



**COFFEE AND ITS CONSTITUENTS AS
POTENTIAL TREATMENTS FOR
METABOLIC SYNDROME**

A Thesis submitted by

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In loving memory of my late brother

Dr Rohit Bhandarkar

Abstract

Metabolic syndrome is the constellation of metabolic disorders such as central obesity, dyslipidaemia, insulin resistance, impaired glucose tolerance and hypertension. This combination increases the risk of development of cardiovascular disease, fatty liver and type 2 diabetes. The prevalence of metabolic syndrome is increasing worldwide.

Diet is important in the development of metabolic syndrome assisted by increased oxidative stress and inflammation. Plant-based diets provide potential therapeutic approaches to metabolic syndrome. Testing requires an appropriate animal model that mimics the human syndrome. In this project, I used a diet-induced obese rat model for examining the nutraceutical effects of some potential functional foods. To induce metabolic syndrome, young male Wistar rats were fed with a high-carbohydrate high-fat diet for 16 weeks while corn starch served as control diet. The high-carbohydrate high-fat diet induced an impaired glucose tolerance, insulin resistance, obesity, elevated blood pressure, dyslipidaemia, cardiovascular remodelling such as hypertrophy and fibrosis, increased cardiac stiffness, hepatic disorders such as inflammation and steatosis, along with elevated plasma markers of liver function.

Dietary interventions were given for the last 8 weeks only, as a reversal protocol. Interventions included green coffee extract (5%), decaffeinated green coffee extract (5%), chlorogenic acid (100 mg/kg/day), coffee pulp (5%), spent coffee (5%) and fish oils (3%). Green coffee with or without caffeine attenuated body weight and reduced cardiovascular disorders such blood pressure and cardiac stiffness, and improved heart and liver structure without improving glucose homeostasis or plasma lipid concentrations. Coffee pulp and spent coffee considered as waste products of coffee manufacturing industries attenuated cardiovascular remodelling and non-alcoholic fatty liver disease. Both waste products reduced body weight, improved glucose tolerance and decreased abdominal fat. Chlorogenic acid was present in all coffee products. Intervention with chlorogenic acid decreased body weight and visceral fat accumulation, improved heart and liver structure and function but did not improve glucose tolerance.

Prostate cancer patients treated with testosterone deprivation therapy, either through orchidectomy or leuprolide injection, show increased obesity. In high-

carbohydrate, high-fat fed rats, leuprolide treatment worsened metabolic syndrome symptoms and cardiovascular function, and orchidectomy produced greater responses. In H-fed leuprolide-treated rats, Omacor (a mixture of ethyl esters of EPA and DHA) decreased systolic blood pressure and left ventricular diastolic stiffness, reduced infiltration of inflammatory cells and collagen deposition in the heart, reduced lipid accumulation and inflammatory cell infiltration without improving liver damage. Thus, fish oils may provide an option to reduce metabolic syndrome while leuprolide treatment continues in patients with prostate cancer.

My studies show the promising potential of functional foods against life-style associated metabolic disorders. In particular, widely used beverage coffee showed relevant actions against most signs of metabolic syndrome. Further, waste products from coffee production are a potential source for new interventions in diet-induced cardiovascular and metabolic diseases.

Keywords

metabolic syndrome; obesity; cardiovascular remodelling; high-carbohydrate, high-fat diet; coffee; fish oil; coffee pulp; spent coffee; chlorogenic acid

Certification of Thesis

This Thesis is the work of **Nikhil Bhandarkar** except where otherwise acknowledged, with the majority of the authorship of the papers presented as a Thesis by Publication undertaken by the Student. The work is original and has not previously been submitted for any other award, except where acknowledged.

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Chapter 6. Attenuation of metabolic syndrome by EPA/DHA ethyl esters in testosterone deficient high-carbohydrate, high-fat diet-fed rats. Nikhil Bhandarkar, Senthil Arun Kumar, Jared Martin, Lindsay Brown, Sunil K Panchal

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Chapter 1. Introduction

Metabolic syndrome, cardiovascular disease and coffee as a functional food

Metabolic syndrome: prevalence and consequences

Obesity is characterised by the excessive accumulation of fat in adipose tissue [1]. The prevalence of obesity has escalated to epidemic levels globally. According to the World Health Organization (WHO), approximately 600 million people or 13% of the global population are categorised as obese and it is estimated that the number of newly diagnosed obese individuals has doubled since 1980 [2]. In the US, the prevalence of obesity was estimated at 35% among males and 40.4% among females in 2013-2014 [3]. Metabolic syndrome is the constellation of metabolic disorders such as dyslipidaemia, hypertension, visceral obesity, insulin resistance and impaired glucose tolerance [4]. Metabolic syndrome increases the risk of developing cardiovascular disease, type 2 diabetes and non-alcoholic fatty liver disease [5]. The association between obesity and metabolic syndrome was demonstrated in the National Health and Nutritional Examination Survey (NHANES) III study in which 5% of normal weight, 22% of overweight and 60% of obese subjects were classified as having metabolic syndrome in the United States [6].

Increasing poor nutrition, high-calorie diets, sedentary lifestyle and the comparative ageing of populations promotes an increase in the prevalence of metabolic syndrome [7]. Thus, targeting metabolism or energy homeostasis could be an approach to prevent or reverse metabolic syndrome and its consequences [8]. Due to the improvements in medical management, hypertriglyceridaemia and hypertension were reduced leading to reduction in metabolic syndrome from about 25.5% in 1999 to 22.9% in 2010 in the United States. However, abdominal circumference and hyperglycaemia steadily increased over the same period of time [9]. Cardiovascular diseases are one of the leading causes of mortality and morbidity worldwide [10, 11]. Patients with metabolic syndrome have higher risks to develop cardiovascular morbidity and mortality as well as type 2 diabetes. The prevalence of cardiovascular diseases was approximately three times higher in individuals with metabolic syndrome than individuals without it [12]. Despite some promising developments, metabolic syndrome remains one of the predominant causes of morbidity and mortality worldwide.

Chronic inflammation is closely linked to the development of obesity, insulin resistance, oxidative stress, cardiovascular disease and type 2 diabetes, characterised by abnormal secretion of cytokines and activation of the inflammatory signalling cascades [8]. Adipocytes modify metabolism by secreting non-esterified fatty acids, hormones such as leptin and adiponectin and pro-inflammatory cytokines [13, 14]. Excessive adipocyte accumulation triggers increased secretion of several pro-inflammatory adipokines such as interleukin (IL)-6, tumour necrosis factor (TNF), resistin, angiotensinogen, C-reactive protein (CRP), plasminogen activator inhibitor-1 (PAI-1) and fibrinogen causing prothrombotic alterations, increases in chronic low-grade inflammation and cardiovascular complications such as atherogenesis [15, 16]. These adipokines also regulate pathways that are associated with atherosclerosis, insulin resistance, hypertension and endothelial dysfunction [15, 16]. For instance, TNF is an adipokine that upregulates inflammatory variations in the vascular tissue, causing insulin resistance by inhibition of the insulin receptor signalling pathway and reduced the PPAR- γ expression level [15, 16].

Major components of metabolic syndrome

Metabolic syndrome is linked to hypertension and is responsible for myocardial dysfunctions, characterised by extended contraction and relaxation time, reduced rate of myocardial contraction and relaxation and reduced myocardial function [17, 18]. Several mechanisms have been proposed for cardiovascular abnormalities linked to metabolic syndromes, such as disrupted energy production because of reduced mitochondrial respiration and pyruvate dehydrogenase activity, oxidative stress, dyslipidaemia, mitochondrial leakage, reduced autophagy healing, endoplasmic reticulum stress and interrupted intracellular Ca^{2+} ion signalling [19, 20]. In a transgenic mouse model of lipotoxic diabetic cardiomyopathy, heart tissue showed elevated fatty acid intake, reduced diastolic sarcomere length, diminished myofilament Ca^{2+} response, higher β -MHC expression and extended diastole [21]. Several therapeutic options have been developed for the management of cardiovascular disease associated with metabolic syndrome, but they are not yet making a difference. This could be due to lack of clear understanding of metabolic syndrome associated pathogenesis in cardiovascular remodelling [22, 23].

Metabolic syndrome impairs cardiovascular function by the combination of atherogenic dyslipidaemia and hyperglycaemia due to the chronic pro-inflammatory conditions resulting in obesity, hypertension and prothrombotic state (Figure 1) [24, 25]. Subjects classified as having metabolic syndrome and diagnosed with at least 4 of 5 metabolic syndrome parameters have been shown to have 3-fold increased risk of cardiovascular diseases in a 5-year follow-up [26]. Atherogenic dyslipidaemia consists of elevations of lipoproteins such as apolipoprotein B (ApoB), elevated serum triglycerides, a high concentration of small LDL particles, and low concentration of high-density lipoproteins (HDL) [27]. The elevated levels of triglycerides and small LDL and low levels of HDL-cholesterol, collectively known as the lipid triad are the fundamental reasons for causing cardiovascular disorders such as angina pectoris, unstable angina, myocardial infarction or cardiac-related deaths [28].

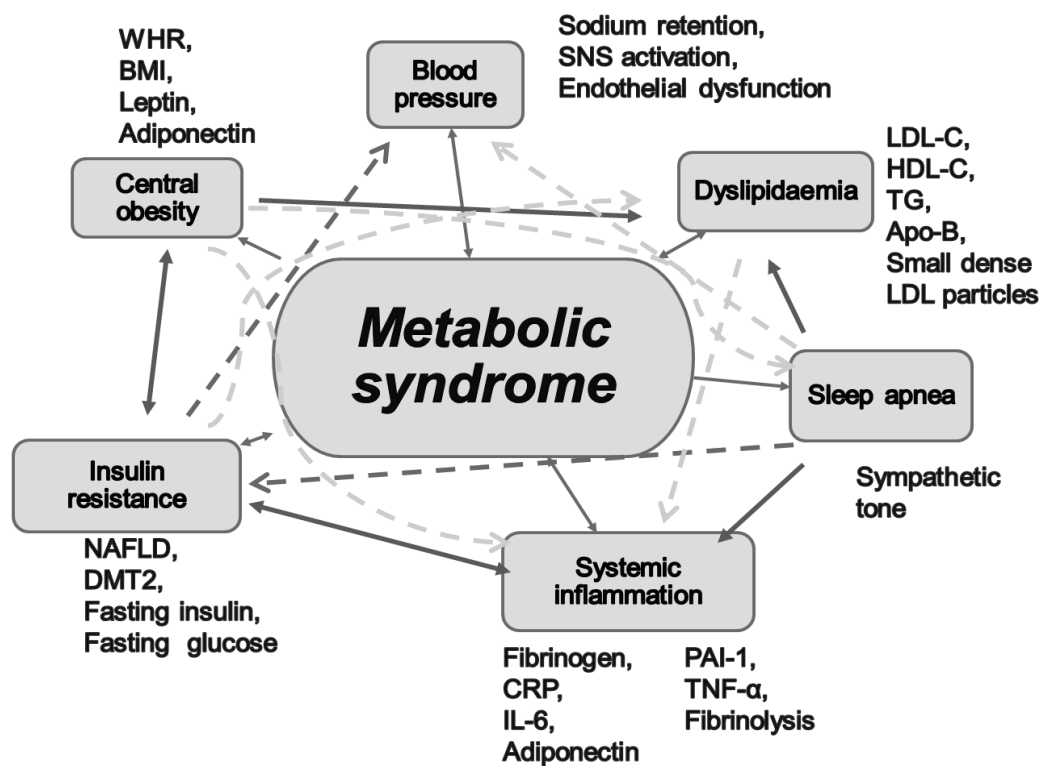


Figure 1. Components of metabolic syndrome [29]

Obesity

Obesity is responsible for triggering insulin resistance resulting in the pathogenesis of cardiovascular disease as insulin resistance promotes atherogenic dyslipidaemia [30, 31]. Insulin resistance is described as reduced sensitivity to circulating insulin. In diet-induced obesity, excess energy intake leads to an overload of adipocytes beyond their lipid storage capacity, causing excessive release of lipids into the bloodstream. High levels of circulating non-esterified fatty acids suppress glucose uptake by skeletal muscles [32]. Malfunction of fatty acid metabolism causes excess accumulation of lipids in myocytes, hepatocytes, and β -cells of pancreas, in turn, causing insulin resistance as well as impairment of β -cell function; these effects are noted as lipotoxicity [33, 34].

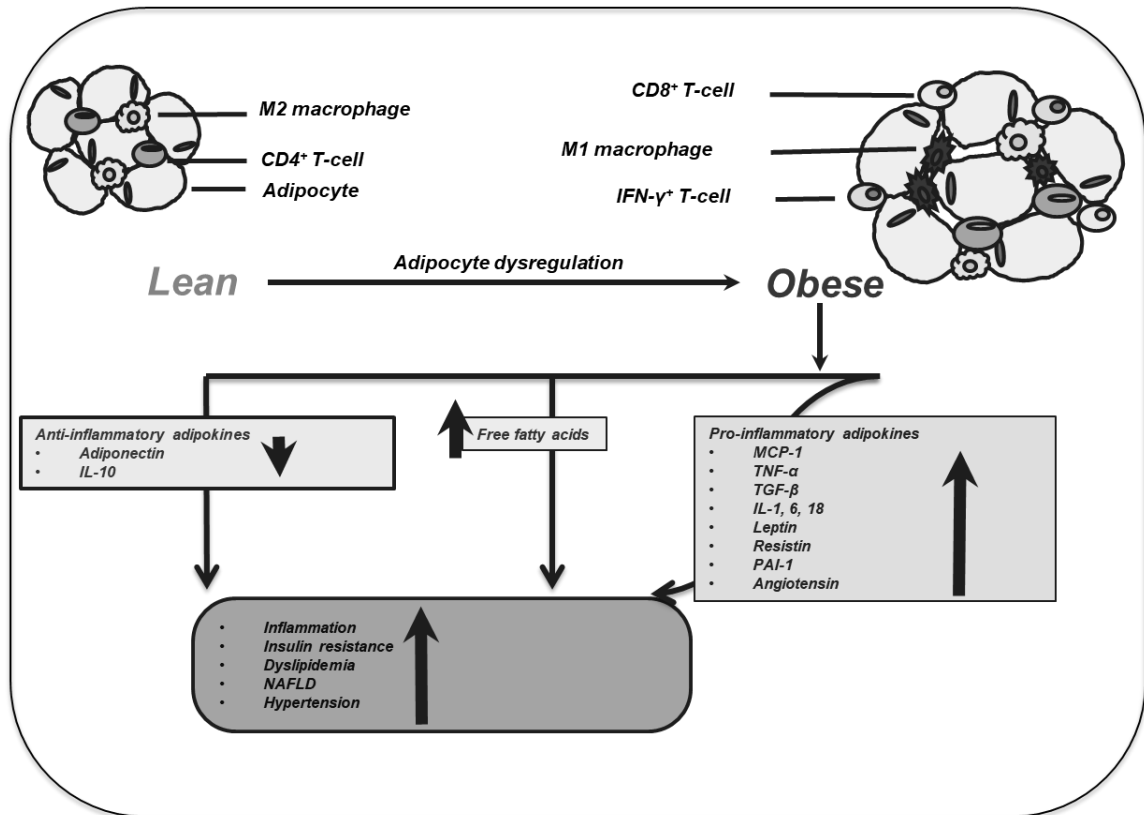


Figure 2. Chronic low-grade inflammation in obesity [35]

Lipotoxicity reduced insulin-mediated glucose metabolism in myocytes through inhibition of pyruvate dehydrogenase, consequently decreasing glucose oxidation and accumulation of glycolytic intermediates [36]. In such circumstances, pancreatic β -cells release more insulin to suppress the hyperglycaemia in the insulin-resistant state

essentially causing hyperinsulinemia. Sustained insulin resistance results in clinical symptoms of metabolic syndrome [37, 38] and may lead to cardiovascular events and type 2 diabetes [39]. Apart from glucose intolerance, dyslipidaemia and obesity play a crucial role in the excessive release of cytokines, leptin, TNF, IL-6 and decreased production of adiponectin (Figure 2) [40].

Insulin resistance

The insulin signalling pathway plays a fundamental role in glucose metabolism. Insulin binds to the receptor, a ligand-activated tyrosine kinase and insulin-associated tyrosine phosphorylation trigger activation of two corresponding pathways: the phosphoinositide 3-kinase (PI3K) pathway and the mitogen-activated protein (MAP) kinase pathway. In the insulin resistance state, the PI3K-Akt pathway is disturbed, while the MAP kinase pathway performs normally. Thus, insulin resistance creates disproportion between these two parallel pathways. Inhibition of the PI3K-Akt pathway causes a decrease in endothelial nitric oxide production, leading to endothelial dysfunction, and a decline in GLUT4 translocation, prominently resulting in reduced glucose uptake by myocytes [41]. Opposite to this, the MAP kinase pathway remains unaltered functionally. Consequently, there is a continuous release of endothelin-1 (ET-1), a stimulus of growth factors such as transforming growth factor- β (TGF- β), platelet-derived growth factor (PDGF), basic fibroblast growth factor (bFGF) and vascular cell adhesion molecules (VCAMs), which are crucial for development of an atherogenic plaque [42]. Therefore, insulin resistance induces vascular abnormalities, predisposing to atherosclerosis [43]. However, an insulin-resistant subject is not always categorised as clinically obese and such a subject may have an irregular distribution of accumulated adipose tissue, described as upper body fat [44]. Irrespective of the relationship between body fat and insulin resistance, insulin resistance has a strong correlation with progression in central obesity and thus, with metabolic syndrome [43].

Oxidative stress

Oxidative stress is the common factor that occurs during obesity, insulin resistance, impaired glucose tolerance, type 2 diabetes and cardiovascular disease [45].

Oxidative stress is a disturbed equilibrium between tissue oxidants (free radicals or reactive oxygen species (ROSs)) and antioxidants. Oxidative stress plays an important role in the development of obesity-associated comorbidities such as cardiovascular disease and diabetes [46]. In conditions of excessive energy intake, the abundance of substrates stimulates the activity of the citric acid cycle leading to the production of excessive mitochondrial NADH (mNADH) and ROS [47]. Free fatty acids and glucose are the primary energy sources in the skeletal muscle cells and adipocytes. Breakdown of either substrates mitochondrial acetyl-CoA and NADH [48]. When this acetyl-CoA enters citric acid cycle, it produces isocitrate which then combines with NAD^+ generating NADH with the help of catalytic activity of isocitrate dehydrogenase. When excessive NADH is generated and not eliminated by oxidative phosphorylation, it causes an increased mitochondrial proton gradient and the shift of a single charged electron to oxygen, thereby destabilising it and consequently creating free radical species such as superoxide anion (Figure 3) [49].

The overproduction of ROS is the initial and significant step that stimulates other pathways engaged in the pathogenesis of endothelial dysfunction during hyperglycaemia [50, 51]. Excessive ROS such as superoxide combines with nitric oxide, triggering formation of the strong oxidant peroxynitrite [52] that can alter DNA sequences [51] and the damaged DNA initiates the activation of the nuclear enzyme poly(ADP-ribose) polymerase [51] which initiates DNA repair, a process that requires NAD^+ as a substrate. The process of DNA repair may cause scarcity of intracellular NAD^+ . Restricted availability of NAD^+ slows the rate of glycolysis, electron transport and ATP generation, and promotes ADP-ribosylation of GAPDH [51], triggering acute endothelial dysfunction. Free fatty acids may function similarly [53] to increase oxidative stress and cause endothelial dysfunction. However, antioxidants can reverse consequences of oxidative stress in humans [54, 55].

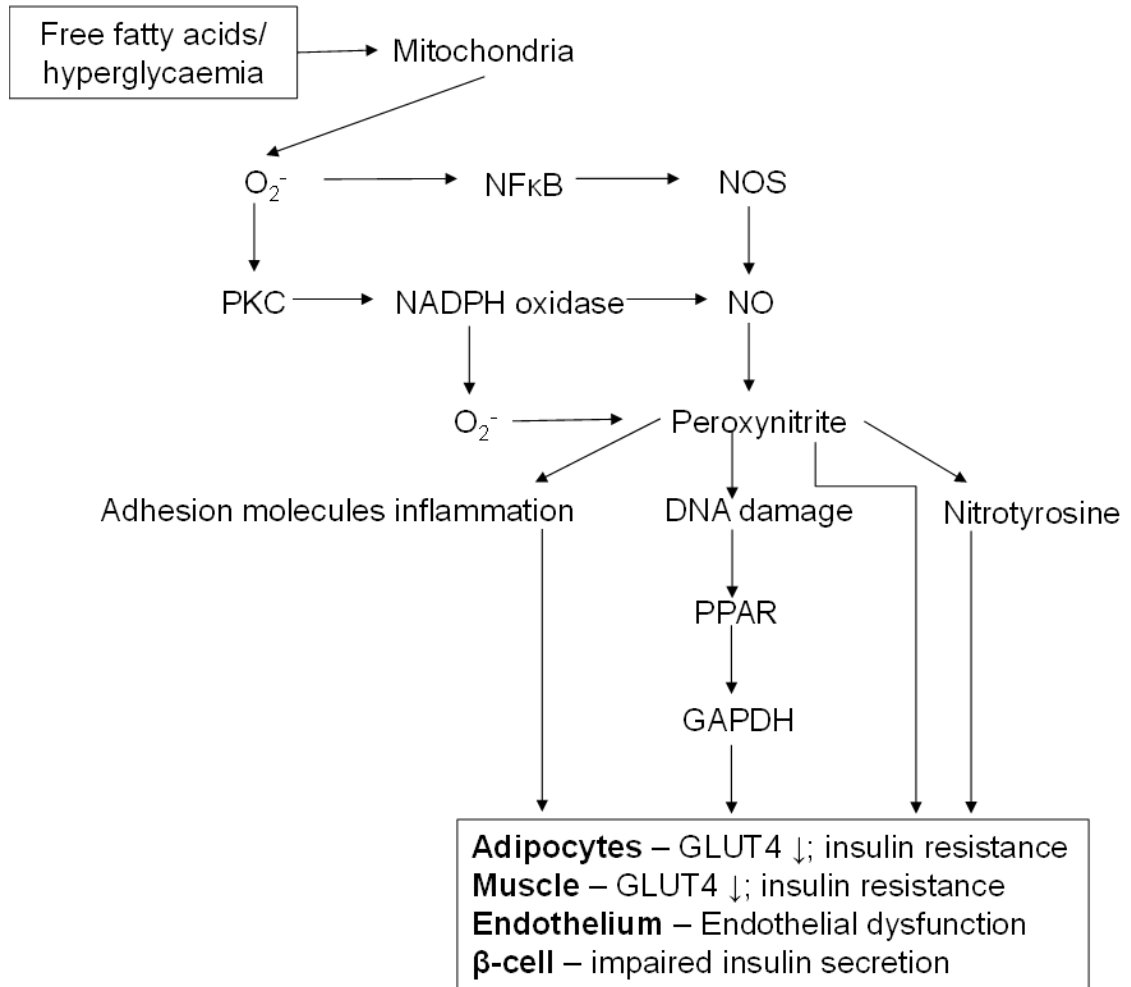


Figure 3. Association between oxidative stress and metabolic abnormalities [49].

Obesity and cardiovascular disease

Obesity has several severe impacts on cardiovascular structure and function and hence it is one of the prime reasons for cardiovascular disease-associated morbidity (Figure 4) [56]. Obesity escalates cardiac output, as well as total blood volume, thus increasing cardiac workload. In obesity, increased stroke volume causes an elevation in cardiac output associated with higher sympathetic activation which subsequently increases heart rate [57]. Obese subjects are more prone to develop hypertension than the lean persons as weight gain is associated with impaired control of arterial pressure [58]. Thus, obese and overweight subjects generally end up with left ventricular dilatation [56]. Regardless of age and sex, obesity increases the probability of left ventricular hypertrophy and other structural irregularities such as concentric remodelling and concentric left

ventricular hypertrophy [59]. Additionally, obesity causes left atrial enlargement because of elevated blood volume and eccentric left ventricular diastolic loading [60]. Such abnormalities contribute to impairments such as atrial fibrillation due to left atrial enlargement and heart failure [61]. Further, abnormal left ventricular structure increases the occurrence of frequent and complex ventricular arrhythmias [62].

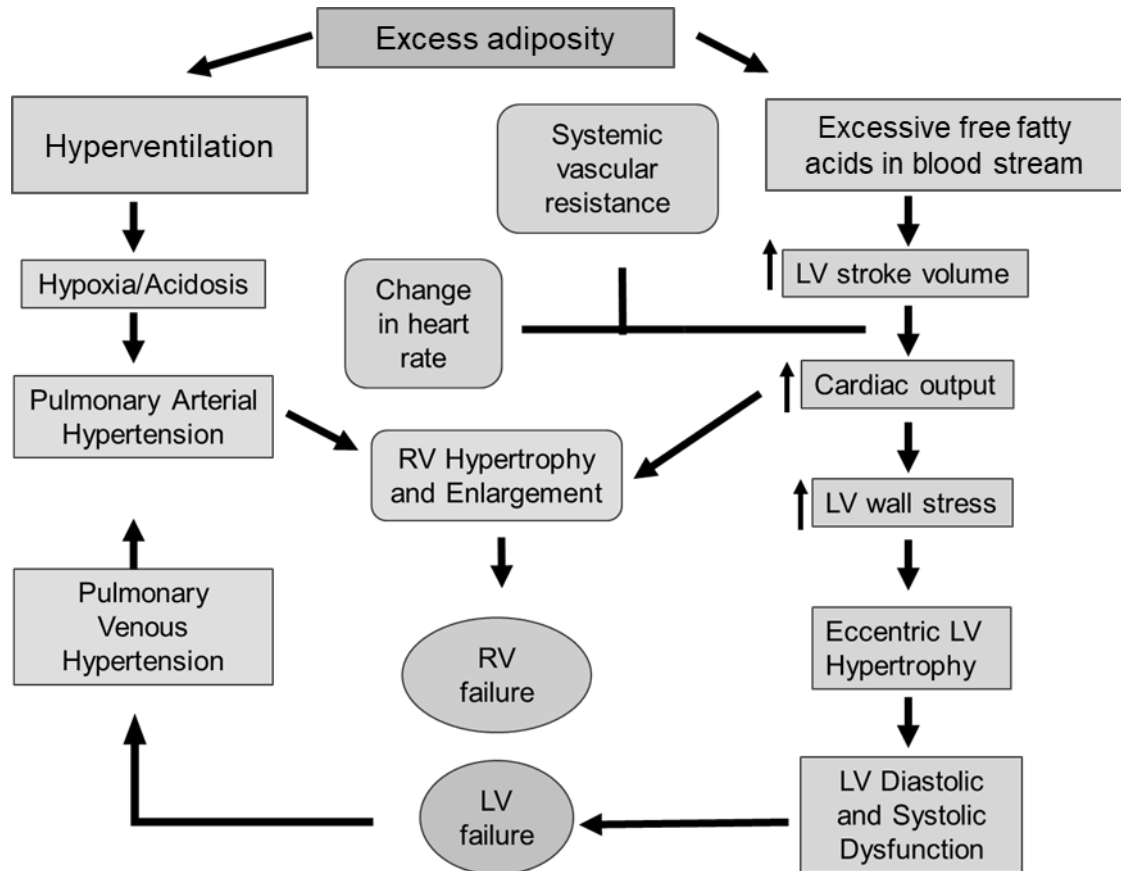


Figure 4. Association of obesity and cardiomyopathy. LV = left ventricular; RV = right ventricular [56].

Thus, obesity as the major component of metabolic syndrome affects the cardiovascular system and increases the prevalence of hypertension, dyslipidaemia and type 2 diabetes. Similarly, obesity plays a crucial role in atherosclerosis and cardiovascular disease [63].

Cardiovascular disease

High circulating concentrations of lipids and inflammatory cytokines lead to the atherosclerotic condition. In cardiovascular conditions such as unstable angina and

myocardial infraction, elevated concentrations of CRP and IL-6 have been associated with worsening of prognosis [64]. Moreover, other inflammatory markers such as IL-7, IL-8, soluble CD40 ligand and CRP-related protein pentraxin 3 increased during cardiovascular events [65-67]. Following are the cardiovascular outcomes that can develop as a result of uncontrolled metabolic abnormalities.

Hypertension

Hypertension is one of the predominant disease conditions associated with metabolic disorders such as obesity, glucose intolerance and dyslipidaemia [68]. Renin-angiotensin-aldosterone (RAAS) system-associated pathway AT₁R (Angiotensin II type 1 receptor) and mineralocorticoid receptor initiate NADPH oxidase pathway and promote the production of ROS. NADPH oxidase, as well as the mitochondrial generation of ROS, causes oxidative stress in the myocardium. This ROS also triggers insulin resistance, endothelial dysfunction and atherosclerosis [69]. The RAAS pathway is activated by hyperglycaemia and hyperinsulinaemia, leading to increased expression of angiotensinogen, angiotensin II (AT II) and the AT₁ receptors. Mutually, these play a crucial role in the progress of hypertension in a subject with insulin resistance [70]. There is also evidence that insulin resistance and hyperinsulinaemia lead to activation of the sympathetic nervous system, causing higher sodium reabsorption in the nephron, higher cardiac output and vasoconstriction of arteries initiating severe hypertension [71]. Moreover, it has been recently reported that adipose tissue also generates aldosterone in response to ATII [72]. Hence, the adipose tissue could be a location for renin-angiotensin-aldosterone system occurrence.

Chronic systemic inflammation or obesity has an important role in the development and progression of hypertension. A strong correlation exists between obesity and higher blood concentrations of inflammatory cytokines such as TNF- α , IL-1 β , IL-6 and CRP. Subsequently higher concentrations of inflammatory cytokines (TNF-alpha and IL-1 β) increase arterial stiffness, thus low-grade inflammation may play a role in the elevation of blood pressure [73, 74]. Increasing IL-6 concentrations in blood trigger elevation of soluble intercellular adhesion molecule-1 (sICAM-1) which is associated with elevated systolic and diastolic blood pressure, pulse pressure and mean arterial

pressure [75]. Plasma IL-6 was higher in women with elevated systolic and diastolic blood pressure, while in men, IL-6 was correlated with impaired glucose tolerance [74]. Several studies have shown that hypertension declined with the reduction in body weight regardless of the mechanisms involved [76]. Moreover, studies have reported the reduction in systolic blood pressure of 1-4 mmHg and diastolic blood pressure of 1-2 mmHg per kilogram of body weight lost in ~50% of the subjects tested [77].

Stroke

The published studies have shown that approximately 43% of people with metabolic syndrome are at the risk of stroke [78]. Obesity, insulin resistance and ROS are associated with increased inflammation that causes the development of the atherosclerotic cerebrovascular disease. Chronic low-grade inflammation present during atherogenesis of cerebral arteries is accelerated by several acute phase reactants such as CRP and fibrinogen. CRP is associated with an increasing risk of stroke events [79, 80]. Further, elevated CRP concentrations increased the incidence of systematic atrial fibrillation that escalates the risk of stroke [81, 82].

At cellular and molecular levels, insulin resistance has a role in the development of pathophysiology of vascular diseases such as stroke. Hyperglycaemia leads to increased oxidative stress by several mechanisms, for example, autoxidation of glucose produces ROS, a leading cause of oxidative stress [83]. In addition, advanced glycation end products, a heterogeneous group of compounds synthesised by non-enzymatic reactions between reducing sugars and free amino groups of proteins, are able to bind to cell surface receptors or cross-link with proteins, causing a modification of their structure and function [84]. Advanced glycation end products have the potential to induce insulin resistance and lead to increased oxidative stress and inflammation [85]. Furthermore, NADPH oxidase activation also initiates release of ROS [86]. NADPH oxidase catalyses the reduction of molecular oxygen causing superoxide radical formation [87]. In case of obesity, higher production of ROS occurs, especially in adipocytes, mediated by increased expression of NADPH oxidase. Further, treatment with NADPH oxidase inhibitors reduced ROS production in adipocytes, reduced the dysregulation of adipokines and improved metabolic syndrome parameters such as dyslipidaemia, obesity and insulin resistance

[88]. Nitric oxide is essential for prevention of cardiac ischaemia caused by oxidative stress and inhibits platelet aggregation thereby playing an important role in the response to cardiovascular disease [89]. Overproduction of nitric oxide can also cause adverse cardiac events such as atherosclerosis [90].

Hypertrophic cardiomyopathy

Obesity is strongly associated with heart failure and has been shown to cause deleterious effects on diastolic function [91, 92]. Cardiac hypertrophy is an adaptive response to physiological stress such as pressure or volume stress, mutations in myocyte proteins or metabolic disruptions. However, cardiac hypertrophy could be maladaptive and leads to cardiac dysfunction [93, 94]. Obesity could be a leading cause of severe hypertrophy by several mechanisms such as elevated sympathetic tone, elevated leptin concentrations, myocardial infiltration, insulin resistance and increased renin-angiotensin activity [95]. Obese individuals with hypertrophic cardiomyopathy had considerably higher left ventricular mass, left ventricular mass index and increased chances of progressing to severe cardiac events [96]. Moreover, the same study showed that obese subjects had a higher obstruction in the left ventricular outflow tract [96]. Prevalence of left ventricular obstruction was more than two-fold higher in obese individuals having hypertrophic cardiomyopathy compared to normal subjects and it was directly associated with elevated left ventricular mass [97, 98].

Coronary artery disease

Inflammation stimulates impaired arterial function both acutely as well as in chronic situation [99]. Chronic low-grade inflammation and oxidative stress contribute significantly to the development of atherosclerosis by contributing to the early stage of elevated endothelial permeability as well the creation of mature atherosclerotic plaque and plaque rupture [100]. Inflammatory biomarkers such as CRP have a crucial role in coronary artery disease risk [101]. Likewise, elevated concentrations of circulating soluble adhesion molecules such as sICAM-1, soluble VCAM-1 and soluble P-selectin contribute to coronary artery disease pathogenesis [102]. However, different inflammatory biomarkers cause an increased risk of coronary artery disease through various

mechanisms. Elevated CRP and IL-6 have been found in subjects with stable angina and myocardial infarction with worsening prognosis [67]. Moreover, other inflammatory biomarkers such as fibrinogen, IL-7, IL-8 and soluble CD40 ligand are also increased in cardiovascular disease patients [103].

Regardless of gender, weight-stable obese individuals have a higher risk of arrhythmias and sudden cardiac death [104]. In both genders, the rate of sudden cardiac arrest was approximately 40 times higher in obese individuals than lean individuals [95]. Further, for subjects aged between 35-44 years, mortality due to cardiac arrest is 6 times higher in men with severe obesity than in women of a similar age group. Hyperglycaemia associated with metabolic syndrome decreases nitric oxide availability resulting in elevated vasomotor tone and ventricular irregularity [105, 106]. Likewise, increased free fatty acids cause increased plasma catecholamine resulting in altered cardiac repolarisation [107]. Specifically, long-chain saturated fatty acids have an important role in the occurrence of ventricular arrhythmias leading to myocardial infarction [108]. The saturated fatty acid, palmitate, caused a reduction in mitochondrial respiration in cardiomyocytes and promoted production of excessive total cellular ROS as well as mitochondrial ROS [109]. Excessive ROS production caused partial depolarisation of the mitochondrial inner membrane and subsequently triggered mitochondrial calcium overload by raising sarcoplasmic reticulum calcium leakage, that leads to arrhythmias and heart failure [110].

Dietary interventions against metabolic syndrome

Previous studies have shown that moderately modified lifestyle and dietary interventions can reduce the risk for metabolic diseases such as diabetes [111-113]. Diet and physical activity promote weight loss, favourably modulating parameters of metabolic syndrome. Epidemiological studies have reported that polyphenol intake has a positive impact on cardiovascular and hepatic disorders [114-116]. In addition, human studies with polyphenol-rich diets have concluded a favorable effect of polyphenols on risk factors of cardiovascular health such as blood pressure [117], serum lipid profile [118] and insulin sensitivity [119]. Studies using animal models have demonstrated that polyphenols improve insulin sensitivity [120], have anti-obesity effects by reducing body fat [121] and

are suggested to reduce plasma concentrations of triglycerides [122, 123] and cholesterol [124].

Traditionally, coffee is recommended for health management as it is rich in bioactive substances with a wide range of physiological effects [125-127]. Key bioactive substances include caffeine, chlorogenic acid and the diterpenes, cafestol and kahweol. The chemistry and biochemistry of coffee have been extensively described [128]. Coffee undergoes a chemical transformation during the roasting process of green bean, and the type of bean (arabica vs robusta), the degree of roasting, and brewing techniques including coffee grind setting and brew type will all have an impact on the biochemical configuration of the final beverage [129, 130].

Coffee as a functional food for metabolic syndrome

Coffee is one of the most commonly consumed beverages in the world. It is a healthier alternative to alcoholic beverages such as beer and spirits [131]. Coffee beans of two main species, *Coffea canephora* (also known as robusta) and *C. Arabica*, are commercially available for the preparation of coffee. Both beans are well-known for their pleasant aroma and enjoyable taste [132]. The range of compounds in coffee may provide beneficial effects against metabolic syndrome [133]. Coffee is the second-most traded commodity in the world after oil [134]. The consumption of coffee has doubled in the last four decades [135]. Coffee consumption was 6.947 million tonnes in 2010 [136] and it is estimated that consumption will increase to 9 million tonnes in 2019 [135]. In 2010, Brazil was the major producer of coffee contributing about 19% of the world's coffee production. The United States was the major coffee consumer with about 16% of world consumption [136].

