Introduction

Chronic endogenous glucocorticoid excess in persons with Cushing’s disease is associated with metabolic dysfunction characterized by central obesity, insulin resistance, and dyslipidemia (increased fasting plasma triglyceride (TG) and total and LDL-cholesterol concentrations and decreased HDL-cholesterol concentration) (1). Subclinical hypercortisolemia often occurs in subjects with features of the metabolic syndrome, and it has been suggested that it may be, at least in part, responsible for the development of these metabolic abnormalities. However, the metabolic effects of glucocorticoid administration to mimic subclinical glucocorticoid excess have not been evaluated.

Methods: We used stable isotope-labeled tracer methods in conjunction with magnetic resonance techniques to measure the effect of glucocorticoid excess within the physiological range (~0.7 mg dexamethasone/day for 3 weeks) on glucose and free fatty acid (FFA) rates of appearance (Ra) into plasma, intrahepatic triglyceride (TG) content, very low-density lipoprotein (VLDL)-TG and VLDL-apolipoprotein B-100 (apoB-100) kinetics and plasma lipoprotein subclass concentrations, and particle sizes in nine overweight and obese individuals.

Results: Dexamethasone treatment led to a very small but significant increase in body weight (from 87.4 ± 7.1 to 88.6 ± 7.2 kg; P = 0.003) and increased HDL-cholesterol (from 45.9 ± 2.8 to 55.1 ± 4.6 mg/dl; P = 0.037) and HDL particle (from 33.7 ± 2.2 to 41.4 ± 4.2 nmol/l; P = 0.023) concentrations in plasma but had no effect on intrahepatic TG content, glucose and FFA Ra in plasma, hepatic VLDL-TG and VLDL-apoB-100 secretion rates and mean residence times in the circulation, plasma TG and LDL-cholesterol concentrations, and plasma lipoprotein particle sizes.

Conclusion: Subclinical hypercortisolemia does not have significant adverse metabolic consequences.
We therefore used a combination of stable isotopically labeled tracer and nuclear magnetic resonance techniques to evaluate the effect of chronic glucocorticoid excess on: i) peripheral adipose tissue lipolysis (FFA rate of appearance (Ra) in plasma); ii) intrahepatic TG content; iii) hepatic VLDL-TG and VLDL-apoB-100 secretion rates and VLDL-TG and VLDL-apoB-100 mean residence times in the circulation; iv) hepatic glucose production rate and hepatic insulin sensitivity (hepatic insulin sensitivity index, HISI); and v) plasma lipoprotein subclass concentrations and particle sizes in overweight and obese but otherwise healthy (i.e. no evidence of features of the metabolic syndrome) individuals.

Materials and methods

Subjects

Nine individuals (six women and three men; age: 40.8 ± 4.4 years; body mass index: 31.7 ± 2.0 kg/m²; percent body fat: 35.0 ± 2.6%; mean ± s.e.m.) volunteered for the study. All subjects completed a comprehensive medical examination, which included a detailed history and physical examination, a resting electrocardiogram, standard blood tests, and an oral glucose tolerance test. None of the subjects had elevated morning plasma cortisol concentrations or signs of hypercortisolemia, evidence of illness, or were taking medications known to affect carbohydrate or lipid metabolism. Body composition was determined by using dual-energy X-ray absorptiometry (Hologic QDR 1000/w; Hologic, Inc., Waltham, MA, USA). The study was approved by the Institutional Review Board of Washington University School of Medicine in St Louis. Written informed consent was obtained from all subjects before participation in the study.

Experimental protocol

All subjects completed two metabolic studies: one before and one after low-dose dexamethasone administration for 3 weeks (0.013 mg/kg fat-free mass daily). The specified dose of dexamethasone (~0.7 mg/day) is equivalent to ~20 mg/day hydrocortisone, the dose clinically used for glucocorticoid replacement therapy (14). Before and on the last day of treatment, subjects were admitted to the Clinical Research Unit in the late afternoon. At 1900 h, they consumed a standard meal containing 12 kcal/kg adjusted body weight (15): 55% of total energy was provided as carbohydrate, 30% as fat, and 15% as protein. Subjects then rested in bed and fasted (except for water) until completion of the study the next day.

