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CLINICAL SCIENCES

# Increased Cholesteryl Ester Transfer Protein Activity in Impaired Glucose Tolerance: Relationship to High Density Lipoprotein Metabolism

Jens Pietzsch, Katja Fuecker<sup>1</sup>

Pathological Biochemistry Group, Department of Positron Emission Tomography, Institute of Bioinorganic and Radiopharmaceutical Chemistry, Research Center Rossendorf, Dresden; and <sup>1</sup>Lipoprotein Laboratory, Department of Internal Medicine III, Carl Gustav Carus Medical Faculty, Dresden Technical University, Dresden, Germany

**Aim.** To investigate the mechanisms and dynamics of cholesteryl ester (neutral lipid) transfer protein activity in subjects with impaired glucose tolerance.

**Methods.** Eighty six Caucasian subjects were recruited by advertisement from the local population between January 1998 and December 1999. The activity of cholesteryl ester transfer protein in 44 non-obese subjects with impaired glucose tolerance (plasma triglycerides  $1.56 \pm 0.64 \text{ mmol/L}$ ; high density lipoprotein (HDL) cholesterol  $0.96 \pm 0.25 \text{ mmol/L}$ ; and plasma insulin  $78 \pm 8 \text{ pmol/L}$ ) and in 42 normoglycemic controls (plasma triglycerides  $0.88 \pm 0.41 \text{ mmol/L}$ ; HDL cholesterol  $1.48 \pm 0.29 \text{ mmol/L}$ ; and plasma insulin  $38 \pm 14 \text{ pmol/L}$ ) was measured with a new fluorometric assay.

**Results.** Cholesteryl ester transfer protein activity was increased in subjects with impaired glucose tolerance by 47% ( $39.5 \pm 7.8 \text{ vs} 26.8 \pm 6.8 \text{ nmol/mL} \times h-1$ ; t-test, p < 0.05). Linear regression analysis showed that cholesteryl ester transfer protein activity in subjects with impaired glucose tolerance significantly correlated with the following parameters: plasma triglycerides (r=0.614, p<0.05), HDL-triglycerides (r=0.595, p<0.05), percentage of HDL-triglyceride (r=0.667, p<0.05), percentage of HDL cholesterol ester (r=-0.751, p<0.01), percentage of HDL phospholipid (r=0.648, p<0.05), 2-h-insulin (r=0.668, p<0.05), and 2-h-proinsulin (r=0.658, p<0.01). In a subgroup of 13 subjects with impaired glucose tolerance, cholesteryl ester transfer protein activity correlated with HDL apoA-I fractional catabolic rate (r=0.701, p<0.01). In normoglycemic subjects, significant correlations were found only between cholesteryl ester transfer protein activity and percentage of HDL-triglycerides (r=0.541, p<0.05), percentage of HDL cholesterol ester (r=0.642, p<0.05), and HDL apoA-I fractional catabolic rate (r=0.639, p<0.01), 2-h-proinsulin (r=0.642, p<0.05), and HDL apoA-I fractional catabolic rate (n=10, r=0.587, p<0.05).

**Conclusion.** Cholesteryl ester transfer is important for HDL composition and HDL catabolism both in normoglycemic subjects and subjects with impaired glucose tolerance. Under insulin resistant conditions, increased cholesteryl ester transfer protein activity modulates HDL metabolism more drastically than in normoglycemic conditions. This modulation may be explained by increased availability of triglyceride-rich lipoproteins for neutral lipid exchange in subjects with impaired glucose tolerance.

*Key words:* apolipoproteins; carrier proteins; glucose intolerance; hyperlipidemia; insulin resistance; lipoproteins, HDL; metabolism

