### **ORIGINAL CONTRIBUTIONS**





# Effects of Sleeve Gastrectomy on Nonalcoholic Fatty Liver Disease in an Obese Rat Model

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Published online: 26 December 2017 © Springer Science+Business Media, LLC, part of Springer Nature 2017

### Abstract

**Background/Aim** Laparoscopic sleeve gastrectomy (SG) is an increasingly used bariatric surgery, which is reported to be effective for nonalcoholic fatty liver disease (NAFLD). Recently, activation of farnesoid X receptor (FXR), which is a nuclear receptor of bile acid (BA), was reported to contribute to the resolution of NAFLD. However, it is unclear whether SG has an effect on expression of FXR in the liver. We aimed to investigate the expression of FXR and its related factors in the liver after SG and to clarify the relationship between changes in FXR expression and NAFLD in an obese rat model.

**Methods** Thirty male Zucker fatty rats were divided into three groups: sham-operated (SO) control, pair-fed (PF) control, and SG. Eight weeks after the surgery, metabolic parameters, plasma levels of total BA and liver enzymes, liver triglyceride (TG) content, and mRNA expression of FXR and its related factors, such as small heterodimer partner (SHP) and peroxisome proliferator-activated receptor  $\alpha$  (PPAR $\alpha$ ), were measured.

**Results** Metabolic parameters in the SG group were significantly improved compared with the SO group. Liver enzymes and TG were significantly lower in the SG group than in the SO group. Plasma levels of BA were significantly higher in the SG group than in the SO and PF groups. mRNA expression of FXR, SHP, and PPAR $\alpha$  in the liver was significantly higher in the SG group than in the SO group.

**Conclusions** These results suggest that the effects of SG on NAFLD should be associated with the expression of the FXR pathway in the liver in a Zucker fatty rat model.

Keywords Sleeve gastrectomy · NAFLD · Bariatric surgery · Farnesoid X receptor · Bile acid

### Introduction

The prevalence of obesity has increased globally in recent decades [1]. Obesity is reported to be associated with significantly higher all-cause mortality compared with normal weight and is now one of the greatest public health problems on a worldwide scale [2]. Although bariatric surgery, as well as nonsurgical treatments, is available for obesity, bariatric surgery is more effective in losing body weight and resolving obesity-related comorbidities, such as diabetes and hyperlipidemia [3, 4]. Recently, sleeve gastrectomy (SG) has become a markedly increased bariatric procedure used worldwide [5, 6]. We previously reported improvements in glucose and lipid metabolisms in an obese diabetic rat model [7, 8]. In addition, we also reported changes in the hypothalamic feeding center following SG in a diet-induced obese rat model [9]. Recently, SG was reported to improve nonalcoholic fatty liver disease (NAFLD) in a diet-induced NAFLD rat model [10].

NAFLD is defined by the presence of hepatic steatosis either by imaging or by histology and exclusion of secondary hepatic fat accumulation such as alcohol consumption, use of medication, or hereditary disorders [11]. Risk factors associated with NAFLD include obesity, diabetes mellitus, dyslipidemia, and hypertension [12, 13]. Recently, the prevalence of NAFLD has increased with the rise of obesity and metabolic syndrome. In addition, NAFLD increases overall mortality deriving from cardiovascular disease and liver-related diseases, such as liver cirrhosis and hepatocellular carcinoma,

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and has become a serious public health issue [14]. Treatment of NAFLD includes intensive lifestyle intervention, such as dietary restrictions, exercise, and drug therapy. Johnson et al. reported that regular aerobic exercise reduced hepatic lipids in obesity even in the absence of body weight reduction [15]. Promrat et al. reported that weight reduction achieved through lifestyle intervention led to improvement of liver histology in nonalcoholic steatohepatitis (NASH) [16]. Thiazolidinediones and vitamin E have been reported to significantly resolve NAFLD [17, 18]. Recently, a farnesoid X receptor (FXR) agonist, obeticholic acid, significantly resolved NAFLD and insulin sensitivity in a randomized controlled clinical trial in patients with type 2 diabetes [19, 20]. It was also demonstrated that bariatric surgery is effective in patients with NAFLD and morbid obesity [21, 22].

FXR, first cloned in 1995 from rat liver cDNA, belongs to a family of nuclear hormone receptors and is highly expressed in the liver, intestines, and kidney [23]. Bile acid (BA) is a physiological ligand of FXR [24]. It was demonstrated that expression of FXR and activation of FXR by BA in the liver were associated with resolution of hepatic steatosis [25, 26]. Serum BA was reported to increase after SG, and BA may have the potential to contribute to resolution of morbid obesity and its comorbidities in animals and humans [27, 28]. However, it is unclear how SG influences the expression of FXR and its related factors in the liver and resolves NAFLD. Therefore, the aims of this study were to evaluate the expression of FXR and its related factors after SG and to clarify the relationship between changes in the expression and NAFLD in a Zucker fatty rat model.