The following morning, at ~0530 h, one catheter was inserted into a forearm vein to administer stable isotope-labeled tracers (purchased from Cambridge Isotope Laboratories, Andover, MA, USA) and a second catheter was inserted into a contralateral hand vein to obtain arterialized blood samples by heating the hand to 55 °C using a thermostatically controlled box. Blood samples were obtained immediately before the tracer administration to determine plasma substrate, insulin and cytokine concentrations, and background glycerol, palmitate, and leucine tracer-to-tracee ratio (TTR). At 0700 h, a bolus of 0.9% NaCl solution containing [1,1,2,3,3-2H5]glycerol (50 μmol/kg body weight), [5,5,5-2H3]leucine (7.2 μmol/kg), and [6,6-2H2]glucose (22 μmol/kg) was administered through the catheter in the forearm vein, followed by constant infusions of [5,5,5-2H3]leucine (0.12 μmol/kg per min for 12 h) and [6,6-2H2]glucose (0.22 μmol/kg per min for 4 h), dissolved in 0.9% NaCl solution, and [2,2-2H2]palmitate (0.035 μmol/kg per min for 12 h), dissolved in 25% human albumin solution. Blood samples were obtained at 5, 15, 30, 60, 90, and 120 min and then every hour for 10 h to determine free glycerol, glucose, palmitate, and leucine TTRs in plasma and in VLDL-TG and VLDL-apoB-100. Intrahepatic TG content was determined by using magnetic resonance spectroscopy (3T Siemens Magnetom Trio scanner; Siemens, Erlanger, Germany).

Sample collection and storage

Blood samples were collected in chilled tubes containing EDTA to determine plasma lipid concentrations and substrate TTRs. To determine plasma glucose, insulin, and cytokine concentrations, blood samples were collected in chilled tubes containing heparin. Samples were placed on ice and plasma was separated by centrifugation within 30 min of collection. Aliquots of plasma were kept in the refrigerator for immediate isolation of VLDL and measurement of plasma apolipoprotein B-100 concentration. The remaining plasma samples were stored at −80 °C until the final analyses were performed. The VLDL fraction was isolated from plasma by ultracentrifugation (16, 17). Aliquots of the VLDL fraction were used for measuring VLDL-apoB-100 concentration immediately after collection; the remaining samples were stored at −80 °C until final analyses.

Sample processing and analyses

Plasma concentrations of total TG, total cholesterol, HDL-cholesterol, and LDL-cholesterol were measured by standard hospital laboratory methods using a Hitachi 917 autoanalyzer (Hitachi). Plasma lipoprotein particle concentrations and average lipoprotein sizes were determined by using proton nuclear magnetic resonance spectroscopy (LipoScience, Raleigh, NC, USA) (18, 19). Plasma glucose concentration was determined by using an automated glucose analyzer (YSI 2300 STAT plus; Yellow Springs Instrument Co., Yellow Springs, OH, USA). Plasma insulin concentration was measured
by using an automated chemiluminescent immunoanalyzer (IMMULITE; Siemens Healthcare Diagnostic, Los Angeles, CA, USA). Plasma-FFA concentrations were quantified by gas chromatography (GC; Hewlett-Packard 5890-II, Palo Alto, CA, USA) after adding heptadecanoic acid to plasma as an internal standard (20). Plasma apoB-100 and VLDL-apoB-100 concentrations were measured by using a turbidimetric immunoassay (Kamiya Biomedical Co., Seattle, WA, USA). VLDL-TG concentration was measured by using a colorimetric enzymatic kit (Sigma–Aldrich Co.). The concentrations of tumor necrosis factor alpha (TNF-α), interleukin 6 (IL6), and cortisol were measured by using commercially available ELISAs (R&D Systems, Minneapolis, MN, USA). Plasma ACTH concentration was measured by using a commercially available immunoassay on the Immulite 2000 system at the Mayo Clinic in Rochester, MN, USA.

