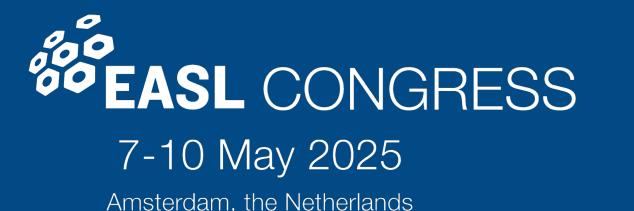
# Transcriptomic and metabolic insights into hyperammonemia: the complementary therapeutic roles of toll-like receptor 4 inhibitor and ornithine phenylacetate



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# Introduction

- Hyperammonemia is a critical issue in patients levels predict mortality and a reduction in ammo
- Despite this, no effective therapies are available
- Previous studies have identified toll-like recepto hyperammonemia by modulating mitochondrial

Aim

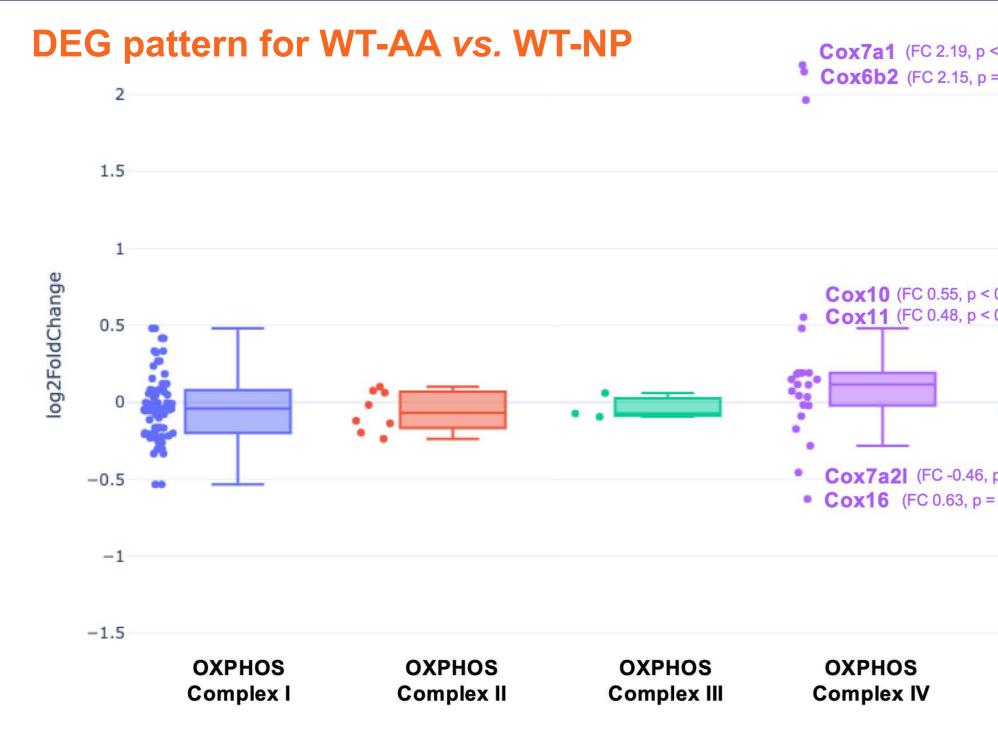
 To explore the therapeutic potential of TLR4 ant **Ornithine Phenylacetate (OP)** through transcri treatment strategies.

## Methods

- A mouse model of chronic hyperammonemia (V was studied<sup>1,2</sup>.
- Ammonia levels were significantly elevated in V both TAK-242 and OP treatment effectively re levels.
- Liver transcriptomic analysis was performed usi to investigate key metabolic pathways, including tricarboxylic acid cycle, oxidative phosphorylatic complexes, and oxidative stress pathways.

# Main findings

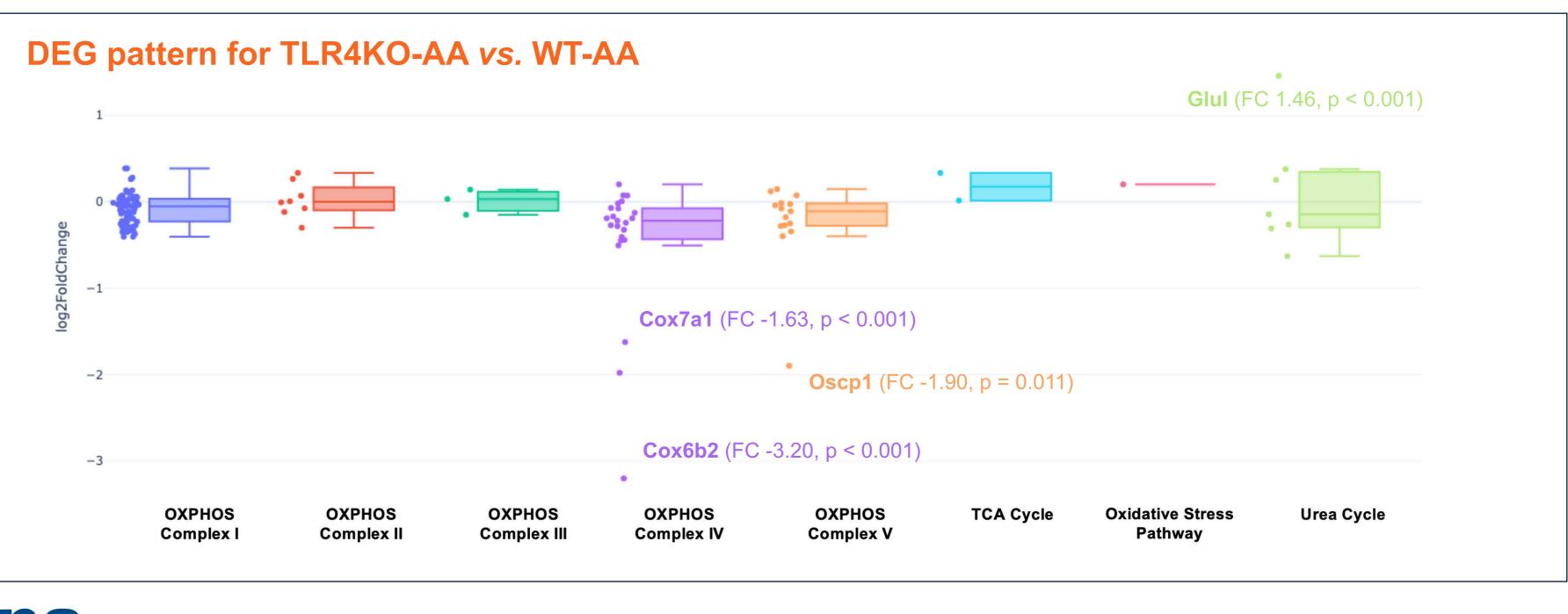
### Hyperammonemia significantly impacted



