

Skeletal muscle lipase content and activity in obesity and type 2 diabetes.

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Skeletal Muscle Lipase Content and Activity in Obesity and Type 2 Diabetes

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Context: The obese insulin-resistant state is characterized by elevated lipid storage in skeletal muscle tissue.

Objective: We tested whether differences in muscle triacylglycerol (TAG) and diacylglycerol (DAG) lipase content and activity are associated with incomplete *in vivo* lipolysis and lipid accumulation.

Design and Patients: Two case-control studies were conducted on skeletal muscle biopsies from lean ($n = 13$) and obese ($n = 10$) men (study 1) and from 11 nonobese type 2 diabetic (T2D), obese T2D, and healthy normoglycemic men (study 2).

Main Outcome Measures: Skeletal muscle lipase protein content and activity and muscle lipid content (TAG and DAG) were determined.

Results: Skeletal muscle hormone-sensitive lipase protein content was lower (0.39 ± 0.07 vs. 1.00 ± 0.19 arbitrary units; $P = 0.004$) and adipose triglyceride lipase protein content was higher in obese men compared with lean controls (2.17 ± 0.40 vs. 0.42 ± 0.23 arbitrary units; $P = 0.008$). This apparent difference in lipase content was accompanied by a 60% lower ratio of DAG to TAG hydrolase activity in the obese men (11.4 ± 2.3 vs. 26.5 ± 7.3 nmol/h · mg; $P = 0.045$), implying incomplete lipolysis. Lower hormone-sensitive lipase and higher adipose triglyceride lipase content was confined to obesity *per se*, because it was observed solely in obese T2D men but not in healthy normoglycemic controls and nonobese T2D men. Muscle total DAG content was not higher in obese men but was even lower (6.2 ± 0.7 vs. 9.4 ± 0.9 $\mu\text{mol/mg}$ dry weight; $P = 0.017$). TAG content did not differ between groups (84.7 ± 18.9 vs. 70.4 ± 12.4 $\mu\text{mol/mg}$ dry weight; $P = 0.543$).

Conclusions: Our data do not support an important role of total muscle DAG content in the development of insulin resistance in obese men. (*J Clin Endocrinol Metab* 95: 5449–5453, 2010)

The obese insulin-resistant state is characterized by disturbed lipid metabolism including increased circulating fatty acid levels (*i.e.* lipid overflow), resulting in an elevated fatty acid supply to peripheral tissues like skeletal

muscle (1). As a result, ectopic fat deposition may occur in skeletal muscle, causing lipotoxicity. There is a strong link between increased intramuscular triacylglycerol (TAG) storage and skeletal muscle insulin resistance in obesity and type 2

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Abbreviations: ATGL, Adipose triglyceride lipase; AU, arbitrary units; BMI, body mass index; CGI-58, comparative gene interaction-58; DAG, diacylglycerol; DAGH, DAG hydrolase; DGAT-1, DAG acyltransferase-1; DGK δ , DAG kinase δ ; HbA1c, glycosylated hemoglobin; HOMA_{IR}, homeostasis model assessment for insulin resistance; HSL, hormone-sensitive lipase; MUFA, monounsaturated fatty acid; PLIN1, perilipin 1; PUFA, polyunsaturated fatty acid; SFA, saturated fatty acid; TAG, triacylglycerol; TAGH, TAG hydrolase; T2D, type 2 diabetic.

diabetes (2). However, it has become increasingly apparent that im accumulation of less abundant lipid metabolites, like diacylglycerol (DAG), rather than TAG *per se* may be causally related to skeletal muscle insulin resistance (3). Previously, we showed that under fasting conditions, total glycerol release was reduced across the forearm muscle of obese *vs.* lean men, indicating that obesity may be accompanied by disturbances in skeletal muscle lipolysis (4).

