

ORIGINAL INVESTIGATION

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# Eight week exposure to a high sugar high fat diet results in adiposity gain and alterations in metabolic biomarkers in baboons (*Papio hamadryas* sp.)

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## Abstract

**Background:** Baboons (*Papio hamadryas* Sp.) develop features of the cardiometabolic syndrome and represent a clinically-relevant animal model in which to study the aetiology of the disorder. To further evaluate the baboon as a model for the study of the cardiometabolic syndrome, we developed a high sugar high fat diet and hypothesized that it could be used to induce adiposity gain and affect associated circulating biomarkers.

**Methods:** We developed a diet enriched with monosaccharides and saturated fatty acids that was composed of solid and liquid energy sources. We provided a group of baboons (n = 9) *ad libitum* access to this diet for 8 weeks. Concurrently, a control group (n = 6) was maintained with *ad libitum* access to a low sugar low fat baseline diet and normal water for 8 weeks. Body composition was determined by dual-energy X-ray absorptiometry and circulating metabolic biomarkers were measured using standard methodology before and after the 8 week study period.

**Results:** Neither body composition nor circulating biomarkers changed in the control group. Following the 8 weeks, the intervention group had a significant increase in fat mass ( $1.71 \pm 0.98$  vs.  $3.23 \pm 1.70$  kg,  $p = 0.004$ ), triglyceride ( $55 \pm 13$  vs.  $109 \pm 67$  mg/dL,  $p = 0.006$ ), and leptin ( $1.19 \pm 1.40$  vs.  $3.29 \pm 2.32$  ng/mL,  $p = 0.001$ ) and a decline in adiponectin concentrations ( $33530 \pm 9744$  vs.  $23330 \pm 7863$  ng/mL,  $p = 0.002$ ). Percentage haemoglobin A<sub>1c</sub> ( $4.0 \pm 0.3$  vs.  $6.0 \pm 1.4$ ,  $p = 0.002$ ) also increased in the intervention group.

**Conclusions:** Our findings indicate that when exposed to a high sugar high fat diet, young adult male baboons develop increased body fat and triglyceride concentrations, altered adipokine concentrations, and evidence of altered glucose metabolism. Our findings are in keeping with observations in humans and further demonstrate the potential utility of this highly clinically-relevant animal model for studying diet-induced metabolic dysregulation.

## Background

The cardiometabolic syndrome represents one of the most significant current global public health issues [1]. Manifestations of the cardiometabolic syndrome include increased central adiposity, insulin resistance, and

dyslipidaemia [2]. Collectively these and other related abnormalities represent an increased risk for the development of chronic diseases such as type 2 diabetes and coronary heart disease [3]. The aetiology of the cardiometabolic syndrome is not completely understood. Studies in humans and in animal models have suggested that exposure to Western diets- particularly those containing sugar-sweetened beverages, contribute to its development [4-8]. The mechanisms underlying the

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contribution of diet to the cardiometabolic syndrome have yet to be fully elucidated.

Efforts to determine the mechanisms underlying the dietary inducement of the cardiometabolic syndrome could be enhanced through the use of more clinically-relevant animal models. The baboon (*Papio hamadryas* sp.) is an established non-human primate model with a long history in biomedical research [9,10]. Baboons are a long-lived old world monkey species (*Catarhini*) known to consume diets containing plants, fruits, and some animal matter [11,12]. Wild baboons feeding on human waste foods manifest metabolic abnormalities consistent with human obesity and the cardiometabolic syndrome [13,14]. Our previous work has shown that captive baboons are susceptible to obesity even when maintained on a diet low in fat and simple carbohydrates [15]. We, along with others, have also described impaired insulin signalling at the whole body and molecular level [16] and, in some cases overt diabetes [17] in spontaneously obese baboons.

