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Postprandial changes in cardiometabolic disease risk in young Chinese men following isocaloric high or low protein diets, stratified by either high or low meal frequency - a randomized controlled crossover trial

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Abstract

Background: Cardio-Metabolic Disease (CMD) is the leading cause of death globally and particularly in Asia. Postprandial elevation of glycaemia, insulinaemia, triglyceridaemia are associated with an increased risk of CMD. While studies have shown that higher protein intake or increased meal frequency may benefit postprandial metabolism, their combined effect has rarely been investigated using composite mixed meals. We therefore examined the combined effects of increasing meal frequency (2-large vs 6-smaller meals), with high or low-protein (40 % vs 10 % energy from protein respectively) isocaloric mixed meals on a range of postprandial CMD risk markers.

Methods: In a randomized crossover study, 10 healthy Chinese males (Age: 29 ± 7 years; BMI: 21.9 ± 1.7 kg/m²) underwent 4 dietary treatments: CON-2 (2 large Low-Protein meals), CON-6 (6 Small Low-Protein meals), PRO-2 (2 Large High-Protein meals) and PRO-6 (6 Small High-Protein meals). Subjects wore a continuous glucose monitor (CGM) and venous blood samples were obtained at baseline and at regular intervals for 8.5 h to monitor postprandial changes in glucose, insulin, triglycerides and high sensitivity C-reactive protein (hsCRP). Blood pressure was measured at regular intervals pre- and post- meal consumption. Urine was collected to measure excretion of creatinine and F_2 -isoprostanes and its metabolites over the 8.5 h postprandial period.

Results: The high-protein meals, irrespective of meal frequency were beneficial for glycaemic health since glucose incremental area under the curve (iAUC) for PRO-2 ($185 \pm 166 \text{ mmol.min.L}^{-1}$) and PRO-6 ($214 \pm 188 \text{ mmol.min.L}^{-1}$) were 66 and 60 % lower respectively (both p < 0.05), compared with CON-2 ($536 \pm 290 \text{ mmol.min.L}^{-1}$). The iAUC for insulin was the lowest for PRO-6 ($13.7 \pm 7.1 \text{ U.min.L}^{-1}$) as compared with CON-2 ($28.4 \pm 15.6 \text{ U.min.L}^{-1}$), p < 0.001. There were no significant differences in postprandial responses in other measurements between the dietary treatments. (Continued on next page)

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Conclusions: The consumption of composite meals with higher protein content, irrespective of meal frequency appears to be beneficial for postprandial glycemic and insulinemic responses in young, healthy Chinese males. Implications of this study may be useful in the Asian context where the consumption of high glycemic index, carbohydrate meals is prevalent.

Trial registration: NCT02529228.

Keywords: High protein, Meal frequency, Cardiometabolic disease risk

Background

Increases in postprandial glucose, insulin and lipids have been linked to increased cardiometabolic disease (CMD) risk [1-3], even though the majority of epidemiological and intervention trials have relied on measuring these parameters in the fasted state. Given that most individuals spend majority of their waking hours in the postprandial state, eating and digesting food, fasting measurements alone may not, in its entirety, represent the metabolic processes indicative of CMD risk [4]. Moreover, Asians are at a greater risk of CMD than other populations such as the Caucasians [5] and several studies reported Asians having higher levels of fasting and postprandial glucose and insulin responses [6, 7], higher levels of HbA_{1C} [8] and increased type 2 diabetes risk compared with Caucasians [9, 10]. Asians also have a lower metabolic capacity than Caucasians to handle an excess dietary load [11]. Therefore alterations in meal composition and patterns are warranted in order to improve postprandial metabolism amongst Asians.

Increasing meal frequency has long been suggested by several authors to improve metabolic health and weight control [12-14]. It has been shown that eating smaller and frequent meals seems to better maintain blood glucose, blood lipid and incretin hormone concentrations, with lower variations throughout the day [14, 15]. However, the evidence thus far relating to the effects of meal frequency alone on glycaemic response remain equivocal with some studies showing a greater postprandial glucose responses with increased meal frequency [16, 17], while other studies reporting lower blood glucose responses with higher meal frequencies in non-diabetics [14] as well in diabetics [18, 19]. There have also been studies reporting no difference in postprandial glycaemia with increased meal frequency [20, 21]. However, the compositions of the meals were rather heterogenous between these studies and this may explain the disparate findings observed between studies investigating the effect of meal frequency alone.

