

# Low-energy diets differing in fibre, red meat and coffee intake equally improve insulin sensitivity in type 2 diabetes: a randomised feasibility trial

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## Abstract

**Aims/hypothesis** Epidemiological studies have found that a diet high in fibre and coffee, but low in red meat, reduces the risk for type 2 diabetes. We tested the hypothesis that these

nutritional modifications differentially improve whole-body insulin sensitivity (primary outcome) and secretion.

**Methods** Inclusion criteria were: age 18–69 years, BMI  $\geq 30$  kg/m<sup>2</sup>, type 2 diabetes treated with diet, metformin or

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acarbose and known disease duration of  $\leq 5$  years. Exclusion criteria were:  $\text{HbA}_{1c} > 75$  mmol/mol (9.0%), type 1 or secondary diabetes types and acute or chronic diseases including cancer. Patients taking any medication affecting the immune system or insulin sensitivity, other than metformin, were also excluded. Of 59 patients (randomised using randomisation blocks [four or six patients] with consecutive numbers), 37 (54% female) obese type 2 diabetic patients completed this controlled parallel-group 8-week low-energy dietary intervention. The participants consumed either a diet high in cereal fibre (whole grain wheat/rye: 30–50 g/day) and coffee ( $\geq 5$  cups/day), and free of red meat (L-RISK,  $n=17$ ) or a diet low in fibre ( $\leq 10$  g/day), coffee-free and high in red meat ( $\geq 150$  g/day) diet (H-RISK,  $n=20$ ). Insulin sensitivity and secretion were assessed by hyperinsulinaemic–euglycaemic clamp and intravenous glucose tolerance tests with isotope dilution. Whole-body and organ fat contents were measured by magnetic resonance imaging and spectroscopy.

**Results** Whole-body insulin sensitivity increased in both groups (mean [95% CI]) (H-RISK vs L-RISK: 0.8 [0.2, 1.4] vs 1.0 [0.4, 1.7]  $\text{mg kg}^{-1} \text{min}^{-1}$ ,  $p=0.59$ ), while body weight decreased ( $-4.8\%$  [ $-6.1\%$ ,  $-3.5\%$ ] vs  $-4.6\%$  [ $-6.0\%$ ,  $-3.3\%$ ], respectively). Hepatic insulin sensitivity remained unchanged, whereas hepatocellular lipid content fell in both groups ( $-7.0\%$  [ $-9.6\%$ ,  $-4.5\%$ ] vs  $-6.7\%$  [ $-9.5\%$ ,  $-3.9\%$ ]). Subcutaneous fat mass ( $-1,553$  [ $-2,767$ ,  $-340$ ]  $\text{cm}^3$  vs  $-751$  [ $-2,047$ ;  $546$ ]  $\text{cm}^3$ , respectively) visceral fat mass ( $-206$  [ $-783$ ,  $371$ ]  $\text{cm}^3$  vs  $-241$  [ $-856$ ,  $373$ ]  $\text{cm}^3$ , respectively) and muscle fat content ( $-0.09\%$  [ $-0.16\%$ ,  $-0.02\%$ ] vs  $-0.02\%$  [ $-0.10\%$ ,  $0.05\%$ ], respectively) decreased similarly. Insulin secretion remained unchanged, while the proinflammatory marker IL-18 decreased only after the L-RISK diet.

**Conclusions/interpretation** No evidence of a difference between both low-energy diets was identified. Thus, energy restriction per se seems to be key for improving insulin action in phases of active weight loss in obese type 2 diabetic patients, with a potential improvement of subclinical inflammation with the L-RISK diet.

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**Keywords** Diet · Inflammation · Insulin sensitivity · Type 2 diabetes · Weight loss

### Abbreviations

AIR Acute insulin response  
BCAA Branched chain amino acids

DHBA	3,5-Dihydroxybenzoic acid
DHPPA	3-(3,5-Dihydroxyphenyl)-propanoic acid
DI	Disposition index
EGP	Endogenous glucose production
FA	Fatty acids
GDRS	German Diabetes Risk Score
HCL	Hepatocellular lipid content
H-RISK	Higher-risk diet: low fibre coffee-free, high red meat diet
IMCL	Intramyocellular lipids
IQR	Interquartile range
jMRUI	Java-based Magnetic Resonance User Interface
L-RISK	Low-risk diet: high whole grain fibre and coffee, red meat-free diet
PKC $\theta$	Protein kinase-C theta
$R_d$	Glucose disappearance
SCAT	Truncal subcutaneous adipose tissue
TE	Echo time
TE eff	Effective echo time
TR	Repetition time
VAT	Visceral adipose tissue
$\dot{V}\text{CO}_2$	Carbon dioxide production
$\dot{V}\text{O}_2$	Oxygen consumption

### Introduction

Several clinical scores predicting the future risk of type 2 diabetes have been developed comprising classical risk factors such as age, sex and BMI, in addition to other parameters [1–3]. The Finnish diabetes risk score integrates dietary components such as fibre [1], while the German Diabetes Risk Score (GDRS) also includes coffee and red meat consumption [2]. Results of observational studies support the protective effects of high fibre and coffee intake and reduced red meat consumption [4–6], but randomised intervention trials are needed to exclude unrecognised confounders and to reduce diabetes-related mortality in the future [7].

Few randomised controlled trials have tested the effects of dietary components on key mechanisms underlying the pathogenesis of type 2 diabetes, such as impaired insulin sensitivity and secretion. Isoenergetic diets with higher fibre intake improved insulin sensitivity in overweight glucose-tolerant humans [8], while no change in insulin sensitivity was observed in other studies comparing whole grain with refined grain intake [9]. Regular coffee drinking did not affect glucose homeostasis despite its anti-inflammatory effects [10], while a single cup of coffee even worsened postprandial glucose excursion in individuals with and without type 2 diabetes [11]. This may have resulted from adrenaline

(epinephrine) release and adenosine receptor inhibition followed by impaired insulin sensitivity [11].

