



Original article

Metabolic adaptations in severe obesity: Insights from circulating oxylipins before and after weight loss



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SUMMARY

Background: The relationship between lipid mediators and severe obesity remains unclear. Our study investigates the impact of severe obesity on plasma concentrations of oxylipins and fatty acids and explores the consequences of weight loss.

Methods: In the clinical trial identifier NCT05554224 study, 116 patients with severe obesity and 63 overweight/obese healthy controls matched for age and sex ($\approx 2:1$) provided plasma. To assess the effect of surgically induced weight loss, we requested paired plasma samples from 44 patients undergoing laparoscopic sleeve gastrectomy one year after the procedure. Oxylipins were measured using ultra-high-pressure liquid chromatography coupled to a triple quadrupole mass spectrometer via semi-targeted lipidomics. Cytokines and markers of interorgan crosstalk were measured using enzyme-linked immunosorbent assays.

Results: We observed significantly elevated levels of circulating fatty acids and oxylipins in patients with severe obesity compared to their metabolically healthier overweight/obese counterparts. Our findings indicated that sex and liver disease were not confounding factors, but we observed weak correlations in plasma with circulating adipokines, suggesting the influence of adipose tissue. Importantly, while weight loss restored the balance in circulating fatty acids, it did not fully normalize the oxylipin profile. Before surgery, oxylipins derived from lipoxygenase activity, such as 12-HETE, 11-HDoHE, 14-HDoHE, and 12-HEPE, were predominant. However, one year following laparoscopic sleeve gastrectomy, we observed a complex shift in the oxylipin profile, favoring species from the cyclooxygenase pathway, particularly proinflammatory prostanoids like TXB₂, PGE₂, PGD₂, and 12-HHTrE. This transformation appears to be linked to a reduction in adiposity, underscoring the role of lipid turnover in the development of metabolic disorders associated with severe obesity.

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Conclusions: Despite the reduction in fatty acid levels associated with weight loss, the oxylipin profile shifts towards a predominance of more proinflammatory species. These observations underscore the significance of seeking mechanistic approaches to address severe obesity and emphasize the importance of closely monitoring the metabolic adaptations after weight loss.

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Abbreviations			
AA	arachidonic acid	HODE	hydroxyoctadecadienoic acid
ACN	acetonitrile	HOMA-IR	homeostatic model assessment-insulin resistance
ALA	alpha-linolenic acid	HOTrE	hydroxyoctadecatrienoic acid
BHT	butylated hydroxytoluene	IL-10	interleukin-10
BMI	body mass index	LA	linoleic acid
BS	bariatric surgery	LDA	linear discriminant analysis
CCL2	C-C Motif Chemokine Ligand 2	LOX	lipoxygenase
COX	cyclooxygenase	LSG	laparoscopic sleeve gastrectomy
CYP450	cytochrome p450	MeOH	methanol
DHA	docosahexaenoic acid	MUFA	monounsaturated fatty acid
DiHDPE	dihydroxydocosapentaenoic acid	NAFLD	non-alcoholic fatty liver disease
DiHETE	dihydroxyeicosatetraenoic acid	NEFAs	non-esterified fatty acids
DiHETrE	dihydroxyeicosatrienoic acid	oxoODE	oxooctadecadienoic acid
DiHODE	dihydroxyoctadecadienoic acid	PGD	prostaglandin D
DiHOME	dihydroxyoctadecenoic acid	PGE	prostaglandin E
EDTA	ethylenediaminetetraacetic acid	PGF	prostaglandin F
ELISA	enzyme-linked immunoassay	PGH	prostaglandin H
EPA	eicosapentaenoic acid	HPLA2	phospholipase A2
EpODE	epoxyoctadecadienoic acid	PLSDA	partial least square discriminant analysis
EpOME	epoxyoctadecenoic acid	PUFA	polyunsaturated fatty acid
FGF	fibroblast growth factor	QqQ/MS	triple quadrupole mass spectrometry
HDoHE	hydroxydocosahexaenoic acid	SFA	saturated fatty acid
HEPE	hydroxyeicosapentaenoic acid	T2DM	type 2 diabetes mellitus
HETE	hydroxyeicosatetraenoic acid	TriHOME	trihydroxyoctadecenoic acid
HHTrE	hydroxyheptadecatrienoate	TNF- α	tumor necrosis factor- α
		TXB2	thromboxane B2
		UHPLC	ultra-high pressure liquid chromatography

1. Introduction

A prominent epidemiological challenge in contemporary times is the escalating incidence of obesity, attaining epidemic proportions [1]. Individuals with obesity share a common metabolic foundation, and their clinical status is frequently influenced by environmental factors like an unbalanced dietary regimen or a sedentary way of life [2]. Obesity is intricately linked with the manifestation of chronic diseases, encompassing type 2 diabetes mellitus, nonalcoholic fatty liver disease, cardiovascular disorders, and metabolic syndrome [3]. Several observations support the hypothesis that lipid mediators interact with obesity and influence weight changes [4–8]. Excess body fat can lead to tissue damage, inflammation, and disruptions in lipid metabolism, with higher body mass index (BMI, Kg/m²) exacerbating these complications.

Individuals with severe obesity, characterized by a BMI over 40 kg/m², who underwent weight loss surgery provide a unique perspective to investigate these phenomena. Bariatric surgery is a widely used surgical procedure for the treatment of morbid obesity and its associated comorbidities. Patients who benefit from this procedure not only reduce weight but also improve insulin resistance, hepatic alterations, and hypertension.

Lipids are versatile molecules that regulate oxidative and inflammatory stress [9]. Among them, oxylipins are bioactive oxygenated metabolites of fatty acids with similar biochemical

transport and are established as essential mediators in inflammatory diseases. Animal studies have shown that oxylipins can contribute to metabolic dysfunction in obesity [10–13]. The analytical limitations and the complex enzymatic and non-enzymatic pathways involved [14] complicate the examination of human oxylipins. To avoid limitations, we propose focusing on a specific subset of plasma oxylipins that are consistently detected, easily quantifiable, and resistant to freezing from the total pool of oxylipins in human blood. By taking this approach, we can reliably assess human oxylipin levels considering their potential source from specific reactions on polyunsaturated fatty acids (PUFAs) catalyzed by cyclooxygenases (COX), lipoxygenases (LOX), and cytochromes P450 (CYP450) [10,15–17].

In this study, we extensively investigated the relationship between plasma oxylipins and various factors that could influence our results. Severe obesity often leads to the accumulation of lipids in alternative tissues, especially the liver, triggering metabolic stress and the abnormal secretion of hormones known as organokines [18–20]. Although the link between oxylipins and organokines has yet to be fully comprehended, animal studies suggest their involvement in inflammation, insulin resistance, and energy balance [21,22]. To scrutinize these effects in humans, we measured specific hormones in the blood, including adiponectin, leptin, fibroblast growth factor (FGF)-19, and FGF-21 [23–26], as well as irisin and galectin-3, both associated with inflammation and fibrosis [27,28].