It is widely known that Asian diets are particularly rich in refined carbohydrates which may exacerbate the risk of type 2 diabetes and CVD [22]. A recent review has highlighted that the prevalence of diabetes worldwide also parallels the increases in carbohydrate intake over the same period [23]. Therefore, modulating dietary carbohydrate content and composition has been suggested to be an effective way of decreasing glycaemia [24, 25]. Recent evidence suggests that consuming higher amounts of protein can lead to a lower CMD risk [26–29]. In particular, greater protein to carbohydrate ratios in the diet have been shown result in better glycaemic control [29–32].

Therefore the primary aim of this study was to investigate whether altering protein composition with or without increasing meal frequency can improve postprandial glucose and/or insulin response in young, healthy, Chinese males. Since postprandial glycemic control is also associated with changes in triglyceride, hs-CRP and markers of oxidative stress [2], as secondary outcomes, we have also measured postprandial changes in these markers in plasma as a result of the acute dietary modulations.

Methods

Participants

This randomized control trial was conducted at the Clinical Nutrition Research Centre (CNRC) within the Singapore Institute of Clinical Sciences (SICS), Agency of Science Technology and Research (A*STAR), Singapore, between October 2014 to July 2015. The study received ethical approval from the Domain Specific Review Board of the National Healthcare Group in Singapore (NHG DSRB Reference No. 2014/01054). All subjects gave their written informed consent before participating in the study. Research procedures and trial protocols were followed in accordance to good clinical practice (GCP) guidelines and with the ethical standards in concordance to the Declaration of Helsinki, 1983. This trial is registered within clinicaltrials.gov under trial registration no. NCT02529228.

Healthy Chinese males between the ages of 21-40 years were recruited for this trial. The exclusion criteria for the study were: 1) smokers, 2) BMI > 25 kg/m^2 ; 3) fasting blood glucose > 5.5 mmol/l; 4) blood pressure > 130/90 mmHg; 5) body weight change > 3 kg within the previous 2 months; 6) having taken part in any dietary intervention trials or having followed a special dietary

habit (e.g., vegetarian, Atkins, weight reducing diets) within 3 months of study participation; 7) suffering from any endocrine (thyroid dysfunction) or metabolic (dyslipidemia, diabetes, metabolic syndrome) disorders; and 8) alcohol use > 4 days per week. Volunteers who met the initial inclusion/exclusion criteria at screening were invited to take part in the study. A total of 10 volunteers were recruited and there were no withdrawals from the study. Participant characteristics are summarized in Table 1.

Design

The study was a randomized, crossover trial with four acute dietary conditions, comparing postprandial effects of varying two parameters of: 1) meal frequency and 2) dietary protein composition, in a 2-by-2 factorial design. Participants were studied under 4 separate 1-day isocaloric nutritional interventions: CON-2 (2 large Low-Protein meals :10 % Protein, 57 % Carbohydrate, 38 % Fat); CON-6 (6 Small Low-Protein meals with identical nutrient composition as CON-2); PRO-2 (2 Large High-Protein meals :38 % Protein, 30 % Carbohydrate, 32 % Fat); PRO-6 (6 Small High-Protein meals with identical nutrient composition as PRO-2). The order of the sessions was randomized with a washout period of at least 3 days between interventions. Participants refrained from alcohol consumption and vigorous exercise 48-h before each trial session. The evening before each session, participants came to the clinical suite for the insertion of a continuous glucose monitor (CGM) (Medtronics, USA) under the abdominal subcutaneous adipose tissue by a trained researcher and also for the consumption of a standardized evening meal. Participants fasted for 12-h following their evening meal up to the start of each session. The CGM measured interstitial glucose levels in 5 min intervals for the entire duration of the sessions.

Study protocol

The schematic of the study protocol is shown in Fig. 1. On the morning of each test day, the participants arrived

Table 1 Participant's characteristics

		Range	
Characteristic	Mean (SD)	Min	Max
Age (Years)	28.8 (6.6)	22	40
Height (m)	1.76 (0.04)	1.71	1.84
Body Mass (kg)	68.2 (7.8)	58.1	83
Body Mass Index (kg/m²)	21.9 (1.7)	19.5	24.5
Umbilical Waist Circumference (cm)	80.5 (7.9)	70	92
Body Fat %	17.0 (4.7)	8.9	22.6
Systolic Blood Pressure (mmHg)	120.4 (5.7)	113	130
Diastolic Blood Pressure (mmHg)	76.1 (6.4)	65	85