The first step in skeletal muscle TAG hydrolysis is catalyzed by adipose triglyceride lipase (ATGL), which is activated by its coactivator comparative gene interaction-58 (CGI-58), as reviewed previously (5). ATGL provides DAG substrate for the subsequent action of hormone-sensitive lipase (HSL), resulting in the conversion of DAG into monoacylglycerol. In contrast to HSL, ATGL exhibits high substrate specificity for the hydrolysis of TAG (6). HSL has also TAG hydrolase (TAGH) activity, although lower than its DAG hydrolase (DAGH) activity (7). ATGL-deficient animals display improved glucose tolerance and enhanced insulin sensitivity despite TAG accumulation in multiple tissues (*e.g.* adipose tissue and skeletal muscle) (8), suggesting TAG storage is rather protective for the development of insulin resistance. In contrast, HSL-deficient mice show increased DAG storage in adipose tissue and skeletal muscle and signs of impaired insulin sensitivity (9).

In the present study, we hypothesized that differences in lipase expression, and TAGH relative to DAGH activity, might be associated with incomplete lipolysis and storage of lipid metabolites in skeletal muscle tissue of obese as opposed to nonobese, insulin-resistant men. Therefore, lipase protein content and activity and lipid content (TAG and DAG) were measured in muscle biopsies obtained from male obese and nonobese type 2 diabetic (T2D) subjects as well as from a lean normoglycemic control group.

Subjects and Methods

Study 1 is a follow-up on a recent stable isotope study by our group in which we showed a blunted glycerol release across the forearm muscle of obese insulin-resistant men, suggesting impaired lipolysis (4). Biopsies from these subjects were used to determine muscle lipase protein content, activity, and lipid content (TAG and DAG). In study 2, we wanted to elucidate whether possible differences in lipase content are related to insulin sensitivity or obesity *per se*. Therefore, we took muscle biopsies from male obese T2D [age 60.6 ± 1.1 yr, body mass index (BMI) 34.9 ± 0.8 kg/m², homeostasis model assessment for insulin resistance (HOMA_{IR}) 7.8 ± 0.8 , glycosylated hemoglobin (HbA_{1c}) $7.9 \pm 0.5\%$] and nonobese T2D (age 61.0 ± 2.0 yr, BMI 26.1 ± 0.4 kg/m², HOMA_{IR} 6.2 ± 1.1 , HbA_{1c} $7.4 \pm 0.2\%$) subjects and a normoglycemic healthy control group (age 56.9 ± 1.2 yr, BMI 25.8 ± 0.9 kg/m², HOMA_{IR} 2.3 ± 0.4 , HbA_{1c} $5.4 \pm$

0.2%), 11 each. All subjects were matched for age, and the nonobese T2D men were matched for BMI with the normoglycemic control group. Biopsies were used to determine lipase content using Western blot analysis. The Medical Ethical Review Committee of Maastricht University and Virga Jessa Hospital Hasselt approved the study protocol, and clinical investigations were performed according to the declaration of Helsinki.

Skeletal muscle biopsies were taken from the vastus lateralis muscle under local anesthesia of the skin and fascia (xylocaine; AstraZeneca, Zoetermeer, The Netherlands) using the needle biopsy technique (10). Muscle biopsies were immediately frozen in liquid nitrogen and stored at -80 C until further analysis.

Muscle tissue was freeze-dried and dissected free of all visible adipose tissue, connective tissue, and blood under a microscope and was subsequently homogenized (1:80 by volume), as described previously (4). Solubilized muscle tissue protein were separated on a 12% SDS-PAGE, transferred to a nitrocellulose membrane (Hybond-ECL; Amersham Biosciences, Freiburg, Germany), and incubated with primary antibodies rabbit anti-ATGL (catalog no. 2138; Cell Signaling Technology, Beverly, MA) and rabbit anti-CGI-58 (catalog no. NB110-41576; Novus Biologicals, Littleton, CO), and the rabbit anti-HSL antibody was a kind gift from Prof. Dr. Cecilia Holm, Lund University, Sweden, as previously described (4). In a pilot experiment, an adipose tissue sample was loaded as positive control showing a clear band at the expected height for perilipin 1 (PLIN1) (~ 58 kDa); no band was detected in our muscle samples. This indicates that the guinea pig anti-PLIN1 antibody (catalog no. GP33; Progen Biotechnik GmbH, Heidelberg, Germany) is sensitive and useful for assessing potential adipocyte contamination in our muscle samples.