Impaired glucose tolerance is characterized by insulin resistance and hyperinsulinemia and is associated with dyslipidemia, a condition characterized by increased plasma triglycerides, decreased high density lipoprotein (HDL) levels, and a preponderance of smaller, more dense low density lipoprotein (LDL) particles (1-3). Our previous studies focused on the regulation of HDL metabolism in subjects with impaired glucose tolerance (4,5). By investigating in vivo kinetics of HDL apolipoproteins (apo) A-I and A-II using a stable isotope approach, we demonstrated that increased catabolism of HDL apoA-I in subjects with impaired glucose tolerance was related to the increased triglyceride content of HDL particles and accounted for lower HDL levels (4,5). Furthermore, we showed that the impairment of HDL catabolism was evident in subjects with impaired glucose tolerance even when triglyceride concentrations were in the upper normal range (plasma triglyceride levels between 1.5 and 2.3 mmol/L). In this context, increased fasting cholesteryl ester transfer protein activity seemed to be directly associated with HDL apoA-I catabolism and plasma HDL triglycerides (4,5). Cholesteryl ester transfer protein facilitates the exchange of neutral lipids among plasma lipoproteins, which results in net transfer of cholesteryl esters from HDL to triglyceride-rich lipoproteins, e.g., very low density lipoproteins (VLDL), in exchange for triglycerides. This redistribution process should have consequences for HDL composition and consequently for the HDL metabolism (6,7). Indeed, increased cholesteryl ester transfer protein activity in humans was reported in several metabolic disorders with low HDL cholesterol (9-13), such as type 1 diabetes (8), overt type 2 diabetes (9), obesity (10), nephrotic syndrome (11), acromegaly (12), and hypercholesterolemia (13).

However, little data are available regarding direct association between cholesteryl ester transfer protein activity and the metabolic decompensation of HDL metabolism under prediabetic conditions, e.g., in subjects with impaired glucose tolerance. Our study was designed to explore further the mechanisms and dynamics of cholesteryl ester transfer and their specific link with low plasma HDL concentrations and an altered HDL composition under insulin resistant/hyperinsulinemic conditions. We aimed at finding associations between cholesteryl ester transfer protein activity and parameters of carbohydrate and lipid metabolism in a carefully selected group of normotriglyceridemic or mildly hypertriglyceridemic patients with impaired glucose tolerance.

## **Subjects and Methods**

#### Subjects

Eighty six Caucasian subjects were recruited by advertisement from the local population between January 1998 and December 1999. All subjects consented to participate in the study, which was approved by the local Medical Ethics Committee. The study group consisted of 44 non-obese subjects (24 women and 20 men), aged 19 to 63 years (median, 46.5 years), with impaired glucose tolerance and normal or moderately increased plasma triglyceride (TG) concentration. The control group included 42 healthy, normolipidemic subjects (22 women and 20 men), aged 19 to 63 years (median, 40.5 years), with normal glucose tolerance (Table 1). There was no evidence of thyroid, liver, cardiovascular, hematologic, or renal abnormalities in any of the subjects, and none were taking medications. All subjects were non-smokers. The diagnosis of impaired glucose tolerance was based on two consecutive oral glucose tolerance tests (oGTT; 75 g oral glucose load) according to the World Health Organization (WHO) guidelines and criteria (two 2-h post-glucose values between 7.8 and 11.1 mmol/L) (14).

#### Experimental Protocol

Blood (10-20 mL) was collected into vacuum tubes containing EDTA at a final concentration of 0.1%. Plasma was recovered after centrifugation at 4 °C. Sodium azide (0.01%) and aprotinin (200 KIU/mL) were added immediately. Plasma samples were stored at 4 °C and used to determine plasma glucose, plasma lipids, and concentrations of apoB-100, apoA-I, apoA-II, and apoE. In addition, very low density lipoproteins (VLDL; d < 1.006 kg/L), LDL (1.006 < d < 1.063 kg/L), and HDL (1.063 < d < 1.210 kg/L) were isolated from 0.5 mL plasma by sequential very fast ultracentrifugation by means of an OptimaTM TLX table top ultracentrifuge with fixed angle rotor TLA-120.2 at 120,000 rev/min (Beckman instruments, Palo Alto, CA, USA) (15,16). HDL prepared by this very fast ultracentrifugation technique is free of albumin and other protein contaminations, which allows correct compositional analysis (15,16). Lipid and protein constituents in plasma and lipoprotein fractions were measured as previously described (15). For the determination of insulin, proinsulin, and cholesteryl ester transfer protein activity, plasma samples (VLDL-plus LDL-free plasma for measurement of cholesteryl ester transfer protein activity) were shock-frozen in liquid nitrogen and