To further evaluate the potential of the baboon model for the study of diet-induced cardiometabolic syndrome, we developed an experimental diet with a fatty acid and monosaccharide composition similar to that of a typical fast food diet. We hypothesized that we could induce adiposity gain and influence associated metabolic biomarkers using this diet. Herein, we report changes in fat mass, adipokines, triglyceride, and percentage haemoglobin A<sub>1C</sub> (%HbA<sub>1C</sub>) in baboons after only 8 weeks of *ad libitum* consumption of this diet.

## Methods

### Animals

Fifteen male baboons (*Papio hamadryas* Sp.) from the Southwest National Primate Research Center (San Antonio, TX) colony were studied. All animals were group-housed in outdoor enclosures according to established National Research Council guidelines. To minimize disruption to social rankings and normal behaviours, two established groups of animals were selected for the study. A Veterinarian performed standard health assessments, including blood chemistry and hematologic profiles, on all animals before the start of the study. Animals had no prior significant medical history. All study procedures were approved by the Institutional Animal Care and Use Committee of the Southwest Foundation for Biomedical Research, San Antonio, TX.

### Design

At baseline all animals had been consuming the commercially available 5LE0 solid feed (LabDiet, PMI, St. Louis, MO) *ad libitum*. This feed is high in complex carbohydrates and low in fat. Animals also had *ad libitum* access to normal water (Table 1). Following a 12 hour overnight fast, animals were sedated with ketamine

HCl (10 mg/Kg) before undergoing baseline body composition assessment and collection of blood samples. Each group was then randomly assigned to receive one of two diets for 8 weeks:

1) Control group (n = 6) - continued *ad libitum* access to the low sugar low fat baseline diet and normal water; and

2) Intervention group (n = 9) - *ad libitum* access to the palatable high sugar high fat experimental diet containing solid and liquid energy sources and normal water.

Twelve-hour fasted blood samples and body composition assessments were again collected after 8 weeks of dietary exposure.

### Experimental diet

Preliminary experiments were undertaken on singly-housed baboons to determine the palatability of various high saturated fatty acid solid feed preparations. The feed found to be most palatable was prepared using 73% Purina Monkey Chow 5038 (a grain-based meal), 7% lard, 4% Crisco, 4% coconut oil, 10.5% flavoured high fructose corn syrup, and 1.5% water. Vitamins and mineral preparations were added to match the micronutrient composition of the baseline feed. Palatability was enhanced using non-caloric artificial fruit flavours and by lightly baking the feed for 12 minutes at 300°F. Liquid calories were offered to the intervention group in a beverage that was prepared using water, high fructose corn syrup (2.83 Kcal/g, 76% sugar, 41.8% fructose, 34.2% dextrose, ISOSWEET 5500, Staley, Decatur, IL), and artificial fruit flavouring. The sweetened water contained 110 mL of high fructose corn syrup per litre. A summary of the nutritional composition of the solid feed and the sweetened water is given in Table 1.

### Body composition

Body weight was measured on an electronic scale (GSE 665, Texas Scales Inc., Cibolo, TX). Dual energy X-ray absorptiometry (DXA) body composition scans were undertaken using a Lunar Prodigy densitometer (GE Healthcare, Madison, WI). Animals were placed in the supine position on the DXA bed and extremities were positioned within the scanning region. Scans were analyzed using encore2007 software version 11.40.004 (GE Healthcare, Madison, WI). Total body, trunk region (torso), and limb (arm + leg) region compositions were determined. The coefficients of variation for total fat mass and total lean mass for two replicate scans in 3 baboons were found to be 2.2 and 2.3%, respectively.

### Glucose, %HbA<sub>1C</sub>, serum lipid, and lipoprotein cholesterol measurements

Fasting plasma concentrations of glucose, triglyceride, total cholesterol, LDL cholesterol, and HDL-cholesterol

**Table 1 Energy composition of diets**

	Energy density	%Kcal as fat	%Kcal as carbohydrate	%Kcal as sugar	%Kcal as protein
5LE0 solid chow <sup>1</sup>	3.26 <sup>3</sup>	13.81	67.16	2.95	19.03
High fat chow <sup>2</sup>	4.33 <sup>3</sup>	38.46	51.00	10.72	10.53
Sweetened water <sup>1</sup>	0.42 <sup>4</sup>	0	100.00	100.00	0

