

## ORIGINAL ARTICLE

## Surgical removal of inflamed epididymal white adipose tissue attenuates the development of non-alcoholic steatohepatitis in obesity

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**BACKGROUND:** Non-alcoholic fatty liver disease (NAFLD) is strongly associated with abdominal obesity. Growing evidence suggests that inflammation in specific depots of white adipose tissue (WAT) has a key role in NAFLD progression, but experimental evidence for a causal role of WAT is lacking.

**METHODS:** A time-course study in C57BL/6J mice was performed to establish which WAT depot is most susceptible to develop inflammation during high-fat diet (HFD)-induced obesity. Crown-like structures (CLS) were quantified in epididymal (eWAT), mesenteric (mWAT) and inguinal/subcutaneous (iWAT) WAT. The contribution of inflamed WAT to NAFLD progression was investigated by surgical removal of a selected WAT depot and compared with sham surgery. Plasma markers were analyzed by enzyme-linked immunosorbent assay (cytokines/adipokines) and lipidomics (lipids).

**RESULTS:** In eWAT, CLS were formed already after 12 weeks of HFD, which coincided with maximal adipocyte size and fat depot mass, and preceded establishment of non-alcoholic steatohepatitis (NASH). By contrast, the number of CLS were low in mWAT and iWAT. Removal of inflamed eWAT after 12 weeks (eWATx group), followed by another 12 weeks of HFD feeding, resulted in significantly reduced NASH in eWATx. Inflammatory cell aggregates (−40%;  $P < 0.05$ ) and inflammatory genes (e.g., TNF $\alpha$ , −37%;  $P < 0.05$ ) were attenuated in livers of eWATx mice, whereas steatosis was not affected. Concomitantly, plasma concentrations of circulating proinflammatory mediators, viz. leptin and specific saturated and monounsaturated fatty acids, were also reduced in the eWATx group.

**CONCLUSIONS:** Intervention in NAFLD progression by removal of inflamed eWAT attenuates the development of NASH and reduces plasma levels of specific inflammatory mediators (cytokines and lipids). These data support the hypothesis that eWAT is causally involved in the pathogenesis of NASH.

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## INTRODUCTION

Non-alcoholic fatty liver disease (NAFLD) is a significant health problem and the most common form of chronic liver disease worldwide.<sup>1,2</sup> The prevalence of NAFLD parallels the steady increases in the rates of obesity, and consumption of saturated fat is positively associated with the risk of NAFLD.<sup>3</sup> Clinicopathologically, NAFLD comprises a wide spectrum of liver damage ranging from bland steatosis (NAFL) to non-alcoholic steatohepatitis (NASH), fibrosis and ultimately cirrhosis.<sup>4</sup> Bland steatosis is benign whereas NASH is characterized by hepatocyte injury, TNF $\alpha$ -mediated inflammation,<sup>4</sup> and a high risk of liver-related morbidity and mortality.<sup>1</sup>

The pathogenesis of NAFLD is not fully understood, and the factors that contribute to disease progression from bland steatosis to NASH remain enigmatic. Epidemiological and human observational studies do provide indications that progressive NAFLD is strongly associated with white adipose tissue (WAT) inflammation, insulin resistance and elevated circulating levels of inflammatory mediators including certain adipokines and lipids.<sup>5–10</sup> Furthermore, longitudinal rodent studies demonstrated that high-fat diet (HFD)-induced expression of inflammatory genes in WAT precedes the development of NASH in disease models with obesity,

suggesting a potential role of inflamed WAT in NAFLD progression.<sup>11</sup> Also, the severity of the NAFLD pathology appears to be closely linked to WAT dysfunction, that is, hypertrophy of adipocytes combined with macrophage infiltration, formation of crown-like structures (CLS) and enhanced expression of inflammatory genes.<sup>12</sup> Hence it has been postulated that obesity-induced inflammation in WAT is critical for the development of NAFLD,<sup>13,14</sup> but experimental evidence for an involvement of WAT is still lacking.

WAT is a complex endocrine organ that is composed of different depots, among which are the intra-abdominal (e.g., epididymal and mesenteric) and subcutaneous (e.g., inguinal) WAT depots.<sup>15,16</sup> These depots are thought to have different roles in energy storage and inflammation<sup>13,15,16</sup> and may thus have different contributions to the pathogenesis of NAFLD. The temporal development of inflammation (i.e., CLS formation) in WAT depots has not been systematically investigated and it is not known whether a particular depot is more prone to become inflamed during HFD-induced obesity and associated NAFLD.

The present time-course study analyses HFD-evoked changes in epididymal (eWAT), mesenteric (mWAT) and inguinal WAT (iWAT) with specific emphasis on adipocyte hypertrophy and WAT

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inflammation (CLS formation). To that end, a cohort of male C57BL/6J mice was treated with HFD for a period of 24 weeks. Groups of mice were killed at regular intervals, and compared with chow controls. Longitudinal histological analyses revealed that a particular depot (eWAT) is highly susceptible to develop inflammation with pronounced CLS formation after already 12 weeks. In a separate experiment, the inflamed eWAT depot of obese HFD-fed mice was surgically removed (after 12 weeks on a HFD) to examine a potential role of eWAT in the subsequent development of NASH. Our results provide evidence that inflamed eWAT has an important role in the pathogenesis of NASH. Analysis of adipokines and circulating lipids by lipidomics supports the view that circulating inflammatory factors derived from eWAT mediate NASH development.

## MATERIALS AND METHODS

### Animals and housing

Animal experiments were approved by an independent Animal Care and Use Committee and were in compliance with European Community specifications for the use of laboratory animals.

**Time-course cohort study.** Male 9-week-old wild-type C57BL/6J mice ( $n=84$ ) were obtained from Charles River Laboratories (L'Arbresle Cedex, France). After an acclimatization period of 3 weeks on chow diet (R/M-H, Ssniff Spezialdiäten GmbH, Soest, Germany; containing 33 kcal% protein, 58 kcal% carbohydrate and 9 kcal% fat), mice were matched into seven groups of  $n=12$  mice each based on body weight. One group was killed after matching to define the starting condition of the experiment ( $t=0$ ). Three groups were treated with HFD (D12451, Research Diets Inc., New Brunswick, USA; with 20 kcal% protein, 35 kcal% carbohydrate and 45 kcal% lard fat) and three control groups remained on chow. Mice had *ad libitum* access to food and water, and groups were killed after 6, 12 and 24 weeks on diet, respectively. Plasma samples were collected after 5 h fasting at 4-week intervals. Animals were killed by CO<sub>2</sub> asphyxiation, a serum sample was collected by heart puncture, and liver, eWAT, mWAT and iWAT were isolated. A part of the tissues was fixed in formalin and paraffin embedded for histological analysis; another part was snap frozen in liquid nitrogen and stored at  $-80^{\circ}\text{C}$  for real-time PCR (RT-PCR).