Greater consumption of red or processed meat could decrease insulin sensitivity because of its high protein and fat content, both of which can inhibit insulin action via mTOR, toxic lipid intermediates, inflammatory pathways or oxidative stress [12–14], and elevated branched chain amino acids (BCAA) are predictors of incident type 2 diabetes [15]. On the other hand, both amino acids and fatty acids (FA) can enhance insulin secretion and thereby counteract a reduction in insulin sensitivity [16, 17]. At present, no controlled trials have reported the effects of red meat consumption on insulin sensitivity and glucose metabolism.

We hypothesised that a dietary intervention combining the nutrients that best predict a lower diabetes risk according to the GDRS would improve insulin sensitivity and subclinical inflammation beyond the effects of weight loss in short-term, well-controlled type 2 diabetes. Thus, this randomised controlled parallel group feasibility trial examines the effects of a moderate energy-restricted diet, as recommended by the ADA, with either increased fibre and coffee consumption and a lack of red meat (low-risk diet, L-RISK) or with increased red meat consumption, low in fibre and lacking coffee (higher-risk diet, H-RISK) on whole-body insulin sensitivity (primary endpoint), insulin secretion, body and tissue fat content and markers of inflammation.

## Methods

**Volunteers** All participants gave written informed consent before their inclusion in the study (Clinicaltrials.gov registration number: NCT01409330), which was performed according to the Declaration of Helsinki (2008 version) and approved by the local ethics board. Inclusion criteria were: age 18–69 years, BMI  $\geq 30$  kg/m<sup>2</sup>, type 2 diabetes treated with diet, metformin or acarbose and known disease duration of  $\leq 5$  years. Exclusion criteria were: HbA<sub>1c</sub>  $> 75$  mmol/mol (9.0%), type 1 or secondary diabetes types and acute or chronic diseases including cancer. Patients taking any medication affecting the immune system or insulin sensitivity, other than metformin, were also excluded. Screening included medical history, laboratory tests, anthropometry and 12-lead ECG. All participants were randomly allocated to one intervention at one centre using randomisation blocks (four or six patients) with consecutive numbers. Patients were instructed to maintain their medication and physical activity assessed by weekly protocols.

**Experimental design** After randomisation, participants maintained their dietary habits during the run-in period (19  $\pm$  8 days), and were monitored by daily dietary weighing protocols to calculate average energy intake (Prodi expert 6.1, Nutri-Science, Hausach, Germany). Metformin was stopped 3 days before and during the experimental tests, which were

performed on 3 days within a single week [18]. After an overnight fast of 10–12 h, the participants reported to one of the three clinical research centres (Düsseldorf, Berlin-Potsdam, Heidelberg) for metabolic tests. During the intervention, all participants received individually calculated daily diet sheets providing a 1,256 kJ reduction in total daily energy intake and a constant macronutrient distribution (50% of energy from carbohydrates, 30% from fat, 20% from protein). The L-RISK diet contained 30–50 g/day cereal fibre from wheat and rye (100 g of wholegrain crispbread and 250–300 g of wheat/rye wholegrain bread),  $\geq 5$  cups of coffee/day containing 7–8 g coffee powder each and no red meat (although poultry was allowed). The H-RISK diet contained  $\leq 10$  g/day whole grain fibre,  $\geq 150$  g/day red meat (beef) and no coffee or tea. Patients were asked to document any changes and return the completed sheet to monitor compliance. The weekly visits also included monitoring of body weight, hip and waist circumference, blood pressure and blood sampling. After 8 weeks of intervention, the participants again stopped metformin for 3 days and underwent tests identical to those at baseline.

**Modified Botnia clamp** After fasting blood sampling, participants received a primed (0.36 mg [fasting blood glucose in mmol/l]  $\times$  [5 mmol/l]<sup>-1</sup>  $\times$  [body weight in kg]<sup>-1</sup> min<sup>-1</sup> for 5 min), continuous intravenous infusion (0.036 mg  $\times$  [body weight in kg]<sup>-1</sup> min<sup>-1</sup>) of [6,6-<sup>2</sup>H<sub>2</sub>]glucose (<sup>2</sup>H glucose, 99% enriched; CIL, Andover, MA, USA) started at -120 min to assess endogenous glucose production (EGP) [18]. At 0 min, an intravenous glucose bolus (1 mg/kg body weight in a 30% [wt/vol.] solution containing 1.98% of [6,6-<sup>2</sup>H<sub>2</sub>]glucose) was injected within 60 s and blood samples were obtained every 2 min for the first 10 min, and then every 10 min up to 1 h to assess glucose-induced insulin secretion. The hyperinsulinaemic-euglycaemic clamp was then performed for 3 h using a primed-continuous insulin infusion (10 mU [body weight in kg]<sup>-1</sup> min<sup>-1</sup> for 10 min, followed by 1.5 mU [body weight in kg]<sup>-1</sup> min<sup>-1</sup>; Insuman Rapid, sanofi-aventis, Frankfurt am Main, Germany) to assess insulin sensitivity. A variable infusion of 20% glucose (wt/vol.; B. Braun, Melsungen, Germany) enriched with 2% [6,6-<sup>2</sup>H<sub>2</sub>]glucose (wt/vol.) was administered to maintain blood glucose at 5 mmol/l [18]. Plasma glucose was measured at 5 min intervals.

**Indirect calorimetry** Indirect calorimetry was performed in the canopy mode (Vmax Encore 29n, CareFusion, Höchberg, Germany) during fasting and steady-state clamp conditions for 20 min followed by a post-exposure recalibration procedure [19]. Oxygen consumption ( $\dot{V}O_2$ ) and carbon dioxide production ( $\dot{V}CO_2$ ) were measured and substrate oxidation rates were calculated as previously described [13]. Non-oxidative glucose disposal was

calculated as the difference between glucose disappearance ( $R_d$ ) and carbohydrate oxidation.

