

Metabolism of Genetically Obese Rats on Normal or High-Fat Diet

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Summary. Adult genetically obese *fafa* rats showed a high level of lipogenesis from glucose in liver but not in adipose tissue; pancreatic content and serum levels of insulin were elevated. Glucose uptake and insulin sensitivity were decreased in muscle. *fafa* rats and their lean littermates fed a high-fat diet showed increased fat deposits. Serum insulin levels were not significantly affected by diet in either group. The larger the fat cells were, the more actively they utilised glucose; insulin sensitivity was influenced both by diet and cell size. Control rats made obese by a high caloric diet did not show insulin resistance in muscle. — The data indicate

that in adult obesity in these rats, even in the presence of marked hyperinsulinism, increased lipogenesis in adipose tissue is not a prerequisite. Rather, fat storage is a consequence of increased uptake of circulating triglycerides. On a diet rich in carbohydrate, adipose tissue fatty acids were mainly of hepatic origin; on a high-fat diet they were of dietary origin.

Key words: High-fat diet, rat, nutritional obesity, genetic obesity, adipose tissue, liver, muscle metabolism, insulin sensitivity, pancreatic glucagon, pancreatic and plasmatic insulin.

Obesity is associated with hyperinsulinism i.e. increased insulin secretion and high plasma insulin levels. This hyperinsulinism is partly related to a high carbohydrate intake [1]. Long term feeding of a high fat diet to genetically obese *obob* mice (a) markedly increases their obesity, (b) has no effect on their hyperglycemia, but (c) decreases serum insulin levels [2]. Moreover, obesity can be induced in lean animals by feeding a high-fat-low-carbohydrate diet given *ad lib.* [3]. This kind of experimental obesity is characterized by normal levels of circulating insulin *in vivo*, and *in vitro* by a decreased insulin response to glucose [4]. These data indicate that obesity *per se* is not closely related to hyperinsulinism.

In the present investigation, genetically obese *fafa* rats were studied. They are known to present with a high degree of hyperinsulinism [5, 6], high levels of serum lipids [7] and an elevated lipogenesis of adipose tissue [8]. The aim of the study was to investigate the

Materials and Methods

Five month old male genetically obese *fafa* rats [7] and their lean littermates (FaFa or *Fafa*) were fed *ad lib* for 7 months either a control diet T (9% of calories as lipid, 22% as protein, and 69% as carbohydrate), or a high-fat (72% of calories as lipid, 22% as protein, and 16% as carbohydrate) diet S [3]. Four groups of rats were obtained. Animals were killed by decapitation in the fed state in the morning. Blood was collected and serum glucose was assayed by the glucose oxidase method (Boehringer Mannheim Test). Serum insulin was measured by the method of Rosselin *et al.* [9]. The pancreatic content of insulin was determined as previously described [4]. Pancreatic glucagon was assayed according to Jarrouse *et al.* [10].

The incorporation of glucose- ^{14}C into CO_2 , lipids or glycogen in liver slices, hemidiaphragms and peri-

Table 1. Pancreatic insulin and glucagon, serum insulin and glucose of lean Fa or genetically obese fa rats fed a control T or a high-fat diet S. The animals were maintained for 7 months on the diets T or S until the age of 12 months, time of sacrifice

Groups	n	Body weight (g)	Serum glucose (mg/100 ml)	Serum insulin ($\mu\text{U}/\text{ml}$)	Pancreatic weight (mg)	Stored insulin (U per pancreas)	Stored glucagon (μg per pancreas)
Fa T	8	530 \pm 12.2	132 \pm 5.6	138 \pm 11.5	1404 \pm 77.7	1.85 \pm 0.149	8.37 \pm 0.440
Fa S	9	566 \pm 16.1	119 \pm 3.8	167 \pm 42.3	1341 \pm 65.2	1.48 \pm 0.185	7.06 \pm 0.293
fa T	6	672 \pm 36.4	134 \pm 4.7	536 \pm 101	1314 \pm 97.5	3.22 \pm 0.549	5.22 \pm 0.400
fa S	6	885 \pm 31.0 ^b	121 \pm 3.1	422 \pm 69.9	1414 \pm 64.0	3.15 \pm 0.387	6.58 \pm 0.386 ^a