<sup>1</sup>From chemical analysis of macronutrient composition provided by manufacturer, <sup>2</sup>from chemical analysis of nutrient composition from Covance Laboratories (Madison, WI), <sup>3</sup>Kcal/g, <sup>4</sup>Kcal/mL.

were determined using an ACE clinical analyzer (Alfa Wasserman Diagnostic Technologies LLC; West Caldwell, NJ). Total haemoglobin and %HbA<sub>1c</sub> were determined from whole blood using the ACE clinical analyzer.

#### Plasma insulin, C-reactive protein, and adipokine measurements

Plasma insulin, leptin, and total adiponectin concentrations were determined using commercially available radioimmunoassay kits [Millipore Inc., Billerica, MA]; the intra-assay CVs were 2.2%, 2.2%, and 5.2%, respectively. Plasma C-reactive protein was measured using the Monkey C-reactive protein ELISA kit [Life Diagnostics Inc., West Chester, PA].

#### Statistical analyses

Data from each diet group was analyzed independently; pre- and post-intervention means were evaluated using two-tailed paired samples *t*-tests. Variable values were converted to common logarithms to correct for deviations from the normal distribution when necessary. Statistical significance was set at *P* < 0.05. Statistical analyses were undertaken using SYSTAT version12 [Chicago, IL].

#### Results

Basic descriptive characteristics of the study animals are presented in Table 2. All animals passed physical examinations and had normal blood chemistry and hematologic profiles at the study outset. None of the groups had a statistically significant change in body weight, although weight change did approach statistical significance in the intervention group (*p* = 0.09).

No differences were observed between baseline and 8 week body composition and metabolic variables in the control group (Table 3). Animals in the intervention

**Table 2 Descriptive statistics**

	Controls (n = 6)	Intervention (n = 9)
Age at baseline (yrs.)	7.4 ± 0.3	7.8 ± 0.5
Baseline weight (Kg)	26.6 ± 2.8	29.0 ± 3.0
8-week weight (Kg)	26.5 ± 2.4	30.1 ± 3.3

Data are mean ± SD. Controls had continued *ad libitum* access to the baseline diet and water. Intervention group had *ad libitum* access to the experimental diet.

group gained fat mass, trunk fat mass, and percentage fat mass (Table 4). The proportion of total fat mass gain that occurred in the trunk region was 75.5 ± 31.5%. Triglyceride, leptin, and %HbA<sub>1c</sub> also increased significantly from baseline in this group (Table 4). In addition, adiponectin concentrations decreased significantly after the 8 week feeding period. Fasting glucose, insulin, C-reactive protein, total-, LDL-, and HDL-cholesterol concentrations were not significantly affected (Table 4).

#### Discussion

Baboons spontaneously develop obesity, insulin resistance, dyslipidaemia, and type 2 diabetes. We hypothesized that we could induce adiposity gain and alter circulating metabolic biomarkers in baboons by exposing a group to an experimental diet designed to be representative of a typical fast food diet containing solid and liquid energy sources. The animals gained fat mass, particularly in the trunk region and had significant increases in triglyceride, leptin, and %HbA<sub>1c</sub>, and a decline in adiponectin concentrations. Our findings

