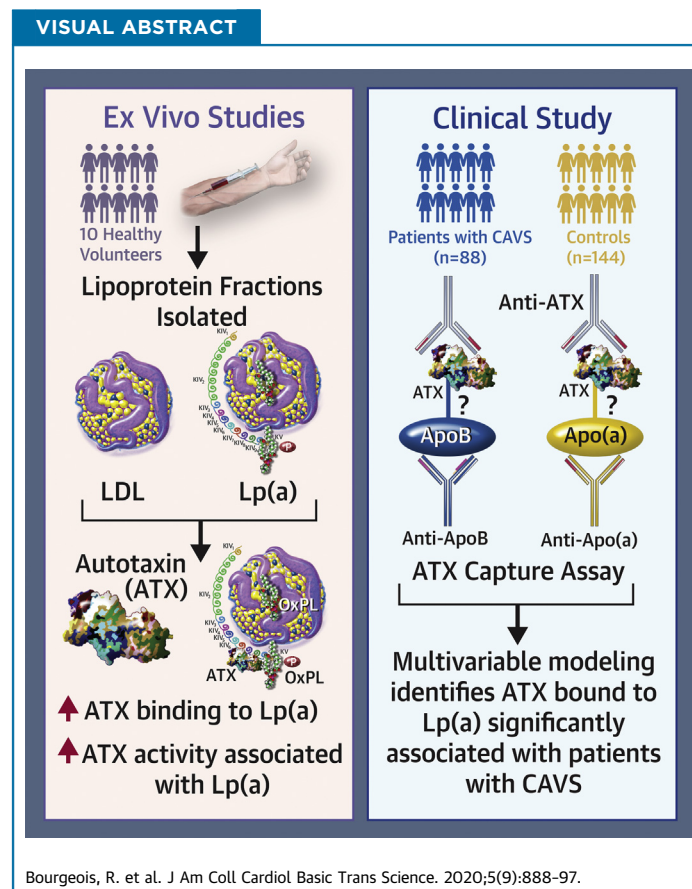


CLINICAL RESEARCH

Interaction of Autotaxin With Lipoprotein(a) in Patients With Calcific Aortic Valve Stenosis



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HIGHLIGHTS

- Lipoprotein(a) is an important carrier of autotaxin and lysophosphatidic acid in the bloodstream.
- Autotaxin might physically interact with apolipoprotein(a).
- The plasma concentration of autotaxin transported by lipoprotein(a) may be an important predictor of calcific aortic valve stenosis.

SUMMARY

Our objectives were to determine whether autotaxin (ATX) is transported by lipoprotein(a) [Lp(a)] in human plasma and if it could be used as a biomarker of calcific aortic valve stenosis (CAVS). We first found that ATX activity was higher in Lp(a) compared to low-density lipoprotein fractions in isolated fractions of 10 healthy participants. We developed a specific assay to measure ATX-Lp(a) in 88 patients with CAVS and 144 controls without CAVS. In a multivariable model corrected for CAVS risk factors, ATX-Lp(a) was associated with CAVS ($p = 0.003$). We concluded that ATX is preferentially transported by Lp(a) and might represent a novel biomarker for CAVS. (*J Am Coll Cardiol Basic Trans Science* 2020;5:888-97) © 2020 The Authors. Published by Elsevier on behalf of the American College of Cardiology Foundation. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Calcific aortic valve stenosis (CAVS) is the most prevalent heart valve disease, affecting 2% to 3% of individuals aged 65 years or older. The main risk factors for CAVS are age, sex, obesity, hypertension, type 2 diabetes (T2D), low-density lipoprotein cholesterol (LDL-C), and smoking (1). The main driving force of CAVS is the calcification of the aortic valve, which leads to a reduction of the aortic valve effective orifice area (2,3). Currently, the only treatment known to improve outcomes in CAVS patients is the surgical removal of the aortic valve.

Over the past 20 years, clinical and molecular studies suggested a potential role for lipoprotein(a) [Lp(a)] in the aortic valve calcification process (4-8). Seeking to identify genetic variants associated with aortic valve calcium accumulation, Thanassoulis et al. (9) identified a variant at the *LPA* locus on chromosome 6, which is strongly associated with plasma Lp(a) levels, as the top candidate locus. Lp(a) was also shown to be an important predictor of valvular outcomes in patients with mild-to-moderate CAVS (10,11). Besides its important role as a carrier of oxidized phospholipids, the molecular mechanisms through which Lp(a) might

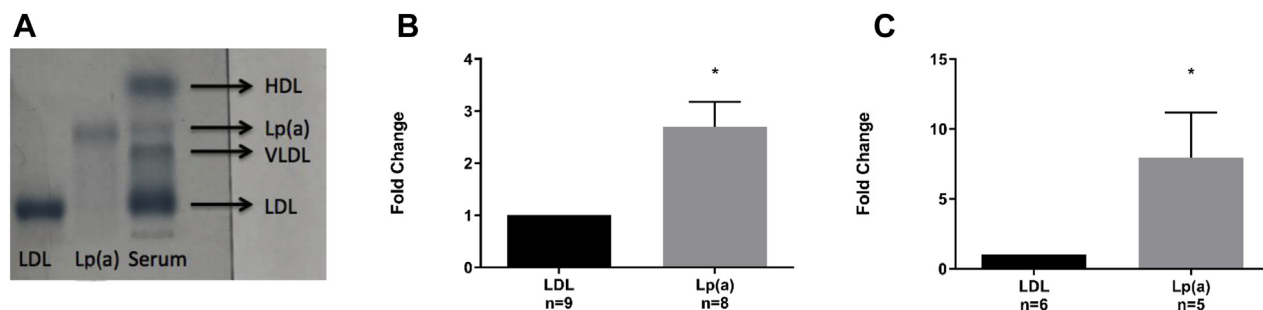
cause aortic valve calcium accumulation are poorly understood.