**Surgical removal of epididymal adipose tissue depot (eWAT).** In a separate HFD feeding experiment the contribution of eWAT to NASH development was analyzed. Male 9-week-old wild-type C57BL/6J mice (Charles River Laboratories) were acclimatized for 3 weeks and matched into two groups ( $n=15$  per group), after 12 weeks of HFD feeding based on body weight and fasting plasma insulin concentrations. All mice were injected subcutaneously with carprofen analgesic ( $5\text{ mg kg}^{-1}$ ) 30 min before surgery and anesthetized with isoflurane during surgery. In the eWATx group, both eWAT fat pads were surgically removed through a mid-ventral abdominal incision as described.<sup>17</sup> Testes were visualized and the attached epididymal fat pads were carefully removed and weighed, without damaging the testicular blood supply. In the sham group, a mid-ventral incision was made and the epididymal fat pads were visualized, that is, fat pads were pulled out, but were left intact and placed back inside the peritoneal cavity. One animal from the eWATx group died during surgery and was therefore excluded from the study. Daily food intake and body weight regain were evaluated to determine recovery from surgery. After surgery, mice continued HFD feeding for another 12 weeks and were then killed for histological evaluation of livers.

**Histological, biochemical, lipidomic and gene expression analyses**  
A detailed description of biochemical, lipidomic and gene expression analyses is provided as Supplementary 1. For histological analysis of livers, 5- $\mu\text{m}$ -thick cross-sections were stained with Hematoxylin and Eosin. NAFLD was scored blindly using a general scoring system for rodent models, which is based on the human NAS grading criteria.<sup>18</sup> Briefly, micro- and macro-vesicular steatosis were separately scored and expressed as a percentage of the cross-sectional area. Hepatic inflammation was analyzed by counting the number of inflammatory foci per field at  $\times 100$  magnification (view size  $3.1\text{ mm}^2$ ) in five different fields per specimen. For WAT, paraffin-embedded cross-sections (5  $\mu\text{m}$  thick) were stained with Hematoxylin-Phloxine-Saffron for quantification of adipocyte size and CLS

using an Olympus BX51 microscope and Cell'D software (Olympus, Zoeterwoude, the Netherlands). The number of CLS was counted in five fields ( $\times 100$  magnification) per mouse and depot, and data were expressed as number of CLS per 1000 adipocytes.

### Statistical analysis

All data are presented as mean  $\pm$  s.e.m. Significance of differences of continuous variables between HFD- and chow-fed animals was tested using Student's *t*-test. Changes over time between the different HFD groups ( $t=0, 6, 12$  and 24 weeks) were statistically analyzed by one-way analysis of variance and Tukey *post-hoc* test (normally distributed variables). Non-normally distributed variables were tested by non-parametric Kruskal-Wallis test followed by Mann-Whitney *U*-tests. Statistical significance of differences between SHAM and eWATx was tested using unpaired one-sided *t*-tests. Paired two-sided *t*-tests were used to calculate the significance of induction of inflammatory mediators in plasma (fatty acids and adipokines) between 12 and 24 weeks (i.e., before and after surgery) within each group. Results were considered statistically significant at  $P < 0.05$ . Analyses were performed using Graphpad Prism software (version 6, Graphpad Software Inc., La Jolla, USA).

## RESULTS

### Time-resolved analysis of HFD-induced obesity, hyperinsulinemia and hyperglycemia in a cohort of mice

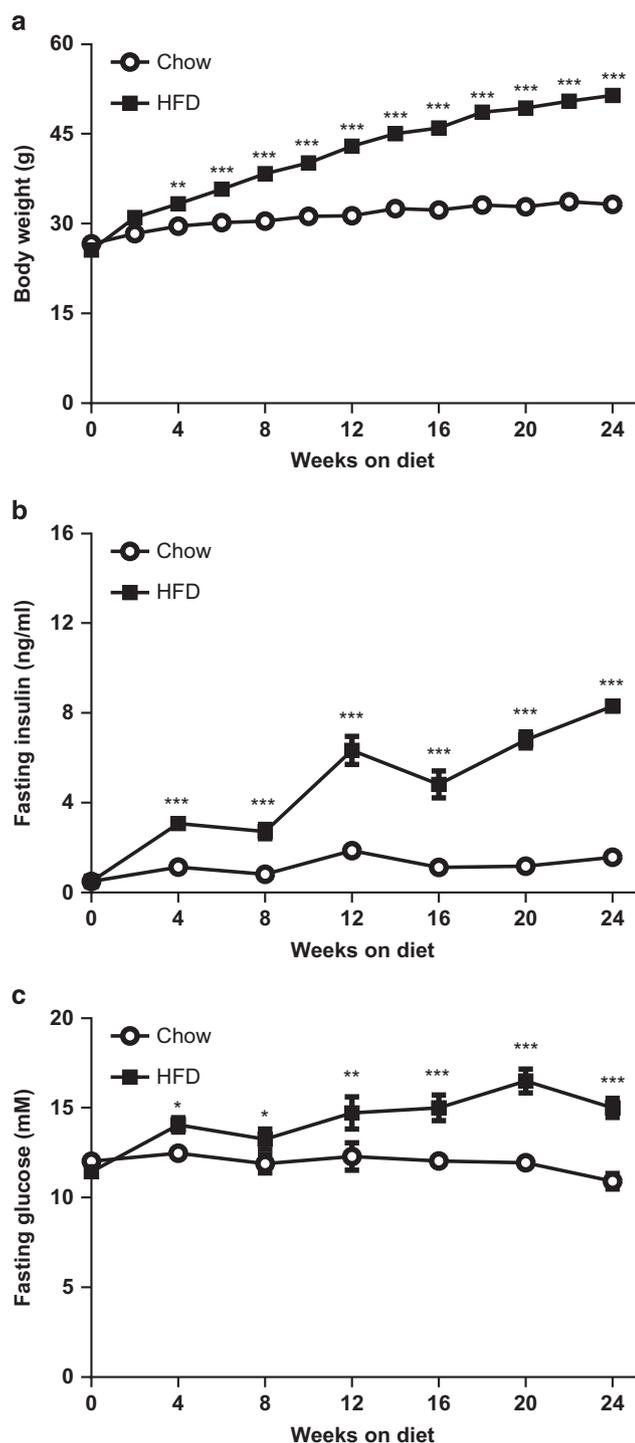
Mice had an average body weight of  $26.2 \pm 1.0\text{ g}$  at the start of the experiment ( $t=0$ ). Body weight was significantly higher in mice on a HFD compared with control mice already after 4 weeks of diet feeding and HFD-fed mice reached a final body weight of  $51.6 \pm 0.8\text{ g}$  versus  $33.3 \pm 0.7\text{ g}$  in chow-fed control mice at 24 weeks of diet feeding (Figure 1a). HFD feeding significantly increased fasting plasma insulin ( $8.2 \pm 0.2\text{ ng ml}^{-1}$ ) compared with chow ( $1.6 \pm 0.2\text{ ng ml}^{-1}$ ; Figure 1b). The HFD effect on insulin was accompanied by a significant increase in fasting plasma glucose ( $15.1 \pm 0.5\text{ mM}$ ) compared with chow ( $10.9 \pm 0.4\text{ mM}$ ; Figure 1c).

Plasma triglyceride concentrations were comparable between HFD- and chow-treated groups and decreased slightly over time (not shown). Altogether, these data show that HFD-treated mice developed obesity, hyperinsulinemia and hyperglycemia, all of which are associated with NAFLD.