There exists limited understanding regarding the plasma oxylipin profile in morbid obesity, how this profile relates to pertinent metabolic biomarkers, and how this is changed after patients have undergone bariatric surgery. Consequently, this investigation aimed to delineate the plasma oxylipin profile in individuals afflicted with severe obesity, discern any significant associations between this profile and metabolic markers and comorbidities, and ascertain the impact of weight reduction and metabolic readjustment occurring 12 months post-bariatric surgery.

2. Materials and methods

2.1. Human subjects, study design, and ethical considerations

We conducted a pilot investigation involving individuals afflicted by severe obesity at the Bariatric Surgery Unit of Hospital Universitari Sant Joan, Reus.

Our approach for sample size determination ($n = 116$) and study design was guided by prior research findings [29,30] and adhered to established protocols. Among eligible participants, either Roux-en-Y gastric bypass ($n = 59$) or laparoscopic sleeve gastrectomy (LSG) ($n = 57$) were independently indicated. We did not find differences between both groups, which were combined. By utilizing the results of nonalcoholic fatty liver scores (NAS) on liver biopsies obtained during the procedure, we separated patients with and without nonalcoholic steatohepatitis (NASH) ($n = 61$ with $NAS \geq 5$ and $n = 55$ with $NAS \leq 2$, respectively) to assess the effect of liver lesion in plasma oxylipins. To maximize the differences in liver disease, we excluded patients with intermediate damage ($NAS = 3$ or 4). We also collected paired, before and one year after surgery, bioelectric impedance data using a Tanita TBF-420 scale. Although not initially designed in the study, we collected under permission paired computerized tomography (CT) scans from several patients retrospectively to assess variations in adiposity observed with imaging techniques. We excluded patients under 18 years of age, those with excessive alcohol consumption, acute illness, malignancies, previous or current major cardiovascular events, and advanced liver disease. Our study received approval from the Institutional Review Board (EPIMET PI21/00510_083 and PL4NASH 112/2021), and all participants provided written informed consent. We collected plasma samples from ongoing research (ClinicalTrials.gov NCT05542224) while adhering to Good Clinical Practice Guidelines and the Declaration of Helsinki. The samples were collected the previous day before surgery, processed within the first 2 h, and frozen (-80°C) until analysis. We specifically selected patients undergoing LSG to study post-surgery effects in a 1-year follow-up study and plasma was collected equally. It was important to ensure the fasting state for at least 12 h and the lack of physical activity during the previous 30 min. We retained data and paired plasma samples from 44 participants who adhered to the proposed schedules for this period, resulting in an attrition rate of 17.5%. Patients were weight stable in this particular point of observation. Patients usually cease active weight loss approximately 6–8 months after surgery. A few patients may continue this process after 1 year and even some re-gain some weight but none was included in this cohort. Though susceptible to debate, we assume a static picture in the interpretation.

In addition, we obtained age- and sex-matched samples from a cross-sectional survey (GENUP 2021) conducted by the Institut d'Investigació Sanitària Pere Virgili designed to monitor health and nutritional status. We ruled out liver disease in these volunteers through laboratory values and diagnostic medical ultrasonography without liver biopsy. These samples came from 63 otherwise healthy overweight/obese participants who served as controls.

2.2. Biopsies sampling and quantification of cytokines and organokines in plasma

We collected intraoperative hepatic, visceral, and subcutaneous adipose tissue biopsies from the patients that underwent bariatric surgery, as per protocol. We used a standardized method to evaluate the alterations in liver tissue, which involved scoring fibrosis, steatosis, lobular inflammation, and ballooning in liver portions obtained through wedge resection [31–33]. Adipose tissue samples were examined to assess adipocyte size and fibrosis severity [34,35]. We used staining techniques such as Hematoxylin and eosin, Masson's trichrome, and Sirius Red to aid in these evaluations. The scores ranged from 0 to 2 or 3 for liver samples and from 0 to 3 for adipose tissue samples. We diagnosed hyperlipidemia, type 2 diabetes mellitus, and hypertension using established protocols and anthropometric and routine laboratory methods [36,37]. We measured specific cytokines C-C motif chemokine Ligand 2 (CCL2), interleukin-10 (IL-10), and tumor necrosis factor- α (TNF- α) in plasma samples using ELISAs from Peptotech (London, UK). Additionally, we measured circulating levels of irisin, FGF-19, FGF-21, leptin, adiponectin, and galectin 3 using ELISAs from R&D Systems (Minneapolis, USA).

2.3. Identification and quantification of fatty acids and oxylipin metabolites

We employed previously established procedures to identify and quantify fatty acids and oxylipin metabolites [38–42]. The measurement of oxylipins presented challenges due to their diverse chemical structures and isomers and their susceptibility to degradation and auto-oxidation. We provided access to further experimental data, reference spectra, and analytical procedures through the Metabolights database and repository under the study number MTBLS7758 (www.ebi.ac.uk/metabolights/MTBLS7758). We cross-referenced our obtained concentrations with The Human Metabolome Database to validate their accuracy (<https://hmdb.ca/>).

The process involved multiple sample preparations and runs in a chromatography system comprising a 1290 Infinity II ultra-high pressure liquid chromatograph (UHPLC) coupled with a 6490 triple quadrupole (QqQ) mass spectrometer (MS) operating in negative electrospray mode. We mixed 100 μL of plasma with 400 μL of methanol for fatty acid measurement, including a mixture of labeled fatty acids as internal standards. After centrifugation and drying 300 μL of the extract on a SpeedVac® vacuum concentrator, we reconstituted it with 100 μL of methanol for chromatographic analysis.

We combined 250 μL of plasma with 100 μL labeled oxylipins as a standard internal mixture and 500 μL of 0.001 M butylated hydroxytoluene in methanol for oxylipin measurement. Labeled internal standards enabled us to account for inter-sample differences resulting from the extraction procedure and analytical signal drift. After protein precipitation and centrifugation, we collected and purified the supernatant onto the SPE cartridge, using acetonitrile: methanol (9:1, v/v) for oxylipin elution. The samples were then dried under N_2 flow and reconstituted in water: methanol (1:1, v/v) for chromatographic analysis. To quantify specific oxylipins (Table S1), we utilized materials obtained from Cayman Chemical (Ann Arbor, MI) and generated ten seriated concentrations with good linearity using a standard internal mixture. We used stock standards dissolved in methanol for calibration curves. To ensure measurement reproducibility, we injected lipid extracts from a pool of different samples twice daily and after every 20 analyses as a quality control measure. We followed a semi-targeted approach for molecules without available standards, quantifying them using the most similar oxylipin molecule.