Autotaxin (ATX), an adipose tissue-derived phospholipase D, converts lysophosphatidylcholine (LysoPC) into lysophosphatidic acid (LysoPA), a bioactive phospholipid that can act as a signaling molecule promoting inflammation, fibrosis, and cell motility (12). LysoPA can also induce an osteogenic transition in valve interstitial cells through its binding to the LPAR1 receptor (13). Bouchareb et al. (14) and Torzewski et al. (15) have previously shown that ATX can associate with Lp(a). However, important questions remain with regard to the nature of the association between ATX and Lp(a) and its potential clinical significance. For instance, it is not known if ATX binds to apolipoprotein(a) or if the activity of ATX (or LysoPA levels) is higher on Lp(a) compared to other lipoproteins with similar size and density such as LDL. Additionally, whether the circulating levels of ATX bound to Lp(a) are associated with CAVS is also unknown. The objective of this study was to gain new knowledge about the potential interaction of ATX with atherogenic lipoproteins and to determine whether a newly developed biomarker,

ABBREVIATIONS AND ACRONYMS

- apo(a)** = apolipoprotein(a)
- apoB** = apolipoprotein B
- ALR** = adiponectin-to-leptin ratio
- ATX** = autotaxin
- ATX-apo(a)** = ATX carried by Lp(a)
- ATX-apoB** = ATX carried by apoB-containing lipoproteins
- BMI** = body mass index
- CAD** = coronary artery disease
- CAVS** = calcific aortic valve stenosis
- HDL** = high-density lipoprotein
- LDL** = low-density lipoprotein
- Lp(a)** = lipoprotein(a)
- LysoPA** = lysophosphatidic acid
- LysoPC** = lysophosphatidylcholine
- OxPLs** = oxidized phospholipids

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FIGURE 1 Autotaxin Activity and Lysophosphatidic Acid Content in Lipoprotein(a) Versus Low-Density Lipoproteins

Purity confirmation by Sebia Hydragel (A), measurement of autotaxin (ATX) activity (B), and lysophosphatidic acid (LysoPA) content (C) in lipoprotein (a) [Lp(a)] and low-density lipoprotein (LDL) of healthy participants. ATX activity (relative light absorbent unit) and LysoPA content (μM) were normalized by cholesterol content ($\mu\text{mol/l}$). Asterisk (*) indicates values significantly different from LDL ($p = 0.016$ for ATX activity [B] and $p = 0.031$ for LysoPA content [C]). VLDL = very low density lipoprotein.

ATX bound to Lp(a) [ATX-apo(a)], is associated with the presence of CAVS.

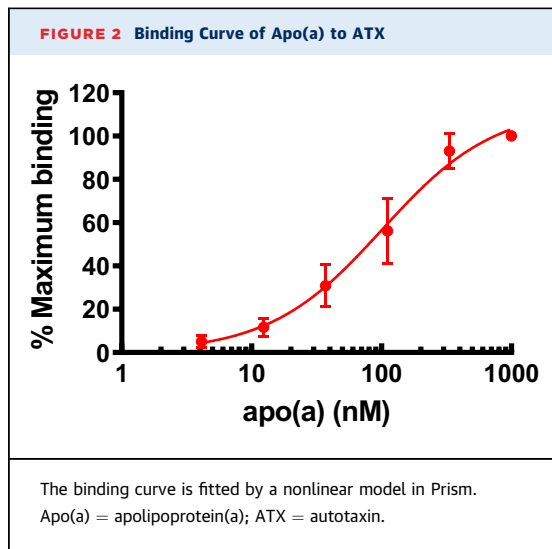
METHODS

STUDY PARTICIPANTS. A series of consecutive patients with mild to severe CAVS who did not undergo aortic valve replacement were recruited at the echocardiography laboratory of the Québec Heart and Lung Institute (QHLI). Exclusion criteria included the presence of mitral valve stenosis, mitral insufficiency (\geq moderate), aortic insufficiency (\geq moderate) and heart failure (ejection fraction $<40\%$). Cases were excluded if they had CAVS of rheumatic etiology or if they had any type of cancer that required radiotherapy in the thoracic area (breast, trachea, bronchus, or lung cancer) before the diagnosis of CAVS. Women were also excluded if they were pregnant or lactating. Monthly interrupted time series were performed to recruit individuals without CAVS undergoing Doppler echocardiography (controls with normal aortic valves). Additional controls were also recruited through the echocardiography laboratory and advertisements at the QHLI. Previous and current medical history included history of smoking, documented diagnoses of hypertension (patients receiving antihypertensive medications or having known but untreated hypertension [blood pressure $\geq 140/90$ mm Hg]), diabetes (fasting glucose ≥ 7 mmol/l or treatment with antidiabetic medication), and detailed information on current medication was collected. Body weight, height, and waist circumference were measured following standardized procedures. Blood pressure and heart rate were also assessed. Peak aortic jet velocity, aortic valve area, peak and mean

