# High-density lipoprotein subfractions display proatherogenic properties in overweight and obese children

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**BACKGROUND:** In adults, obesity-driven inflammation can lead to increased cardiovascular disease (CVD). However, information regarding childhood obesity and its inflammatory sequelae is less well defined. Serum amyloid-A (SAA) is an inflammatory molecule that rapidly associates with highdensity lipoproteins (HDLs) and renders them dysfunctional. Therefore, SAA may be a useful biomarker to identify increased CVD potential in overweight and obese children.

**METHODS:** Young Hearts 2000 is a cross-sectional cohort study in which 92 children who were obese were matched for age and sex with 92 overweight and 92 lean children. HDL<sub>2</sub> and HDL<sub>3</sub> (HDL<sub>283</sub>) were isolated from plasma by a three-step rapid-ultracentrifugation procedure. SAA was measured in serum and HDL<sub>283</sub> by an enzyme-linked immunosorbent assay procedure, and the activities of cholesterol ester transfer protein (CETP) and lecithin cholesteryl acyltransferase (LCAT) were measured by fluorimetric assays.

**RESULTS:** Trends across the groups indicated that SAA increased in serum and  $HDL_{283}$  as BMI increased, as did  $HDL_2$ -CETP and HDL<sub>2</sub>-LCAT activities.

**CONCLUSION:** These results have provided evidence that overweight and obese children are exposed to an inflammatory milieu that impacts the antiatherogenic properties of HDL and that could increase CVD risk. This supports the concept that it is important to target childhood obesity to help minimize future cardiovascular events.

Cardiovascular disease (CVD) is a leading cause of mortality and morbidity in the developed world (1), and its incidence is augmented with a concomitant increase in BMI (2,3). Furthermore, the global prevalence of childhood obesity has reached significant proportions, with more than one-third of children and adolescents being classed as either overweight or obese (4); this has led to fears that younger generations may experience the cardiovascular-associated complications identified in obese adult populations (3,5).

This is especially relevant because the obesity-related pathologies of atherosclerosis are known to begin in childhood (6) and track through into adult life (7), thus exposing these individuals to a continuous inflammatory milieu over their life-course. Conversely, minimizing obesity-related inflammation in young people may have far-reaching cardioprotective effects (5). This is important because obesity is characterized by adiposopathy and increased inflammation (8). One molecule that is released from hypertrophic adipocytes and that has been shown to predict CVD, is serum amyloid-A (SAA) (9,10). Furthermore, SAA has been suggested as a candidate molecule linking increased fat load and its comorbidities (11). Once in the circulation, SAA rapidly associates with high-density lipoproteins (HDLs), especially the smaller of the HDL subfractions, HDL<sub>2</sub>, rendering them dysfunctional (12) with numerous proinflammatory actions (13,14). Dysfunctional HDL has reduced reverse cholesterol transport capabilities (15) and antioxidant abilities (16). In addition, two enzymes involved in HDL remodeling, cholesteryl ester transfer protein (CETP) and lecithin cholesterol acyl transferase (LCAT), are altered to an atherogenic phenotype in the presence of SAA (16,17).

However, information regarding obesity-driven inflammation in younger generations is limited, with only a few studies having examined this process in serum. To date, studies have mainly focused on the more traditional markers of inflammation such as C-reactive protein, tumor necrosis factor- $\alpha$ , and interleukin-6 and -8 (18–22) showing consistent associations with obesity. We have also examined several of these inflammatory molecules in the cohort described here, showing that low-grade inflammation was associated with increased CVD risk (23). To our knowledge, only three studies have examined SAA in children. In the first, the authors found that SAA was not affected by the presence of increased weight, which we suggest may due to the young age of their prepubertal children (<10 y) (24). In a second study, both groups were obese, therefore limiting the likelihood of identifying any differences in SAA (25). On the other hand, in a third study, SAA was shown to be directly related to increased adiposity in a group of children with a mean age of 12 y (26). To date, an examination of the influence of SAA-related inflammation on HDL function in children is missing from the literature; this study addresses this shortcoming by examining age- and sex-matched obese, overweight, and normal-weight children. Inflammation was assessed by measuring SAA and its association with HDL's two major subfractions, that is, HDL<sub>2</sub> and HDL<sub>3</sub> (HDL<sub>283</sub>), and its effect on the functioning of HDL<sub>283</sub> was examined by measuring the activities of CETP and LCAT. These analyses will assist in

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determining if the function of HDL is altered to a more proatherogenic phenotype in overweight and obese children.

