## Abstract

Metabolic pathways in adipose tissue affect the whole-body energy homeostasis. *De novo* lipogenesis and futile metabolic cycling based on lipolysis and fatty acid re-esterification which is engaged in regulation of fatty acid level in bloodstream are occuring there. These processes are partly regulated by nuclear receptor PPAR $\gamma$ . Mitochondrial biogenesis and oxidative phosphorylation in adipocytes are controlled by interacting of PPAR $\gamma$  with transcriptional coactivators PGC-1 $\alpha$  and PGC-1 $\beta$ . The aims of this thesis were to find out whether PGC-1 $\beta$  is connected with regulation of futile cycling and *de novo* lipogenesis in white adipose tissue and also how specific inactivation of PGC-1 $\beta$  gene in adipose tissue affects phenotype of mice during short-term cold exposure or treatment based on high fat diet enriched by *n*-3 polyunsaturated fatty acids in combination with mild calorie restriction.

The results show that inactivation of PGC-1 $\beta$  probably does not affect futile cycling based on lipolysis and fatty acid re-esterification. In mice with PGC-1 $\beta$  ablation compensation in weight of brown adipose tissue was observed as well as increase in the gene expression of nuclear receptors PPAR, transcriptional coactivator PGC-1 $\alpha$  and UCP1 during cold exposure. Even though the inactivation of PGC-1 $\beta$  in brown adipose tissue is compensated by upregulation of gene expression and also by increase of UCP1 protein, tissue does not appear to be fully functional.