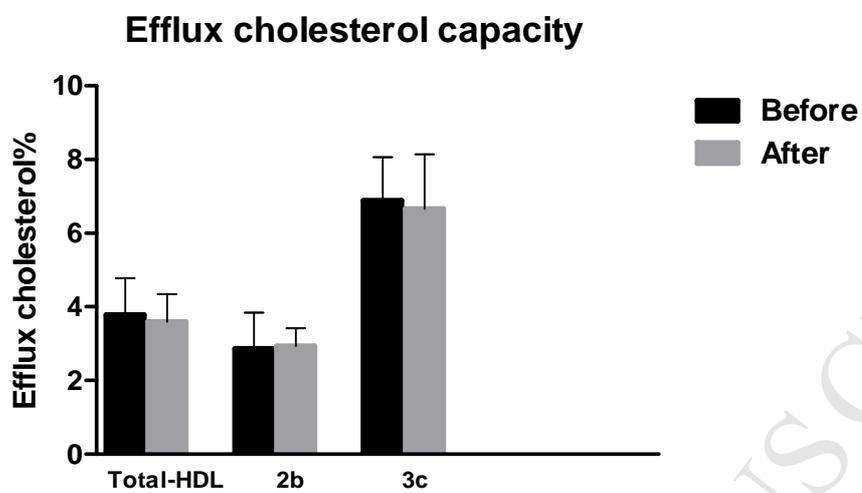
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658 Figure 2. Total mass and chemical composition of HDL subpopulations expressed as
 659 mg/dL (A-F) and as weight percentage of total mass (G-K) before and after atorvastatin
 660 treatment in patients with T2DM.

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663 Figure 3.- Cholesterol efflux from THP-1 cells to total HDL and to HDL2b and 3c

664 subpopulations before and after atorvastatin treatment in patients with T2DM.

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Highlights

1. Cholesterol Efflux capacity of HDL is altered in subjects with type 2 diabetes
2. Low levels of HDL-Cholesterol and mild hypertriglyceridemia are typical in T2DM
3. Statin effects on HDL biological activities are contradictory across studies
4. Different classes of statins could have differentiated effects on HDL