transvalvular gradients, and aortic valve morphology were measured by Doppler echocardiography (16-18). Patients were classified as having mild, moderate, or severe CAVS based on peak aortic jet velocity, aortic valve pressure gradient, and aortic valve area as suggested by the American College of Cardiology/American Heart Association Task Force on Practice Guidelines (19). We isolated Lp(a) from 9 healthy volunteers (5 men and 4 women) without aortic stenosis and with high Lp(a) level (>125 nmol/l). The study protocols were approved by the Ethics Committee of the QHLI and all patients signed a written informed consent.

LABORATORY DATA. Overnight fasting blood samples were collected in ethylenediaminetetraacetic acid tubes and immediately processed in the clinical biochemistry laboratory for the measurement of glucose, total cholesterol, LDL-C, high-density lipoprotein cholesterol (HDL-C), and triglyceride levels. Plasma samples were also stored at -80°C and used for measurement of Lp(a) and ATX levels. Plasma Lp(a) levels were measured by turbidimetric assay using the Tina-quant Lipoprotein(a) Gen.2 system (Cobas integra 400/800, Roche Mannheim, Germany). Plasma adiponectin and leptin concentrations were measured by enzyme-linked immunosorbent assay (ELISA) according to the manufacturer's instructions (R and D Systems, Minneapolis, Minnesota). ATX mass was also measured in plasma by ELISA according to the manufacturer's instructions (Echelon Biosciences, Salt Lake City, Utah) (intra-assay coefficient of variation [CV] of $<2\%$).

LDL AND LP(A) ISOLATION AND SEPARATION. To isolate and separate Lp(a) and LDL, blood was



collected into BD-Vacutainer tubes (Becton, Dickinson and Co., Franklin Lakes, New Jersey). Serum was then separated by centrifugation at 3,000 g for 15 min. Lipoproteins were separated into 3 fractions by ultracentrifugation for 9 h at 100,000 g. Lp(a)-containing fractions (assessed by Sebia Hydrigel) were subjected to size exclusion chromatography with Tris-HCl buffer. Lp(a)-containing fractions were identified by western blotting. Positive Lp(a) fractions were then subjected to ion exchange chromatography and LDL and Lp(a) were separated using a NaCl gradient. After purification, samples were concentrated and equilibrated at 150 mM NaCl and purity was confirmed by Sebia Hydrigel (Figure 1A). Lp(a) and LDL fractions were free of exosomes as shown by western blot using CD63 and CD9 antibodies (data not shown). Cholesterol content of lipoprotein subfractions was measured using a commercial kit (Randox, Crumlin, United Kingdom). LysoPA was measured in plasma by ELISA according to the manufacturer's instructions (Echelon Biosciences, Salt Lake City, Utah).

ATX ACTIVITY. Autotaxin activity in isolated fractions of LDL and Lp(a) was determined by assessing the cleavage of the ATX-specific fluorogenic substrate FS-3 (Echelon Biosciences, L-2000). Assays were conducted in a final volume of 210 μ l comprising 180 μ l assay buffer (50 mM Tris-HCl pH 8.0 containing 1 mg/ml bovine serum albumin [BSA], 140 mM NaCl, 5 mM KCl, 1 mM MgCl₂, and 1mM CaCl₂), 5 μ l of FS-3 (final concentration 5 μ M) and 25 μ l of each sample. The reaction was allowed to proceed at 37 °C for 17 h. Fluorescence was measured with a FilterMax F3 microplate reader (Molecular Devices, Wokingham,

United Kingdom). ATX activity was quantified by measuring the fluorescence increase at 528 nm with excitation at 485 nm.

CAPTURE ASSAY. The detection of ATX on lipoproteins containing apo(a) [Lp(a)] or apoB-100 was measured with an antibody capture assay. Microtiter 96-well medium binding plates (Greiner Bio-One, Kremsmunster, Austria) were coated with anti-apoB-100 (ab20737, Abcam, Cambridge, United Kingdom) or anti-apo(a) (MABS1284, Millipore, Burlington, Massachusetts) antibodies overnight at 4°C (1 μ g/ml, 50 μ l/well). Excess material was removed. Plates were then blocked for 2 h with PBST -5% BSA. After washing, plasma was added at 1:5 dilution (phosphate-buffered saline solution with Tween [PBST] -1% BSA; 50 μ l/well) for 2 h at 37°C. After washing, rabbit anti-human ATX antibody (PA5-12478, ThermoFisher, Waltham, Massachusetts) (4 μ g/ml; 50 μ l/well), was added and incubated for 2 h. Horseradish peroxidase-labeled goat anti-rabbit antibody (AP132P, Millipore) (0.03 μ g/ml; 50 μ l/well) was added overnight at 4°C. After a final washing step, 100 μ l of o-phenylenediamine dihydrochloride (OPD) (34005, ThermoFisher) was added for 30 min, followed by 100 μ l of sulfuric acid (2.5 M) to stop the reaction. Absorbance was read on a spectrophotometer at 492 nm according to the manufacturer's instructions. Results were expressed in relative light absorbance units after subtraction of the background (PBST-1% BSA).