We excluded oxylipins with low concentrations, known variation after storage, and those resulting in more than 10% missing values or outliers. Our analysis encompassed data for 51 omega-3 and omega-6 fatty acid-derived oxylipins (Figure S1). We observed that the proposed plasma signature remained consistent across the measurements, with the time elapsed after venipuncture (<2 h until freezing) and adherence to a strict plasma processing protocol emerging as critical factors.

Our study exclusively examined non-esterified oxylipins and fatty acids without the inclusion of a hydrolysis step. It is important to clarify this point because if we do not avoid an hydrolysis step circulating lipoproteins represent a major source of esterified lipids that become non-esterified in vitro, i.e., artificially. Then, separation between these pools is impossible and adds considerable confusion. This approach allowed us to comprehensively explore metabolic adaptations and the potential use of oxylipins as a biomarker. We did not measure esterified oxylipins and fatty acids in lipoproteins, which would require additional analytical steps irrelevant to our present aim. Furthermore, it is crucial to note that the plasma albumin concentration was average in all fasting state samples.

2.4. Statistical methods and analysis

We utilized various software tools and packages, such as GraphPad Prism 6.01, R program, and related packages (ggplot2, ggpubr, dplyr, ggrepel, corrplot, and Performance Analytics) in our data analysis. We employed parametric or non-parametric tests based on the distribution of our data to determine the appropriate statistical test for bivariate comparisons and correlations. Unless stated otherwise, we presented numerical values as a median with an interquartile range. MetaboAnalyst 5.0 was used for machine learning algorithms, and MassHunter Quantitative Analysis B.07.00 was used for lipid analysis with a calibration curve generated from the corresponding standard. We applied the Benjamini and Hochberg procedure to control the false discovery rate and adjust for multiple comparisons.

3. Results

3.1. Influence of sex and liver disease on plasma oxylipin signature

We found that sex-associated differences in PUFAs and oxylipins in plasma were negligible (Fig. 1A, Table S2). Patients in the cohort were diagnosed with metabolic syndrome using the American Heart Association's criteria. Liver biopsies in patients with severe obesity revealed the potential health implications of ectopic fat accumulation [43,44]. Patients with NASH had more metabolic imbalances than those without NASH. However, we found no significant links between fatty liver disease severity, age, comorbidities, medication, BMI, or other anthropometric measurements (Table 2, Figure S2). Although plasma PUFAs were significantly lower in NASH patients, plasma oxylipin levels did not differ between those with and without NASH (Fig. 1B, Table S3). We also observed that plasma oxylipins were not predictive of NASH, and there were no significant changes in the circulating oxylipin signature across different histologic features (Figure S3). Our data firmly establish that sex and liver disease are not confounding factors in assessing the circulating oxylipin signature.

3.2. Impact of cytokine and organokine levels on the oxylipin signature

Patients with severe obesity exhibit a higher prevalence of low-grade chronic inflammation compared to overweight/obese individuals, as indicated by elevated levels of C-reactive protein,

CCL2, and TNF- α , and reduced IL-10 levels in the bloodstream (Table 1). However, these cytokines did not provide insight into plasma fatty acid or oxylipin signatures in severe obesity. It is essential to note that plasma organokine levels do not necessarily reflect interactions within different tissues, but severe obesity can exacerbate abnormal secretion. Compared to overweight or obese individuals, those with severe obesity had higher circulating levels of leptin and lower FGF-19 levels, likely indicating an influence from liver and gut but there were no significant differences in the circulating levels of other selected organokines. Similarly, the impact of metabolic comorbidities on plasma organokine levels was not significant (Figures S4 and S5). Additionally, the relationships between organokines and oxylipins were weak and only exhibited marginal statistical significance (Figures S6 and S7). These findings underscore the complexity of the metabolic phenotype in individuals with severe obesity.

3.3. Elevated oxylipin levels in severe obesity

Our study revealed that patients with severe obesity exhibit significantly higher levels of circulating saturated, mono-unsaturated, and polyunsaturated fatty acids compared to healthy overweight or less severely obese individuals, with excess adiposity playing a pivotal role. The study observed differences in both ω -6 and ω -3 polyunsaturated fatty acids. Notably, patients with severe obesity exhibited a unique plasma oxylipin profile, with most species showing significantly higher concentrations (Fig. 2 A-C, Table S4). Our analysis showed that lipoxygenase-derived oxylipins are the dominant species in the plasma of patients with severe obesity, with specific metabolites (12-HETE, 11-HDoHE, 14-HDoHE, and 12-HEPE) key in establishing the circulating oxylipins profile, as depicted in Fig. 2 D, E. The Receiver Operating Characteristic (ROC) curve based on comparing these metabolite levels indicated high accuracy in differentiating between the two groups with an Area under the Curve value of 0.937 (Figure S8). Importantly, metabolic comorbidities do not significantly impact the plasma oxylipin profile (Figure S9), indicating that lipoxygenase activity could contribute to lipotoxicity in severe obesity mainly due to inefficient adipose tissue expansion and ectopic lipid accumulation.

3.4. Changes in plasma oxylipin composition one year after weight loss surgery. The potential role of major variations in adiposity

One year after weight loss surgery, patients experienced a significant reduction in excess fat and BMI by 42% and 29.1%, respectively ($p < 0.0001$) (Table 3). The surgery also improved glucose and lipid metabolism, arterial hypertension, and decreased low-grade chronic inflammation, reducing the need for medication. Additionally, transaminase activity decreased, indicating relief in liver damage (Table 3). We observed a consistent decrease in plasma fatty acids after weight loss, likely due to metabolic adaptations (Fig. 3). Furthermore, the levels of plasma oxylipins increased following the surgery. One year after surgery, there was a notable increase in total circulating oxylipin levels, altering the plasma signature. Statistical analysis revealed significant differences between signatures, making before and one year after surgery profiles distinguishable in linear discriminant analysis. However, some differences were inconsistent, and we did not notice any apparent association with improving comorbidities. While plasma levels of individual ω -3 PUFAs-derived oxylipins remained stable or slightly decreased, those derived from ω -6 PUFAs increased. Although plasma levels of oxylipins derived from linoleic acid remained unchanged one year after surgery, those derived from arachidonic acid were significantly overrepresented. The most dominant species identified in the signature were COX-derived oxylipins,