The chemical composition of coffee is affected by factors including harvesting and post-harvest techniques, climate, physical conditions of soil and tree and genetic aspects of coffee seeds [137]. These components also vary between *C. canephora* and *C. arabica*. Coffee is a complex mixture of caffeine (0.5-2.8%), chlorogenic acid (4.80-6.41%), polyphenols, caffeic acid, lactones and diterpenoid alcohols including cafestol and kahweol, as well as niacin and the vitamin B₃ precursor, trigonelline. It also has ample

amounts of vitamin B₃, magnesium and potassium [137, 138]. During the roasting process of coffee beans, chlorogenic acid undergoes chemical transformations such as isomerisation, hydrolysis and degradation, causing reduction of chlorogenic acid concentrations and conversion into quinolactones and melanoidins [128, 139]. The roasting process initiates the browning reaction and gives various different compounds of varying molecular weights such as aldehydes, ketones, dicarbonyls, acrylamides, heterocyclic amines and melanoidins that impact flavour [140]. Melanoidins account for 25-29% (w/w of dry matter) of roasted coffee beans [141]. These compounds are high molecular weight, water-soluble, brown-colored nitrogenous compounds with antioxidant, antimicrobial and anti-hypertensive activities [142, 143]. Melanoidins link or bind with chlorogenic acid non-covalently and regulate gut microbiota growth to produce anti-inflammatory effects [139]. Although coffee is rich in caffeine, chlorogenic acid, trigonelline, diterpenes, and melanoidins, which collectively or individually can positively influence health, the availability of these compounds depends on how the coffee beans are processed and consumed.

There are several types of coffee beans generated depending upon processing of green coffee beans. A green coffee bean does not have the flavour qualities of a roasted bean and it's green in color with a grassy smell. Roasting process induced the aroma and flavour in beans depending on temperature and time taken for processing. It creates fresh roast flavour with different chemical components in roasted beans. The coffee extraction and brewing process only extracts part of the coffee components, so that the resulting spent coffee ground may be a source of useful products as healthy bioactive compounds, including trigonelline and melanoidins [139, 144].

Green coffee

Green coffee beverages accelerate glucose metabolism and lead to reduced body weight and prevent obesity [145]. Green coffee extract has the potential to inhibit fat absorption and control lipid metabolism in livers of mice [146]. Likewise, green coffee has anti-hypertensive properties against high blood pressure in rats [147].

Green coffee is the mixture of complex chemicals that impart flavour, aroma and texture to coffee beverages after roasting. Along with carbohydrates, proteins, lipids and minerals, it is also a rich source of phenolic acids such as chlorogenic acid and alkaloids such as caffeine [148]. Health beneficial effects from green coffee are mainly associated with the presence of phenolic compounds, especially chlorogenic acid, that have been recognised for their antioxidant properties [149] associated with its ability to scavenge ROS. In addition, chlorogenic acid and caffeine inhibited tumour promotion and inflammation cascade by restricting the pro-oxidant enzyme lipoxygenase for metabolising arachidonic acid [150]. Moreover, green coffee phytochemicals have demonstrated potential to decrease visceral adipose tissue accumulation and body weight gain with several studies supporting green coffee as a weight loss supplement in animal models [151, 152].

Chronic intake of coffee has been related inversely to body weight gain in humans [153]. Regular coffee intake has also been associated with modulation of glycaemic markers [154]. Likewise, green coffee intake reduced weight gain, where caffeine and other bioactive compounds present in coffee play a crucial role as thermogenic agents [155]. In human study, green coffee has shown potential for modulating hormone release and glucose tolerance [156]. The possible mechanism could be the delay in glucose absorption where green coffee causes the absorption of glucose in distal colon instead of the proximal colon. This results in the reduction in postprandial plasma concentrations of glucose and blood lipids [157]. Additionally, this modulation increased the diversion of the Na⁺ electrochemical gradient, and consequently reduction in intestinal brush border membrane glucose uptake [156]. Green coffee was also shown to regulate glucose homeostasis by inhibiting hepatic glucose-6-phosphatase [158]. Through suppression of hepatic triglyceride accumulation, green coffee could also decrease obesity [151]. Green coffee also altered adipokines secretion, prevented fat accumulation via downregulation of lipogenesis while upregulating lipid oxidation and expression of PPAR- α in the liver [159].

Roasted coffee

While green coffee beverages have a mild, green and bean-like taste and aroma, the desirable properties of coffee are acquired during the roasting process. During the roasting process, the coffee beans are heated from 200-250°C and process duration can last between 0.75 to 25 minutes, the optimal period being 1.5 - 6 minutes depending upon the specifications of the end product i.e., light, medium or dark roasted beans. Roasting process causes several physical and chemical modifications leading to a shift in colour from green to brown. The chemical components such as protein, amino acids, polysaccharides, trigonelline, chlorogenic acid and water undergo complex processes to form melanoidins. Melanoidins are a product of Maillard reaction through caramelisation of polysaccharides and amino acids with other coffee components during heating of coffee beans [160]. Roasted coffee is a rich source of bioactive compounds especially methylxanthines and polyphenols. The roasted coffee extract has demonstrated anti-inflammatory and anti-oxidant properties with great potential for reducing ROS and inhibitory action against linoleic acid peroxidation and lipoxygenase [161]. In a human study, chlorogenic acid from roasted coffee caused a reduction in oxidative stress-induced DNA and macromolecular damage and increased antioxidant defense capacity [162]. In addition, the efficiency of antioxidant properties of roasted coffee is shown as the increases in glutathione S-transferase activity *in vitro* as well as in rats [163]. Further, roasted coffee components increased transcription and nuclear translocation of Nrf2 in HT-29 cells [164]; this modification induced by coffee components prevented colon cell damage from oxidative stress caused by ROS. Therefore, roasted coffee components reduced oxidative stress and restrict the progress of metabolic syndrome associated complications such as cardiovascular disease.

Decaffeinated coffee

Coffee beverage is generally popular for caffeine intake, and caffeine's stimulatory effect makes it the most recognised and most studied component in coffee. Studies have shown that caffeine has favorable effects on cardiovascular function [165, 166]. However, caffeine intake causes nervousness, irritability, headache and troubled sleeping patterns in many individuals [167]. Further, coffee consists of hundreds of other bioactive

compounds such as caffeic acid, chlorogenic acid and trigonelline, making it difficult to isolate the health effects of caffeine from those of the other ingredients.

Decaffeinated coffee is consumed by individuals who have problems in tolerating caffeine. According to the United States Department of Agriculture, decaffeinated coffee is not necessarily absolutely caffeine-free. In fact, coffee only requires being ~97% caffeine-free in order to be classified as decaffeinated coffee, although Brazilian regulations only permit a maximum of 0.1% of the residual caffeine in decaffeinated coffee [168]. The process of decaffeination causes an alteration in decaffeinated coffee bean and depending upon coffee variety, coffee components such as chlorogenic acid and other components increase or decrease [149]. Several studies have reported health benefits from the decaffeinated coffee intake. In a clinical trial, decaffeinated green coffee reduced weight in overweight volunteers treated with 400 mg/day for 60 days [169]. Another study stated that drinking 0.3% decaffeinated green coffee extract in water for 20 weeks reduced fat accumulation and insulin resistance in mice fed on a high-fat diet [170]. Decaffeinated coffee extract downregulates TLR4-mediated pro-inflammatory pathways and stimulates GLUT4 translocation to the plasma membrane in adipocytes [170]. Similarly, ingestion of decaffeinated coffee for 20 weeks after inducing obesity using a high-fat diet over a 10 week period resulted in a decrease in blood glucose concentrations and improvement in insulin tolerance. The coffee extract also promoted recovery from liver remodelling, showing a decrease in steatosis, and inhibited iNOS expression and restored the insulin-inducing Akt phosphorylation [171].

Coffee wastes

Almost 50% of the coffee bean is discarded as waste during the process of commercial production up to and including the consumption by the end-user. The waste is generated from production (e.g. coffee pulp, cherry husks and defective beans), from roasting industries (e.g. coffee silver skin), from soluble coffee industry (industrial spent coffee) and also directly by daily coffee consumers after coffee brewing (spent coffee) [172]. These coffee wastes have high concentrations of bioactive compounds including caffeine and polyphenols [173].

Coffee wastes have been used mainly for commercial processes such as the production of ethanol, butanol, hydrogen, organic acids, glycerol and hydroxymethylfurfural. Coffee wastes are also used for composting in several agro-industrial sectors [134]. Despite the high bioactive content of coffee wastes, only partial progress has been accomplished in its application as a functional food [174]. Coffee waste such as coffee husks, skin and pulp contain four major classes of polyphenols, flavan-3-ols, hydroxycinnamic acids, flavonols and anthocyanidins which include chlorogenic acid, epicatechin, catechin, rutin and ferulic acid. Coffee husks, skin, and pulp are potential sources of 5-feruloyl-quinic acid and anthocyanins such as cyanidin-3-rutinoside and cyanidin-3-glucoside [172].

Coffee mucilage and parchment include several bioactive components such as amino acids, polysaccharides, pectic constituents and a trace amount of ash in addition to α -cellulose, hemicellulose and lignin [134]. However, research on the functional characteristics of this coffee portion has not been performed yet. Roasted coffee silverskin, a by-product of roasted coffee beans, is a rich source of dietary fibre, accounting for up to 60% of the by-product, 14% of which are soluble dietary fibre [175]. The roasted coffee silverskin also contained a minor amount of phenolic compounds with demonstrated antioxidant properties. The antioxidant properties may be attributable to melanoidins. Additionally, roasted coffee silverskin stimulated the preferential growth of *Bifidobacterium* rather than *Clostridia* and *Bactericides* spp. Hence, the roasted coffee silverskin could be used as a functional ingredient due to its high content of soluble fibre, its potential antioxidant activity and the prospective prebiotic properties [175].

Coffee components

Caffeine

Caffeine is a well-known stimulant of the central nervous system, acting as an antagonist to adenosine receptors. Adenosine receptors are present and expressed in the central nervous system, the vascular endothelium, heart, liver, adipose tissues and muscles. Thus, caffeine triggers a broad range of responses in the body [138]. It produces many biological functions including an acute elevation of heart rate, stimulation of the

central nervous system, increased carbohydrate metabolism and decreased inflammation, as well as increasing metabolic rate and causing diuresis [176-178]. Caffeine (~30 mg/kg/day) in diet-induced obese rats reduced abdominal fat, total body fat, systolic blood pressure and improved glucose tolerance, cardiovascular and hepatic structure and function [179]. Caffeine promoted the gene expression of carnitine acyltransferase in the liver of obese mice [180]. This increase in the expression of carnitine acyltransferase may be responsible for the reduction of adipose tissue weight through increased fat oxidation and hence suppression of body weight gain. Caffeine also increased energy expenditure [181]. Regular intake of caffeine decreased the risk of insulin resistance by decreasing the sympathetic nervous system response [182].

Chlorogenic acid

Chlorogenic acids are a group of compounds that are esters of caffeic acid and quinic acid [183] and may have health benefits due to their antioxidant and anti-inflammatory activities. Only one-third of ingested chlorogenic acid is absorbed in the small intestine and some amount that escapes metabolism enters the colon [184]. In the colon, the microbiota hydrolyses chlorogenic acid into caffeic acid and quinic acid [184]. These microbial metabolites undergo absorption or are changed by human enzymes and finally get excreted in urine [183]. Negligible amounts (0.8%) of intact chlorogenic acid are found in human urine. Chlorogenic acid has bacteriostatic properties towards a wide range of Gram-positive and Gram-negative bacteria, such as *E. coli* and *Salmonella* spp. [185] while specifically promoting the growth of *Bifidobacterium* spp., *Lactobacillus/Enterococcus* spp., *Bactericides* spp. and *C. coccoides–E.rectale* [184]. As almost two-thirds of the dietary chlorogenic acid reaches the colon and undergoes metabolism, it may have some effects in the colon, especially on the gut microflora, which may help in attenuating obesity [186].

Chlorogenic acid inhibited liver glucose-6-phosphatase activity, suggesting that it can reduce liver glucose output [158]. Chlorogenic acid from decaffeinated coffee downgraded the expression of sodium-dependent glucose transporters in brush border membranes of the small intestine thus suggesting its potential to reduce glucose absorption [187]. Similarly, chlorogenic acid from green coffee extract (0.25 - 1% diet for 6 weeks)

lowered blood pressure in Spontaneously Hypertensive Rats [147]. Chlorogenic acid increased mRNA expression of PPAR- α and liver X receptor- α , the key mediators of lipid metabolism [188]. Chlorogenic acid (100 mg/kg/day in C57BL/6 mice for 15 weeks) prevented diet-induced obesity and metabolic syndrome by inhibiting expression of PPAR- α and by reducing inflammation [189]. Chlorogenic acid helps to delay glucose absorption from the gut into the bloodstream. Hence, chlorogenic acid could be a potential dietary candidate for the host to manage metabolic disorders such as obesity, increased blood glucose concentrations and cardiac complications.

Melanoidins

During the roasting of coffee beans, melanoidins are formed by the non-enzymatic browning reaction as a chemical reaction between reducing polysaccharides, amino acids and phenolic compounds [190]. Moreover, melanoidins are formed during the heating process of a broad range of foodstuffs including bread, malt, meat and tomato sauce [191]. The formation of melanoidins leads to increased antioxidant capacity for coffee beans, but this occurs only at a particular phase of coffee roasting [192]. Studies of coffee revealed that some positive effects on pathophysiology which are not due to caffeine can be credited to polyphenols and melanoidins [193]. The molecular structure of melanoidins are yet to be defined and thus, they are generally defined as macromolecular nitrogenous browning compounds [194]. However, some studies have shown that roasting process causes covalently linkage between arabinogalactans and proteins to produce arabinogalactan proteins [129] that have hydrophobic characteristics and ion-chelating abilities [195]. Roasting process leads to degradation of the proteins present in green coffee beans, including arginine, cysteine, lysine, and serine, and the degradation products contribute to the formation of complex melanoidins [196]. Moreover, chlorogenic acid and their derivatives are fused in coffee melanoidins [197]. The intact chlorogenic acids are incorporated into the melanoidin structures, combined through the caffeic acid moiety by mainly non-ester linkages [130]. In summary, polysaccharides, proteins and phenolic compounds such as chlorogenic acids are involved in the formation of coffee melanoidins.

An *in vitro* study by stimulating gastrointestinal enzymatic digestion showed that the coffee melanoidins are largely resistant to digestion in the human gut [198] suggesting that melanoidins could have prebiotic activity. Melanoidins have the potential to alter human gut microflora as fermentation of melanoidins supports the growth of residential bacteria [199]. The gut bacteria may also partly metabolise the melanoidins in the large intestine, hence, this might play a crucial role in the chelating and/or release of other, potentially harmful dietary components. For instance, a study of coffee melanoidins with human subjects has reported for growth in the number of *Bacteroides* and *Prevotella* bacteria identified in faecal samples [200]. In *in vivo* studies, melanoidins have increased the quantity of *Bifidobacterium spp.*, known for their probiotic activity [201, 202]. *In vitro* fermentation of melanoidins for 24 hours with faecal bacteria from human subjects showed its anti-oxidant properties [203] indicating a potential role for coffee melanoidins in the protection against oxidative stress caused by free radical species in the colon, resulting in the reduction of gut dysbiosis.

The anti-oxidant properties of melanoidins have been implicated in their protective effects on human hepatoma HepG2 cells against oxidative stress where melanoidins reduced glutathione peroxidase and glutathione reductase [204], indicators of attenuation in oxidative stress, as melanoidins contain hydroxyl groups, which serves as a hydrogen donor, and are able to scavenge free radical species [205]. Melanoidins inhibits secondary lipid oxidation products that contribute to health benefits, products from lipid oxidation are involved in the development of cardiovascular diseases [206]. Thus, in addition to the antioxidant property of melanoidins, the available studies suggested that ingested melanoidins from regular coffee intake could possibly provided the protection against inflammation and promote the growth of selective gut microbiota [191].

Other coffee components

Along with chlorogenic acid, caffeine and melanoidins, coffee is also a source of other bioactive components such as trigonelline and diterpenes (cafestol and kahweol). These compounds from coffee have the ability to protect cells from ROS-induced oxidative stress by inducing the transcription of genes in the Nrf2 antioxidant response element signalling pathway [164].

Trigonelline is an alkaloid and derivative of nicotinic acid. After demethylation, it is converted to nicotinic acid which is present at approximately 1–3 mg/240 ml brewed coffee [187]. During roasting of coffee beans, trigonelline is also transformed into its intermediate, N-methylpyridinium, which improves the reducing and total antioxidant capacity of trigonelline [207]. N-methylpyridinium is converted to nicotinic acid which has the potential for anti-inflammatory properties through down-regulation of MCP-1 and has increased adiponectin in adipocytes treated with TNF [208]. Decomposition of trigonelline was greater at higher roasting temperature generating higher amounts of nicotinic acid [209].

Diterpens such as cafestol and kahweol have a negative impact on cholesterol concentrations. Diterpenes could be a possible reason for increased plasma lipid concentrations in coffee drinkers as a high concentration of diterpenes have been reported to induce increased total cholesterol and LDL-cholesterol [187]. Coffee also increased serum LDL through increased expression of cholesterol esterase transfer protein assisted by cafestol [210]. Diterpenes have shown antioxidant properties and cells treated with diterpenes showed decreased DNA damage because of the ROS scavenging activity of cafestol and kahweol [211]. Also, kahweol and cafestol treatment protected NIH3T cells challenged with hydrogen peroxide [211]. Kahweol conferred an anti-inflammatory effect by down-reregulating the expression of cyclooxygenase-2 and MCP-1 in HUVECS and decreased the inducible nitric oxide synthase in rats treated with carrageenan [212]. A recent study has also shown beneficial effects of cafestol against diabetes in KKAY mice [213].

Thus, metabolic syndrome is strongly associated with the increasing prevalence of cardiovascular diseases and associated comorbidity. Cardiovascular diseases are reliant on reversible metabolic abnormalities such as oxidative stress, dyslipidaemia and impaired glucose tolerance, which are associated with metabolic syndrome. Coffee and its phytochemical components exhibit the potential against oxidative stress, inflammation and insulin resistance and thus, coffee beverages could be a potential dietary intervention for cardiovascular disease. It is difficult to determine what component of coffee could be responsible for beneficial health effects because of the complex chemical composition of

coffee beans as well as coffee waste. By investigating responses to coffee products and coffee components, beneficial effects of coffee waste and coffee components can be explored. Clearly, all investigations into the health benefits of coffee intake must consider several parameters such as coffee variety, coffee beans, roasting method, sample size, brewing course, and a dose of individual coffee components administered and the duration of treatment. In order to understand the specific roles of individual components of coffee in cardiovascular health, properly controlled experiments have been carried out to determine preclinical outcomes.

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Chapter 2.

Green coffee improves diet-induced metabolic syndrome in rats

Green coffee improves diet-induced metabolic syndrome in rats

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Abstract

Abdominal obesity is associated with the development of metabolic disorders, type 2 diabetes and cardiovascular diseases. This study assessed the effects of green coffee on metabolic variables in obesity induced by high-carbohydrate, high-fat diet in rats. Male Wistar rats (8-9 weeks old, 340 ± 5 g, $n = 72$) were divided into 6 groups and fed with either corn starch diet (16 weeks) (C), corn starch diet with 5% green coffee (CGC) or decaffeinated green coffee extract (CDC) in food for the last 8 weeks, high-carbohydrate high-fat diet (16 weeks) (H), or high-carbohydrate high-fat diet with 5% green coffee (HGC) or decaffeinated green coffee extract (HDC) in food for the last 8 weeks. The high-carbohydrate high-fat diet-fed rats showed signs of metabolic syndrome as changes in cardiovascular and liver structure and function. Green coffee attenuated body weight (C 385 ± 5 g, H 563 ± 14 g, HGC 515 ± 16 g, HDC 522 ± 15 g), feed efficiency (C 0.11 ± 0.01 kJ/g, H 0.37 ± 0.02 kJ/g, HGC 0.31 ± 0.02 kJ/g, HDC 0.32 ± 0.02 kJ/g), systolic blood pressure (C 128 ± 1 mmHg, H 147 ± 1 mmHg, HGC 130 ± 2 mmHg, HDC 128 ± 2 mmHg) and improved structure of heart and liver without improving obesity, glucose sensitivity or dyslipidaemia. The chlorogenic acid in green coffee could be the major bioactive compound to attenuate diet-induced abnormalities in heart and liver structure and function.

Keywords: Green coffee, Obesity, Metabolic syndrome, High-carbohydrate, high-fat diet

Introduction

Functional foods have been proposed as interventions to prevent or reverse metabolic syndrome [1], a constellation of metabolic disorders including obesity, hypertension, impaired glucose tolerance, insulin resistance, dyslipidaemia and fatty liver as a major risk factor for cardiovascular disease and type 2 diabetes [1]. Obesity and insulin resistance are strongly linked with metabolic syndrome in the general population [2, 3]. As part of a healthy diet, green coffee could be a potential functional food for metabolic syndrome [4]. Green coffee extract decreased obesity and blood lipid changes by mechanisms including inhibition of adipogenesis, scavenging of reactive oxygen species, and reduced triglycerides and glucose concentrations in rats and adipocytes in culture [5]. Coffee components such as chlorogenic acids, caffeine, trigonelline, and diterpenes reach active concentrations in the human body [6]. A cross-sectional study in humans concluded that green coffee intake was inversely associated with some parameters of metabolic syndrome and cardiovascular mortality [7]. Likewise, green coffee improved weight loss [8] and protected against the development of non-alcoholic fatty liver disease (NAFLD) [9]. A meta-analysis of studies on weight loss with green coffee showed an average loss of 2.47 kg, but all studies had a high risk of bias and were of poor methodological quality [10]. However, the extent of the responses on the range of disorders in metabolic syndrome and the compounds responsible for those effects have not been determined.

The processing of coffee berries into coffee involves six main steps of pulping, fermentation, drying, hulling, roasting then grinding [11]. Green coffee products, usually sold as capsules, come from coffee beans that have not been roasted, so these products

contain different proportions of components compared to roasted or black coffee [12]. The moderate intake of green coffee, 4-6 cups/day, delivers bioactivity of some coffee components such as caffeine [13], chlorogenic acids [13], trigonelline [14], and diterpenes [15] in human body [16,6]. Caffeine intake has been demonstrated to decrease metabolic syndrome [17], however, caffeine consumption is also reported for adverse outcomes consistent with the physiologic effects among adults and young adults [18,19]. Esters of caffeic acid and chlorogenic acid (caffeoylquinic acids) are well demonstrated for health benefits such as improvement in cardiovascular disease and type 2 diabetes [20]. Chlorogenic acids show potential anti-oxidant and anti-inflammatory effects through reducing liver mRNA expression of TNF-alpha, IL-6 and IL-1beta that provide protection against lipid accumulation of Sprague-Dawley rats fed a high-fat diet (HFD) [21]. Chlorogenic acids (50 mg/kg for 42 days) increased the plasma lipid metabolism in rats by reducing the concentrations of free fatty acids and triglycerides and regulating the multiple components in hepatocytes through AMPK pathway [22], which indicates that chlorogenic acids could be a promising ingredient in diet for obesity management. Along with chlorogenic acids, caffeic acids, trigonelline and cafestol provided the synergistic effects and improved the insulin sensitivity in the treated rats, and increased plasma concentrations of adiponectin [23]. Further, cafestol (a diterpene present in coffee brew) has insulinotropic effects on pancreatic β -cells, induces glucose uptake in human myoblasts [24], and increases glycaemic regulation *in vivo* [25].

This study has determined whether green coffee and its decaffeinated product can improve signs of metabolic syndrome when given as 8-week dietary interventions in rats fed a high-carbohydrate, high-fat diet for 16 weeks to mimic human metabolic syndrome

[13]. Cardiovascular function was studied *in vivo* by measurement of systolic blood pressure and *ex vivo* in isolated Langendorff heart preparations, isolated thoracic aortic rings together with histology. Liver structure and function were measured by histology and plasma biochemistry. Metabolic parameters related to obesity and glucose tolerance were also evaluated.

Methods

Analysis of green coffee

Extractions were prepared in 3:2 ethanol:water mixture. Briefly, 1 g of coffee (*Arabica spp.*) was dissolved in 50 mL of ethanol:water mixture, sonicated for 15 minutes and an aliquot of the supernatant was taken for analysis by the HPLC using an Agilent 1100 series system coupled with a mass spectrometer for further peak confirmation or identification as required. The HPLC system consisted of a diode array detector (G4212B), binary pump (G4220A), an autosampler (G4226A), a vacuum degasser and a column oven with an MSD (G1946D) detector also present. The chromatography was performed on a Phenomenex luna C18 (2) HPLC column (100 x 4.6 mm) using a gradient method of water and acetonitrile with 0.005% trifluoroacetic acid over 28 minutes. The optimal solvent gradient for separation of target constituents were starting at 10% acetonitrile which was increased as a gradient to 30% acetonitrile over 10 minutes, then to 95% acetonitrile over 8 minutes, at a flow rate of 0.75 mL/minute and an injection volume of 5 μ L. Calibration standards of trigonelline, caffeine and chlorogenic acid were prepared in 60:40 ethanol:water, at concentrations from 0.01 to 1 mg/mL, 0.005 to 0.5 mg/mL and 0.004 to 1 mg/mL for each of these standards, respectively. Specific detection and calibration

curves for each compound were performed at 254 nm, 280 nm and 330 nm, respectively. Quantification was performed using the Chemstation Software based on reference standards, peak area and sample dilution at specific wavelengths for each compound [14].

Rats, diets and treatment

Male Wistar rats (8–9 weeks old, 330–340 g, n = 72) were obtained from the Animal Resource Centre, Perth. Rats were divided into 6 groups (n = 12 rats/group): C, corn starch diet-fed rats; CGC, corn starch diet + regular green coffee extract (5% in food); CDC, corn starch diet + decaffeinated green coffee extract (5% in food); H, high-carbohydrate, high-fat diet-fed rats; HGC, high-carbohydrate, high-fat diet-fed rats + regular green coffee extract (5% in food); and HDC, high-carbohydrate, high-fat diet-fed rats + decaffeinated green coffee extract (5% in food).

C and H rats were fed with corn starch and high-carbohydrate, high-fat diets for 16 weeks, respectively [13]. Treatment groups were fed with either corn starch (C) or high-carbohydrate, high-fat diets (H) for the first 8 weeks and then the respective diets were supplemented with treatments for the last 8 weeks. Extracts of regular green coffee or decaffeinated green coffee were prepared by mixing ground coffee (50 g) with hot water but not boiling water (100 mL) and then filtering after 5 minutes of mixing to obtain 50 mL of extract. This extract was then mixed with the food by replacing 50 mL water per kg food.

Cornstarch diet contained 570 g of corn starch, 155 g of powdered rat food, 25 g of HMW salt mixture and 250 mL of water per kilogram of diet. High-carbohydrate, high-fat diet contained 175 g of fructose, 395 g of sweetened condensed milk, 200 g of beef

tallow, 155 g of powdered rat food, 25 g of HMW salt mixture and 50 mL of water per kilogram of diet [13]. Drinking water containing 25 % (w/v) fructose was provided to all H groups. C rat groups received normal drinking water. Rats were individually housed under temperature controlled, 12-hour-light/dark conditions and given free access to food and water [13].

Physiological parameters

Body weight and food and water intakes were measured daily [13]. Abdominal circumference and body length were measured using a standard measuring tape under light anaesthesia with Zoletil (tiletamine 10 mg/kg and zolazepam 10 mg/kg, intraperitoneal; Virbac, Peakhurst, Australia) [13]. Body mass index was calculated as body weight (in grams)/[body length (in cm)]² [13]. Feed efficiency was calculated as [mean body weight gain (in grams)/daily energy intake (in kJ)] [13].

Systolic blood pressure was determined under light sedation with Zoletil (tiletamine 10 mg/kg and zolazepam 10 mg/kg, intraperitoneal; Virbac, Peakhurst, Australia), using an MLT1010 Piezo-Electric Pulse Transducer and inflatable tail-cuff connected to an MLT844 Physiological Pressure Transducer and PowerLab data acquisition unit (ADInstruments, Sydney, Australia) [13].

Dual-energy X-ray absorptiometric measurements were carried out at the end of the protocol with a Norland XR46 DXA instrument (Norland Corp, Fort Atkinson, WI). These scans were evaluated using the manufacturer's suggested software for use in laboratory animals (Small Subject Analysis Software, version 2.5.3/1.3.1; Norland Corp)

[13]. The precision error of lean mass for replicate measurements, with repositioning, is 3.2%.

Oral glucose tolerance tests were performed after determining overnight fasting blood glucose concentrations in tail vein blood using Medisense Precision Q.I.D. glucose meters (Abbot Laboratories, Bedford, MA). For overnight fasting, rats were deprived of all types of diets for 12 hours. Fructose-supplemented drinking water in H diet-fed groups was replaced with normal drinking water for the overnight food deprivation period. Rats were given a glucose load of 2 g/(kg body weight) as 40% glucose solution via oral gavage and blood glucose concentrations were measured again 30, 60, 90 and 120 minutes after oral glucose administration [13].

Euthanasia

Rats were euthanized with Lethobarb (pentobarbitone sodium, 100 mg/kg, intraperitoneal; Virbac, Peakhurst, Australia). After euthanasia, heparin (200 IU; Sigma-Aldrich Australia, Sydney, NSW, Australia) was injected through the right femoral vein. The abdomen was then opened and blood (~5 mL) was withdrawn from the abdominal aorta and collected into heparinised tubes. Blood was centrifuged at $5000 \times g$ for 10 minutes to obtain plasma. Plasma samples were stored at -20°C .

Hearts were then removed from rats for isolated Langendorff heart preparation [13]. These hearts were perfused with modified Krebs–Henseleit bicarbonate buffer bubbled with 95% O_2 –5% CO_2 and maintained at 35°C . Isovolumetric ventricular function was measured by inserting a latex balloon catheter into the left ventricle connected to a Capto SP844 MLT844 physiological pressure transducer and Chart software on a Maclab

system. Left ventricular end-diastolic pressure values were measured during pacing of the heart at 250 beats per minute using an electrical stimulator. End-diastolic pressures were obtained from 0 to 30 mmHg for the calculation of diastolic stiffness constant (κ , dimensionless) [13].

Thoracic aortic rings (~ 4 mm in length) were suspended in an organ bath filled with Tyrode physiological salt solution bubbled with 95% O₂–5% CO₂, maintained at 35°C and allowed to stabilise at a resting tension of ~10 mN. Cumulative concentration-response curves (contraction) were obtained for noradrenaline and cumulative concentration-response curves (relaxation) were obtained for acetylcholine and sodium nitroprusside after submaximal (~70%) contraction to noradrenaline [13].

After isolated heart perfusion studies, hearts were separated into left ventricles (with septum) and right ventricles and weighed. Livers were isolated and weighed. Retroperitoneal, epididymal and omental abdominal fats were removed separately and weighed. These organ weights were normalised against the tibial length at the time of organ removal and expressed as mg/mm of tibial length [13].

Histology

Hearts and livers were removed from the rats (n = 4) soon after euthanasia and these organs were fixed in 10% neutral buffered formalin. The samples were then dehydrated and embedded in paraffin wax. Thin sections (5 μ m) were cut and stained with haematoxylin and eosin stain to study infiltration of inflammatory cells ($\times 20$) and for determining fat vacuoles ($\times 40$) in the liver. Picrosirius red stain was used to study left ventricular collagen deposition ($\times 20$) [13].

Plasma biochemistry

Plasma activities of aspartate transaminase, alanine transaminase and alkaline phosphatase, and plasma concentrations of total cholesterol, triglycerides and non-esterified fatty acids were measured [13].

Statistical analysis

Data are presented as mean \pm SEM. Results were tested for variance using Bartlett's test and variables that were not normally distributed were transformed (using log 10 functions) prior to statistical analyses. Groups were tested for effects of diet, treatment and their interactions by two-way analysis of variance. When the interaction and/or the main effects were significant, means were compared using Newman-Keuls multiple comparison post hoc tests. A *P* value <0.05 was considered statistically significant. All statistical analyses were performed using GraphPad Prism version 5.0 for Windows (San Diego, CA, USA).

Results

Green coffee analysis

The contents and doses of chlorogenic acid, caffeine and trigonelline in green coffee are given in Table 1.

Table 1: Green and decaffeinated green coffee

<i>Components</i>	<i>Green coffee (mg/ml)</i>	<i>Decaffeinated green coffee (mg/ml)</i>	<i>CGC (mg/kg/day)</i>	<i>CDC (mg/kg/day)</i>	<i>HGC (mg/kg/day)</i>	<i>HDC (mg/kg/day)</i>
Chlorogenic acid	2.58	1.3	118 ± 5	55 ± 5	69 ± 2	34 ± 1
Caffeine	0.7	0	32 ± 2	0	19 ± 2	0
Trigonelline	0.85	0.9	39 ± 2	38 ± 4	23 ± 1	24 ± 1

Dietary intake, body composition and metabolic parameters

Green coffee treatment in corn starch rats (CGC, CDC rats) did not alter physiological parameters or plasma biochemistry compared to C rats (Tables 2 and 3). Green coffee treatment in HGC and HDC rats decreased the rate of body weight gain without affecting food, water or energy intakes, by reducing feed efficiency compared to H rats (Table 2). Body mass index reduced in HGC rats but did not change in HDC rats compared to H rats. However, abdominal circumference, basal or final blood glucose

concentrations, the area under the curve, plasma lipid concentrations, and total lean and fat mass did not change in HGC and HDC rats compared to H rats. Retroperitoneal fat was decreased while epididymal fat and omental fat were unchanged in HGC and HDC rats compared to H rats (Table 2).

Cardiovascular and liver function

Green coffee did not change cardiovascular or liver parameters in CGC and CGC compared to C rats (Table 2). Systolic blood pressure, left ventricular stiffness, fibrosis and inflammation were increased in H rats compared to C rats; in HGC and HDC rats, both parameters were reduced compared to H rats (Table 2). Green coffee treatments did not change fibrosis or inflammation in hearts of CGC (Figure 1B and 1H) or CDC (Figure 1C and 1I) rats compared to C rats (Figure 1A and 1G) while decreasing fibrosis and inflammation in HGC rats (Figure 1E and 1K) and HDC rats (Figure 1F and 1L) compared to H rats (Figure 1D and 1J). Green coffee treatment did not alter aortic responses in C or H rats (Figure 2A, 2B and 2C). Green coffee treatments did not change inflammation and fat deposition in livers of CGC (Figure 3B and 3H) or CDC (Figure 3C and 3I) rats compared to C rats (Figure 3A and 3G) while decreasing inflammation and fat deposition in livers of HGC rats (Figure 3E and 3K) and HDC rats (Figure 3F and 3L) compared to H rats (Figure 3D and 3J).

Table 2: Green and decaffeinated green coffee on physiological and metabolic variables

<i>Variables</i>	<i>C</i>	<i>CGC</i>	<i>CDC</i>	<i>H</i>	<i>HGC</i>	<i>HDC</i>	<i>P value</i>		
							<i>Diet</i>	<i>Treatment</i>	<i>Diet × Treatment</i>
Initial body weight, g	340 ± 1	339 ± 1	340 ± 1	340 ± 1	337 ± 1	339 ± 1	0.23	0.12	0.61
Final body weight, g	385 ± 5 ^c	384 ± 6 ^c	402 ± 9 ^c	563 ± 14 ^a	515 ± 16 ^b	522 ± 15 ^b	<0.0001	0.12	0.035
Body weight gain (8-16 weeks), %	7.7 ± 1.4 ^c	10.5 ± 0.8 ^c	7.6 ± 1.3 ^c	20.7 ± 1.8 ^a	16.0 ± 1.0 ^b	16.7 ± 1.6 ^b	<0.0001	0.33	0.028
Water intake, mL/d	25.1 ± 3.5	27.5 ± 2.6	30.2 ± 1.9	32.2 ± 2.5	34.6 ± 2.7	31.5 ± 1.1	0.054	0.56	0.42
Food intake, g/d	35.2 ± 1.3 ^a	34.2 ± 1.8 ^a	33.0 ± 0.9 ^a	27.2 ± 0.7 ^b	26.5 ± 0.9 ^b	25.5 ± 0.6 ^b	<0.0001	0.22	0.98
Energy intake, kJ/d	395 ± 15 ^b	365 ± 6 ^b	371 ± 10 ^b	612 ± 17 ^a	621 ± 22 ^a	578 ± 10 ^a	<0.0001	0.13	0.20

Feed efficiency, kJ/g	0.11 ± 0.01 ^c	0.11 ± 0.01 ^c	0.15 ± 0.02 ^c	0.37 ± 0.02 ^a	0.31 ± 0.02 ^b	0.32 ± 0.02 ^b	<0.0001	0.19	0.038
Body mass index, g/cm ²	0.63 ± 0.02 ^d	0.65 ± 0.01 ^d	0.71 ± 0.02 ^c	0.85 ± 0.02 ^a	0.77 ± 0.03 ^b	0.81 ± 0.02 ^a	<0.0001	0.19	0.038
Abdominal circumference, cm	18.8 ± 0.3 ^b	18.3 ± 0.1 ^b	18.8 ± 0.3 ^b	23.3 ± 0.7 ^a	21.7 ± 0.7 ^a	22.0 ± 0.6 ^a	<0.0001	0.19	0.038
Whole-body lean mass, g	297 ± 4	273 ± 7	312 ± 11	278 ± 8	278 ± 9	295 ± 18	0.20	0.051	0.39
Whole-body fat mass, g	68.7 ± 4.8 ^b	89.6 ± 6.9 ^b	74.1 ± 4.8 ^b	252 ± 17 ^a	209 ± 17 ^a	224 ± 29 ^a	<0.0001	0.70	0.12
Bone mineral density, g/cm ²	0.174 ± 0.003 ^b	0.168 ± 0.003 ^b	0.176 ± 0.002 ^{ab}	0.185 ± 0.003 ^a	0.182 ± 0.002 ^a	0.183 ± 0.004 ^a	<0.0001	0.24	0.52
Bone mineral content, g	11.2 ± 0.2 ^c	11.3 ± 0.3 ^c	11.7 ± 0.3 ^c	17.4 ± 0.5 ^a	15.5 ± 0.7 ^b	15.4 ± 0.8 ^b	<0.0001	0.16	0.034

Basal blood glucose concentrations, mmol/L	4.1 ± 0.2^b	4.3 ± 0.2^b	4.0 ± 0.2^b	5.1 ± 0.2^a	5.4 ± 0.3^a	4.8 ± 0.2^a	<0.0001	0.13	0.79
Blood glucose concentrations at 120 minutes, mmol/L	4.9 ± 0.1^b	4.7 ± 0.1^b	5.0 ± 0.1^b	5.7 ± 0.3^a	5.8 ± 0.1^a	6.0 ± 0.1^a	<0.0001	0.22	0.62
Area under the curve, mmol/L-minute	715 ± 20^b	661 ± 10^b	707 ± 17^b	805 ± 18^a	832 ± 12^a	778 ± 20^a	<0.0001	0.52	0.009
Retroperitoneal fat, mg/mm tibial length	136 ± 12^c	192 ± 16^c	179 ± 15^c	484 ± 54^a	380 ± 28^b	393 ± 29^b	<0.0001	0.59	0.010

Epididymal fat, mg/mm tibial length	87.2 ± 6^b	112 ± 9^b	97.2 ± 10.7^b	226 ± 23^a	220 ± 21^a	246 ± 15^a	<0.0001	0.59	0.37
Omental fat, mg/mm tibial length	130 ± 8^b	136 ± 8^b	126 ± 7^b	251 ± 16^a	224 ± 16^a	252 ± 20^a	<0.0001	0.67	0.27
Total abdominal fat, mg/mm tibial length	353 ± 18^b	439 ± 30^b	402 ± 27^b	961 ± 89^a	824 ± 56^a	892 ± 52^a	<0.0001	0.86	0.08
Plasma total cholesterol, mmol/L	1.64 ± 0.08	1.64 ± 0.08	1.53 ± 0.04	1.58 ± 0.08	1.60 ± 0.06	1.55 ± 0.05	0.64	0.45	0.83

Plasma triglycerides, mmol/L	0.53 ± 0.06 ^d	0.54 ± 0.07 ^d	0.80 ± 0.06 ^{cd}	1.30 ± 0.16 ^b	1.23 ± 0.16 ^{bc}	2.80 ± 0.22 ^a	<0.0001	<0.0001	<0.0001
Plasma NEFA, mmol/L	1.4 ± 0.2 ^b	1.4 ± 0.2 ^b	1.9 ± 0.3 ^b	3.8 ± 0.3 ^a	3.7 ± 0.3 ^a	4.4 ± 0.2 ^a	<0.0001	0.052	0.93

Values are mean ± SEM, n=6-12. Means in a row with superscripts without a common letter differ significantly, $P < 0.05$. C, corn starch diet-fed rats; CGC, corn starch diet-fed rats treated with green coffee extract; CDC, corn starch diet-fed rats treated with decaffeinated green coffee extract; H, high-carbohydrate, high-fat diet-fed rats; HGC, high-carbohydrate, high-fat diet-fed rats treated with green coffee extract; HDC, high-carbohydrate, high-fat diet-fed rats treated with decaffeinated green coffee extract.