### HFD feeding induces liver steatosis by week 12, which progresses to NASH

Livers collected at  $t=0, 6, 12$  and 24 weeks of diet feeding were analyzed for steatosis and presence of inflammatory cell aggregates to evaluate development of NASH (representative images shown in Figure 2a). HFD feeding resulted in modest steatosis by week 12, which intensified significantly toward the end of the study, whereas chow-fed mice showed no steatosis and normal liver histology at all the time points (not shown). Quantification of distinct forms of steatosis, that is, micro- and macrovesicular steatosis, demonstrated a gradual increase in HFD-induced microvesicular steatosis over time (Figure 2b). By contrast, macrovesicular steatosis (a hallmark of overt human NASH<sup>4</sup>) had hardly developed by week 12, but was significantly increased in week 24 (Figure 2c). HFD-induced liver steatosis can be attributed to significant increases in liver triglycerides as measured biochemically in corresponding liver homogenates (Figure 2d).

Lobular inflammation was specifically induced by HFD (Figure 2e), not by chow, and showed a similar time pattern as macrovesicular steatosis. More specifically, the number of inflammatory cell aggregates (an indicator of lobular inflammation<sup>18</sup>) remained low until week 12 and increased significantly by week 24. HFD-induced lobular inflammation was accompanied by significantly increased TNF $\alpha$  and MCP-1/Ccl2 gene expression in livers at  $t=24$  weeks (Figures 2f and g).



**Figure 1.** Time-course analysis of the effect of HFD on body weight and metabolic parameters. (a) HFD feeding increased body weight compared with chow control diet. HFD feeding gradually increased fasting plasma concentrations of (b) insulin and (c) glucose compared with chow. Data are expressed as mean  $\pm$  s.e.m. ( $n = 12$  per group per time point), \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$  versus chow control.

Taken together, these data demonstrate that 12 weeks of HFD feeding resulted in bland steatosis, which progressed to NASH by week 24 as demonstrated by the establishment of pronounced macrovesicular steatosis and lobular inflammation.

### eWAT depot is prone to develop HFD-induced inflammation

We next examined whether the eWAT, mWAT and iWAT depots would differ in their susceptibility to develop inflammation during HFD feeding and we defined the time point at which CLS formation started in the various depots. HFD feeding led to an increase in mass of eWAT, mWAT and iWAT depots over time (Figure 3a), whereas the mass of the depots remained unchanged on chow (not shown). eWAT mass increased strongly at week 6 and reached a maximum already in week 12 ( $2.4 \pm 0.1$  g). By contrast, the mass of mWAT and iWAT increased continuously over time until the end of the experiment (Figure 3a). In all WAT depots, adipocytes increased in size during HFD feeding, indicating that adipocyte expansion is a generic response of all WAT depots. However, the adipocytes of eWAT rapidly reached a maximal size in week 6 (Figure 3b). Adipocytes in the other depots were still smaller at this time point and their size increased more slowly and gradually until the end of the study. Notably, in week 6 first CLS were observed in eWAT specifically and their numbers increased greatly in week 12 when the maximal capacity of eWAT seemed to be reached (viz. maximal mass and maximal adipocyte size; Figure 3c). CLS formation in eWAT was more rapid and pronounced than in mWAT and iWAT (Figures 3c and d), showing that eWAT is most prone to develop HFD-induced inflammation. In mWAT, CLS numbers increased later than in eWAT (by week 24) and when maximal average adipocyte size was reached, essentially as observed in eWAT.

These observations show that the expandability of a depot (and its adipocytes) is limited and that this depot-specific restriction seems critical for the development of inflammation.

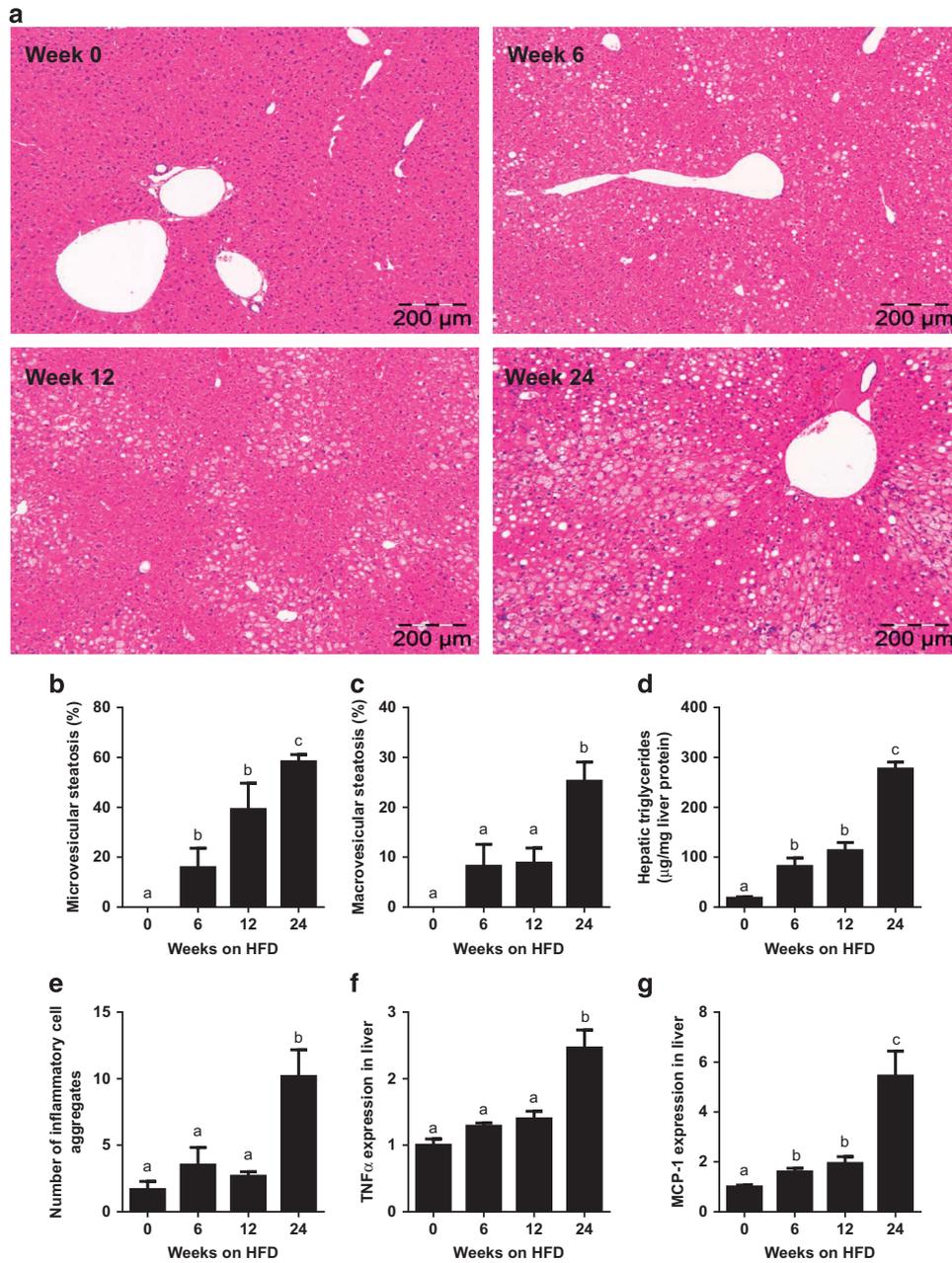
In all, our time-resolved histological analyses show that HFD-induced WAT inflammation starts in a specific depot (eWAT) and increases strongly when eWAT has expanded maximally (at  $t = 12$  weeks). Importantly, eWAT inflammation coincides with bland steatosis and hence precedes the development of NASH.