Table 1. Characteristics and biochemical parameters (mean ± SD) of 42 subjects with impaired glucose tolerance (IGT) and 44 subjects with normal glucose tolerance (NGT)<sup>3</sup>

subjects with normal glucose tolerance	(NGI)*		
Subject characteristics	NGT group	IGT group	р
Sex:			
women/men	22/20	24/20	
Age (years)	$40 \pm 15$	$48 \pm 12$	NS <sup>+</sup>
Body mass index (kg/m <sup>2</sup> )	$24.2 \pm 2.4$	$25.8 \pm 2.4$	NS
Fasting plasma glucose (mmol/L)	$5.35 \pm 0.48$	$6.04 \pm 0.74$	< 0.001
2-h-post glucose (mmol/L)	$5.58 \pm 1.16$	$8.86 \pm 0.86$	< 0.001
Fasting insulin (pmol/L)	$38 \pm 14$	$78 \pm 8$	< 0.001
2-h-post insulin (pmol/L)	$259 \pm 111$	$610 \pm 196$	< 0.001
Fasting proinsulin (pmol/L) <sup>‡</sup>	$1.85 \pm 0.69$	$2.83 \pm 1.36$	< 0.001
2-h-post proinsulin (pmol/L) <sup>‡</sup>	$10.1 \pm 5.7$	$20.5 \pm 12.8$	< 0.001
Fasting free fatty acids (mmol/L)	$0.41 \pm 0.20$	$0.61 \pm 0.17$	0.008
HbA1c (%) <sup>‡</sup>	$5.2 \pm 0.4$	$5.8 \pm 1.2$	NS
Triglycerides (mmol/L) <sup>‡</sup>	$0.88 \pm 0.41$	$1.56 \pm 0.64$	0.020
VLDL triglycerides (mmol/L) <sup>‡</sup>	$0.52 \pm 0.11$	$0.98 \pm 0.34$	0.020
Total cholesterol (mmol/L)	$4.92 \pm 0.66$	$5.18 \pm 0.76$	NS
HDL cholesterol (mmol/L)	$1.48 \pm 0.29$	$0.96 \pm 0.25$	0.015
ApoA-I (g/L)	$1.46 \pm 0.09$	$1.00 \pm 0.29$	0.030
ApoA-II (g/L)	$0.24 \pm 0.04$	$0.22 \pm 0.03$	NS
ApoB-100 (g/L)	$0.95 \pm 0.36$	$1.48 \pm 0.29$	0.029
ApoE (g/L)	$0.056 \pm 0.011$	$0.078 \pm 0.021$	0.045
CETP activity (nmol/mL h <sup>-1</sup> )	$26.8 \pm 6.8$	$39.5 \pm 7.8$	0.017
HDL composition:			
HDL triglycerides (mmol/L) <sup>‡</sup> (%) <sup>§</sup>	$0.12 \pm 0.06$ (3.20 ± 1.10)	$0.21 \pm 0.08$ (4.90 ± 1.91)	0.020 (0.020)
HDL free cholesterol (mmol/L) (%)	$0.21 \pm 0.04$ (2.48 ± 0.62)	$0.23 \pm 0.04$ (2.72 ± 0.15)	NS (NS)
HDL cholesteryl esters (mmol/L) (%)	$1.05 \pm 0.07$ (13.87 ± 1.81)	$0.81 \pm 0.08$ (10.34 ± 1.59)	0.030 (0.001)
HDL phospholipids (mmol/L) (%)	$1.20 \pm 0.36$ (27.30 ± 5.80)	$1.46 \pm 0.24$ (34.70 ± 2.61)	0.015 (0.008)
HDL apoA-I (g/L) (%)	$1.34 \pm 0.08$ (43.20 ± 2.36)	$0.94 \pm 0.11$ (30.54 ± 4.42)	< 0.001 (< 0.001)
HDL apoA-II (g/L) (%)	$0.24 \pm 0.02$ (7.20 ± 0.70)	$0.22 \pm 0.02$ (6.91 ± 0.62)	NS (NS)
HDL apoE (g/L) (%)	$0.011 \pm 0.005 \ (0.30 \pm 0.10)$	$0.016 \pm 0.011 \ (0.50 \pm 0.20)$	NS (NS)
HDL residual protein (g/L) (%)	$0.09 \pm 0.06$ (2.40 ± 1.50)	$0.31 \pm 0.09  (9.30 \pm 2.70)$	0.007 (<0.001)

\*The values of glucose and insulin were based on the mean of two consecutive oral glucose tolerance test (oGTT); the diagnosis of impaired glucose tolerance was based on two 2-h-post glucose values between 7.8 and 11.1 mmol/L, according to the World Health Organization guidelines. Abbreviatons: VLDL – very low density lipoproteins; HDL – high density lipoproteins; CETP – cholesteryl ester transfer protein. <sup>†</sup>NS – not significant (p>0.05).