in a fasting state. Throughout each test session, lasting 8.5 h during the day (0800-1630 h), participants stayed within a clinical suite and predominantly remained in a seated posture doing passive work (i.e., watching TV, using computers or reading quietly). The temperature of the suite was kept under thermo-neutral conditions (airconditioned to keep within 22-24 degree Celsius ((°C)) throughout the entire study period. At the start of each test day, an indwelling venous cannula was inserted into the participant's antecubital vein by a trained nurse for the collection of baseline and subsequent regular blood sampling. This cannula was kept patent by flushing with 3 ml non-heparinized saline. After the fasting baseline blood sample was taken, participants consumed high or low protein isocaloric mixed meals in either a 2-meal or 6-meal frequency pattern over the subsequent 8.5 h period. The total energy content of the meals (approximately 2000 kcals) were designed to provide a small caloric surplus based on two-thirds (from breakfast and lunch) of the daily energy requirements for an average sedentary male (Singapore National Nutrition Survey, 2010) [33]. This would reflect the typical overconsumption of calories in modern day society. For the 2-meal trials (CON-2 and PRO-2), test meals consisted of 1 large breakfast meal and 1 large lunch meal, whereas for the 6-meal trials (CON-6 and PRO-6), the breakfast and lunch were each divided into 3 equal, isocaloricl portions. Meal times were scheduled at equidistant time points (Fig. 1); that is, during the 2-meal sessions, meals were provided at 4.5-h intervals (0800 and 1230-h), and during the 6-meal sessions, meals were served at 1.5-h intervals (0800, 0930, 1100, 1230, 1400, and 1530). Two of the meals during the 6-meal trials were provided at the same time as the meals during the 2-meal conditions (i.e., 0800 and 1230). Postprandial blood samples and blood pressure measurements were taken at 1 to 1.5-h intervals. Participants also collected their entire urine passed during the 8.5-h postprandial period for the analyses of urinary F₂-isoprostanes, its metabolites and creatinine excretion.

The energy and the macronutrient compositions, along with the food amounts and constituents and the estimated glycemic indices (GIs) of the test meals are detailed in Table 2. The estimated GIs of each composite meal were calculated according to the method by Hatonen KA et al. [34]. The GI of individual foods used for the study were derived using published GI data of foods same or similar foods [35, 36]. Based on these calculations for the low-protein CON-2/CON-6 meals, the overall GI was estimated to be 62, whereas, for the high-protein PRO-2/PRO-6 meals, the overall GI was estimated to be 31. Volunteers were required to consume all of their meals as provided and no other food or drinks were permitted. Researcher surveillance of study participants during meal

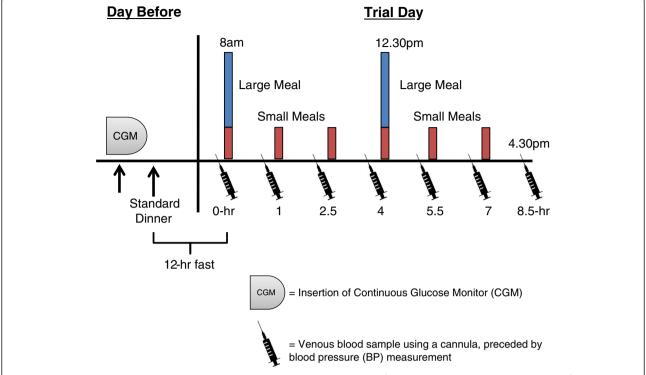


Fig. 1 Study Protocol: For each dietary condition, participants came to the clinical trial facilities on 2 consecutive days. The day before the main trial day was for the insertion of a continuous glucose monitor and the consumption of a standardized evening meal. On the main trial day, BP and venous blood samples were collected at regular intervals, as shown, for up to 8.5 h following the first meal of each dietary condition

Table 2 Nutritional information of test-meals

CON-2, CON-6 dietary conditions

Meal Constituents: Breakfast – White Bread (171 g), Margarine (30 g), Malted Cocoa Drink (375 g)

Lunch - Chicken Soup Noodles (225 g), Cashew Nuts (30 g)

	Breakfast	Lunch	Total
Energy (kcal)	889	1127	2016
Protein (g)	23 g (10 %)	26 g (9 %)	49 (10 %)
Carbohydrate (g)	130 g (57 %)	134 g (42 %)	264 (52 %)
Fat (g)	32 g (33 %)	53 g (49 %)	85 (38 %)
Estimated Glycaemic Index	73.4	50.3	61.7

PRO-2, PRO-6 dietary conditions

Meal Constituents: Breakfast – Milk (300 g), Ice-Cream (150 g), Whey powder (91.2 g), Cashew Nuts (30 g), Lunch – Braised Chicken (330 g), Prawn Dumplings (120 g), Malted Cocoa Drink (375 g)