^a ($p < 0.05$) and ^b ($p < 0.01$) degree of statistical significance in differences versus the corresponding control group (Fa T or fa T).

locus of fat synthesis in the obese animals and their lean littermates, and the possible influence of different diets. Special attention was given to the role of insulin and of fat cell size in this model of hereditary and/or dietary obesity.

genital adipose tissue pieces, was measured as described elsewhere [11]. Insulin (1 mU/ml) was added to the incubation medium in the case of adipose tissue and muscle. Measurements of adipose tissue cellularity have been described [3].

Results

Body weight was significantly increased by the combination of high-fat diet and the obesity genes of *fafa* rats (Table 1).

Serum glucose tended to decrease slightly in rats on high fat diet. Pooling obese (*fafa*) and nonobese (FaFa) type rats, the high fat diet reduced blood glucose levels significantly ($p < 0.01$). Serum insulin was not significantly changed by high fat diet. The most striking observation was a four-fold increase in circulating insulin levels in the obese group on control diet.

As shown in Table 1, the pancreas of obese rats contained twice as much insulin as those of lean animals ($p < 0.01$); this was not affected by diet. Pancreatic glucagon content was similar in all groups.

Adipose Tissue Metabolism

Fat cell volume was 2.2 fold increased in the epididymal adipose tissue of genetically obese animals regardless of diet composition. In the lean rats however, high fat diet induced a 50% increase in fat cell volume above control. Changes in cell number were not observed in this site [12]. Fig. 1 shows the basal incorporation of glucose- ^{14}C into CO_2 , total glycerol and total fatty acids (FA) of adipose tissue fragments. All results are expressed per fat cell. Total dpm recovered as $^{14}\text{CO}_2$ and labeled lipids were twice as high ($p < 0.01$) in the obese rats indicating an increased glucose uptake in these animals. High fat diet increased this value as well ($p < 0.01$) but to a lesser extent. Whereas FA incorporated only 5–10% of total ^{14}C uptake, the bulk of radioactivity was found in the glycerol moiety. Fig. 1 also shows that in the *basal* state, labeled glycerol was augmented by both genetic obesity and high fat diet. By contrast, ^{14}C incorporation into FA was increased only by genetic obesity, but not by diet.

The effect of insulin added *in vitro* (1 mU/ml) was most marked for FA synthesis (Fig. 1c). Adipose tissue of animals on control diet was significantly ($p < 0.05$) more responsive. The same was true for $^{14}\text{CO}_2$ production. By contrast, insulin-induced glycerol synthesis was insignificant whatever the group. When the results were related to fat cell surface, most of these differences were maintained.

Liver Metabolism

Fig. 2 shows ^{14}C -glucose incorporation into CO_2 and phospholipids as well as into the glycerol and FA moieties of tissue triglyceride (TG) of incubated liver slices. Most of the label was recovered in CO_2 and the bulk of changes in glucose uptake was reflected in alteration of FA synthesis (Fig. 2d).

High fat diet had an impressive effect as it reduced glucose uptake into TG-FA by 90%. All other parameters measured were reduced by this diet. Genetic

obesity, by contrast, markedly increased glucose uptake into liver tissue.

Muscle Metabolism

Glucose uptake and its conversion into CO_2 and/or glycogen of hemi-diaphragms are depicted in Fig. 3. About 9/10 of glucose taken up was converted into glycogen.

In the basal state it appears clear that fat diet and to a lesser extent genetic obesity decreased all three: glucose uptake, CO_2 production and glycogen synthesis ($p < 0.01$). Glucose uptake and glycogen synthesis in diaphragms of *fafa* rats were not further reduced by fat diet.