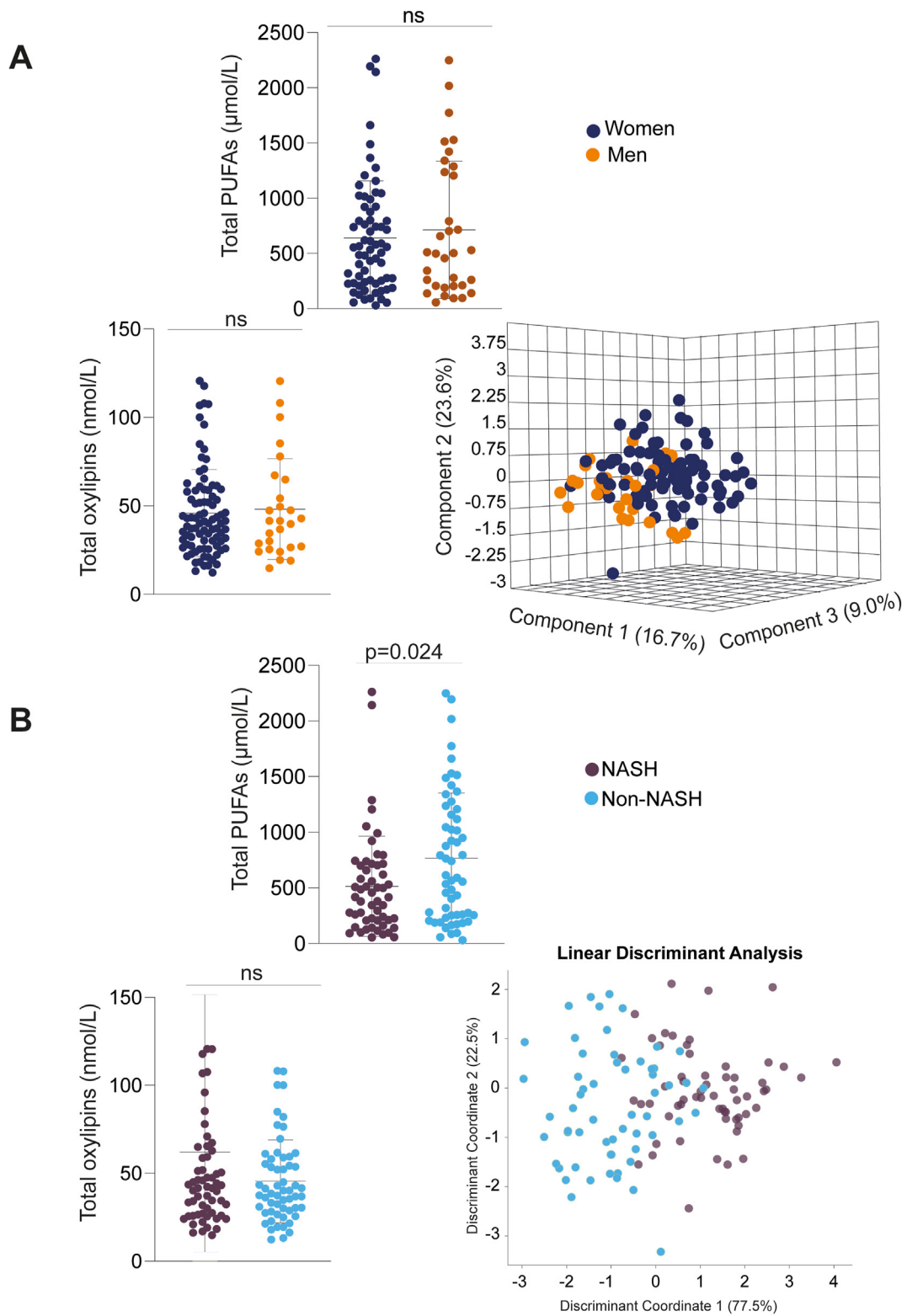


Fig. 1. Plasma levels of polyunsaturated fatty acids and oxylipins in patients with severe obesity: Impact of sex differences and NASH. (A) The results showed that sex was not a significant factor affecting measured variables. (B) In NASH patients, plasma PUFAs showed a significant decrease, whereas plasma oxylipin levels remained unchanged regardless of NASH status.

Table 1
Severe obesity-associated metabolic changes.

	Healthy overweight obese (n = 63)	Severe Obesity (n = 116)
Female, n (%)	50 (79.4)	88 (75.9)
Age (years)	44.5 (38.0–58.8)	50.0 (41.0–56.0)
BMI (kg/m ²)	27.1 (24.9–31.1)	43.4 (40.0–48.7)*
Type 2 diabetes mellitus, n (%)	1 (2.0)	36 (31.0)*
Arterial hypertension, n (%)	14 (28.0)	68 (58.6)*
Dyslipidemia, n (%)	3 (4.8)	40 (34.5)*
Medication		
Sulfonylureas, n (%)	–	4 (3.4)
Insulin, n (%)	–	14 (12.1)
Diuretics, n (%)	7 (11.1)	16 (13.8)
ACEIs, n (%)	3 (4.8)	26 (22.4)*
Biochemical variables		
Glucose (mmol/L)	4.5 (4.3–4.9)	6.7 (5.7–8.5)*
Insulin (mmol/L)	49.4 (30.7–78.9)	82.7 (42.7–121.4)*
HOMA-IR	1.4 (0.9–2.2)	3.5 (1.6–5.9)*
Triglycerides (mmol/L)	1.0 (0.7–1.4)	1.4 (1.1–1.9)*
Total Cholesterol (mmol/L)	4.9 (4.5–5.8)	3.8 (3.5–4.5)*
HDL-cholesterol (mmol/L)	1.5 (1.3–1.8)	0.9 (0.8–1.1)*
LDL-cholesterol (mmol/L)	3.1 (2.5–3.7)	2.2 (1.8–2.9)*
VLDL-cholesterol (mmol/L)	0.4 (0.3–0.6)	0.6 (0.5–0.8)
NEFAs (μmol/L)	234.4 (100.7–466.5)	2339.9 (1083.7–5033.5)*
CCL2 (pg/mL)	21.2 (10.1–36.7)	30.1 (12.4–46.1)*
IL-10 (pg/mL)	450.2 (201.3–854.4)	80.1 (62.3–85.7)*
TNF-α (pg/mL)	3.8 (2.9–7.9)	7.9 (5.7–13.6)*
CRP (mg/dL)	1.1 (0.4–2.3)	9.4 (5.3–14.3)*
Transaminases (μKat/L)		
ALT	0.2 (0.2–0.4)	0.6 (0.4–0.9)*
AST	0.3 (0.3–0.4)	0.6 (0.4–0.8)*
GGT	0.2 (0.1–0.3)	0.4 (0.3–0.6)*

Values are provided as n (percentage) or median (interquartile range). Statistical analyses performed by the Student's t-test (quantitative) or χ^2 -square test (qualitative), *p < 0.05. ACEIs: angiotensin-converting-enzyme inhibitor; ALT: alanine aminotransferase; AST: aspartate aminotransferase; BMI: body mass index; CCL2: CC-chemokine ligand2; CRP: c-reactive protein; GGT: gamma-glutamyl transferase; IL-10: interleukin 10; NEFAs: non-esterified fatty acids; TNF- α : tumor necrosis factor alpha.

Table 2
Hepatic features in patients with severe obesity.