<sup>‡</sup>Log-transformed value.

Mean lipoprotein mass composition: the mass percentage of each lipid and protein constituent from the total HDL mass (15).

HDL residual protein consists of other HDL protein constituents, e.g., apolipoproteins C-I to C-III.

stored in cryovials at -80 °C. Plasma insulin and proinsulin were determined by specific enzyme immunoassay (BioSource, Ratingen, Germany; within-day precision < 3.8%, between-day precision <6.5%; no cross reaction with human proinsulin and insulin, respectively). Cholesteryl ester transfer protein activity was measured as published elsewhere, with some modifications (4). In brief, HDL-like particles containing 0.1  $\mu$ mol/mL of the fluorescent cholesteryl ester analog N-(7-Nitrobenz-2-oxa-1,3-diazole)-23,24-dinor-5-cholen-22-amine-3ß-ol (NBD-CE, WAK Medical, Bad Soden, Germany) were used as donor particles and VLDL isolated from pooled normal fasting plasma were used as acceptor particles. The fluorescent cholesteryl linoleate is present in a self-guenched state when contained within the core of the donor particle. The cholesteryl ester transfer is determined by an increase in fluorescence intensity (excitation at 465 nm, and emission at 535 nm) as the cholesteryl linoleate is removed from the donor particle and transferred to the acceptor particle in the presence of a cholesteryl ester transfer protein-containing sample. The reaction was linear and the amount of cholesteryl ester transferred was derived from a standard curve by use of different concentrations of NBD-CE. The within-day and between-day precision of the assay was <6.0% and <7.8%, respectively. The use of VLDL- and LDL-free plasma samples substantially eliminates the influence of the endogenous VLDL and LDL pools on the cholesteryl ester transfer activity measurement ex vivo. Cholesteryl ester transfer protein activity is expressed as nmol/mL×h<sup>-1</sup> (4). Accordingly, in this assay, cholesteryl ester transfer protein activity was only influenced by the cholesteryl ester transfer protein concentration, therefore reflecting likely the cholesteryl ester transfer protein mass

#### Statistical Analysis

We used SPSS 11.0.1 software package (SPSS Inc., Chicago, IL, USA) for all statistical analyses. Descriptive data were expressed as arithmetic means  $\pm$  standard deviations (SD). The distribution of values was assessed by Kolmogorov-Smirnov test for homogeneity of variances and parameters were transformed logarithmically if necessary. Metabolic variables of two study groups were compared by t-test. For correlation analyses within the total study population, Spearman's rank correlation coefficient (r) was used. Additionally, we calculated partial correlation coefficients within the study groups using sex, age, and body mass index as covariates.

# **Results**

#### **Biochemical Characteristics**

Subjects with impaired glucose tolerance and their controls were well-matched for age, sex, and body mass index (Table 1). The concentration of glycosylated hemoglobin (HbA1c) was below 6% in both groups. Fasting values for plasma glucose, free fatty acids, insulin, and proinsulin, and 2-hour postchallenge values (after 75 g oral glucose load) for glucose, free fatty acids, insulin, and proinsulin were significantly increased in subjects with impaired glucose tolerance (Table 1). These values were considered as highly indicative for an insulin resistant/hyperinsulinemic state of the subjects with impaired glucose tolerance. In subjects with impaired glucose tolerance, plasma triglycerides were higher by 77% (p < 0.05), apoB-100 concentrations by 56% (p<0.05), and apoE concentrations by 39% (p<0.05), compared with control group. However, all these parameters were still in the upper normal range. VLDL triglycerides were higher by 88% (p < 0.05) in subjects with impaired glucose tolerance. The plasma levels of total cholesterol showed no differences between subjects with impaired glucose tolerance and controls.