### Surgical removal of inflamed eWAT attenuates NASH development

To examine whether eWAT is causally involved in the progression of liver steatosis to NASH, we performed a separate HFD feeding experiment in which eWAT was surgically removed in one group (eWATx) and compared with a SHAM surgery control group (SHAM). The surgery was performed after week 12 of HFD feeding, that is, the time point at which livers in the above time-course experiment were steatotic and eWAT was inflamed.

Body weight at the time of surgery was comparable between SHAM and eWATx groups (SHAM:  $42.3 \pm 1.2$  g, and eWATx:  $42.3 \pm 0.9$  g; Figure 4a). On average  $1.9 \pm 0.1$  g of eWAT was removed and this reduction in fat mass was reflected in the body weight of eWATx mice the day after surgery (SHAM:  $41.6 \pm 1.7$  g and eWATx:  $39.5 \pm 1.2$  g, not shown). CLS were abundantly present in this tissue, confirming pronounced inflammation at the time of surgery (Figure 4b). Food intake was comparable between the eWATx and SHAM group throughout the study. The total body weight at the end of the experiment was  $47.1 \pm 1.3$  g in SHAM and  $46.7 \pm 0.9$  g in eWATx, and whole-body fat mass determined by EchoMRI LLC (Houston, TX, USA) was slightly lower in eWATx mice (Figure 4c, not significant), whereas lean mass was comparable between the groups (SHAM:  $29.1 \pm 0.6$  g vs eWATx:  $29.7 \pm 0.4$  g, not significant). Fasting plasma glucose concentrations increased during the experiment, essentially as observed in the time-course study, and hyperglycemia was comparable in both groups (SHAM:  $14.6 \pm 0.7$  mM vs eWATx:  $15.2 \pm 0.6$  mM; not significant).

Isolation of individual fat depots after sacrifice in week 24 showed that eWAT mass was significantly reduced in eWATx mice (Figure 4d). The mass of mWAT, iWAT (Figure 4d), retroperitoneal WAT and brown adipose tissue (not shown) was comparable between both groups, indicating that these depots did not compensate for the removed eWAT. Also, the



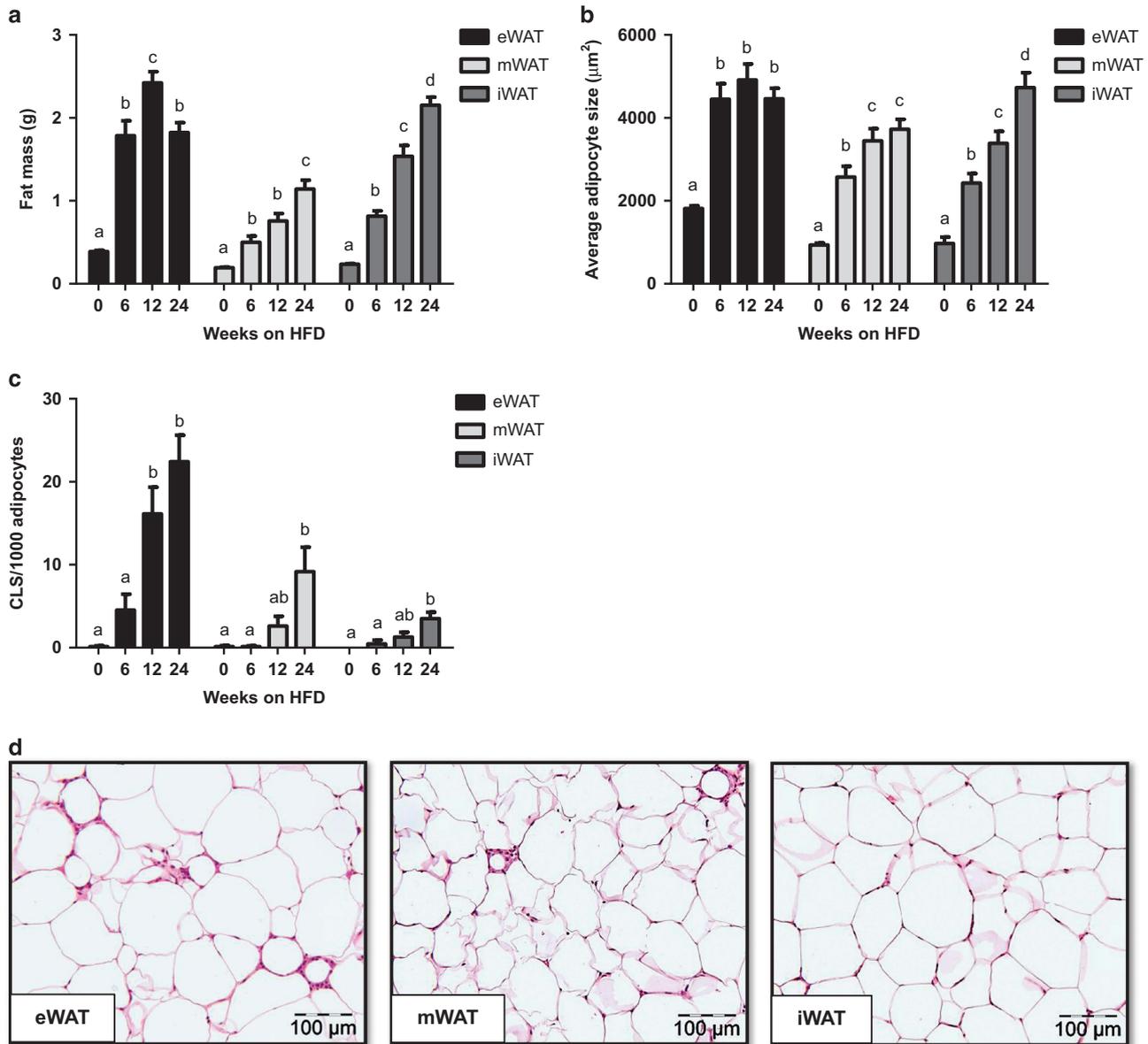
**Figure 2.** Time-resolved development of NAFLD induced by HFD feeding. (a) Representative photomicrographs of Hematoxylin and Eosin-stained liver cross-sections of mice treated with HFD for 0, 6, 12 or 24 weeks (magnification  $\times 100$ ). Histological analysis of (b) microvesicular and (c) macrovesicular steatosis as percentage of the cross-sectional area ( $n=6-12$  per group per time point). (d) Biochemical quantification of hepatic triglyceride content ( $n=11-12$  per group). (e) Development of lobular inflammation in liver over time defined as the number of inflammatory cell aggregates ( $n=6-12$  per group per time point). (f) Gene expression of TNF $\alpha$  and (g) MCP-1 in liver over time. Data ( $n=8$  per group) are expressed as fold change in gene expression relative to  $t=0$ . Data are expressed as mean  $\pm$  s.e.m. <sup>a,b,c</sup>Mean values with unlike letters differ significantly from each other ( $P < 0.05$ ).

weight of heart and kidneys was comparable between the groups (not shown).