TAGH and DAGH activities were measured on muscle tissue homogenates, as described previously (11). Total lipids were extracted from freeze-dried muscle tissue using the method of Folch *et al.* (12). TAG and DAG fatty acid methyl esters were separated by capillary gas liquid chromatography using a $50\text{-m} \times 0.25\text{-mm}$ CP-sil 88 silica column (Varian, Middelburg, The Netherlands) with helium as carrier gas at a flow of 130 kPa. The column oven was maintained at 165 C for 10 min and increased at a rate of 5 C/min to 190 C. This temperature was maintained for 15 min. The temperature was increased to 230 C with a flow rate of 2 C/min for DAG and 5 C/min for TAG analysis. For DAG, this temperature was maintained for 22 min, whereas for TAG, this temperature was maintained for 7 min. All statistical analyses were performed using SPSS for Macintosh (version 16.0; SPSS Inc., Chicago, IL).

Results

Study 1

Muscle HSL protein content was significantly lower in obese compared with lean individuals [0.39 ± 0.07 *vs.* 1.00 ± 0.19 arbitrary units (AU); ANOVA group-effect $P = 0.004$, Fig. 1A]. In contrast to a lower HSL content, muscle ATGL protein content was significantly higher in obese individuals (2.17 ± 0.40 *vs.* 0.42 ± 0.23 AU; ANOVA group-effect $P = 0.008$; Fig. 1A). CGI-58 protein expression was comparable between groups (0.85 ± 0.19 *vs.* 1.09 ± 0.27 AU; $P = 0.484$). PLIN was not detectable

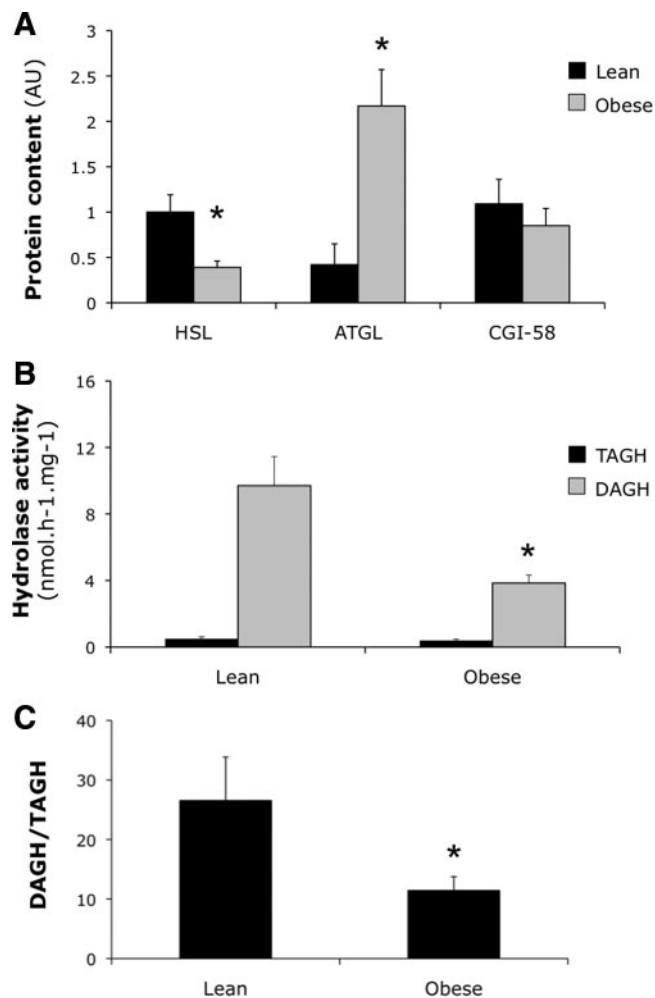


FIG. 1. Skeletal muscle lipase protein content and activity in lean vs. obese insulin-resistant men (study 1). A, HSL (ANOVA group-effect $P = 0.004$), ATGL (ANOVA group-effect $P = 0.008$), and CGI-58 protein content in lean vs. obese men participating in study 1. α -Actin was used to correct for differences in protein loading. B, TAGH and DAGH activity in skeletal muscle of lean and obese men participating in study 1. C, Ratio of DAGH to TAGH activity was calculated as marker for complete TAG hydrolysis. Values are mean \pm SEM. *Post hoc* unpaired t test statistics: *, $P < 0.05$ obese compared with lean.

in our muscle samples, which excludes potential contamination via adipocyte infiltration (data not shown).