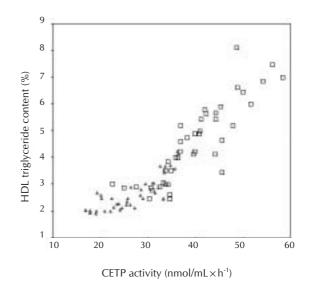
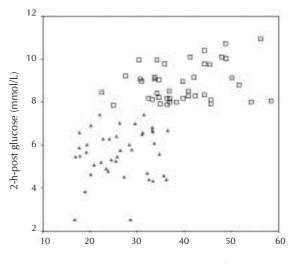


Figure 1. Relationship between cholesteryl ester transfer protein (CETP) activity and high density lipoprotein (HDL) triglyceride (%) in 86 subjects with impaired glucose tolerance (squares) or normal glucose tolerance (triangles).



CETP activity (nmol/mL×h<sup>-1</sup>)

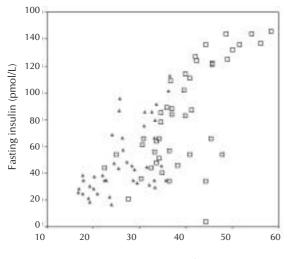
**Figure 2.** Relationship between cholesteryl ester transfer protein (CETP) activity and 2-h-post glucose in 86 subjects with impaired glucose tolerance (squares) or normal glucose tolerance (triangles).

## Cholesteryl Ester Transfer Protein Activity

The rate of cholesteryl linoleate transfer from HDL particles to VLDL was significantly higher (47%) in the impaired glucose tolerance group than in the control group.

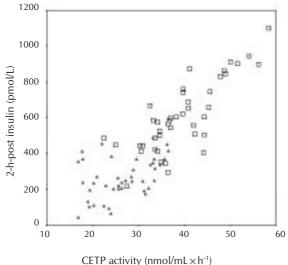
## HDL Composition

Plasma apoA-I and HDL cholesterol concentrations were 32% and 35% lower, respectively, in subjects with impaired glucose tolerance (Table 1). Among HDL constituents, concentrations of HDL apoA-I and HDL cholesteryl esters were significantly lower, whereas concentrations of HDL triglycerides and HDL phospholipids were significantly higher in



CETP activity  $(nmol/mL \times h^{-1})$ 

**Figure 3.** Relationship between cholesteryl ester transfer protein (CETP) activity and fasting insulin in 86 subjects with impaired glucose tolerance (squares) or normal glucose tolerance (triangles).



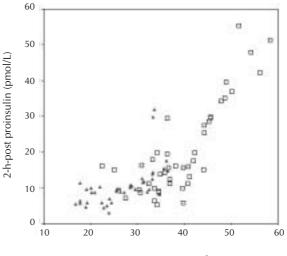
CETP activity ( $nmol/mL \times n^{-1}$ )

**Figure 4.** Relationship between cholesteryl ester transfer protein (CETP) activity and 2-h-post insulin in 86 subjects with impaired glucose tolerance (squares) or normal glucose tolerance (triangles).

subjects with impaired glucose tolerance. The concentrations of free cholesterol, apoA-II, and apoE in HDL did not differ between the two groups. Furthermore, HDL particle composition, expressed as the mass percentage of each lipid and protein constituent from the total HDL mass, was substantially altered in subjects with impaired glucose tolerance (Table 1). This compositional misbalance consequently led to HDL particles that were significantly enriched with triglycerides and phospholipids, but depleted in their cholesterol ester and apoA-I content.

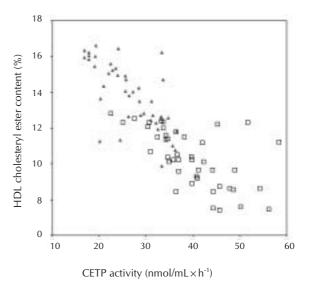
# **Correlation Analysis**

In the total study population of 86 participants, cholesteryl ester transfer protein activity correlated di-



CETP activity (nmol/mL  $\times$  h<sup>-1</sup>)

**Figure 5.** Relationship between cholesteryl ester transfer protein (CETP) activity and 2-h-post proinsulin in 86 subjects with impaired glucose tolerance (squares) or normal glucose tolerance (triangles).