Table 3: Green and decaffeinated green coffee on cardiovascular and liver function

<i>Variables</i>	<i>C</i>	<i>CGC</i>	<i>CDC</i>	<i>H</i>	<i>HGC</i>	<i>HDC</i>	<i>P value</i>		
							<i>Diet</i>	<i>Treatment</i>	<i>Diet × Treatment</i>
Systolic blood pressure, mmHg	128 ± 1 ^b	129 ± 1 ^b	127 ± 2 ^b	147 ± 1 ^a	130 ± 2 ^b	128 ± 2 ^b	<0.0001	<0.0001	<0.0001
Left ventricular diastolic stiffness constant (κ)	24.1 ± 1.2 ^b	22.1 ± 0.4 ^b	22.1 ± 0.5 ^b	27.5 ± 2.2 ^a	22.4 ± 0.6 ^b	21.5 ± 0.6 ^b	0.28	0.002	0.20
Left ventricular + Septum wet weight, mg/mm tibial length	20.6 ± 1	20.1 ± 1.1	22.1 ± 0.8	24.1 ± 2.0	20.3 ± 0.8	22.7 ± 0.6	0.12	0.09	0.29

Right ventricular wet weight, mg/mm tibial length	4.8 ± 0.4	4.9 ± 0.4	5.1 ± 0.2	5.5 ± 0.8	4.6 ± 0.4	4.6 ± 0.4	0.89	0.62	0.33
Liver wet weight, mg/mm	220 ± 7 ^b	225 ± 7 ^b	254 ± 5 ^b	336 ± 19 ^a	308 ± 17 ^a	329 ± 8 ^a	<0.0001	0.11	0.17
Plasma aspartate transaminase activity, U/L	69.1 ± 1.8	76.7 ± 3.2	75.2 ± 3	81.6 ± 6.1	70.7 ± 5.8	74.9 ± 5.1	0.58	0.93	0.12
Plasma alanine transaminase activity, U/L	29.8 ± 2.6	31.8 ± 2.5	30.5 ± 2.3	37.6 ± 3.4	31.9 ± 2.7	31.1 ± 2	0.19	0.54	0.26

¹Values are mean ± SEM, n=6-10. Means in a row with superscripts without a common letter differ significantly, $P < 0.05$. C, corn starch diet-fed rats; CGC, corn starch diet-fed rats treated with green coffee extract; CDC, corn starch diet-fed rats treated with decaffeinated green coffee extract; H, high-carbohydrate, high-fat diet-fed rats; HGC, high-carbohydrate, high-fat diet-fed rats treated with green coffee extract; HDC, high-carbohydrate, high-fat diet-fed rats treated with decaffeinated green coffee extract.

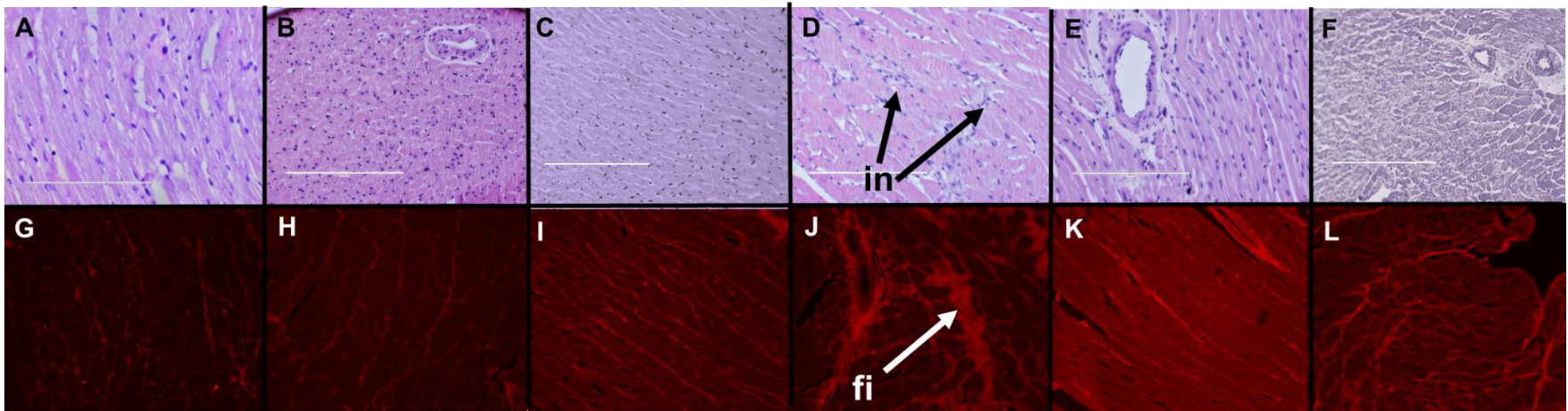


Figure 1. Effects of green coffee extracts on inflammation and fibrosis in the heart. Haematoxylin and eosin staining of left ventricle showing infiltration of inflammatory cells (A–F, inflammatory cells marked as “in”) from C (A), CGC (B), CDC (C), H (D), HGC (E) and HDC (F) rats. Picrosirius red staining of left ventricle showing collagen deposition (G–L, fibrosis marked as “fi”) from C (G), CGC (H), CDC (I), H (J), HGC (K) and HDC (L) rats. C, corn starch diet-fed rats; CGC, corn starch diet-fed rats treated with regular green coffee extract; CDC, corn starch diet-fed rats treated with decaffeinated green coffee extract; H, high-carbohydrate, high-fat diet-fed rats; HGC, high-carbohydrate, high-fat diet-fed rats treated with regular green coffee extract; HDC, high-carbohydrate, high-fat diet-fed rats treated with decaffeinated green coffee extract.

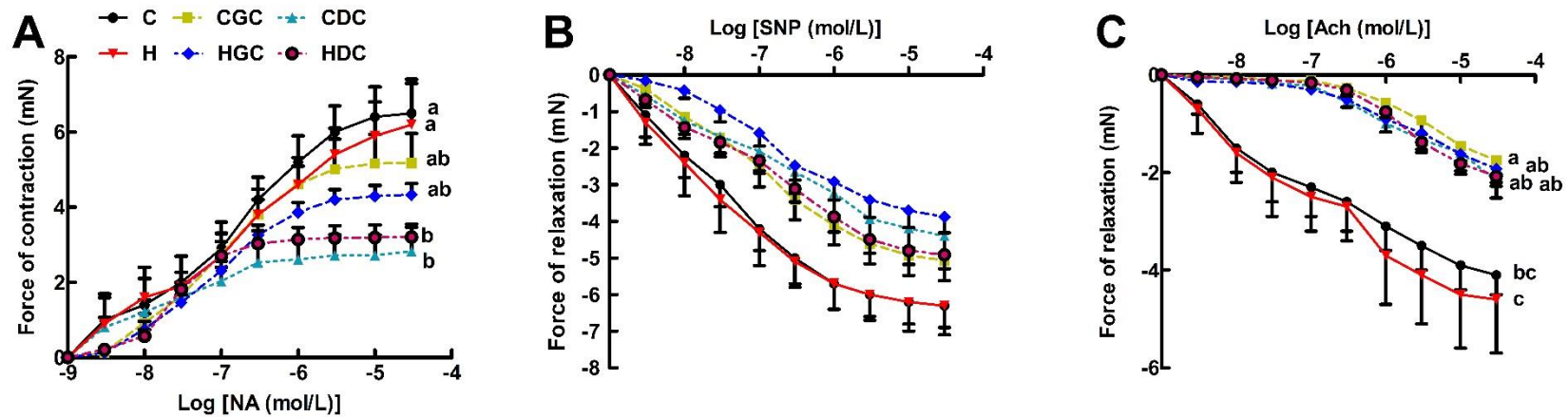


Figure 2. Effects of regular green coffee and decaffeinated green coffee extract on (A) noradrenaline-induced contraction, (B) sodium nitroprusside-induced relaxation and (C) acetylcholine-induced relaxation in thoracic aortic preparation from C, CGC, CDC, H, HGC and HDC rats. Values are mean \pm SEM, $n = 8-12$. Endpoint means without a common letter differ, $P < 0.05$. C, corn starch diet-fed rats; CGC, corn starch diet-fed rats treated with regular green coffee extract; CDC, corn starch diet-fed rats treated with decaffeinated green coffee extract; H, high-carbohydrate, high-fat diet-fed rats; HGC, high-carbohydrate, high-fat diet-fed rats treated with regular green coffee extract; HDC, high-carbohydrate, high-fat diet-fed rats treated with decaffeinated green coffee extract.

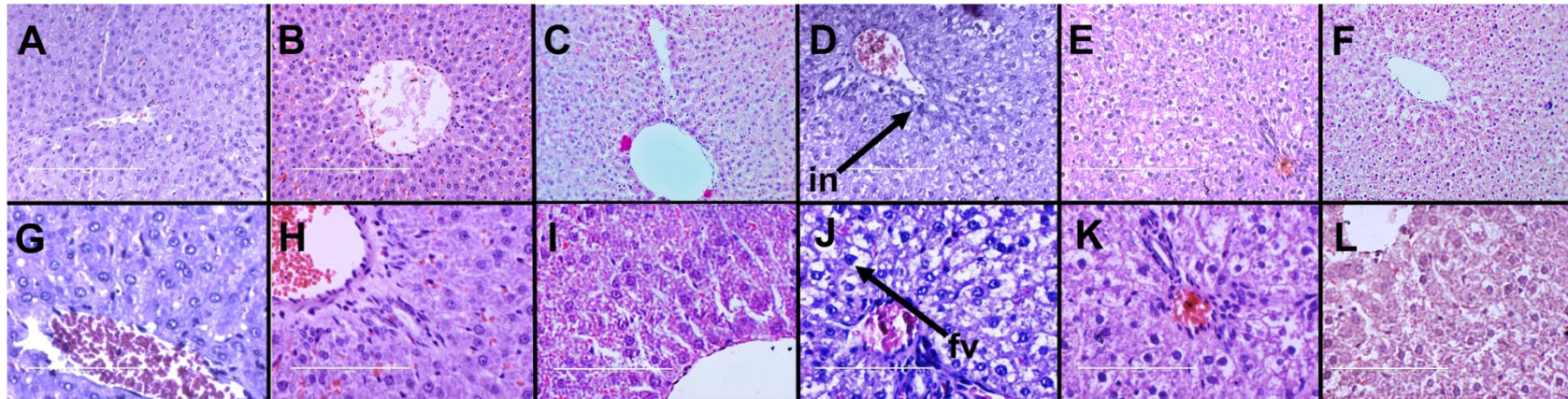


Figure 3. Effects of green coffee extracts on inflammation and fat deposition in the liver. Haematoxylin and eosin staining of liver showing infiltration of inflammatory cells (A–F, inflammatory cells marked as “in”) from C (A), CGC (B), CDC (C), H (D), HGC (E) and HDC (F) rats and fat deposition (G–L, enlarged fat vacuoles as “fv”) from C (G), CGC (H), CDC (I), H (J), HGC (K) and HDC (L) rats. C, corn starch diet-fed rats; CGC, corn starch diet-fed rats treated with regular green coffee extract; CDC, corn starch diet-fed rats treated with decaffeinated green coffee extract; H, high-carbohydrate, high-fat diet-fed rats; HGC, high-carbohydrate, high-fat diet-fed rats treated with regular green coffee extract; HDC, high-carbohydrate, high-fat diet-fed rats treated with decaffeinated green coffee extract.

Discussion

In this study, green coffee extract and decaffeinated green coffee extract decreased body weight and visceral fat accumulation, and improved heart and liver structure and function in obese, hypertensive rats fed a high-carbohydrate, high-fat diet, without improving glucose tolerance or dyslipidaemia. High-carbohydrate, high-fat diet-fed rats are an appropriate preclinical model for mimicking human metabolic syndrome as they develop relevant symptoms including visceral obesity, impaired glucose tolerance, dyslipidaemia, hypertension, cardiovascular remodelling and liver abnormalities [13].

Obesity is the critical factor of metabolic syndrome associated with high calorie intake and sedentary life lifestyle [15, 16]. In contrast to the few studies in green coffee, roasted (black) coffee has been more widely studied so that it has been proposed as a functional food to reverse the health complications caused by obesity [17]. Coffee contains around 1000 possibly biologically active components, including caffeine, chlorogenic acid, caffeic acid, hydroxyhydroquinone, trigonelline, diterpenes such as kahweol and cafestol, lignin and melanoidins [18]. Chronic black coffee consumption has been associated with a reduction in the development of chronic diseases such as cardiac disorders [19], hepatic diseases [20] and type 2 diabetes [21]. The anti-obesity effect of black coffee has been associated with reduced lipid accumulation in cells by reduction of adipogenesis and modifications of transcription factors and lipogenesis-related proteins in the adipose tissue of animal models [22] but could also be associated with altered glucose homeostasis, antioxidant activity and inflammatory biomarkers [21]. Individual components including caffeine [23], chlorogenic acid [24] and trigonelline [25] may improve glucose tolerance and reduce the risk of type 2 diabetes progression [26] but the

diterpenes may increase serum lipid concentrations in humans [27]. However, the type of coffee and the method of preparation determines the effects of coffee on lipoproteins [28]. In our previous study, roasted coffee extract in obese rats improved cardiovascular and hepatic structure and function with no improvements in obesity or blood concentrations of triglycerides and non-esterified fatty acids [23]. In contrast, caffeine at the same dose as in coffee extract decreased body weight and abdominal obesity [29]. This clearly indicated that caffeine was only partly contributing to the responses from coffee extract.

This study determined whether the unroasted green coffee has potential for health improvement, rather than black coffee. Decaffeinated green coffee supplementation reduced postprandial hyperglycaemia in rats and humans [30]. An intake of 200 mg/kg/day green coffee extract for 6 weeks reduced obesity and dyslipidaemia in mice [31] and anti-inflammatory effects of green coffee extract in rats were demonstrated by decreasing lipopolysaccharide-induced leukocyte migration in the peritonitis test [32]. Likewise, decaffeinated green coffee reduced obesity and fat accumulation with improving insulin sensitivity by 0.3% diet supplementation for 11 weeks in high-fat diet-fed mice [33]. In humans, a randomised crossover study showed that green coffee consumption reduced body weight, body mass index, fat accumulation and hypertension [34]. A 22-week crossover study of green coffee extract in overweight humans reduced body weight and heart rate, and body mass index shifted back to normal [34]. Meta-analysis of human clinical trials showed reduced body weight with green coffee extract compared with placebo [10].

The observation in our study was that green coffee extract reduced feed efficiency. The reduced feed efficiency indicates higher satiety level from diet supplemented with

green coffee. This outcome pointed out that green coffee extract delivered a similar energy intake as their control, and revealed a decrease in feed efficiency and hence, body weight. The decreased feed efficiency could result from elevated energy expenditure and thermogenesis. Chlorogenic acid, the hydroxycinnamic acid derivative present in green coffee extract, enhanced energy expenditure through the upregulation of adenosine monophosphate activated protein kinase (AMPK) by inhibiting cAMP phosphodiesterase, with the subsequent increase in fatty acid oxidation [35]. The proposed mechanism of action is the inhibition of the enzyme pancreatic lipase, which leads in a decrease in lipid absorption [36] resulting in weight reduction.

Green coffee contains a wide range of potentially bioactive compounds. In this study, green coffee contained higher concentrations of chlorogenic acid compared to the decaffeinated green coffee with no caffeine present in decaffeinated green coffee. Trigonelline was equivalent in the two types of green coffee. As the responses of green and decaffeinated green coffee were quite similar, caffeine could be ruled out for contribution towards the responses observed in obese rats. Further, the dose of chlorogenic acid was almost half in decaffeinated green coffee (~34 mg/kg/day) compared to green coffee (~69 mg/kg/day). In addition, previous studies used a dose of 100 mg/kg/day of chlorogenic acid to suggest its beneficial effects [37, 38]. This suggests that chlorogenic acid is one of several active components in green coffee to attenuate metabolic syndrome.

Trigonelline is also a putative bioactive compound in green coffee with the dose around 25 mg/kg/day from both types of green coffee interventions. A higher dose of trigonelline (40 mg/kg/day) for 48 weeks in a high-carbohydrate, high-fat diet reduced body weight, lowered blood glucose and HbA1c via increasing serum GLP-1

concentrations and GLP-1R expression in rats [39]. At the same dose, trigonelline for 4 weeks in rats improved insulin sensitivity by reducing the availability of free fatty acids, TNF α and IL-6 in serum [40]. A limitation of our study was the limited availability and high cost of pure trigonelline to test its responses.

The responses of green and decaffeinated coffee suggest that chlorogenic acid and trigonelline may have additive or synergistic effects towards attenuating symptoms of metabolic syndrome. Further, coffee also contains diterpenoids, cafestol and kahweol, that were not measured and may have their role in either improving metabolic complications [41] or have detrimental effects as in previous study where non-filtered, boiled coffee containing increased diterpenoids increased serum cholesterol concentrations and hence the risk of cardiovascular disease [42].

Conclusions

Green coffee with or without caffeine attenuated obesity and reduced cardiovascular disease components without changes in glucose homeostasis and plasma lipid concentrations in high-carbohydrate, high-fat diet-induced metabolic syndrome in rats. Both green coffee products improved structure and function of the heart and liver in these rats. Results from this study suggest that chlorogenic acid and trigonelline, rather than caffeine, may serve as the bioactive components of green coffee. Following well-designed human studies with green coffee [8, 43], green coffee may serve as a simple and safe dietary supplement to manage weight and blood pressure that may prove very effective in combination with healthy lifestyles in obese hypertensive patients.

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Authors' contribution: S.K.P. and L.B. designed the research protocol; S.K.P., L.B., and N.B. interpreted the results; S.K.P. and N.B. conducted the animal experiments and analysed the data; S.K.P., L.B., and N.B. wrote the manuscript; P.M. analysed the components of coffee samples; and S.K.P. had primary responsibility for the final content. All authors read and approved the final manuscript.

Compliance with ethical Standards

Conflict of Interest The authors declare that they have no conflict of interest.

Statement on the Welfare of Animals All experimental protocols were approved by the University of Southern Queensland Animal Ethics Committee under the guidelines of the National Health and Medical Research Council of Australia. The approval number for this project was 13REA005. This article does not contain any studies with human participants performed by any of the authors.

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Chapter 3.

Chlorogenic acid attenuates metabolic syndrome in diet-induced obese rats

Chlorogenic acid attenuates metabolic syndrome in diet-induced obese rats

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Abstract

Chlorogenic acid as a constituent of coffee is consumed regularly in the human diet. Chlorogenic acid intake has been associated with decreased risk of cardiovascular disease and type 2 diabetes. This study investigated whether chlorogenic acid improves cardiovascular, liver and metabolic responses in a diet-induced rat model of metabolic syndrome induced by a high-carbohydrate, high-fat diet. Male Wistar rats (8-9 week old, 335 ± 5 g, $n = 48$) were divided into 4 groups and fed with either corn starch diet (16 weeks), corn starch diet with 2 g/kg chlorogenic acid in food for the last 8 weeks, high-carbohydrate, high-fat diet (16 weeks), or high-carbohydrate, high-fat diet with 2 g/kg chlorogenic acid in food for the last 8 weeks. In high-carbohydrate, high-fat diet-fed rats, chlorogenic acid reduced energy intake and food efficiency to reduce visceral fat (especially retroperitoneal fat) and abdominal circumference; reversed the elevated systolic blood pressure and attenuated left ventricular diastolic stiffness while reducing collagen deposition and infiltration of inflammatory cells in the left ventricle. Chlorogenic acid decreased inflammation and fat deposition in the liver along with reduction in plasma liver enzyme activities of obese rats but did not change the plasma lipid profile. These results suggest that chronic dietary chlorogenic acid may attenuate inflammation as well as cardiovascular, liver, and metabolic abnormalities induced by high-carbohydrate, high-fat diet.

Introduction

Coffee, a complex mixture of almost 1000 compounds, may decrease the risk of developing metabolic syndrome, for example, reducing the risk of type 2 diabetes and decreasing triglyceride concentrations [1]. An important component of coffee is a class of compounds known as chlorogenic acids, which are isomeric hydroxycinnamic esters of quinic acid [2]; one of the major compounds from this class is 5-caffeoyl-quinic acid [3]. Closely related compounds are also present in coffee beans, including the 3- and 4-isomers (isochlorogenic acids A and B), as well as 3,4-, 3,5-, and 4,5-diquinyl esters of cinnamic acid [4]. Chlorogenic acid intake is common and the content in one cup of coffee varies between 70 and 350 mg [3]. In the stomach, chlorogenic acid is absorbed directly whereas in the small and large intestines, it is broken down to caffeic acid and quinic acid before absorption [3].

Patients with metabolic syndrome have an increased risk of developing cardiovascular disease and diabetes [5]. Consumption of chlorogenic acids has been associated with decreased risk of these two chronic and common diseases [3]. These therapeutic responses to chlorogenic acids have been attributed to their antioxidant and anti-inflammatory properties [3, 6], modulation of glucose and lipid metabolism [7] by increased mRNA expression of PPAR- α and liver X receptor α as key mediators of lipid metabolism [8, 9], inhibition of glucose-6-phosphatase activity to reduce hepatic glucose output [10], and decreased expression of sodium-dependent glucose transporters in brush border membranes of the small intestine to reduce glucose absorption [11].

Our previous study with coffee extract in high-carbohydrate, high-fat diet-fed rats did not show any changes in obesity but improved heart and liver structure and function [12]. Following this, we matched the dose of caffeine from the coffee extract study to show that this dose of caffeine reversed obesity along with other improvements from coffee extract [13].

Thus, the current study investigated whether chlorogenic acid has beneficial or opposing effects to caffeine during an 8-week diet intervention in rats fed a high-carbohydrate, high-fat diet. After treatment with chlorogenic acid, metabolic parameters related to obesity and glucose tolerance, and the structure and function of the cardiovascular system and liver were examined. Cardiovascular function was studied in isolated Langendorff heart preparations, isolated thoracic aortic rings and by measurement of systolic blood pressure together with histopathology. Liver structure and function were measured by plasma biochemistry and histopathology.

Methods and materials

Rats, diets, and treatments

All experimental protocols were approved by the University of Southern Queensland Animal Ethics Committee under the guidelines of the National Health and Medical Research Council of Australia. Male Wistar rats (8–9 weeks old, 340 ± 2 g, $n = 48$) were obtained from Animal Resource Centre, Murdoch, WA, Australia. Rats were randomly divided into 4 groups for the 16 week feeding protocol: corn starch diet-fed rats (C, $n = 12$), corn starch diet + chlorogenic acid (CC, $n = 12$; 2 g/kg of food for the final 8 weeks); high-carbohydrate, high-fat diet-fed rats (H; $n = 12$); and high-carbohydrate, high-fat diet + chlorogenic acid (HC; $n = 12$, 2 g/kg of food for the final 8 weeks). C diet contained 570 g corn starch, 155 g powdered rat food, 25 g HMW salt mixture, and 250 g water per kilogram of diet. H diet contained 175 g fructose, 395 g sweetened condensed milk, 200 g beef tallow, 155 g powdered rat food, 25 g HMW salt mixture, and 50 g water per kilogram of diet [14]. Drinking water with 25% w/v fructose was provided to H and HC groups. C and CC rats were given normal drinking water. All the rats were individually housed under temperature-controlled, 12-hour-light/dark conditions and given *ad libitum* access to food and water [14].

Physiological parameters

Body weight, food, and water intakes were measured daily [14]. Abdominal circumference and body length (nose to anus) were measured using a standard measuring tape under light anesthesia with Zoletil (tiletamine 10 mg/kg, zolazepam 10 mg/kg, intraperitoneally) [14]. Body mass index was calculated as body weight (in grams)/[body length (in cm)]² [14]. Feed efficiency was calculated as [mean body weight gain (in grams)/daily energy intake (in kJ)] [14].

Systolic blood pressure measurements

Systolic blood pressure was determined under light sedation with Zoletil (tiletamine 10 mg/kg, zolazepam 10 mg/kg, intraperitoneally), using an MLT1010 Piezo-Electric Pulse Transducer and inflatable tail-cuff connected to an MLT844 Physiological Pressure Transducer and PowerLab data acquisition unit [14].

Body composition measurement

Dual-energy X-ray absorptiometric measurements were carried out with a Norland XR36 DXA instrument (Norland Corp, Fort Atkinson, WI). These scans were evaluated using the manufacturer's suggested software for use in laboratory animals (Small Subject Analysis Software, version 2.5.3/1.3.1; Norland Corp). The precision error of lean mass for replicate measurements, with repositioning, was 3.2% [15]. Visceral adiposity index (%) was analyzed as ([retroperitoneal fat (g) + omental fat (g) + epididymal fat (g)] / [body weight (g)]) × 100.

Oral glucose tolerance test

Oral glucose tolerance tests were performed on rats following a 12-hour food deprivation during which fructose-supplemented drinking water in H and HC groups was

replaced with normal drinking water [14]. After determining basal blood glucose concentrations in tail vein blood using Medisense Precision Q.I.D. glucose meters, rats were given a glucose load of 2 g/kg body weight as 40% glucose solution via oral gavage and blood glucose concentrations were measured again 30, 60, 90, and 120 minutes after oral glucose administration [14].

Terminal experiments

Rats were euthanased with Lethobarb (pentobarbitone sodium, 100 mg/kg, intraperitoneally). After euthanasia, heparin (200 IU) was injected through the right femoral vein. The abdomen was then opened and blood (~5 mL) was withdrawn from the abdominal aorta and collected into heparinized tubes. Blood was centrifuged at $5000 \times g$ for 15 minutes to obtain plasma. Plasma was stored at -20°C for further characterization. Hearts were then removed from rats for isolated Langendorff heart preparation [14].

Isolated Langendorff heart preparation

Hearts isolated from euthanased rats were perfused with modified Krebs–Henseleit bicarbonate buffer bubbled with 95% O_2 –5% CO_2 and maintained at 35°C . Isovolumetric ventricular function was measured by inserting a latex balloon catheter into the left ventricle connected to a Capto SP844 MLT844 physiological pressure transducer and Chart software on a Maclab system. All left ventricular end-diastolic pressure values were measured during pacing of the heart at 250 beats per minute using an electrical stimulator. End-diastolic pressures were obtained from 0 to 30 mmHg for the calculation of diastolic stiffness constant (κ , dimensionless) [14].

Isolated tissue reactivity

Thoracic aortic rings (~4 mm in length) were suspended in an organ bath filled with Tyrode physiological salt solution bubbled with 95% O₂–5% CO₂ maintained at 35°C and allowed to stabilize at a resting tension of ~10 mN. Cumulative concentration–response curves (contraction) were obtained for noradrenaline and cumulative concentration–response curves (relaxation) were obtained for acetylcholine and sodium nitroprusside after submaximal (~70%) contraction to noradrenaline [14].

Organ weights

After isolated heart perfusion studies, the heart was separated into left ventricle (with septum) and right ventricle and weighed separately. The liver was isolated and weighed. Retroperitoneal, epididymal, and omental abdominal fat pads were removed separately and weighed. These organ weights were normalized against the tibial length at the time of organ removal and expressed as mg/mm of tibial length [14].

Histology

Heart and liver were removed from rats soon after euthanasia and fixed in 10% neutral buffered formalin. The samples were then dehydrated and embedded in paraffin wax. Thin sections (~5 μ m) of heart and liver were cut and stained with hematoxylin and eosin to study infiltration of inflammatory cells and for determining fat vacuoles in liver. Heart sections were also stained with picosirius red stain to study collagen distribution [14].

Plasma biochemistry

Plasma activities of aspartate transaminase (AST) and alanine transaminase (ALT) and concentrations of total cholesterol, triglycerides, and non-esterified fatty acids (NEFA) in plasma were measured [14].

Statistical analysis

All data are presented as mean \pm SEM. Results were tested for variance using Bartlett's test and variables that were not normally distributed were transformed (using log 10 functions) prior to statistical analyses. C, CC, H, and HC groups were tested for effects of diet, treatment, and their interactions by 2-way Analysis of Variance. When the interaction and/or the main effects were significant, means were compared using Newman-Keuls multiple comparison *post hoc* tests. A *P* value <0.05 was considered significant. All statistical analyses were performed using GraphPad Prism version 5.0 for Windows (San Diego, CA, USA).

Results

Dietary intake, body composition, and plasma biochemistry

At the end of 16 weeks, H rats had higher body weight compared to C rats while chlorogenic acid reduced the body weight in HC rats (Table 1). Water intakes were similar between C and CC rats and between H and HC rats. Food intakes was lower in H rats compared to C rats. Chlorogenic acid increased food intake in CC rats compared to C rats while it did not change food intake in HC rats compared to H rats. The food intake in CC rats was higher than HC rats; hence, the chlorogenic acid intake was higher in CC rats compared to HC rats. Energy intake was higher in H rats than in C rats. Chlorogenic acid increased energy intake in CC rats compared to C rats while it did not change energy intake in HC rats compared to H rats (Table 1). Feed efficiency, body mass index, and abdominal circumference were higher in H rats than

in C rats whereas chlorogenic acid reduced these parameters in HC rats compared to H rats with no effects in CC rats compared to C rats on these parameters at the end of 16 weeks (Table 1). Whole-body lean mass was similar in all diet groups. Whole-body fat mass was higher in H rats than in C rats and was decreased with chlorogenic acid in HC rats compared to H rats (Table 1). Basal blood glucose concentrations were higher in H rats than in C rats and reduced by chlorogenic acid in both HC and CC rats compared to H and C rats, respectively. However, chlorogenic acid failed to reduce the 120-minute blood glucose concentrations and hence area under the curve during oral glucose tolerance test (Table 1). Plasma total cholesterol concentrations were unchanged between the groups while chlorogenic acid was unable to reduce the plasma concentrations of triglycerides and non-esterified fatty acids in CC or HC rats (Table 1). Retroperitoneal, omental, epididymal, and total abdominal fat pads were higher in H rats than in C rats. These fat depots were unchanged in CC rats compared to C rats while chlorogenic acid reduced total abdominal fat through reduction in retroperitoneal fat in HC rats compared to H rats. Visceral adiposity followed the same trend between the groups as total abdominal fat (Table 1).

Cardiovascular structure and function

Histopathological analysis of left ventricle indicated increased infiltration of inflammatory cells in H rats (Figure 1C) compared to C rats (Figure 1A). Chlorogenic acid reduced infiltration of inflammatory cells into the left ventricle of HC rats (Figure 1D) than in H rats. Picrosirius red staining of left ventricle suggested higher collagen deposition in H rats (Figure 1G) compared to C rats (Figure 1E). Chlorogenic acid reduced collagen deposition in the left ventricle of HC rats (Figure 1H) than in H rats. Noradrenaline-induced contraction and acetylcholine-induced relaxation were lower in H rats than in C rats and chlorogenic acid treatment did not change these responses (Figure 2A and 2C). There were no differences in sodium nitroprusside-induced vascular relaxation in isolated thoracic aortic rings between the

groups (Figure 2B). Systolic blood pressure and diastolic stiffness were higher in H rats than in C rats and chlorogenic acid reduced both systolic blood pressure and diastolic stiffness in HC rats compared to H rats (Table 1). LV and RV wet weights were unchanged between the groups (Table 1).

Liver structure and function

Increased inflammatory cell infiltration and fat deposition were observed in H rats (Figure 1K) than in C rats (Figure 1I). Chlorogenic acid reduced infiltration of inflammatory cells and fat deposition in livers of HC rats (Figure 1L) compared to H rats. Liver wet weight was higher in H rats than in C rats and it was unchanged by chlorogenic acid treatment in both CC and HC rats (Table 1). Plasma ALT activities were higher in H rats compared to C rats and reduced in HC rats. Plasma AST activity was unchanged between H and C rats and reduced in HC rats compared to H rats (Table 1).

Discussion

In this study, chlorogenic acid decreased body weight and visceral fat accumulation, and improved heart and liver structure and function in obese, hypertensive rats fed a high-carbohydrate, high-fat diet, without improving glucose tolerance. These responses of chlorogenic acid, one of the major components of coffee, are an important aspect in functional foods research as many people consume chlorogenic acid and related compounds on a daily basis through diet, mainly coffee. Further, the responses to chlorogenic acid in this study were obtained in a high-carbohydrate, high-fat diet-fed rats as an appropriate model for human metabolic syndrome developing similar symptoms including visceral obesity along with impaired glucose tolerance, dyslipidemia, hypertension, endothelial dysfunction, and liver abnormalities [14].

Coffee consumption in human studies, mostly cross-sectional analyses, has been associated with decreases in components of metabolic syndrome including waist circumference, systolic and diastolic blood pressures, and blood concentrations of triglyceride and fasting glucose [1]. Coffee consumption has also been associated with decreased risk of type 2 diabetes and cardiovascular disease [1]. Coffee is a widely-used source of bioactive compounds such as caffeine, chlorogenic acid, trigonelline, cafetstol, and kahweol [1]. Caffeine and chlorogenic acid are two major components of coffee that have been associated with beneficial health effects, especially against metabolic syndrome [1]. Our previous study with coffee extract treatment in obese rats (5% coffee extract in food for 8 weeks) showed improved cardiovascular and hepatic structure and function with no improvements in obesity or blood concentrations of triglycerides and NEFA [12]. Caffeine treatment (~28 mg/kg/day) in obese rats for 8 weeks showed improved structure and function of heart and liver, with reduction in obesity but no improvements in blood concentrations of cholesterol, triglycerides, and NEFA [13]. Caffeine inhibited A₁-adenosine receptors in the hypothalamus to suppress appetite and increase energy expenditure in order to reduce diet-induced obesity in mice [16].

Chlorogenic acid has been associated with decreases in blood pressure and improvements in endothelial function, lipid metabolism, and carbohydrate metabolism [3, 17, 18]. The dose of chlorogenic acid in the high-carbohydrate, high-fat diet-fed rats was ~100 mg/kg in this study. In mice, no improvements in body weight or insulin resistance were observed with lower doses of ~50 mg/kg/day for 12 weeks or ~30 mg/kg/day for 8 weeks [19, 20]. The doses of CGA used in our study are comparable with those obtained in a typical cup of coffee. However, the relationship is not straightforward as these values fluctuate markedly depending on the type of coffee and the preparation method [21]. Typical values in green coffee bean are of the order of 8 % for CGA so the doses used in our study are achievable from the daily consumption levels.

Intestinal absorption of chlorogenic acid has been controversial. In some studies, chlorogenic acid has been found in urine after coffee or pure chlorogenic acid consumption [22-26] while chlorogenic acid was not detected in other studies in plasma, urine, or bile [27-33]. One study suggested direct absorption of chlorogenic acid in the gut of 8% [27]. Further, metabolism of chlorogenic acid is quite variable in different individuals and it may cause the variation in the responses to chlorogenic acid [34]. After ingestion of coffee, various derivatives of chlorogenic acid were found in the urine. Further, chlorogenic acid was broken down to caffeic acid and hence derivatives of caffeic acid were also found in the urine [35]. These metabolites derived from chlorogenic acid are further metabolized into compounds including *m*-coumaric acid and hydroxy-derivatives of phenylpropionic, benzoic and hippuric acids [26, 36, 37]. Chlorogenic acids and their metabolites scavenge free radical oxygen species, with dihydrocaffeic acid showing the strongest effect against oxidative stress-induced inflammation [38]. Further, metabolites of chlorogenic acid, but not chlorogenic acid itself, improved the growth of *Bifidobacterium* spp. and moderated the *Firmicutes/Bacteroidetes* ratio [39]. These changes in gut microbiota may modulate the metabolic changes observed in obesity. This may be one of mechanisms of action of chlorogenic acid in improving metabolic changes in obesity in this study.