Histological analysis of livers revealed that eWATx mice exhibited a similar degree of micro- and macro-vesicular steatosis compared with SHAM mice (Figures 5a and c). Biochemical analysis of liver triglycerides in liver homogenates showed no significant difference between the groups (SHAM:  $160.8 \pm 18.9 \mu\text{g mg}^{-1}$  liver protein vs eWATx:  $170.5 \pm 16.2 \mu\text{g mg}^{-1}$  liver protein; not significant). Minor liver lipids (cholesteryl esters and free cholesterol) were also comparable between the groups (data not shown). Remarkably, eWATx livers displayed a significantly reduced number

of inflammatory cell aggregates, indicating attenuated NASH development upon eWAT removal (Figure 5d). Reduced liver inflammation was substantiated by a significantly decreased gene expression of TNF $\alpha$  (Figure 5e) and MCP-1 (trend  $P=0.08$ ; Figure 5f) in eWATx livers.

Collectively, these results show that surgical removal of inflamed eWAT significantly attenuates the development of NASH in the absence of an effect on hyperglycemia and demonstrates a causal role of eWAT in the pathogenesis of NAFLD. We next analyzed circulating factors that could mediate this effect on liver.

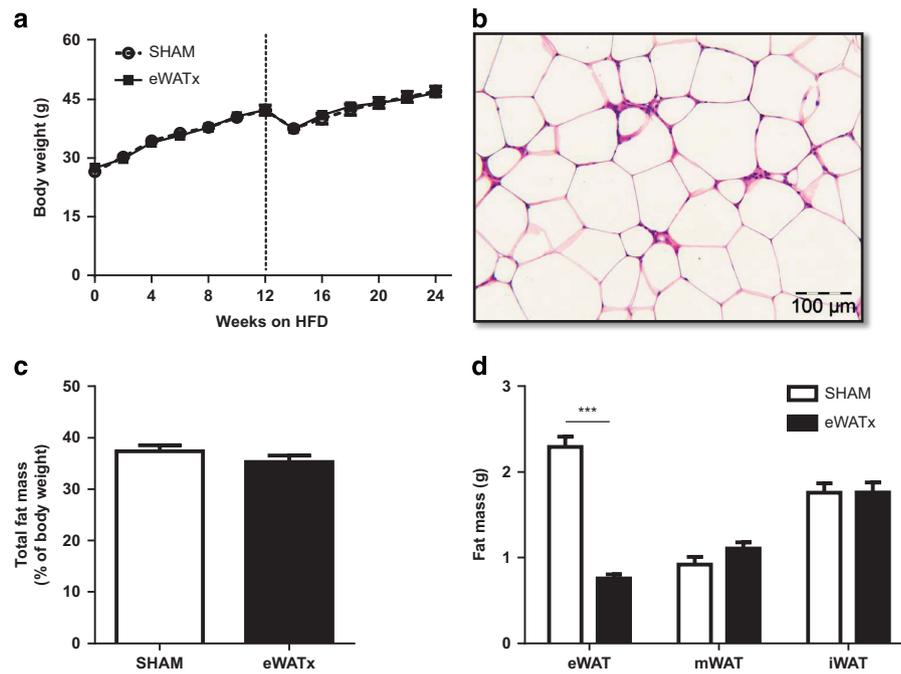


**Figure 3.** Effect of HFD feeding on the quantity and inflammatory state of the eWAT, mWAT and iWAT depot during HFD feeding time-course experiment ( $n = 12$  per group per time point). **(a)** WAT mass of eWAT, mWAT and iWAT depot during HFD feeding time-course experiment ( $n = 12$  per group per time point). **(b)** Development of adipocyte cell size of the different WAT depots quantified by morphometric analysis of Hematoxylin-Phloxine-Saffron (HPS)-stained sections. **(c)** Quantitative analysis of the number of crown-like structures (CLS) in the different WAT depots over time ( $n = 8-12$  per group per time point). **(d)** Representative images of HPS-stained cross-sections of eWAT, mWAT and iWAT after 24 weeks of HFD (magnification  $\times 200$ ). Data are expressed as mean  $\pm$  s.e.m. <sup>a,b,c,d</sup>Mean values with unlike letters differ significantly from each other ( $P < 0.05$ ).

#### Surgical removal of eWAT affects circulating levels of proinflammatory mediators

The effects of eWAT removal on adipokines and lipids associated with NAFLD development were assessed. In eWATx, the IL-6 serum concentrations were slightly lower ( $5.5 \pm 2.6$  pg ml<sup>-1</sup>) than in SHAM ( $7.0 \pm 2.5$  pg ml<sup>-1</sup>; not shown) but the difference was not statistically significant. The eWATx and SHAM groups also had comparable plasma concentrations of MCP-1 ( $86.6 \pm 11.7$  pg ml<sup>-1</sup> vs  $80.8 \pm 13.3$  pg ml<sup>-1</sup>; not significant) and adiponectin ( $13.6 \pm 1.1$  µg ml<sup>-1</sup> vs  $14.3 \pm 1.2$  µg ml<sup>-1</sup>; not significant). By contrast, plasma leptin concentrations increased significantly in SHAM mice and this increase was not observed in eWATx mice (Figure 6a). Plasma leptin levels also correlated positively ( $r^2 = 0.7$ ;  $P = 0.01$ ) with hepatic MCP-1 expression, suggesting a link to hepatic inflammation.

We next profiled the circulating (proinflammatory) lipids using lipidomics to define the most abundant lipid species in plasma (Figure 6b), and subsequently analyzed whether eWATx removal affected these lipids (Table 1). Saturated fatty acids (SFA) were the most abundant lipid species after 12 weeks of HFD feeding (before surgery), followed by monounsaturated fatty acids (MUFA), and n-6 and n-3 polyunsaturated fatty acids (PUFA; Figure 6b). Within the SFA, C16:0 (palmitic acid), C18:0 (stearic acid) and C14:0 (myristic acid) were most abundant. Within the class of MUFA C18:1n9 (oleic acid) and C16:1n7 (palmitoleic acid) were circulating at high levels, and within the PUFA, C20:4n3 (eicosatetraenoic acid) and C18:2n6 (linoleic acid) were most abundant. Table 1 shows that the level of total SFA, and palmitic acid in particular ( $+1.4$ ;  $P < 0.05$ ), increases in SHAM, whereas such an increase was not observed in eWATx. Myristic acid and several