Steatosis, n (%)	
<5%	20 (17.2)
5–33%	62 (53.4)
34–66%	26 (22.4)
>66%	8 (6.9)
Lobular inflammation, n (%)	
No foci	4 (3.5)
<2 foci per 200 × field	68 (59.1)
2–4 foci per 200 × field	41 (35.7)
>4 foci per 200 + field	2 (1.7)
Hepatocellular ballooning, n (%)	
None	32 (27.8)
Few cells	52 (45.2)
Many cells	31 (27.0)
Fibrosis, n (%)	
None (stage 0)	1 (0.9)
Perisinusoidal or periportal (stage 1)	28 (24.6)
Perisinusoidal and portal (stage 2)	53 (46.5)
Bridging fibrosis (stage 3)	30 (26.3)

Values are provided as n (percentage).

including TXB2, PGE2, PGD2, and 12-HHTre (Fig. 4, Table S5). Whether this response has any pathobiological significance is unclear and requires further research effort and other design study. In particular, we interpret these changes as the result of metabolic dysfunction caused by the accumulation and subsequent reduction in adiposity. As an observational study, paired biopsies after surgery were considered ethically unacceptable but biopsies obtained during the procedure revealed major structural changes between

subcutaneous and omental visceral adipose tissue. Notably, adipocytes in visceral adipose tissue were more fibrotic and significantly smaller in size. Similarly, CT scans one year after surgery were only indicated in a limited number of patients but illustrated major variations in adiposity (Fig. 5, Figure S10). These findings emphasize the need for further research and consideration in postoperative care.

4. Discussion

Our research underscores the importance of evaluating the impact of severe obesity and weight loss surgery on circulating fatty acids and oxylipins. It is imperative to take early therapeutic actions to prevent severe obesity, which can lead to adipose tissue dysfunction and harm other bodily tissues [30,45]. We carefully examined factors that could have affected our findings and their implications. Previous animal studies have suggested that there may be differences in oxylipin metabolism in males and females in various tissues, including the liver, depending on age, inflammation, and enzyme expression [46–48]. However, we found no evidence that these differences extend to blood plasma levels. Our analysis showed that all patients had at least two features indicating metabolic dysfunction: arterial hypertension, hypertriglyceridemia, low HDL-C, prediabetes, insulin resistance, and subclinical inflammation. However, we found no links between plasma oxylipins and these comorbidities. Sex and liver disease did not affect the plasma oxylipin profile of patients with severe obesity. Nevertheless, the disrupted endocrine patterns in leptin and FGF-19 responsiveness suggest metabolic crosstalk between adipose tissue, liver, and gut dysfunctions. In addition, we found no strong associations between organokines and oxylipins in the

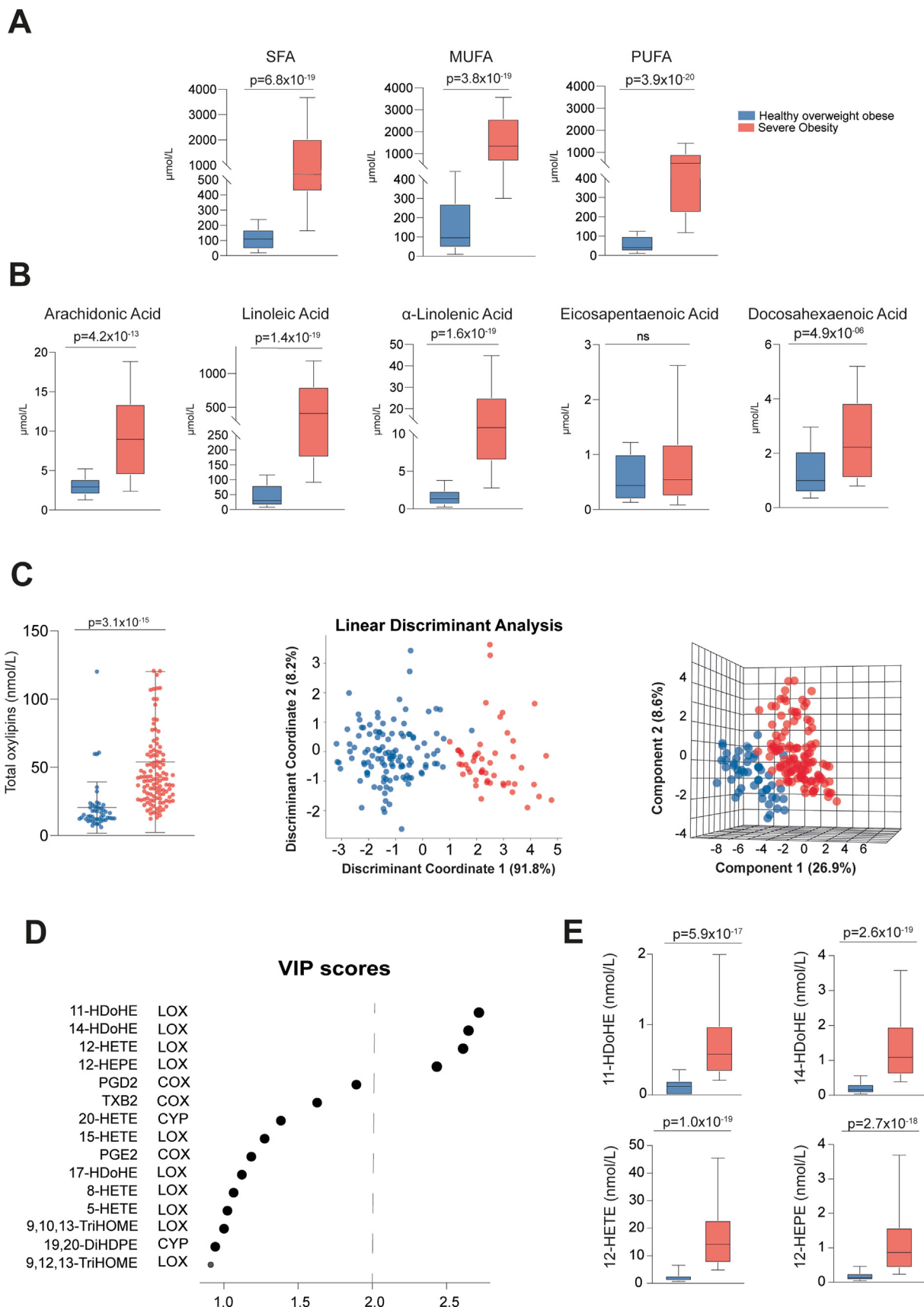


Fig. 2. Severe obesity leads to an elevated concentration of fatty acids and oxylipins into the bloodstream. (A) Plasma levels of saturated, monounsaturated, and polyunsaturated fatty acids were elevated. (B) All five selected PUFAs exhibited an increase in plasma concentration. However, the ω -3 eicosapentaenoic acid level difference was not statistically significant. (C) By performing a specific measurement of plasma oxylipins, we observed a substantial increase in most species (see also Table S4). The discriminant analysis indicated

Table 3
Bariatric surgery-associated metabolic changes.