**Figure 6.** Relationship between cholesteryl ester transfer protein (CETP) activity and HDL cholesteryl ester (%) in 86 subjects with impaired glucose tolerance (squares) or normal glucose tolerance (triangles).

rectly with plasma triglycerides (r=0.675; p<0.05), HDL triglycerides (r=0.818; p<0.01), HDL triglyceride content (r=0.927; p<0.001; Fig. 1), HDL phospholipid content (r=0.655; p<0.05), as well as with 2-h-post glucose (r=0.651; p<0.001; Fig. 2), fasting insulin (r=0.724; p<0.001; Fig. 3), 2-h-post insulin (r=0.802; p<0.001; Fig. 4), and 2-h-post proinsulin (r=0.651; p<0.001; Fig. 5). An inverse association was found between the cholesteryl ester transfer protein activity and HDL cholesteryl ester content (r=-0.688; p<0.05) and HDL cholesteryl ester content (r=-0.849; p<0.001; Fig. 6). To compensate for possible effects of age, body mass index, and sex, partial correlation coefficients were calculated for both **Table 2.** Partial correlation analysis in subjects with impaired glucose tolerance (IGT) and those with normal glucose tolerance (NGT), taking into account age, sex, and body mass index

Cholesteryl ester	Partial correlation coefficients					
transfer protein	NGT		IGT			
(CETP) activity versus:	(n = 42)	р	(n = 44)	р		
Triglycerides	_		0.614	0.038		
HDL triglycerides	-		0.595	0.012		
HDL triglyceride (%)*	0.541	0.024	0.667	0.014		
HDL cholesteryl ester (%)*	-0.639	0.001	-0.751	0.009		
HDL phospholipids (%)*			0.648	0.023		
2-h-post glucose	-		0.601	0.048		
Fasting insulin	-		0.681	0.039		
2-h-post insulin	-		0.668	0.033		
2-h-post proinsulin	0.587	0.045	0.658	0.004		
*Lipoprotein mass composition: the mass percentage of each lipid and protein constituent from the total HDL mass (%) (15).						

groups separately, with these parameters used as covariates. In subjects with impaired glucose tolerance, differences remained statistically significant for correlation between cholesteryl ester transfer protein activity and plasma triglycerides, HDL triglycerides, HDL triglyceride content, HDL cholesteryl ester content, HDL phospholipid content, 2-h-post glucose, fasting insulin, 2-h-post insulin, and 2-h-post proinsulin (Table 2). In contrast, only the HDL triglyceride and cholesteryl ester content, as well as the 2-h-post proinsulin levels remained associated with cholesteryl ester transfer protein activity in controls (Table 2).

## Discussion

This study showed that fasting cholesteryl ester transfer protein activity, measured by a new fluorescent spectrophotometric method, was significantly increased in normotriglyceridemic or mildly hypertriglyceridemic subjects with impaired glucose tolerance. The assay measured cholesteryl ester transfer protein activity as the specific cholesteryl linoleate transfer from HDL (donor) particles to VLDL (acceptor) particles. By using this method and VLDL- and LDL-depleted plasma as the cholesteryl ester transfer protein source, we avoided the influence of endogeneous lipoproteins. Thus, the cholesteryl ester transfer protein activity reflected the cholesteryl ester transfer protein mass. Observations regarding cholesteryl ester transfer protein activity may differ depending on the assays used (6). However, the increased cholesteryl ester transfer protein activity we showed is in agreement with our previous data and partly with the results of several recent studies in which a similar assay system to measure cholesteryl ester transfer protein activity was used (4, 12, 17, 18).

The activity of cholesteryl ester transfer protein was directly associated with parameters highly indicative for an insulin resistant/hyperinsulinemic condition (19-21). Among these, fasting insulin, 2-h-post insulin, and 2-h-post proinsulin showed the highest statistical power. On the other hand, no association between free fatty acids and cholesteryl ester transfer protein activity could be found. This is of special interest, because there is experimental and clinical evidence that free fatty acids may exert direct effects on cholesteryl ester transfer protein concentrations and also modulate cholesteryl ester transfer rates through alterations in the concentration and composition of plasma lipoprotein substrates (6,7,11,22).

