



Introduction

✓ Drug-induced steatosis is a side effect associated with various drugs arising from the excessive accumulation of fats in liver cells. This effect can interfere with normal liver function and progress into more severe anomalies (1). \checkmark Specific drugs causing this problem include Tetracycline, Valproic Acid, Amiodarone, Methotrexate etc. (2). • Receptor tyrosine kinases (RTKs) play critical for various cellular processes and cell types including hepatocytes. Thus, it is worthwhile to investigate whether particular RTK plays a crucial role in regulating lipid metabolism in hepatocytes, making them potential therapeutic targets for treating drug-induced steatosis. \checkmark Based on primary screening, we identify that AXL inhibition alleviates steatosis in hepatocyte cell lines.

✓ To study effect of AXL inhibitors on tetracycline induced steatosis and its underlying pathway

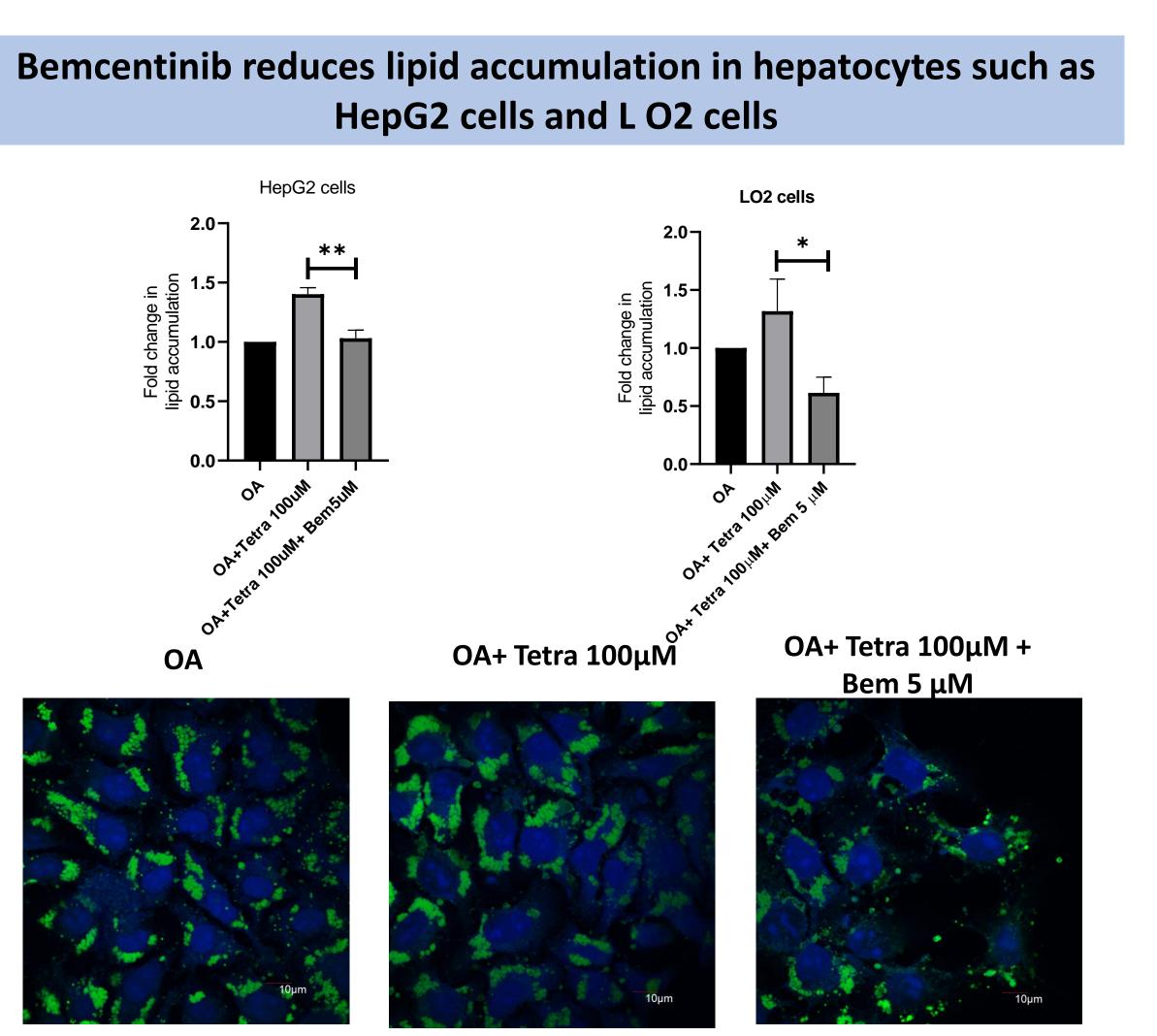


Fig.1 Effect of AXL inhibition on lipid accumulationA. Effect of Bemcentinib on tetracycline induced lipid accumulation in HepG2 and L 02 cells. B. Confocal image of HepG2 cells treated with OA, OA with Tetra100µM, OA with Tetra100µM in presence of Bemcentinib 5 μM. OA: Oleic Acid, Tetra: Tetracycline, Bem: Bemcentinib. One –way ANOVA, n= at least 3, *p<0.05, ***p<0.005, ****p<0.0001

Conclusion and Discussion

✓ AXL inhibition alleviates drug induced steatosis and could be explored as therapeutic strategy to manage steatosis. ✓AXL inhibition alters the regulation of lipid metabolism. It reduces *de novo* lipogenesis and fatty acid uptake, while induces β-oxidation. \checkmark AXL inhibition modulates transcriptional factors such as SREBP1c and PPAR α involved in *de novo* lipogenesis and β -oxidation via ERK and mTOR pathway.