	Before surgery (n = 44)	One year after surgery (n = 44)
Female, n (%)	31 (70.5)	—
Age (years)	53 (49–61)	—
BMI (kg/m ²)	46.0 (41.7–50.3)	32.6 (30.6–36.4)*
Fat mass (%)	50.3 (48.9–54.3)	33.7 (29.2–42.5)*
Visceral fat rating	18.0 (15.0–21.0)	9.5 (6.0–12.5)*
Type 2 diabetes mellitus, n (%)	18 (40.4)	16 (36.4)*
Arterial hypertension, n (%)	31 (70.5)	26 (59.1)*
Dyslipidemia, n (%)	20 (45.5)	9 (20.5)*
Medication		
Sulfonylureas, n (%)	3 (6.8)	—
Insulin, n (%)	9 (20.5)	2 (4.5)*
Diuretics, n (%)	12 (27.3)	1 (2.3)*
ACEIs, n (%)	15 (34.1)	1 (2.3)*
Biochemical variables		
Glucose (mmol/L)	7.1 (6.1–9.7)	4.7 (4.3–5.4)*
Insulin (mmol/L)	86.1 (52.1–118.8)	50.9 (37.3–74.6)*
HOMA-IR	3.7 (2.4–5.8)	1.6 (1.2–2.2)*
Triglycerides (mmol/L)	1.4 (1.2–2.0)	0.8 (0.7–1.4)*
Total Cholesterol (mmol/L)	3.8 (3.3–4.7)	4.2 (3.7–4.9)*
HDL-cholesterol (mmol/L)	0.9 (0.8–1.0)	1.5 (1.3–1.7)*
LDL-cholesterol (mmol/L)	2.1 (1.7–2.9)	2.0 (1.7–2.3)
VLDL-cholesterol (mmol/L)	0.7 (0.6–0.9)	0.4 (0.3–0.6)*
NEFAs (μmol/L)	2339.9 (1083.7–5033.5)	395.7 (164.4–757.8)*
CCL2 (pg/mL)	30.0 (13.1–15.2)	29.8 (13.4–42.5)
IL-10 (pg/mL)	85.8 (79.8–95.9)	180.2 (150.5–210.0)*
TNF-α (pg/mL)	8.1 (6.3–12.9)	4.1 (3.8–5.7)*
CRP (mg/dL)	9.3 (4.9–12.3)	1.3 (0.3–2.1)*
Transaminases (μKat/L)		
ALT	0.7 (0.5–1.0)	0.2 (0.2–0.3)*
AST	0.7 (0.4–0.9)	0.3 (0.2–0.3)*
GGT	0.4 (0.3–0.6)	0.3 (0.2–0.3)*

Values are provided as n (percentage) or median (interquartile range). Statistical analyses performed by the Student’s t-test (quantitative) or χ -square test (qualitative), *p < 0.05. ACEIs: angiotensin-converting-enzyme inhibitor; ALT: alanine aminotransferase; AST: aspartate aminotransferase; BMI: body mass index; CCL2: CC-chemokine ligand2; CRP: c-reactive protein; GGT: gamma-glutamyl transferase; IL-10: interleukin 10; NEFAs: non-esterified fatty acids; TNF-α: tumor necrosis factor alpha.

plasma of patients with severe obesity. This finding indicates that the relationship between organokines and oxylipins observed in animal models and isolated tissues [22–28] may not be apparent in human circulation.

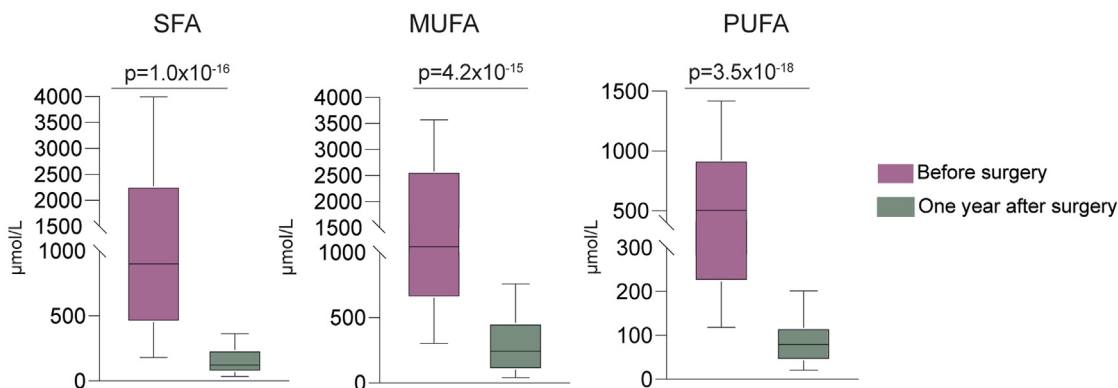
Some of our observations in severe obesity suggest that changes in the amount and function of adipose tissue are likely responsible for the metabolic response of circulating fatty acids and oxylipins. Patients with severe obesity have significantly higher levels of plasma oxylipins and fatty acids than those who are overweight or less severely obese. However, the response to weight loss varies, and while plasma fatty acid levels may return to typical values partially, the same was not valid for plasma oxylipins. The distinct plasma oxylipin signature before surgery may represent an adaptive response to sustained metabolic dysfunction over time. The lipoxygenase-derived oxylipins were the most dominant species. Although their biological significance in different tissues is still unclear, there is evidence that they regulate adipocyte lipid storage and adipokine production [20,49,50]. Moreover, they competitively inhibit metabolic pathways leading to self-limiting factors in inflammation [51–54].

After losing weight, we observed a reversal in the effects of severe obesity, and we confirmed a significant improvement in metabolic dysfunction [55,56]. However, there was an increase in plasma oxylipin levels, and we noted qualitative differences in the

plasma oxylipin signature. Following surgery, arachidonic acid derivatives dominated the circulating oxylipin signature, specifically significant increases in prostaglandins and leukotrienes. Although studies in mice have shown that dysfunctional oxylipin metabolism occurs exclusively in adipose tissue, no guidance is available for humans. Our investigation revealed significant alterations in visceral adipose tissue (VAT) following bariatric surgery, underscoring its importance in the context of severe obesity. Post-surgery assessments showed a notable reduction in VAT and a substantial decrease in subcutaneous adipose thickness, resulting in a marked decrease in overall body fat percentage. Further analysis of VAT biopsies exhibited reduced adipocyte size. We could not determine the size of the adipose tissue with CT scan because the image we obtained was not standardized. These findings suggest that bariatric surgery impacts total body fat and induces profound changes in the composition and quality of VAT. These observations emphasize the interplay between VAT alterations and metabolic improvements observed post-bariatric surgery [57,58]. Our findings highlight the contribution of oxylipin trafficking to the metabolic adaptation in adipose tissue after weight loss. We confirm previous data that suggest PGD2 plays a potential role in driving weight gain [13], and the increase in TXB2 levels may indicate adipose tissue remodeling [59]. However, studies on the correlation between plasma levels of primary prostanoid metabolites and

a distinct profile in individuals with severe obesity. (D) Docosanoids and eicosanoids are overrepresented among the measured oxylipins in comparison to octanoids. (E, F) Variable importance in the projection of the partial least squares-discriminant analysis identified the critical role of lipoxygenase-derived metabolites. We used the Mann–Whitney U test to assign statistical significance to comparisons. DiHDPE: dihydroxydocosapentaenoic acid; DiHODE: dihydroxyoctadecadienoic acid; HDOHe: hydroxydocosahexaenoic acid; HEPE: hydroxyeicosapentaenoic acid; HETE: hydroxyeicosatetraenoic acid; MUFA: monounsaturated fatty acids; PUFA: polyunsaturated fatty acid; SFA: saturated fatty acid; TriHOME: trihydroxyoctadecanoic acid.