**Laboratory analyses** Blood samples were immediately chilled and centrifuged, and supernatants stored at  $-20^\circ\text{C}$  until analysis. Whole-blood glucose concentration was measured at the bedside (EKF biosen C-Line glucose analyser, EKF Diagnostics, Barleben, Germany) [20]. Serum triacylglycerols, cholesterol and liver enzymes were analysed using a Cobas c311 analyser (Roche Diagnostics, Mannheim, Germany). FA, C-peptide, insulin, glucagon, glucagon-like peptide 1 (GLP-1) and gastric inhibitory peptide (GIP) were measured as previously described [13]. Serum cytokines were assayed using the Quantikine HS (IL-6) and Quantikine (total adiponectin, IL-1RA) ELISA kits (R&D Systems, Wiesbaden, Germany) and an IL-18 ELISA kit (MBL, Nagoya, Japan) [21]. Serum amino acids were measured by GC-MS using an EZ:faast kit (Phenomenex, Torrance, CA, USA) with Norvaline as internal standard. Plasma coffee-derived compounds (caffeine and its metabolites, theophylline, theobromine and paraxanthine, and the polyphenols caffeic acid and its metabolites, dihydrocaffeic acid, ferulic acid, dihydroferulic acid, isoferulic acid, dihydroisoferulic acid, dihydro-3-coumaric acid, 3-(3,4-dimethoxyphenyl)propionic acid, 3,4-dimethylcaffeic acid and 3-coumaric acid) were determined using HPLC and GC-MS [10]. The two main urinary alkylresorcinol metabolites 3,5-dihydroxybenzoic acid (DHBA) and 3-(3,5-dihydroxyphenyl)-propanoic acid (DHPPA) were analysed using GC-MS [22]. Blood atom per cent enrichment of  $^2\text{H}$  was measured by GC-MS (Agilent 6890 and 5975, Santa Clara, CA, USA) as previously described [13].

**Magnetic resonance imaging and spectroscopy** The experiments were performed on a clinical 3 T (Philips X-series Achieva, Best, the Netherlands) or a 1.5 T (Magnetom, Siemens Healthcare, Erlangen, Germany) whole-body magnet using standardised procedures [23].

Hepatocellular lipid content (HCL) was quantified from non-water and water-suppressed  $^1\text{H}$  MR spectra using stimulated echo acquisition mode (STEAM) and absolute concentrations were expressed as per cent hepatocellular lipids relative to water content [24]. Truncal subcutaneous (SCAT) and visceral (VAT) adipose tissue were measured by whole-body imaging using transverse multislice turbo spin echo (TSE) sequences (repetition time/effective echo time:  $\text{TR}/\text{TE}_{\text{eff}}=400\text{--}510/38$  ms for 3 T and  $\text{TR}/\text{TE}=490/12$  ms for 1.5 T) with a turbo factor of 7 and using the quadrature body coil [25].

Intramyocellular lipids (IMCL) were measured in the left calf using FLEX-S coils (Philips Healthcare, Best, the Netherlands) and a single-voxel point-resolved spectroscopy (PRESS) sequence ( $\text{TR}/\text{TE}=2,000/32$  ms) with a voxel size of

2 cm placed within the soleus and tibialis anterior muscles.  $^1\text{H}$  MR spectra were processed using Java-based Magnetic Resonance User Interface (jMRUI; [www.mruui.uab.es/mruui/](http://www.mruui.uab.es/mruui/); version 4.0) and IMCL quantified from the peak areas of the signal from middle chain methylene groups ( $-\text{CH}_2-$ , at 1.3 ppm) relative to water signals in the same voxel.

**Calculations and statistical analyses** EGP before the start of the clamp (min  $-30$  to 0), whole-body insulin sensitivity ( $R_d$ , primary outcome) and hepatic insulin sensitivity (per cent EGP suppression by insulin) were determined from the tracer infusion rate of D-[6,6- $^2\text{H}_2$ ] glucose and its enrichment of hydrogen bound to carbon 6 divided by the mean per cent enrichment of plasma D-[6,6- $^2\text{H}_2$ ]glucose [13]. The acute insulin response (AIR; pmol/l), as a measure of first phase insulin secretion during IVGTT, and the disposition index (DI), an index of beta cell function corrected for insulin sensitivity, were calculated as described previously [26]. For the feasibility study, change in whole-body insulin sensitivity was the primary outcome to obtain effect size and variance estimates for both diets; all other analyses have been predefined secondary exploratory analyses carried out on potential mechanisms.

Normally distributed parameters are presented as means  $\pm$  SD, otherwise as median (interquartile range [IQR]). Non-normally distributed data were  $\log_e$ -transformed to achieve near-normal distribution and compared using a two-tailed Student's *t* test for unpaired samples for comparisons between groups and paired analysis (before and after intervention). Mean change within-group is given with 95% CI, and the difference of the within-group changes were compared using general linear models with estimating least square means and 95% CI adjusted for the respective baseline value. Pearson correlation coefficients were calculated. Statistical significance of differences was set at  $p<0.05$ . Calculations were performed using GraphPad Prism version 6.03 (GraphPad Software, La Jolla, CA, USA) and SAS Version 9.2 TS Level 2M3 (SAS Institute, Cary, NC, USA).

## Results

**Study population** Between April 2011 and October 2013, 216 patients were contacted. Of these, 90 did not meet the inclusion criteria, 33 declined to participate and 34 were excluded for other reasons. Of the remaining 59 randomised participants (30 in the L-RISK group, 29 in the H-RISK group), 22 patients (13 in L-RISK, nine in H-RISK) discontinued the study during the run-in or treatment phase or were excluded from the analysis (electronic supplementary material [ESM] Fig. 1). The groups did not differ in baseline characteristics (Table 1).