The TTRs of glycerol, palmitate, leucine, and glucose in plasma, glycerol and palmitate in VLDL-TG, and leucine in VLDL-apoB-100 were determined by GC–mass spectrometry (MS; MSD 5973 System, Hewlett-Packard) as described previously (16, 17, 20, 21, 22). Plasma-free palmitate and palmitate in VLDL-TG were analyzed as methylesters. The heptafluorobutyryl derivative was formed for the analysis of glycerol and glucose in plasma and glycerol and VLDL-TG, the N-heptafluorobutyryl N-propyl ester derivative was used for leucine in VLDL-apoB-100, and the t-butyldimethylsilyl derivative was used for free leucine in plasma.

**Calculations**

The fractional turnover rate (FTR) of VLDL-TG and VLDL-apoB-100 was determined by fitting the glycerol and leucine TTR time courses in plasma and VLDL-TG and VLDL-apoB-100 respectively, to multicompartmental models as described previously (16, 17, 20, 21, 22). The secretion rates of VLDL-TG (μmol/l plasma per min) and VLDL-apoB-100 (nmol/l plasma per min), which represent the amount of VLDL-TG and VLDL-apoB-100 (i.e. VLDL particles) secreted by the liver per unit of plasma, were calculated by multiplying the FTR of VLDL-TG and VLDL-apoB-100 (i.e. VLDL particles) by the plasma concentrations of VLDL-TG (μmol/l) and VLDL-apoB-100 (nmol/l) respectively. The mean residence times (MRT) of VLDL-TG and VLDL-apoB-100, which represent the time VLDL-TG and VLDL-apoB-100 remain in the circulation after secretion by the liver, were calculated as 1/FTR.

Palmitate Ra in plasma was calculated by dividing the palmitate tracer infusion rate by the average plasma palmitate TTR value between 60 and 240 min. Total FFA Ra was calculated by dividing palmitate Ra by the proportional contribution of palmitate to total plasma FFA concentration (23). The relative contribution of systemic plasma FFA and nonsystemic fatty acids to total VLDL-TG production was calculated on the basis of isotopic dilution upon fitting the palmitate TTR in plasma and VLDL-TG to a multicompartmental model (16, 17, 22). Nonsystemic fatty acids in VLDL-TG are derived from the pools of fatty acids that are not labeled with tracer during the infusion period, including fatty acids derived from: i) lipolysis of preexisting, slowly turning over lipid stores in the liver and tissues draining directly into the portal vein; ii) lipolysis of plasma lipoproteins that are taken up by the liver; and iii) hepatic de novo lipogenesis.

Glucose Ra into plasma was calculated by dividing the glucose tracer infusion rate by the average plasma glucose TTR between 210 and 300 min (22). The HSI was calculated as the inverse of the product of glucose Ra and plasma insulin concentration (24, 25). The adipose tissue insulin sensitivity index (ATISI) was calculated as the inverse of the product of FFA Ra and plasma insulin concentration (26).

**Results**

Dexamethasone treatment caused a very small but significant increase in body weight (from 87.4 ± 7.1 to 88.6 ± 7.2 kg; P = 0.003) but did not affect intrahepatic TG content (before: 5.1 ± 1.9%; after: 3.6 ± 0.9%; P = 0.261). Morning plasma cortisol concentration decreased by ~50% after dexamethasone treatment; ACTH, TNF-α, and IL6 concentrations remained unchanged (Table 1). Glucose Ra (Fig. 1) and glucose concentration (Table 1) in plasma were not different before or after dexamethasone treatment. FFA Ra in plasma was not affected (P = 0.185) by dexamethasone treatment (Fig. 1), but plasma FFA concentration was reduced by ~20% (Table 1). Plasma insulin concentration increased by ~60% (Table 1). Accordingly, the HSI and the ATISI (Fig. 1) declined by ~50 and 25%, respectively, after dexamethasone treatment, but these differences failed to reach significance at the P < 0.05 level (P = 0.066 and 0.086 respectively).

There were no significant differences (P ≥ 0.3) in VLDL-TG and VLDL-apoB-100 secretion rates and MRTs before or after dexamethasone treatment (Fig. 2). The contribution of systemic plasma FFA and nonsystemic fatty acids to total VLDL-TG production was not affected by dexamethasone treatment: systemic plasma FFA
Table 1 Metabolic profile before and after 3 weeks of low-dose dexamethasone treatment. Values are mean±S.E.M. or median (quartiles).