A



B

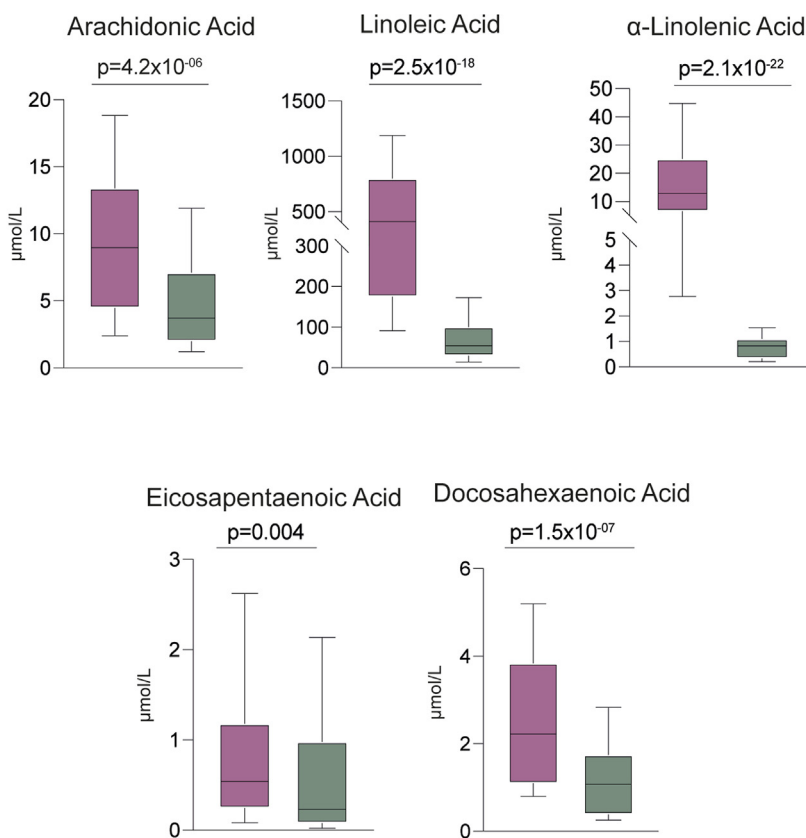


Fig. 3. Plasma fatty acids decreased after weight loss. Although patients remained obese one year after the surgery, plasma fatty acids decreased significantly (A), including those characterized as precursors of oxylipins (B). We utilized the Mann–Whitney U test to determine the statistical significance of the comparisons before and one year after surgery. MUFA: monounsaturated fatty acids; PUFA: polyunsaturated fatty acids; SFA: saturated fatty acids.

cardiovascular risk have yielded contradictory results. TXB2 stimulates platelet aggregation [60–62], while PGE2 promotes anti-inflammatory macrophage polarization and reduces atherosclerosis progression [63]. Nevertheless, the necessary research to understand the mechanisms behind these effects in humans is complex and raises considerable ethical concerns.

We identify limitations that may improve the design of future research. The study is observational and solely focuses on patients

with severe obesity. Patients remained obese after the one-year observation period, and it is unclear if patients achieved complete metabolic homeostasis. Variations in the gut microbiota may affect the relationship between obesity and metabolic dysfunction. Our findings on oxylipins suggest that patients who have undergone bariatric surgery need monitoring for the long term. Ongoing research will explore the mechanisms contributing to the excessive accumulation and management of adipose tissue metabolism.