BINDING ASSAY. High-binding capacity microtiter 96-well plates (polystyrene, Greiner Bio-One) were coated overnight at 4°C with recombinant ATX (5 μ g/ml) in carbonate buffer. Excess material was removed, and plates were blocked with PBST-1% BSA (150 μ l/well) for 1 h. After washing, increasing concentrations of PBST-diluted apo(a) (0.05 μ M to 1 μ M; 100 μ l/well) were added and incubated for 2 h at room temperature. After washing, apo(a) antibody (1 μ g/ml, 100 μ l/well) was added for 1 h at room temperature, followed by the secondary horseradish peroxidase-conjugated antibody for 45 min (1/10,000 dilution; 100 μ l/well). Following PBST-1% BSA washes, absorbance was read at 405 nm. Purified apo(a) was obtained following the previously described method (20). Plasmid encoding a 17-kringle-containing form of recombinant apo(a) was constructed and transfected into human embryonic kidney (HEK 293) cells as previously described (20,21). Recombinant-apo(a) was purified from the conditioned medium of stably-expressing cell lines by lysine-Sepharose (GE Healthcare, Chicago, Illinois) affinity chromatography as previously reported (20).

TABLE 1 Clinical Characteristics of Study Participants

	Total N = 232	Without CAVS N = 144	With CAVS N = 88	p Value
Clinical				
Age, yrs	64.8 ± 8.9	62.7 ± 7.5	68.3 ± 9.8	<0.001
Male	133 (57.3)	74 (51.4)	59 (67.1)	0.019
BMI, kg/m ²	28.8 ± 5.5	27.6 ± 5.0	30.8 ± 5.8	<0.001
Waist circumference, cm	101.1 ± 14.8	97.2 ± 13.2	107.5 ± 15.2	<0.001
Hypertension	116 (50.0)	49 (34.0)	67 (76.1)	<0.001
Diabetes	47 (20.3)	20 (13.9)	27 (30.7)	0.002
Obesity (BMI ≥30 kg/m ²)	86 (37.1)	40 (27.8)	46 (52.3)	<0.001
Medication				
Statins	121 (52.2)	53 (36.8)	68 (77.3)	<0.001
Calcium-channel inhibitors	46 (19.8)	12 (8.3)	34 (38.6)	<0.001
ACE inhibitors	44 (19.0)	20 (13.9)	24 (27.3)	0.013
ARA	51 (22.0)	19 (13.2)	32 (36.4)	<0.001
Bicuspid AV	15 (6.5)	1 (0.7)	14 (15.9)	<0.001
CAVS severity				<0.001
mild	26 (11.2)	0 (0)	26 (11.2)	
moderate	42 (18.1)	0 (0)	42 (18.1)	
severe	20 (8.6)	0 (0)	20 (8.6)	
Laboratory Data				
LDL cholesterol, mg/dl	93.3 ± 39.3	101.8 ± 39.3	79.3 ± 35.3	<0.001
HDL cholesterol, mg/dl	53.0 ± 17.5	53.7 ± 17.7	51.8 ± 17.1	0.423
Triglycerides, mg/dl	156.9 ± 80.0	160.4 ± 77.3	151.1 ± 84.2	0.390
Lp(a), nmol/l	73.3 ± 101.4	63.3 ± 87.9	89.6 ± 119.0	0.055
ATX, ng/ml	266.9 ± 115.4	264.2 ± 79.7	271.4 ± 157.7	0.645
ATX-apo(a)	0.14 ± 0.14	0.12 ± 0.12	0.17 ± 0.17	0.034
ATX-apoB	0.07 ± 0.12	0.07 ± 0.10	0.08 ± 0.14	0.257
Values are mean ± SD or n (%)				
ACE = angiotensin-converting enzyme; apo(a) = apolipoprotein(a); apoB = apolipoprotein B-100; ARA = angiotensin II receptor antagonists; ATX = autotaxin; BMI = body mass index; CAVS = calcific aortic valve stenosis; HDL = high-density lipoprotein; LDL = low-density lipoprotein; Lp(a) = lipoprotein(a).				

STATISTICAL ANALYSIS. Continuous data are expressed as mean ± SD and were compared between 2 groups using Student's *t*-test; and categorical data are presented as counts with percentage and compared using the chi square test. Normality of continuous variables was assessed by the Shapiro-Wilk test and log-transformed variables were entered in the model if they were not normally distributed. Univariable and multivariable logistic regression models were performed to examine the association between ATX-apo(a) and the presence of CAVS. The variables selected for the models were influenced by CAVS case-control status ($p < 0.05$). Results are reported as odds ratios (ORs) with 95% confidence intervals (CIs). Logistic regression was used to determine the respective contribution of ATX-apo(a) and body mass index (BMI) based on the cutpoints that were identified by sensitivity and specificity analyses (area under the receiver-operating characteristic curve for CAVS presence). Spearman's correlation coefficient (r_s) was used

to compare 2 continuous variables. Statistical analyses were performed using JMP software (V.14). A p value ≤ 0.05 was considered statistically significant.