# RESULTS

# Subject Characteristics

Subject characteristics are presented in **Table 1**. The three groups were matched for age and gender, and, as expected, BMI increased across the groups (normal weight > overweight > obese,  $P \le 0.001$ ). Similarly to BMI, triglyceride levels also increased across the groups ( $P \le 0.001$ ), and total cholesterol was higher in the overweight and obese children as compared with the normal-weight children ( $P \le 0.05$  for both comparisons). In addition, HDL cholesterol decreased with increasing BMI ( $P \le 0.001$ ).

## Serum Analyses

Both high-sensitive C-reactive protein and SAA increased with increasing BMI (**Table 2**;  $P \le 0.001$  for both analytes). However, the activities of serum CETP and LCAT were unaffected by BMI (**Table 2**), although there was a trend for the activity of serum CETP to increase as BMI increased (P = 0.076).

## HDL, and HDL, Analyses

SAA associated with HDL<sub>2&3</sub> followed a similar trend to that of serum SAA, in which it increased within both HDL<sub>2&3</sub>, as BMI increased (**Table 3**,  $P \leq 0.001$  for both subfractions). In addition, SAA associated with HDL<sub>2</sub> was significantly higher in the obese children as compared with both the overweight

## Table 1. Subject characteristics

Weight category ( $n = 92$ per group)	Normal-weight	Overweight	Obese	Pª
Males (%)	45.7	45.1	45.7	0.996
Age (years)	13.2 (1.5)	13.2 (1.5)	13.2 (1.5)	0.995
BMI (kg/m²)	19.0 (2.2) <sup>b</sup>	24.4 (1.7) <sup>c</sup>	30.4 (2.8) <sup>d</sup>	<0.001
Cholesterol (mmol/l)	4.01 (0.81) <sup>b</sup>	4.40 (0.69) <sup>c</sup>	4.23 (0.67) <sup>c</sup>	<0.002
Triglyceride (mmol/l)	0.65 (0.48, 0.86) <sup>b</sup>	0.80 (0.61, 1.02) <sup>c</sup>	1.02 (0.73, 1.45) <sup>d</sup>	<0.001
HDL-Chol (mmol/l)	1.43 (0.32) <sup>b</sup>	1.25 (0.31) <sup>c</sup>	1.11 (0.24) <sup>d</sup>	<0.001

Results are mean (SD), where data were normally distributed, or for triglycerides are geometric mean (interquartile range), where data were not normally distributed and were logarithmically transformed. Differences in percentage of males between groups were calculated using a  $\chi^2$  test.

HDL-Chol, high-density lipoprotein cholesterol.

<sup>a</sup>Significance level for difference between groups, calculated using one-way ANOVA. <sup>bcd</sup>Values with different superscripts indicate differences between groups (Newman–Keuls *post hoc* test).

#### Table 2. Serum analyses of systemic inflammation and lipoprotein-associated enzymes

Weight category ( $n = 92$ per group)	Normal-weight	Overweight	Obese	Pª
CRP (µg/l)	0.18 (0.09, 0.32) <sup>b</sup>	0.41 (0.24, 0.86) <sup>c</sup>	0.93 (0.49,1.60) <sup>d</sup>	<0.001
SAA (µg/l)	5,086 (2,693; 7,783) <sup>b</sup>	8,219 (4,203; 15,614) <sup>c</sup>	11,980 (7,878; 19,411) <sup>d</sup>	<0.001
CETP (µmol/l)	279 (106)	292 (91)	311 (82)	0.076
LCAT (ratio)	1.04 (0.18)	1.03 (0.18)	1.05 (0.17)	0.616

Results are mean (SD), where data were normally distributed, or for SAA are geometric mean (interquartile range), where data were not normally distributed and were logarithmically transformed, or for CRP are median (interquartile range), where logarithmic transformations did not normalize the distributions.

CETP, cholesterol ester transfer protein; CRP, C-reactive protein; LCAT, lecithin cholesteryl acyltransferase; SAA, serum amyloid-A.

<sup>a</sup>Significance level for difference between groups, calculated using one-way ANOVA. <sup>bcd</sup>Values with different superscripts indicate differences between groups (Newman–Keuls *post hoc* test).