	Breakfast	Lunch	Total
Energy (kcal)	910	998	1908
Protein (g)	89 g (39 %)	90 g (36 %)	179 (38 %)
Carbohydrate (g)	66 g (30 %)	75 g (31 %)	141 (30 %)
Fat (g)	31 g (31 %)	37 (33 %)	68 (32 %)
Estimated Glycaemic Index	37.1	24.9	30.6

Macronutrient information obtained from nutrition labels on constituent foods

times and collection of empty food containers ensured dietary compliance. Participants were asked to consume each meal within 15-mins of serving. There were no complaints of any gastrointestinal or other discomforts after consuming the test meals. Plain drinking water was provided ad libitum during first test day and this volume was consumed for the remaining test sessions. All blood samples were collected in K-EDTA blood vacutainer tubes (Becton Dickinson Diagnostics, USA) and were centrifuged at 4500 rpm for 10 mins at 4 °C. The plasma so obtained was immediately stored at -80 °C until batch analysis of separate analytes at the completion of the clinical trial. Subcutaneous interstitial fluid glucose readings were obtained every 5-min for the entire 8.5 h postprandial period using the CGM. Appropriate calibrations with capillary blood glucose were also made for this purpose. Plasma samples were assayed for insulin, triacylglycerol (TAG), and high-sensitivity C-reactive protein (hs-CRP) using an immunochemistry analyzer COBAS e411 (Roche, HITACHI, USA). Urinary F₂-Isoprostanes (5-F_{2t}-isoprostane and 15-F_{2t}-isoprostane) and its metabolites (2,3-dinor-15-F_{2t}-isoprostane and 2,3-dinor-5,6-dihydro-15-F_{2t}-isoprostane) were measured using a LC-MS/ MS method [37, 38] and creatinine was measured using the COBAS e411 analyser (Roche, Hitachi, USA).

Statistical analyses

The initial sample size was calculated to detect a difference between dietary conditions in the primary outcome measures, being both postprandial glycemia and insulinemia, using primary data obtained from a similar dietary intervention by Holmstrup M et al. [17]. A sample size of 9 was estimated to detect a difference with a 80 % power and a significance (α) level set at 0.05 using a crossover design. The incremental area under the curve (iAUC) for postprandial CGMS glucose, plasma insulin, triglycerides and hs-CRP as well as blood pressure (BP) were calculated using the trapezoidal method [39] and compared between dietary treatments using a one-way ANOVA. The concentrations of glucose, insulin, TAG, hsCRP and BP were compared between treatments and over time using two-way repeated measures ANOVA. Where appropriate, Bonferroni post hoc comparisons were used. Data are presented as means (standard deviations, SD) unless otherwise defined. Significance was set at p < 0.05. Statistical analyses were carried out using Statistical Package for the Social Sciences statistical software, version 23.0 (IBM Inc.).

Results

Figure 2 displays the 8.5 h postprandial response, depicted as mean incremental areas under the curve (iAUCs) for glucose, insulin, triacylglycerol (TAG), systolic and diastolic blood pressure for each of the 4 dietary conditions. One-way repeated measures ANOVA for the mean iAUCs showed significant differences overall, between the dietary treatments for glucose (P = .003, $\eta^2 = .399$), insulin (P = .001, $\eta^2 = .433$) and TAG (P = .026, $\eta^2 = .287$). Table 3 summarizes the pairwise comparisons of the means between dietary conditions across time as well as the differences in the iAUCs for each dietary treatment and their respective effect size (d) as compared with CON-2.

Postprandial glucose, insulin and triglycerides

For CGMS gluclose, the iAUCs for PRO 2 and PRO 6 were significantly lower than that of both CON 2 and CON 6 (all P < 0.05) when compared using least significant differences. The glucose iAUCs for CON-6, PRO-2 and PRO-6 were 11, 66 and 60 % lower respectively, as compared to CON-2. For plasma insulin, the iAUC for PRO 2 and PRO 6 were significantly lower than CON 2 (P < 0.05) when compared using least significant differences. The insulin iAUC for CON-6, PRO-2 and PRO-6 were 30, 36 and 52 % lower respectively, compared to CON-2. For plasma TAG, the iAUC for CON-6 was 24 % lower respectively compared to the CON-2 trial. The iAUC for PRO-2 and PRO-6 were slightly higher (10 and 16 % respectively), compared to the CON-2 although these differences were not statistically significant.

