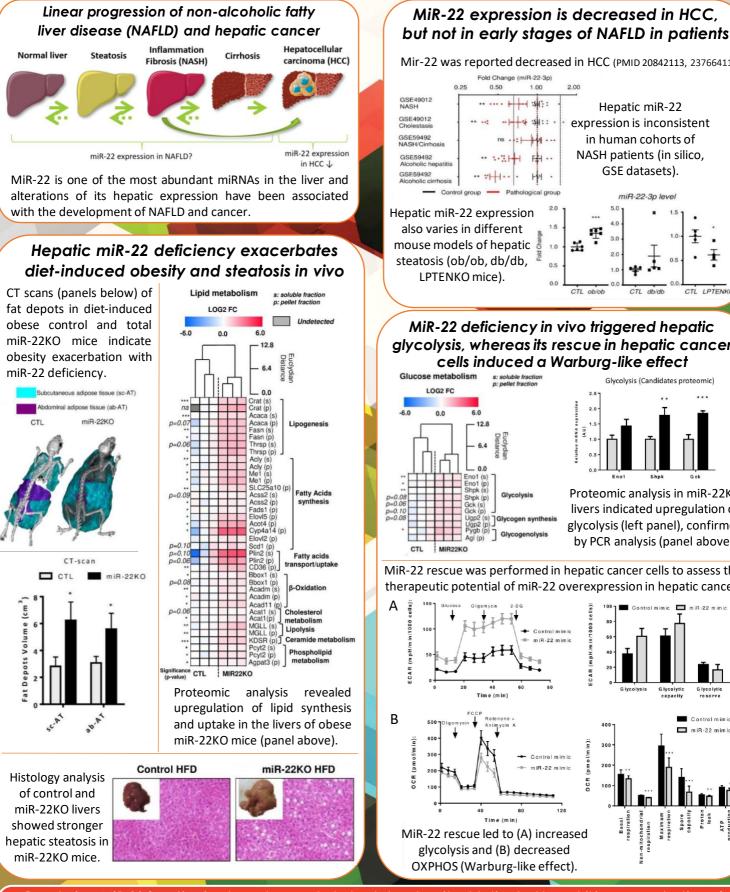
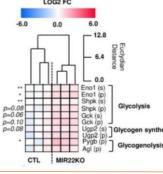
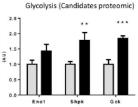
## MiR-22 differentially regulates energy metabolism in normal versus hepatic cancer cells

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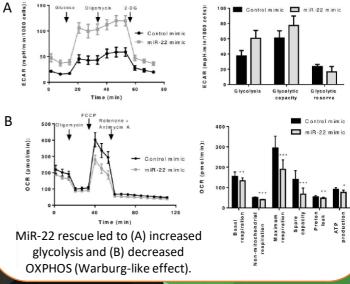
Mir-22 was reported decreased in HCC (PMID 20842113, 23766411) Fold Change (miR-22-3p) 0.50 1.00 \*\* \*\*\*\* +++++\* :#13 : Hepatic miR-22 expression is inconsistent in human cohorts of :-+++ ... NASH patients (in silico, F#1 .. 1.21 GSE datasets). .. ... HH miR-22-3p level 5.0 Hepatic miR-22 expression 4.0 1.5 also varies in different E. mouse models of hepatic 3.0 1.0 **-T**-2.0 steatosis (ob/ob. db/db. 0.5 1.0 LPTENKO mice). CTL LPTENKO CTL db/dt MiR-22 deficiency in vivo triggered hepatic glycolysis, whereas its rescue in hepatic cancer cells induced a Warburg-like effect Glycolysis (Candidates proteomic)





Proteomic analysis in miR-22KO livers indicated upregulation of glycolysis (left panel), confirmed by PCR analysis (panel above).

MiR-22 rescue was performed in hepatic cancer cells to assess the therapeutic potential of miR-22 overexpression in hepatic cancer.



Conclusion: MiR-22 function is relevant on a whole-body level, as its ablation led to a striking exacerbation of diet-induced obesity and hepatic steatosis. MiR-22 deletion further led to a metabolic reprograming in the liver (glycolysis and lipogenesis 1). Nevertheless, its rescue (overexpression) in a hepatic cancer cell line resulted in a Warburg-like metabolic switch, calling for caution when considering this miRNA as a therapeutic target in HCC. Gjorgjieva et al., JPM, 2020