### **Materials and Methods**

### Animals

Thirty male Zucker fatty rats (8 weeks old) were obtained from Charles River Japan, Inc. (Saga, Japan) and housed in individual cages under controlled temperature  $(24 \pm 2 \text{ °C})$ , at  $50 \pm 10\%$  humidity, and a 12 h-light cycle (7:00 am–7:00 pm) with ad libitum access to standard rat chow (CE-2, Clea Japan, Tokyo, Japan) and tap water. Fourteen days before surgery, the rats were acclimated to their local facilities. This study was approved by the Animal Committee of Oita University (Oita, Japan) and conformed to the Guidelines for Animal Experimentation of Oita University. All applicable institutional and/or national guidelines for the care and use of animals were followed.

### Surgical Procedure

The rats were divided into three groups—sham-operated (SO) control group (n = 10), pair-fed (PF) control group (n = 10),

and SG group (n = 10)—and were fasted for 24 h before surgery, which was performed under anesthesia (4% sevoflurane; Maruishi Pharmaceutical Co., Osaka, Japan). SG was performed as described previously [7, 29]. Briefly, the greater curvature from the antrum to the fundus across the forestomach and glandular stomach was incised, and approximately 90% of the forestomach and 70% of the glandular stomach were removed. The incision line in the stomach was then closed using 5-0 PDS® in three layers, to create the gastric sleeve. The SO and PF control rats underwent laparotomy, and their stomachs were elevated and returned to the abdominal cavity. The SO and SG groups had free access to standard rat chow for 8 weeks after surgery. The amount of food in the PF group was yoked ad lib to the weekly intake observed in the SG group as previously described [9]. Body weight and food intake were measured (Animal Scale; Clare, Tokyo, Japan) weekly, in all the groups (at 10:00 am). Livers and blood samples were collected 8 weeks after surgery.

### **Biochemical Tests**

Blood glucose, total cholesterol (TC), triglyceride (TG), free fatty acid (FFA), aspartate transaminase (AST), and alanine transaminase (ALT) levels were estimated using an H7180 automatic biochemical analyzer (Hitachi, Tokyo, Japan). Enzyme-linked immunosorbent assay (ELISA) kits were used to evaluate plasma insulin level (rat insulin ELISA kit; Shibayagi, Gunma, Japan), high molecular weight adiponectin (mouse/rat high molecular weight adiponectin ELISA kit; Shibayagi, Gunma, Japan), glucagon-like peptide-1 (GLP-1) (YK160 GLP-1 EIA; Yanaihara Institute Inc., Shizuoka, Japan), ghrelin (YK251 Rat GIP (Active) ELISA kit; Yanaihara Institute Inc.), and total BA (total bile acids assay half kit; Cosmobio, Tokyo, Japan). To evaluate GLP-1, total blood was treated with dipeptidyl dipeptidase (DDP)-IV inhibitor at the moment of extraction. DDP-IV is a peptide in the blood that causes the degradation of GLP-1. To evaluate insulin resistance, the homeostasis model assessment ratio (HOMA-R) was calculated by the formula: HOMA-R = fasting glucose (mmol/L) × fasting insulin ( $\mu$ U/ mL)/22.5 [30].

### **Liver TG Content**

Liver TG was measured as described previously [8]. In brief, liver samples (200 mg) were homogenized using a tissue homogenizer and then centrifuged at  $10,000 \times g$  for 10 min at 4 °C. The TG content of the samples was then determined using a commercial kit (Triglyceride E-test kit; Wako Pure Chemical Industries, Ltd., Osaka, Japan).

### **Liver Histological Analysis**

Histological changes in the liver were evaluated using hematoxylin and eosin (H&E) and oil red O staining under a light microscope. After freezing at -80 °C, the liver was stained with oil red O (oil red O for microscopy Certistain, Merck, Germany) to reveal intracellular lipids. A pathologist who was blinded to the study evaluated all histological sections at  $\times$  200 magnification.