**Macronutrient intake** At baseline, total energy and macronutrient intake was comparable between groups, although red and white meat intake was higher in the L-RISK group (Table 1). During the intervention period, the average total energy intake decreased similarly in both groups (Table 2).

**Table 1** Patient characteristics

Characteristic	H-RISK	L-RISK
<i>n</i> (male/female)	20 (9/11)	17 (8/9)
Age (years)	53±10	55±7
Weight (kg)	103±16	106±19
BMI (kg/m <sup>2</sup> )	34.1 (31.8, 36.9)	35.4 (33.1, 38.2)
Waist circumference (cm)	112±11	114±12
Hip circumference (cm)	113±14	117±11
Systolic BP (mmHg)	135±17	135±14
Diastolic BP (mmHg)	87±14	85±10
Triacylglycerols (mmol/l)	1.50 (1.12, 2.79)	1.55 (1.35, 2.16)
Total cholesterol (mmol/l)	5.56 (4.94, 6.13)	5.28 (4.80, 5.80)
LDL-cholesterol (mmol/l)	3.45 (2.78, 3.99)	3.34 (2.80, 3.79)
HDL-cholesterol (mmol/l)	1.18 (1.06, 1.42)	1.22 (1.10, 1.40)
ALT (μkat/l)	0.58 (0.36, 0.79)	0.50 (0.41, 0.68)
AST (μkat/l)	0.43 (0.33, 0.54)	0.37 (0.29, 0.46)
Fasting glucose (mmol/l)	6.7 (5.5, 7.6)	5.9 (5.3, 6.8)
Fasting insulin (pmol/l)	104.4 (80.4, 147.1)	90.1 (72.5, 152.9)
HbA <sub>1c</sub> (mmol/mol)	45 (41, 52)	48 (45, 52)
HbA <sub>1c</sub> (%)	6.3 (5.9, 6.9)	6.5 (6.3, 6.9)
Diabetes duration (years)	0.8 (0.5, 2.1)	1.9 (0.7, 3.2)
Metformin treatment (yes/no) <sup>a</sup>	16/4	13/4
Total energy intake (kJ/day)	8,881±1,778	9,402±2,084
Protein intake (% of energy)	17.3±3.8	19.3±5.2
Carbohydrate intake (% of energy)	45.3±7.3	45.9±6.8
Fat intake (% of energy)	39.5±7.6	36.9±8.5
Cereal fibre intake (g/day)	8.7 (6.5, 11.9)	9.2 (7.6, 13.0)
Red meat intake (g/day)	77 (57, 134)	109 (70, 160)
White meat intake (g/day)	19 (6, 41)	53 (34, 80)
Plasma caffeine (μmol/l)	3.0 (1.5, 4.0)	3.4 (2.1, 4.3)
24 h urinary AR metabolite excretion (μmol/24 h)	27.8 (14.8, 48.4)	28.8 (19.3, 53.2)
Total serum amino acids (mmol/l)	2.71±0.50	2.80±0.43
BCAA (μmol/l)	494±142	518±100
Histidine (μmol/l)	42±11	48±16
IL-6 (pg/ml)	2.40 (1.92, 3.32)	2.70 (2.13, 3.94)
IL-18 (pg/ml)	300 (216, 380)	248 (192, 301)
Adiponectin (μg/ml)	3.64 (2.49, 5.42)	4.85 (3.79, 7.56)
IL-1RA (pg/ml)	594 (390, 863)	717 (413, 1,197)

Values given are means±SD or median (IQR)

<sup>a</sup>No patient received any glucose-lowering treatment other than metformin

ALT, alanine aminotransferase; AR, alkylresorcinol; AST, aspartate aminotransferase

Carbohydrate intake increased in both groups, while the decrease in fat intake and increase in protein intake was more pronounced in the L-RISK compared with the H-RISK group ( $p<0.05$ ). Cereal fibre intake from wheat and rye was comparable at baseline (Table 1) and increased by  $19.8\pm 10.0$  g/day in the L-RISK group, but decreased by  $5.0\pm 4.7$  g/day in the H-RISK group ( $p<0.0001$  vs L-RISK) during the intervention (Fig. 1a). Plasma caffeine concentration increased in the L-RISK group by  $2.7\pm 2.9$  μmol/l and decreased in the H-RISK group by  $2.5\pm 2.2$  μmol/l ( $p<0.0001$  vs L-RISK; Fig. 1b). The 24 h urinary excretion of DHBA and DHPPA (biomarkers of whole grain wheat and rye intake) slightly decreased by  $10.7\pm 12.9$  μmol/24 h in the H-RISK group and increased by  $60.7\pm 59.7$  μmol/24 h in the L-RISK group ( $p<0.0001$  vs H-RISK; Fig. 1c). Red meat intake decreased in the L-RISK group to 0 g/day, consistent with the protocol, but increased in the H-RISK group by  $99\pm 40$  g/day ( $p<0.0001$  vs L-RISK; Fig. 1d). Intake of white meat was lower in the H-RISK group at baseline (Table 1) and decreased to 0 g/day, while it increased in the L-RISK group by  $72\pm 50$  g/day ( $p<0.0001$  vs H-RISK; Fig. 1d). Total serum amino acid and BCAA concentrations did not change during the intervention, but histidine increased in the H-RISK group ( $p<0.001$  vs L-RISK; Table 2).