<table>
<thead>
<tr>
<th></th>
<th>Before</th>
<th>After</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Glucose (mg/dl)</td>
<td>94.1±3.0</td>
<td>92.3±1.8</td>
<td>0.451</td>
</tr>
<tr>
<td>Insulin (mU/ml)</td>
<td>8.5±1.7</td>
<td>11.9±2.3</td>
<td>0.048</td>
</tr>
<tr>
<td>Free fatty acids (µmol/l)</td>
<td>506±47</td>
<td>390±40</td>
<td>0.003</td>
</tr>
<tr>
<td>Total cholesterol (mg/dl)</td>
<td>164±10</td>
<td>168±15</td>
<td>0.514</td>
</tr>
<tr>
<td>HDL-cholesterol (mg/dl)</td>
<td>45.9±2.8</td>
<td>55.1±4.6</td>
<td>0.037</td>
</tr>
<tr>
<td>LDL-cholesterol (mg/dl)</td>
<td>99.4±8.9</td>
<td>96.6±10.7</td>
<td>0.620</td>
</tr>
<tr>
<td>Total TG (mg/dl)</td>
<td>91.8±20.1</td>
<td>83.8±17.3</td>
<td>0.330</td>
</tr>
<tr>
<td>VLDL-TG (mg/dl)</td>
<td>30.9±4.9</td>
<td>29.3±7.0</td>
<td>0.820</td>
</tr>
<tr>
<td>Total apoB-100 (mg/dl)</td>
<td>66.0±8.3</td>
<td>65.8±8.9</td>
<td>0.980</td>
</tr>
<tr>
<td>VLDL-apoB-100 (mg/dl)</td>
<td>2.6±0.5</td>
<td>2.6±0.6</td>
<td>0.896</td>
</tr>
<tr>
<td>Tumor necrosis factor α (pg/ml)</td>
<td>0.8 (0.7–1.2)</td>
<td>1.0 (0.4–1.2)</td>
<td>0.374</td>
</tr>
<tr>
<td>Interleukin 6 (pg/ml)</td>
<td>2.7 (1.6–3.2)</td>
<td>2.4 (1.5–4.5)</td>
<td>0.214</td>
</tr>
<tr>
<td>Cortisol (µg/dl)</td>
<td>5.76±0.49</td>
<td>2.97±0.66</td>
<td>0.008</td>
</tr>
<tr>
<td>ACTH (pg/ml)</td>
<td>21.8±2.0</td>
<td>15.2±3.6</td>
<td>0.125</td>
</tr>
</tbody>
</table>

accounted for 58±4% of all fatty acids in VLDL-TG before and 55±5% after dexamethasone treatment (P = 0.485).

The concentrations of total and LDL-cholesterol, total and VLDL-TG, and total VLDL-apoB-100 in plasma were not affected by dexamethasone treatment, whereas plasma HDL-cholesterol concentration increased by ~20% (Table 1). VLDL, intermediate-density lipoprotein (IDL), and LDL particle concentrations and average particle sizes were not different before or after dexamethasone treatment, but total HDL particle concentration increased by ~20%, mainly because of more medium-sized particles that therefore did not affect the average HDL size (Table 2).

Discussion

Endogenous glucocorticoid excess has been proposed to be at least partially responsible for the metabolic abnormalities associated with obesity (2, 3). In this study, we evaluated the effect of treatment using dexamethasone, a synthetic glucocorticoid, on metabolic function in overweight and obese individuals with normal plasma cortisol concentrations at baseline. In our study, we chose to administer dexamethasone orally to induce glucocorticoid excess not only in the systemic circulation but also in the portal system because the metabolic syndrome in obese persons is characterized by increased glucocorticoid signaling in peripheral and visceral adipose tissues due to hyperactivity of the hypothalamic–pituitary–adrenal axis and increased 11β-hydroxysteroid dehydrogenase type 1 activity in adipose tissue (peripheral and visceral) and possibly also liver; in addition, portal vein cortisone provides a substrate for liver cortisol production (27, 28, 29). We found that dexamethasone treatment led to a slight increase in body weight and a slight decrease in hepatic and adipose tissue insulin sensitivity (evident by the unchanged glucose and FFA Ra in plasma in the face of greater plasma insulin concentration) but had no effect on intrahepatic TG content and basal rates of hepatic VLDL-TG and VLDL-apoB-100 secretion, VLDL-TG and VLDL-apoB-100 mean residence times in the circulation, plasma TG and LDL-cholesterol concentrations, and circulating lipoprotein particle sizes. In addition, dexamethasone treatment had some beneficial metabolic effects by increasing plasma HDL-cholesterol and particle concentrations.

