

Hepatic fatty acid profile in the rat model of NAFLD: influence of sex and diet

Kristina Starčević¹ (kristina.starcevic@vef.hr), Petra Roškarić¹, Josip Barišić², Tomislav Mašek³ (tomislav.masek@vef.hr)

¹Department of Chemistry and Biochemistry, Faculty of Veterinary Medicine, University of Zagreb, 10000 Zagreb, Croatia.

²Laboratory for Biotechnology in Aquaculture, Division of Materials Chemistry, Ruđer Bošković Institute, 10000 Zagreb, Croatia

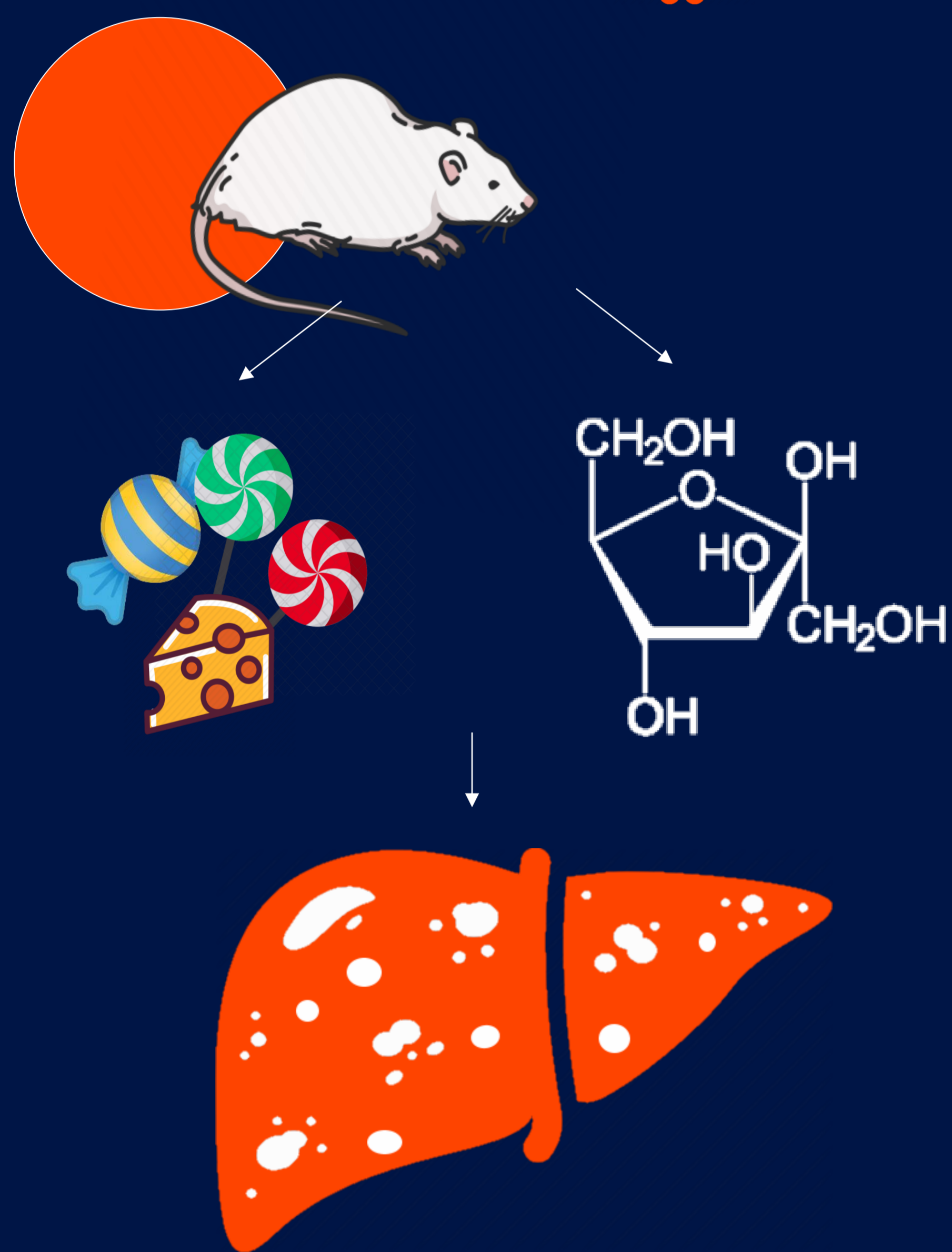
³Department of Animal Nutrition and Dietetics, Faculty of Veterinary Medicine, University of Zagreb, 10000 Zagreb, Croatia, tomislav.masek@vef.hr