In line with a lower HSL protein content, muscle DAGH activity, presumably for a large part reflecting HSL activity, was significantly lower in obese compared with lean men (3.85 ± 0.48 vs. 9.71 ± 1.73 nmol/h \cdot mg; group-effect $P = 0.004$; Fig. 1B). Moreover, DAGH activity was positively correlated with HSL content in skeletal muscle of lean and obese men ($r = 0.45$ and $r = 0.36$, respectively; $P < 0.05$). In contrast to a reduced DAGH activity, skeletal muscle total TAGH activity was not significantly affected in our obese men (obese vs. lean, 0.40 ± 0.08 vs. 0.50 ± 0.12 nmol/h \cdot mg; $P = 0.468$; Fig. 1B). Also, HSL-independent TAGH activity, measured after addition of the selective HSL inhibitor BAY [4-isopropyl-3-methyl-2-[1-(3-(*S*)-methyl-piperidin-1-yl)-methanoyl]-2-*H*-

isoxazol-5-yl], was comparable between groups (data not shown). A strong positive correlation between TAGH activity and ATGL content was observed in lean ($r = 0.63$; $P < 0.01$) but not in obese men. The ratio of DAGH to TAGH activity, a marker for complete TAG hydrolysis, was 60% lower in obese individuals (11.4 ± 2.3 vs. 26.5 ± 7.3 ; $P = 0.04$; Fig. 1C).

Total muscle TAG content did not differ between lean and obese men (84.7 ± 18.9 vs. 70.3 ± 12.4 μ mol/mg dry weight; $P = 0.543$) and correlated positively with TAGH activity ($r = 0.55$; $P < 0.05$). Moreover, there was no significant difference in total and percentage of saturated fatty acid (SFA), monounsaturated fatty acid (MUFA), and polyunsaturated fatty acid (PUFA) species in TAG across groups. The TAG C16:1 n-7 to C16:0 ratio (estimated $\Delta 9$ -desaturase activity) was greater in obese compared with lean men (0.172 ± 0.017 vs. 0.117 ± 0.009 ; $P = 0.025$). Surprisingly, total DAG (obese vs. lean, 6.2 ± 0.7 vs. 9.4 ± 0.9 μ mol/mg dry weight; $P = 0.017$) and absolute values of SFA, MUFA, and PUFA species in DAG were lower in muscle from obese men by more than 30%, despite their lower DAGH activity. The percentage of SFA ($P = 0.700$), MUFA ($P = 0.963$), and PUFA ($P = 0.437$) in total DAG was comparable between groups. In line, skeletal muscle DAGH activity showed no significant correlation with total DAG content in lean and obese men.

Study 2

Obese T2D subjects showed the highest muscle ATGL protein content (0.53 ± 0.06 AU; Fig. 2), whereas nonobese normoglycemic controls showed an intermediate value (0.28 ± 0.05 AU), and the lowest ATGL protein content was observed in nonobese T2D men (0.13 ± 0.04 AU) (ANOVA group-effect $P = 0.021$; Fig. 2). Skeletal muscle HSL protein content showed the opposite pattern; *post hoc* analysis showed

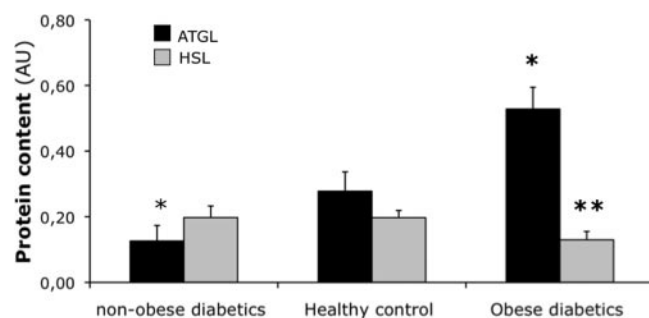


FIG. 2. Skeletal muscle lipase protein content in obese T2D and nonobese T2D men (study 2). The graph depicts ATGL (black bars) and HSL (gray bars) protein content in nonobese T2D, healthy normoglycemic controls, and obese T2D (ANOVA group-effect $P = 0.021$). α -Actin was used to correct for differences in protein loading. Values are mean \pm SEM. *Post hoc* unpaired t test statistics: *, $P < 0.05$, ATGL compared with lean healthy control; **, $P < 0.05$, HSL compared with lean healthy control.

that HSL content was significantly lower in obese T2D compared with nonobese T2D and normoglycemic controls (0.13 ± 0.05 vs. 0.21 ± 0.03 AU; $P < 0.05$; Fig. 2). PLIN was not detectable in our muscle samples, which excludes potential contamination via adipocyte infiltration (data not shown).