**Body weight and body composition** In line with the study protocol, the relative reduction in body weight during the run-in period was  $<5\%$  of screening weight in all patients. During the intervention period, the average body weight reduction (mean [95% CI]) was  $-4.8\%$  ( $-6.1\%$ ,  $-3.5\%$ ) in the H-RISK group and  $-4.6\%$  [ $-6.0\%$ ,  $-3.3\%$ ] in the L-RISK group when compared with the run-in period (Fig. 2a) with no difference between groups ( $p=0.87$ ). Waist circumference decreased in both groups ( $p=0.17$ , H-RISK vs L-RISK; Fig. 2b). In addition, the reduction of HCL in the H-RISK ( $-7.0\%$  [ $-9.6\%$ ,  $-4.5\%$ ]) and L-RISK ( $-6.7\%$  [ $-9.5\%$ ,  $-3.9\%$ ]) groups was similar (mean difference  $-0.3\%$  [ $-4.1\%$ ,  $3.5\%$ ],  $p=0.87$ ; Fig. 2c). We observed no differences between H-RISK and L-RISK groups for a change in SCAT ( $-1,553$  [ $-2,767$ ,  $-340$ ] cm<sup>3</sup> vs  $-751$  [ $-2,047$ ,  $546$ ] cm<sup>3</sup>), VAT ( $-206$  [ $-783$ ,  $371$ ] cm<sup>3</sup> vs  $-241$  [ $-856$ ,  $373$ ] cm<sup>3</sup>) and IMCL ( $-0.09\%$  [ $-0.16\%$ ,  $-0.02\%$ ] vs  $-0.02\%$  [ $-0.10\%$ ,  $0.05\%$ ]), although the within-group reduction in the H-RISK group was more pronounced for SCAT and IMCL in soleus muscle (Fig. 2d–f).

**Tissue-specific insulin sensitivity**  $R_d$  was comparable at baseline in the H-RISK and L-RISK groups ( $5.6\pm 0.5$  vs  $5.8\pm 0.4$  mg kg<sup>-1</sup> min<sup>-1</sup>,  $p=0.91$ ) and improved similarly (H-RISK vs L-RISK:  $0.8$  [0.2, 1.4] vs  $1.0$  [0.4, 1.7] mg kg<sup>-1</sup> min<sup>-1</sup>; mean difference  $-0.2$  [ $-1.1$ ,  $0.6$ ] mg kg<sup>-1</sup> min<sup>-1</sup>,  $p=0.59$ ; Fig. 3a). Non-oxidative glucose utilisation increased in the H-RISK compared with the L-RISK group (mean difference  $1.18$  [0.31, 2.05] mg kg<sup>-1</sup> min<sup>-1</sup>,  $p=0.01$ ), while glucose oxidation

**Table 2** Changes during intervention period

	H-RISK <sup>a</sup>	L-RISK <sup>a</sup>	Mean difference <sup>b</sup>
Total energy intake (kJ/day)	-1,558 (-1,849, -1,268)	-1,194 (-1,509, -879)	-365 (-795, 65)
Fat intake from energy (%)	-6.7 (-7.9, -5.5)	-8.8 (-10.2, -7.5)	2.1 (0.3, 4.0)*
Carbohydrate intake from energy (%)	4.1 (2.5, 5.7)	5.7 (3.9, 7.4)	-1.5 (-3.9, 0.9)
Protein intake from energy (%)	2.5 (2.1, 2.9)	3.1 (2.7, 3.6)	-0.7 (-1.3, -0.1)*
Triacylglycerols (mmol/l) <sup>c</sup>	-0.22 (-0.40, -0.03)	-0.26 (-0.47, -0.06)	0.05 (-0.23, 0.32)
Total cholesterol (mmol/l) <sup>c</sup>	-0.04 (-0.09, 0.02)	-0.08 (-0.14, -0.02)	0.04 (-0.04, 0.13)
LDL-cholesterol (mmol/l) <sup>c</sup>	-0.001 (-0.068, 0.067)	-0.084 (-0.161, -0.007)	0.083 (-0.019, 0.186)
HDL-cholesterol (mmol/l) <sup>c</sup>	-0.01 (-0.08, 0.05)	-0.01 (-0.14, 0.01)	0.05 (-0.05, 0.15)
ALT (μkat/l) <sup>c</sup>	-0.32 (-0.50, -0.14)	-0.31 (-0.50, -0.12)	-0.01 (-0.27, 0.25)
AST (μkat/l) <sup>c</sup>	-0.21 (-0.34, -0.08)	-0.18 (-0.32, 0.04)	-0.03 (-0.22, 0.17)
Fasting glucose (mmol/l) <sup>c</sup>	-0.13 (-0.18, -0.07)	-0.07 (-0.13, -0.002)	-0.06 (-0.15, 0.03)
Fasting insulin (pmol/l) <sup>c</sup>	-0.18 (-0.35, -0.01)	-0.20 (-0.38, -0.02)	0.02 (-0.23, 0.27)
HbA1c (mmol/mol) <sup>c</sup>	-0.07 (-0.10, -0.03)	-0.10 (-0.14, -0.06)	0.03 (-0.02, 0.09)
Total serum amino acids (mmol/l)	0.07 (-0.09, 0.24)	-0.02 (-0.21, 0.16)	0.10 (-0.15, 0.35)
BCAA (μmol/l) <sup>c</sup>	-43 (-77, -9)	-57 (-97, -18)	14 (-38, 67)
Histidine (μmol/l)	13.8 (8.6, 19.0)	-1.6 (-7.6, 4.5)	15.4 (7.3, 23.5)***

<sup>a</sup> Mean change (95% CI) within-group

<sup>b</sup> Mean difference (95% CI) of the within-group change adjusted for respective baseline values using general linear model

<sup>c</sup> Not-normally distributed data have been  $\log_e$ -transformed to achieve near-normal distribution; the mean difference of the  $\log_e$ -transformed within-group change is given