## Table 3. HDL<sub>2</sub>- and HDL<sub>3</sub>-associated inflammation and HDL<sub>2</sub>- and HDL<sub>3</sub>-associated enzymes

Weight category ( $n = 92$ per group)	Normal-weight	Overweight	Obese	Pª
SAA HDL <sub>2</sub>	0.80 (0.00, 1.63) <sup>b</sup>	0.97 (0.00, 2.31) <sup>b</sup>	2.23 (1.25, 4.16) <sup>c</sup>	<0.001
SAA HDL <sub>3</sub>	0.96 (0.41, 1.38) <sup>b</sup>	1.50 (0.75, 2.15) <sup>c</sup>	1.87 (1.08, 2.98) <sup>c</sup>	<0.001
CETP HDL <sub>2</sub>	1.00 (0.64) <sup>b</sup>	1.28 (0.73) <sup>b</sup>	1.67 (1.43) <sup>c</sup>	<0.001
CETP HDL <sub>3</sub>	0.09 (0.03)	0.09 (0.03)	0.10 (0.03)	0.330
LCAT HDL <sub>2</sub>	0.04 (0.03) <sup>b</sup>	0.06 (0.05) <sup>c</sup>	0.08 (0.07) <sup>c</sup>	<0.001
LCAT HDL <sub>3</sub>	0.49 (0.21)	0.52 (0.24)	0.53 (0.23)	0.470

Results are mean (SD) where data were normally distributed, or for SAA in HDL<sub>26.3</sub> are geometric mean (interquartile range), where data were not normally distributed and were logarithmically transformed. SAA µg/mg protein. CETP µmol/mg protein. LCAT ratio/mg protein in HDL<sub>2</sub> and ratio/mg protein X103 in HDL<sub>3</sub>.

CETP, cholesterol ester transfer protein; HDL, high-density lipoprotein; LCAT, lecithin cholesteryl acyltransferase; SAA, serum amyloid-A.

<sup>a</sup>Significance level for difference between groups, calculated using one-way ANOVA. <sup>bac</sup>Values with different superscripts indicate differences between groups (Newman–Keuls *post hoc* test).

and normal-weight children ( $P \le 0.05$  for both comparisons). SAA associated with HDL<sub>3</sub> was significantly higher in both the overweight and obese children as compared with the normal-weight children ( $P \le 0.05$  for both comparisons).

With regard to trends in the activities of CETP and LCAT within HDL<sub>283</sub>, these were also found to increase as BMI increased, although this was only statistically significant for HDL<sub>2</sub> (**Table 3**,  $P \le 0.001$  for both CETP and LCAT). In addition, the activity of HDL<sub>2</sub>-CETP was significantly higher in the obese children as compared with the overweight and normal-weight children ( $P \le 0.05$  for both comparisons), whereas the activity of HDL<sub>2</sub>-LCAT was significantly higher in both the obese and overweight children as compared with the normal-weight children ( $P \le 0.05$  for both comparisons).

## Serum and HDL<sub>283</sub> Correlations

Correlations in the complete cohort of 275 children (mean BMI, 24.6 (SD 5.18) kg/m<sup>2</sup>) showed that SAA positively correlated with HDL<sub>283</sub>-SAA (HDL<sub>2</sub>, r = 0.250,  $P \le 0.001$ ; HDL<sub>3</sub>, r = 0.604,  $P \le 0.001$ ). In addition, there were weak positive correlations between HDL<sub>2</sub>-SAA and the activity of HDL<sub>2</sub>-LCAT (r = 0.151, P = 0.022) and between HDL<sub>3</sub>-SAA and the activity of HDL<sub>3</sub>-CETP (r = 0.130, P = 0.036).

Correlations in each of the three weight groups showed that:

- 1. In the lean group (n = 92, mean BMI 19.9 (SD 2.2) kg/m<sup>2</sup>), SAA was positively correlated with HDL<sub>3</sub>-SAA ( $r = 0.410, P \le 0.001$ ).
- 2. In the overweight group (n = 92, mean BMI 24.4 (SD 1.7) kg/m<sup>2</sup>), SAA was positively correlated with HDL<sub>3</sub>-SAA ( $r = 0.680, P \le 0.001$ ).
- 3. In the obese group (n = 92, mean BMI 30.4 (SD 2.8) kg/m<sup>2</sup>), SAA was positively correlated with HDL<sub>2&3</sub>-SAA (r = 0.458,  $P \le 0.001$ ; r = 0.554,  $P \le 0.001$ , respectively). In addition, in this group, HDL<sub>2</sub>-SAA positively correlated with the activity of CETP and LCAT in HDL<sub>2</sub> (r = 0.325, P = 0.011; r = 0.356, P = 0.005, respectively). HDL<sub>3</sub>-SAA positively correlated with the activity of HDL<sub>3</sub>-CETP (r = 0.260, P = 0.014).