Discussion

The present study shows a lower HSL and higher ATGL protein content in skeletal muscle of obese insulin-resistant men. This apparent difference in lipase protein content in skeletal muscle of obese men extends our previous results, which showed a lower HSL content in skeletal muscle of obese men (4). Interestingly, we showed for the first time that this apparent difference in lipase content is related to the obese phenotype because it was observed solely in obese subjects and not in nonobese normoglycemic and T2D controls. Moreover, given the close relationship between mRNA expression of these two lipases, it might be possible that there is some posttranslational regulatory difference that results in their different relative protein expression, which should be explored in more detail in future research (13).

This marked difference in muscle lipase content in obese insulin-resistant men was accompanied by a lower DAGH activity, resulting in a 60% lower ratio of DAGH to TAGH activity, suggesting incomplete TAG hydrolysis (14). This is consistent with our previous observed reduction in HSL serine phosphorylation (presumably reflecting lower HSL activity) and lower forearm muscle glycerol release after an overnight fast in obese insulin-resistant compared with lean men (4). Surprisingly, in contrast to a lower DAGH activity, we paradoxically observed no DAG accumulation in skeletal muscle of obese insulin-resistant men compared with the lean group, but rather a 30% reduction. Although numerous studies have shown elevated DAG levels in skeletal muscle from insulin-resistant rodents and humans (3, 15), there is actually no consensus as to whether this is relevant to the development of skeletal muscle insulin resistance *in vivo* in humans. Our paradoxical results could be partially attributed by a rapid conversion of DAG back to TAG by DAG acyltransferase-1 (DGAT-1) activity in skeletal muscle of obese insulin-resistant men. It has been shown that DGAT-1 overexpression in skeletal muscle decreases DAG and increases TAG content and protects against high-fat diet-induced insulin resistance in rodents (16, 17). However, no difference in muscle DGAT-1 protein content is observed between lean and obese women (18). Alternatively, DAG kinase δ (DGK δ) can phosphorylate DAG to form phosphatidic acid, a lipid second messenger. In contrast to an

elevated DGK activity, however, reduced DGK δ expression and activity has been observed in skeletal muscle from obese T2D patients (19). Our results do not fully preclude the possibility that alterations in DAG species could contribute to insulin resistance. However, we also observed lower SFA, MUFA, and PUFA species in DAG in muscle from obese men. Future research should elucidate whether specific DAG species or stereoisomers and its cellular localization are related to the development of skeletal muscle insulin resistance.

Because our present result largely judges against DAG, it might be speculated that ceramide accumulation plays a more significant role in human obesity and the development of insulin resistance (20). Increased skeletal muscle ceramide content, however, has been observed in some (21–23) but not all human studies with obese insulin-resistant subjects (24). Mechanistically, funneling off the lipid precursors from the ATGL TAGH step to ceramide formation, or alternatively *de novo* ceramide synthesis by increasing serine palmitoyl transferase activity are two possibilities that should be investigated in more detail in future research.

In summary, skeletal muscle HSL protein content is lower and ATGL protein content is higher in the obese state. This difference in lipase content is accompanied by a lower DAGH and a normal TAGH activity, implying incomplete lipolysis. Lower HSL and higher ATGL muscle content is confined to obesity *per se*, because it is solely observed in obese diabetic men but not in nonobese diabetic and healthy normoglycemic men. Importantly, total DAG content is not elevated in skeletal muscle of obese insulin-resistant subjects and was even lower when compared with lean men. Our data do not support the proposed key role of muscle total DAG content in the development of human insulin resistance and suggest that other lipid species (*e.g.* ceramides) or possibly specific DAG species and stereoisomers might be of greater importance, at least in obese insulin-resistant men.

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