### Quantitative Real-Time PCR for mRNA Quantification of FXR, SHP, PPARα, LXRα, and SREBP1c

Total RNA isolation was performed as described previously [9]. Quantitative real-time polymerase chain reaction (PCR) was performed as described previously with a Light Cycler system (Roche Diagnostics, Lewes, East Sussex, UK) [9]. The sequences of the primers used are listed in Table 1. Data were analyzed using the LightCycler analysis software (Roche Diagnostics), and a standard curve correlating cycle number with the amount of products formed was plotted for each sequence of interest. mRNA expression of FXR, small hetero-dimer partner (SHP), peroxisome proliferation-activated receptor  $\alpha$  (PPAR $\alpha$ ), liver X receptor  $\alpha$  (LXR $\alpha$ ), and sterol regulatory element-binding protein 1c (SREBP1c) was then normalized to rat glyceraldehyde-3-phosphate dehydrogenase (GAPDH).

### **Statistical Analysis**

All data are expressed as mean  $\pm$  standard deviation. All data were evaluated using one-way analysis of variance with Bonferroni correction for multiple comparisons. A *P* value < 0.05 was considered to be statistically significant. Statistical analysis was performed using the Statistical Package for the Social Sciences (SPSS) II software (SPSS, Inc., Chicago, IL, USA).

Table 1	Oligonucleotides	used in quantitative	real-time PCR
	0		

FXR	(Forward)	5'-CCACGACCAAGCTATGCAG-3'
	(Reverse)	5'-TCTCTGTTTGCTGTATGAGTCCA-3'
SHP	(Forward)	5'-GCAGCACTGCCTGGAGTC-3'
	(Reverse)	5'-GTGTGCAATGTGGCAGGA-3'
PPARα	(Forward)	5'-TGCGGACTACCAGTACTTAGGG-3'
	(Reverse)	5'-GGAAGCTGGAGAGAGGGTGT-3'
LXRα	(Forward)	5'-CAGGAAGAGATGTCCTTGTGG-3'
	(Reverse)	5'-TCTTCCACAACTCCGTTGC-3'
SREBP1c	(Forward)	5'-ACAAGATTGTGGAGCTCAAGG-3'
	(Reverse)	5'-TGCGCAAGACAGCAGATTTA-3'

### Results

# Changes in Body Weight, Food Intake, and Liver Weight

All animals survived the operations. Body weight after surgery was significantly lower in the PF and SG groups than the SO group; however, there was no significant difference in weight between the PF and SG groups (Fig. 1a). Weekly food intake except 8 weeks after surgery was significantly decreased in the PF and SG groups compared with the SO group (Fig. 1b). Liver weight 8 weeks after surgery was significantly lower in the PF and SG groups than the SO group, and there was no significant difference between the PF and SG groups (Fig. 1c).

# Changes in Metabolic Parameters and Hormones in the Plasma

The mean plasma levels of glucose, TC, TG, FFA, insulin, HOMA-R, adiponectin, GLP-1, ghrelin, AST, ALT, and total BA 8 weeks after surgery are shown in Table 2. Levels of TG, HOMA-R, AST, and ALT in the SG group were significantly lower than the SO group; however, there were no significant differences between the SO and PF groups. Significant differences in TC and FFA were only seen between the SO and PF groups. Levels of insulin and ghrelin were significantly lower in the SG group than the SO and PF groups. Conversely, levels of adiponectin, GLP-1, and total BA were significantly higher in the SG group than the SO and PF groups.

# Tissue TG Content and Histological Changes in the Liver

Liver TG content 8 weeks after surgery was significantly lower in the SG group than the SO group, and there were no significant differences between SO and PF groups (Fig. 2). Oil red O staining revealed that intracellular lipid in the liver was markedly decreased in the SG group compared with the SO group and slightly decreased compared with the PF group (Fig. 3).

# mRNA Expression of FXR, SHP, PPAR $\alpha$ , LXR $\alpha$ , and SREBP1c in the Liver

mRNA expression of FXR, SHP, and PPAR $\alpha$  in the liver was significantly higher in the SG group than the SO group (Fig. 4a–c). Conversely, mRNA expression of LXR $\alpha$  and SREBP1c were significantly lower in the SG group than the SO group (Fig. 4d, e). There were no significant differences in mRNA expression of FXR and its related factors between the SG and PF groups, but the SG group had better lipid metabolism in the liver than the PF group.





Fig. 1 Changes in body weight and food intake, and liver weight. a Changes in body weight after surgery in the sham-operated (SO) control, pair-fed (PF) control, and sleeve gastrectomy (SG) groups.