#### DISCUSSION

To our knowledge, this is the first report to show an association between the inflammatory molecule SAA and  $HDL_{283}$  in overweight and obese children. Furthermore, this study is also the first study to identify changes within  $HDL_{283}$  within this cohort, which augmented the atherogenic potential of these subfractions. Therefore, these results significantly add to and are in support of our previous findings, in which low-grade inflammation was associated with increased CVD risk in overweight and obese children (23).

The inflammatory molecule SAA has consistently been shown to be a sensitive marker of obesity-related inflammation in adult populations (11,27), although its role in children is largely undefined. However, one study (24) found little utility for SAA as a marker of obesity-related inflammation, which we suggest may be due to the prepubertal status of their cohort (children <10 y). This lack of effect was also described by Neuman *et al.* (25), however, the BMIs of their study groups were similar, and would therefore display a similar level of obesity-related inflammation. However, our results were in support of Gómez-Ambrosi *et al.* (26), who reported that SAA was associated with increasing BMI in children. Therefore, we suggest that SAA is a sensitive marker for detecting obesity-related inflammation in children, similar to that identified in adults (11,27).

In addition to this, we have shown that this increase in SAA augmented its association with HDL; we found that SAA positively correlated with HDL<sub>3</sub>-SAA in the overweight group (r = 0.680,  $P \le 0.001$ ) and with HDL<sub>2&3</sub>-SAA in the obese group (HDL<sub>2</sub> r = 0.458,  $P \le 0.001$ ; HDL<sub>3</sub> r = 0.554,  $P \le 0.001$ ), which supports the concept that SAA primarily associates with HDL in the circulation (15). In adults, this association renders HDL proatherogenic, with a loss or reduction of many of its functions, including its reverse cholesterol transport and anti-oxidant properties (15,28–30). Therefore, we suggest that this would also be the case in SAA-enriched HDL from overweight and obese children; in support of this concept, we have shown that the activities of HDL<sub>2</sub>-CETP and HDL<sub>2</sub>-LCAT were also influenced in these overweight and obese cohorts.

CETP is an important enzyme that is involved in the remodeling of HDL and apo B lipoproteins (16,31). However, an increase in its activity is considered proatherogenic because it can reduce HDL concentrations and influence adipocyte morphology (16,32,33). Therefore, our results suggest that these overweight and obese children were exposed to a similar CETP-mediated proatherogenic phenotype to that of overweight and obese adults, which may be negated by weight loss as documented in adults (34).

With regard to LCAT, an increase in the activity of this enzyme is seen as antiatherogenic because it is responsible for esterifying free cholesterol within pre- $\beta$ -HDL and HDL<sub>2</sub>, with HDL, being the final mature product. Therefore, LCAT is more closely associated with the former HDL particles, whereas HDL, contains minimum LCAT levels. However, this is not the case in HDL, from obese adults, where in fact LCAT is increased, which suggests that these subjects have more immature HDL, particles than are found in lean subjects (35). Therefore, our results support this finding; we also showed that LCAT's activity was increased in HDL, from both the overweight and obese children vs. their lean comparators, therefore suggesting that their HDL, was more immature. However, this finding is not consistent with the concept that LCAT's activity is normally reduced in SAA-enriched HDL (28), which suggests that obesity, independent of SAA, had influenced the activity of LCAT. However, this anomaly would require further investigation, and a mass assay for LCAT, alongside this activity assay may be more informative.

Overall, this study has provided evidence that children who are overweight or obese are exposed to an inflammatory milieu that impacts functional aspects of  $HDL_{283}$  and would potentially increase their CVD risk. Therefore, the growing epidemic of childhood obesity needs to be addressed to help prevent premature mortality in later life.