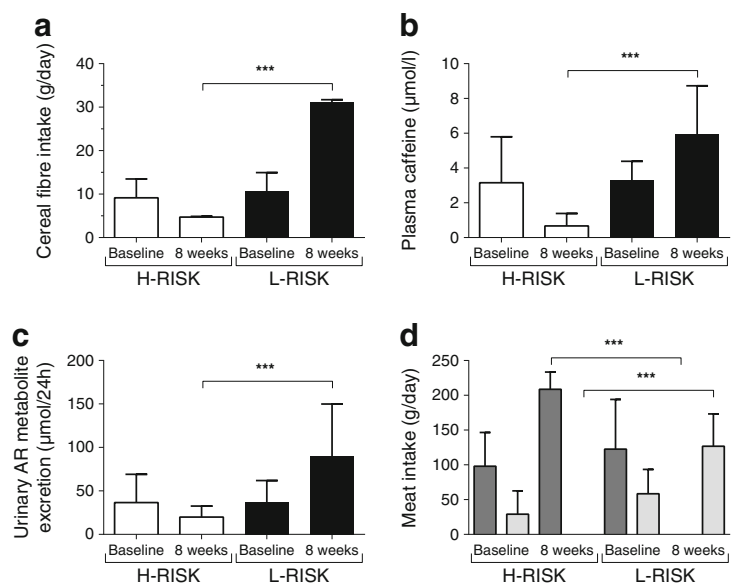
\* $p < 0.05$ , \*\*\* $p < 0.001$

ALT, alanine aminotransferase; AST, aspartate aminotransferase

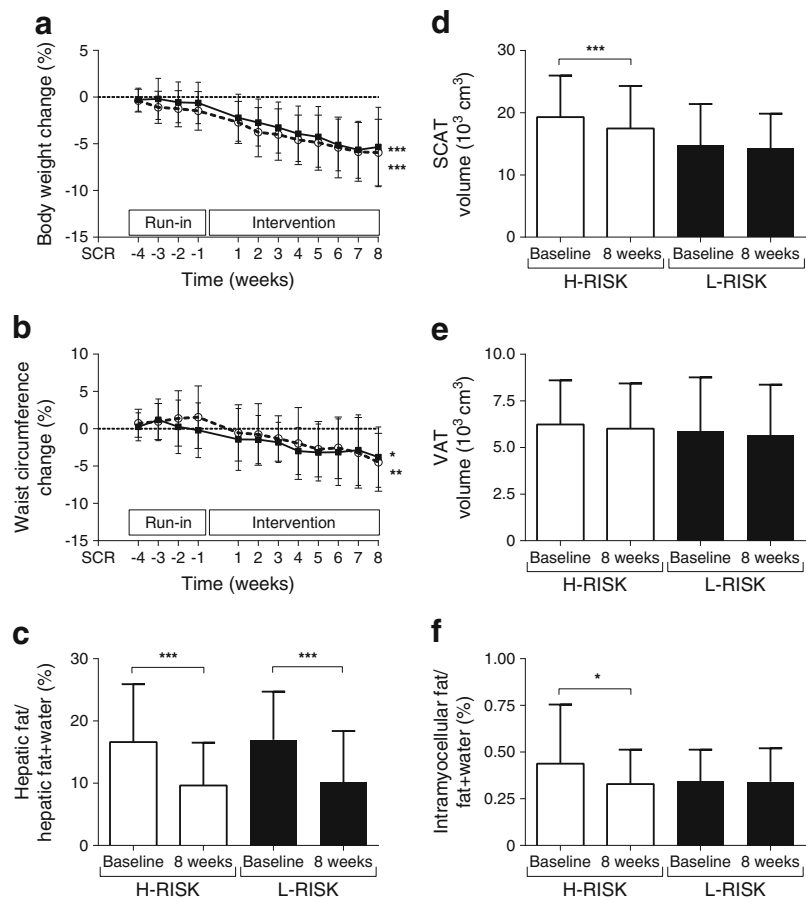
did not change (Fig. 3b). Lipid oxidation was suppressed during the clamp, but did not change in either group after the intervention (Fig. 3c). Hepatic insulin sensitivity did not differ between groups before or after the intervention (Fig. 3d). Despite comparable weight loss in both groups, improvement in  $R_d$  associated with a greater weight loss in the H-RISK group, but not in the L-RISK group (Table 3).

**Insulin secretion** Fasting insulin levels did not change during the intervention period, while fasting blood glucose concentrations declined in both groups (Table 2). AIR tended to be lower after the H-RISK diet (mean difference  $-0.05 [-0.10, 0.01]$  pmol/l,  $p = 0.09$ ; Fig. 3e) and correlated with change in  $R_d$  (Table 3), while total insulin response did not change (data not shown). DI did not change (mean difference  $-0.21 [-0.46,$

**Fig. 1** Total cereal fibre intake (a), plasma caffeine concentration (b), 24 h urinary alkylresorcinol (AR) metabolite excretion (c) and intake of red meat (dark grey) or white meat (light grey) (d) in the L-RISK and H-RISK groups during the intervention period. Total whole grain fibre, 24 h urinary AR metabolite excretion and meat intake are expressed as means  $\pm$  SD. Mean values have been compared between groups using Student's *t* test for unpaired samples with Welch's correction, and not-normally distributed values were  $\log_e$ -transformed. \*\*\* $p < 0.001$



**Fig. 2** Changes in body weight (a), waist circumference (b), hepatic fat content (c), SCAT (d), VAT (e) and intramyocellular fat content in M. soleus muscle (f) during run-in and intervention periods in L-RISK (black/filled circles with solid line) and H-RISK (white/open circles with dotted line) groups. Weight and waist circumference change is given as mean per cent change $\pm$ SD to body weight/waist circumference at screening visit (SCR). Truncal SCAT and VAT volume has been calculated from MRI images, hepatic fat content and intramyocellular fat content in soleus muscle have been assessed by  $^1\text{H}$ -magnetic resonance spectroscopy (MRS) and are expressed as means $\pm$ SD. Changes at the end of the intervention period have been compared with the end of the run-in period/baseline assessment using Student's *t* test for paired samples; not-normally distributed values were  $\log_e$ -transformed. \* $p<0.05$ , \*\* $p<0.01$ , \*\*\* $p<0.001$



0.04]AU,  $p=0.09$ ) despite a tendency towards an increase after the L-RISK diet (mean change 0.20 [0.02, 0.39]AU), but not after the H-RISK diet ( $-0.01$  [ $-0.18$ ; 0.16]AU; Fig. 3f).