### Discussion

Liver steatosis results from disruption of hepatic TG metabolism and is influenced by various factors, such as insulin, GLP-1, and adiponectin. Insulin induces enhanced expression of SREBP1c in the hepatocyte and hyperinsulinemia with the insulin resistance enhanced hepatic steatosis [31]. GLP-1

Table 2Metabolic parametersand hormones 8 weeks aftersurgery

\*P < 0.05 versus the SO group. **b** Weekly food intake after surgery in the SO, PF, and SG groups. \*P < 0.05 versus the SO group. **c** Liver weight after surgery in the SO, PF, and SG groups. \*P < 0.05

agonists significantly reduce hepatic expression of SREBP-1c and increase expression of PPAR $\alpha$  [32]. Hepatic steatosis is enhanced in adiponectin knockout mice compared with wild-type mice [33]. FXR is also associated with hepatic TG metabolism, and FXR activation suppresses liver steatosis by downregulating lipogenesis and promoting TG oxidation. FXR induces SHP, which in turn suppresses the expression

	SO	PF	SG
Glucose (mg/dL)	$193.8\pm89.9$	167.8 ± 31.4	$161.2 \pm 14.2$
TC (mg/dL)	$207.2\pm48.1$	$163.4 \pm 17.2*$	$140.0\pm18.8^*$
TG (mg/dL)	$822.1\pm458.4$	$596.5 \pm 151.2$	$271.5 \pm 120.7*$
FFA (µEQ/L)	$810.0\pm479.2$	$365.0 \pm 104.1 *$	$360.6 \pm 158.6 *$
Insulin (µIU/L)	$9.6\pm3.2$	$8.1 \pm 2.3$	$4.2 \pm 2.9^{*^{\#}}$
HOMA-R	$4.7\pm2.7$	$3.5 \pm 1.3$	$1.7 \pm 1.1*$
Adiponection (ng/mL)	$32.5 \pm 12.7$	$19.9\pm8.5$	$76.2 \pm 43.9^{*\#}$
GLP-1 (pg/mL)	$5.8\pm2.4$	$6.7\pm7.0$	$31.8 \pm 31.0^{\ast \#}$
Ghrelin (fmol/mL)	$15.1 \pm 6.5$	$15.6\pm11.0$	$3.8\pm1.7^{\ast\#}$
AST (IU/L)	$262.7\pm155.4$	$158.9\pm53.0$	$98.8\pm26.6*$
ALT (IU/L)	$148.8\pm57.5$	$113.5 \pm 51.8$	$63.5 \pm 7.4*$
TBA (µmol/L)	$2.3 \pm 1.2$	$2.8 \pm 1.2$	$6.6 \pm 5.3^{*\#}$

*TC* total cholesterol, *TG* triglyceride, *FFA* free fatty acid, *HOMA-R* homeostasis model assessment ratio, *GLP-1* glucagon-like peptide-1, *AST* aspartate transaminase, *ALT* alanine transaminase, *TBA* total bile acid, *SO* shamoperated, *PF* pair-fed, *SG* sleeve gastrectomy

\*P < 0.05 versus the SO group, #P < 0.05 versus the PF group



Fig. 2 Tissue triglyceride (TG) content in the liver. SO, sham-operated; PF, pair-fed; and SG, sleeve gastrectomy. \*P < 0.05

of SREBP1c. SREBP1c is a critical transcription factor that regulates hepatic TG synthesis by inducing key enzymes involved in lipogenesis. Watanabe et al. reported that activation of SHP by FXR suppresses LXR $\alpha$ , which is a promoter of SREBP1c [26]. In addition to suppressing lipogenesis, FXR promotes TG oxidation through the activation of PPAR $\alpha$  [34].

Losing weight is the gold standard treatment of NAFLD. Bariatric surgery, which can achieve weight loss, has been reported to resolve NAFLD in animals and humans. We previously demonstrated that SG significantly resolved hepatic steatosis and increased hepatic expression of PPAR $\alpha$  in an obese diabetic rat model [8]. Myronovych et al. compared SG with SO and PF controls in high-fat diet-induced obese mice. SG and PF mice lost equal weight after surgery, but SG mice had the lowest hepatic TG content [28]. Recently, Talavera-Urquijo et al. reported that SG was superior to a very low-calorie diet in point of insulin resistance and cardiovascular risk markers related to NAFLD [10]. Meta-analyses have reported the effects of bariatric surgery on NAFLD in clinical setting. Bower et al. reported that bariatric surgery was associated with a significant resolution in not only histological features of NAFLD, including steatosis, fibrosis, hepatocyte, ballooning, and lobular inflammation, but also biochemical markers including AST and ALT [21]. Mummadi et al. reported that pooled proportions of patients with resolution of steatosis, steatohepatitis, and fibrosis were 91.6, 81.3, and 65.5%, respectively, and with complete resolution of NASH was 69.5% [22]. As for the effects of SG on NAFLD, Algooneh et al. demonstrated that the rate of complete resolution of NAFLD was 56% postoperatively, and a significant resolution was related to achievement of more than 50% excess weight loss [35]. Although the effects of bariatric surgery on NAFLD have been reported, the mechanism remains unclear.