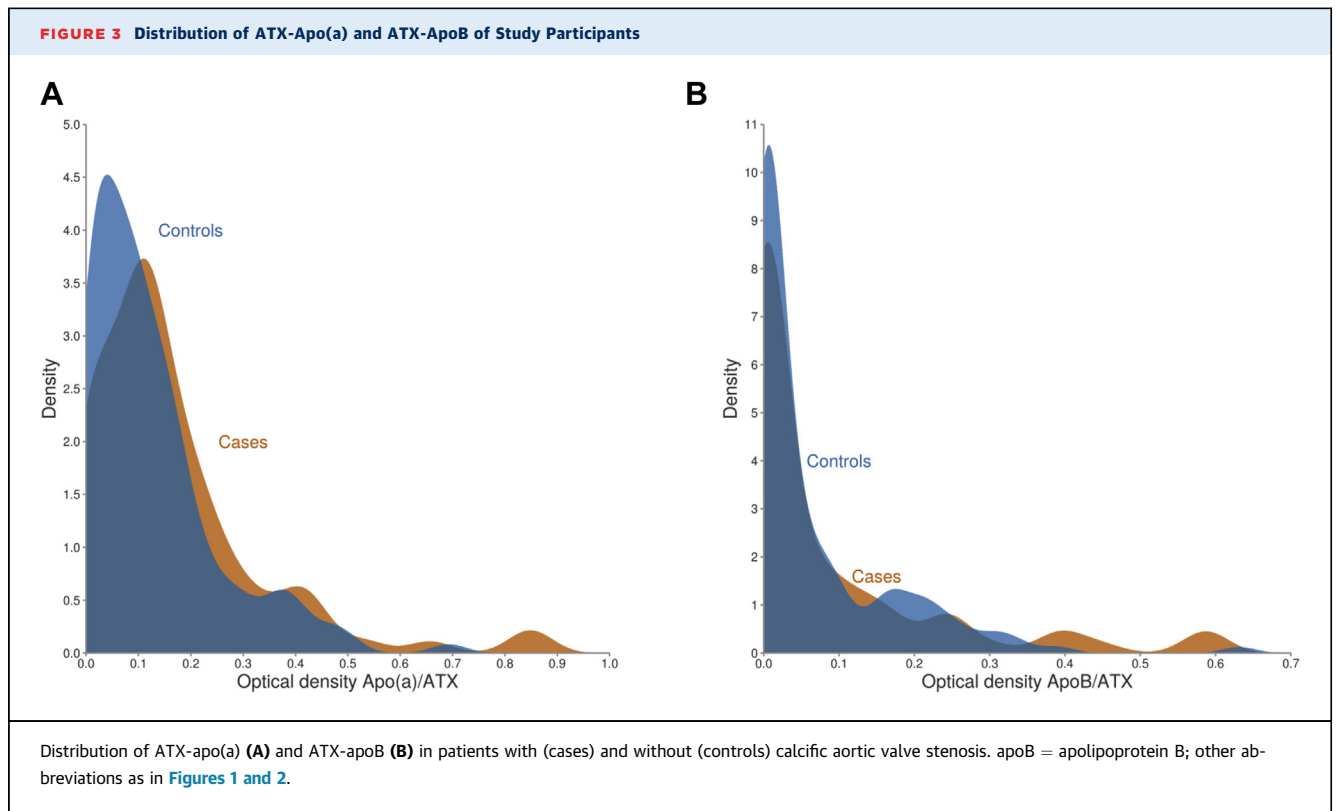
RESULTS

COMPARISON OF ATX ACTIVITY BETWEEN LDL AND LP(A). We first undertook to establish a reliable and reproducible method for LDL and Lp(a) fraction separation validated using the Sebia Hydragel results (Figure 1A). To investigate the preferential association of ATX with Lp(a), we measured ATX activity in isolated fractions of LDL and Lp(a) from healthy participants. ATX activity was significantly higher in Lp(a) compared to the LDL fraction (Figure 1B). In addition, LysoPA content was also significantly higher in the Lp(a) fraction compared to the LDL fraction (Figure 1C).

INVOLVEMENT OF APO(A) MOIETY IN ATX BINDING. Although a physical association between Lp(a) and ATX has been previously reported, it is presently unknown whether the interaction occurs within the LDL component or with apo(a). By using recombinant ATX coated in 96 well plates we assessed using a binding assay the interaction with purified apo(a). In this assay, we observed a dose-dependent interaction with a plateau at a concentration of $\sim 1,000$ nM of apo(a) (Figure 2).

MEASUREMENT OF ATX BOUND TO APO(A) IN PLASMA. Having established that ATX activity is higher in isolated Lp(a) fractions, we sought to establish a quantitative assay to measure ATX-apo(a) in the plasma. We developed a capture assay to measure ATX bound to apo(a) and apoB (see Methods). In 232 subjects, including control and CAVS cases (Table 1), we found that the distribution of both ATX-apo(a) and ATX-apoB were skewed to the left (Figure 3). ATX-apo(a) correlated significantly ($r_s = 0.553$; $p < 0.0001$) (Spearman correlation) with ATX-apoB, but not with Lp(a) or total ATX level measured by ELISA (Figure 4).