Postprandial hs-CRP, blood pressure

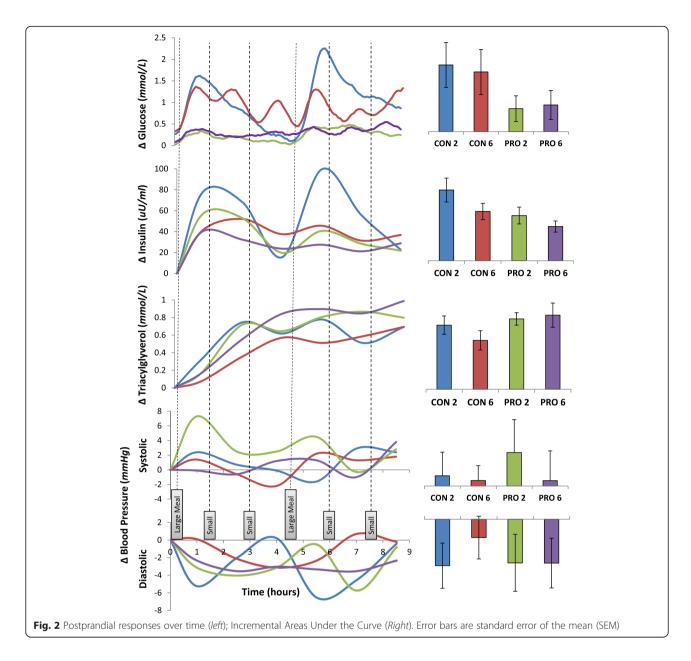
Plasma hs-CRP levels were measured at 0, 1, 4, 5.5 and 8.5-h to obtain an indication of differences in postprandial inflammatory response between the dietary conditions. No statistical significant differences were found. These time points for measurements were selected based on the highest and the lowest fluctuations of the glucose measurements (CGM data) from baseline in order to assess whether postprandial glycaemic variation was associated with changes in hs-CRP levels. There were also no significant differences in systolic or diastolic blood pressure between different dietary conditions.

Urinary total F2-isoprostanes and creatinine excretion

The mean total urinary excretions of 5-F_{2t}-isoprostane, the sum of 15-F_{2t}-isoprostane metabolites (2,3-dinor-15-F_{2t}-isoprostane and 2,3-dinor-5,6-dihydro-15-F_{2t}-isoprostane) and creatinine over the entire 8.5 h postprandial period for each dietary condition is shown in Table 4. Surprisingly, 15-F_{2t}-isoprostane (parent compound) being the well-noted biomarker of oxidative stress in vivo [40] was found in trace amounts or not detectable in the urine samples of the subjects. Urinary creatinine excretion of 15-F_{2t}-isoprostane metabolites during PRO-2 and PRO-6 were significantly higher than CON-2 (p < 0.05). There being differences in creatinine excretion between the different dietary conditions, excretion of F2-isoprostanes or its metabolites could not be adjusted for creatinine excretion. We did not find any statistical difference between 5-F_{2t}-isoprostane between the different dietary conditions, although the sum of 15-F_{2t}-isoprostane metabolites were significantly higher in PRO-2 and PRO-6 than CON-2 (p < 0.05).

Discussion

To our best knowledge, this is first study to have investigated the combined effects of increasing meal frequency and/or protein composition on range of postprandial cardiometabolic markers in an Asian population. Moreover, in most studies investigating influence of high or low protein with/without changes in meal frequency on postprandial metabolic responses used either supplemental meal replacement powders or liquid diets [17, 41–43], whereas, we have used mixed-composite meals which are better representation of a regular diet. This is also the first study using the continuous glucose monitoring (CGM) to determine the combined effects of increased meal frequency with or without higher protein content to determine PPG response. This having collected glucose data every 5 min over the 8.5 h postprandial period should be more accurate than using capillary/venous glucose samples, which are generally collected at greater sampling intervals. From the previous literature, it appears though that the beneficial effects of increasing meal frequency in



improving PPG might be more relevant in those with impaired glucose tolerance (IGT) and diabetes than in healthy individuals [18–21]. In our study we have also demonstrated that increasing meal frequency *per se* did not have a major influence on postprandial glucose control. However, as expected, increasing protein content of the meals (from 10 % of energy to ~40 % of energy) led to significant reductions in PPG by about 60 %, as well as maintained it within 0.5 mmol/l of baseline, irrespective of changes in meal frequency. This was likely to be due to the lower glycemic index (GI) as well as the lower glycemic load (GL) of the high-protein (PRO-2/PRO-6) meals, as compared to the low-protein (CON-2/CON-6) meals. Therefore for the

first time, we have been able to show that the GI/GL of the diet seems more important in controlling PPG irrespective of meal frequency. Increased dietary protein intake have been shown to reduce postprandial glycaemic response in several studies both acutely [44, 45] and chronically [32]. On the other hand, postprandial insulinaemic (PPI) responses were lower with increasing meal frequency, independently of meal composition. For example, CON-6 dietary condition had approximately 30 % lower postprandial iAUC of insulin compared with CON-2 dietary condition, whereas, PRO-6 dietary condition had 24 % lower iAUC compared with PRO-2 and these may be physiologically relevant. Similar observations of lower postprandial