Background



Nonalcoholic fatty liver disease (NAFLD) is an important health disorder with the increasing incidence in Western countries. High-fructose and cafeteria diet rodent models have been important source of data on the pathophysiological mechanisms of NAFLD. Therefore, our aim was to investigate the differences in the hepatic fatty acid profile and the influence of diet and sex in these models.

Experimental design



Results

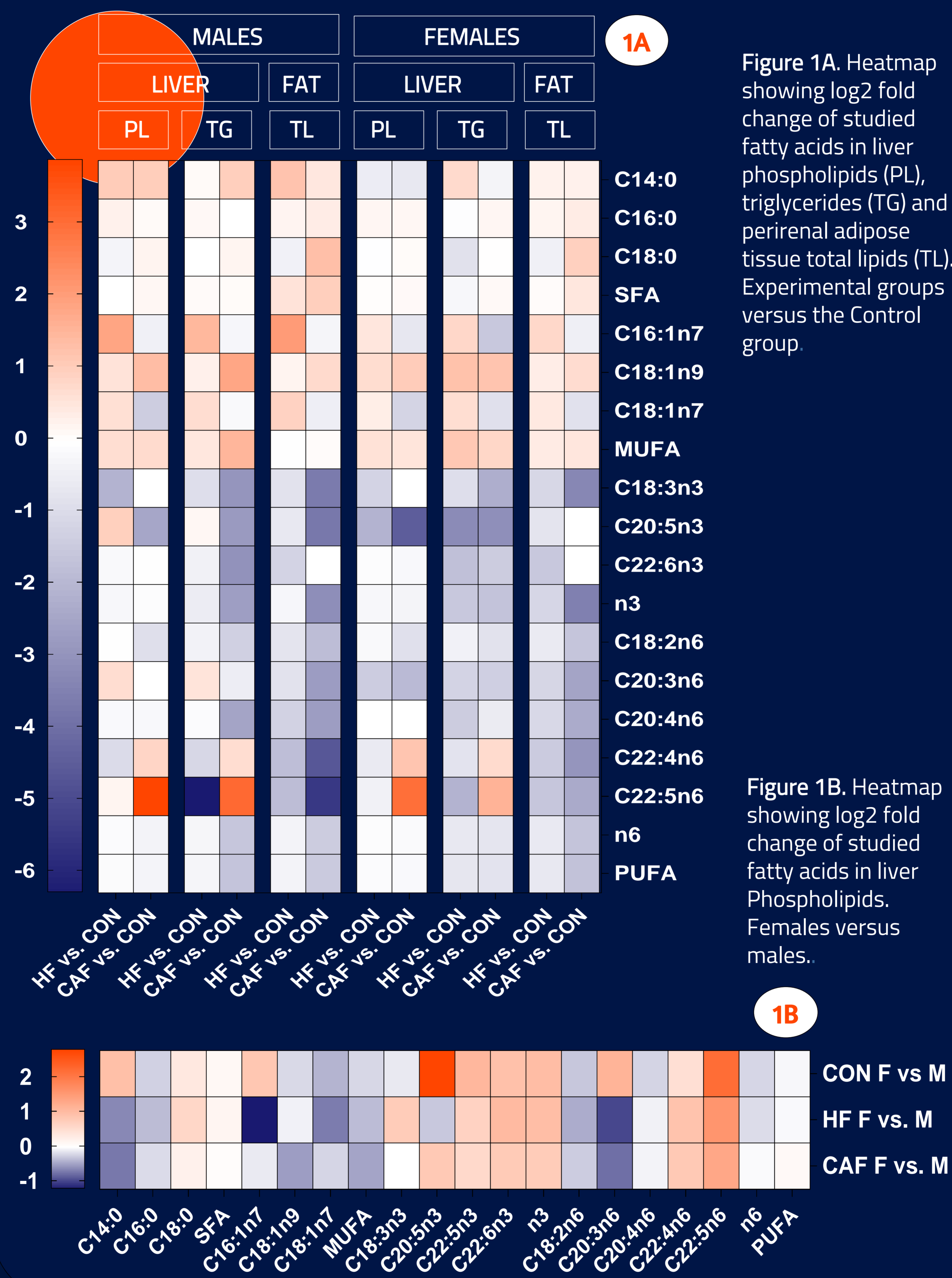


Figure 1A. Heatmap showing log₂ fold change of studied fatty acids in liver phospholipids (PL), triglycerides (TG) and perirenal adipose tissue total lipids (TL). Experimental groups versus the Control group.

Figure 1B. Heatmap showing log₂ fold change of studied fatty acids in liver Phospholipids. Females versus males.