ASSOCIATION BETWEEN ATX AND CAVS. We next evaluated whether the ATX capture assay on lipoproteins is a predictor of CAVS. Clinical and laboratory characteristics of patients with and without CAVS are presented in Table 1. Significant differences in some of the clinical risk factors (age, sex, BMI, and hypertension) were noted with patients in the CAVS group, having a higher proportion of males, patients with hypertension, T2D, and obesity. The proportion of patients treated with statins, calcium channel



inhibitors, angiotensin-converting enzyme (ACE) inhibitors and angiotensin receptor blockers (ARB) was also higher in the CAVS group. Also, patients with CAVS had higher levels of ATX-apo(a). Levels of circulating ATX and ATX-apoB were similar between the 2 groups.

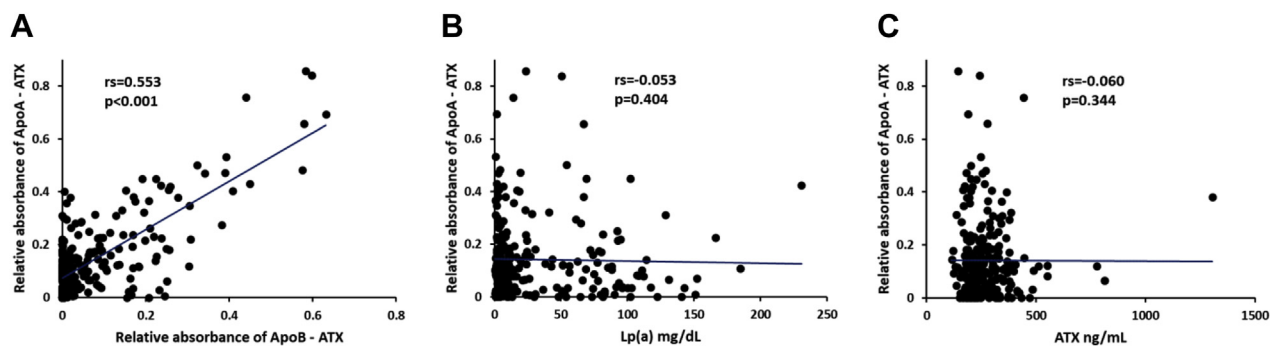
For ATX-apo(a), we selected a cutpoint to simultaneously maximize the sensitivity and specificity (area under the receiver-operating characteristic curve) of the model. This was not done for ATX-apoB because we found no difference between CAVS case and controls. In a multivariable model (model 1), corrected for age, sex, BMI, T2D, hypertension, pharmacotherapy (calcium channel blockers and statins), HDL-C, and corrected LDL-C, we found that circulating ATX-apo(a) was associated with CAVS (Table 2). In a second model (model 2) where log [Lp(a)] was added into model 1, ATX-apo(a) remained associated with the risk of CAVS. We ran the same analyses replacing BMI with the adiponectin-to-leptin ratio (a marker of adipose tissue dysfunction) and found comparable results (data not shown).

CONTRIBUTIONS OF ATX-APO(A) AND BMI TO CAVS. Given that circulating ATX is primarily secreted by adipose tissue, we examined the

possible joint contributions of ATX-apo(a) and obesity on the CAVS risk. We then performed exploratory analyses with groups based on cutpoints ATX-apo(a) and BMI that showed the highest sensitivity and specificity according to the area under the receiver-operating characteristic curve. After corrections (same as model 2) we found that, compared to patients with lower ATX-apo(a) and BMI (referent), patients with higher ATX-apo(a) and BMI had a substantially increased risk of CAVS presence (OR: 8.15; 95% CI: 2.81 to 22.77; $p < 0.001$) (Figure 5). Finally, we sought to determine whether patients with T2D had higher or lower levels of ATX-apo(a) or ATX-apoB compared to patients without T2D and found no differences. We also separated our study population according to BMI and according to the adiponectin-to-leptin ratio (equal or above versus lower than top tertile) and again, we found no differences (data not shown).

DISCUSSION

Over the past 5 years, a series of genetic association, observational, and mechanistic studies have provided evidence that Lp(a) may represent a causal risk factor for CAVS. The mechanisms through which Lp(a) causes CAVS remain elusive. In this study, we aimed to

FIGURE 4 Spearman Correlation Between Plasma Levels of Biomarkers

Spearman correlation between plasma levels of ATX-apo(a) and ATX-apoB (A), ATX-apo(a) and Lp(a) (B), ATX-apo(a) and ATX (C) Abbreviations as in Figures 1 to 3.

further document the relationship between Lp(a) and CAVS risk by studying the interaction between Lp(a) and ATX. We first showed that ATX activity and LysoPA content were significantly elevated in the Lp(a) compared to LDL fraction. Additional in vitro experiments revealed that ATX binds to the apo(a) moiety of the Lp(a) particles at very low (nM) concentrations. We then conducted a clinical investigation with a case-control design showing that circulating ATX bound to Lp(a) was associated with CAVS. The combination of an elevated ATX- apo(a) and an elevated BMI was associated with CAVS. Although additional studies will be needed to confirm these findings, this analysis suggests that ATX-apo(a) may

represent a potential link between obesity, Lp(a), and CAVS and a new biomarker of CAVS risk.