The adverse effects of glucocorticoids on body weight and insulin sensitivity are well established (3, 30) and confirmed in our study. Nevertheless, plasma glucose and FFA Ra into plasma, which are tightly controlled by insulin (31, 32), did not increase in response to dexamethasone, presumably due to a compensatory increase in insulin secretion (33). This is consistent with the results from the previous studies, which demonstrated that large doses of glucocorticoids (30–50 mg/day prednisone, equivalent to six to ten times the dose of dexamethasone given in our study (14)) do not affect adipose tissue lipolysis and hepatic glucose production (34, 35). However, others have shown that acutely, within hours to a few days, glucocorticoid administration at even larger doses increases glucose and FFA Ra in plasma (4, 5, 33, 36). This may be the result of a dose-dependent effect of glucocorticoids on pancreatic β-cell function, leading to proportionally smaller compensatory increases in circulating insulin (37). Subclinical hypercortisolemia per se therefore does not disturb plasma glucose and FFA homeostasis. However, in susceptible persons, it might accelerate the progression of insulin resistance to type 2 diabetes mellitus (38).
We are not aware of any studies that evaluated the effects of exogenous glucocorticoid administration on VLDL kinetics in human subjects, except in those with Cushing’s syndrome which found that hepatic VLDL-TG production rate is greater in patients with endogenous hypercortisolemia than in healthy subjects (39). These results, however, are difficult to interpret because of multiple potential confounding factors. Our data suggest that low-dose glucocorticoid administration does not stimulate VLDL-TG or VLDL-apoB-100 secretion from the liver and does not adversely affect plasma TG or apoB-100 homeostasis. In addition, the low-dose dexamethasone treatment used in our study increased plasma HDL-cholesterol concentration by ~20%. This was linked to an increase in the total number of circulating HDL particles, predominantly due to increased medium-sized HDL. These findings are consistent with the results from studies in which glucocorticoids were administered as anti-inflammatory and immunosuppressive agents in patients with rheumatic and pulmonary diseases, which indicate that the glucocorticoid-induced increase in HDL-cholesterol concentration (9, 10, 11) is predominantly due to an increase in the HDL_2 subfraction (large- and medium-sized HDL) but not in the HDL_1 subfraction (small HDL) (40, 41).

The mechanisms responsible for the glucocorticoid-induced increase in HDL particle and HDL-cholesterol concentrations are not entirely clear but most likely include decreased cholesterol ester transfer protein expression and increased secretion of apoA-I (the main protein of HDL particles) from the liver (42). We can exclude the possibility of reduced cholesterol transfer to VLDL (in exchange for TG) due to faster removal of VLDL-TG from the circulation, which has been proposed to occur in response to glucocorticoid-induced stimulation of lipoprotein lipase (LPL) (43, 44, 45). In our study, dexamethasone administration had no effect on VLDL-apoB-100 or VLDL-TG mean residence times in the circulation. The apparent discrepancy between the dexamethasone-induced stimulation of ex vivo measured adipose tissue LPL in humans and rodents (43, 44, 45) and our results are most likely related to the fact that ex vivo-measured tissue or plasma LPL mass or activity provide information regarding the maximal capacity for LPL-mediated lipolysis but do not reflect in vivo LPL activity that is affected by a variety of factors such as apolipoproteins (apoA-I and apoC-I) present on the particle surface, as well as local FFA concentration, blood flow through the tissue capillary beds, etc. (46). In fact, marked differences in post-heparin plasma LPL activity have been demonstrated in the absence of detectable changes in plasma TG clearance rates in vivo (47).

