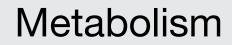


### Translational

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## Contribution of very low-density lipoprotein triglyceride fatty acids to postabsorptive free fatty acid flux in obese humans

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#### ARTICLE INFO

Article history: Received 15 July 2013 Accepted 13 September 2013

Keywords: Isotope dilution [U-<sup>13</sup>C]oleate [<sup>14</sup>C]triolein Hypertriglyceridemia

#### ABSTRACT

*Objective.* In the fasting state, plasma free fatty acids (FFA) are thought to derive almost exclusively from adipose tissue lipolysis. However, there are mixed reports as to whether the spillover of fatty acids (FA) from very low-density lipoprotein triglyceride (VLDL–TG) hydrolysis contributes significantly to the plasma FFA pool. Because substantial VLDL–TG fatty acid spillover into the plasma FFA pool would profoundly impact the interpretation of isotope dilution measures of FFA flux, we investigated the contribution of VLDL–TG spillover to plasma FFA appearance.

Materials/Methods. Eighteen obese adults (15 women) participated in these studies. Each volunteer received a primed, continuous infusion of their own ex-vivo labeled ([ $1^{-14}$ C]triolein) VLDL–TG and a continuous infusion of [U- $^{13}$ C]oleate (8 nmol  $\cdot$  kg fat free mass $^{-1} \cdot$  min $^{-1}$ ) to measure VLDL–TG and FFA rate of appearance (Ra), respectively. The presence of  $^{14}$ C-oleate in the plasma FFA–oleate pool was used to calculate the contribution of spillover from VLDL–TG–oleate to the plasma FFA–oleate Ra.

Results. The spillover rate of VLDL–TG–oleate into plasma FFA–oleate was  $6 \pm 2 \mu$ mol/min (7%  $\pm 2\%$  of [<sup>14</sup>C]oleate from VLDL–TG) and FFA–oleate flux was 240  $\pm 61 \mu$ mol/min. Thus, only 3%  $\pm 1\%$  of total plasma FFA–oleate appearance could be accounted for by VLDL–TG spillover.

*Conclusion.* The contribution of VLDL–TG spillover to the total plasma FFA pool is negligible and will not materially affect the interpretation of FFA flux measures as an index of adipose tissue lipolysis.

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### 1. Introduction

Increased free fatty acids (FFA) derived from excess adipose tissue lipolysis are considered a contributor, if not a primary mediator, of insulin resistance and hypertriglyceridemia [1]. Traditionally, isotopically labeled FFA tracer infusions have been used to measure effective adipose tissue lipolysis under the assumption that the FFA appearing in systemic circulation were a direct result of adipocyte release [2–4]. However, exceptions to this have been observed in that chylomicrontriglyceride fatty acid (TGFA) spillover may contribute 20%– 30% of plasma FFA under postprandial conditions [5–7]. This may vary depending on measurement time following the meal and appears to be greater in adipose tissue than skeletal muscle [8,9]. Although it is generally thought that postabsorptive very low-density lipoprotein triglyceride (VLDL–TG) spillover of FFA is < 5% based upon studies of dogs [10], one group reported that fasting VLDL–TG spillover was ~70% in the

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0026-0495/\$ – see front matter © 2014 Elsevier Inc. All rights reserved. http://dx.doi.org/10.1016/j.metabol.2013.09.008

Abbreviations: FFA, free fatty acids; FA, fatty acid; TG, triglyceride; VLDL, very low density lipoprotein; BMI, body mass index; SA, specific activity.

venous plasma of subcutaneous adipose tissue in humans [11], albeit using different methodology. Given these discrepant findings and the huge impact that a large VLDL–TG spillover would have on interpretation of FFA flux results, we designed a study using independent VLDL–TG and FFA tracers to measure spillover in humans with elevated VLDL–TG.

In order to circumvent the problem of chylomicron–TGFA spillover influencing interpretation of postprandial FFA flux measurements, both an FFA tracer infusion and a meal tracer can be used [5]. Similarly, VLDL–TG spillover needs to be assessed by infusing both a VLDL–TG tracer and an FFA tracer. We employed the ex-vivo labeled VLDL–TG tracer methodology developed for humans by Gormsen et al. [12] to measure both VLDL–TG turnover and the contribution of VLDL–TG spillover to FFA flux, measured using a stable isotope FFA tracer infusion. Using this combination of approaches we were able to directly measure the contribution of postabsorptive VLDL–TGFA spillover to the plasma FFA pool.

#### 2. Methods

#### 2.1. Participants

Eighteen obese (BMI > 35 kg/m<sup>2</sup>) adults (15 women) aged 18-55 years scheduled for bariatric surgery participated in this study, which is part of a larger study of the relationship between hepatic fat content and VLDL-TG turnover. Participants taking lipid lowering medications (e.g. fibrates, statins, niacin) could only be included if it was deemed safe for them to discontinue their use 4 weeks prior to the study. Potential volunteers receiving beta-blockers must have been able to safely discontinue their use 3 days prior to the study in order to participate. Exclusion criteria included type 2 diabetes treated with oral medications or insulin, type 1 diabetes, presence or history of liver disease other than non-alcoholic fatty liver disease, use of nicotine, alcohol consumption exceeding 20 g/day, and the use of medications known to affect lipid metabolism that could not be discontinued prior to the study. The study was approved by the Mayo Clinic Institutional Review Board and written informed consent was obtained from all volunteers.