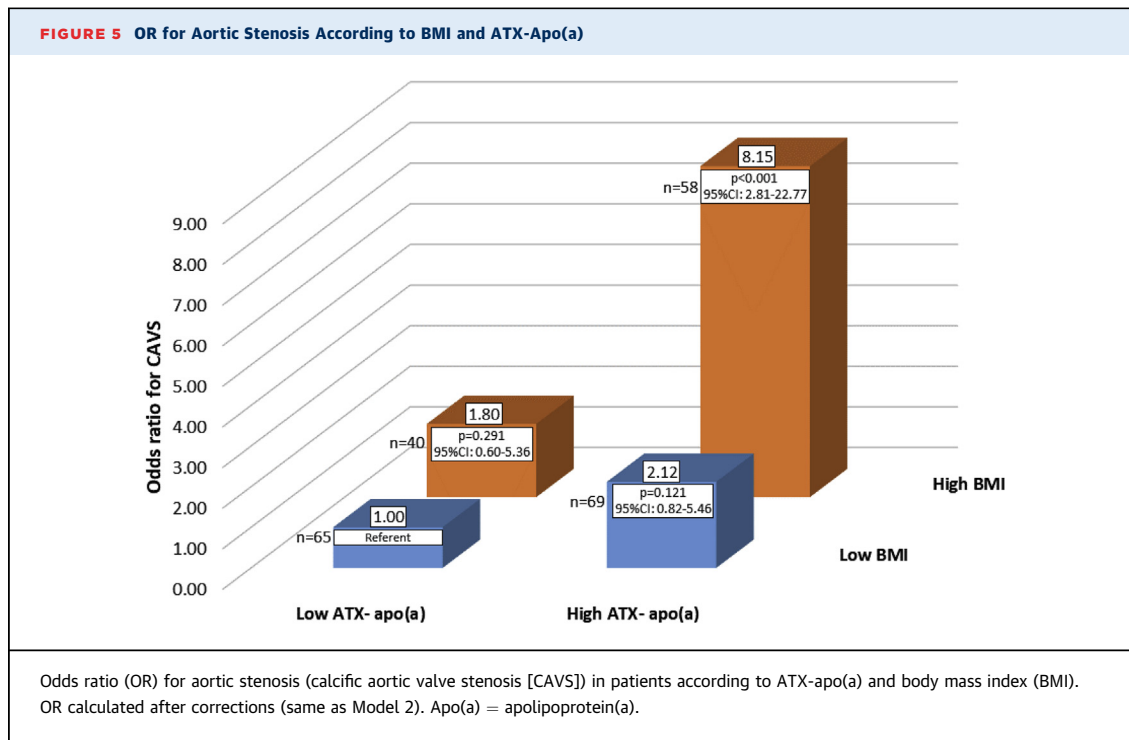
ATX was first identified as a motility factor in tumorigenesis (22,23). It is a secreted glycoprotein that belongs to the ectonucleotide pyrophosphatase/phosphodiesterase (NPP) family. ATX converts LysoPC into LysoPA, which promotes cell motility, inflammation, fibrosis, and calcification (24,25). Several tissues or organs may contribute to circulating levels of ATX or LysoPA. Adipose tissue-specific deletion of ATX resulted in a 40% reduction of plasma LysoPA levels (26-28). Rancoule et al. (29) have also reported evidence that ATX expression is considerably higher in the visceral adipose tissue of obese compared to non-

TABLE 2 Univariable and Multivariable Logistic Regression Analyses for CAVS Presence

	Univariable Analysis		Model 1*		Model 2†	
	OR (95% CI)	p Value	OR (95% CI)	p Value	OR (95% CI)	p Value
Age, yrs	1.08 (1.05-1.12)	<0.001	1.05 (1.01-1.10)	0.017	1.05 (1.01-1.10)	0.018
Male	1.92 (1.11-3.35)	0.019	1.53 (0.73-3.20)	0.254	1.53 (0.73-3.21)	0.256
BMI, kg/m ²	1.12 (1.06-1.18)	<0.001	1.10 (1.02-1.18)	0.012	1.10 (1.02-1.18)	0.013
Diabetes, yes	2.74 (1.43-5.28)	0.002	0.60 (0.23-1.57)	0.294	0.60 (0.23-1.57)	0.295
Hypertension, yes	6.19 (3.40-11.26)	<0.001	1.41 (0.43-4.70)	0.574	1.41 (0.42-4.70)	0.575
CA channel inhibitors, yes	6.93 (3.34-14.38)	<0.001	3.52 (1.35-9.19)	0.008	3.52 (1.34-9.25)	0.009
Statins, yes	5.84 (3.20-10.67)	<0.001	2.35 (0.94-5.89)	0.066	2.35 (0.92-5.98)	0.072
ACE inhibitors, yes	2.33 (1.19-4.52)	0.013	1.62 (0.48-5.49)	0.437	1.62 (0.48-5.49)	0.437
ARA, yes	3.76 (1.96-7.20)	<0.001	2.34 (0.69-7.95)	0.169	2.34 (0.69-7.96)	0.170
Corrected LDL-C, mg/dl	0.96 (0.95-0.98)	<0.001	1.00 (0.97-1.02)	0.715	1.00 (0.96-1.03)	0.775
HDL-C, mg/dl	0.99 (0.98-1.01)	0.419	1.02 (0.99-1.04)	0.196	1.02 (0.99-1.04)	0.196
Log(Triglycerides, mg/dl)	0.69 (0.39-1.20)	0.187	0.61 (0.27-1.38)	0.232	0.61 (0.27-1.38)	0.232
Log(Lp(a), nmol/l)	1.10 (0.93-1.31)	0.272	1.00 (0.74-1.34)	0.982	—	—
ATX- apo(a)>0.096	2.10 (1.21-3.64)	0.007	2.80 (1.39-5.66)	0.003	2.80 (1.39-5.66)	0.003