In our study, chlorogenic acid did not reverse the glucose intolerance in obese rats. In a previous study, 90 mg/kg/day chlorogenic acid in Sprague–Dawley rats improved fasting glucose, glucose tolerance, and fasting serum insulin concentrations through differential expression of SGLT1, GLUT2, and proglucagon in the intestinal segments [40]. However, this study confirmed the outcomes from our study in terms of body weight and abdominal fat reduction [40].

We suggest that the reduced inflammation following chlorogenic acid treatment decreased the signs of the metabolic syndrome in diet-induced obese rats. In animal models, high-fat diets are associated with increased chylomicrons in the intestine, which promote systemic uptake of gut microbiota-derived lipopolysaccharides causing metabolic endotoxemia to trigger the low-grade inflammation that characterizes obesity [41, 42]. In obese humans, circulating endotoxin was 20% higher than in lean patients and 125% higher in a type 2 diabetic patient compared to lean individual [43]. Diets rich in fat or carbohydrate triggered systemic lipopolysaccharides secretion and increased expression of TLR-4, NF- κ B, TNF- α , and IL-6, which are components engaged in pathways that also participate in the regulation of insulin secretion [44]. Thus, further studies on finding specific targets for the suppression of inflammation by chlorogenic acid are warranted.

Increased plasma ALT and AST activities, inflammatory cell infiltration and lipid deposition in the liver are important markers of fatty liver [45]; these changes were reduced by chlorogenic acid in this study. A possible explanation could be the activation of AMP kinase [46]. AMP kinase is an energy-sensing enzyme with a crucial role in the regulation of carbohydrate and lipid metabolism in response to cellular stress [47]. Chlorogenic acid as an anti-inflammatory agent reduced liver fibrosis by decreasing inflammatory cytokines, TLR-4, iNOS, COX-2, and NF- κ B [48]. Further, chlorogenic acid intervention attenuated diet-induced liver abnormalities by downregulating the expression of the PPAR- γ to reduce fibrosis and accumulation of fat [49] with decreased expression of macrophage mRNA such as TNF- α , MCP-1, and chemokines associated with inflammatory cascade pathways. Consistent with our results, chlorogenic acid reduced fat deposition in adipose tissue by decreasing mRNA expression of the lipogenic enzymes (fatty acid synthase, acetyl-CoA carboxylase, and stearoyl-CoA desaturase) in the liver [50]. Overall, the reduction of expression in

inflammation-associated proteins could attenuate lipid-related disorders such as obesity, hypertension, and metabolic abnormalities.

In conclusion, high-carbohydrate, high-fat-diet-induced metabolic syndrome in rats was attenuated by chlorogenic acid. As coffee is the major dietary source of chlorogenic acid, this compound could be one of the most effective bioactive constituents in coffee leading to improved cardiovascular, liver, and metabolic functions. Production of chlorogenic acid-enriched coffee could be an effective intervention strategy with improved compliance. Thus, a clinical trial with chlorogenic acid-enriched coffee in mildly hypertensive, overweight, or obese patients is warranted.

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Figure legends

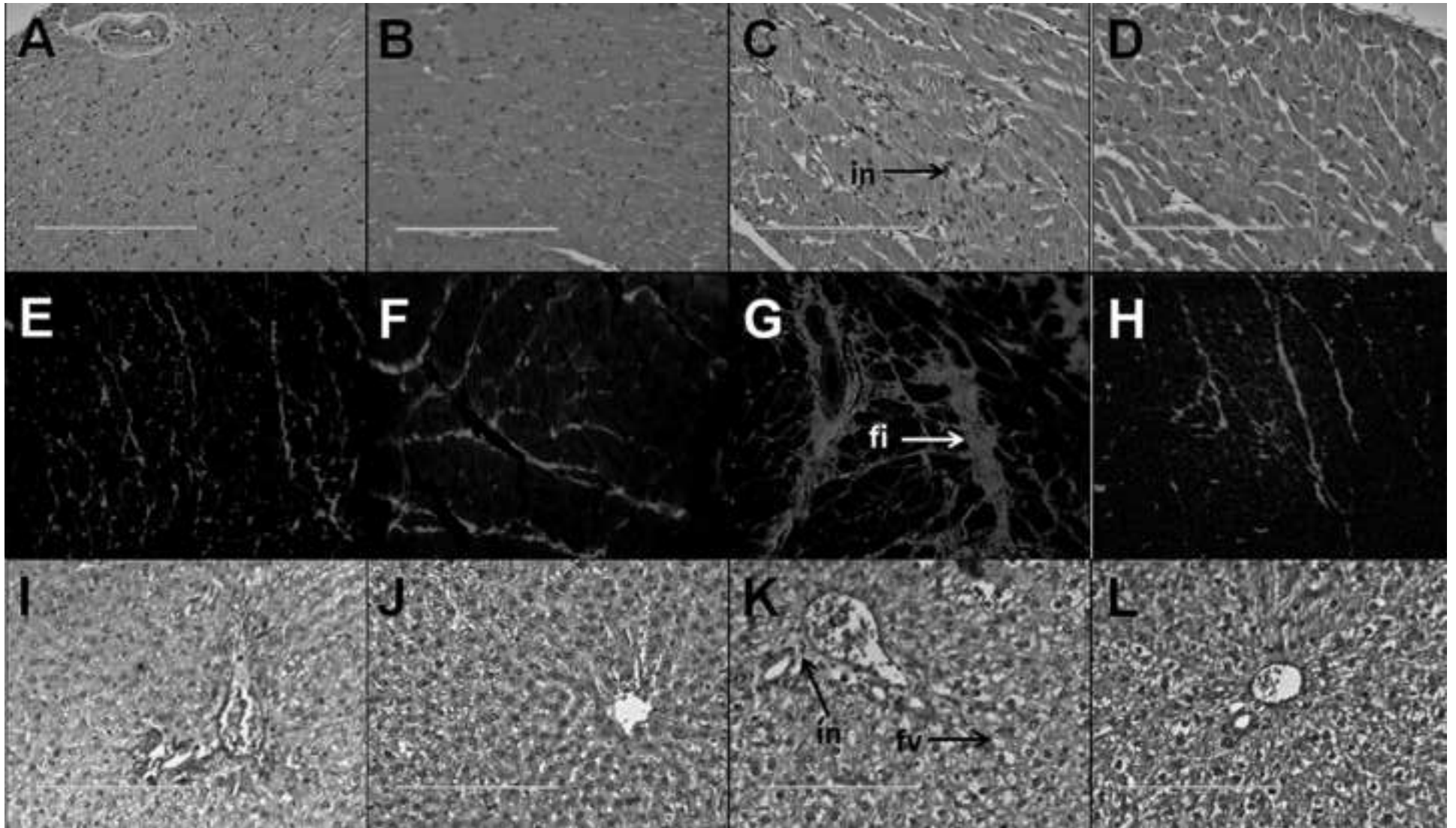
Figure 1. Effects of chlorogenic acid on structure of heart and liver at the end of 16 weeks. Hematoxylin and eosin staining of left ventricle showing inflammatory cells (“in”) in corn starch diet-fed rats (*A*), corn starch diet-fed rats supplemented with chlorogenic acid (*B*), high-carbohydrate, high-fat diet-fed rats (*C*), and high-carbohydrate, high-fat diet-fed rats supplemented with chlorogenic acid (*D*). Picrosirius red staining of left ventricle showing fibrosis (“fi”) in corn starch diet-fed rats (*E*), corn starch diet-fed rats supplemented with chlorogenic acid (*F*), high-carbohydrate, high-fat diet-fed rats (*G*), and high-carbohydrate, high-fat diet-fed rats supplemented with chlorogenic acid (*H*). Hematoxylin and eosin staining of liver showing enlarged fat vacuoles (“fv”) and inflammatory cells (“in”) from rats fed with corn starch diet (*I*), corn starch diet-fed rats treated with chlorogenic acid (*J*), high-carbohydrate, high-fat diet-fed rats (*K*), and high-carbohydrate, high-fat diet-fed rats treated with chlorogenic acid (*L*).

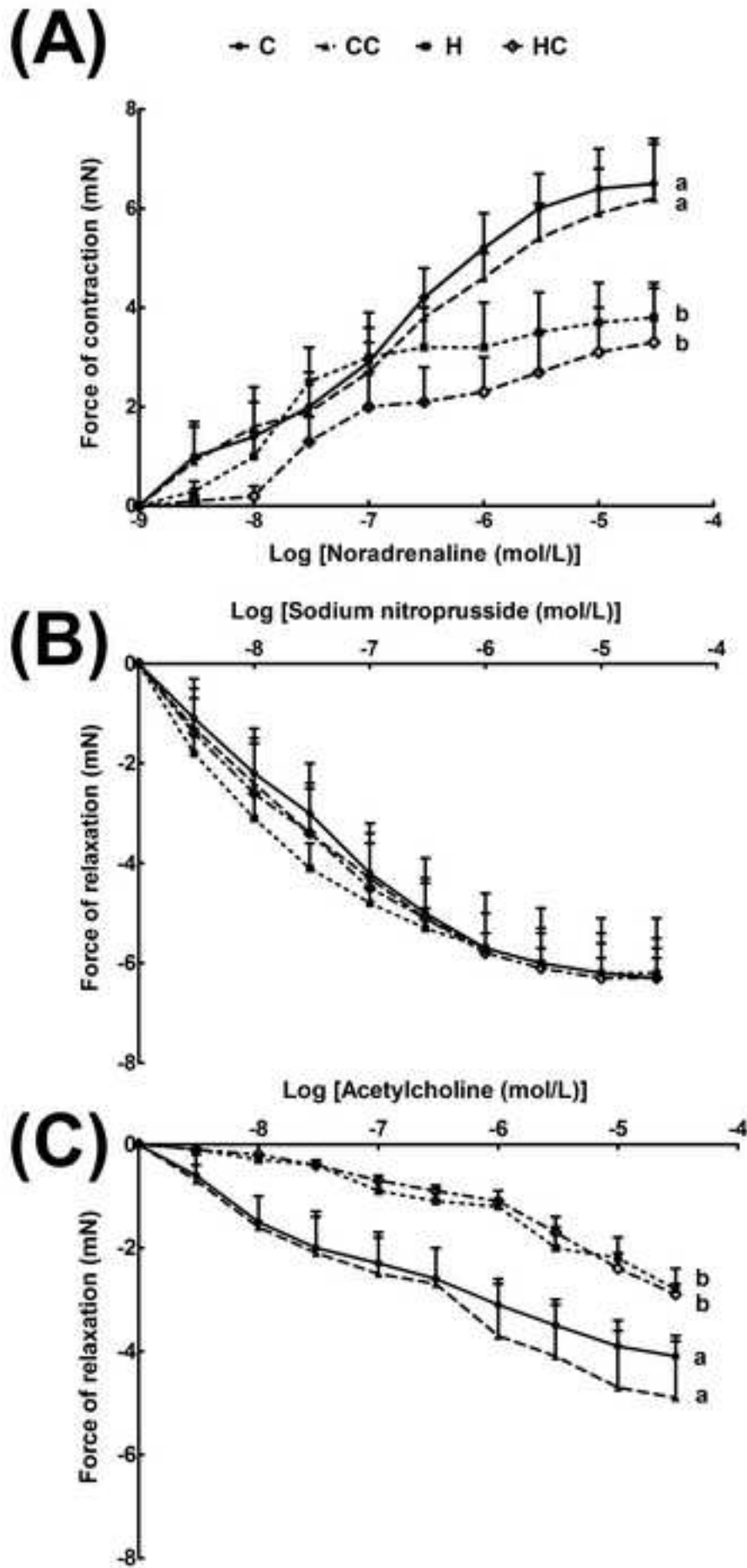
Figure 2. Effects of chlorogenic acid on thoracic aortic responses to noradrenaline (*A*), sodium nitroprusside (*B*), and acetylcholine (*C*) at the end of 16 weeks. Values are presented as mean \pm SEM ($n = 8-10$). End-point means without a common alphabet significantly differ, $P < 0.05$. *C*, corn starch diet-fed rats; *CC*, corn starch diet-fed rats treated with chlorogenic acid; *H*, high-carbohydrate, high-fat diet-fed rats; *HC*, high-carbohydrate, high-fat diet-fed rats treated with chlorogenic acid.

Table 1. Effects of Chlorogenic acid treatment on physiological, compositional, and metabolic parameters at the end of 16 weeks

Variables	C	CC	H	HC	P value		
					Diet	Chlorogenic acid	Diet × Chlorogenic acid
Initial body weight, g	342 ± 1	341 ± 2	340 ± 1	340 ± 2	0.34	0.75	0.75
Final body weight, g	385 ± 5 ^c	380 ± 5 ^c	563 ± 14 ^a	498 ± 13 ^b	<0.0001	0.002	0.006
Water intake, mL/d	27.3 ± 0.8 ^a	27.2 ± 1.1 ^a	24.7 ± 0.8 ^{ab}	23.6 ± 0.8 ^b	0.001	0.50	0.58
Food intake, g/d	32.3 ± 0.8 ^b	36.1 ± 0.9 ^a	25.1 ± 0.9 ^c	23.4 ± 0.6 ^c	<0.0001	0.20	0.002
Chlorogenic acid intake, mg/kg/day	-	173 ± 3	-	103 ± 2	-	-	-
Energy intake, kJ/d	363 ± 9 ^c	406 ± 11 ^b	541 ± 13 ^a	508 ± 14 ^a	<0.0001	0.68	0.003
Feed efficiency, kJ/g	0.12 ± 0.01 ^c	0.10 ± 0.01 ^c	0.41 ± 0.03 ^a	0.31 ± 0.02 ^b	<0.0001	0.003	0.045
Body mass index, g/cm ²	0.63 ± 0.02 ^c	0.63 ± 0.01 ^c	0.85 ± 0.02 ^a	0.75 ± 0.02 ^b	<0.0001	0.008	0.008
Abdominal circumference, cm	18.8 ± 0.3 ^b	18.6 ± 0.2 ^a	23.3 ± 0.5 ^a	21.8 ± 0.2 ^a	0.002	0.007	<0.0001
Whole-body lean mass, g	297 ± 4	293 ± 9	291 ± 6	300 ± 15	0.96	0.79	0.50
Whole-body fat mass, g	69 ± 5 ^c	55 ± 8 ^c	223 ± 11 ^a	187 ± 10 ^b	<0.0001	0.008	0.22
Basal blood glucose, mmol/L	4.14 ± 0.24 ^c	3.65 ± 0.14 ^d	5.25 ± 0.16 ^a	4.66 ± 0.06 ^b	<0.0001	0.002	0.76
Blood glucose at 120 minutes, mmol/L	4.86 ± 0.14 ^b	4.90 ± 0.16 ^b	5.82 ± 0.31 ^a	5.62 ± 0.09 ^a	<0.0001	0.68	0.54
Blood glucose AUC, mmol/L × minutes	715 ± 20 ^b	761 ± 23 ^{ab}	805 ± 18 ^a	819 ± 23 ^a	0.001	0.16	0.45
Plasma total cholesterol, mmol/L	1.70 ± 0.10	1.70 ± 0.06	1.58 ± 0.08	1.56 ± 0.09	0.1242	0.9045	0.9045
Plasma triglyceride, mmol/L	0.56 ± 0.06 ^b	0.64 ± 0.06 ^b	1.30 ± 0.16 ^a	1.23 ± 0.16 ^a	<0.0001	0.9661	0.5249
Plasma NEFA, mmol/L	1.54 ± 0.22 ^b	1.86 ± 0.23 ^b	3.83 ± 0.27 ^a	3.37 ± 0.27 ^a	<0.0001	0.7784	0.1224
Retroperitoneal fat, mg/mm tibial length	136 ± 12 ^c	146 ± 12 ^c	484 ± 54 ^a	279 ± 18 ^b	<0.0001	0.0012	0.0004
Epididymal fat, mg/mm tibial length	87.2 ± 6 ^b	97.0 ± 8.0 ^b	226 ± 23 ^a	189 ± 16 ^a	<0.0001	0.3550	0.1168
Omental fat, mg/mm tibial length	130 ± 8 ^b	127 ± 9 ^b	251 ± 16 ^a	226 ± 13 ^a	<0.0001	0.2401	0.3485
Total abdominal fat, mg/mm tibial length	353 ± 18 ^c	371 ± 27 ^c	961 ± 89 ^a	694 ± 30 ^b	<0.0001	0.0101	0.0038
Visceral adiposity index, %	4.33 ± 0.19 ^c	4.65 ± 0.28 ^c	8.91 ± 0.65 ^a	6.90 ± 0.30 ^b	<0.0001	0.038	0.0052
Systolic blood pressure, mmHg	134 ± 1 ^b	131 ± 1 ^b	142 ± 2 ^a	132 ± 1 ^b	0.002	<0.0001	0.012
LV + septum wet weight, mg/mm tibial length	20.6 ± 1.0	20.5 ± 1.0	24.1 ± 2.0	22.7 ± 1.1	0.051	0.58	0.63
RV wet weight, mg/mm tibial length	4.85 ± 0.45	4.48 ± 0.37	5.49 ± 0.75	5.25 ± 0.38	0.18	0.56	0.89
LV diastolic stiffness constant (κ)	22.5 ± 0.8 ^b	23.0 ± 0.8 ^b	28.4 ± 1.1 ^a	24.2 ± 0.9 ^b	0.0004	0.049	0.014
Liver, mg/mm tibial length	220 ± 7 ^b	209 ± 9 ^b	336 ± 19 ^a	304 ± 10 ^a	<0.0001	0.0676	0.3565
Plasma ALT activity, mmol/L	28.6 ± 2.6 ^b	32.5 ± 3.7 ^b	41.7 ± 3.8 ^a	29.5 ± 1.8 ^b	0.031	0.039	0.013
Plasma AST, mmol/L	71.6 ± 3.0 ^a	71.0 ± 3.7 ^a	81.1 ± 6.2 ^a	60.5 ± 2.3 ^b	0.5812	0.0074	0.0090

Values are mean ± SEM, $n = 8-10$. Mean values within a row with unlike superscript letters are significantly different, $P < 0.05$. ALT, alanine transaminase; AST, aspartate transaminase; AUC, area under the curve; C, corn starch diet-fed rats; CC, corn starch diet-fed rats treated with chlorogenic acid; H, high-carbohydrate, high-fat diet-fed rats; HC, high-carbohydrate, high-fat diet-fed rats treated with chlorogenic acid; LV, left ventricle; NEFA, non-esterified fatty acids; RV, right ventricle.





Chapter 4.

Coffee pulp attenuates metabolic syndrome in diet-induced obese hypertensive rats

Abstract

Coffee pulp is a major by-product of the coffee industry and is a rich source of chlorogenic acid, caffeine and dietary fibre. However, there is very limited data available on applications of coffee pulp in human health. In this study, we demonstrated the pathophysiological and metabolic effects of coffee pulp on a rat model of human metabolic syndrome. 8-9 weeks old male Wistar rats were divided into four groups. Two groups of rats were fed on corn starch diet while the other two groups were fed on high-carbohydrate, high-fat diet for 16 weeks. One group from each diet was given 5% freeze-dried coffee pulp mixed in the food for the last 8 weeks of the protocol. High-carbohydrate, high-fat diet-fed rats developed the symptoms of metabolic syndrome including abdominal obesity, dyslipidaemia, impaired glucose tolerance, and cardiovascular and hepatic complications. Coffee pulp attenuated obesity and hypertension, and improved glucose tolerance, cardiovascular and hepatic dysfunction while reducing blood lipid concentrations without affecting plasma total cholesterol concentrations or vascular reactivity. This study suggested that coffee pulp could be used as a functional food for managing obesity-associated metabolic, cardiovascular and hepatic abnormalities.

Introduction

Obesity has become a major health concern globally [1]. Sedentary lifestyle, lack of physical activity and chronic consumption of excess energy are the prominent causes of the escalating prevalence of obesity [2] as approximately 38% of the world's population is overweight and 20% is obese [3]. Obesity as a component of metabolic syndrome, a constellation of metabolic disorders such as hypertension, central obesity, dyslipidaemia, impaired glucose tolerance and insulin resistance, further increases the risk of developing type 2 diabetes, cardiovascular diseases and non-alcoholic fatty liver disease [4]. Under chronic excess energy intake, excess energy is stored in adipocytes leading to an imbalance in metabolic functions, causing oxidative stress through the disproportionate production of reactive oxygen species, subsequently activating low-grade chronic inflammatory signaling pathways and insulin resistance [5]. Previous studies with functional foods and nutraceuticals have shown antioxidant and anti-inflammatory responses against obesity and metabolic syndrome leading to decreased risks of developing type 2 diabetes, cardiovascular diseases and non-alcoholic fatty liver disease [6].

Coffee is a highly consumed beverage around the world [7], and reports show that coffee is associated with improved glucose tolerance [8], obesity [9], oxidative stress [10], impaired glucose tolerance [11], dyslipidaemia [12], type 2 diabetes [13] and cardiovascular disease [14]. Coffee is an excellent source of phytochemicals including caffeine, chlorogenic acid and fibre that are responsible for delivering the health beneficial effects [15, 16]. High consumption of coffee requires the processing of a large number of coffee berries which generates coffee pulp as the waste from processing. Coffee pulp accounts for ~29% (w/w) of the dry weight of the whole berry. This waste leads to the challenge of disposing of it without environmental contamination [17]. The dry matter of coffee pulp contains carbohydrates (15.7%), proteins (17.4%), crude fibre (14.1%), cellulose (20.7%), hemicellulose (3.6%), lignin (14.3%) [18] and minerals (especially potassium), along with tannins, polyphenols such as chlorogenic acid, and caffeine [17]. HPLC analysis of coffee pulp showed phenolic compounds such as chlorogenic acid (5-caffeoylquinic acid, 42.2%), epicatechin (21.6%), isochlorogenic acid I (5.7%),

isochlorogenic acid II (19.3%), isochlorogenic acid III (4.4%), catechin (2.2%), rutin (2.1%), protocatechuic acid (1.6%) and ferulic acid (1.0%) [19]. Hence, the use of coffee pulp may be considered as a potential approach for management of metabolic complications associated with obesity as there are many reports of health benefits from these individual components of coffee pulp. However, there is insufficient evidence for the health effects of coffee pulp as a dietary supplement.

This study investigated the effects of freeze-dried coffee pulp as an intervention for the last 8 weeks to rats fed a high-carbohydrate, high-fat diet for 16 weeks to mimic human metabolic syndrome [20]. After treatment with coffee pulp, the structure and the function of the cardiovascular system and liver were investigated. Cardiovascular function was studied in isolated Langendorff heart preparations, isolated thoracic aortic rings and by measurement of systolic blood pressure together with histology. Liver structure and function were measured by plasma biochemistry and histology. Metabolic parameters related to obesity and glucose tolerance were also evaluated [20].

Methods

Preparation of coffee pulp powder

Coffee pulp sample was collected from Mountain Top Coffee Farm and Mill, Nimbin, Australia in December 2016. The sample was freeze-dried at School of Agriculture and Food Sciences, University of Queensland, Gatton, Australia and stored at 4°C until analysis.

Characterisation of coffee pulp freeze-dried powder

Extracts of coffee pulp powder were prepared in 3:2 ethanol:water mixture. Briefly, 1 g of powder was dissolved in 50 mL of ethanol:water mixture, sonicated for 15 minutes and an aliquot of the supernatant was taken for analysis by HPLC using an Agilent 1100 series system coupled with a mass spectrometer for further peak confirmation or identification as required. The HPLC system consisted of a diode array detector (G4212B), binary pump (G4220A), an autosampler (G4226A), a vacuum degasser and a column oven with an MSD (G1946D) detector also present. The chromatography was

performed on a Phenomenex luna C18 (2) HPLC column (100 x 4.6 mm) using a gradient method of water and acetonitrile with 0.005% trifluoroacetic acid over 28 minutes. The optimal solvent gradient for separation of target constituents started with 10% acetonitrile which was increased as a gradient to 30% acetonitrile over 10 minutes, then to 95% acetonitrile over 8 minutes, at a flow rate of 0.75 mL/minute and an injection volume of 5 μ L. Calibration standards of trigonelline, caffeine and chlorogenic acid were prepared in 3:2 ethanol:water, at concentrations from 0.01 to 1 mg/mL, 0.005 to 0.5 mg/mL and 0.004 to 1 mg/mL for each of these standards, respectively. Specific detection and calibration curves for each compound were performed at 254 nm, 280 nm and 330 nm, respectively. Quantification was performed using the Chemstation Software based on reference standards, peak area and sample dilution at specific wavelengths for each compound. Mr Peter Mouatt, Senior Analytical Officer from Southern Cross University, played a major role in the analysis of components.

Rats, diets and treatments

All experimental protocols were approved by the University of Southern Queensland Animal Ethics Committee under the guidelines of the National Health and Medical Research Council of Australia. Male Wistar rats (8–9 weeks old, 340 ± 1 g, $n = 48$) were obtained from Animal Resource Centre, Perth, Australia. Rats were divided into 4 groups for the 16 week feeding protocol: corn starch diet-fed rats (C; $n = 12$), corn starch diet + coffee pulp powder (CCP, $n=12$; 5% in food for the final 8 weeks); high-carbohydrate, high-fat-diet-fed rats (H; $n = 12$) and high-carbohydrate, high-fat diet + coffee pulp powder (HCP, $n=12$; 5% in food for the final 8 weeks). Corn starch diet contained 570 g corn starch, 155 g powdered rat food, 25 g Hubbel, Mendel & Wakeman salt mixture and 250 g water per kilogram of diet. High-carbohydrate, high-fat diet contained 175 g fructose, 395 g sweetened condensed milk, 200 g beef tallow, 155 g powdered rat food, 25 g Hubbel, Mendel & Wakeman salt mixture and 50 g water per kilogram of diet [20]. Drinking water with 25% (w/v) fructose was provided to H and HCP groups. C and CCP groups were given normal drinking water. The rats were individually housed under temperature-controlled, 12-hour light/dark conditions and given *ad libitum* access to food and water [20].

Physiological parameters

Body weight, food and water intakes were measured daily. Abdominal circumference and body length (nose to anus) were measured using a standard measuring tape under light anaesthesia with Zoletil (tiletamine 10 mg/kg, zolazepam 10 mg/kg, intraperitoneal). Body mass index was calculated as body weight (in grams)/[body length (in cm)]². Feed efficiency was calculated as [mean body weight gain (in grams)/daily energy intake (in kJ)] [20].

Systolic blood pressure

Systolic blood pressure was measured under light anaesthesia with Zoletil (tiletamine 10 mg/kg, zolazepam 10 mg/kg, intraperitoneally), using an MLT1010 Piezo-Electric Pulse Transducer and inflatable tail-cuff connected to an MLT844 Physiological Pressure Transducer and PowerLab data acquisition unit [20].

Body composition measurement

Dual-energy X-ray absorptiometric measurements were carried out at the end of the protocol with a Norland XR46 DXA instrument (Norland Corp, Fort Atkinson, WI). These scans were evaluated using the manufacturer's suggested software for use in laboratory animals (Small Subject Analysis Software, version 2.5.3/1.3.1; Norland Corp) [21]. The precision error of lean mass for replicate measurements, with repositioning, is 3.2%.

Oral glucose tolerance test

Oral glucose tolerance tests were performed after determining overnight fasting blood glucose concentrations in tail vein blood using Medisense Precision Q.I.D. glucose meters. For overnight fasting, rats were deprived of food for 12 hours. Fructose-supplemented drinking water in all H diet-fed groups was replaced with normal drinking water for the overnight food deprivation period. Basal blood glucose concentrations were measured followed by administration of glucose load 2 g/kg body weight as 40% glucose

solution via oral gavage. Blood glucose concentrations were then measured 30, 60, 90 and 120 minutes after oral glucose administration [20].

Metabolic rate assessment

Indirect calorimetry was applied to determine oxygen consumption and carbon dioxide production using a 4-chamber Oxy-Max system (Columbus Instruments, Columbus, OH) with one rat per chamber. Rats were given *ad libitum* access to food and water during the measurements. Oxygen consumption (V_{O_2}) and carbon dioxide production (V_{CO_2}) were measured individually from each chamber. The respiratory exchange ratio was calculated by Oxy-Max software (v. 4.86). Energy expenditure was determined by assessment of the exchange of oxygen for carbon dioxide that occurs during the metabolic processing of food [22].

Terminal experiments

Rats were euthanased with Lethobarb (pentobarbitone sodium, 100 mg/kg, intraperitoneally). After euthanasia, heparin (200 IU) was injected through the right femoral vein. The abdomen was then opened and blood (~5 mL) was withdrawn from the abdominal aorta and collected into heparinised tubes. Blood was centrifuged at $5000 \times g$ for 10 minutes to obtain plasma. Plasma was stored at -20°C for further characterisation. Hearts were then removed from rats for isolated Langendorff heart preparation [20].

Isolated Langendorff heart preparation

Hearts isolated from euthanased rats were perfused with modified Krebs–Henseleit bicarbonate buffer bubbled with 95% O_2 –5% CO_2 and maintained at 35°C . Isovolumetric ventricular function was measured by inserting a latex balloon catheter into the left ventricle connected to a Capto SP844 MLT844 physiological pressure transducer and Chart software on a Maclab system. Left ventricular end-diastolic pressure values were measured during pacing of the heart at 250 beats per minute using an electrical stimulator. End-diastolic pressures were obtained from 0 to 30 mmHg for the calculation of diastolic stiffness constant (κ , dimensionless) [20].

Vascular reactivity

Thoracic aortic rings (~4 mm in length) were suspended in an organ bath filled with Tyrode physiological salt solution bubbled with 95% O₂–5% CO₂ maintained at 35°C and allowed to stabilise at a resting tension of ~10 mN. Cumulative concentration-response curves (contraction) were obtained for noradrenaline (1×10⁻⁹ – 3×10⁻⁶ M) and cumulative concentration-response curves (relaxation) were obtained for acetylcholine (1×10⁻⁹ – 3×10⁻⁶ M) and sodium nitroprusside (1×10⁻⁹ – 3×10⁻⁶ M) after submaximal (~70%) contraction to noradrenaline [20].

Analysis of intestinal transit time

Rats were orally gavaged with 3 mL of 0.05% phenol red solution 20 minutes before euthanasia. After euthanasia, the entire region from the stomach to small intestine was removed from its mesenteric attachment immediately. The length of small intestine was measured from the pyloric sphincter to the ileocecal junction. The endpoint of phenol red transit in the small intestine was visualised using a few drops of 0.1 M sodium hydroxide [23]. The intestinal transit for each rat was determined by the following formula:

$$\text{Intestinal transit (\%)} = (\text{the total distance traveled by phenol red solution/total length of small intestine}) \times 100$$

Organ weights

After isolated heart perfusion studies, the heart was separated into left ventricle (with septum) and right ventricle and weighed. The liver was isolated and weighed. Retroperitoneal, epididymal and omental fat pads were removed separately and weighed. These organ weights were normalised against tibial length at the time of organ removal and expressed as mg/mm of tibial length [20].

Histology

Hearts and livers were removed from the rats soon after euthanasia and fixed in 10% neutral buffered formalin. These samples were then dehydrated and embedded in

paraffin wax. Thin sections (5 μm) were cut and stained with hematoxylin and eosin to study infiltration of inflammatory cells (heart and liver) and fat deposition (liver) and with picrosirius red (heart) to study collagen deposition [20].

Plasma biochemistry

Plasma activities of aspartate transaminase, alanine transaminase and alkaline phosphatase, and plasma concentrations of total cholesterol, triglycerides and non-esterified fatty acids were measured [20].

Statistical analysis

All data are presented as mean \pm SEM. Results were tested for variance using Bartlett's test and variables that were not normally distributed were transformed (using log 10 functions) prior to statistical analyses. C, CCP, H and HCP groups were tested for effects of diet, treatment and their interactions by 2-way analysis of variance. When the interaction and/or the main effects were significant, means were compared using Newman-Keuls multiple comparison *post hoc* tests. A *P* value of <0.05 was considered significant. All statistical analyses were performed using GraphPad Prism version 5.0 for Windows (San Diego, CA, USA).

Results

Coffee pulp intake

The caffeine and trigonelline doses (mg/kg) were higher in CCP rats than in HCP rats due to higher intake of food by rats fed with corn-starch supplemented with treatments (Table 1).

Table 1. Intakes in coffee pulp powder diets

<i>Component</i>	<i>CCP</i>	<i>HCP</i>
Caffeine intake, mg/kg/day	24.5 ± 2.2	17.5 ± 2.9
Chlorogenic acid intake, mg/kg/day	4.2 ± 0.4	3.0 ± 0.5
Trigonelline intake, mg/kg/day	14.4 ± 1.3	10.2 ± 1.7

Dietary intake, body composition and plasma biochemistry

Body weight was similar between C and CCP rats. H rats had increased body weight compared to C rats while coffee pulp treatment decreased the body weight in HCP rats compared to H rats (Table 2). Food intake was lower in H rats compared to C rats and coffee pulp did not change food intake in both CCP and HCP rats (Table 2). Water intake was similar between C, H and HCP rats while CCP rats had higher water intake. Energy intake was higher in H rats compared to C rats. Coffee pulp did not change energy intake in CCP rats compared to C rats whereas it was higher in HCP rats compared to H rats (Table 2). Feed efficiency and abdominal circumference were similar between C and CCP rats but coffee pulp increased body mass index in CCP rats compared to C rats. Feed efficiency, body mass index and abdominal circumference were higher in H rats than in C rats. Coffee pulp reduced these parameters in HCP rats compared to H rats (Table 2). Heat produced and respiratory exchange ratio were unchanged by coffee pulp in both CCP and HCP rats. Basal blood glucose concentrations were higher in H rats compared to C rats and coffee pulp reduced basal blood glucose concentrations in both CCP and HCP rats compared to C and H rats, respectively. Area under the curve was higher in H rats compared to C rats while it was decreased in both CCP and HCP rats compared to C and

H rats, respectively (Table 2). Retroperitoneal, epididymal, omental and total abdominal fat pads were unchanged in CCP rats compared to C rats but were higher in H rats than in C rats. Coffee pulp reduced total abdominal fat in HCP rats through reduction in omental and epididymal fat pads (Table 2). Whole-body lean mass was similar between C and H rats and coffee pulp only increased it in HCP rats. Whole-body fat mass was higher in H rats than in C rats and it was decreased with coffee pulp in HCP rats compared to H rats (Table 2). Plasma total cholesterol concentrations were unchanged between the groups while coffee pulp reduced plasma concentrations of triglycerides and non-esterified fatty acids in CCP and HCP rats compared to C and H rats, respectively (Table 2).

Cardiovascular structure and function

Histopathological analysis of left ventricle indicated increased infiltration of inflammatory cells in H rats (Figure 1C) compared to C rats (Figure 1A). Coffee pulp reduced infiltration of inflammatory cells into the left ventricle of HCP rats (Figure 1D) than in H rats. Picrosirius red staining of left ventricle showed higher collagen deposition in H rats (Figure 1G) compared to C rats (Figure 1E). Coffee pulp reduced collagen deposition in the left ventricle of HCP rats (Figure 1H) compared to H rats. Noradrenaline-induced contraction and acetylcholine-induced relaxation were lower in H rats than in C rats and coffee pulp treatment did not change these responses (Figure 2A and 2C). There were no differences in sodium nitroprusside-induced vascular relaxation in isolated thoracic aortic rings between the groups (Figure 2B). Systolic blood pressure and left ventricular diastolic stiffness were higher in H rats than in C rats and coffee pulp reduced these parameters in HCP rats compared to H rats (Table 3). LV and RV wet weights were unchanged by coffee pulp treatment (Table 3).