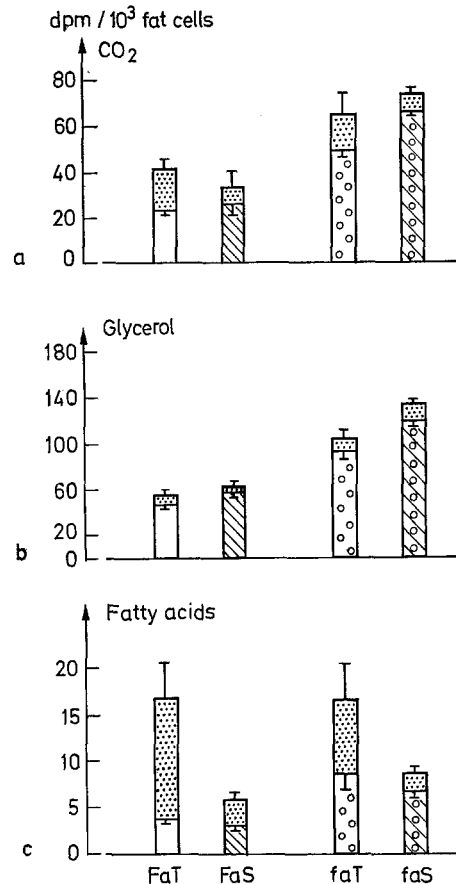


Fig. 1. Glucose ^{14}C incorporation into CO_2 (a), total glycerol (b), total fatty acids (c) of adipose tissue fragments incubated with or without insulin (1 mU/ml). Mean \pm SEM. \square lean control rats; \square genetically obese rats; \square rats fed a high-fat diet S. Fat cell volume ($10^3 \mu^3$) and number of rats in each group: Fa T: 766 ± 73 (8); Fa S: 1114 ± 143 (7); fa T: 1673 ± 139 (6); fa S: 1663 ± 55 (6)

The addition of 1 mU insulin per ml of medium increased glucose uptake and glycogen synthesis in all groups. The effect of insulin on muscle was reduced by obesity but not by diet.

Discussion

The cause of obesity is still poorly understood. Various animal models have been studied to shed light onto possible pathogenic mechanisms. The present study deals with the obesity of the genetically obese *fafa* rats and the influence of diet composition.

As Table 1 shows, the genetically obese animals were heavier than their lean siblings. A diet with a

partly because in those animals, the diet was begun earlier in life than in the present study [13, 14].

In order to analyse the possible mechanisms involved in the pathogenesis of obesity in these *fafa* rats, the three main organ systems involved in diabetic metabolism have been studied *in vitro*, namely adipose tissue, muscle and liver.

Adipose tissue of the adult genetically obese rats showed a much higher glucose uptake, the bulk of