*Model 1: Age, sex, BMI, diabetes, hypertension, calcium channel inhibitor, statins, ACE inhibitors, ARA, corrected LDL-C, HDL, triglycerides, ATX- apo(a). †Model 2: Model 1 + Lp(a).

CA = calcium; CI = confidence interval; HDL-C = high-density lipoprotein cholesterol; LDL-C = low-density lipoprotein cholesterol; OR = odds ratio; other abbreviations as in Table 1.



obese individuals. We have also reported that bariatric surgery is associated with reduction in ATX plasma levels in patients with severe obesity (30). In addition, 2 large prospective studies (24) have reported that excess body weight and abdominal obesity were associated with CAVS incidence.

Lp(a) is an important carrier of oxidized phospholipids (OxPLs). LysoPA, which is also generated during the oxidation of lipoproteins [LDL or Lp(a)] (31-33), is a bioactive lipid that promotes a pro-inflammatory state and drives the mineralization of the aortic valve (14). In human explanted CAVS tissues, ATX activity and LysoPA levels were increased by several-fold compared to noncalcified aortic valves (14). The pro-inflammatory effect as well as the mineralization associated with LysoPA occurs through an RHOA/NF-κB/ bone morphogenetic 2 (BMP2) pathway (13). In a mouse model of CAVS, the exogenous administration of LysoPA exacerbated the mineralization of the aortic valve (14). However, because mice do not express apo(a), the interaction between Lp(a), apo(a), and ATX needs to be documented in humans. Our data suggest protein-protein interaction between ATX and apo(a) both in vitro and within human plasma. These findings are in line with a higher activity of ATX and an elevated LysoPA content in the Lp(a) compared to the LDL fraction (14).

A recent study by Zheng et al. (34) showed the role of Lp(a) in aortic valve microcalcification and CAVS

progression in a combined analysis of the SALTIRE (Study Investigating the Effect of Drugs Used to Treat Osteoporosis on the Progression of Calcific Aortic Stenosis) and Ring of Fire studies. In that analysis, similar to our results, no correlations were observed between circulating levels of Lp(a) and ATX-apo(a). That analysis also reported that participants with higher ATX-apo(a) levels have faster hemodynamic progression of CAVS. However, this finding was limited to patients with lower Lp(a) levels. Altogether, although replication of our work will be required, our results as well as those of Zheng et al. (34) suggest that ATX-apo(a) might be relevant for both the initiation and the progression of CAVS.

STUDY LIMITATIONS. Lp(a) is a significant risk factor for coronary artery disease (CAD) and CAVS. Nsaibia et al. (35) has previously shown that circulating ATX activity was higher in patients with CAD and CAVS versus patients with only CAD. In the present work, we emphasized that ATX-Lp(a) is a predictor of CAVS. As this work is descriptive and included a small number of individuals, we believe that prospective studies are needed to further investigate the potential causal role of ATX-apo(a) on the development of CAVS. Although our data suggest that ATX-apo(a) was associated with the presence of CAVS, we could not determine whether ATX activity specifically in the Lp(a) or LDL fraction were higher in patients with CAVS compared to controls. Additionally, although

our results support the notion that the circulating ATX-apo(a) complex may arise from adipose, this has yet to be experimentally demonstrated. In this regard, Otto et al. (36) identified fibrofatty (intracellular neutral lipid) degeneration of aortic valves with aging, including the presence of adipose cells interposed between the fibrosa and ventricularis layers of the leaflet. Whether these cells secrete ATX is unknown at the moment.

CONCLUSIONS

The present study showed that ATX activity and LysoPA contents are higher in Lp(a) compared to LDL fraction. ATX bound to Lp(a) is strongly associated with the presence of CAVS. Hence, the pathophysiologic role of Lp(a) in aortic valve disease could be attributable, at least in part, to the unique proteome and lipidome of the Lp(a) particle.

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PERSPECTIVES

COMPETENCY IN MEDICAL KNOWLEDGE: Lp(a) carries ATX and LysoPA in the bloodstream and may contribute to CAVS.

TRANSLATIONAL OUTLOOK: Further studies should determine whether lowering of Lp(a) and Lp(a)-related molecules such as ATX and LysoPA may prevent the development and/or progression of CAVS.

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