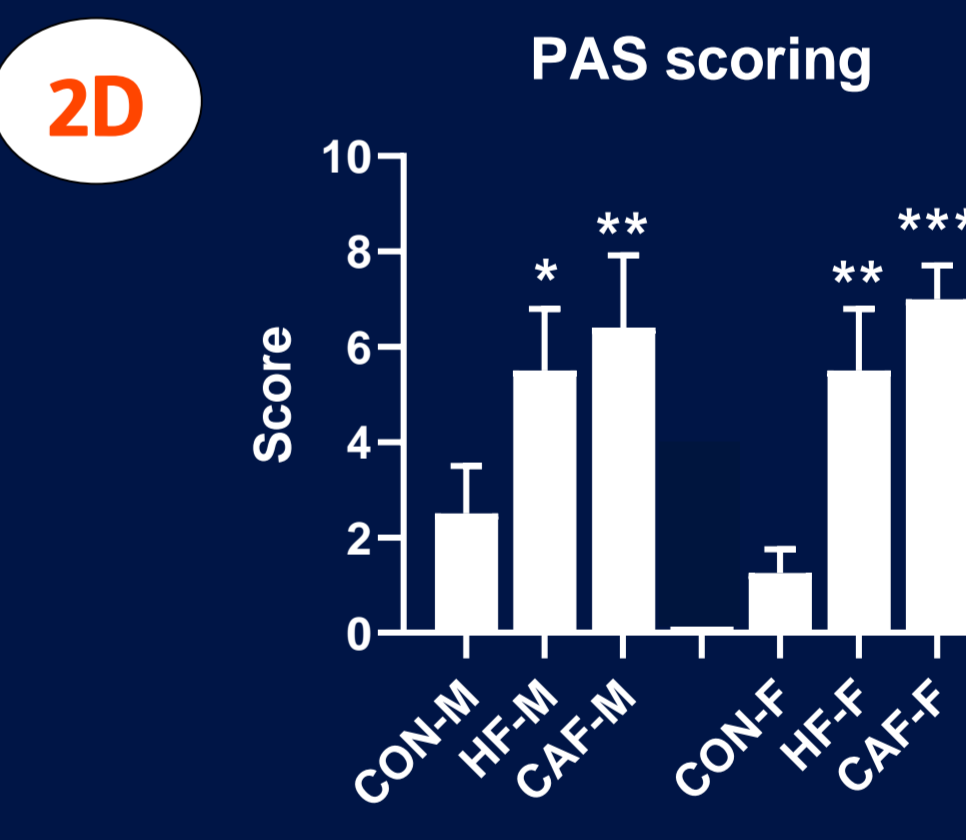
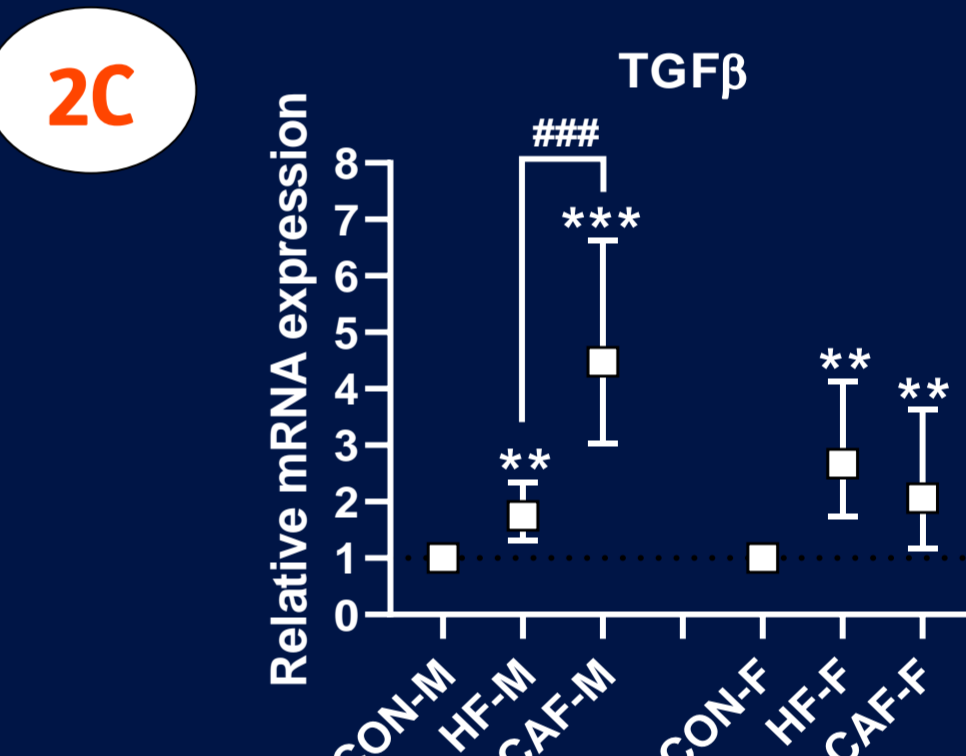
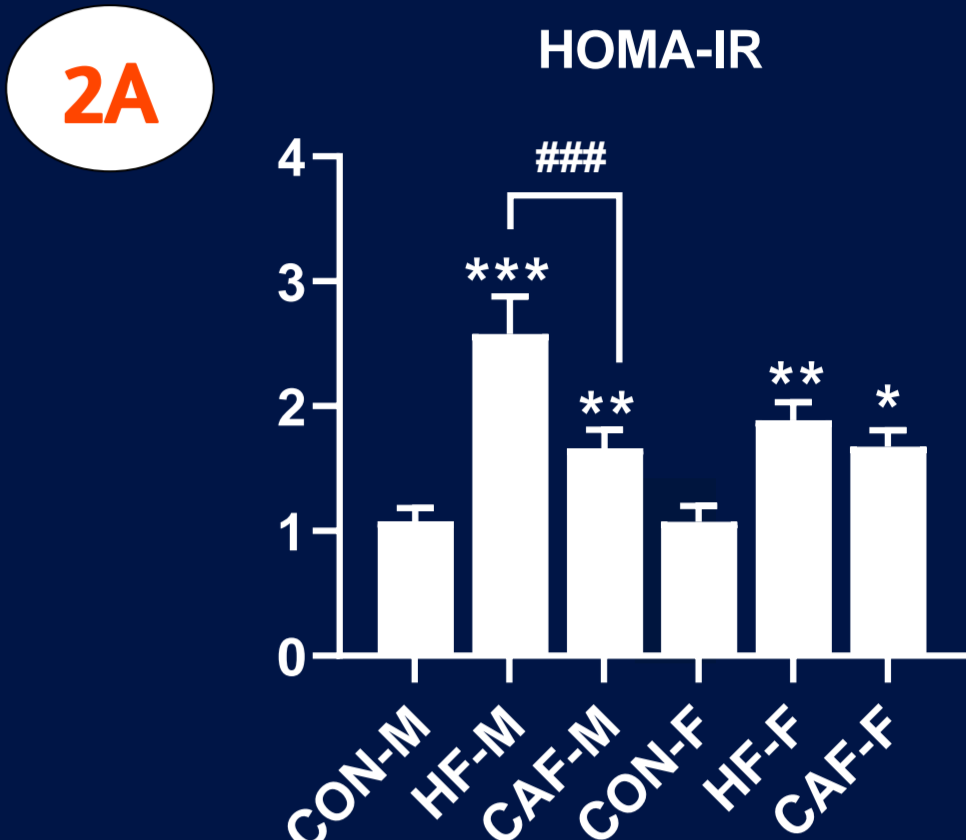


Figure 2. Different features of NAFLD in treated rats. Increased HOMA-IR index (A). Lipid peroxidation investigated as malondialdehyde concentration (B). Increased expression of inflammation markers (TGFβ) (C) and PAS staining quantification (D).