Table 2. Coffee pulp on physiological and metabolic variables

<i>Variables</i>	<i>C</i>	<i>CCP</i>	<i>H</i>	<i>HCP</i>	<i>P values</i>		
					<i>Diet</i>	<i>Intervention</i>	<i>Diet × Intervention</i>
Initial body weight, g	340 ± 1	338 ± 1	339 ± 1	339 ± 1	1.00	0.27	0.27
Final body weight, g	383 ± 2 ^c	386 ± 6 ^c	555 ± 9 ^a	494 ± 9 ^b	<0.0001	0.0002	<0.0001
Body weight gain (8-16 week), %	6.0 ± 2.0 ^c	-1.0 ± 1.0 ^d	23.0 ± 2.0 ^a	10.0 ± 1.0 ^b	<0.0001	<0.0001	0.06
Food intake, g/day	35.2 ± 1.3 ^a	34.2 ± 0.6 ^a	27.2 ± 0.7 ^b	28.0 ± 1.5 ^b	<0.0001	0.93	0.42
Water intake, g/day	31.0 ± 2.0 ^{ab}	36.8 ± 2.1 ^a	26.6 ± 1.6 ^b	32.2 ± 1.7 ^{ab}	0.02	0.004	0.96
Energy intake, kJ/d	405 ± 15 ^c	384 ± 10 ^c	556 ± 19 ^b	616 ± 17 ^a	<0.0001	0.22	0.013
Feed efficiency, kJ/g	0.10 ± 0.01 ^c	0.12 ± 0.02 ^c	0.39 ± 0.02 ^a	0.25 ± 0.02 ^b	<0.0001	0.001	0.0001
Heat, kcal	3.36 ± 0.17 ^b	3.14 ± 0.31 ^b	4.14 ± 0.10 ^a	4.51 ± 0.16 ^a	<0.0001	0.71	0.13
Respiratory exchange ratio	1.01 ± 0.01 ^a	0.95 ± 0.04 ^{ab}	0.89 ± 0.01 ^b	0.89 ± 0.01 ^b	0.0003	0.19	0.29
Body mass index, g/cm ²	0.61 ± 0.01 ^d	0.66 ± 0.01 ^c	0.83 ± 0.02 ^a	0.75 ± 0.01 ^b	<0.0001	0.26	<0.0001
Abdominal circumference, cm	18.4 ± 0.2 ^c	17.8 ± 0.2 ^c	23.2 ± 0.4 ^a	20.2 ± 0.2 ^b	<0.0001	<0.0001	<0.0001
Basal blood glucose concentrations, mmol/L	3.9 ± 0.2 ^b	2.9 ± 0.1 ^c	4.9 ± 0.3 ^a	3.2 ± 0.1 ^c	0.002	<0.0001	0.08
Area under the curve, mmol/L.min	692 ± 28 ^b	421 ± 13 ^d	790 ± 23 ^a	571 ± 13 ^c	<0.0001	<0.0001	0.21
Retroperitoneal fat (mg/mm)	182 ± 21 ^b	196 ± 14 ^b	534 ± 53 ^a	458 ± 22 ^a	<0.0001	0.30	0.14
Epididymal fat (mg/mm)	90 ± 8 ^c	86 ± 10 ^c	222 ± 17 ^a	183 ± 10 ^b	<0.0001	0.07	0.13
Omental fat (mg/mm)	130 ± 8 ^c	117 ± 11 ^c	263 ± 17 ^a	213 ± 12 ^b	<0.0001	0.015	0.14
Total Fat pads (mg/mm)	402 ± 30 ^c	399 ± 28 ^c	1019 ± 81 ^a	855 ± 33 ^b	<0.0001	0.08	0.09
Whole-body lean mass, g	283 ± 5 ^b	295 ± 4 ^{ab}	286 ± 11 ^b	314 ± 7 ^a	0.15	0.01	0.30
Whole-body fat mass, g	83 ± 7 ^c	77 ± 7 ^c	235 ± 17 ^a	168 ± 11 ^b	<0.0001	0.003	0.01
Bone mineral content, g	11.8 ± 0.4 ^b	12.2 ± 0.5 ^b	17.4 ± 0.6 ^a	15.8 ± 0.5 ^a	<0.0001	0.24	0.06
Bone mineral density, g/cm ²	0.172 ± 0.002 ^c	0.188 ± 0.002 ^b	0.185 ± 0.002 ^b	0.197 ± 0.005 ^a	0.0002	<0.0001	0.47
Intestinal transit, %	70.4 ± 7.2 ^b	52.2 ± 7.5 ^b	89.1 ± 4.9 ^a	58.2 ± 8.0 ^b	0.010	0.002	0.40

Plasma total cholesterol concentrations, mmol/L	1.84 ± 0.08	1.71 ± 0.07	1.71 ± 0.10	1.84 ± 0.13	1.0000	1.0000	0.18
Plasma triglyceride concentrations, mmol/L	0.57 ± 0.05 ^b	0.33 ± 0.04 ^c	1.17 ± 0.10 ^a	0.61 ± 0.07 ^b	<0.0001	<0.0001	0.03
Plasma non-esterified fatty acids concentrations, mmol/L	1.37 ± 0.16 ^b	0.76 ± 0.14 ^c	3.57 ± 0.18 ^a	1.53 ± 0.20 ^b	<0.0001	<0.0001	<0.0001

Values are mean ± SEM, n = 8–12. Means in a row with unlike superscripts differ significantly, $P < 0.05$. C, corn starch diet-fed rats; CCP, corn starch diet-fed rats supplemented with coffee pulp; H, high-carbohydrate, high-fat diet-fed rats; HCP, high-carbohydrate, high-fat diet-fed rats treated with coffee pulp.

Table 3. Coffee pulp on cardiovascular and hepatic parameters

Variable	C	CCP	H	HCP	P values		
					Diet	Intervention	Diet × Intervention
Systolic blood pressure, mmHg	125 ± 1 ^b	125 ± 1 ^b	145 ± 2 ^a	129 ± 1 ^b	<0.0001	<0.0001	<0.0001
Diastolic stiffness constant, κ	21.9 ± 0.5 ^b	20.48 ± 0.4 ^b	29.3 ± 1.9 ^a	21.0 ± 0.3 ^b	<0.0001	0.0004	0.002
LV + septum wet weight, mg/mm tibial length	21.1 ± 0.7 ^b	20.4 ± 0.7 ^b	24.7 ± 1.5 ^a	23.5 ± 0.8 ^{ab}	0.002	0.34	0.80
RV wet weight, mg/mm tibial length	4.97 ± 0.38	4.44 ± 0.20	5.64 ± 0.52	5.15 ± 0.20	0.06	0.16	0.96
Liver wet weight, mg/mm tibial length	230 ± 6 ^c	245 ± 10 ^c	336 ± 13 ^a	293 ± 10 ^b	<0.0001	0.17	0.006
Plasma aspartate transaminase activity, mmol/L	79.1 ± 4.8 ^b	94.7 ± 3.9 ^{ab}	82.2 ± 4.4 ^b	103 ± 6.7 ^a	0.25	0.0007	0.60
Plasma alanine transaminase activity, mmol/L	33.0 ± 4.2	37.4 ± 2.8	35.2 ± 3.2	42.1 ± 3.5	0.33	0.12	0.72

Values are mean ± SEM, n = 6–10. Means in a row with unlike superscripts differ significantly, $P < 0.05$. C, corn starch diet-fed rats; CCP, corn starch diet-fed rats treated with coffee pulp; H, high-carbohydrate, high-fat diet-fed rats; HCP, high-carbohydrate, high-fat diet-fed rats treated with coffee pulp.

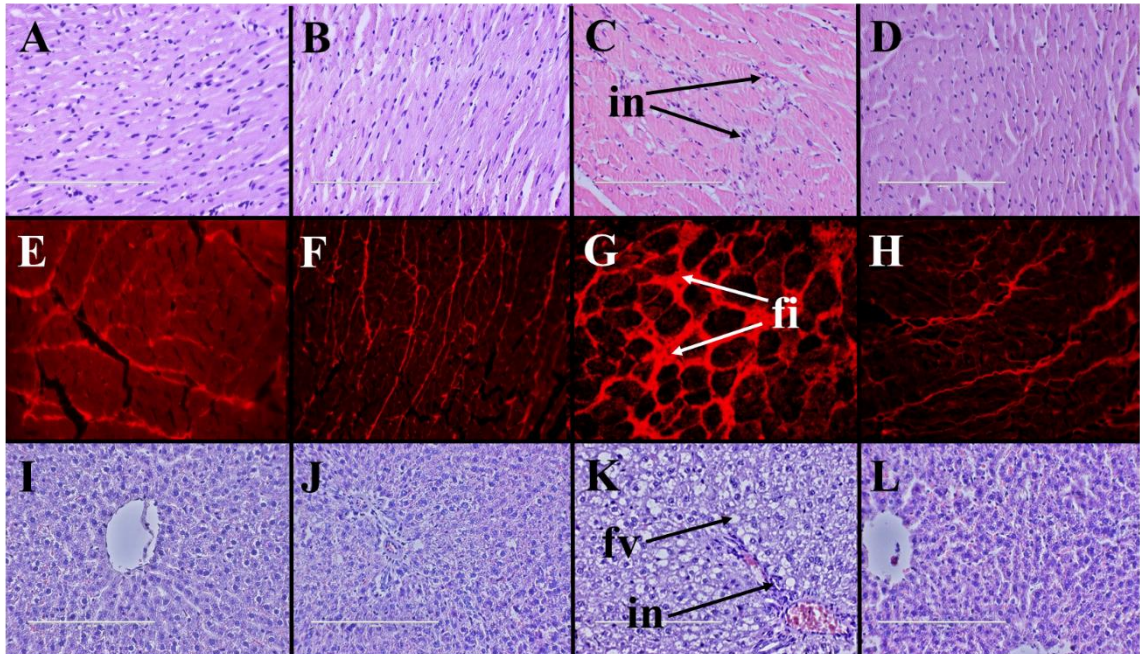


Figure 1. Effects of coffee pulp on the heart and liver structure. Haematoxylin & eosin staining showing infiltration of inflammatory cells (“in”) in hearts from C (A), CCP (B), H (C) and HCP (D) and enlarged fat vacuoles (“fv”) in livers from C (I), CCP (J), H (K) and HCP (L) ($\times 20$). Picosirius red staining showing fibrosis (“fi”) in heart from C (E), CCP (F), H (G) and HCP (H) ($\times 20$). C, corn starch diet-fed rats; CCP, corn starch diet-fed rats treated with coffee pulp; H, high-carbohydrate, high-fat diet-fed rats; HCP, high-carbohydrate, high-fat diet-fed rats treated with coffee pulp.

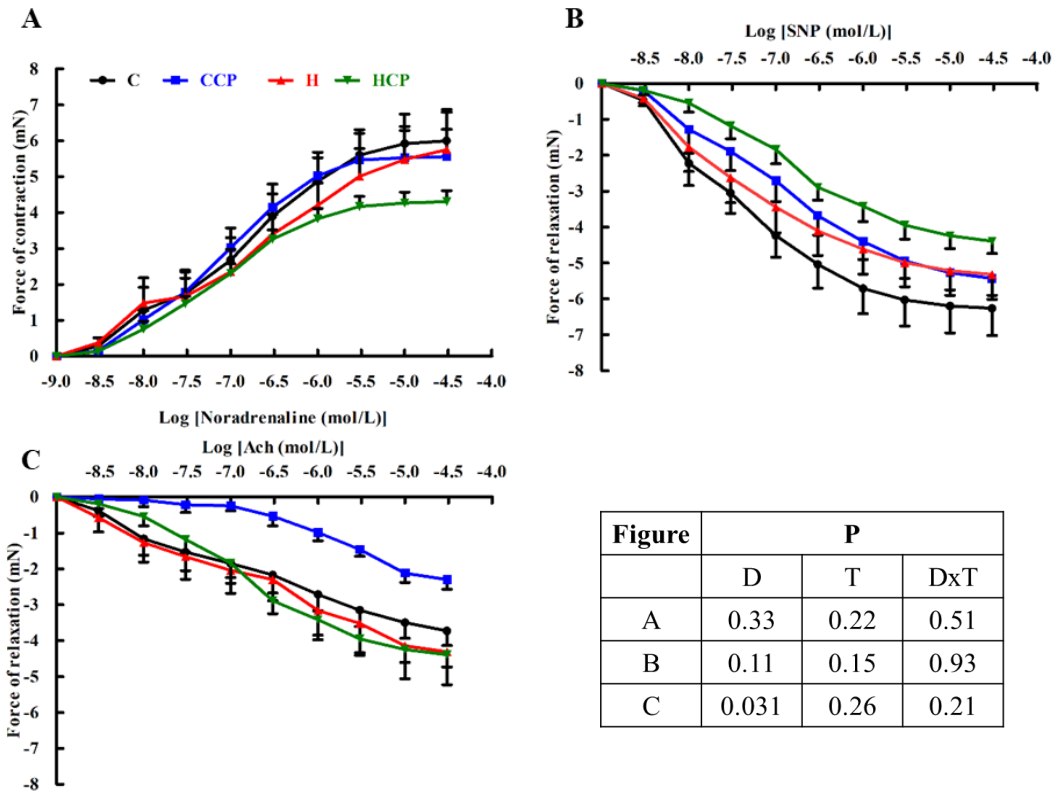


Figure 2. Effects of coffee pulp on noradrenaline-induced contraction (A), sodium nitroprusside-induced relaxation (B) and acetylcholine-induced relaxation (C) in thoracic aorta ring, prepared from C, CCP, H and HCP rats. Values are mean \pm SEM, $n = 8-10$. End-point means without a common letter differ, $P < 0.05$. C, corn starch diet-fed rats; CCP, corn starch diet-fed rats treated with coffee pulp; H, high-carbohydrate, high-fat diet-fed rats; HCP, high-carbohydrate, high-fat diet-fed rats treated with coffee pulp.

Liver structure and function

Liver wet weight was higher in H rats than in C rats and coffee pulp decreased the liver wet weight of HCP rats compared to H rats (Table 3). Liver wet weight was unchanged by coffee pulp treatment between C and CCP rats. Increased inflammatory cell infiltration and fat deposition were observed in livers from H rats (Figure 1K) compared to C rats (Figure 1I). Coffee pulp reduced infiltration of inflammatory cells and fat deposition in livers of HCP rats (Figure 1L) compared to H rats. Plasma alanine transaminase activity was unchanged between the groups. Plasma aspartate transaminase activity was unchanged between H and C rats and increased in HCP rats compared to H rats (Table 3).

Discussion

In this study, coffee pulp reduced body weight and central obesity, improved glucose tolerance along with improved heart and liver structure and function on diet-induced obese rats, fed with the high-carbohydrate high-fat diet. Moreover, coffee pulp increased lean mass and bone mineral density in obese rats. These responses of coffee pulp, a major waste product of the coffee industry, demonstrated the potential application of coffee pulp as a functional food containing bioactive components such as caffeine, chlorogenic acid and trigonelline. These effects of coffee pulp in our diet-induced rat model of metabolic syndrome suggest that this intervention may provide benefits in the complex condition of human metabolic syndrome. The rat model of diet-induced metabolic syndrome used in this study has been validated for the signs of metabolic syndrome including abdominal obesity, impaired glucose tolerance, dyslipidaemia, elevated blood pressure, cardiovascular remodelling and fatty liver [20].

During coffee processing, by-products such as pulp are generated and every 2 tons of coffee creates 1 ton of coffee pulp, a leading cause of serious environmental issue [24, 25]. Therefore, alternative application of coffee pulp is attracting interest. Coffee pulp has been used for production of cascara, a refreshing beverage containing 226 mg/L caffeine [26]. Dry mass of coffee pulp has been used for bioethanol production as it contained polysaccharides including arabinose, galactose, glucose, xylose and mannose at concentrations of 5.8, 5.2, 20.2, 4.2 and 4.7% in dry mass, respectively [27]. Moreover,

coffee pulp consisting of proteins 12%, fibre 21%, caffeine 1.3% and 1% polyphenol [28] and coffee pulp components such as caffeine and chlorogenic acid have antioxidant properties that may counterbalance oxidative stress [29].

Meta-analysis of coffee consumption has reported the inverse relationship with components of metabolic syndrome including abdominal obesity, hypertension, insulin resistance, impaired glucose tolerance, dyslipidaemia and liver complications [30, 31]. Treatment with coffee extract attenuated high-fat diet-induced metabolic disorders and decreased body weight, adipose tissue, and plasma concentrations of glucose, free fatty acid, cholesterol and insulin in mice [32]. In the high-carbohydrate, high-fat fed rat model used in this study, coffee extract [33] and caffeine [34] have previously attenuated metabolic syndrome in these obese hypertensive rats. This study was conducted based on the concept that coffee pulp is a rich source of the same bioactive compounds that are present in coffee prepared as a beverage. This study supported the inverse relationship between consumption of coffee bioactive compounds and cardiovascular disease and type 2 diabetes [35].

Caffeine treatment (~28 mg/kg/day) in obese rats for 8 weeks improved structure and function of heart and liver, with a reduction in obesity except for dyslipidaemia [34]. Caffeine antagonised A₁-adenosine receptors in the hypothalamus to suppress appetite and promote energy use that reduced diet-induced obesity in mice [36]. Chlorogenic acid treatment (5 mg/kg/day) for 3 weeks in obese rats improved glucose tolerance, decreased plasma and liver lipid profile, and recovered mineral pool distribution [37]. Chlorogenic acid (5 mg/kg/day) for 45 days in rats reduced blood lipids including cholesterol, free fatty acids, triglycerides, phospholipids, LDL-cholesterol, VLDL- cholesterol and increased concentrations of HDL- cholesterol and decreased HMG-CoA reductase activity and lipid synthesising enzymes [38]. The study has suggested that diet-induced inflammation in obesity could be managed by the anti-inflammatory property of chlorogenic acid [39] through the increased production of adiponectin and PPAR γ . Likewise, chlorogenic acid reduced the intracellular ROS production and genes associated with ROS-producing enzymes such NOX2, NOX4 and iNOS [40]. Similarly, chlorogenic acid improved lipid metabolism disorders by modifying the expression of PPAR α and LXR α , which contribute

to several intracellular signalling pathways [40]. Consistent with our study, coffee pulp has been reported as a rich source of chlorogenic acid and caffeine, where coffee pulp showed anti-oxidant properties due to their presence [24]. However, in *Chapter 3*, I showed that a much higher dose of chlorogenic acid (approximately 100 mg/kg/day) was required to attenuate inflammation as well as cardiovascular, liver and metabolic abnormalities induced by the high-carbohydrate, high-fat diet. This suggests that chlorogenic acid is not the major bioactive component of coffee pulp.

Coffee pulp used in this study also contained trigonelline, the second most abundant alkaloid in green coffee beans. Trigonelline, as an anti-oxidant, attenuated endoplasmic reticulum-associated stress and oxidative stress-triggered damage in pancreas and adipocytes [41]. Trigonelline (40 mg/kg/day) treatment in the diet for 8 weeks reduced serum activity of aspartate transaminase and aspartate transaminase, and serum concentrations of total cholesterol and LDL-cholesterol and decreased non-alcoholic fatty liver diseases in rats fed with a high fat diet [42]. Trigonelline reduced lipid accumulation by restricting adipocyte differentiation by the PPARgamma cascade. This study suggested that trigonelline inhibits adipocyte differentiation by downregulating fatty acid synthase and GLUT-4 transporter in muscles and adipocytes of mice [43]. These studies suggest that components from coffee pulp as a waste product of the coffee industry, especially trigonelline, have the potential to attenuate metabolic syndrome.

Besides bioactive compounds, the coffee pulp also contained fibre [18] and fibre reduced obesity-associated health disorders through modulating the gut microbiota [44] and reduced body fat, an improved hepatic function that is advantageous to host health [45]. Altered gut microbiota composition disrupts gut barrier junction and promotes the production of lipopolysaccharides by gram-negative bacteria in the gut. This modulation induces gut permeability that enables the translocation of whole bacteria or endotoxic bacterial components into metabolic active tissues and leads to endotoxaemia condition [46]. Endotoxaemia triggers the chronic low-grade inflammation that characterises obesity [47]. Endotoxaemia also induces increased production of nitric oxide metabolites, nitrite and nitrate, increases reactive oxygen species, and increases plasma free fatty acids, tumor necrosis factor and interleukin-6 [48]. Our study with coffee pulp intervention

reduced plasma concentrations of triglycerides and non-esterified fatty acids along with the reduction in body fat mass. These changes may be assisted by healthy gut microbiota that modulate the metabolic changes observed in obesity. Crude fibre accounted for approximately 33.6% of coffee pulp [49] and fibre has prebiotic effects on gut microbiota leading to decreased food intake, weight gain and adiposity, and increased circulating satiety hormones GLP-1 and PYY and colonic fermentation [50]. Colonic fermentation of fibre produces butyrate, acetate and propionate and other dietary secondary metabolites that serve as a substrate for *de novo* lipogenesis in the liver, whereas propionate can be used for gluconeogenesis [51]. Moreover, short-chain fatty acids are also ligands of free fatty acid receptors [52]. Activation of free fatty acid receptors promotes expression and secretion of enteroendocrine hormones such as glucagon-like-peptide 1 or peptide YY which is responsible for satiety [53]. Using the same rat model as in this chapter, treatment with the prebiotic inulin attenuated abdominal obesity, hypertension, increases in inflammatory cell infiltration in the heart and liver, left ventricular diastolic stiffness, lipid droplets in the liver and plasma lipids as well as impaired glucose and insulin tolerance [54]. Dietary fibre has demonstrated prebiotic and anti-bacterial properties, leading to growth of health beneficial gut microbiota and their metabolites [55]. Therefore, I suggest that reduced inflammatory markers and increased colonic production of short-chain fatty acids with the increased intake of dietary fibre from coffee pulp could have contributed to the reduction of the signs of metabolic syndrome in this study.

In conclusion, this chapter provides evidence that coffee pulp can deliver similar results on both pathophysiology and metabolic variables as products from coffee beans. These positive outcomes correlated with reduced plasma lipid concentrations and improvement of glucose tolerance and contributed to reduced obesity, dyslipidaemia and hyperglycaemia. This chapter suggests that dietary intervention with coffee pulp has therapeutic potential to reverse the signs of metabolic syndrome such as obesity, cardiovascular diseases and hepatic abnormalities. Further, the most plausible bioactive components of coffee pulp are trigonelline and fibre. Human clinical trials are necessary to determine whether coffee pulp supplementation in the diet will attenuate or reverse metabolic disorders associated with metabolic syndrome, particularly obesity and hypertension.

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Chapter 5.

**Spent coffee attenuates metabolic syndrome in diet-induced
obese rats**

Abstract

Spent coffee is the by-product of coffee brewing process containing chlorogenic acid, caffeine, trigonelline and melanoidins. However, limited data has been reported for its health benefits. In this study, we investigated responses to spent coffee in a diet-induced rat model of metabolic syndrome. 8-9 weeks old male Wistar rats were divided into four groups. Two groups of rats were fed on corn starch diet with normal drinking water while the other two groups were fed on high-carbohydrate, high-fat diet with 25% fructose in drinking water for 16 weeks. One group from each diet was supplemented with 5% dried spent coffee in the food for the last 8 weeks of protocol. High-carbohydrate, high-fat diet-fed rats developed symptoms of metabolic syndrome including abdominal obesity, impaired glucose tolerance, dyslipidaemia and cardiovascular and hepatic damages. Spent coffee intervention reduced body weight, abdominal fat pads, total body fat mass, systolic blood pressure and concentrations of plasma triglycerides and non-esterified fatty acids along with improvements in glucose tolerance, heart and liver structure and function. Spent coffee increased the diversity of gut microbiota and ratio between Bacteroidetes/Firmicutes. Thus, spent coffee has potential for further testing as a functional food in managing human obesity and metabolic syndrome.

Introduction

Coffee is one of the most consumed beverages and it is the second most important commodity of world trade after crude oil (1). Green coffee beans are an excellent source of compounds such as carbohydrates (59–61%), lipids (10–16%), proteins (10%), chlorogenic acid (7–10%), minerals (4%), caffeine (1–2%), trigonelline (1%) and free amino acids (<1%). The roasting process of green coffee beans causes significant loss of some bioactive compounds including carbohydrates, proteins, chlorogenic acid, free amino acids, lipids, minerals, caffeine and trigonelline (2). During the roasting process, melanoidins are formed as the products of Maillard reaction (3). Melanoidins are nitrogenous non-enzymatic browning compounds that consist of several components contributing to polymerisation reactions (4). The exact chemical structure of coffee melanoidins is still unknown due to variations based on temperature and chemical components present in the complex structure of melanoidins. A potential structure of coffee melanoidins has been provided using sugars, glycoside linkages, amino acids and phenolic compounds (5). Melanoidins may serve as a prebiotic as the melanoidins reach the colon without metabolism in the small intestine. In colon, melanoidins are fermented by gut microbiota, potentially supporting the growth of important gut microbiota (6, 7).

Following the coffee brewing process, spent grains are discarded as a by-product. This discarded spent coffee contributes to approximately 50% of the total coffee weight (8). Thus, a huge quantity of spent coffee is generated as waste during the making of coffee brew. Spent coffee consists of chlorogenic acid, caffeine, trigonelline and melanoidins and hence it may have potential against metabolic syndrome (9). Studies have shown the presence of other bioactive compounds such as diterpenes, lignins, cellulose,

hemicellulose and other polysaccharides in spent coffee (10, 11). The current applications of spent coffee include biodiesel production, as a precursor for the production of activated carbon, as compost and as a sorbent for metal ions removal (12, 13). Very few studies have demonstrated health benefits of spent coffee. Spent coffee exhibited free radical scavenging properties against oxidative stress developed as an outcome of mitochondrial dysfunction (14). Spent coffee restricted the inflammatory activity in J774A.1 mouse macrophage cell line and tumourigenesis activity in P388 leukaemia cell line (15). Melanoidins have previously shown antioxidant and anti-inflammatory activities along with some prebiotic actions on gut microbiota (16). Caffeine, trigonelline and chlorogenic acid found in spent coffee, have already shown potential against various metabolic diseases (17-22). Thus, this project identified the combined effects of these components with melanoidins as spent coffee is generated in large quantities throughout the world and currently is thrown away as a waste material.

The aim of this study was to investigate whether the spent coffee can reverse obesity-associated metabolic changes and changes in structure and function of the heart and liver in high-carbohydrate, high-fat diet-fed rats. High-carbohydrate, high-fat diet-fed rats demonstrated the symptoms of metabolic syndrome with metabolic abnormalities, cardiovascular remodelling and non-alcoholic steatohepatitis (23). Following spent coffee intervention for the last 8 weeks of the 16 week protocol, the structure and function of the heart were characterised through isolated Langendorff heart preparation and histopathology whereas the structure and function of the liver were characterised through histopathology and plasma biochemical studies. Metabolic changes were also characterised through measurement of glucose tolerance, plasma lipids and body

composition. Further, gut microbiota composition was analysed through diversity profiling of faeces isolated from rat colon.

Methods and materials

Spent coffee collection and storage

Spent coffee was collected from the Campus Catering People café at the University of Southern Queensland, Toowoomba, Australia in September 2016. Spent coffee was dried at 60°C until constant weight was achieved. Dried spent coffee was stored at 4°C in dry conditions until analysis of samples and use as dietary intervention.

Characterisation of spent coffee

Spent coffee was prepared in 3:2 ethanol:water mixture. Briefly, 1 g of spent coffee was dissolved in 50 mL of ethanol:water mixture, sonicated for 15 minutes and an aliquot of the supernatant was taken for analysis by HPLC using an Agilent 1100 series system coupled with a mass spectrometer for further peak confirmation or identification as required. The HPLC system consisted of a diode array detector (G4212B), binary pump (G4220A), an autosampler (G4226A), a vacuum degasser and a column oven with an MSD (G1946D) detector also present. The chromatography was performed on a Phenomenex luna C18 (2) HPLC column (100 x 4.6 mm) using a gradient method of water and acetonitrile with 0.005% trifluoroacetic acid over 28 minutes. The optimal solvent gradient for separation of target constituents started with 10% acetonitrile which was increased as a gradient to 30% acetonitrile over 10 minutes, then to 95% acetonitrile over 8 minutes, at a flow rate of 0.75 mL/minute and an injection volume of 5 µL. Calibration

standards of trigonelline, caffeine and chlorogenic acid were prepared in 60:40 ethanol:water, at concentrations from 0.01 to 1 mg/mL, 0.005 to 0.5 mg/mL and 0.004 to 1 mg/mL, respectively. Specific detection and calibration curves for each compound were performed at 254 nm, 280 nm and 330 nm, respectively. Quantification was performed using the Chemstation Software based on reference standards, peak area and sample dilution at specific wavelengths for each compound. Mr Peter Mouatt, Senior Analytical Officer from Southern Cross University, played a major role in the analysis of components.

Rats, diets and treatments

All experimental protocols were approved by the University of Southern Queensland Animal Ethics Committee under the guidelines of the National Health and Medical Research Council of Australia. Male Wistar rats (8–9 weeks old, 340 ± 1 g, $n = 48$) were obtained from Animal Resource Centre, Perth, Australia. Rats were divided into 4 groups for the 16 week feeding protocol: corn starch diet-fed rats (C; $n = 12$), corn starch diet + spent coffee (CSC, $n = 12$; 5% in food for the final 8 weeks); high-carbohydrate, high-fat diet-fed rats (H; $n = 12$), high-carbohydrate high-fat diet + spent coffee (HSC, $n = 12$; 5% in food for the final 8 weeks).

Corn starch diet contained 570 g cornstarch, 155 g powdered rat food, 25 g HMW salt mixture and 250 g water per kilogram of diet. High-carbohydrate, high-fat diet contained 175 g fructose, 395 g sweetened condensed milk, 200 g beef tallow, 155 g powdered rat food, 25 g HMW salt mixture and 50 g water per kilogram of diet. Drinking water with 25% (w/v) fructose was provided to H and HSC groups. C and CSC groups

were given normal drinking water. Rats were individually housed under temperature-controlled, 12-hour-light/dark conditions and given *ad libitum* access to food and water (23).

Physiological parameters

Body weight, food and water intakes were measured daily. Abdominal circumference and body length were measured using a standard measuring tape under light anaesthesia with Zoletil (tiletamine 10 mg/kg, zolazepam 10 mg/kg, intraperitoneal). Body mass index was calculated as body weight (in grams)/[body length (in cm)]². Feed efficiency was calculated as [mean body weight gain (in grams)/daily energy intake (in kJ)] (23).

Systolic blood pressure

Systolic blood pressure was measured under light anaesthesia with Zoletil (tiletamine 10 mg/kg, zolazepam 10 mg/kg, intraperitoneally), using an MLT1010 Piezo-Electric Pulse Transducer and inflatable tail-cuff connected to an MLT844 Physiological Pressure Transducer and PowerLab data acquisition unit (23).

Body composition measurement

Dual-energy X-ray absorptiometric measurements were carried out at the end of the protocol with a Norland XR46 DXA instrument (Norland Corp, Fort Atkinson, WI). These scans were evaluated using the manufacturer's suggested software for use in laboratory animals (Small Subject Analysis Software, version 2.5.3/1.3.1; Norland Corp)

(24). The precision error of lean mass for replicate measurements, with repositioning, is 3.2%.

Oral glucose tolerance test

Oral glucose tolerance tests were performed after determining overnight fasting blood glucose concentrations in tail vein blood using Medisense Precision Q.I.D. glucose meters. For overnight fasting, rats were deprived of food for 12 hours. Fructose-supplemented drinking water for H and HSC groups was replaced with normal drinking water for the overnight food deprivation period. Following basal blood glucose concentration measurement, rats were administered 2 g/kg body weight of glucose as 40% glucose solution via oral gavage. Blood glucose concentrations were measured 30, 60, 90 and 120 minutes after oral glucose administration (23).

Metabolic rate assessment

Indirect calorimetry was applied to determine oxygen consumption and carbon dioxide production using a 4-chamber OxyMax system (Columbus Instruments, Columbus, OH) with one rat per chamber. Rats were given *ad libitum* access to food and water during the measurement. Oxygen consumption and carbon dioxide production were measured individually from each chamber. The respiratory exchange ratio was evaluated by OxyMax software (v. 4.86). Energy expenditure was determined by assessment of the exchange of oxygen for carbon dioxide that occurs during the metabolic processing of food (25).

Terminal experiments

Rats were euthanased with Lethabarb (pentobarbitone sodium, 100 mg/kg, intraperitoneally). After euthanasia, heparin (~200 IU) was injected into the right femoral vein. The abdomen was then opened and blood (~5 mL) was withdrawn from the abdominal aorta and collected into heparinised tubes. Blood was centrifuged at $5000 \times g$ for 10 minutes to obtain plasma. Plasma was stored at -20°C for further characterisation. Hearts were then removed from rats for isolated Langendorff heart preparation (23).

Isolated Langendorff heart preparation

Hearts isolated from euthanased rats were perfused with modified Krebs–Henseleit bicarbonate buffer bubbled with 95% O_2 –5% CO_2 and maintained at 35°C . Isovolumetric ventricular function was measured by inserting a latex balloon catheter into the left ventricle connected to a Capto SP844 MLT844 physiological pressure transducer and Chart software on a Maclab system. All left ventricular end-diastolic pressure values were measured during pacing of the heart at 250 beats per minute using an electrical stimulator. End-diastolic pressures were obtained from 0 to 30 mmHg for the calculation of diastolic stiffness constant (κ , dimensionless) (23).

Vascular reactivity

Thoracic aortic rings (~4 mm in length) were suspended in an organ bath filled with Tyrode physiological salt solution bubbled with 95% O_2 –5% CO_2 maintained at 35°C and allowed to stabilise at a resting tension of ~10 mN. Cumulative concentration–response curves (contraction) were obtained for noradrenaline (1×10^{-9} – 3×10^{-5} M) and cumulative concentration–response curves (relaxation) were obtained for acetylcholine

(1×10^{-9} – 3×10^{-5} M) and sodium nitroprusside (1×10^{-9} – 3×10^{-5} M) after submaximal (~70%) contraction to noradrenaline (23).

Analysis of intestinal transit

Rats from each diet group were orally gavaged with 3 mL of 0.05% phenol red solution 20 minutes before euthanasia. After euthanasia, the entire region from the stomach to small intestine was removed from its mesenteric attachment immediately. The length of small intestine was measured from the pyloric sphincter to the ileo-caecal junction. The endpoint of phenol red transit in small intestine was visualised by adding a few drops of 0.5 M NaOH. The intestinal transit for each rat was determined by following formula (26):

$$\text{Intestinal transit (\%)} = (\text{the total distance traveled by phenol red solution} / \text{total length of small intestine}) \times 100$$

Organ weights

After isolated heart perfusion studies, the heart was separated into left ventricle (with septum) and right ventricle and weighed. The liver was isolated and weighed. Retroperitoneal, epididymal and omental fat contents were removed separately and weighed. These organ weights were normalised against tibial length at the time of organ removal and expressed as mg/mm of tibial length (23).

Histology

Hearts and livers were removed from the rats soon after euthanasia and fixed in 10% neutral buffered formalin. These samples were then dehydrated and embedded in paraffin wax. Thin sections (5 μm) were cut and stained with haematoxylin and eosin to study infiltration of inflammatory cells (heart and liver) and fat deposition (liver) and with picosirius red (heart) to study collagen deposition (23).

Plasma biochemistry

Plasma activities of aspartate transaminase, alanine transaminase and alkaline phosphatase, and plasma concentrations of total cholesterol, triglycerides and non-esterified fatty acids were measured (23).

Gut microbiota diversity profiling

Immediately following euthanasia and organ removal, two to three fecal pellets were collected from the colon of rats and stored at -80°C in nuclease-free tubes. Diversity profiling was performed as described previously (25). In brief, the V3-V4 region of the 16S rRNA gene was chosen for amplification. The primers applied were F341 (5'-CCTAYGGGRBGCASCAG-3') and R806 (5'-GGACTACNNGGGTATCTAAT-3'). PCR amplicons were created using AmpliTaq Gold 360 mastermix (Life Technologies, Scoresby, VIC, Australia) for the primary PCR. A secondary PCR to index the amplicons was operated with TaKaRa Taq DNA Polymerase (Clontech, Mountain View, CA, USA). The subsequent amplicons were measured by fluorometry (Invitrogen Picogreen, Mount Waverley, VIC, Australia) and normalized. The equimolar pool was then calculated by

qPCR (KAPA) followed by sequencing on the Illumina MiSeq (San Diego, CA, USA) with 2 x 300 base pairs paired-end chemistry.

Paired-ends reads were assembled by aligning the forward and reverse reads using PEAR (version 0.9.5) (27). Primers were detected and trimmed. Trimmed sequences were processed using Quantitative Insights into Microbial Ecology (QIIME 1.8) (28) USEARCH (version 7.1.1090) (29, 30) and UPARSE software (31). Using USEARCH, sequences were quality filtered, full length duplicate sequences were eliminated and organized by abundance. Singletons or unique reads in the data set were discarded. Sequences were clustered followed by chimera filtered using “rdp_gold” database as the reference. To obtain the number of reads in each OTU, reads were mapped back to OTUs with a minimum identity of 97%. Using QIIME, taxonomy was assigned using Greengenes database (version 13_8, Aug 2013) (32).

A heat map was constructed using R statistical software according to the developer’s instructions to visualise the relative abundance of each bacterial species and their respective phyla.

Statistical analysis

All data are presented as mean \pm SEM. Results were tested for variance using Bartlett’s test and variables that were not normally distributed were transformed (using log 10 functions) prior to statistical analyses. C, CSC, H and HSC groups were tested for effects of diet, treatment and their interactions by two-way analysis of variance. When the interaction and/or the main effects were significant, means were compared using Newman-Keuls multiple comparison *post hoc* test. A *P* value of less than 0.05 was

considered significant. All statistical analyses were performed using GraphPad Prism version 5.0 for Windows (San Diego, CA, USA).

Results

Spent coffee composition

Spent coffee contained caffeine (7 mg/g w/w), chlorogenic acid (4 mg/g w/w) and trigonelline (1.3 mg/g w/w). The average intake of these compounds over the 8-16 week period was in CSC rats was caffeine (41.9 ± 0.5 mg/kg/day), chlorogenic acid (8.9 ± 0.1 mg/kg/day) and trigonelline (12.1 ± 0.2 mg/kg/day) and in HSC rats was caffeine (17.7 ± 0.4 mg/kg/day), chlorogenic acid (3.8 ± 0.1 mg/kg/day) and trigonelline (5.1 ± 0.1 mg/kg/day).

Dietary intake, body composition and plasma biochemistry

Spent coffee treatment in CSC rats did not change body weight even after increasing food, water and energy intakes compared to C rats. Thus, feed efficiency and body mass index were unchanged between C and CSC rats (Table 1). Abdominal circumference and abdominal fat pads did not change between C and CSC rats while basal blood glucose concentrations and area under the curve were reduced with spent coffee treatment in CSC rats compared to C rats (Table 1). Whole-body lean mass, whole-body fat mass, bone mineral content and bone mineral density did not change between C and CSC rats (Table 1). Spent coffee did not change plasma concentrations of total cholesterol, triglycerides and non-esterified fatty acids in CSC rats compared to C rats (Table 1). Respiratory exchange ratio and heat produced did not change between C and CSC rats (Table 1).