**Table 3 Body composition and metabolic variables at baseline and after 8-weeks in the control group**

	Baseline	8 weeks	p value
Total body fat mass (Kg)	1.50 ± 0.68	1.46 ± 0.37	0.74
Total body lean mass (Kg)	23.5 ± 2.3	23.2 ± 2.3	0.68
Percentage fat mass	5.7 ± 2.1	5.6 ± 1.2	0.70
Trunk fat mass (Kg)	0.89 ± 0.56	0.89 ± 0.28	0.93
Trunk lean mass (Kg)	12.3 ± 1.4	12.2 ± 1.4	0.91
Limb fat mass (g)	453 ± 102	426 ± 77	0.13
Limb lean mass (kg)	8.9 ± 1.1	8.9 ± 1.0	0.36
Bone mineral content (g)	1104 ± 69	1095 ± 107	0.66
Fasting glucose (mg/dL)	88.8 ± 3.4	97.0 ± 10.7	0.20
Fasting insulin (μU/mL)	11.4 ± 4.3	25.4 ± 23.4	0.34
%Hemoglobin A <sub>1c</sub>	4.4 ± 0.3	5.7 ± 2.7	0.21
Total cholesterol (mg/dL)	91.2 ± 15.6	85.7 ± 15.7	0.27
LDL cholesterol (mg/dL)	35.7 ± 10.7	34.2 ± 9.3	0.67
HDL cholesterol (mg/dL)	44.2 ± 10.6	40.3 ± 9.1	0.29
Triglyceride (mg/dL)	55.5 ± 10.3	55.8 ± 15.0	0.89
Leptin (ng/mL)	0.99 ± 0.42	1.06 ± 0.23	0.58
Adiponectin (ng/mL)	29556 ± 4153	27930 ± 5505	0.13
C-reactive protein (ng/mL)	44.8 ± 30.6	47.2 ± 42.0	0.80

Data are mean ± SD. Controls had continued *ad libitum* access to the baseline diet and water. Means comparisons were made using paired *t*-tests. Variables were log<sub>10</sub> transformed before statistical analyses were undertaken, n = 6.

**Table 4 Body composition and metabolic variables at baseline and after 8-weeks in the intervention group**

	Baseline	8 weeks	p value
Total body fat mass (Kg)	1.71 ± 0.98	3.23 ± 1.70	0.004
Total body lean mass (Kg)	25.7 ± 2.7	24.8 ± 2.6	0.003
Percentage fat mass	5.9 ± 3.3	10.8 ± 5.4	0.002
Trunk fat mass (Kg)	1.04 ± 0.84	2.38 ± 1.45	0.004
Trunk lean mass (Kg)	13.3 ± 1.3	12.9 ± 1.3	0.11
Limb fat mass (g)	489 ± 132	626 ± 204	0.02
Limb lean mass (kg)	9.8 ± 1.2	9.6 ± 1.2	0.22
Bone mineral content (g)	1237 ± 122	1242 ± 124	0.72
Fasting glucose (mg/dL)	95.4 ± 16.6	87.6 ± 5.1	0.22
Fasting insulin (μU/mL)	18.4 ± 16.1	18.9 ± 8.8	0.95
%Hemoglobin A <sub>1c</sub>	4.0 ± 0.3	6.0 ± 1.4	0.002
Total cholesterol (mg/dL)	103.3 ± 28.7	108.2 ± 20.9	0.22
LDL cholesterol (mg/dL)	38.0 ± 12.5	37.9 ± 14.5	0.71
HDL cholesterol (mg/dL)	55.0 ± 17.8	48.7 ± 13.7	0.42
Triglyceride (mg/dL)	55 ± 13	109 ± 67	0.006
Leptin (ng/mL)	1.19 ± 1.40	3.29 ± 2.32	0.001
Adiponectin (ng/mL)	33530 ± 9744	23330 ± 7863	0.002
C-reactive protein (ng/mL)	53.0 ± 48.8	81.5 ± 83.1	0.44

Data are mean ± SD. Intervention group had *ad libitum* access to the experimental diet and water. Means comparisons were made using paired *t*-tests. Variables were log<sub>10</sub> transformed before statistical analyses were undertaken, n = 9.

demonstrate that, in as little as 8 weeks, baboons exposed to a high sugar high fat diet gained fat mass and manifested biochemical alterations consistent with the early stages of metabolic dysregulation.