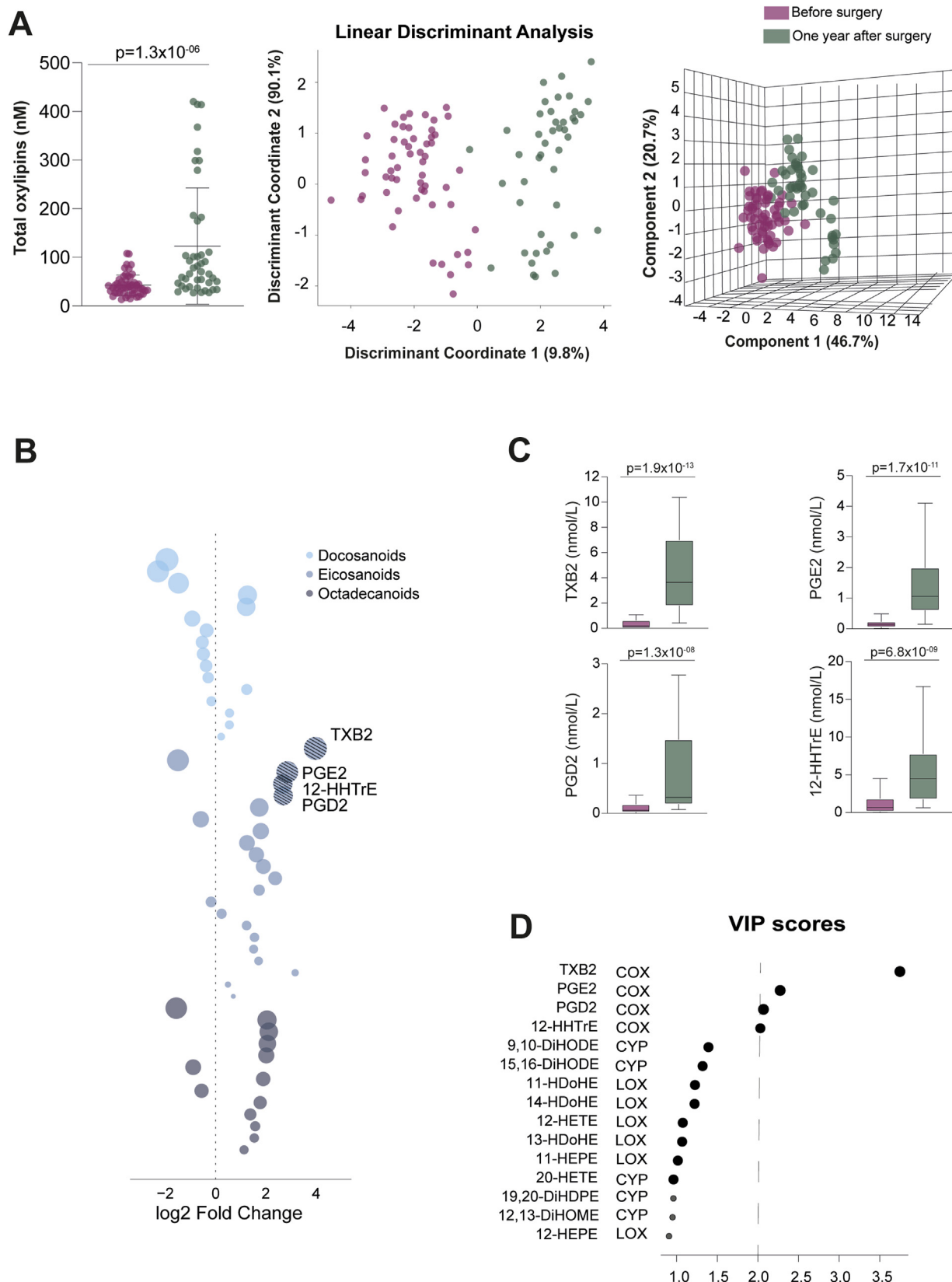


Fig. 4. Surgical weight loss led to an increase in the levels of oxylipins in plasma, which caused changes in the oxylipin profile. The analysis showed a clear separation before and after the surgery (A), with some oxylipin species being decreased, particularly within docosanoids and octadecanoids, while prostaglandins and leukotrienes increased significantly (B). The most dominant species identified by the random forest analysis were cyclooxygenase-derived metabolites from arachidonic acid (C, D). We used the Mann–Whitney U test to determine the statistical significance of the comparisons. HHTrE: hydroxyheptadecatrienoate; PGD2: prostaglandin D2; PGE2: prostaglandin E2; TXB2: thromboxane B2.

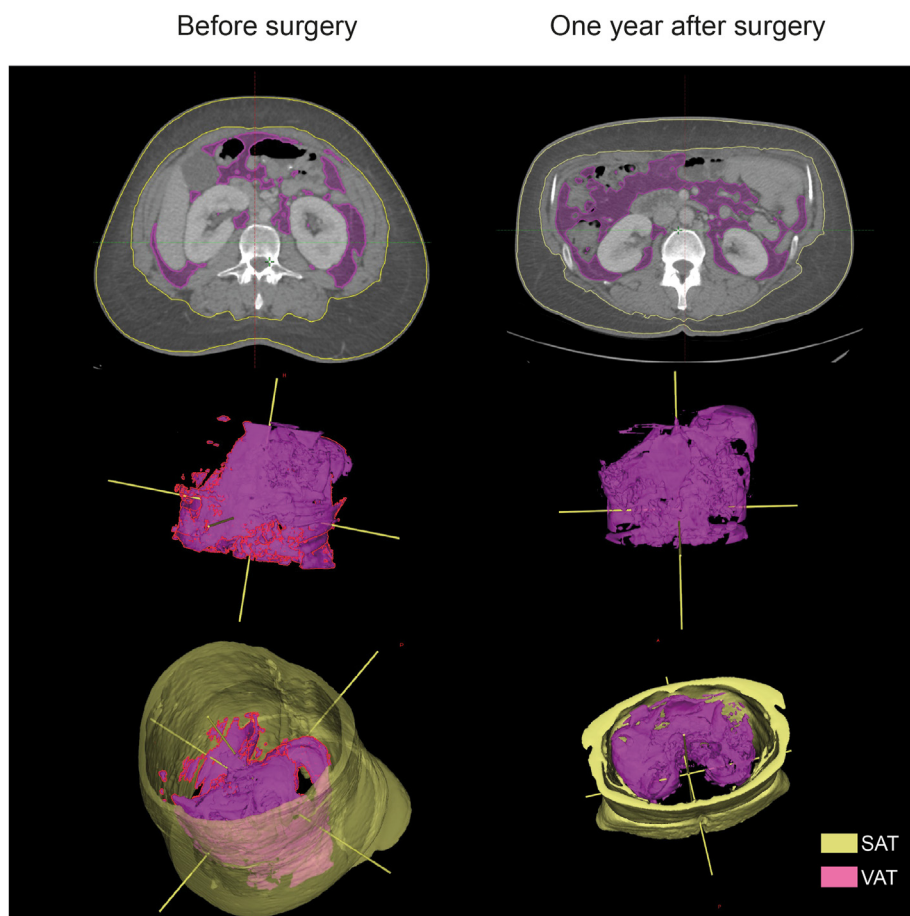


Fig. 5. Computed-tomography scans illustrate the dramatic reduction in adipose tissue one year after surgery.

Other vital questions include why weight loss maintenance becomes easier over time, why macrophage infiltration into adipose tissue does not reverse after weight loss, and how metabolic adaptations persist longer in adipocytes than in other cell types [64–70].

5. Conclusions

Metabolic imbalances are a common consequence of severe obesity. Our research suggests that in patients with this condition, there is an altered balance in circulating fatty acids and oxylipins, leading to higher levels than in overweight/obese individuals who are metabolically healthy. The activity of LOX plays a crucial role in determining the types of oxylipins present in the bloodstream of patients with severe obesity, providing a distinct signature. After weight loss surgery, we observed a shift in the oxylipin signature, with COX-derived species becoming more predominant after one year. This finding was unexpected and requires further investigation to understand the complex changes in metabolic dysfunction in adipose tissue during obesity and weight loss. Effectively addressing metabolic dysfunction in severe obesity is a challenging but essential step in improving the health outcomes of those affected. Future studies aim to address this issue comprehensively.

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Conflict of interest

The authors declare that there are no competing interests in disclosing personal or financial interests, and there are no external factors potentially affecting the objectivity and quality of the information.

Authorship contribution statement

The contributions of the authors to this study are as follows: Andrea Jiménez-Franco and Helena Castañé: Conceptualization, methodology, formal analysis, data curation, and writing drafts. Cristian Martínez-Navidad, Cristina Placed-Gallego, and Marta Canela-Capdevila: Data management, software development, and laboratory tasks. Salvador Fernández-Arroyo and Iris Samarra: Formal analysis and data curation. Meritxell Arenas, Antonio Zorzano, and María Isabel Hernández-Alvarez: Conceptualization and methodology. Daniel del Castillo and Marta Paris: Patient management, supervision, and adherence to ethical standards. Javier A Menendez: Intellectual contribution. Jordi Camps: Supervision, project management, final approval. Jorge Joven: Conceptualization, data curation, revision and editing, and funding.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.clnu.2023.12.002>.

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