#### 2.2. Protocol

Approximately one week prior to the inpatient study visit, each participant was seen at the Mayo Clinic outpatient Clinical Research Unit where a 100 mL fasting blood sample was drawn under sterile conditions to be used for ex-vivo VLDL-TG labeling with [1-<sup>14</sup>C]triolein (PerkinElmer, Boston, MA) and body composition was measured using dual-energy X-ray absorptiometry (Lunar iDXA, GE Healthcare, Madison, WI). The following week participants were admitted to the Mayo Clinic inpatient clinical research unit the evening before the VLDL-TG turnover study. They were provided with a meal at ~ 1800 h and then remained fasting except for water overnight. The following morning an intravenous catheter was placed in a forearm vein for tracer infusions, and a retrograde intravenous catheter was placed in the contralateral hand vein to allow for collection of arterialized blood using the heated (55 °C) hand vein technique [13]. At ~0700 h, after collecting a baseline blood sample, a primed, continuous infusion of ex-vivo labeled [1-<sup>14</sup>C]VLDL–TG was started to measure VLDL–TG turnover [14]. Simultaneously, a continuous infusion of [U-<sup>13</sup>C]oleate (8 nmol  $\cdot$  kg fat free mass<sup>-1</sup>  $\cdot$  min<sup>-1</sup>, Isotec, Sigma-Aldrich, Miamisburg, OH) was started to measure FFA kinetics. Blood samples were obtained at 30, 60, 90, 150, 180, 210 and 240 min for measuring VLDL–TG concentration and specific activity (SA) and plasma oleate concentration and enrichment.

#### 2.3. Ex-vivo VLDL-TG tracer preparation

Details regarding ex-vivo labeling of VLDL-TG with [1-<sup>14</sup>C] triolein have previously been published [14]. In brief, the plasma was separated by centrifugation and aliquoted into three sterile vials to which an average of  $38 \pm 4 \,\mu\text{Ci}$  of  $[1-^{14}\text{C}]$ triolein (~12–13  $\mu$ Ci per vial) was added. The plasma was then shaken at room temperature for two hours at 225 rpm (Barnstead Lab-Line MaxQ 4000 E-class shaker, Melrose Park, IL). To isolate the VLDL-TG fraction, 2.5 mL of plasma was layered beneath 3.5 mL of 0.9% normal saline (d = 1.006 g/ml) in a sterile 6 mL centrifuge tube (Quick-Seal, Beckman Coulter, Fullerton, CA), which was centrifuged at 45,000 rpm for 18 h at 4 °C (50.3 Ti rotor, Optima<sup>TM</sup>, LE-80 K, Beckman Instruments, Spinco Division, Palo Alto, CA). After centrifugation, the top 2-2.5 mL containing the VLDL-TG fraction was removed, filtered (Millex GV 0.22 µm filter, Millipore, Billerica, MA) and mixed with normal saline to achieve the volume needed for infusion. The SA of the final infusate was determined by liquid scintillation counting and was used to calculate the total activity infused (average  $\pm$  SD: 13  $\pm$  4  $\mu Ci)$  and the infusion rate. All preparations involving transferring and handling of samples were performed under sterile conditions, and all samples were tested for pyrogens and sterility before infusion.

## 2.4. Plasma and VLDL–TG oleate SA, concentration and enrichment

The VLDL–TG fraction was isolated from plasma samples by density gradient ultracentrifugation as described above. The tubes are sliced with the Beckman tube slicer 2 cm from the top and the VLDL fraction was aspirated into a pre-weighed tube, which was then reweighed in order to calculate the total volume. For each time point, a 1 mL sample of the VLDL fraction was analyzed for specific activity by liquid scintillation counting and 0.5 mL sample was analyzed for TG concentration (mmol/L; Cobas Integra® 400 plus, Roche Diagnostics, Indianapolis, IN) from which total plasma VLDL–TG (µmol/L) was calculated. Total VLDL–TG SA and concentration were calculated using the average steady-state values.

To determine the oleate SA and concentration in VLDL–TG, 1 mL of the VLDL fraction from the 210 time point during infusion was extracted with chloroform:methanol (2:1) and the TG fraction was collected using solid phase extraction columns (Supelco Supelclean LC-NH2 columns, Sigma-Aldrich, St. Louis, MO). After hydrolyzing the TG fraction with methanol:heptane (80:20) and sodium hydroxide, the FA were extracted using Dole solution, derivatized and analyzed by HPLC to determine oleate SA [3]. Download English Version:

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