AXL: a potential target to alleviate Tetracycline-induced liver steatosis.

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Objective

Results



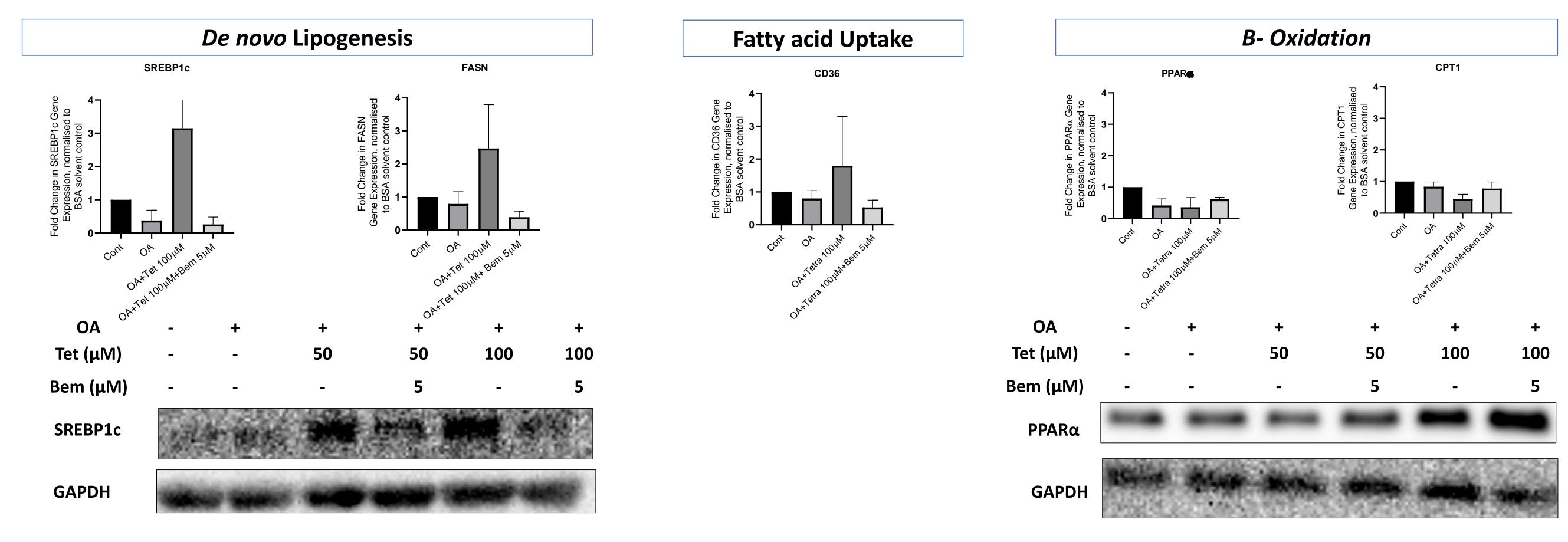


Fig. 2 Effect of AXL inhibition on lipid metabolism in HepG2 cells. A. Effect of Bemcentinib on genes involved in lipid metabolism was analysed using RT-PCR. Bemcentinib reduced gene expression of SREBP1c and FASN involved in *de-novo* lipogenesis. It also reduced expression of gene CD36 which is involved in fatty acid uptake. Bemcentinib, induced expression of genes involved in β-oxidation such as PPARα and CPT1. B. Effect of AXL inhibition on expression of transcriptional factors such as SREBP1c and PPARα normalized to GAPDH. OA: Oleic Acid, Tetra: Tetracycline, Bem: Bemcentinib. One –way ANOVA, n= at least 3 for SREBP1c, FASN and n=2 for CPT1, PPARα and CD36for RT-PCR. N=2 for immunoblotting analysis. *p<0.05, ***p<0.005, ****p<0.0001

This study provides a prototype approach to explore other RTKs for steatosis induced by various other drugs

Materials and Methods

- ✓ HepG2 cells were pre-treated with Oleic Acid (0.075mM) and Tetracycline (100µM) to mimic drug-induced steatosis. \checkmark Effect of AXL on steatosis was investigated using AXL inhibitor, Bemcentinib at concentration 5 μ M along with Oleic acid and Tetracycline.
- Social Section Applies Appl was used for qualitative analysis of lipid accumulation.
- Y RT-PCR was done to quantify relative mRNA expression of genes involved in lipid metabolism such as SREBP1c, FASN, CD36, PPARα and CPT1 normalized to GAPDH
- ✓ Immunoblotting assays were performed to observe changes on protein expression of transcriptional factors such as SREBP1c and GAPDH

Fatty acid uptake and induces β-oxidation

Reference

(1) Cataldi M. *et.al*. Adv. Ther. 2021 (2) Kolaric, T. O.; Nincevic, V. and Kuna, L. et.al. J. Clin. Transl. Hepatol. 2021.

Acknowledgement

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PPARα. Changes in activation of mTOR and ERK pathway were analysed by immunoblotting assay. All samples were normalised to

Bemcentinib reduces activation of mTOR and ERK pathway which may regulate lipid metabolism							
	-		+ 50 -	+ 50 E	+ 100	+ 100 E	
Bem (µM)	-	-	-	5	-	5	
mTOR p mTOR	5		-	-			
GAPDH	-			-		6660	
ERK				=			
p ERK		-	-	-			
GAPDH		-				and the second	

Fig. 3 Effect of AXL inhibition on mTOR and ERK pathway OA: Oleic Acid, Tetra: Tetracycline, Bem: Bemcentinib. One –way ANOVA. n=2 for immunoblotting analysis. *p<0.05, ***p<0.005, ****p<0.0001