Baboons are found primarily in the East African savannah although populations are known to extend over much of the continent. They are diurnal and terrestrial animals that have been noted as indiscriminate consumers of vegetable and animal matter. Among wild baboons feeding on human foods, some were shown to have significantly greater body mass, leptin, insulin, and glucose concentrations relative to their normal diet-consuming counterparts [13]. Given the relatively recent common ancestry, baboons and humans are genetically, anatomically, and physiologically very similar. Baboons and humans also appear to develop similar metabolic disorders; earlier studies have documented spontaneous obesity, insulin resistance, and a form of adult-onset diabetes in captive baboons [15-19]. We, along with others, have described genetic effects on obesity and cardiovascular disease risk factors in keeping with observations in human populations [15,18,20]. In addition, baboons are known to manifest dyslipidemia and develop fatty arterial lesions when exposed to diets high in saturated fatty acids and cholesterol [21-23]. Combined, these observations suggest that the baboon model could prove very useful for furthering our understanding of the aetiology of the cardiometabolic syndrome.

We provided a group of young male adult baboons *ad libitum* access to a palatable high sugar high fat experimental diet for 8 weeks, while a control group of similar age and weight were maintained on the baseline low sugar low fat diet. The experimental diet was composed of a solid feed with a macronutrient composition similar to that of a typical fast food burger and French fries and a sweetened drink with a monosaccharide composition similar to that of a typical soft drink.

Body composition did not change in the control group. The intervention group gained fat mass but lost total body lean mass, and as a result, significant changes in body weight did not occur. The majority of the gain in fat mass occurred in the trunk region. This pattern is consistent with the adiposity pattern that is associated with central obesity in human males [24] and with fat gain patterns observed in moderate fat-fed dogs [25]. It is not clear why the animals lost lean body mass and we are unaware of similar findings in other overfeeding studies [26-28]. The animals had *ad libitum* access to water throughout the study period and it is unlikely that the diet had an effect on hydration status. The protein content of the experimental solid feed was less than that of the baseline feed, but would have been adequate to maintain lean mass. However, it is possible that the animals' protein intake was diminished as a significant quantity of their energy intake came from the sweetened water.

Fasting plasma glucose and insulin concentrations were not significantly affected in the intervention group. Nevertheless, mean %HbA<sub>1c</sub> increased by almost 70%. This apparent discrepancy may have several causes. It is possible that hepatic insulin resistance, reflected primarily by fasting glucose and insulin concentrations, had not developed after only 8 weeks. Alternatively, ketamine, used to sedate the animals before the blood draws, is known to have effects on glucose and insulin concentrations [29] and may have obscured the diet's effects on fasting glucose and insulin concentrations. The acute changes in glucose resulting from the ketamine sedation would not have affected %HbA<sub>1c</sub>. If the former explanation is true, then the increase in glycated haemoglobin must have resulted from elevated post-prandial glucose concentrations in the presence of emerging peripheral tissue insulin resistance. Direct assessments of hepatic and peripheral tissue insulin resistance would be required to address these issues. Regardless, the rise in %HbA<sub>1c</sub> suggests that alterations in glucose homeostasis may have developed after only 8 weeks of consuming the diet. In support, the decline in adiponectin concentrations is consistent with a worsening of glucose homeostasis [30]. Notably, these baboons had a very low body fat percentage at baseline. It is conceivable that a larger effect on glucose metabolism

could have been observed if the animals had greater body fat at baseline.