Furthermore, we found cholesteryl ester transfer protein activity to be directly associated with parameters of HDL particle composition, particularly with the HDL triglyceride and cholesteryl ester content. In this context, the use of very fast ultracentrifugation technique was essential for the present analysis, because it provided albumin-free HDL particles, which allowed the exact calculation of HDL particle composition. In addition, there was a strong relationship between HDL phospholipid content and cholesteryl ester transfer protein activity. This seems to reflect the specific phospholipid transfer activity of cholesteryl ester transfer protein (6).

When the study groups were analyzed separately, most of the associations found persisted in subjects with impaired glucose tolerance, taking into account sex, age, and body mass index. In contrast, only the HDL triglyceride and cholesteryl ester content, as well as the 2-h-post proinsulin concentrations remained associated with cholesteryl ester transfer protein activity in controls. There was no correlation between VLDL triglycerides and cholesteryl ester transfer protein activity in either group, confirming the results of other studies (9). The correlations discussed above suggest that cholesteryl ester transfer protein is likely to be an important modulator of HDL particle composition under both normal and insulin resistant conditions, even when triglyceride concentrations are normal or mildly increased.

Data in this study substantially support our previous findings on increased in vivo catabolism of HDL apoA-I in subjects with impaired glucose tolerance (4). Those former kinetic data suggested that conformational changes in the HDL particles due to their increased triglyceride and phospholipid content as well as decreased cholesteryl ester content favored their direct catabolism. Hypothetically, the enrichment with triglycerides in particular could make the HDL a favored substrate of the hepatic lipase (7,23). Hepatic lipase hydrolyzes HDL triglycerides and subsequently the HDL particle size decreases. ApoA-I is weakly bound to these small, cholesteryl ester-depleted HDL particles and, consequently, the apoA-I pool is easily dissociable and more rapidly catabolized. This is in accordance with the current hypotheses on the metabolic reason of the preponderance of small, dense LDL under diabetic and prediabetic conditions (24-26). Ambrosch and colleagues (18) found increased cholesteryl ester transfer protein activity in insulin resistant persons who were identified among young, clinically healthy subjects by the clamp technique. They reported a strong association of increased cholesteryl ester transfer protein activity with decreasing LDL particle diameter in these subjects (18).

The association between cholesteryl ester transfer protein and phospholipid content in HDL is suspected to be an indicator of the known intrinsic phospholipid transfer activity of cholesteryl ester transfer protein (7). However, we did not investigate the impact of another lipid transfer protein – the phospholipid transfer protein – on the metabolism of HDL. There is experimental and clinical evidence that phospholipid transfer protein activity is altered in insulin-resistant states, with further consequences for HDL composition (27).

In conclusion, our observations suggest that neutral lipid transfer mediated by cholesteryl ester transfer protein is crucial for the modification of HDL composition in subjects with impaired glucose tolerance. Under insulin resistant/hyperinsulinemic conditions, increased cholesteryl ester transfer protein activity modulates HDL metabolism more drastically than in normoglycemic conditions. This modulation may be explained by the increased availability of triglyceride-rich lipoproteins for neutral lipid exchange in subjects with impaired glucose tolerance. As a consequence, HDL cholesterol is decreased in impaired glucose tolerance conditions. Low HDL is an integral part of the atherogenic lipoprotein phenotype in diabetic and prediabetic states (1,4). However, the mechanisms by which cholesteryl ester transfer protein is regulated in its activity or synthesis, particularly in insulin-resistant subjects, have to be further elucidated. We suggest that insulin has a direct influence on the cholesteryl ester transfer protein in vivo, but under insulin resistant conditions this specific insulin action can be diminished. This would be in agreement with clinical and experimental evidence that the higher the plasma insulin concentration (more insulin resistant), the greater the fractional catabolic rate of HDL and the lower the plasma HDL cholesterol concentrations in diabetic and prediabetic states (4,28, 29). Additional work, including experimental and clinical use of recently developed cholesteryl ester transfer protein inhibitors (30), is needed to further understand the consequences of increased cholesteryl ester transfer for the development of the atherogenic lipoprotein phenotype in diabetic and prediabetic states.

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## Correspondence to:

Jens Pietzsch Pathological Biochemistry Group Department Positron Emission Tomography Institute of Bioinorganic and Radiopharmaceutical Chemistry Rossendorf Research Center P.O. Box 51 01 19 D-01314 Dresden Germany *j.pietzsch@fz-rossendorf.de*