**Circulating cytokines** Levels of the proinflammatory cytokine IL-18 was reduced after the L-RISK diet compared with the H-RISK diet (mean difference 0.23 [0.04, 0.42]pg/ml,  $p<0.05$ ; Table 4), while all other pro- or anti-inflammatory cytokines remained unchanged. Changes in cytokine levels did not correlate with changes in insulin sensitivity or body weight (data not shown).

## Discussion

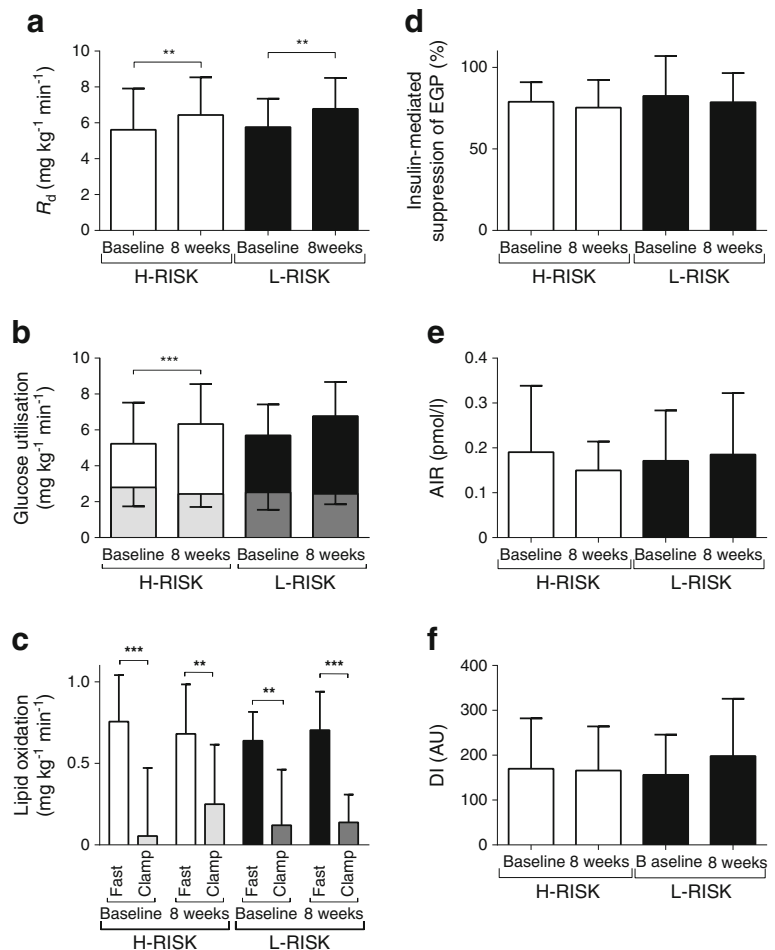
This study shows that moderate energy restriction for 8 weeks irrespective of the intake of cereal fibre from wheat and rye, or intake of coffee or red meat: (1) reduces body weight, (2) increases whole-body insulin sensitivity at least partly by enhanced non-oxidative glucose utilisation, and (3) decreases HCL without affecting hepatic insulin sensitivity in type 2 diabetes. While only the L-RISK diet tended to improve

subclinical inflammation, no differences between groups were observed in the reduction of body and muscle fat content or with respect to beta cell function.

Similar improvements in insulin resistance together with comparable weight loss for both diets suggests that energy restriction exclusively explains the findings. Energy restriction results in weight loss due to a reduction in body fat content and, to a lesser extent, in muscle mass. These effects associate with enhanced tissue-specific glucose uptake by increased insulin-stimulated Akt2 phosphorylation [27], potentially because of a lowering of cellular lipid intermediates—possibly reflected by decreased IMCL as observed after H-RISK diet—reduced subsequent protein kinase-C theta (PKC $\theta$ ) activation and increased IRS-1 tyrosine phosphorylation [28]. Furthermore, the reduction in body weight and several fat depots correlated with insulin sensitivity only in the H-RISK group. This indicates that mechanisms other than simple weight loss may be operative.

Notably, the improved insulin resistance in the L-RISK group could occur independently of any changes in body weight or inflammatory markers, as reported for increased cereal fibre intake over 72 h [8]. The mechanisms responsible could be alterations to gut microbiota improving systemic inflammation independently of weight loss [29], altered digestion and/or absorption of protein subsequently affecting

**Fig. 3** Changes in whole-body insulin sensitivity (a), oxidative (light/dark grey) and non-oxidative (black/white) glucose utilisation (b), lipid oxidation rate (c), hepatic insulin sensitivity (d), AIR (e) and DI (f) before and after the intervention period. Whole-body insulin sensitivity was measured as  $R_d$  ( $\text{mg} [\text{kg body weight}]^{-1} \text{min}^{-1}$ ); glucose utilisation and lipid oxidation rates were calculated by indirect calorimetry; hepatic insulin sensitivity was measured as insulin-mediated suppression of EGP (%); AIR (calculated during the first 10 min of IVGTT), and DI (calculated as  $\text{AIR} \times R_d$ ) are given as mean  $\pm$  SD. Changes before vs end of intervention period or during fasting and clamp conditions have been compared using Student's *t* test for paired samples; not-normally distributed values were  $\log_e$ -transformed. L-RISK group in black, H-RISK group in white.  $**p < 0.01$ ,  $***p < 0.001$



the amino acid profile [12], or altered rates of colonic fermentation with production of short-chain FA [29]. Chronic coffee

**Table 3** Determinants of improvement of whole-body insulin sensitivity

	H-RISK Correlation coefficient	L-RISK Correlation coefficient
Body weight change (%)	-0.61**	-0.05
SCAT change (%)	-0.23	-0.18
VAT change (%)	-0.21	0.22
HCL change (%)	-0.02	0.25
<sup>a</sup> IMCL change (%)	-0.23	0.46
Plasma caffeine change ( $\mu\text{mol/l}$ )	0.38	0.40
HbA1c change (%)	-0.41	0.06
AIR change ( $\text{pmol/l}$ )	-0.54*	-0.22