Spent coffee treatment in HSC rats decreased body weight even after increasing food, water and energy intakes compared to H rats. Thus, feed efficiency and body mass index were decreased in HSC rats compared to H rats (Table 1). Abdominal circumference, abdominal fat pads, basal blood glucose concentrations and area under the curve were reduced in HSC rats compared to H rats (Table 1). Whole-body lean mass, bone mineral content, bone mineral density, respiratory exchange ratio and heat produced did not change between H and HSC rats while whole body fat mass was reduced in HSC rats compared to H rats (Table 1). Spent coffee did not change plasma concentrations of total cholesterol but reduced plasma concentrations of triglycerides and non-esterified fatty acids in HSC rats compared to H rats (Table 1).

Cardiovascular and hepatic function

In CSC rats, systolic blood pressure and left ventricular diastolic stiffness constant did not change with spent coffee treatment compared to C rats (Table 2). Spent coffee treatment did not induce inflammation or fibrosis in hearts of CSC rats (Figure 1B and 1F) compared to C rats (Figure 1A and 1E). Spent coffee did not change aortic responses to noradrenaline, sodium nitroprusside or acetylcholine in CSC rats compared to C rats (Figure 2A, 2B and 2C). Left ventricular wet weight decreased while right ventricular wet weights were unchanged in CSC rats compared to C rats (Table 2). Liver wet weight, and plasma activities of aspartate transaminase and alanine transaminase were unchanged between C and CSC rats (Table 2). Spent coffee did not induce inflammation and fat deposition in livers of CSC rats (Figure 3B and 3F) compared to C rats (Figure 3A and 3E).

In HSC rats, systolic blood pressure and left ventricular diastolic stiffness constant decreased with spent coffee treatment compared to H rats (Table 2). Spent coffee treatment reduced inflammation and fibrosis in hearts of HSC rats (Figure 1D and 1H) compared to H rats (Figure 1C and 1G). Spent coffee did not change aortic responses to noradrenaline, sodium nitroprusside or acetylcholine in HSC rats compared to H rats (Figure 2A, 2B and 2C). Left ventricular wet weight decreased while right ventricular wet weight was unchanged in HSC rats compared to H rats (Table 2). Liver wet weight decreased while plasma activities of aspartate transaminase and alanine transaminase were unchanged between H and HSC rats (Table 2). Spent coffee reduced inflammation and fat deposition in livers of HSC rats (Figure 3D and 3H) compared to H rats (Figure 3C and 3G).

Table 1. Response to spent coffee on physiological and metabolic variables

<i>Variables</i>	C	CSC	H	HSC	P		
					Diet	Spent coffee	Diet × spent coffee
Initial body weight, g	340 ± 1	341 ± 1	339 ± 1	339 ± 1	0.13	0.61	0.61
Final body weight, g	383 ± 2 ^c	374 ± 5 ^c	555 ± 9 ^a	481 ± 12 ^b	<0.0001	<0.0001	<0.0001
Food intake, g/day	35.7 ± 0.6 ^c	44.7 ± 0.4 ^a	26.0 ± 0.4 ^d	36.7 ± 0.6 ^b	<0.0001	<0.0001	0.01
Water intake, g/day	26.8 ± 0.4 ^c	34.6 ± 0.4 ^b	27.7 ± 0.5 ^c	36.5 ± 0.5 ^a	<0.0001	<0.0001	0.01
Energy intake, kJ/day	378 ± 8 ^d	443 ± 7 ^c	558 ± 7 ^b	640 ± 10 ^a	<0.0001	<0.0001	0.30
Feed efficiency, kJ/g	0.11 ± 0.01 ^c	0.07 ± 0.01 ^c	0.39 ± 0.02 ^a	0.22 ± 0.03 ^b	<0.0001	<0.0001	0.002
BMI, g/cm ²	0.61 ± 0.01 ^c	0.63 ± 0.01 ^c	0.83 ± 0.02 ^a	0.76 ± 0.02 ^b	<0.0001	0.13	0.01
Abdominal circumference, cm	18.4 ± 0.2 ^c	18.8 ± 0.2 ^c	23.2 ± 0.4 ^a	21.5 ± 0.2 ^b	<0.0001	0.015	0.0002

Basal blood glucose concentrations, mmol/L	3.9 ± 0.2 ^b	2.6 ± 0.1 ^d	4.9 ± 0.3 ^a	3.3 ± 0.1 ^c	0.0002	<0.0001	0.47
Area under the curve, mmol/L·minutes	692 ± 28 ^b	400 ± 10 ^c	790 ± 23 ^a	682 ± 15 ^b	<0.0001	<0.0001	<0.0001
Retroperitoneal fat, mg/mm	182 ± 21 ^c	165 ± 13 ^c	534 ± 53 ^a	404 ± 33 ^b	<0.0001	0.04	0.10
Epididymal fat, mg/mm	90 ± 8 ^c	66 ± 8 ^c	222 ± 17 ^a	142 ± 16 ^b	<0.0001	0.0003	0.04
Omental fat, mg/mm	130 ± 8 ^c	108 ± 3 ^c	263 ± 17 ^a	221 ± 20 ^b	<0.0001	0.031	0.49
Total abdominal fat, mg/mm	402 ± 30 ^c	339 ± 21 ^c	1019 ± 81 ^a	767 ± 63 ^b	<0.0001	0.007	0.091
Whole-body lean mass, g	283 ± 5	308 ± 6	286 ± 11	309 ± 8	0.80	0.052	0.90
Whole-body fat mass, g	83 ± 7 ^c	49 ± 4 ^c	235 ± 17 ^a	180 ± 21 ^b	<0.0001	0.002	0.45
Bone mineral content, g	11.8 ± 0.4 ^b	11.4 ± 0.2 ^b	17.4 ± 0.6 ^a	15.9 ± 0.8 ^a	<0.0001	0.08	0.31
Bone mineral density, g/cm ²	0.172 ± 0.002 ^b	0.175 ± 0.002 ^b	0.185 ± 0.002 ^a	0.188 ± 0.003 ^a	<0.0001	0.19	1.00
Respiratory exchange ratio	1.01 ± 0.01 ^a	1.01 ± 0.01 ^a	0.89 ± 0.01 ^b	0.88 ± 0.02 ^b	<0.0001	0.70	0.70
Heat, kJ	3.36 ± 0.17 ^{bc}	2.94 ± 0.09 ^c	4.14 ± 0.10 ^a	3.61 ± 0.19 ^b	<0.0001	0.003	0.70

Plasma total cholesterol, mmol/L	1.84 ± 0.08	1.85 ± 0.06	1.71 ± 0.10	1.92 ± 0.08	0.72	0.20	0.24
Plasma triglycerides, mmol/L	0.57 ± 0.05 ^c	0.45 ± 0.05 ^c	1.17 ± 0.10 ^a	0.81 ± 0.1 ^b	<0.0001	0.004	0.12
Plasma non-esterified fatty acids, mmol/L	1.37 ± 0.16 ^{bc}	0.94 ± 0.15 ^c	3.57 ± 0.18 ^a	1.76 ± 0.18 ^b	<0.0001	<0.0001	0.0002

Values are mean ± SEM, n = 8–10. Means in a row with superscripts without a common letter differ significantly, P < 0.05. C, corn starch diet-fed rats; CSC, corn starch diet-fed rats treated with spent coffee; H, high-carbohydrate, high-fat diet-fed rats; HSC, high-carbohydrate, high-fat diet-fed rats treated with spent coffee.

Table 2. Effects of spent coffee on cardiovascular and liver function

Variable	C	CCR	H	HCR	<i>p</i>		
					Diet	Spent coffee	Diet × Spent coffee
Systolic blood pressure, mmHg	125 ± 1 ^b	127 ± 2 ^b	145 ± 2 ^a	127 ± 2 ^b	<0.0001	<0.0001	<0.0001
Left ventricular diastolic stiffness constant (κ)	20.7 ± 0.4 ^b	20.8 ± 0.4 ^b	26.7 ± 1.3 ^a	20.8 ± 0.2 ^b	0.0002	0.0002	0.0002
Left ventricle + septum wet weight, mg/mm	21.6 ± 0.8 ^a	19.0 ± 1.3 ^b	24.7 ± 1.5 ^a	19.5 ± 1.2 ^b	0.16	0.004	0.30
Right ventricle wet weight, mg/mm	4.97 ± 0.38	4.91 ± 0.78	5.64 ± 0.52	6.02 ± 0.76	0.15	0.80	0.72
Liver wet weight, mg/mm	230 ± 6 ^c	245 ± 10 ^c	336 ± 13 ^a	293 ± 10 ^b	<0.0001	0.17	0.01
Plasma aspartate transaminase, U/L	79.1 ± 4.8	80.7 ± 3.0	82.2 ± 4.4	81.1 ± 5.7	0.73	0.96	0.79
Plasma alanine transaminase, U/L	27.7 ± 1.8 ^{ab}	24.7 ± 1.9 ^b	35.2 ± 3.2 ^a	29.1 ± 1.8 ^{ab}	0.02	0.07	0.52

Values are mean \pm SEM, n = 8–10. Means in a row with superscripts without a common letter differ significantly, $P < 0.05$. C, corn starch diet-fed rats; CSC, corn starch diet-fed rats treated with spent coffee; H, high-carbohydrate, high-fat diet-fed rats; HSC, high-carbohydrate, high-fat diet-fed rats treated with spent coffee.

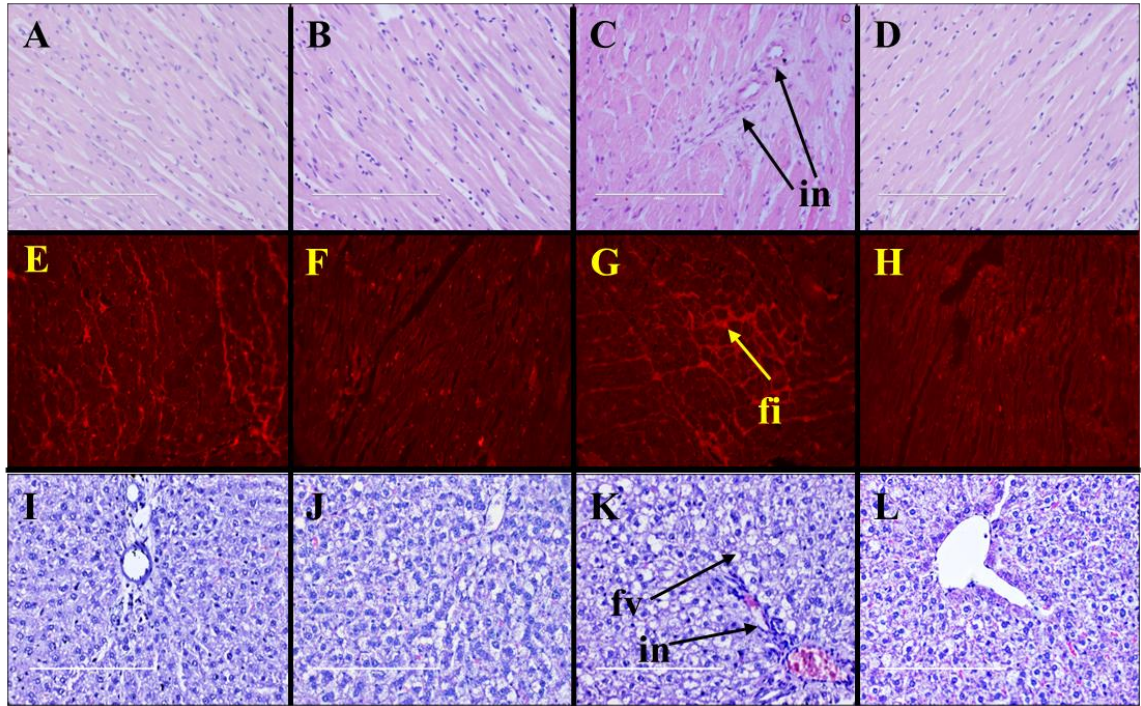


Figure 1. Effects of spent coffee in the heart and liver. H&E staining showing infiltration of inflammatory cells (in) and enlarged fat vacuoles (fv) in heart (A-D) from C (A), CSC (B), H (C), and HSC (D) and liver (I-L) from C (I), CSC (J), H (K) and HSC (L) ($\times 20$). Picrosirius red staining showing fibrosis (fi) in heart (E-H) from C (E), CSC (F), H (G), and HSC (H) ($\times 20$). C, corn starch diet-fed rats; CSC, corn starch diet-fed rats treated with spent coffee; H, high-carbohydrate, high-fat diet-fed rats; HSC, high-carbohydrate, high-fat diet-fed rats treated with spent coffee.

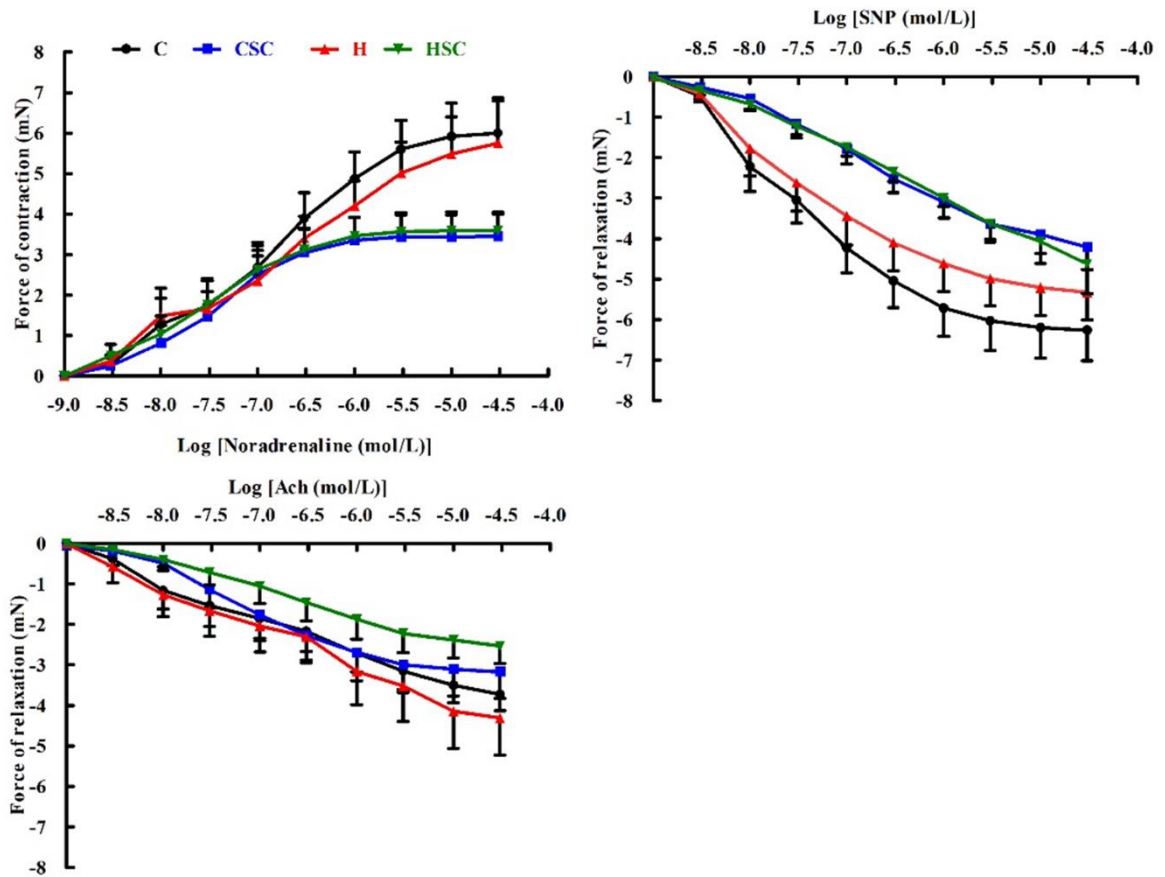


Figure 2. Effects of spent coffee on noradrenaline-induced contraction (A), sodium nitroprusside-induced relaxation (B) and acetylcholine-induced relaxation (C) in thoracic aorta ring, prepared from C, CSC, H and HSC rats. Values are mean \pm SEM, $n = 6-10$. End-point means without a common letter differ, $P < 0.05$. C, corn starch diet-fed rats; CSC, corn starch diet-fed rats treated with spent coffee; H, high-carbohydrate, high-fat diet-fed rats; HSC, high-carbohydrate, high-fat diet-fed rats treated with spent coffee.

Gut microbiome

Figure 3 shows the heat map of bacterial species abundance. In all groups, the major gastrointestinal bacterial phyla, Firmicutes and Bacteroidetes, were predominant (Figure 4). Gut bacterial composition of groups are shown in Figure 4. HSC rats had higher abundance of Bacteroidetes compared to H rats while it did not change between C and CSC rats. Conversely, abundance of Firmicutes remained unchanged among all diet groups (Figure 5A). Spent coffee increased the Bacteroidetes/Firmicutes ratio in CSC and HSC rats compared to C and H rats respectively (Figure 5B). Based on the Shannon diversity index, there was an increase in diversity between the treatment groups CSC and HSC rats compared to control groups C and H rats. (Figure 5C).

The relative abundance of all species was compared between the treatment and control groups to evaluate the effect of spent coffee at species level. A cut-off point of 1% abundance in C rats was applied to enhance confidence. The abundance of two species from phylum Bacteroidetes (*Bacteroides* sp., and an unspecified species from the S24-7 family) was increased and two species from phylum Firmicutes (*Turicibacter* sp. and an unspecified species from Clostridiaceae family) was decreased by spent coffee (Figure 5D). Compared to C rats, there was significant increase in *Bacteroides* sp without changing in the S24-7 family species, *Turicibacter* sp. and an unspecified species from Clostridiaceae family in CSC rats. HSC rats had higher abundance of *Bacteroides* sp., S24-7 family species, *Turicibacter* sp. and an unspecified species from Clostridiaceae family in compared to H rats (Figure 5D).

carbohydrate, high-fat diet-fed rats; HSC, high-carbohydrate, high-fat diet-fed rats treated with spent coffee.

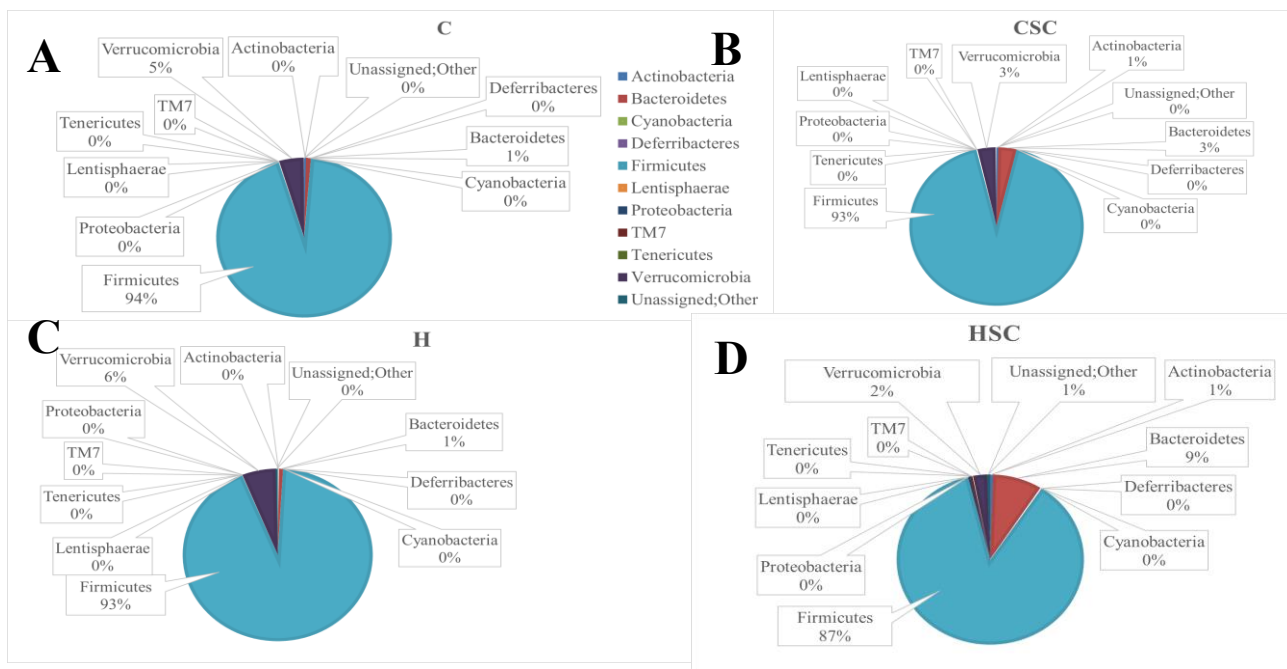


Figure 4. Effect of spent coffee on gut microbiota diversity profiles. Alpha diversity with the diet-group. The relative abundance of each phylum presented as a percentage of the total population for each treatment group. C, corn starch diet-fed rats; CSC, corn starch diet-fed rats treated with spent coffee; H, high-carbohydrate, high-fat diet-fed rats; HSC, high-carbohydrate, high-fat diet-fed rats treated with spent coffee.

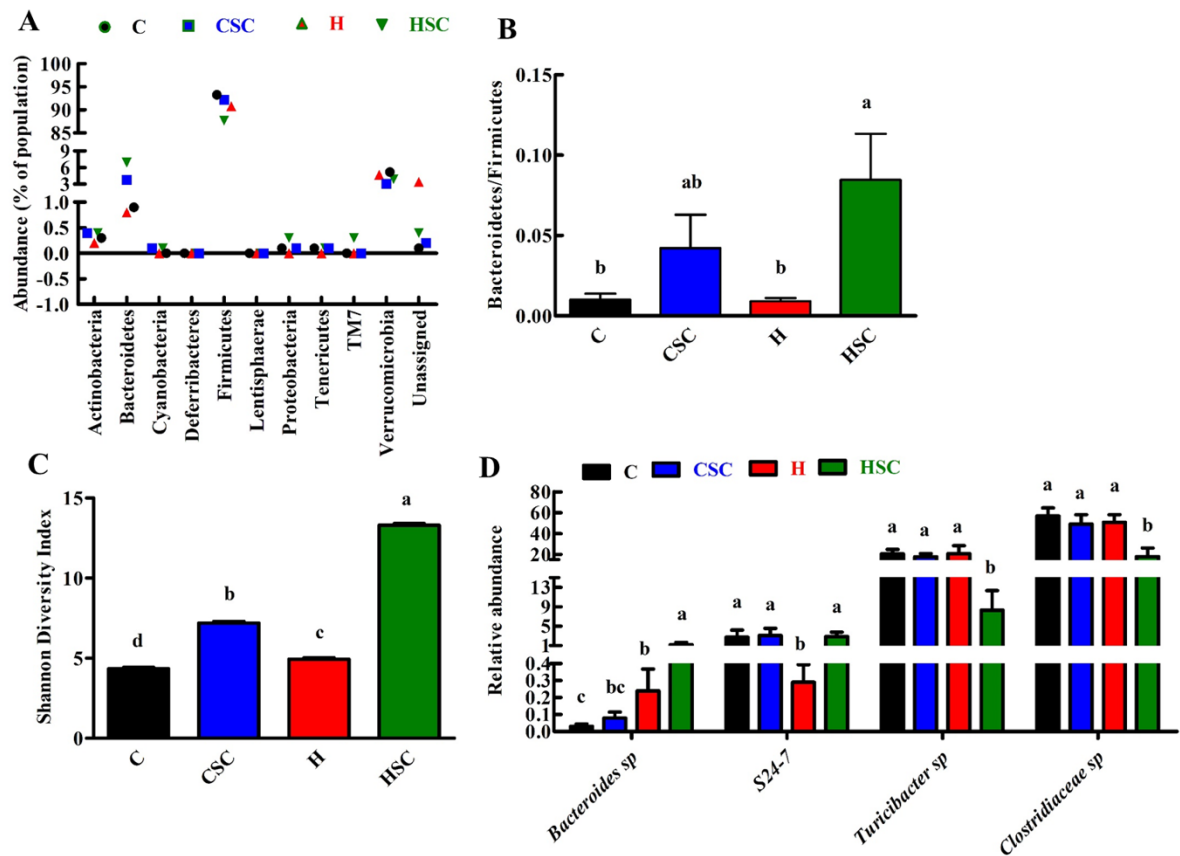


Figure 5. Effect of spent coffee on gut microbiota diversity profiles. A: The relative abundance of each phylum presented as a percentage of the total population for each treatment group. B: Bacteroidetes/Firmicutes ratio. C: Shannon diversity index. The index was determined from the means of abundance for each treatment group. D: Relative abundance of species that were differentially regulated by diet. Only species whose mean abundance for the C diet group was equal to or higher than 1% were plotted in order to enhance confidence. C, corn starch-rich diet-fed rats; CSC, corn starch-rich diet-fed rats treated with spent coffee; H, high-carbohydrate, high-fat diet-fed rats; HSC, high-carbohydrate, high-fat diet-fed rats treated with spent coffee..

Discussion

In this study, spent coffee that is usually discarded after brewing was used as the possible intervention in diet-induced metabolic syndrome in rats. Spent coffee decreased abdominal obesity, systolic blood pressure, plasma triglycerides and plasma non-esterified fatty acids along with improvements in glucose tolerance, cardiovascular and hepatic structure and function in diet-induced obese rat. In our previous studies, we reported that coffee extract (33) and caffeine (34), improved glucose tolerance and cardiovascular structure and function. In these studies, caffeine decreased obesity while coffee extract failed to decrease obesity. Caffeine, through its inhibitory effects on A₁R adenosine receptors (34, 35), decreased obesity while coffee extract was unable to decrease obesity.

Spent coffee contained significant amounts of caffeine, chlorogenic acid and trigonelline, thus suggesting potential of these components in attenuating obesity and metabolic syndrome. However, the dose of caffeine in obese rats (~18 mg/kg/day) was lower than the dose of the caffeine in our previous study (~28 mg/kg/day) (34). This suggests that these 3 components may play a role in attenuating metabolic syndrome but are not the component causing significant response. Spent coffee has also been shown to contain melanoidins, large molecular complexes that still requires further work for elucidating their structure. Roasting process of coffee beans reduced chlorogenic acid (36) while producing melanoidins (37). Melanoidins are formed as the product of Maillard reactions during roasting process using amino acids, polysaccharides and phenolic acids such as chlorogenic acid (5, 38). Similarly, spent coffee includes melanoidins with polysaccharides such as arabinogalactans and galactomannans, which have been shown to have prebiotic activity (39). Melanoidins from roasted coffee have shown several

biological activities such as antioxidant (40), anti-inflammatory (41) and antihypertensive (42). Coffee melanoidins are metabolised by the gut microbiota and produces fermentation products in the colon (43). Previous studies have concluded that coffee melanoidins, through their prebiotic activity, produced acetate and propionate from polysaccharides including galactomannans and arabinogalactans (44, 45). These fermentation products may have a role in improving the quality of gut microbiota. Previous *in vitro* studies have supported the improvements in gut microbiota with the use of melanoidins (46). In this study, we have identified the role of melanoidins from spent coffee in improving the gut microbiota in obese rats thus attenuating metabolic complications. Gut microbiota diversity has increased in spent coffee-treated group of rats. The changes observed in gut microbiota at the species level included increased Bacteroidetes species - *Bacteroides sp*, *S24-7*, and decreased Firmicutes species - *Turicibacter sp*, and *Clostridiaceae sp* in spent coffee-treated group of H rats. Obesity is associated with the modulation of gut microbiota where free fatty acids promotes growth of Firmicutes and consequently reduces Bacteroidetes count (47). The positive modulation in gut microbiota by coffee pulp treatment could be associated with the dietary fibres present in coffee pulp as dietary fibres have been reported for delivering health beneficial (48). Dietary fibre improved the water content of faeces, decreased intestinal transit time and shifted the composition of the gut microbiota positively and enhanced the concentration of short-chain fatty acids in the faeces of mice (48). Thus, fibre from spent coffee could be considered as a prebiotic which could be possible reason for the positive modulation in gut microbiota. Further, this study with spend coffee showed reduction in blood lipids and these changes were not seen in any of our previous studies with coffee (33) or caffeine (34).

Spent coffee, generated in large amounts throughout the world on a daily basis, has served as a functional food against diet-induced obesity in rats in this study. Spent coffee reduced decreased systolic blood pressure as found in our previous study with coffee extract and caffeine (33, 34). This may suggest a role of caffeine and other phenolic acids such as chlorogenic acid in causing these effects. Further, spent coffee also improved glucose tolerance in treatment groups. This study supported our previous studies where coffee (49) and caffeine (34) diet-supplements improved glucose tolerance and insulin sensitivity. Spent coffee still includes an appreciable amount of caffeine. Caffeine induced thermogenesis effects that facilitate energy expenditure and weight management (50). Moreover, the combination of caffeine and chlorogenic acid produced synergistic effects against lipogenesis in mice where caffeine + chlorogenic acid was found to reduce serum and hepatic total cholesterol and triglycerides concentrations (51). This synergetic effect occurs by inhibition of FAS and down-regulation of FAS protein in liver as well as upregulation of AMPK protein expression in adipose tissue (51). Thus, the higher energy expenditure triggered by synergistic effects of caffeine and chlorogenic acid could be the possible reason in our study where elevated food intake delivered the reduction in body weight, feed efficiency and whole-body fat mass.

A high-fat diet is associated with inflammation and modulating energy homeostasis, particularly diets rich in saturated fatty acids (C14:0, C16:0 and C18:0), and caused higher availability of free fatty acids that serve as a ligand to TLR4 (52). Activated TLRs trigger phosphorylation of JNK and binding of activator protein-1 that serve as a transcriptional activator of pro-inflammatory cytokines such as TNF, MCP-1, IFN- γ and IL-6 (53). Thus, we speculated that spent coffee may reduce activation of TLR4-mediated

signalling pathway and accordingly prevented the expression of pro-inflammatory cytokines, confirmed by reduced inflammation in heart and liver. The phosphorylated JNK inhibits IRS-1 by phosphorylating its serine unit. Phosphorylated IRS-1 prevents GLUT4 translocation resulting in insulin resistance state (54). In this study, spent coffee may have improved glucose tolerance by reducing JNK activation and stimulating GLUT4 translocation leading to decreased basal blood glucose concentration and improved glucose response at the end of glucose tolerance test.

Spent coffee improved lipid profile by reducing triglycerides and non-esterified fatty acids in plasma. Increased concentrations of these lipid components are associated with development of obesity, cardiovascular complications and fatty liver (55). Reduction in the lipid components in the blood may suggest the role of spent coffee in improving overall metabolic status thus contributing towards the improvement in metabolic syndrome-associated complications. These improvements in the metabolic status can be linked to the improved gut microbiota observed with the spent coffee treatment. This is the first study to demonstrate the potential of spent coffee in decreasing obesity, and improving obesity-related complication. As we are throwing away useful bioactive components that are present in spent coffee, this is an important finding to warrant future human studies for using food wastes in improving human metabolic health, and so possibly decrease the incidence of obesity in the world.

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Chapter 6.

Attenuation of metabolic syndrome by EPA/DHA ethyl esters in testosterone-deficient high-carbohydrate, high-fat diet-fed rats

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3 *Attenuation of metabolic syndrome by EPA/DHA ethyl esters in testosterone-deficient*
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5 *high-carbohydrate, high-fat diet-fed rats*
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23 esters
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26 **Running title:** EPA/DHA ethyl esters, testosterone and obesity
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1
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3 ***Abstract***
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6 Inducing testosterone deficiency by orchidectomy or gonadotrophin-releasing
7 hormone agonists is the standard treatment of prostate cancer but this treatment may cause
8 metabolic disorders including insulin resistance, hyperinsulinemia, dyslipidemia, central
9 obesity, cardiovascular diseases, and type 2 diabetes. This study measured responses to
10 testosterone deficiency in a diet-induced model of metabolic syndrome induced by a high-
11 carbohydrate, high-fat (H) diet in rats. We then tested whether eicosapentaenoic acid
12 (EPA)/docosahexaenoic acid (DHA) ethyl esters (Omacor) reversed these metabolic changes.
13 Male Wistar rats (8-9 weeks old) were divided into 8 groups with 4 groups fed corn starch
14 (C) and 4 groups fed H diet. For each diet, 1 group received diet only; 1 group was
15 orchidectomized, 1 group was given leuprolide acetate (2 mg/kg every 4th week) and the last
16 group was treated with leuprolide and their diet was supplemented with 3% Omacor. The
17 protocol was for 16 weeks. Leuprolide treatment worsened the metabolic syndrome
18 symptoms and cardiovascular function, and orchidectomy produced greater responses. In H-
19 fed leuprolide-treated rats, Omacor decreased systolic blood pressure and left ventricular
20 diastolic stiffness, reduced infiltration of inflammatory cells and collagen deposition in the
21 heart, reduced lipid accumulation and inflammatory cell infiltration without improving liver
22 damage. These results suggest that Omacor supplementation is associated with decreased
23 inflammation associated with H diets and so could attenuate metabolic complications in
24 prostate cancer patients with induced testosterone deprivation.
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Introduction

Metabolic syndrome is the constellation of insulin resistance, impaired glucose tolerance, dyslipidemia, hypertension, and obesity which increases risk for cardiovascular disease and type 2 diabetes, with an increasing prevalence in the last few decades (O'Neill & O'Driscoll, 2015). Metabolic syndrome, potentially caused by imbalances in energy intake and expenditure, increases morbidity and mortality and is one of the leading preventable causes of death (Zhang, *et al.*, 2017). Further, low testosterone concentrations have been associated with many of these complications including insulin resistance, hyperinsulinemia, dyslipidemia, and cardiovascular disorders (Dockery, *et al.*, 2003; Haffner, *et al.*, 1993; Simon, *et al.*, 1997; Stellato, *et al.*, 2000). Prostate cancer is the second most commonly diagnosed cancer world-wide; in men with prostate cancer, the presence of metabolic syndrome was associated with worse oncologic outcomes, in particular with more aggressive tumor features and biochemical recurrence (Di Francesco & Tenaglia, 2017; Gacci, *et al.*, 2017). The major therapies for prostate cancer rely on induction of testosterone deficiency by orchidectomy or gonadotrophin-releasing hormone (GnRH) agonists such as leuprolide acetate (Lepor & Shore, 2012). However, GnRH agonists increased the risk of development of diabetes and cardiovascular disease, increased fat mass, and decreased lean mass (Goldenberg, *et al.*, 2011; Smith, *et al.*, 2008; Van Londen, *et al.*, 2008).

Diet interventions targeting these metabolic syndrome parameters would be a potential approach to decrease the risk of cardiovascular disease in prostate cancer survivors treated with induced testosterone deficiency, and possibly slow down tumor progression. The health benefits of omega-3 fatty acids such as eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA) in metabolic syndrome and cardiovascular diseases have been reviewed (Bennacer, *et al.*, 2017; Guo, *et al.*, 2017; Tortosa-Caparros, *et al.*, 2017). Omega 3-

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3 fatty acids (EPA and DHA, 1.8 g/day for 26 weeks) decreased expression of genes associated
4 with inflammation and atherogenesis-associated pathways (Bouwens, *et al.*, 2009). Omega-3
5 fatty acids reduced inflammatory markers and cardiovascular disease risk factors (Calder,
6 2013; Mori & Beilin, 2004). Mixed EPA:DHA interventions decreased inflammation and
7 cardiovascular disease risk components with elevated antioxidant enzymes activities
8 (Dasilva, *et al.*, 2016). In a randomized controlled trial, daily doses of 300 mg EPA and 200
9 mg of DHA for 8 weeks reduced high-sensitivity C reactive protein, fasting blood glucose
10 and triglyceride concentrations in hypertensive and/or diabetic obese patients (Ellulu, *et al.*,
11 2016).

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14 In this study, we firstly investigated whether orchidectomy worsened metabolic
15 syndrome in rats fed a high-carbohydrate, high-fat diet. This diet mimics the changes
16 observed in humans with metabolic syndrome (Panchal, *et al.*, 2011). Secondly, we
17 investigated whether rats with testosterone deficiency following 4-weekly leuprolide acetate
18 injections developed similar pathophysiological changes to the high-carbohydrate, high-fat
19 diet as the rats with orchidectomy. Thirdly, we investigated whether a commercially-
20 available mixture of ethyl esters of EPA and DHA (Omacor) reversed the cardiovascular,
21 hepatic, and metabolic parameters in these leuprolide-treated high-carbohydrate, high-fat
22 diet-fed rats. Our hypothesis was that this mixture of EPA and DHA esters has potential as a
23 treatment for metabolic syndrome in prostate cancer patients treated long-term with
24 leuprolide acetate.