**Figure 4.** Effect of surgical removal of eWAT on body weight and other WAT depots. Mice were fed a HFD and, on average, 1.9 g of eWAT was carefully removed after 12 weeks of HFD. **(a)** Body weight development of the eWATx and SHAM group. The dashed line indicates the time point of surgery. **(b)** Representative image of a Hematoxylin-Phloxine-Saffron-stained eWAT cross-section showing presence of inflammatory cells and CLS at time of removal (magnification  $\times 200$ ). **(c)** Analysis of total fat mass measured by EchoMRI LLC in week 24 of HFD. **(d)** Mass of eWAT, mWAT and iWAT isolated at the end of the study (24 weeks of HFD) in eWATx and SHAM group. The eWAT mass of eWATx mice was significantly lower than in SHAM. Data also show that mWAT and iWAT did not compensate for the removed eWAT. Data are expressed as mean  $\pm$  s.e.m. ( $n = 14$ –15 per group), \*\*\* $P < 0.001$  versus SHAM.

of the less abundant SFA (C20:0, C22:0 and C24:0) decreased significantly after eWAT removal. Furthermore, total MUFA increased strongly over time in SHAM (+3.7;  $P < 0.05$ ), mainly due to significant rises in palmitoleic, oleic and eicosenoic acids. By contrast, there was no significant change over time in any of the MUFA in the eWATx group. Total PUFA levels increased in both, SHAM (+2.0) and eWATx (+1.5). The observed changes in SFA and MUFA support the notion that removal of eWAT prevents the development of a proinflammatory state.

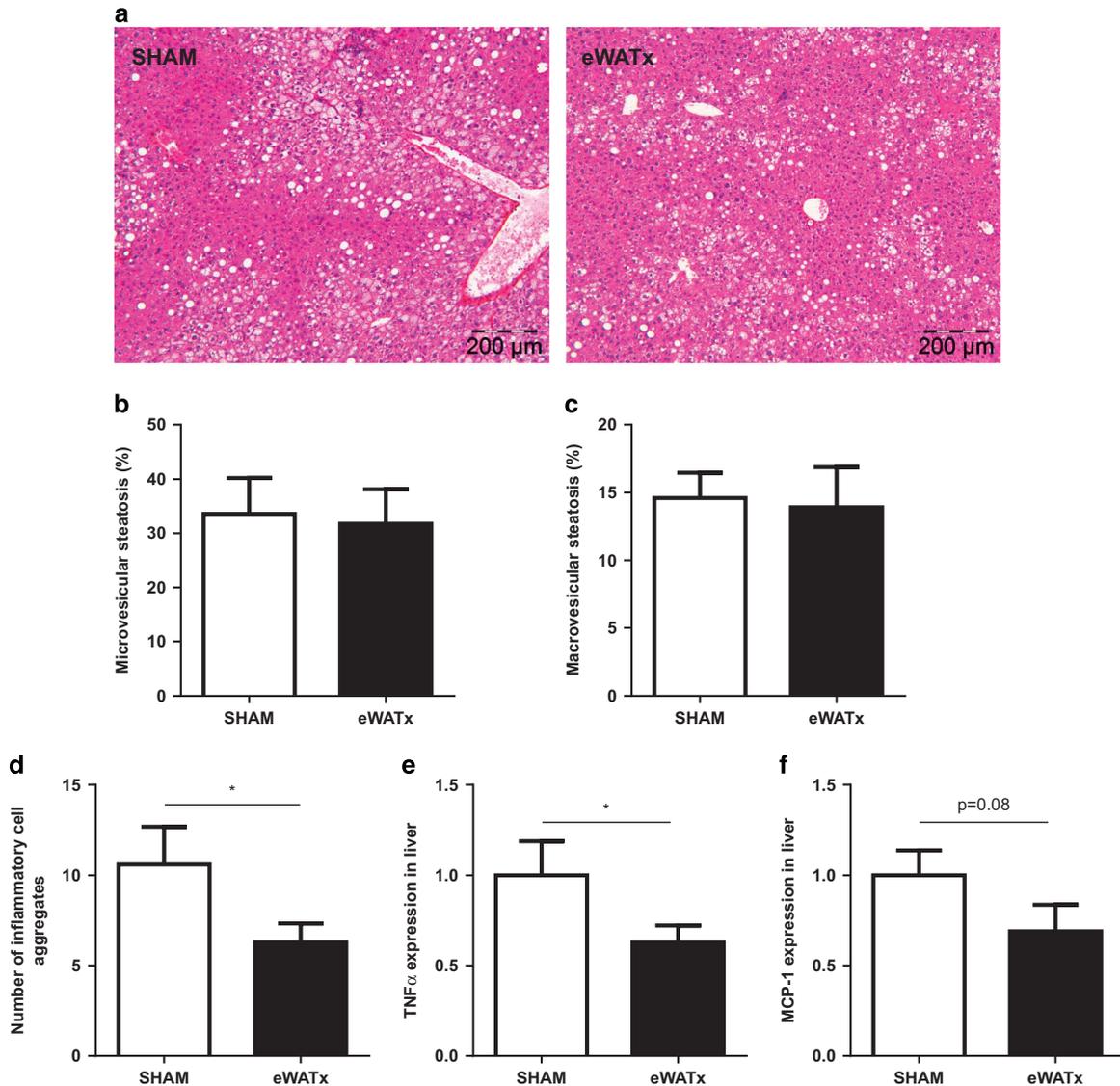
## DISCUSSION

NAFLD is strongly associated with obesity, but the pathogenesis of the disease, and in particular the role of WAT, is poorly understood. It has been proposed that inflammation in WAT may have a critical role in obesity-induced NAFLD development,<sup>4,13,19,20</sup> but evidence for causality is lacking. This study shows that HFD-induced inflammation in eWAT develops more rapidly than in mWAT or iWAT, and that this inflammation precedes overt NASH. Notably, pronounced CLS formation was observed in eWAT once the adipocytes of this tissue did not further increase in size and the depot had reached a maximal mass (week 12 of HFD feeding). A subsequent experiment showed that removal of the inflamed eWAT depot at week 12 attenuates liver inflammation and reduces the development of NASH. Removal of eWAT affected the circulating levels of specific proinflammatory mediators, among which are leptin and specific lipids (e.g., palmitic acid), providing a rationale for the observed hepatoprotective effect.

The time-course analysis of HFD-induced NAFLD shows that a particular intra-abdominal depot in mice, eWAT, is prone to develop tissue inflammation characterized by presence of macrophages and CLS. Consistent with this finding, other groups

have reported that eWAT of obese HFD-treated mice exhibits a higher number of CLS than mesenteric and subcutaneous (inguinal) WAT depots.<sup>21,22</sup> Differences in macrophage content appear to exist already in lean C57BL6 mice: Altinas *et al.*<sup>21</sup> showed that the subcutaneous depot differs from the intra-abdominal depots with respect to immune cell composition and density. For instance, the density of solitary adipose tissue macrophages in subcutaneous WAT is much lower than in intra-abdominal depots, such as eWAT.<sup>21</sup> Hence, the relatively high number of solitary adipose tissue macrophages in eWAT may predispose this depot to develop CLS more rapidly in response to HFD than other depots analyzed in this study. In obese subjects, CLS are also more prevalent in abdominal (omental) WAT than in subcutaneous WAT<sup>6,23</sup> suggesting that in humans, intra-abdominal depots are also more prone to become inflamed than subcutaneous depots, and that our observations are not restricted to mice.