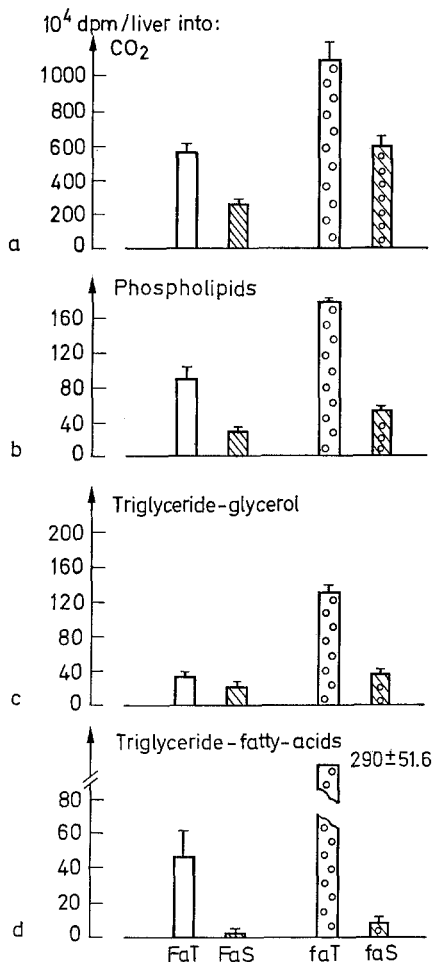


Fig. 2. Glucose $U^{14}C$ incorporation into CO_2 (a), phospholipids (b) and triglycerides (c, d) of liver slices incubated *in vitro*. Mean \pm SEM. \square lean controls rats, \boxtimes rats on a high fat diet S; \square genetically obese rats. Weight (g) of the livers, (n° of rats in each group): Fa T 17.3 ± 0.65 (7); Fa S 15.3 ± 0.52 (8); fa T 24.1 ± 2.61 fa S 23.3 ± 0.90

high fat content increased not only the fat stores of the genetically obese, but also of the lean rats [12]. Thus, body weight appears to be influenced by at least two factors, the genetic background and the composition of the diet. Similar results have been described in *obob* mice [2]. Studies done with Swiss mice and Wistar rats showed a more marked effect of diet [3, 13],

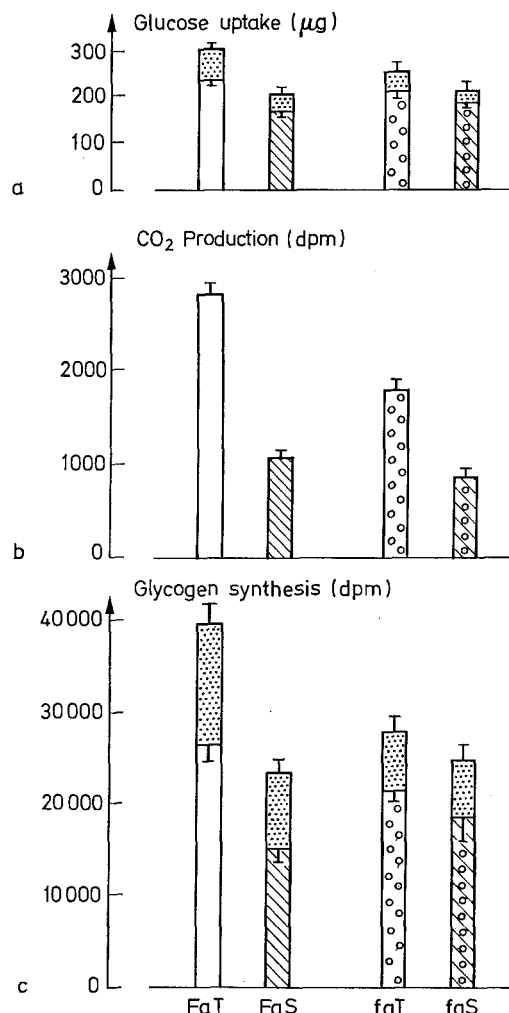


Fig. 3. Glucose uptake and glucose $U^{14}C$ incorporation into CO_2 and glycogen per \log_{10} 100 mg of hemidiaphragms incubated with \boxtimes or without insulin (1 mU/ml). Mean \pm SEM. \square Control rats, \boxtimes rats on a high fat diet S \square genetically obese rats. Weight (mg) of the 2 hemidiaphragms (N° of rats): Fa T 741 ± 43.2 (7); Fa S 738 ± 33.7 (8); fa T 592 ± 25.5 (8); fa S 675 ± 15.5 (6)

which was incorporated into glycerol (Fig. 1). FA synthesis was not particularly marked. However, the effect of insulin added *in vitro* was most striking in increasing FA synthesis and by far the most impressive in the animals (lean or obese genetically) on control diet. In muscle, as one might expect, about 90% of glucose taken up was incorporated into glycogen. High

fat diet decreased glucose uptake and, thereby, glycogen synthesis and CO₂ production.