Table 3 Postprandial responses across time

Time (h)	0	1	2.5	4	5.5	7	8.5	iAUC	Overall p (Diet)	D
Glucose (mmol/L)			-							
CON-2	4.7 ± 0.6	6.1 ± 1.1^{a}	5.4 ± 0.6^{a}	4.8 ± 0.8^{a}	6.8 ± 1.0^{a}	5.7 ± 0.5	5.4 ± 0.8	536 ± 290	.003	
CON-6	5.0 ± 0.8	6.2 ± 1.1	5.9 ± 0.5^{b}	5.8 ± 0.6^{b}	6.1 ± 0.5^{a}	5.7 ± 0.7^{a}	6.1 ± 1.1	480 ± 292		- 0.19
PRO-2	4.8 ± 0.4	5.2 ± 0.4^{b}	5.0 ± 0.4^{a}	4.9 ± 0.4^{a}	5.2 ± 0.6^{b}	5.2 ± 0.7	5.0 ± 0.8	185 ± 166		- 1.21
PRO-6	4.7 ± 0.4	5.2 ± 0.4	5.0 ± 0.4^{a}	5.1 ± 0.4^{a}	5.1 ± 0.5^{b}	5.1 ± 0.6^{b}	5.2 ± 0.7	214 ± 188		- 1.11
Insulin (µU/ml)										
CON-2	6.1 ± 3.3	84.3 ± 45.3^{a}	74.0 ± 43.6^{a}	21.6 ± 16.8	105.6 ± 61.0^{a}	62.0 ± 46.4	28.6 ± 17.9	28.4 ± 15.6 a	.001	
CON-6	8.1 ± 5.0	49.7 ± 19.5	60.1 ± 31.9	45.6 ± 29.2^{a}	53.7 ± 33.2	39.5 ± 22.0	44.8 ± 24.9	19.8 ± 10.3		- 0.57
PRO-2	6.0 ± 4.2	63.3 ± 41.2	58.7 ± 36.2	25.6 ± 18.5^{b}	46.6 ± 22.5	34.4 ± 24.9	27.8 ± 25.9	18.1 ± 10.6		- 0.68
PRO-6	7.1 ± 3.6	47.6 ± 28.4^{b}	39.4 ± 21.3^{b}	30.8 ± 16.3	34.6 ± 12.6^{b}	28.3 ± 15.4	36.0 ± 25.0	13.7 ± 7.1^{b}		- 0.97
TAG (mmol/L)										
CON-2	0.9 ± 0.3	1.2 ± 0.4	1.6 ± 0.6	1.5 ± 0.6	1.6 ± 0.5	1.4 ± 0.6	1.6 ± 0.4	294 ± 133	.026	
CON-6	1.1 ± 0.5	1.1 ± 0.5	1.4 ± 0.5	1.6 ± 0.6	1.6 ± 0.6	1.6 ± 0.5	1.8 ± 0.6	224 ± 139		- 0.52
PRO-2	0.9 ± 0.3	1.1 ± 0.3	1.6 ± 0.6	1.5 ± 0.4	1.7 ± 0.6	1.7 ± 0.5	1.7 ± 0.6	322 ± 91		0.21
PRO-6	1.0 ± 0.5	1.2 ± 0.5	1.6 ± 0.6	1.9 ± 0.8	1.9 ± 0.7	1.9 ± 0.7	2.0 ± 0.7	340 ± 175		0.34
hs-CRP (mg/L)										
CON-2	0.74 ± 1.41	0.79 ± 1.44	-	0.74 ± 1.44	0.69 ± 1.39	-	0.72 ± 1.40	-0.5 ± 90	.17	
CON-6	0.80 ± 1.04	0.81 ± 1.00	-	0.73 ± 0.87	0.66 ± 0.81	-	0.55 ± 0.70	-50 ± 115		- 0.55
PRO-2	0.64 ± 0.86	0.69 ± 0.84	-	0.70 ± 0.82	0.65 ± 0.77	-	0.63 ± 0.71	12 ± 71		0.14
PRO-6	0.12 ± 0.18	0.13 ± 0.21	-	0.15 ± 0.23	0.18 ± 0.24	-	0.17 ± 0.25	19 ± 35		0.22
Systolic BP (mmHg)										
CON-2	120 ± 10	122 ± 10	121 ± 11	120 ± 7	118 ± 8	123 ± 9	122 ± 9	459 ± 3343	.84	
CON-6	118 ± 6	120 ± 6	118 ± 9	116 ± 6	121 ± 8	120 ± 6	120 ± 6	249 ± 2101		-0.06
PRO-2	116 ± 8	123 ± 8	118 ± 11	118 ± 12	120 ± 13	116 ± 6	119±8	1501 ± 4658		0.31
PRO-6	119 ± 8	119 ± 7	118 ± 12	120 ± 9	120 ± 7	118 ± 11	123 ± 9	245 ± 4209		-0.06
Diastolic BP (mmHg)										
CON-2	78 ± 8	73 ± 8	76 ± 9	78 ± 7	71 ± 8	73 ± 8	77 ± 5	-1560 ± 2399	.86	
CON-6	76 ± 6	76 ± 9	74 ± 7	73 ± 6	73 ± 4	76 ± 7	75 ± 6	-614 ± 2272		-0.39
PRO-2	78 ± 9	75 ± 5	74 ± 7	75 ± 9	77 ± 14	72 ± 8	77 ± 8	-1466 ± 3024		-0.04
PRO-6	79 ± 6	77 ± 10	76 ± 6	76 ± 6	76 ± 8	76 ± 8	77 ± 6	-1473 ± 2642		-0.04