We found that the number of CLS in eWAT increased strongly once this depot had reached a maximal mass and concomitantly no further increase in adipocyte size was observed. In line with this, other groups reported that the weight of eWAT does typically not exceed  $\sim 2.5$  g.<sup>12,22,24,25</sup> This limitation in eWAT mass has been observed with different diets and in different strains of mice, pointing to a generic threshold of eWAT independent of the experimental conditions employed. Consistent with this, Virtue and Vidal-Puig<sup>26</sup> proposed that organisms possess a maximum capacity for adipose expansion, and failure in the capacity for adipose tissue expansion, rather than obesity *per se*, may underlie the development of inflammation. Indeed, also in the case of mWAT, CLS numbers increased at week 24, that is, after the average adipocyte size had reached a maximum. Little is known about the mediators that control WAT expansion during diet-induced obesity. It is possible that localized cytokine production limits further WAT expansion: Salles *et al.*<sup>27</sup> showed that TNF $\alpha$



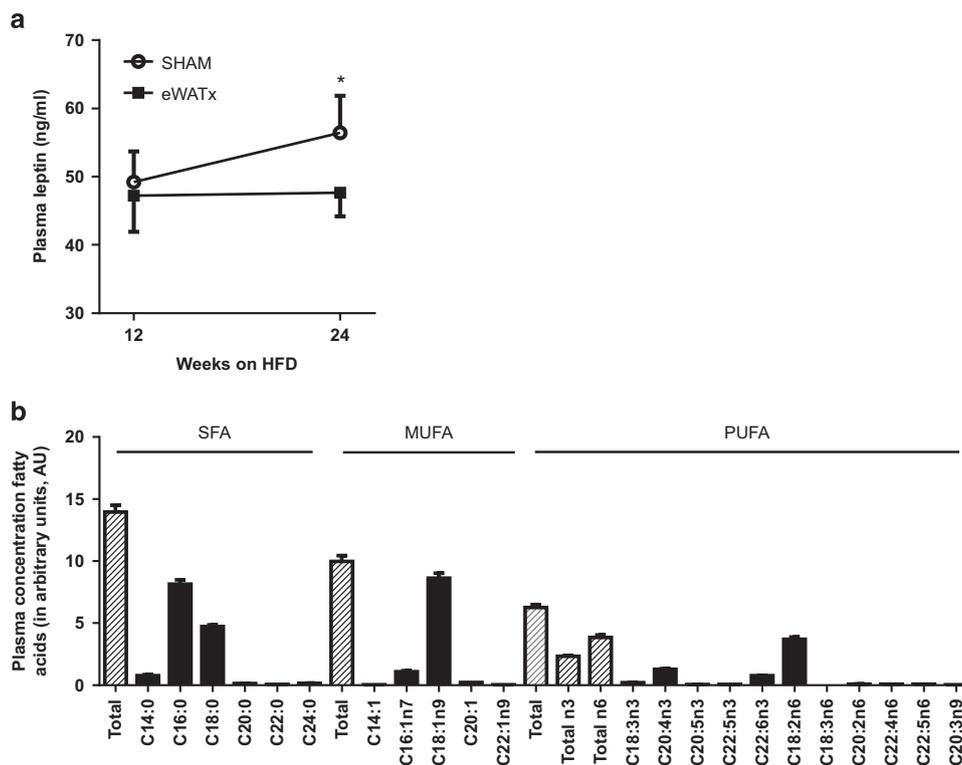
**Figure 5.** Effect of surgical removal of eWAT on NAFLD development. **(a)** Representative images of Hematoxylin and Eosin-stained liver sections (magnification  $\times 100$ ). **(b)** Quantification of microvesicular steatosis and **(c)** macrovesicular steatosis as percentage of the cross-sectional liver area ( $n=14-15$  per group). **(d)** Number of inflammatory cell aggregates in livers of eWATx and SHAM mice. Liver gene expression of **(e)** TNF $\alpha$  and **(f)** MCP-1 in eWATx and SHAM. Real-time PCR data are expressed as fold change in gene expression relative to SHAM ( $n=8$  per group). Data are expressed as mean  $\pm$  s.e.m., \* $P < 0.05$ .

knockout mice have twofold more eWAT mass than wild-type mice during HFD feeding. In support of this notion, increased TNF $\alpha$  gene expression in eWAT was observed after attainment of maximal adipocyte size in an experiment conducted under conditions comparable to those applied herein.<sup>22</sup> Together, our time-resolved analysis of the inflammatory component in diet-induced obesity shows that WAT inflammation develops sequentially across depots.

C57BL/6 mice constitute a frequently used model to study diet-induced obesity and associated comorbidities. For the interpretation of these studies, it is important to recognize that development of inflammation upon HFD feeding is a very complex and dynamic process involving multiple tissues, including WAT and liver.<sup>11,15</sup> As demonstrated herein, the different WAT depots become inflamed at specific time points during HFD feeding, rather than simultaneously. Because animal studies often analyze a single-WAT depot (frequently the eWAT) at one

particular time point during HFD feeding, conclusions about the condition 'adipose tissue in general' or the inflammatory state in other depots should be made with caution. Our results support a more comprehensive analysis of WAT (with precise specification of the intra-abdominal depots analyzed), and advocate the study of the crosstalk between organs in NAFLD. For instance, between 12 and 24 weeks of HFD, that is, the period in which eWAT was inflamed and did not further expand, we observed a pronounced increase of triglyceride concentrations in the liver. This supports the concept that, once the expansion limit of a particular WAT depot has been reached, adipose tissue ceases to store energy efficiently and lipids begin to accumulate as ectopic fat in other tissues.<sup>26</sup>

In patients, the accumulation of intra-abdominal WAT is strongly associated with progressive NASH.<sup>5</sup> Hepatic inflammation and fibrosis augmented incrementally with increases in intra-abdominal fat mass. Importantly, intra-abdominal fat of patients



**Figure 6.** Effect of surgical removal of eWAT on circulating inflammatory mediators and lipids. **(a)** Plasma concentrations of leptin before surgery at 12 weeks of HFD and at the end of the experiment (24 weeks) in eWATx and SHAM groups. Data are expressed as mean  $\pm$  s.e.m., \* $P < 0.05$  according to paired Student's *t*-test. **(b)** Profiling of plasma lipids after 12 weeks of HFD by lipidomic analysis. Fasting plasma was collected before surgery. The levels of saturated free fatty acids (SFA), monounsaturated fatty acids (MUFA) and polyunsaturated fatty acids (PUFA) are shown and the most abundant lipid species of each category are indicated. Data are expressed as mean  $\pm$  s.e.m. and as arbitrary units (AU) relative to internal standard.

was directly associated with liver inflammation and fibrosis independent of insulin resistance and hepatic steatosis.<sup>5,7</sup> Consistent with this, removal of eWAT in the present study attenuated NASH development without an effect on hyperinsulinemia, hyperglycemia and liver steatosis. A possible explanation for the strong association between abdominal WAT mass and NASH severity in the liver may lie in the anatomical distance between both tissues. Inflammatory mediators from the intra-abdominal depots can reach the liver relatively easily (venous drainage via the portal vein).<sup>28</sup> By contrast, associations between systemically drained adipose tissue depots (e.g., the deep layer of subcutaneous WAT in the abdominal area) and NASH are rare,<sup>29</sup> suggesting that these depots only have a minor role in the pathogenesis of NASH.