Values given are Pearson correlation coefficients (*r*) between the absolute change of the respective variable and change in  $R_d$  during intervention; not-normally distributed parameters were analysed after  $\log_e$ -transformation

<sup>a</sup>IMCL was measured in soleus muscle

\* $p < 0.05$ , \*\* $p < 0.01$

consumption has been reported to have similar effects [30]. On the other hand, coffee can impair insulin sensitivity, while liver steatosis and inflammation is reduced, despite identical and unchanged body weight in spontaneously diabetic mice [31]. Thus, L-RISK and H-RISK diets are likely to improve insulin sensitivity by different mechanisms, which may occur independently of the weight loss resulting from energy restriction.

Interestingly, we found a marked reduction in HCL after both diets, while hepatic insulin sensitivity was unaltered. Although HCL generally correlates with hepatic insulin resistance, skeletal muscle mainly accounts for improvements in whole-body insulin sensitivity [28]. The use of higher insulin doses to assess whole-body insulin sensitivity may have prevented the detection of small differences in hepatic insulin sensitivity. Nevertheless, improved hepatic insulin resistance and reduced HCL do not necessarily occur simultaneously [23]. This paradoxical finding could result from a selective insulin resistance in the liver when insulin fails to inhibit gluconeogenesis and EGP, while continuing to stimulate lipogenesis and HCL synthesis [32]. However, a very low-energy diet improves both hepatic insulin sensitivity and HCL [33], so we cannot rule out the possibility that the reduction in HCL



**Table 4** Changes in biomarkers of subclinical systemic inflammation

	H-RISK <sup>a</sup>	L-RISK <sup>a</sup>	Mean difference <sup>b</sup>
IL-6 (pg/ml)	−0.04 (−0.26, 0.17)	−0.24 (−0.48, 0.002)	0.19 (−0.13, 0.52)
IL-18 (pg/ml)	−0.01 (−0.14, 0.12)	−0.24 (−0.38, −0.19)	0.23 (0.04, 0.42)*
Adiponectin (μg/ml)	−0.04 (−0.20, 0.12)	0.13 (−0.05, 0.31)	−0.17 (−0.41, 0.08)
IL-1RA (pg/ml)	0.05 (−0.13, 0.22)	−0.06 (−0.25, 0.14)	0.10 (−0.16, 0.36)

<sup>a</sup> Mean change (95% CI) within-group

<sup>b</sup> Mean difference (95% CI) of the within-group change adjusted for respective baseline values using general linear model

All data have been log<sub>e</sub>-transformed to achieve near-normal distribution and the mean difference of the log<sub>e</sub>-transformed within-group change is given  
\**p*<0.05

precedes a subsequent improvement in hepatic insulin resistance.

In line with other studies [34], this trial did not detect any effect on insulin secretion although the high cereal fibre intake during the L-RISK diet may have altered the amino acid profile [8, 12, 35] and thereby might affect insulin secretion [16, 36]. The observed mild reduction in the proinflammatory response may also improve insulin secretion following an L-RISK diet [37]. Furthermore, this study found no correlation between changes in insulin sensitivity and insulin secretion in either group. Thus, any adaptive effect of insulin secretion on improved insulin sensitivity was either undetectable because of the short intervention period, or counteracted by modulation of the insulin secretory response by other mechanisms that have been triggered by increased total protein and decreased fat intakes with corresponding changes in amino acids and FA [16].

This study has the strength of being a randomised controlled trial performed in well-defined, intensively phenotyped patients, which is only possible in small-scale studies. This enabled us to uncover subtle changes resulting from the interaction between dietary composition and weight loss, which would not be apparent during short-term trials in larger cohorts. Some limitations need to be considered. First, the small but significant weight loss could mask other specific effects potentially induced by the individual dietary compositions. However, this study was designed as a pretest for larger intervention trials in obese patients with type 2 diabetes, for whom weight reduction remains the primary motivation and goal of participation in the study. Moreover, detailed analysis of metabolic changes suggests different metabolic effects despite the small variation in nutrient composition. This leads to the generation of hypotheses, which need to be addressed in detail in studies using isoenergetic diets. Second, the multimodal intervention does not allow the separation of the specific effects of each dietary component, although it allowed the relevance of the net effect to be analysed in a clinical setting. Third, effects on glycaemic control, as well as the long-term effects of potentially reduced subclinical inflammation, cannot be evaluated owing to the short duration and

small size of the study, although there was a tendency towards greater reduction in HbA<sub>1c</sub>. It is unclear whether a successful lifestyle intervention can reduce negative macrovascular outcomes, a major cause of mortality [7].

In conclusion, an 8 week low-energy diet reduces body weight, HCL and whole-body insulin resistance irrespective of the dietary composition in overt type 2 diabetes. Surprisingly, this short-term intervention failed to confirm previous reports of pronounced effects of high cereal wheat and rye fibre or coffee intake during phases of active weight loss, but uncovered potentially different mechanisms for both diets.

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**Duality of interest** All authors declare that there is no duality of interest associated with this manuscript.

**Contribution statement** MR designed the study and headed the clinical experiments. BN, LZ, AB, SK, PJN, AKF, JBG, GP, IE and RL researched the data. BN and LZ wrote the first draft of the manuscript, BN interpreted the statistical analysis of the data and coordinated the inclusion of specific sections as outlined. AB, PJN, CH, IE and RL conducted and wrote aspects of the laboratory analyses. GP calculated indices of beta cell function and wrote the respective sections. All the authors contributed substantially to aspects of study design or the acquisition of data, contributed to drafting of the article or revised it critically for

important intellectual content and gave final approval to the version to be published. MR is responsible for the integrity of the work as a whole.

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