Liver slices of genetically obese rats showed a six fold incorporation rate into TG-FA compared to that of lean animals. Therefore, the main site for lipogenesis in *adult fafa* rats appears to be the liver and not adipose tissue. On the high fat diet, lipogenesis was reduced by more than 90% in lean and obese animals, indicating that diet is an important regulator of hepatic lipogenesis.

It was of particular interest to study the conditions *in vivo* which may have influenced the behaviour of the organs *in vitro*. Hyperinsulinemia appears to be a characteristic of genetically obese *fafa* rats (Table 1). However, high fat diet reduced serum insulin levels slightly whereas fat stores increased markedly [12]. This may be due partly to the lower carbohydrate intake which could not be avoided with the high fat diet, as long as protein supply had to be constant. It is quite possible that the insulin levels *in vivo* were directly responsible for the behaviour of the isolated organs *in vitro*. Indeed, adipose tissue of *fafa* rats exposed to very high insulin levels *in vivo*, showed higher rates of glucose metabolism (Fig. 1). This was all the more marked in liver slices where up to sixfold differences could be observed (Fig. 2). The connection between hyperinsulinemia and hepatic lipogenesis has been suggested by Steiner [15] and Letarte [16]. In muscle, interestingly, *fafa* rats showed less glucose metabolism. By contrast, although no dramatic changes in insulin levels were obtained by feeding the high fat diet, the glucose uptake decreased markedly in liver and muscle tissue. In adipose tissue such an effect was found only for the insulin response in respect to FA synthesis.

Although the main FA synthesis occurs in the liver, *fafa* rats have an increased adipose mass where glucose is mainly incorporated into α -glycerophosphate to form triglyceride. The FA moiety comes from circulating triglycerides as suggested by the doubled lipoprotein lipase activity found in the fat cells of these *fafa* rats [17]. A similar argument applies to the higher glucose incorporation into triglyceride glycerol of adipose tissue of obese and lean animals fed the high fat diet, i.e. the increased supply of FA (in this case from exogenous sources) needed more α -glycerophosphate for triglyceride synthesis. Thus, feeding a high fat diet seems to speed the effects of ageing [18]. Fatty acid composition of adipose tissue reveals that dietary fatty acids are stored as such in adipose tissue [11]. Human studies with adult obese and lean subjects have lead to similar conclusions: lipogenesis is insignificant in adipose tissue [19, 20].

The role of fat cell size deserves special consideration. Since *fafa* rats have adipocytes of twice the volume of lean rats, one might implicate this fact in the higher glucose uptake of *fafa* adipose tissue. Indeed, when adipocytes of lean rats were increased in volume by high fat diet, glucose uptake of adipose tissue *in*

vitro was equally increased. Similar results have been obtained by Smith in human obesity [20] and by Stern in *fafa* rats [21]. An additional increase of glucose utilisation was observed in the *fafa* rat fed a high-fat diet although cell size did not change any further. Insulin resistance has been suggested to be related to the increased fat cell size by a decreased number of binding sites [22]. The present data confirm this argument in part: *in vitro* added insulin had a smaller effect on tissue of *fafa* rats which have larger adipocytes. However, although high fat diet increased fat cell size and decreased insulin sensitivity in the lean rats, in the obese rats the insulin effect was diminished by high fat diet with no further change in fat cell size. It appears, therefore, that basal glucose uptake and insulin sensitivity are not dependent on fat cell size only.

In conclusion, the data presented here show the importance of diet composition in relation to obesity of genetically obese *fafa* rats. In adults the bulk of adipose tissue FA appears to derive from either dietary fatty acids or from fatty acids synthesized in the liver. Hyperinsulinemia is probably an important factor contributing to obesity in this animal as well as to insulin resistance in muscle. Also, fat cell size induced by either genetic background or by high fat diet appears to be related to increased glucose metabolism and partly to insulin resistance.

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