All values are mean \pm standard deviation. n=10. iAUC: Incremental Area under the Curve for 8.5 h period. D: Effect size Cohen's d, Glass's Δ correction. Figures with differing superscript letters are significantly different: p < 0.05 (after Bonferroni correction)

insulin response with increasing meal frequency have also been observed by several other investigators [16, 18, 21]. Moreover, the combined effects of higher protein intake with increased meal frequency gave rise to the lowest iAUC for postprandial insulin compared with the three other dietary conditions. While it is well known that dietary protein is insulinogenic [46], in this study we have shown that when the same quantity and quality of protein in diet is separated into smaller boluses through increasing meal frequency (PRO-6), the total PPI response was much lower than having the same meal as larger boluses (PRO-2). At the same time, we did not find any major difference in the postprandial glycaemic response between

PRO-2 and PRO-6 dietary conditions, which may indirectly indicate that the insulin sensitivity in conditions of high protein and high meal frequency (PRO-6) might be greater than PRO-2. Although insulin sensitivity was not directly tested in our study, similar observations were made in other longer term studies, where a moderate intake of 20–25 % protein had given rise to better insulin sensitivity than a higher dietary protein (25–30 % protein as energy) intake [47]. It should also be noted that different protein types (e.g., whey, casein, soy) can give rise to variability in postprandial insulinaemic response [45] and this has to be taken into consideration before arriving at any meaningful conclusion regarding dietary protein

Table 4 Total Urinary Excretions of 5-F_{2t}-isoprostanes, 15-F_{2t}-isoprostanes Metabolites and Creatinine over 8.5 h postprandial period

	CON-2 Mean (SD)	CON-6 Mean (SD)	PRO-2 Mean (SD)	PRO-6 Mean (SD)	Overall p (ANOVA)
5-F _{2t} -isoprostanes (μg)	7.8 (13.84)	3.4 (3.96)	2.0 (2.20)	3.2 (4.94)	0.268
Sum of 15-F _{2t} -isoprostane metabolites (µg)	2.4 (1.50)	3.2 (2.89)	4.7 (2.53)	3.4 (1.48)	0.059
Creatinine (mmol)	4.2 (1.0)	5.8 (1.8)	8.4 (4.7)	7.6 (2.8)	0.004

content and insulin sensitivity. Postprandial hypertrigly-ceridaemia is associated with increased CVD and type 2 diabetes risk [48, 49]. Therefore the 24 % lower mean iAUC of postprandial triglyceridaemia (TAG) with increasing meal frequency in the low protein-high carbohydrate (CON-6) condition as compared with CON-2 condition is likely to be physiologically relevant, even though this was not statistically significant due to the large inter-individual variability.

