

The Effect of Chronic High Dose Vitamin A Supplementation on Lipid Metabolism in Adipose Tissue (P02-013-19)

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Objectives: The objective of this study was to assess the impact of high-dose vitamin A (VA) on lipid metabolism. Previously, VA has been shown to enhance fat mobilisation, leading to a reduction in body fat. We hypothesise that hypervitaminosis A will increase expression of genes associated with lipid catabolism.

Methods: To induce chronic hypervitaminosis A, two groups of pigs ($n = 8$) were fed a commercial diet. The treatment group was additionally dosed, daily, with an oral supplement of retinyl propionate of 10,000 $\mu\text{g}/\text{KgBW}$ for 17 weeks. To assess the impact of VA on lipid metabolism, a microarray analysis was performed to identify gene

expression in adipose tissue. Differentially expressed transcripts and pathways were identified using Genespring and mapped to human orthologues for Ingenuity Pathway Analysis (IPA); gene fold changes were confirmed using qRT-PCR. Additionally, an untargeted UPLC-MS lipidomic analysis was carried out in serum samples to identify changes in lipid classes and their metabolites.

Results: In dosed animals, significant increases in plasma retinol (0.66 $\mu\text{mol}/\text{L}$) and liver retinyl ester concentrations (11.98 $\mu\text{mol}/\text{g}$ both $P < 0.001$), as well as an increase in serum NEFA of 92.84 $\mu\text{mol}/\text{L}$ ($P = 0.001$) were observed. Gene expression fold changes in subcutaneous adipose tissue were related to mitochondrial dysfunction and lipid metabolism, including increased expression of MT-CYTB ($\uparrow 4.78\text{x}$, $P < 0.05$) and ATP5A1 ($\uparrow 3.13\text{x}$, $P < 0.05$). Metabolomics confirmed changes in lipids and their metabolites relevant to adipose tissue in blood ($P = 0.05$), namely a decrease in triacylglyceride concentration and increases in acyl carnitine and cardiolipin concentrations.

Conclusions: An integrated pathway is suggested to explain the role of vitamin A in leading to increased lipolysis, β -oxidation and oxidative phosphorylation, but when in excess, markers of mitochondrial dysfunction were observed.

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