Some reports have demonstrated that resolution of NAFLD is associated with FXR activation by BA. In animals, FXRdeficient mice were reported to exhibit hepatosteatosis and hyperlipidemia compared with the wild type [36]. In addition, it was also shown that mice with obesity or aging exhibited hepatosteatosis and lipidemia and decreased expression of FXR [26, 37]. Aguilar-Olivos et al. reported that protein expression of FXR in the liver was decreased in patients with NASH compared with those with simple steatosis, and FXR expression was correlated with progression of NAFLD [38]. The effects of endogenous and synthetic FXR agonists on NAFLD were evaluated in animals and humans. Watanabe et al. reported that the activation of FXR by BA suppressed the expression of SREBP1c and inhibited hepatosteatosis [26]. Pineda et al. reported that BA increased expression of PPAR $\alpha$  through the FXR activation [34]. Furthermore,



Fig. 3 Histological changes of the liver in H&E staining (a) and oil red O staining (b) under a light microscope (× 200). SO, sham-operated; PF, pair-fed; and SG, sleeve gastrectomy



**Fig. 4** mRNA expression of farnesoid X receptor (FXR) (**a**), small heterodimer partner (SHP) (**b**), peroxisome proliferator-activated receptor  $\alpha$  (PPAR $\alpha$ ) (**c**), liver X receptor  $\alpha$  (LXR $\alpha$ ) (**d**), and sterol regulatory-binding protein 1c (SREBP1c) (**e**) in the liver were

quantified by real-time PCR and expressed as a ratio to glyceraldehyde-3-phosphate dehydrogenase (GAPDH). SO, sham-operated; PF, pair-fed; and SG, sleeve gastrectomy. \*P < 0.05

Neuschwander-Tetri et al. evaluated the efficiency of obeticholic acid on NAFLD in a randomized controlled trial and reported that the histological features of NASH in the obeticholic acid group showed more resolution than in the placebo group [20].

There have been some reports about the relationship between SG, FXR, and BA. Ryan et al. demonstrated reduced body weight, altered feeding behavior, improved glucose tolerance, and altered composition of cecal microbial communities after SG in wild-type mice but not FXR knockout mice [39]. Myronovych et al. reported that elevated serum BA after SG in mice changed serum BA composition, which included elevated cholic and tauroursodeoxycholic acids, might be associated with a reduction of hepatic steatosis [28]. Belgaumkar et al. also showed that SG changed serum BA composition and these changes of BA were associated with a reduction of insulin resistance, proinflammatory cytokines, and cytokeratin-18, a marker of hepatocyte apoptosis [40].

In this study, we used Zucker fatty rats as a NAFLD model. Zucker fatty rats are widely used not only as an animal model of genetic obesity and metabolic syndrome but also NAFLD [41]. Liver weight and TG content were significantly lower in the SG group than the SO group but not in the PF group. Similarly, mRNA expression of FXR and its related factors was significantly changed in the SG group compared with the SO group but not the PF group. Therefore, this study demonstrates that SG activated expression of FXR and its related factors in the liver and resolved NAFLD through changed mRNA expression.

Until now, it has been unclear whether expression of FXR is independent of weight loss. However, weight loss by bariatric surgery and diet therapy influences changes of the microbiota, which can affect BA and BA composition [42, 43]. In fact, the BA composition in SG and PF groups was clearly altered compared with that in high-fat native and SO groups [28]. Since the change in BA composition can affect FXR expression [44], it seems that the expression may be partially dependent on weight loss. This study also showed that SG can influence insulin, adiponectin, GLP-1, ghrelin, and BA, independent of weight. However, this study has limitations. First, our results were obtained from a rodent model, which was different from humans. Second, although NAFLD and expression of FXR and its related factors were similarly changed after SG, this study did not functionally evaluate the FXR pathway using the antagonists. Therefore, further experimental and clinical studies are necessary to investigate the association between SG and activation of FXR pathway.

In conclusion, these results suggest that the effects of SG on NAFLD should be associated with the expression of the FXR pathway in the liver in a Zucker fatty rat model.

Acknowledgments We thank Ms. Mayumi Takeda and Yuiko Aso for the technical assistance.x

### **Compliance with Ethical Standards**

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

**Ethical Approval** All applicable institutional and/or national guidelines for the care and use of animals were followed.

**Informed Consent** Does not apply.

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