25 ***Materials and methods***

26 **Rats and diets**

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28 All experimental procedures were approved by the Animal Ethics Committee of the
29 University of Southern Queensland under the guidelines of the National Health and Medical
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3 Research Council of Australia. 8–9 weeks old male Wistar rats (338 ± 1 g, $n = 96$) were
4 obtained from Animal Resource Centre (Murdoch, WA, Australia). Rats were randomly
5 divided into 8 groups each of 12 rats, with 4 groups fed a corn starch diet and 4 groups fed a
6 high-carbohydrate, high-fat diet, as follows:
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- 11 • **C**: Corn starch diet-fed rats for 16 weeks
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- 13 • **COr**: Orchidectomized rats fed corn starch diet for 16 weeks
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- 15 • **CL**: Rats treated with leuprolide acetate and fed corn starch diet for 16 weeks
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- 17 • **CLOm**: Rats treated with leuprolide acetate for 16 weeks and fed corn starch diet for first
18 8 weeks followed by corn starch diet supplemented with 3% Omacor for the final 8 weeks
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- 20 • **H**: High-carbohydrate, high-fat diet-fed rats for 16 weeks
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- 22 • **HOr**: Orchidectomized rats fed high-carbohydrate, high-fat diet for 16 weeks
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- 24 • **HL**: Rats treated with leuprolide acetate and fed high-carbohydrate, high-fat diet for 16
25 weeks
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- 27 • **HLOm**: Rats treated with leuprolide acetate for 16 weeks and fed high-carbohydrate,
28 high-fat diet for first 8 weeks followed by high-carbohydrate, high-fat diet supplemented
29 with 3% Omacor for the final 8 weeks
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32 **Induction of testosterone deficiency**

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43 24 male rats were orchidectomized at the age of 8 weeks. Bilateral orchidectomy was
44 performed under anesthesia induced by intraperitoneal injection of Zoletil (tiletamine 15
45 mg/kg, zolazepam 15 mg/kg; Virbac, Peakhurst, NSW, Australia) combined with Rompun
46 (xylazine 10 mg/kg; Troy Laboratories, Smithfield, NSW, Australia). An incision was made
47 at the midpoint of the scrotum and the underlying tissue, followed by excision of the testicles
48 and part of the spermatic cord. The incision site was sutured and rats were allowed to recover
49 with administration of carprofen (1 mg/kg for 3 days). Before initiation of the experimental
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3 diet, the orchidectomized rats were given standard laboratory chow diet and monitored daily.
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5 In a further 24 age-matched male rats, testosterone deficiency was induced by subcutaneous
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7 injection of 2 mg/kg leuprolide acetate (Lupron Depot, AbbVie, Sydney, NSW, Australia) at
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9 0, 4, 8, and 12 weeks of the protocol.
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11 12 **Rats, diets, and treatments** 13

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15 All rats were individually housed at the University of Southern Queensland animal
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17 house under temperature-controlled, 12-hour-light/dark conditions and were fed *ad libitum*
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19 with their respective diets. C, CO_r, and CL rats were fed with corn starch diet for 16 weeks
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21 while CLO_m rats were fed with corn starch diet for first 8 weeks and then 3% Omacor-
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23 supplemented corn starch diet for the last 8 weeks. C, CO_r, CL, and CLO_m rats received
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25 normal drinking water for the duration of the protocol. H, HO_r, and HL rats were fed with
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27 high-carbohydrate, high-fat diet for 16 weeks while HLO_m rats were fed with high-
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29 carbohydrate, high-fat diet for first 8 weeks and then 3% Omacor-supplemented high-
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31 carbohydrate, high-fat diet for the last 8 weeks. H, HO_r, HL, and HLO_m rats received
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33 drinking water with 25% fructose (w/v) for the duration of the protocol. Corn starch diet
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35 contained 570 g corn starch, 155 g powdered rat food, 25 g Hubble, Mendel and Wakeman
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37 salt mixture, and 250 mL water per kilogram of diet. High-carbohydrate, high-fat diet
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39 contains 175 g fructose, 395 g sweetened condensed milk, 200 g beef tallow, 155 g powdered
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41 rat food, 25 g Hubble, Mendel and Wakeman salt mixture, and 50 mL water per kilogram of
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43 diet (Panchal, *et al.*, 2011). Each gram of Omacor contained 840 mg of the omega-3 fatty
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45 acid ethyl esters comprising 460 mg of EPA ethyl ester and 380 mg of DHA ethyl esters.
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51 **Physiological parameters** 52

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54 Body weight, and food and water intakes of all rats were measured daily (Panchal, *et*
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56 *al.*, 2011). Abdominal circumference and body length (nose to anus) were measured using a
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3 standard measuring tape under light sedation with Zoletil (tiletamine 10 mg/kg, zolazepam 10
4 mg/kg, intraperitoneal). Feed efficiency was calculated as [mean body weight gain (in
5 grams)/daily energy intake (in kJ)] (Panchal, *et al.*, 2011).
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10 **Systolic blood pressure measurements**

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13 Systolic blood pressure was determined every fourth week under light sedation with
14 Zoletil (tiletamine 10 mg/kg, zolazepam 10 mg/kg, intraperitoneal), using an MLT1010
15 Piezo-Electric Pulse Transducer and inflatable tail-cuff connected to an MLT844
16 Physiological Pressure Transducer and PowerLab data acquisition unit (Panchal, *et al.*, 2011).
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18 After blood pressure measurements, a small volume of blood was collected from the tail vein
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20 for measuring plasma total testosterone concentrations.
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27 **Echocardiography**

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30 Echocardiographic examination (Phillips iE33, 12-MHz transducer) was performed in
31 rats at the end of protocol (Panchal, *et al.*, 2011). Briefly, rats were anesthetized using Zoletil
32 (tiletamine 10 mg/kg and zolazepam 10 mg/kg, intraperitoneal) and Ilium Xylazil (xylazine 6
33 mg/kg, intraperitoneal) and positioned in dorsal recumbency before scanning (Panchal, *et al.*,
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45 **Body composition measurement**

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47 Dual-energy X-ray absorptiometric (DXA) measurements were carried out at the end
48 of the protocol with a Norland XR36 DXA instrument (Norland Corp, Fort Atkinson, WI,
49 USA). These scans were evaluated using the manufacturer's suggested software for use in
50 laboratory animals (Small Subject Analysis Software, version 2.5.3/1.3.1; Norland Corp)
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52 (Ward & Battersby, 2009). The precision error of lean mass for replicate measurements, with
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Oral glucose tolerance test

Oral glucose tolerance tests were performed on rats every fourth week following a 12-hour food deprivation when fructose-supplemented drinking water in all H diet-fed groups was replaced with normal drinking water (Panchal, *et al.*, 2011). After determining basal blood glucose concentrations in tail vein blood using Medisense Precision Q.I.D. glucose meters, rats were given a glucose load of 2 g/kg body weight as 40% glucose solution via oral gavage and blood glucose concentrations were measured again 30, 60, 90, and 120 minutes after oral glucose administration (Panchal, *et al.*, 2011).

Terminal experiments

Rats were euthanized with Lethobarb (pentobarbitone sodium, 100 mg/kg, intraperitoneal) before injection of heparin (200 IU) through the right femoral vein. The abdomen was then opened and blood (~5 mL) was withdrawn from the abdominal aorta and collected into heparinized tubes. Blood was centrifuged at $5000 \times g$ for 15 minutes to obtain plasma. Plasma was stored at -20°C for further biochemical characterization. Hearts were then removed for isolated Langendorff heart studies.

Left ventricular function

Isolated Langendorff heart preparations were used to assess left ventricular function of rats (Panchal, *et al.*, 2011). Hearts isolated from euthanized rats were perfused with modified Krebs–Henseleit bicarbonate buffer bubbled with 95% O_2 – 5% CO_2 and maintained at 35°C . Isovolumetric ventricular function was measured by inserting a latex balloon catheter into the left ventricle connected to a Capto SP844 MLT844 physiological pressure transducer and Chart software on a Maclab system. Left ventricular end-diastolic pressure values were measured during pacing of the heart at 250 beats per minute using an

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3 electrical stimulator. End-diastolic pressures were obtained from 0 to 30 mmHg for the
4 calculation of diastolic stiffness constant (κ , dimensionless) (Panchal, *et al.*, 2011).
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8 **Vascular reactivity**

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11 Thoracic aortic rings (~4 mm in length) were suspended in an organ bath filled with
12 Tyrode physiological salt solution bubbled with 95% O₂ – 5% CO₂ maintained at 35°C and
13 the rings were allowed to stabilise at a resting tension of ~10 mN. Cumulative concentration–
14 response curves (contraction) were obtained for noradrenaline and cumulative concentration–
15 response curves (relaxation) were obtained for acetylcholine and sodium nitroprusside after
16 submaximal (~70%) contraction to noradrenaline (Panchal, *et al.*, 2011).
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25 **Organ weights**

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28 After isolated heart perfusion studies, hearts were separated into left ventricles (with
29 septum) and right ventricles and weighed. Livers were isolated and weighed. Retroperitoneal
30 and omental fat pads were removed separately and weighed; epididymal fat pads were
31 removed from rats except orchidectomized rats. Organ weights were normalized against the
32 tibial length at the time of organ removal and expressed as mg/mm of tibial length (Panchal,
33 *et al.*, 2011).
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42 **Histology**

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45 Heart and liver portions were collected and fixed in 10% neutral buffered formalin.
46 The samples were then dehydrated and embedded in paraffin wax. Thin sections (~5 μ m) of
47 heart and liver were cut and stained with hematoxylin and eosin to study infiltration of
48 inflammatory cells and for determining fat vacuoles in liver. Heart sections were also stained
49 with picosirius red stain to study collagen distribution in the heart (Panchal, *et al.*, 2011).
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Plasma biochemistry

Activities of aspartate transaminase (AST) and alanine transaminase (ALT), and concentrations of total cholesterol, triglycerides, and non-esterified fatty acids (NEFA) in plasma were measured (Panchal, *et al.*, 2011). Plasma total testosterone concentrations were measured using commercial kits (Enzo Life Sciences, Farmingdale, NY) according to protocols provided by the manufacturer.

Statistical analysis

All data are presented as mean \pm SEM. Rats were divided in groups of 12 as this group size produced statistically robust data to answer the hypotheses of this project, as in our previous studies (Panchal, *et al.*, 2011; Poudyal, *et al.*, 2013). Results were tested for variance using Bartlett's test and variables that were not normally distributed were transformed (using log 10 function) prior to statistical analyses. C, CO_r, HL, H, HL, and HO_r rats were tested for effects of diet, testosterone deficiency and their interactions by two-way analysis of variance. When the interaction and/or the main effects were significant, means were compared using Newman-Keuls multiple comparison post hoc test. CL, CLO_m, HL, and HLO_m groups were tested for effects of diet + leuprolide, Omacor treatment and their interactions by two-way analysis of variance. When the interaction and/or the main effects were significant, means were compared using Newman-Keuls multiple comparison *post hoc* test. A *P* value of <0.05 was considered significant. All statistical analyses were performed using GraphPad Prism version 6.1 for Windows (San Diego, CA, USA).

Results

Bilateral orchidectomy

Dietary intakes, body composition, and metabolic parameters

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3 Orchidectomy rapidly reduced plasma total testosterone concentrations in both CO_r
4 and HO_r rats compared to C and H rats, respectively, confirming the effectiveness of
5 orchidectomy (Figure 1A). Orchidectomy did not change body weight, feed efficiency, or
6 intakes of food, water, and energy after 16 weeks of feeding C or H diet compared to their
7 respective non-orchidectomized rats (Table 1). Abdominal circumference, basal blood
8 glucose concentrations, and area under the curve for glucose were higher in orchidectomized
9 rats fed either C or H diet compared to their respective controls (Table 1). Plasma
10 concentrations of total cholesterol, triglycerides, and NEFA were unchanged between the
11 controls and their respective orchidectomized rats (Table 1). Orchidectomy reduced total lean
12 mass in CO_r and HO_r rats compared to C and H rats, respectively. In contrast, the total fat
13 mass was unchanged in C and CO_r rats but increased in HO_r rats compared to H rats (Table
14 1). Bone mineral content increased in HO_r rats compared to H rats but was unchanged in CO_r
15 and C rats while bone mineral density was unchanged. Retroperitoneal fat increased in CO_r
16 and HO_r rats compared to C and H rats, respectively, while omental fat did not change
17 between C and CO_r or H and HO_r rats (Table 1).

35 *Cardiovascular and liver functions*

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37 Orchidectomy in CO_r rats increased systolic blood pressure and left ventricular
38 diastolic stiffness while all other cardiovascular parameters were similar to C rats (Table 2).
39 Orchidectomy induced inflammation and fibrosis in hearts of CO_r rats (Figure 2B and 2H)
40 compared to C rats (Figure 2A and 2G). Orchidectomy reduced aortic responses to
41 noradrenaline, sodium nitroprusside, and acetylcholine in CO_r rats compared to C rats
42 (Figure 3A, 3B, and 3C). HO_r rats had lower heart rate and cardiac output along with higher
43 systolic blood pressure than H rats. H diet induced cardiac changes including inflammation
44 (Figure 2D) and fibrosis (Figure 2J) while orchidectomy worsened these changes in HO_r rats
45 (Figure 2E and 2K). H diet reduced the aortic responses to noradrenaline, sodium
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3 nitroprusside, and acetylcholine compared to C rats while HOr rats had similar aortic
4 response as in H rats (Figure 3A, 3B, and 3C).
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7 C diet did not induce inflammation or fat deposition in liver (Figure 4A and 4G),
8 while orchidectomy induced inflammation and fat deposition in livers of COr rats (Figure 4B
9 and 4H). H diet induced inflammation and fat deposition in livers (Figure 4D and 4J) whereas
10 HOr rats showed similar changes to H rats (Figure 4E and 4K). No changes in liver weight
11 and ALT and AST activities were observed between C and COr or H and HOr rats (Table 2).
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18 **Leuprolide treatment**

19 *Dietary intake, body composition, and metabolic parameters*

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22 Leuprolide reduced plasma total testosterone concentrations after 8 weeks from initial
23 injection in CL rats compared to C rats (Figure 1A). Leuprolide treatment did not change
24 body weight, feed efficiency, or water and food intakes in CL rats compared to C rats;
25 however, energy intakes were increased in CL rats compared to C rats after 16 weeks (Table
26 1). Abdominal circumference, basal blood glucose concentrations, and area under the curve
27 for glucose were higher in CL rats compared to C rats (Table 1). Plasma concentrations of
28 total cholesterol, triglycerides, and NEFA were unchanged between CL and C rats.
29 Leuprolide treatment did not change fat mass of CL rats compared to C rats while the lean
30 mass was reduced in CL rats compared to C rats. Bone mineral content and bone mineral
31 density did not change between CL and C rats. Retroperitoneal fat increased in CL rats
32 compared to C rats, while omental and epididymal fat did not change between C and CL rats
33 (Table 1).
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50 Leuprolide reduced plasma testosterone concentrations after 8 weeks from initial
51 injection in HL rats compared to H rats (Figure 1A). Leuprolide treatment did not change
52 body weight, feed efficiency, or water and food intakes in HL rats compared to H rats.
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3 Energy intakes were increased in HL groups compared to H rats after 16 weeks (Table 1).
4 Abdominal circumference and area under the curve for glucose were higher in HL rats
5 compared to H rats while basal blood glucose concentrations did not change in HL rats
6 compared to H rats (Table 1). Plasma concentrations of triglycerides and NEFA were higher
7 in HL rats compared to H rats while plasma concentrations of total cholesterol were not
8 different between H and HL groups. Leuprolide treatment had no effects on the lean or fat
9 mass in HL rats compared to H rats. Bone mineral content decreased in HL rats compared to
10 H rats whereas bone mineral density did not change with leuprolide treatment.
11 Retroperitoneal fat increased in HL rats compared to H rats, while omental and epididymal
12 fat did not change between H and HL rats (Table 1).
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15 *Cardiovascular and liver function*

16 CL rats had increased LVIDd, LVIDs, diastolic and systolic volumes, and systolic
17 wall stress along with decreased fractional shortening and ejection fraction compared to C
18 rats (Table 2). Leuprolide induced inflammation in CL rats (Figure 2C) and cardiac fibrosis
19 (Figure 2I) and reduced aortic responses to noradrenaline and acetylcholine while not
20 changing the response to sodium nitroprusside in CL rats compared to C rats (Figure 3A, 3B,
21 and 3C). Leuprolide did not induce inflammation or fat deposition in livers of CL rats (Figure
22 4C and 4I). No changes in liver weight or ALT and AST activities were observed between C
23 and CL rats (Table 2).
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26 HL rats had decreased heart rate and fractional shortening with increased LVIDd,
27 LVIDs, and systolic volume than H rats (Table 2). HL rats showed inflammation and fibrosis
28 in the heart (Figure 2F and 2L) and had similar aortic responses as in H and HOr rats (Figure
29 3A, 3B, and 3C). HL rats showed inflammation and fat deposition in liver (Figure 4F and 4L)
30 and no changes in liver weight or ALT and AST activities were observed between H and HL
31 rats (Table 2).
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Omacor treatment

Dietary intake, body composition, and metabolic parameters

Plasma total testosterone concentrations were reduced in CLOm rats from 8 weeks after initial injection, as in CL rats, and there were no differences in plasma total testosterone concentrations between CL and CLOm rats at 16 weeks (Figure 1B). Omacor treatment in CLOm rats starting 8 weeks after the initial leuprolide injection decreased body weight even after increasing food and energy intakes. Omacor reduced the feed efficiency, abdominal circumference, basal blood glucose concentrations, and area under the curve in CLOm rats. Plasma concentrations of total cholesterol and NEFA were decreased while triglycerides did not change when compared to CL rats. Total body lean mass, total body fat mass, retroperitoneal fat, and epididymal fat were decreased in CLOm rats compared to CL rats (Table 3).

Omacor treatment did not change body weight and feed efficiency but it increased water intake (fructose-containing water) in HLOm rats compared to HL rats. Food and energy intakes did not change in HLOm rats compared to HL rats after 16 weeks (Table 1). Abdominal circumference did not change in HLOm rats compared to HL rats. The area under the curve for glucose load was higher in HLOm rats compared to HL rats while basal blood glucose concentrations were decreased in HLOm rats compared to HL rats (Table 1). Plasma concentrations of total cholesterol, triglycerides, and NEFA were reduced in HLOm rats compared to HL rats. In contrast to the plasma lipid profile, Omacor decreased lean mass in HLOm rats compared to HL rats while it increased fat mass in HLOm rats compared to HL rats. Bone mineral content was unchanged while bone mineral density increased in HLOm rats. Retroperitoneal fat, epididymal fat, and visceral adiposity index did not change in HL and HLOm rats while omental fat was increased in HLOm rats in comparison to HL rats (Table 3).

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3 *Cardiovascular and hepatic function*
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5 In CLOm rats, blood pressure and left ventricular diastolic stiffness constant
6 decreased with Omacor treatment compared to CL rats. Omacor treatment reduced
7 inflammation and fibrosis in hearts of CLOm rats (Figure 5B and 5F) compared to CL rats
8 (Figure 5A and 5E). Omacor treatment did not improve aortic responses in CLOm rats
9 compared to CL rats (Figure 6A, 6B, and 6C). Liver weights decreased and ALT activity
10 increased in CLOm rats (Table 4). Omacor treatment attenuated inflammation in livers of
11 CLOm rats (Figure 7B) compared to CL rats (Figure 7A).
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20 In HLOm rats, LVIDs, diastolic volume, systolic volume, systolic wall stress,
21 estimated left ventricular mass, right ventricular weight, and left ventricular weight were
22 increased when compared to HL rats. LVPWs, fractional shortening, ejection fraction,
23 systolic blood pressure, and left ventricular diastolic stiffness decreased in HLOm rats when
24 compared with HL rats (Table 4). Omacor treatment reduced inflammation and fibrosis in
25 hearts of HLOm rats (Figure 5D and 5H) compared to HL rats (Figure 5C and 5G). Omacor
26 treatment did not improve aortic responses in HLOm rats compared to HL rats (Figure 6A,
27 6B, and 6C). Omacor treatment attenuated inflammation and fat deposition in livers of
28 HLOm (Figure 7D and 7H) rats compared to HL rats (Figure 7C and 7G). No changes were
29 observed in liver wet weight or plasma ALT and AST activities between HL and HLOm rats
30 (Table 4).
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45 ***Discussion***
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48 Prostate cancer is the second most commonly diagnosed cancer world-wide covering
49 14% of newly diagnosed cases of cancer and it is associated with about 6% total cancer
50 deaths (Jemal, *et al.*, 2011). The association between metabolic syndrome and prostate cancer
51 focused on the function of insulin, IGF-1, and their receptors as strategic factors in
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3 downstream signaling pathways that stimulate tumor growth (Renehan, *et al.*, 2004). The
4 GnRH agonist leuprolide decreased testosterone concentrations to lead to improved health
5 status of prostate cancer patients by restricting the process of tumorigenesis in the prostate
6 leading to tumor regression, easing of urinary symptoms and bone pain, and prolonged
7 survival (You, *et al.*, 2010). However, patients on androgen deprivation therapy show
8 detrimental changes in body composition such as weight gain, loss of muscle mass, increased
9 fat mass, and decreased muscle strength, with increased fasting glucose, triglycerides, and
10 cholesterol concentrations (Macleod, *et al.*, 2015).
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21 In the present study, we firstly demonstrated that orchidectomy is associated with
22 worse pathophysiological signs of metabolic syndrome in rats fed a high-carbohydrate, high-
23 fat diet. These metabolic changes mimic the progression present in humans with metabolic
24 syndrome (Panchal, *et al.*, 2011). Secondly, we demonstrated that rats with testosterone
25 deficiency induced either by leuprolide or orchidectomy developed similar
26 pathophysiological changes when fed with high-carbohydrate, high-fat diet. Noticeably,
27 orchidectomy delivered more severe pathophysiological complications than leuprolide. When
28 leuprolide is given, there is a period of testosterone flare which, in humans, lasts for around
29 1-2 weeks (Damber, *et al.*, 2012). This is probably the reason for the slower decrease in
30 testosterone concentrations in the leuprolide-treated groups than in the orchidectomized
31 groups. It also means that the exposure to testosterone deficiency in leuprolide-treated rats is
32 shorter than in the orchidectomized groups. Further, testosterone deficiency induced cardiac
33 inflammation and fibrosis with liver steatosis in the corn starch diet-fed rats. Thirdly, we
34 demonstrated that a commercially-available mixture of ethyl esters of EPA and DHA
35 (Omacor) reversed the cardiovascular complications such as decreased blood pressure and
36 left ventricular wall stiffness, reduced hepatic damage such as decreased inflammation and
37 fat deposition in liver and decreased plasma lipid concentrations in these leuprolide-treated
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3 high-carbohydrate high-fat diet-fed rats. Thus, the positive changes against metabolic
4 syndrome with the dietary supplementation have validated our hypothesis that this mixture of
5 EPA and DHA ethyl esters has potential as a nutraceutical approach for reducing components
6 of metabolic syndrome in prostate cancer patients treated long-term with leuprolide acetate.
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12 Orchidectomy and treatment with GnRH agonists such as leuprolide are successful
13 interventions for prostate cancer by initiating testosterone deficiency (Marberger, *et al.*,
14 2010), although orchidectomy is more effective for the initiation of glandular apoptosis and
15 atrophy (Cakiroglu, *et al.*, 2016). Testosterone deficiency produces complications including
16 increased central obesity, increased triglycerides concentrations and elevated fasting plasma
17 glucose concentrations (Gandaglia, *et al.*, 2014; Goldenberg, *et al.*, 2011; Sidaway, 2015;
18 Smith, *et al.*, 2002; Wang, *et al.*, 2011). Leuprolide acetate, although associated with these
19 metabolic complications, has shown tolerability in routine clinical use (Tunn, 2011). Thus, it
20 is important to treat complications associated with GnRH agonist therapy in order to provide
21 benefits against prostate cancer while minimising the development of metabolic and
22 cardiovascular complications.
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37 This study compared the physiological changes produced by testosterone deficiency,
38 either by orchidectomy or leuprolide treatment, and whether these changes worsen during a
39 high-carbohydrate, high-fat diet. In this study, corn starch was used as the control diet as corn
40 starch is slowly digestible, low-glycemic carbohydrate (Panchal, *et al.*, 2011). Unlike fructose
41 in high-carbohydrate, high-fat diet, corn starch does not induce clinical signs of metabolic
42 syndrome (Panchal, *et al.*, 2011; Patel, *et al.*, 2009; Thirunavukkarasu, *et al.*, 2004). High-
43 carbohydrate, high-fat diet induced metabolic syndrome with obesity, dyslipidemia,
44 hypertension and impaired glucose tolerance along with changes in liver and heart structure
45 and function (Panchal, *et al.*, 2011). Orchidectomy reduced lean mass while increasing fat
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3 mass to a greater extent than leuprolide, possibly due to the slower onset of testosterone
4 deficiency with leuprolide. Both orchidectomy and leuprolide increased the deposition of
5 retroperitoneal fat, a major component of the abdominal fat. Presence of abdominal obesity is
6 the major metabolic complication in the initiation of inflammation, potentially increasing the
7 risk of cardiovascular disease (Després, 2012). During the protocol, rats were provided with
8 similar environment including cage size and environmental enrichment with no facility to
9 increase physical activity. We have previously shown that exercise in our high-carbohydrate,
10 high-fat diet-fed rats improved metabolic and cardiovascular function (Cameron, *et al.*,
11 2012). This indicates that the basal physical activity in this study is unlikely to change
12 metabolic or cardiovascular function by itself.
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25 Metabolic syndrome, characterized by the presence of central obesity, hypertension,
26 and dyslipidemia as risk factors for type 2 diabetes and cardiovascular disease, is one of the
27 major complications with leuprolide treatment (Dockery, *et al.*, 2003; Haffner, *et al.*, 1993;
28 Simon, *et al.*, 1997; Stellato, *et al.*, 2000). Obesity as the major component of metabolic
29 syndrome is still without an effective, non-invasive treatment free from adverse effects, with
30 functional foods and nutraceuticals, including omega-3 fatty acids, proposed as effective
31 treatment options (Brown, *et al.*, 2015). EPA and DHA have been reviewed extensively for
32 their effectiveness against metabolic syndrome (Bennacer, *et al.*, 2017; Guo, *et al.*, 2017;
33 Tortosa-Caparros, *et al.*, 2017). Omega-3 and omega-6 fatty acids play vital roles in
34 regulating metabolism as well as state of inflammation (Poudyal, *et al.*, 2011; Simopoulos,
35 2016). A balance is required between pro-inflammatory omega-6 and anti-inflammatory
36 omega-3 fatty acids for the body to maintain homeostasis (Simopoulos, 2016). With the
37 increases in omega-6 fatty acid intake in the modern world, dietary supplementation of
38 omega-3 fatty acids will reduce the imbalance between omega-3 and omega-6 intake and
39 hence contribute in alleviating metabolic and inflammation-related complications
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3 (Simopoulos, 2016). For omega-3 fatty acids, animal studies described anti-obesity effects,
4 but human studies do not conclusively suggest this (Buckley & Howe, 2009). Further,
5 conflicting results are available from the randomized trials of omega-3 fatty acids which may
6 have resulted from differences in study design, dosage used, omega-6/omega-3 fatty acid
7 ratio of the background diet, duration of omega-3 fatty acid supplementation, use of other
8 supplements in addition to omega-3 fatty acids, and demographics of the study population
9 (Simopoulos, 2016).
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19 This study used a commercial mixture of the ethyl esters of EPA and DHA, Omacor,
20 to identify the beneficial effects in the treatment of leuprolide-treated obese rats. Omacor
21 treatment improved bone mineral density, plasma concentrations of total cholesterol,
22 triglycerides, and NEFA, and basal blood glucose concentrations. Omega-3 fatty acids
23 attenuated obesity and glucose intolerance and decreased plasma triglycerides in humans
24 (Lorente-Cebrian, *et al.*, 2013; Rafiee, *et al.*, 2016; Skulas-Ray, *et al.*, 2011). Further, EPA
25 and DHA suppressed the production of pro-inflammatory cytokines including IL-6, TNF- α ,
26 IL-1, and IL-2 (Das, 2006). Omega-3 fatty acids upregulated lipoprotein lipase and adipose
27 triglyceride lipase, enzymes catalysing hydrolysis of triglycerides in skeletal muscle and
28 adipose tissue, respectively (Zimmermann, *et al.*, 2004). Further, omega-3 fatty acids played
29 crucial roles in lowering the rate of fatty acid synthesis and glucose metabolism through
30 downregulation of fatty acid synthase in liver (Kim, *et al.*, 1999). Our study results also
31 suggest that omega-3 fatty acid supplementation improved cardiovascular responses
32 including reduced systolic blood pressure and left ventricular diastolic stiffness along with
33 reduced inflammatory cell infiltration and collagen deposition. Further, Omacor reduced lipid
34 accumulation and inflammatory cell infiltration without changing wet weight of liver and its
35 enzyme activities.
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3 The mechanism of action of omega-3 fatty acids is based on their anti-inflammatory
4 responses (Mori & Beilin, 2004). Omega-3 fatty acids protected against metabolic syndrome
5 through their anti-inflammatory and platelet activating properties that enhance endothelial
6 function and normalize blood pressure by restricting the lipogenesis and activation of lipid
7 oxidation (Carpentier, *et al.*, 2006). In our previous study, EPA and DHA individually
8 improved metabolic syndrome in obese rats at the same dose as in this study (3% in food)
9 (Poudyal, *et al.*, 2013). Colon-specific delivery of EPA or DHA increased the release of
10 glucagon-like peptide 1 and insulin with subsequent reduction in glucose concentrations
11 (Morishita, *et al.*, 2008). Increasing adiponectin is one mechanism by which omega-3 fatty
12 acids can improve cardiometabolic profile in people with cardiovascular risk (Barbosa, *et al.*,
13 2017). Plasma adiponectin was associated with insulin sensitivity (Borges, *et al.*, 2017) and
14 reduced plasma adiponectin was a marker of insulin resistance and increased risk of type 2
15 diabetes (Yamauchi, *et al.*, 2001). Both animal and human studies showed that omega-3 fatty
16 acid supplements improved plasma adiponectin concentrations (Barbosa, *et al.*, 2017; Itoh, *et*
17 *al.*, 2007; Mohammadi, *et al.*, 2012; Neschen, *et al.*, 2006). Thus, increasing both glucagon-
18 like peptide 1 secretion and adiponectin production by omega-3 fatty acids could improve
19 both insulin secretion and sensitivity (Bhaswant, *et al.*, 2015) resulting in improved recovery
20 from insulin resistance and dyslipidemia, thus attenuating metabolic syndrome. Measuring
21 glucagon-like peptide 1 and adiponectin before and after intervention with omega-3 fatty
22 acids could support a plausible mechanism of action of EPA and DHA. However, this study
23 was unable to measure these hormonal changes making it a limitation of our study.
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49 **Conclusions**

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52 We demonstrated that orchidectomy and treatment with a GnRH agonist produced
53 similar worsening of metabolic syndrome symptoms and cardiovascular function. Omacor, a
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3 combination of ethyl esters of EPA and DHA, delivered positive physiological and
4 biochemical responses to reduce symptoms of metabolic syndrome. Further, this study is
5 consistent with previous outcomes obtained for omega 3-fatty acids except for attenuating
6 visceral obesity. Reduced systolic blood pressure and left ventricular stiffness reduction were
7 the major cardiovascular outcomes from this study. Given these observations and the ease of
8 administration, clinical trials of Omacor in men with prostate cancer being managed with
9 androgen deprivation therapy are warranted.
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32 Queensland, Gatton, QLD, Australia) for echocardiography and plasma biochemical
33 analyses.
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41 ***Author contribution***

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43 J.M., L.B., and S.K.P. designed the research protocol. S.K.P. performed orchidectomy
44 on rats. N.B., S.A.K., and S.K.P. conducted animal experiments and collected the data. N.B.,
45 S.A.K., and S.K.P. analysed the data and S.K.P. and L.B. interpreted the data. N.B. and
46 S.K.P. wrote the manuscript with S.K.P. and L.B. having primary responsibility for final
47 content. All authors read and approved the final manuscript.
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55 ***Disclosures***

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3 Authors have nothing to disclose.
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3 **Figure legends**
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6 **Figure 1.** Plasma total testosterone concentrations in C, CO_r, CL, H, HO_r, and HL rats (A)
7 and in CL, CLO_m, HL, and HLO_m rats (B). Data are shown as mean ± SEM, n = 6–
8 10/group. End-point means without a common alphabet significantly differ, *P*<0.05.
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13 **Figure 2.** Effects of orchidectomy and leuprolide on the structure of the heart. Top row
14 represents hematoxylin and eosin staining of left ventricle showing inflammatory cell (“in”)
15 while the bottom row represents picosirius red staining of left ventricle showing collagen
16 deposition (“cd”) from C (A, G), CO_r (B, H), CL (C, I), H (D, J), HO_r (E, K), and HL (F, L)
17 rats.
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24 **Figure 3.** Cumulative concentration-response curves for noradrenaline (A), sodium
25 nitroprusside (B), and acetylcholine (C) in thoracic aortic rings from C, CO_r, CL, H, HO_r,
26 and HL rats. Data are shown as mean ± SEM, n = 10–12/group. End-point means without a
27 common alphabet significantly differ, *P*<0.05.
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33 **Figure 4.** Effects of orchidectomy and leuprolide on the structure of the liver. Top row
34 represents hematoxylin and eosin staining of liver showing inflammatory cell (“in”) while the
35 bottom row represents hematoxylin and eosin staining of liver showing fat deposition (“fd”)
36 from C (A, G), CO_r (B, H), CL (C, I), H (D, J), HO_r (E, K), and HL (F, L) rats.
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43 **Figure 5.** Effects of Omacor on leuprolide-induced changes in the structure of the heart. Top
44 row represents hematoxylin and eosin staining of left ventricle showing inflammatory cell
45 (“in”) while the bottom row represents picosirius red staining of left ventricle showing
46 collagen deposition (“cd”) from CL (A, E), CLO_m (B, F), HL (C, G), and HLO_m (D, H) rats.
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52 **Figure 6.** Cumulative concentration-response curves for noradrenaline (A), sodium
53 nitroprusside (B), and acetylcholine (C) in thoracic aortic rings from CL, CLO_m, HL, and
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3 HLOm rats. Data are shown as mean \pm SEM, n = 10–12/group. End-point means without a
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5 common alphabet significantly differ, $P < 0.05$.
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7 **Figure 7.** Effects of Omacor on leuprolide-induced changes in the structure of the liver. Top
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9 row represents hematoxylin and eosin staining of liver showing inflammatory cell (“in”)
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11 while the bottom row represents hematoxylin and eosin staining of liver showing fat
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13 deposition (“fd”) from CL (A, E), CLOm (B, F), HL (C, G), and HLOm (D, H) rats.
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Table 1. Effects of orchidectomy and leuprolide-induced changes in physiological and metabolic parameters

Variables	C	CO _r	CL	H	HO _r	HL	P value		
							Diet	Testosterone deficiency	Diet × testosterone deficiency
Initial body weight (g)	337 ± 1	338 ± 1	337 ± 1	336 ± 1	339 ± 1	338 ± 1	0.68	0.14	0.52
Final body weight (g)	419 ± 5 ^b	429 ± 5 ^b	417 ± 8 ^b	520 ± 6 ^a	522 ± 14 ^a	539 ± 18 ^a	<0.0001	0.71	0.37
Water intake (mL/day)	31.0 ± 2.1 ^a	32.0 ± 2.4 ^a	26.4 ± 2.6 ^{ab}	19.8 ± 1.6 ^{bc}	19.8 ± 1.6 ^{bc}	17.5 ± 1.7 ^c	<0.0001	0.12	0.71
Food intake (g/day)	31.6 ± 2.0 ^a	31.9 ± 2.3 ^a	36.1 ± 2.2 ^a	20.8 ± 1.3 ^b	21.1 ± 1.8 ^b	24.3 ± 1.9 ^b	<0.0001	0.08	0.96
Energy intake (kJ/day)	346 ± 3 ^d	356 ± 3 ^d	401 ± 6 ^c	444 ± 3 ^b	447 ± 5 ^b	507 ± 12 ^a	<0.0001	<0.0001	0.54
Feed efficiency (g/kJ)	0.23 ± 0.01 ^b	0.25 ± 0.02 ^b	0.20 ± 0.02 ^b	0.41 ± 0.01 ^a	0.40 ± 0.03 ^a	0.40 ± 0.03 ^a	<0.0001	0.48	0.51
Abdominal circumference (cm)	18.4 ± 0.1 ^d	20.1 ± 0.4 ^c	20.1 ± 0.2 ^c	22.0 ± 0.2 ^b	23.2 ± 0.5 ^a	23.0 ± 0.3 ^a	<0.0001	<0.0001	0.52
Basal blood glucose concentrations (mmol/L)	4.1 ± 0.1 ^d	4.7 ± 0.1 ^c	4.8 ± 0.1 ^c	5.1 ± 0.1 ^{bc}	5.6 ± 0.1 ^a	5.4 ± 0.2 ^{ab}	<0.0001	<0.0001	0.24
Area under the curve (mmol/L×minutes)	632 ± 21 ^c	776 ± 19 ^b	781 ± 10 ^b	774 ± 16 ^b	854 ± 14 ^a	844 ± 24 ^a	<0.0001	<0.0001	0.07
Total cholesterol (mmol/L)	1.37 ± 0.05 ^b	1.71 ± 0.15 ^{ab}	1.65 ± 0.04 ^{ab}	1.88 ± 0.06 ^a	1.84 ± 0.17 ^a	1.90 ± 0.10 ^a	0.001	0.28	0.20
Triglyceride (mmol/L)	0.60 ± 0.03 ^c	0.56 ± 0.05 ^c	0.60 ± 0.06 ^c	1.42 ± 0.15 ^b	1.06 ± 0.22 ^{bc}	2.02 ± 0.32 ^a	<0.0001	0.019	0.032
NEFA (mmol/L)	1.74 ± 0.16 ^c	2.47 ± 0.17 ^c	2.38 ± 0.19 ^c	3.69 ± 0.37 ^b	3.41 ± 0.43 ^b	4.60 ± 0.35 ^a	<0.0001	0.034	0.09
Whole-body lean mass (g)	312 ± 11 ^a	271 ± 11 ^b	275 ± 5 ^b	314 ± 8 ^a	229 ± 14 ^c	308 ± 9 ^a	0.78	<0.0001	0.002
Whole-body fat mass (g)	114 ± 11 ^c	121 ± 10 ^c	112 ± 8 ^c	191 ± 8 ^b	259 ± 23 ^a	161 ± 15 ^{bc}	<0.0001	0.0007	0.006
Bone mineral content (g)	13.3 ± 0.5 ^b	12.4 ± 0.4 ^{bc}	11.9 ± 0.2 ^{bc}	13.3 ± 0.4 ^b	16.2 ± 0.6 ^a	11.3 ± 0.5 ^c	0.005	<0.0001	<0.0001
Bone mineral density (g/cm ²)	0.183 ± 0.003	0.173 ± 0.004	0.174 ± 0.003	0.183 ± 0.004	0.173 ± 0.003	0.177 ± 0.003	0.72	0.12	0.88
Retroperitoneal fat (mg/mm tibial length)	122 ± 8 ^d	231 ± 18 ^c	220 ± 18 ^c	366 ± 17 ^b	537 ± 26 ^a	479 ± 54 ^a	<0.0001	<0.0001	0.51
Epididymal fat (mg/mm tibial length)	101 ± 4 ^b	-	125 ± 11 ^b	224 ± 17 ^a	-	263 ± 26 ^a	<0.0001	0.07	0.65
Omental fat (mg/mm tibial length)	65 ± 6 ^b	90 ± 6 ^b	83 ± 6 ^b	132 ± 9 ^a	137 ± 13 ^a	147 ± 13 ^a	<0.0001	0.16	0.52

Value are means ± SEM, *n* = 10–12. Mean values in a row with unlike superscript letters are significantly different (*P*<0.05). C, corn starch diet-fed rats; CO_r, orchidectomized rats fed corn starch diet; CL, leuprolide-treated rats fed corn starch diet; H, high-carbohydrate high-fat diet fed rats; HO_r, orchidectomized rats fed high-carbohydrate high-fat diet; HL, leuprolide-treated rats fed high-carbohydrate high-fat diet; NEFA, non-esterified fatty acids.