We cannot exclude the possibility that the low dose of dexamethasone used or the short duration of our study limited the metabolic adverse effects. However, we consider it unlikely because we did observe a decrease in insulin sensitivity and an increase in HDL concentration, which are the commonly observed metabolic side effects of glucocorticoid pharmacotherapy. Furthermore, longer duration of treatment and/or using pharmacological doses would most likely have led to confounding influences (2, 3). Dexamethasone treatment lowered endogenous cortisol release and may have also reduced endogenous androgen production. Nevertheless, total glucocorticoid activity (endogenous plus exogenous) was greater with dexamethasone treatment and it is unlikely that changes in androgen availability confounded our results because we have shown that an approximately sevenfold change in plasma testosterone concentration and the normal physiological variations

### Table 2 Lipoprotein particle concentrations and sizes before and after 3 weeks of low-dose dexamethasone treatment. Values are mean ± S.E.M. or median (quartiles).

<table>
<thead>
<tr>
<th>Lipoprotein</th>
<th>Before</th>
<th>After</th>
<th>P value</th>
</tr>
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<tbody>
<tr>
<td><strong>VLDL particles</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Total (nmol/l)</td>
<td>68.4 ± 12.2</td>
<td>58.0 ± 9.2</td>
<td>0.339</td>
</tr>
<tr>
<td>Large</td>
<td>0.55 (0.13–0.96)</td>
<td>0.61 (0.10–1.39)</td>
<td>0.859</td>
</tr>
<tr>
<td>Medium</td>
<td>26.1 ± 6.3</td>
<td>20.7 ± 4.1</td>
<td>0.300</td>
</tr>
<tr>
<td>Small</td>
<td>40.4 ± 6.1</td>
<td>35.4 ± 6.2</td>
<td>0.406</td>
</tr>
<tr>
<td><strong>Average VLDL size (nm)</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Large</td>
<td>44.4 ± 1.8</td>
<td>46.9 ± 3.2</td>
<td>0.274</td>
</tr>
<tr>
<td>Medium</td>
<td>29.4 ± 12.4</td>
<td>28.9 ± 13.3</td>
<td>0.833</td>
</tr>
<tr>
<td>Small</td>
<td>1139 (853–1327)</td>
<td>1043 (900–1500)</td>
<td>0.260</td>
</tr>
<tr>
<td><strong>IDL particles (nmol/l)</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Large</td>
<td>48.8 ± 47</td>
<td>47.8 ± 40</td>
<td>0.416</td>
</tr>
<tr>
<td>Medium</td>
<td>600 (405–859)</td>
<td>618 (477–910)</td>
<td>0.441</td>
</tr>
<tr>
<td>Small</td>
<td>21.2 ± 0.2</td>
<td>21.1 ± 0.1</td>
<td>0.564</td>
</tr>
<tr>
<td><strong>Average LDL size (nm)</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Large</td>
<td>7.8 (5.7–11.1)</td>
<td>8.9 (6.6–17.3)</td>
<td>0.066</td>
</tr>
<tr>
<td>Medium</td>
<td>2.7 ± 0.6</td>
<td>6.7 ± 1.5</td>
<td>0.010</td>
</tr>
<tr>
<td>Small</td>
<td>22.9 ± 1.5</td>
<td>23.5 ± 2.0</td>
<td>0.769</td>
</tr>
<tr>
<td><strong>Average HDL size (nm)</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Large</td>
<td>9.1 ± 0.1</td>
<td>9.1 ± 0.1</td>
<td>0.939</td>
</tr>
</tbody>
</table>

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in plasma progesterone concentration do not affect VLDL kinetics in obese women (48, 49).

In summary, low-dose dexamethasone treatment in obese subjects led to a small decrease in insulin sensitivity, which was compensated for by an increase in insulin secretion and an increase in HDL-cholesterol and HDL particle concentrations in plasma but had no effect on intrahepatic TG content, glucose or FFA Ra in plasma, or pro-atherogenic lipoprotein concentrations or VLDL metabolism. Subclinical hypercortisolism, therefore, is unlikely to be responsible for the development of the metabolic complications associated with obesity.

Declaration of interest
The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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Author contribution statement
X Wang, D N Reeds, and J Kampelman conducted the experiments and researched the data; X Wang, F Magkos, B W Patterson, and B Mittendorfer analyzed and interpreted the data; X Wang, F Magkos, and B Mittendorfer wrote the manuscript; all authors edited the manuscript; B Mittendorfer wrote the manuscript; all authors edited the manuscript; B Mittendorfer designed the study and had the overall supervision.

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