Table 4. BMI  $(kg/m^2)$  cut-off points for overweight and obese 12- and 15-y-old adolescents

Weight	Overweight		Obese		
	Males	Females	Males	Females	
12 y	21.6	22.1	26.4	27.2	
15 y	23.6	24.2	28.6	29.3	

The cut-off points for overweight and obese adults are 25 and 30 kg/m<sup>2</sup>, respectively. From ref. 36.

## **METHODS**

#### **Study Population**

Articles

The cohort utilized in this analysis has been described elsewhere (23). In brief, the Young Hearts 2000 study was a cross-sectional study that took place between 1999 and 2001. A total of 2,017 subjects, aged 12 and 15 y, were recruited from 36 nationally representative schools from across Northern Ireland. Height and weight measurements were used to assess BMI (kg/m<sup>2</sup>), and this was used to classify each subject as normal-weight, overweight, or obese on the basis of age- and sex-specific cut-off points for BMI linked to adult cut-off points (36) (**Table 4**). The prevalence of overweight and obese subjects in the complete cohort was 16.2 and 4.7%, respectively. All of the obese adolescents were selected (95 subjects). However, due to the inability to obtain blood from three of these subjects, the final subject number was reduced to 92. These were then matched according to age, sex, and smoking status with 92 overweight and 92 normal-weight children.

Ethical approval was obtained from the Research Ethics Committee of Queens University Belfast. Written informed consent was signed by the participants and the participants' parent or guardian. All investigations conformed to the principles in the Declaration of Helsinki.

#### **Blood Sampling**

Fasting peripheral venous blood was obtained by venipuncture and collected into serum or  $K_2$ -EDTA tubes. Serum and plasma were harvested following centrifugation at 1,100g for 10 min at 4 °C within 4h of collection.

#### Lipid Analyses

Serum total cholesterol, HDL cholesterol, and triglycerides were determined using enzymatic assays (Boehringer, Mannheim, Germany) on a Cobas Fara-automated analyzer (Roche Diagnostics, West Sussex, UK).

#### Isolation of HDL, and HDL,

 $\mathrm{HDL}_{283}$  were isolated from plasma by rapid ultracentrifugation according to a method previously established in our laboratory (37), which is a three-step, 6-h procedure. First, crude HDL was isolated by a 2-h rapid sedimentation method, second, this crude HDL was subfractionated into HDL<sub>283</sub> by two separate 2-h sequential rapid flotation ultracentrifugation procedures.

#### **C-Reactive Protein Concentration**

C-reactive protein has previously been reported (23), but for comparison with SAA, its results are presented here. This was measured in plasma by a latex-enhanced immune-turbidimetric assay (Wako Chemicals, Neuss, Germany) using a Cobas Fara automated analyzer.

#### **SAA** Concentration

SAA, in serum and associated with HDL<sub>283</sub>, was analyzed in duplicate by an enzyme-linked immunosorbent assay (Invitrogen, Paisley, UK) per the manufacturer's instructions. Analysis was performed on a TRITURUS-automated ELISA system (Grifols, Ghezzano Pisa, Italy).

#### **CETP and LCAT Activity**

The activity of CETP and LCAT was measured in duplicate in serum and in  $HDL_{283}$  using commercially available fluorometric assays per manufacturer's instructions (CETP and LCAT; Roar Biomedical, New York, NY).

#### **Total Protein Determination**

The protein concentration of  $HDL_{28/3}$  was determined in duplicate by a spectrophotometric assay, as previously described (38). Total protein concentration was utilized to standardize SAA, CETP, and LCAT within  $HDL_{28/3}$ .

#### **Statistical Analysis**

Statistics were completed using the statistical software package SPSS for Windows, version 17 (SPSS, Chicago, IL). Variables were assessed for normality and logarithmically transformed where required. Differences between BMI groups were assessed by a one-way ANOVA followed by the Newman–Keuls multiple range comparison test when the data were normally distributed, or by the Kruskal–Wallis test when the data were not normally distributed, even after logarithmic transformation. Differences between the BMI groups in categorical variables were assessed using  $\chi^2$  test. Correlations between HDL<sub>283</sub>-SAA levels and the activities of LCAT and CETP within HDL<sub>284</sub> were assessed using Pearson's correlation coefficient. All variables were summarized as mean and SD when normally distributed, as geometric mean (interquartile range) when normally distributed after logarithmic transformation, and as median (interquartile range) when logarithmic transformation did not normalize the distribution. The significance level was set as P < 0.05.

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