Table 2. Effects of orchidectomy and leuprolide-induced changes in cardiovascular and hepatic function

Variables	C	CO _r	CL	H	HO _r	HL	P value		
							Diet	Testosterone deficiency	Diet × testosterone deficiency
Heart rate (bpm)	253 ± 23 ^{bc}	227 ± 17 ^c	238 ± 11 ^{bc}	350 ± 10 ^a	282 ± 15 ^{bc}	290 ± 15 ^b	<0.0001	0.01	0.29
LVIDd (mm)	6.64 ± 0.28 ^c	6.63 ± 0.30 ^c	7.70 ± 0.15 ^{ab}	7.09 ± 0.22 ^{bc}	7.23 ± 0.20 ^{bc}	8.24 ± 0.22 ^a	0.007	<0.0001	0.95
LVIDs (mm)	3.11 ± 0.19 ^c	3.68 ± 0.18 ^{bc}	4.52 ± 0.19 ^a	3.22 ± 0.23 ^c	3.48 ± 0.25 ^{bc}	4.14 ± 0.18 ^{ab}	0.35	<0.0001	0.49
IVSd (mm)	1.91 ± 0.12	1.96 ± 0.10	1.77 ± 0.06	2.05 ± 0.08	2.05 ± 0.17	1.82 ± 0.06	0.28	0.10	0.91
IVSs (mm)	3.09 ± 0.26 ^{ab}	3.13 ± 0.20 ^{ab}	2.80 ± 0.08 ^b	3.65 ± 0.09 ^a	3.43 ± 0.20 ^{ab}	3.12 ± 0.11 ^{ab}	0.006	0.047	0.70
LVPWd (mm)	1.70 ± 0.09	1.91 ± 0.12	1.79 ± 0.03	2.04 ± 0.06	2.08 ± 0.15	1.75 ± 0.05	0.05	0.06	0.14
LVPWs (mm)	2.73 ± 0.16 ^b	2.65 ± 0.14 ^b	2.58 ± 0.15 ^b	3.27 ± 0.14 ^a	3.04 ± 0.13 ^{ab}	2.95 ± 0.10 ^{ab}	0.0003	0.23	0.80
Diastolic volume (μL)	317 ± 41 ^c	318 ± 38 ^c	481 ± 26 ^{ab}	427 ± 40 ^{abc}	356 ± 30 ^{bc}	531 ± 47 ^a	0.036	0.0001	0.59
Systolic volume (μL)	34.2 ± 6.1 ^c	55.0 ± 8.2 ^{bc}	100.2 ± 12.8 ^a	38.4 ± 7.0 ^c	49.2 ± 10.5 ^{bc}	77.1 ± 8.6 ^{ab}	0.28	<0.0001	0.33
Stroke volume (μL)	283 ± 38 ^b	263 ± 33 ^b	381 ± 22 ^{ab}	388 ± 35 ^{ab}	306 ± 24 ^b	454 ± 43 ^a	0.009	0.0007	0.65
SBP:LVIDs	42.4 ± 2.9 ^{ab}	40.9 ± 1.8 ^{abc}	32.1 ± 1.5 ^c	48.3 ± 3.8 ^a	48.0 ± 3.0 ^a	36.9 ± 2.4 ^{bc}	0.009	0.0001	0.91
SBP:systolic volume	4826 ± 956 ^{ab}	3016 ± 418 ^{ab}	1599 ± 198 ^b	5562 ± 1462 ^a	4436 ± 840 ^{ab}	2291 ± 479 ^{ab}	0.17	0.001	0.89
ESS:LVIDs	2.40 ± 0.10 ^b	2.44 ± 0.08 ^{ab}	2.84 ± 0.16 ^a	2.30 ± 0.09 ^b	2.69 ± 0.09 ^{ab}	2.56 ± 0.11 ^{ab}	0.63	0.007	0.05
Cardiac output (mL/min)	70.1 ± 9.2 ^b	56.3 ± 7.9 ^b	90.6 ± 7.3 ^b	135.2 ± 10.9 ^a	84.8 ± 5.6 ^b	130.9 ± 12.9 ^a	<0.0001	0.0001	0.14
Relative wall thickness	0.56 ± 0.05	0.58 ± 0.05	0.47 ± 0.02	0.56 ± 0.02	0.60 ± 0.05	0.45 ± 0.02	1.00	0.003	0.87
Systolic wall stress	75.7 ± 7.0 ^b	104.4 ± 8.9 ^{ab}	129.9 ± 12.2 ^a	74.8 ± 7.1 ^b	93.3 ± 6.6 ^b	106.4 ± 7.3 ^{ab}	0.09	<0.0001	0.41
Estimated LV mass (g)	0.81 ± 0.03 ^b	0.89 ± 0.05 ^{ab}	1.01 ± 0.04 ^{ab}	1.16 ± 0.07 ^a	1.08 ± 0.12 ^{ab}	1.07 ± 0.06 ^{ab}	0.006	0.65	0.11
Fractional shortening (%)	53.1 ± 2.3 ^{ab}	47.1 ± 2.1 ^{bc}	41.2 ± 2.1 ^c	56.6 ± 2.5 ^a	49.6 ± 2.7 ^{abc}	47.7 ± 2.3 ^{bc}	0.033	0.0002	0.68
Ejection fraction (%)	89.1 ± 1.6 ^a	84.0 ± 2.0 ^{ab}	79.3 ± 2.0 ^b	91.2 ± 1.5 ^a	86.6 ± 2.1 ^a	85.2 ± 1.7 ^{ab}	0.021	0.0002	0.53
Systolic blood pressure (mmHg)	126 ± 2 ^d	146 ± 1 ^{bc}	144 ± 2 ^c	152 ± 2 ^b	161 ± 2 ^a	150 ± 2 ^{bc}	<0.0001	<0.0001	<0.0001
Right ventricular wet weight (mg/mm tibial length)	2.37 ± 0.19	3.39 ± 0.64	2.76 ± 0.16	4.21 ± 1.14	2.88 ± 0.12	3.19 ± 0.15	0.20	0.85	0.11
Left ventricular + septum wet weight (mg/mm tibial length)	17.6 ± 0.5 ^b	18.1 ± 0.7 ^{ab}	18.1 ± 0.5 ^{ab}	19.5 ± 0.5 ^{ab}	19.9 ± 0.2 ^a	19.1 ± 0.5 ^{ab}	0.003	0.62	0.62
Left ventricular diastolic stiffness constant, κ	23.8 ± 0.7 ^b	29.4 ± 0.6 ^a	27.3 ± 0.5 ^a	29.1 ± 0.7 ^a	29.1 ± 1.1 ^a	27.9 ± 0.7 ^a	0.003	0.002	0.0006
Liver wet weight (mg/mm tibial length)	241 ± 7 ^b	239 ± 7 ^b	250 ± 15 ^b	333 ± 8 ^a	351 ± 11 ^a	325 ± 19 ^a	<0.0001	0.76	0.31
ALT activity (U/L)	25.9 ± 1.6 ^b	29.0 ± 2.6 ^b	27.6 ± 1.9 ^b	41.8 ± 2.0 ^a	39.6 ± 1.6 ^a	40.4 ± 5.9 ^a	<0.0001	0.99	0.68
AST activity (U/L)	64.3 ± 2.9 ^b	78.3 ± 5.5 ^{ab}	64.1 ± 3.5 ^b	88.3 ± 3.6 ^a	95.1 ± 4.3 ^a	86.7 ± 9.4 ^a	<0.0001	0.07	0.78

Value are means ± SEM, *n* = 10–12. Mean values in a row with unlike superscript letters are significantly different (*P* < 0.05). C, corn starch diet-fed rats; CO_r, orchidectomized rats fed corn starch diet; CL, leuprolide-treated rats fed corn starch diet; H, high-carbohydrate high-fat diet fed rats; HO_r, orchidectomized rats fed high-carbohydrate high-fat diet; HL, leuprolide-treated rats fed high-carbohydrate high-fat diet; LVIDd, left ventricular internal diameter during diastole; LVIDs, left ventricular internal diameter during systole; IVSd, interventricular septal thickness during diastole; IVSs, interventricular septal thickness during systole; LVPWd, left ventricular

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posterior wall thickness during diastole; LVPWs. left ventricular posterior wall thickness during systole; ESS, end-systolic stress; AST, aspartate transaminase; ALT, alanine transaminase.

Table 3. Effects of Omacor on leuprolide-induced changes in physiological and metabolic parameters

Variables	CL	CLOm	HL	HLOm	P value		
					Diet + leuprolide	Omacor	(Diet + leuprolide) × Omacor
Initial body weight (g)	337 ± 1	342 ± 2	338 ± 1	339 ± 3	0.61	0.13	0.31
Final body weight (g)	417 ± 8 ^b	319 ± 9 ^c	539 ± 18 ^a	536 ± 19 ^a	<0.0001	0.001	0.002
Water intake (mL/day)	26.4 ± 2.6 ^b	33.2 ± 2.6 ^a	17.5 ± 1.7 ^c	24.5 ± 1.1 ^b	0.0001	0.002	0.96
Food intake (g/day)	36.1 ± 2.2 ^a	38.3 ± 2.8 ^a	24.3 ± 1.9 ^b	26.1 ± 1.1 ^b	<0.0001	0.34	0.93
Energy intake (kJ/day)	401 ± 6 ^c	487 ± 34 ^b	507 ± 12 ^{ab}	560 ± 21 ^a	0.0001	0.002	0.44
Feed efficiency (g/kJ)	0.20 ± 0.02 ^b	-0.03 ± 0.01 ^c	0.40 ± 0.03 ^a	0.35 ± 0.03 ^a	<0.0001	<0.0001	0.0005
Abdominal circumference (cm)	20.1 ± 0.2 ^b	17.0 ± 0.2 ^c	23.0 ± 0.4 ^a	22.0 ± 0.4 ^a	<0.0001	<0.0001	0.002
Basal blood glucose concentrations (mmol/L)	4.8 ± 0.1 ^b	3.5 ± 0.1 ^c	5.4 ± 0.2 ^a	4.5 ± 0.3 ^b	0.0002	<0.0001	0.31
Area under the curve (mmol/L×minutes)	781 ± 10 ^b	599 ± 16 ^c	844 ± 24 ^b	933 ± 40 ^a	<0.0001	0.07	<0.0001
Total cholesterol (mmol/L)	1.65 ± 0.05 ^b	1.04 ± 0.03 ^d	1.89 ± 0.11 ^a	1.41 ± 0.07 ^c	0.0001	<0.0001	0.37
Triglyceride (mmol/L)	0.60 ± 0.06 ^b	0.27 ± 0.02 ^b	1.98 ± 0.31 ^a	0.33 ± 0.04 ^b	<0.0001	<0.0001	0.0004
NEFA (mmol/L)	2.38 ± 0.19 ^b	0.46 ± 0.03 ^c	4.63 ± 0.43 ^a	0.99 ± 0.18 ^c	<0.0001	<0.0001	0.0016
Whole-body lean mass (g)	314 ± 8 ^a	261 ± 10 ^b	308 ± 9 ^a	251 ± 13 ^b	0.44	<0.0001	0.85
Whole-body fat mass (g)	191 ± 8 ^b	59 ± 8 ^c	161 ± 15 ^b	248 ± 27 ^a	<0.0001	0.18	<0.0001
Bone mineral content (g)	0.173 ± 0.003	0.179 ± 0.003	0.177 ± 0.004	0.180 ± 0.001	0.40	0.14	0.62
Bone mineral density (g/cm ³)	12.0 ± 0.2 ^b	10.6 ± 0.5 ^b	11.3 ± 0.5 ^b	16.8 ± 0.9 ^a	<0.0001	0.001	<0.0001
Retroperitoneal fat (mg/mm tibial length)	220 ± 18 ^b	95 ± 14 ^c	479 ± 54 ^a	406 ± 50 ^a	<0.0001	0.015	0.50
Epididymal fat (mg/mm tibial length)	125 ± 11 ^b	67 ± 7 ^c	263 ± 26 ^a	226 ± 26 ^a	<0.0001	0.02	0.59
Omental fat (mg/mm tibial length)	83 ± 6 ^c	79 ± 10 ^c	147 ± 13 ^b	249 ± 20 ^a	<0.0001	0.0007	0.0003

Value are means ± SEM, *n* = 10–12. Mean values in a row with unlike superscript letters are significantly different (*P*<0.05). CL, leuprolide-treated rats fed corn starch diet; CLOm, leuprolide-treated rats fed corn starch diet supplemented with Omacor; HL, leuprolide-treated rats fed high-carbohydrate high-fat diet; HLOm, leuprolide-treated rats fed high-carbohydrate, high-fat diet supplemented with Omacor; NEFA, non-esterified fatty acids.

Table 4. Effects of Omacor on leuprolide-induced changes in cardiovascular and hepatic function

Variables	CL	CLOm	HL	HLOm	P value		
					Diet + leuprolide	Omacor	(Diet + leuprolide) × Omacor
Heart rate	238 ± 11	255 ± 22	290 ± 15	283 ± 11	0.08	0.75	0.44
LVIDd (mm)	7.70 ± 0.15 ^b	7.64 ± 0.17 ^b	8.24 ± 0.22 ^{ab}	8.54 ± 0.16 ^a	0.0002	0.50	0.32
LVIDs (mm)	4.52 ± 0.19 ^b	4.62 ± 0.13 ^b	4.14 ± 0.18 ^b	5.33 ± 0.09 ^a	0.29	0.0001	0.0009
IVSd (mm)	1.77 ± 0.06 ^{ab}	1.68 ± 0.05 ^b	1.82 ± 0.06 ^{ab}	1.91 ± 0.02 ^a	0.008	1.0000	0.08
IVSs (mm)	2.80 ± 0.08 ^b	2.72 ± 0.07 ^b	3.12 ± 0.11 ^a	2.92 ± 0.06 ^{ab}	0.0028	0.10	0.47
LVPWd (mm)	1.79 ± 0.03 ^{ab}	1.67 ± 0.03 ^b	1.75 ± 0.05 ^{ab}	1.84 ± 0.04 ^a	0.10	0.70	0.009
LVPWs (mm)	2.58 ± 0.15 ^b	2.39 ± 0.05 ^b	2.95 ± 0.10 ^a	2.65 ± 0.06 ^b	0.003	0.017	0.58
Diastolic volume (μL)	481 ± 26 ^b	473 ± 34 ^b	531 ± 47 ^b	658 ± 36 ^a	0.002	0.11	0.07
Systolic volume (μL)	100.0 ± 12.8 ^b	104.9 ± 8.9 ^b	77.0 ± 8.6 ^b	159.2 ± 7.8 ^a	0.12	<0.0001	0.0003
Stroke volume (μL)	381 ± 22 ^b	368 ± 25 ^b	454 ± 43 ^{ab}	499 ± 31 ^a	0.002	0.61	0.36
SBP:LVIDs	32.1 ± 1.5 ^b	26.5 ± 0.9 ^c	36.9 ± 2.4 ^a	25.9 ± 0.7 ^c	0.18	<0.0001	0.08
SBP:systolic volume	1599 ± 198 ^{ab}	1216 ± 101 ^b	2291 ± 479 ^a	882 ± 48 ^b	0.50	0.002	0.06
ESS:LVIDs	2.84 ± 0.16	2.55 ± 0.07	2.56 ± 0.11	2.61 ± 0.08	0.33	0.28	0.13
Cardiac output (mL)	90.6 ± 7.3 ^b	94.5 ± 10.8 ^b	130.9 ± 12.9 ^a	141.2 ± 10.0 ^a	0.0001	0.50	0.76
Relative wall thickness	0.47 ± 0.02	0.44 ± 0.01	0.45 ± 0.02	0.44 ± 0.01	0.53	0.21	0.53
Systolic wall stress	129.9 ± 12.2 ^{ab}	117.8 ± 4.0 ^{ab}	106.4 ± 7.3 ^b	139.0 ± 4.0 ^a	0.88	0.19	0.006
Estimated LV mass (g)	1.01 ± 0.04 ^b	0.92 ± 0.05 ^b	1.07 ± 0.06 ^b	1.29 ± 0.05 ^a	0.0001	0.21	0.004
Fractional shortening (%)	41.2 ± 2.1 ^b	39.6 ± 0.7 ^b	47.7 ± 2.3 ^a	37.6 ± 0.8 ^b	0.18	0.0009	0.013
Ejection fraction (%)	79.3 ± 2.0 ^b	77.9 ± 0.7 ^b	85.2 ± 1.7 ^a	75.6 ± 0.9 ^b	0.22	0.0004	0.006
Systolic blood pressure (mmHg)	144 ± 2 ^b	122 ± 1 ^d	150 ± 2 ^a	137 ± 2 ^c	<0.0001	<0.0001	0.016
Right ventricular wet weight (mg/mm tibial length)	2.76 ± 0.16 ^c	3.59 ± 0.29 ^b	3.19 ± 0.15 ^{bc}	4.71 ± 0.22 ^a	0.0008	<0.0001	0.11
Left ventricular + septum wet weight (mg/mm tibial length)	18.1 ± 0.5 ^b	17.2 ± 0.8 ^b	19.1 ± 0.5 ^b	22.9 ± 0.9 ^a	<0.0001	0.044	0.002
Left ventricular diastolic stiffness constant, κ	27.3 ± 0.5 ^a	22.8 ± 1.0 ^{bc}	27.9 ± 0.7 ^a	21.3 ± 1.0 ^c	0.59	<0.0001	0.21
Liver wet weight (mg/mm tibial length)	250 ± 15 ^b	161 ± 7 ^c	325 ± 19 ^a	294 ± 14 ^a	<0.0001	0.0001	0.05
ALT activity (U/L)	27.6 ± 1.9 ^b	41.4 ± 4.4 ^a	34.9 ± 2.4 ^{ab}	39.7 ± 2.8 ^a	0.36	0.004	0.15
AST activity (U/L)	64.1 ± 3.5 ^b	72.5 ± 3.1 ^{ab}	86.7 ± 9.4 ^a	70.0 ± 4.2 ^{ab}	0.08	0.47	0.032

Value are means ± SEM, $n = 10-12$. Mean values in a row with unlike superscript letters are significantly different ($P < 0.05$). CL, leuprolide-treated rats fed corn starch diet; CLOm, leuprolide-treated rats fed corn starch diet supplemented with Omacor; HL, leuprolide-treated rats fed high-carbohydrate high-fat diet; HLOm, leuprolide-treated rats fed high-carbohydrate, high-fat diet supplemented with Omacor; LVIDd, left ventricular internal diameter during diastole; LVIDs, left ventricular internal diameter during systole; IVSd, interventricular septal thickness during diastole; IVSs, interventricular septal thickness during systole; LVPWd, left ventricular posterior wall thickness during diastole; LVPWs, left ventricular posterior wall thickness during systole; ESS, end-systolic stress; AST, aspartate transaminase; ALT, alanine transaminase.

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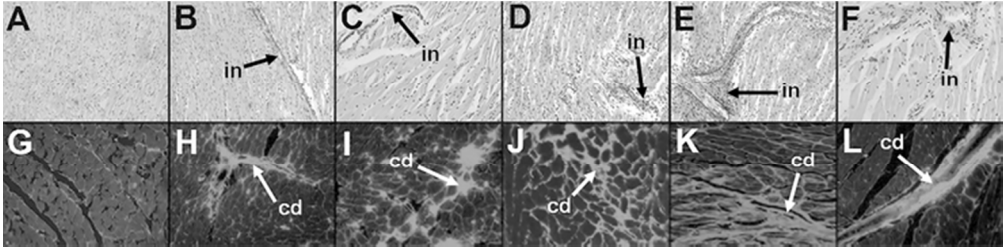


Figure 2. Effects of orchidectomy and leuprolide on the structure of the heart. Top row represents hematoxylin and eosin staining of left ventricle showing inflammatory cell ("in") while the bottom row represents picrosirius red staining of left ventricle showing collagen deposition ("cd") from C (A, G), COr (B, H), CL (C, I), H (D, J), HO (E, K), and HL (F, L) rats.

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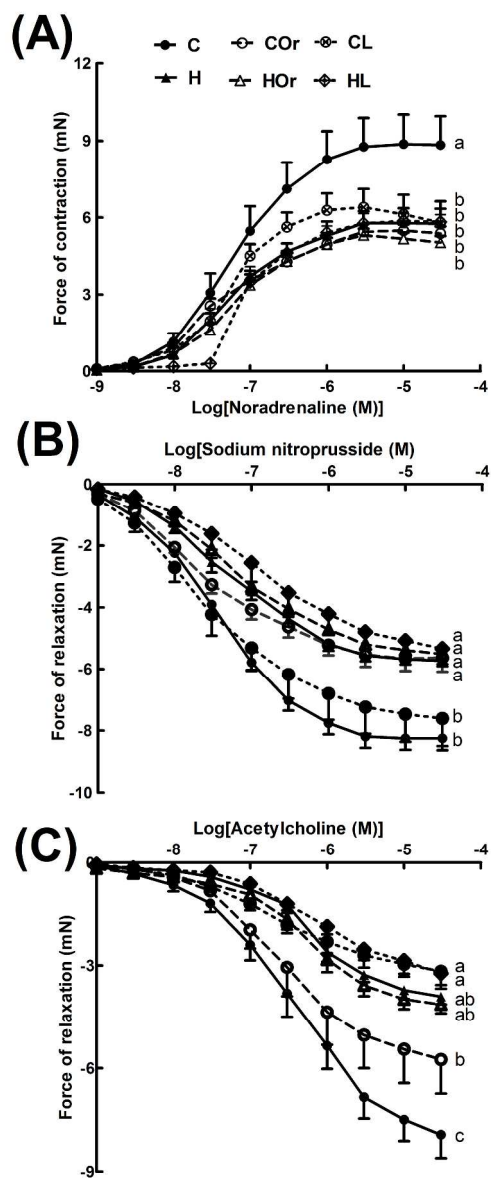


Figure 3. Cumulative concentration-response curves for noradrenaline (A), sodium nitroprusside (B), and acetylcholine (C) in thoracic aortic rings from C, COr, CL, H, HOr, and HL rats. Data are shown as mean \pm SEM, n = 10–12/group. End-point means without a common alphabet significantly differ, $P < 0.05$.

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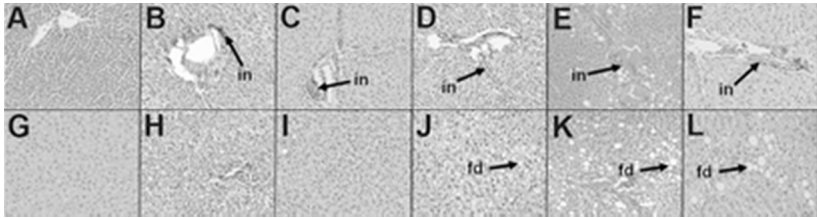


Figure 4. Effects of orchidectomy and leuprolide on the structure of the liver. Top row represents hematoxylin and eosin staining of liver showing inflammatory cell ("in") while the bottom row represents hematoxylin and eosin staining of liver showing fat deposition ("fd") from C (A, G), CO_r (B, H), CL (C, I), H (D, J), HO_r (E, K), and HL (F, L) rats.

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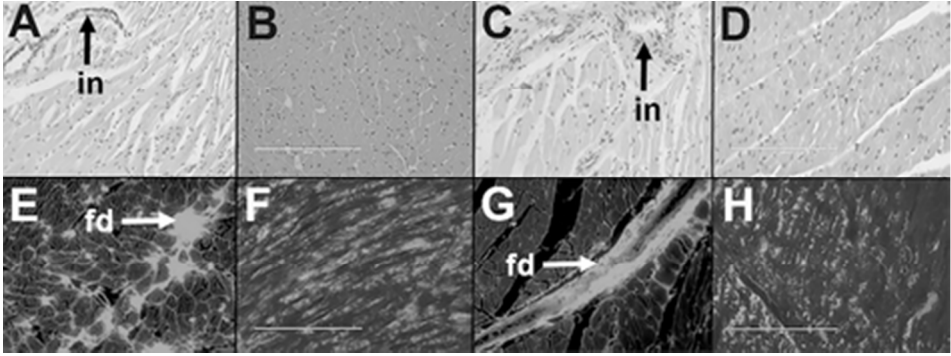


Figure 5. Effects of Omacor on leuprolide-induced changes in the structure of the heart. Top row represents hematoxylin and eosin staining of left ventricle showing inflammatory cell ("in") while the bottom row represents picrosirius red staining of left ventricle showing collagen deposition ("cd") from CL (A, E), CLOm (B, F), HL (C, G), and HLOm (D, H) rats.

40x14mm (300 x 300 DPI)

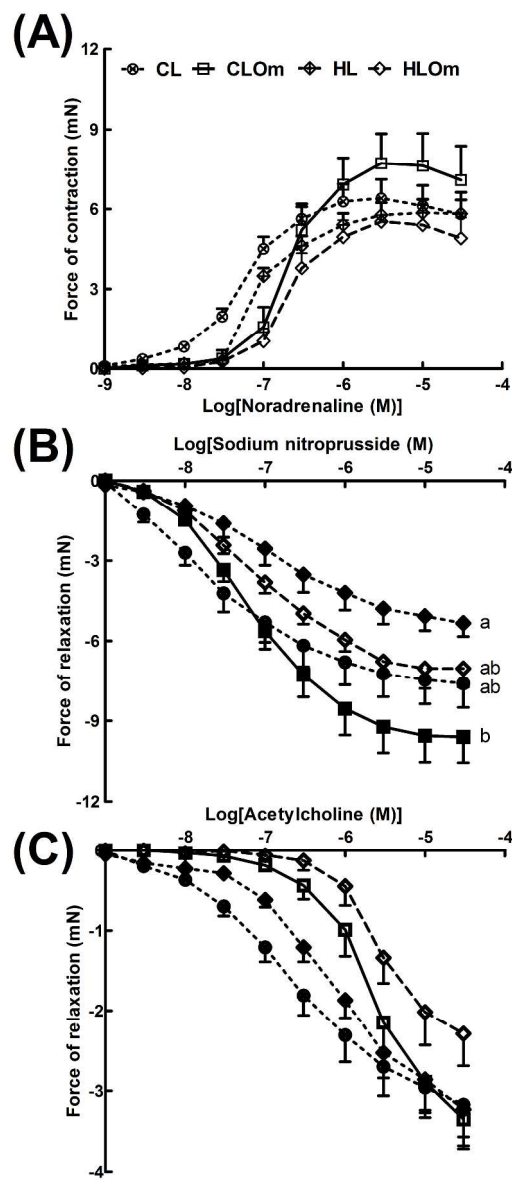


Figure 6. Cumulative concentration-response curves for noradrenaline (A), sodium nitroprusside (B), and acetylcholine (C) in thoracic aortic rings from CL, CLOm, HL, and HLOm rats. Data are shown as mean \pm SEM, n = 10-12/group. End-point means without a common alphabet significantly differ, $P < 0.05$.

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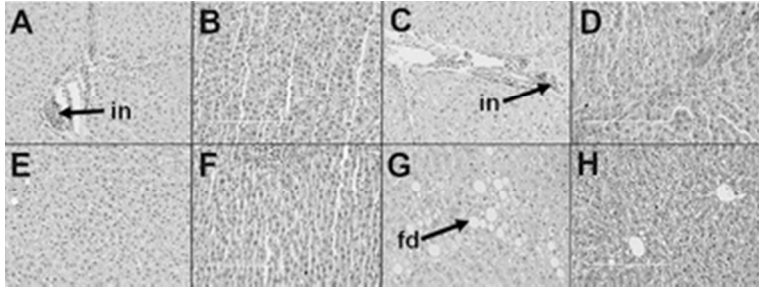


Figure 7. Effects of Omacor on leuprolide-induced changes in the structure of the liver. Top row represents hematoxylin and eosin staining of liver showing inflammatory cell ("in") while the bottom row represents hematoxylin and eosin staining of liver showing fat deposition ("fd") from CL (A, E), CLOm (B, F), HL (C, G), and HLOm (D, H) rats.

32x12mm (300 x 300 DPI)

Chapter 7. Discussion and conclusions

Summary of results: This thesis has emphasised preclinical studies with an appropriate and characterised diet-induced rat model that mimics metabolic syndrome in humans to demonstrate that foods are sources of potentially effective treatments. *Chapter 1* describes metabolic syndrome and its components, metabolic syndrome-associated cardiovascular complications and role of coffee and its components against the metabolic syndrome. Prevalence of metabolic syndrome is increasing worldwide [1]. The constellation of metabolic disorders such as obesity, elevated blood pressure, dyslipidaemia and insulin resistance or impaired glucose tolerance escalates the risk of cardiovascular diseases, hepatic disorders and type 2 diabetes, which can lead to death. Oxidative stress and chronic inflammation have crucial roles in the development of these metabolic consequences. Since coffee and its components show antioxidant and anti-inflammatory properties, it is logical to examine possible beneficial effects of coffee components against metabolic syndrome.

Chapter 2 describes the metabolic, cardiovascular and liver remodeling induced by high-calorie rich diet in rats and development of metabolic syndrome parameters such as abdominal obesity, hypertension, impaired glucose tolerance and dyslipidaemia. Previous studies from our research group in diet-induced obese rats with Colombian coffee extract containing caffeine, chlorogenic acid, trigonelline and diterpenoids showed reversal of the cardiovascular and liver remodelling to reduce hypertension and cardiac stiffness as well as non-alcoholic steatohepatitis, decreased impaired glucose tolerance and hypertension without affecting plasma lipids or abdominal obesity [2]. Further, caffeine produced different responses, especially with pronounced decreases in body weight [3]. My project extended these studies, initially by defining responses to green (unroasted) coffee and decaffeinated green coffee in rats with diet-induced metabolic syndrome. Both coffee products were provided as an extract incorporated in the food. Green coffee reduced body weight, abdominal fats and body mass index without altering plasma lipid contents indicating fat redistribution. Decaffeinated green coffee reduced body weight and retroperitoneal fat without causing change plasma content of lipid. Both decreased infiltration of inflammatory cells in the heart and the liver, and decreased collagen accumulation in the heart wall and fibrosis in the liver. The similarity of the responses suggests that caffeine is not the major bioactive compound in green coffee

reducing metabolic syndrome. Other compounds in green coffee that could have biological effects include chlorogenic acid, trigonelline, diterpenes such as cafestol and kahweol, and phenolic acids. The doses of chlorogenic acid are similar to literature studies which decreased some of the physiological changes in metabolic syndrome [4, 5], so this compound was then investigated.

Chapter 3 describes the responses to chlorogenic acid as an intervention in rats with diet-induced metabolic syndrome. There are several isomers of chlorogenic acid in nature, and my study investigated the most common, 5-O-caffeoylquinic acid. Chlorogenic acid decreased abdominal fat pads and body weight and improved cardiovascular and liver structure and function. This strongly suggests that chlorogenic acid produces at least some of the biological responses to green coffee. The roles of other compounds, especially trigonelline, the diterpenes present in coffee such as cafestol and kahweol, and phenolic acids, need further investigation.

Chapter 4 examines whether coffee waste could become a functional food. In the coffee production process, coffee waste and by-products cause problems of waste management and environmental contamination. High consumption of coffee requires the processing of a large number of coffee berries which generates coffee pulp as the waste from processing. Coffee pulp accounts for ~29% (w/w) of the dry weight of the whole berry. The dry matter of coffee pulp contains carbohydrates (15.7%), proteins (17.4%), crude fibre (14.1%), cellulose (20.7%), hemicellulose (3.6%), lignin (14.3%) [6] and minerals (especially potassium), along with tannins, polyphenols such as chlorogenic acid, and caffeine [7]. Hence, the use of coffee pulp may be considered as a potential approach for management of metabolic complications associated with obesity as there are many reports of health benefits from these individual components of coffee pulp. My study showed that coffee pulp intervention attenuated obesity, hypertension, improved glucose tolerance, cardiovascular and hepatic dysfunction while reducing blood lipid concentrations without affecting plasma total cholesterol concentrations and vascular reactivity.

Chapter 5 investigated the potential therapeutic effects of another major waste product in the production of coffee as a beverage, the spent coffee. This product contains

at least 1000 compounds, including the melanoidins produced during roasting of the beans by caramelisation and Maillard reaction. When given as a dietary intervention in high-carbohydrate, high-fat diet-fed rats, spent coffee reversed the metabolic, cardiovascular and liver abnormalities. Addition of spent coffee normalised blood glucose concentrations and delayed intestinal glucose absorption, reduced the elevated triglycerides and non-esterified fatty acids in plasma, and reversed abdominal obesity. Further, spent coffee reduced systolic blood pressure, cardiac wall stiffness and inflammation in the heart and liver. My study identified that gut microbiota played a major role in improving metabolic syndrome with spent coffee treatment.

Prostate cancer is one of the commonly diagnosed cancers in men and is treated by chemical or surgical induction of testosterone deficiency [8]. However, testosterone deficiency is associated with metabolic disorders such as central obesity, dyslipidaemia, impaired glucose tolerance, hepatic and cardiovascular remodelling [9]. *Chapter 6* has investigated whether omega-3 unsaturated fatty acids are effective as functional foods in testosterone deficiency in rats. Testosterone deficiency produced by castration or treatment with GnRH agonist increased the metabolic disturbances measured in rats fed a high-carbohydrate, high-fat diet. Addition of omega-3 unsaturated fatty acids to the diet decreased systolic blood pressure and left ventricular diastolic stiffness, reduced left ventricular infiltration of inflammatory cells and collagen deposition, reduced liver lipid accumulation and inflammatory cell infiltration without improving liver function. These results suggest that increased omega-3 unsaturated fatty acid supplementation could attenuate metabolic complications in prostate cancer patients with induced testosterone deprivation.

Conclusions: My results from these studies with coffee, coffee waste and fish oil demonstrated the functional food properties where coffee and fish oil-intervention markedly reduced cardiovascular and hepatic disorder. Furthermore, positive outcomes from coffee waste studies introduced an innovation for coffee waste management and has shown the unique concept to apply them as dietary supplements in humans for managing obesity-associated health complications. These studies have shown that these components can improve several parameters of metabolic syndrome, particularly reversing

cardiovascular and adipocytes changes. Moreover, other fruits such as pineapple, apple, potato, strawberry and tomato are also sources of chlorogenic acid. Hence, their by-products such as pineapple waste, apple pomace, potato peel, berry-juice processing waste could hold the relevant functional food properties against metabolic syndrome parameters. Human clinical trial could be the future prospective of coffee component and their wastes to describe their potential advantage against metabolic syndrome for human health. Overall, functional foods such as coffee and fish oil can be therapeutic supplement if taken in regular and proper amount as per individual needs.

Similar to coffee (green or roasted), many fruits and vegetables possess a range of compounds [10]. Some of the components from such fruits and vegetables have shown health benefits, including flavonoids, anthocyanins, ellagic acid and stilbenes [11]. As I have done in my thesis for coffee, many other foods still require extensive research to confirm whether they have sufficient bioactive compounds or whether we are discarding the most important components in food waste. Further, when considering the role of various components from a food, the food matrix should be considered in characterising its health effects [12]. Once health effects are defined for certain foods, food industries need to design the food products, functional foods or nutraceuticals that can attract compliance from the general community.

My thesis has defined responses to green coffee, coffee components and coffee waste products in a validated rat model of metabolic syndrome and obesity. As these interventions share antioxidant and anti-inflammatory activities, these responses may well be extended to other chronic diseases that are associated with oxidative stress and chronic inflammation, for example, inflammatory bowel disease, chronic kidney disease, non-alcoholic fatty liver disease and osteoarthritis. Previous and current studies from our laboratory have characterised rat models of these chronic disease [13-15] (inflammatory bowel disease model – under review).

Another aspect to be explored from my thesis is the advantage of food waste such as coffee pulp and spent coffee. Other food waste such as red grape marc from the wine industry, citrus waste from juice industry and spent cereals from beer brewing are some of the major food wastes from food processing. These wastes generate greenhouse gases

[16] and have an immense environmental footprint. Thus, identifying foods wastes that can provide nutraceuticals will not only provide a cheaper source of bioactive compounds for human health improvement but also reduce the environmental footprint. Once these health effects are defined, it will be important to identify the commercial viability of these products and then develop food products that can be accepted by the general community. For defining health benefits and developing food products, human intervention trials will need to be carried out to define the extent of the responses produced in humans. Success of these trials will help in convincing the community to consume these foods and hence will generate a viable market for food industries.

Although this study has provided evidence for effects of green coffee, coffee components and coffee wastes against diet-induced obesity, this study was unable to confirm molecular mechanisms involved in improvements of obesity-associated metabolic changes. Further, measuring a range of metabolic hormones such as insulin, leptin, adiponectin and resistin could supplement the data provided in this thesis. Further, identifying the role of gut microbiota for each intervention would further support the role of gut microbiota in improving metabolic complications in obesity. Further, sufficient and cost-effective availability of pure trigonelline, cafestol and kahweol would have allowed the study of the effects of these individual components of coffee. This would have further enabled me to extend my study outcomes to individual components.

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