Impact of chronic heart failure on adipose tissue functional plasticity – a role for fatty acids?



UNITED KINGDOM · CHINA · MALAYSIA

Andrew D Yale¹, Malcolm Cobb¹, Alexander Lyon², Alison Mostyn¹

¹Faculty of Medicine and Health Sciences, School of Veterinary Medicine and Science, Sutton Bonington Campus, University of Nottingham, LE12 5RD, ² NIHR Cardiovascular Biomedical Research Unit, Royal Brompton Hospital

Introduction

Chronic heart failure (HF) is a major cause of morbidity and mortality in humans and domestic animals. HF is a systemic condition characterised by chronic inflammation and functional changes in non-cardiovascular tissues; obesity is also associated with low level chronic inflammation, with adipose tissue (AT) exhibiting infiltration of macrophages and alterations in fatty acid (FA) profiles⁽¹⁾.

Our **hypothesis** was that AT from animals with HF will display greater expression of inflammatory mediators and a fatty acid profile consistent with an inflammatory state.

Therefore the **aim** of this project was to investigate expression of regulators of lipid metabolism and inflammation and fatty acid profiles in AT from a rodent model of HF.

Methods

Rats with HF post myocardial infarction were compared with age matched non-failing controls. Intra-abdominal (IA) and subcutaneous (SC) adipose tissue depots were sampled after humane euthanasia, and then processed for analysis.

Adult male Sprague-Dawley rats (N=6) underwent proximal coronary ligation to induce chronic myocardial infarction. Sham ligation was used as control (N=6). Sixteen weeks later assessment of cardiac function completed and rat humanely euthanased and adipose tissue removed and stored at -80°C.

RNA extracted, reverse transcribed and subjected to QPCR. Rat specific primers for several inflammatory mediators and metabolic markers included stearoyl-CoA desaturase-1 (SCD1), interleukin 1 (IL1) and phosphoenolpyruvate carboxykinase (PCK1)

Lipid extracted (Folch method⁽²⁾); triglycerides (TAG) measured colourometrically, fatty acid profiles assessed using gas chromatography.

Statistical significance was assessed using ANOVA with post hoc Tukey tests

Results

<u>Triglycerides</u> - Greater storage of TAG observed in HF intraabdominal AT.

<u>Gene Expression</u> - Reduced expression of SCD1 (enzyme that catalyses desaturation to mono-unsaturates) in HF intra-abdominal AT.

<u>Fatty Acids</u> - FA analysis identified differences in several species; including reductions in C10, C15 and C17 in HF-SC. However, most interesting was a reduction of both C20:5n3 (EPA) and C22:6n3 (DHA) in HF-SC.

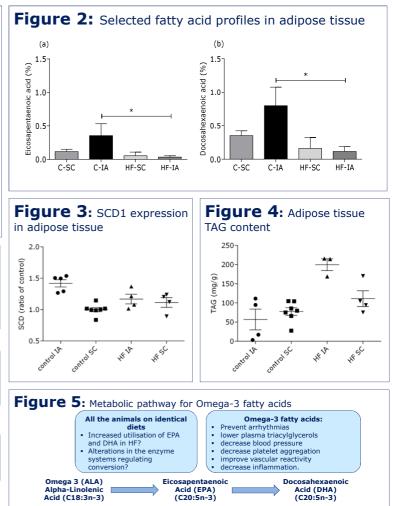


Ő

SHAM

HF

SHAM = sham ligation (control group) MI HF = myocardial infarction heart failure LVEF = left ventricular ejection fraction



Conclusions

Despite a reduction in SCD1 expression, we did not detect any difference in mono-unsaturated FA species in the HF rats.

EPA and DHA are omega 3 FAs with recognised antiinflammatory and cardio-protective effects; a reduction in HF suggests altered uptake and metabolism in these animals.

Future Research

Possible areas for further research are:

- To investigate the impact of SERCA2 gene therapy and whether it can ameliorate the effects of HF on adipose tissue.
- To assess the impact of HF on brown adipose depots given. the emerging roles of BAT in adult mammals.
- To determine whether the FAs act as biomarkers of HF.
- Nutritional intervention

Reference

⁽¹⁾ Cinti S, et al. J Lipid Res. 2005;46.

⁽²⁾ Folch, J., M. Lees and G. H. Sloane Stanley (1957). "A simple method for the isolation and purification of total lipids from animal tissues." <u>J Biol</u> Chem **226**(1): 497-509

⁽³⁾ Lyon, A. R., et al. (2009) Proceedings of the National Academy of Sciences **106**(16): 6854-6859.



Contact For more information please contact: svyay@nottingham.ac.uk