Materials and methods

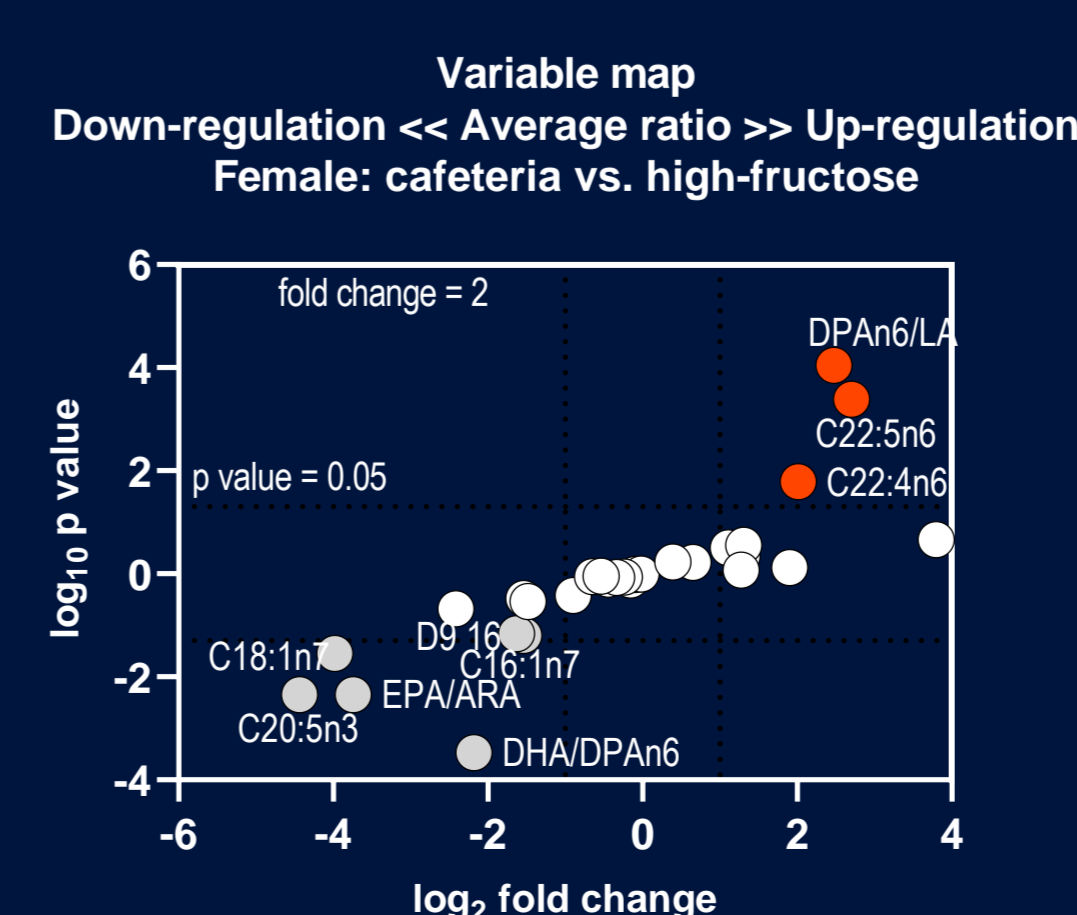
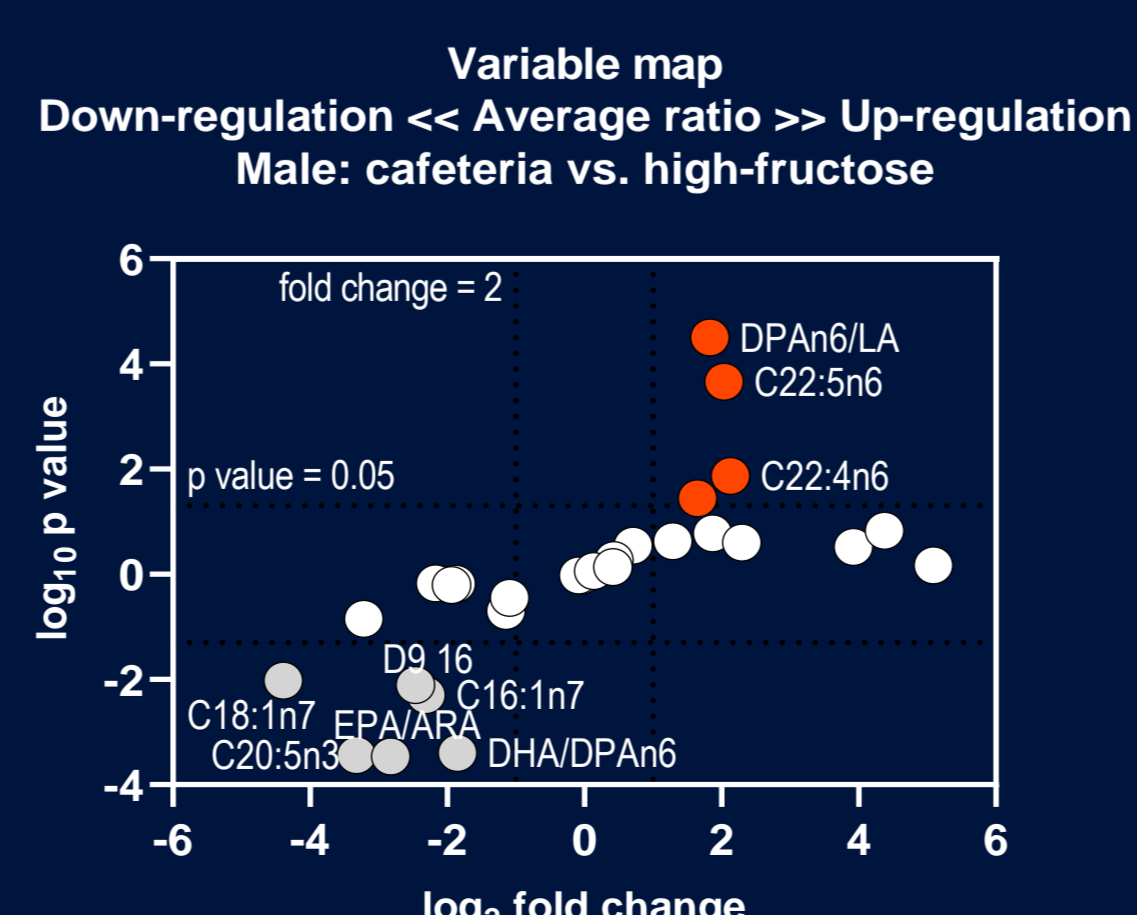


Thirty-six Wistar rats (18 male and 18 female) were divided into the control group (CON), the high fructose group (HF, 15% of fructose in the drinking water) and the cafeteria diet group (CAF, 50% basal diet and 50% cafeteria diet). All dietary treatments lasted for 20 weeks. Liver histopathology was assessed by H&E, PAS and Oil red staining, lipid peroxidation was assessed by measuring MDA-TBARS and 4-HNE and the expression of the inflammation gene markers was quantified by RT-qPCR. The analysis of the hepatic fatty acid composition was performed using gas chromatography after the lipid extraction and methylation. For statistical data analysis, GraphPad 8 was used. Data were compared using the analysis of variance and Tukey post hoc test.

Conclusions



The results showed significant differences in the hepatic fatty acid profile in investigated rat models of NAFLD. The observed differences include fatty acids with important biological effects (e.g. n3 PUFA), which, therefore, must be considered in the investigations of NAFLD.



Acknowledgements



Supported by the Croatian Science Foundation as part of the project (IP-2016-06-3163) awarded to Kristina Starčević.