We did not observe changes in cholesterol concentrations in response to the diet. Nonetheless, fasting triglyceride and leptin concentrations were markedly increased while adiponectin concentrations decreased. Earlier studies in baboons have not described triglyceride changes in animals exposed to high saturated fatty acid diets [31]. However, exposure to high monosaccharide diets have been shown to result in elevated triglycerides in this species [23]. In humans, 4 week consumption of fructose at 17% of total energy intake has been shown to increase fasting triglyceride without influencing cholesterol [32]. Several other studies have also shown triglyceride to be an early responder to high monosaccharide (particularly fructose) overfeeding in humans [33-35]. Adipokines are known to respond to dietary manipulation [36]. Elevated leptin has also been shown to be an early response to high fat and high carbohydrate overfeeding in humans and in rodents [37-39]. Similar findings have been reported for adiponectin in human studies [40]. Interestingly, the triglyceride and leptin responses of the baboons are consistent with those reported in a 4 week overfeeding study in humans in which a solution of fructose was used to deliver the excess dietary energy: fasting insulin and total cholesterol concentrations also remained unaltered in that study [41]. Hence, intake of the monosaccharide enriched sweetened water was likely to be responsible for the triglyceride response observed in this group of baboons. Because of the constraints of the outdoor housing environment, it was not possible to accurately quantify energy intake. Hence, we cannot dismiss the possibility that triglyceride and adipokine responses were attributable to increased total energy intake. This would have to be determined using another experimental design and was not the objective of this study. Nevertheless, our estimates of solid feed and drink consumption suggest that the energy intake of the treatment group was much higher than that of the control group and that a large proportion of the excess energy intake came from the sweetened water (data not shown). Regardless, the alterations in triglyceride, leptin, and adiponectin observed in baboons exposed to our diet are consistent with those seen in human overfeeding studies, particularly those that tested the effects of monosaccharide overfeeding. Taken together, the data suggest that triglyceride, leptin, and adiponectin may be early responders to excessive energy intake and may represent markers of emerging metabolic dysregulation.

Importantly, the short 8 week dietary exposure leaves additional questions, particularly those relating to whether this diet could result in, and if so the time course taken to develop, overt cardiometabolic syndrome or related end-stage diseases. The rate of fat mass accrual suggests that the time taken to become

overtly obese would be rather short. Indeed, studies undertaken in the closely-related rhesus macaque species have documented an earlier onset of type 2 diabetes than in humans [42]. Future studies will be designed to address these questions.

## Conclusions

Our findings indicate that young male baboons exposed to a high sugar high fat diet, composed of solid and liquid energy sources, developed multiple metabolic abnormalities consistent with emergent cardiometabolic syndrome. Our findings further demonstrate the potential utility of this animal model for studying diet-induced metabolic dysregulation.

## List of abbreviations

DXA: dual energy X-ray absorptiometry; %HbA<sub>1c</sub>: percentage haemoglobin A<sub>1c</sub>.

## Acknowledgements

We acknowledge the efforts of Dr. Pat Frost, D.V.M., the Veterinary research technicians, animal caretaker staff, and behavioural staff of the Southwest National Primate Research Centre. We also acknowledge the use of facilities kindly provided by Don Strange of Texas, Inc. This study was funded by the Max and Minnie Tomerlin Voelcker Fund, San Antonio, TX and was conducted using facilities constructed with support from the Research Facilities Improvement Program under grant number C06 RR (numbers 014578, 013556, 015456, 017515) from the National Center for Research Resources and with support from National Institutes of Health grants PO1 HL028972 and P51 RR013986. This work was also supported, in part, by a small equipment grant from the Southwest Foundation for Biomedical Research Founders Council. PBH and JMP were supported by postdoctoral fellowships from the Brackenridge Foundation of San Antonio Texas.

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## Authors' contributions

AGC, JCLA, RAB, and PBH conceived and designed the study. PBH wrote the manuscript. PBH, JCLA, and AGC analyzed the data. JCLA, MGC, RAB, VSV, RES, PBH, and AGC developed and conducted preliminary tests on the diet. JCLA, PBH, MGF, JMP, VSV, MET, VM, SAC, and KH participated in data collection and biochemical analyses. Manuscript edits were performed by JCLA, VSV, MET, KH, SAC, RES, and AGC. Additional consultation on study design and animal behaviour was provided by RES. All authors have read and approved the final manuscript.

## Competing interests

The authors declare that they have no competing interests.

Received: 25 July 2010 Accepted: 29 October 2010

Published: 29 October 2010

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doi:10.1186/1475-2840-9-71

**Cite this article as:** Higgins et al.: Eight week exposure to a high sugar high fat diet results in adiposity gain and alterations in metabolic biomarkers in baboons (*Papio hamadryas* sp.). *Cardiovascular Diabetology* 2010 9:71.