Given that the postprandial state is also associated with an increased oxidative stress and/or inflammation [48, 50], we investigated whether high and low protein diets, in combination with high and low meal frequency patterns can modify markers of postprandial oxidative stress or inflammation. We did not find any significant influence of altering meal frequency per se on these parameters and these findings are similar to other studies where no differences in oxidative stress markers upon changing meal frequency were observed [51]. However, we found that while the excretion 15-F_{2t}-isoprostane metabolites (2,3-dinor-15-F_{2t}-isoprostane and 2,3-dinor-5,6-dihydro-15-F_{2t}-isoprostane) increased with higher protein intake (PRO-2 and PRO-6 combined vs CON-2 and CON-6 combined, p < 0.05), the excretion of 5-F_{2t}isoprostane was in fact lower with higher protein intakes (74 and 59 % lower in PRO-2 and PRO-6 diets respectively as compared to the CON-2 diet), although none of these differences were statistically significant. It should be noted that the urine samples of the subjects in this study had only trace concentrations of 15-F_{2t}-isoprostane compared to 5-F_{2t}-isoprostane; similar findings were reported in other studies [52]. This, in addition to the increased renal clearance, as evidenced by greater creatinine excretion with higher protein diets, explained the higher excretion of 15-F_{2t}-isoprostane metabolites in PRO-2 and PRO-6 diets compared with the low protein (CON-2 and CON-6) diets. We have also compared the ratios of F₂-isoprostane to creatinine excretion for the separate of F₂-isoprostane species between the different dietary conditions, although there were no statistical differences. Thus, increased excretion of 15-F_{2t}-isoprostane metabolites in this study with increased protein intake may not necessarily indicate increased oxidative stress with the higher protein diets, particularly given that the 5-F_{2t}-isoprostane excretion was much lower and that 5-F_{2t}-isoprostane could well be better marker of in vivo oxidative stress under some circumstances [53]. The higher creatinine excretion with increasing protein intake (PRO-2 and PRO-6 vs CON-2 and CON-6) was similar to other observations which could be due to a higher glomerular filtration rate [54, 55]. However, in healthy volunteers without any kidney disorder, there is no evidence to show that having a high protein diet is detrimental to kidney health [56, 57]. While there are some indications that dietary carbohydrate load is associated with increased CRP levels [58], we did not find any effect of varying meal composition in our study on postprandial hsCRP levels. While postprandial hsCRP levels was unaltered with the acute dietary modulations, it remains to be investigated how similar dietary changes may affect hsCRP in the longer term. Finally, one of the major limitations of the study was that the postprandial responses were tested after dietary manipulation for one day only. How longer term dietary manipulations will affect these responses remain to be investigated.

Conclusions

In contrast to previous studies that have used meal replacement liquid based formulas to undertake similar investigations, we have for the first time demonstrated that it is feasible to use solid composite mixed meals to study combined effect of high protein content and increased meal frequency on markers of postprandial cardiometabolic risk. As expected, we found that increasing protein content of meals decreased PPG response. Contrary to the commonly held view that increasing meal frequency attenuates postprandial glycaemia, our results seem to indicate that the overall protein content and the consequent glycemic index and/or glycemic load of the meals may be more important than meal frequency alone. There were some indications that increasing meal frequency may benefit postprandial insulinemia. Increasing meal frequency and/or protein composition did not affect the other postprandial markers of cardiometabolic risk measured in this study. However, these remain to be tested in larger populations using longer term trials. The findings of this study will have important implications in Asians who already consume diets rich in carbohydrates with high glycaemic index. In order to improve their cardiometabolic risk, Asians may increase the protein content of the meal, irrespective of any need to change their meal frequency.

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Abbreviations

ANOVA: Analysis of Variance; A*STAR: Agency for Science Technology and Research, Singapore; CGM: Continuous Glucose Monitor; CON-2: 2 large Low-Protein iso-caloric meals; CON-6: 6 small Low-Protein iso-caloric meals; CMD: Cardiometabolic Disease; CNRC: Clinical Nutrition Research Centre; CVD: Cardiovascular Disease; hs-CRP: High-sensitivity C-reactive Protein; iAUC: incremental area under the curve; PPG: Postprandial glucose; PRO-2: 2 large High-protein iso-caloric meals; PRO-6: 6 small High-protein iso-caloric meals; SD: Standard Deviation; TAG: Triacylglycerol.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

The authors' contributions are as follows: AM, SH and CJH conceived and designed the study. AM carried out the study including statistical analyses of the data. AM and SH wrote major parts of the manuscript. All authors have been involved in the critical analysis of the data and approved the final manuscript.

Acknowledgements

We thank all research volunteers for taking part in this study. We also thank Dr Siew Ling (Agnes) Tey, Mr Zhien Joseph Lim, Mr Kevin Wei Jie Tan, Miss Yu Chin (Rina) Quek and Mr Kin Sum Leung for their technical assistance.

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Received: 7 October 2015 Accepted: 24 February 2016 Published online: 15 March 2016

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