Although the above studies mainly focused on the quantity of adipose tissue, increasing evidence also points to a role of its inflammatory state. Livers of obese subjects with inflamed intra-abdominal (omental) WAT contain more fibro-inflammatory lesions than livers of equally obese subjects without WAT inflammation.<sup>6,7</sup> This observation suggests that inflammation in a specific WAT depot contributes to the inflammatory component in human NASH. The present study supports this view, because surgical removal of inflamed eWAT reduces liver inflammation (~40% less inflammatory aggregates). Of note, intra-abdominal eWAT in mice has no human equivalent and our findings should not be generalized with respect to the role of other WAT depots. Because of the close relationship between visceral obesity and NASH development in patients,<sup>5</sup> it is possible that inflamed visceral WAT depots, such as mWAT, may contribute to NASH in a similar way as eWAT. For instance, mWAT develops similar features of inflammation as observed in eWAT, including formation of CLS during WAT expansion and expression of proinflammatory

mediators (e.g., cytokines, adipokines and fatty acids), in both mice<sup>30,31</sup> and humans.<sup>32,33</sup> Furthermore, proinflammatory mediators that are released by mWAT can reach the liver not only via systemic drainage (such as eWAT) but also via the portal vein, which constitutes a more direct connection to the liver.<sup>28</sup> However, additional studies are needed to investigate the contribution of inflamed mWAT to NASH development.

Specific circulating factors have been proposed as inducers of liver inflammation in NASH.<sup>19,20</sup> Among these mediators are cytokines/adipokines (including IL-6, TNF $\alpha$ , leptin and adiponectin) and specific lipids with reported activities on liver cells.<sup>4,13,14,20</sup> Of the adipokines measured in the present study, only plasma leptin differed between the eWATx and the SHAM group. Several studies have shown that leptin exerts proinflammatory and profibrogenic effects on liver cells.<sup>34,35</sup> For instance, leptin stimulates hepatic stellate cells to express the proinflammatory cytokine MCP-1,<sup>35</sup> a chemotactic factor and critical mediator of lobular inflammation.<sup>36</sup> In line with this, we observed that plasma leptin concentrations correlated with MCP-1 gene expression in the liver. Besides cytokines/adipokines, certain lipid mediators (e.g., SFA with TLR4 binding properties) have also been implicated in the pathogenesis of NASH. For example, *in vitro* studies have shown that SFA, and in particular palmitic acid, can trigger inflammation via the NF- $\kappa$ B pathway and thereby induce TNF $\alpha$  production.<sup>37</sup> In SHAM mice, palmitic acid levels increased significantly between 12 and 24 weeks of HFD feeding, that is, during the progression from NAFL to NASH. This increase was not observed in eWATx mice and, in line with this, hepatic TNF $\alpha$  expression was lower than in SHAM. A total plasma lipid analysis in humans showed that obese subjects with NAFL and NASH have significantly elevated MUFA levels when compared with lean controls.<sup>38</sup> Among these MUFA were palmitoleic acid and oleic

**Table 1.** Change in plasma fatty acid levels in SHAM and eWATx

Fatty acids	SHAM Δ change	eWATx Δ change
<b>Saturated fatty acid (SFA)</b>		
Myristic acid (C14:0)	1.3 ± 0.8	-0.3 ± 1.0
Palmitic acid (C16:0)	0.02 ± 0.1	-0.3 ± 0.2
Stearic acid (C18:0)	1.4 ± 0.5*	-0.1 ± 0.6
Arachidic acid (C20:0)	-0.03 ± 0.3	0.4 ± 0.2
Behenic acid (C22:0)	-0.01 ± 0.0	-0.05 ± 0.0*
Lignoceric acid (C24:0)	-0.01 ± 0.0	-0.04 ± 0.0*
Lignoceric acid (C24:0)	-0.02 ± 0.0	-0.1 ± 0.1*
<b>Monounsaturated acid fatty acid (MUFA)</b>		
Myristoleic acid (C14:1)	3.7 ± 0.9**	0.8 ± 0.8
Palmitoleic acid (C16:1n7)	0.0 ± 0.0	0.0 ± 0.0
Oleic acid (C18:1n9)	0.5 ± 0.2*	0.2 ± 0.1
Eicosenoic acid (C20:1)	3.1 ± 0.8**	0.6 ± 0.7
Erucic acid (C22:1n9)	0.1 ± 0.0*	0.03 ± 0.0
Erucic acid (C22:1n9)	0.004 ± 0.0	-0.003 ± 0.0
<b>Polyunsaturated fatty acid (PUFA)</b>		
Total n6-fatty acids	2.0 ± 0.6*	1.5 ± 0.4*
Total n3-fatty acids	1.7 ± 0.5*	1.0 ± 0.3*
Linoleic acid (C18:2n6)	0.3 ± 0.1*	0.5 ± 0.1*
γ-linolenic acid (C18:3n6)	1.7 ± 0.5*	1.0 ± 0.3*
Eicosadienoic acid (C20:2n6)	0.0 ± 0.0	0.0 ± 0.0
Adrenic acid (C22:4n6)	0.03 ± 0.0*	0.02 ± 0.0*
Docosapentaenoic acid (C22:5n6)	0.02 ± 0.0*	0.03 ± 0.0*
α-linolenic acid (C18:3n3)	0.02 ± 0.0*	0.01 ± 0.0
Eicosatetraenoic acid (C20:4n3)	0.1 ± 0.0	0.2 ± 0.0
Eicosapentaenoic acid (C20:5n3)	0.1 ± 0.1	0.3 ± 0.1*
Docosapentaenoic acid (C22:5n3)	0.01 ± 0.0	0.01 ± 0.0*
Docosapentaenoic acid (C22:5n3)	0.01 ± 0.0*	0.02 ± 0.0*
Docosahexaenoic acid (C22:6n3)	0.01 ± 0.0*	0.1 ± 0.0*
Mead acid (C20:3n9)	0.1 ± 0.0*	0.1 ± 0.0*

Delta (Δ) change in plasma lipids between week 12 and 24, that is, before and after surgery. Data are in arbitrary units (mean ± s.e.m.). \* $P \leq 0.05$  indicates significant changes over time within a group. \*\* $P < 0.05$  indicates significant difference between SHAM ( $n = 11$ ) and eWATx ( $n = 12$ ).

acid, which also increased after surgery in SHAM, whereas they did not change significantly in eWATx. The observed increases in MUFA (both in humans and mice) may be an adaptive response to protect the liver from the lipotoxic effects of SFA (i.e., palmitic acid). As MUFAs themselves can suppress liver inflammation in mice,<sup>39</sup> it is thus likely that increased levels of palmitic acid in SHAM mice are critical for the development of liver inflammation.

Collectively, this study demonstrates that obesity-induced inflammation develops progressively across various WAT depots, starting in eWAT. Surgical excision of inflamed eWAT shows that this depot participates in the development of NASH. Hence, interventions that target WAT may have significant therapeutic benefit for the treatment of NASH in the context of obesity.

## CONFLICT OF INTEREST

The authors declare no conflict of interest.

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