**Urea Cycle** Complex V \* Genes with significant adjusted p-value are highlighted Abbreviations: AA, amino acid; COX, Cytochrome C oxidase; DEG, differentially expressed gene; KO, knock-out; NP, normal protein; OP, Ornithine Phenylacetate; TAK242, toll-like receptor 4 inhibitor; TLR4KO, TLR4 knock-out; WT, wild-type

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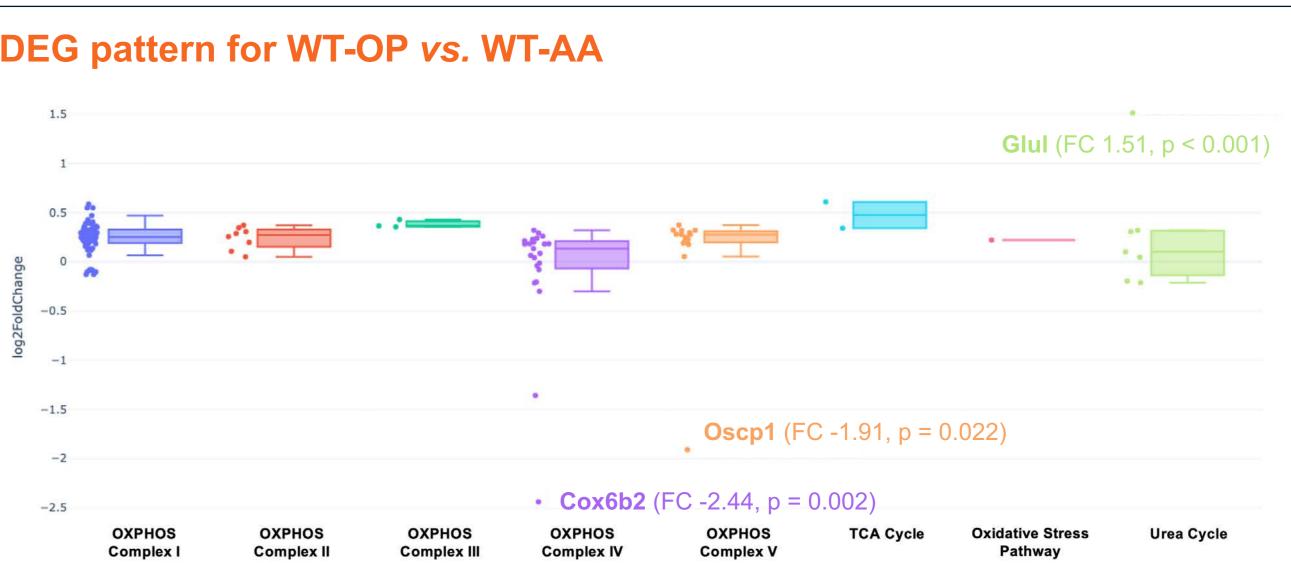
with liver diseases, where elevated ammonia onia levels is linked to improved survival. e, highlighting an unmet clinical need. or 4 (TLR4) as a potential therapeutic target for function <sup>1,2</sup> .	Both TAK-242 and OP treatment reverted Cox6b2 a downregulated in the TAK-242 group, while Oscp1 distinct mechanisms of action for each treatment. DEG pattern for WT-TAK242 vs. WT-AA
tagonist (TAK-242) and the ammonia scavenger	Glul (FC 1.49, p < 0.001)
iptomic analysis to identify pathways for targeted	<ul> <li>Cox7a1 (FC -0.76, p = 0.029)</li> <li>-1</li> <li>-1.5</li> <li>Cox6b2 (FC -1.64, p = 0.031)</li> <li>OXPHOS OXPHOS OXPHOS OXPHOS Complex II</li> <li>OXPHOS Complex II</li> <li>OXPHOS Complex II</li> <li>OXPHOS Complex II</li> </ul>
WT and TLR4KO) MT-AA mice and educed ammonia	<b>3</b> TLR4KO mice exhibited gene expression changes and OP treatment. DEG pattern for TLR4KO-AA vs. WT-AA
sing RNA sequencing on (OXPHOS)	1 and the second secon
d the OXPHOS complex IV and urea cycle.	-2 -3 OXPHOS OXPHOS OXPHOS OXPHOS Complex II Complex III Complex IV
<0.001) = 0.006)	Conclusions
• Ass1 (FC 1.15, p = 0.018) Asl (FC 0.72, p < 0.001) • • • • • • • • • • • • •	<ul> <li>TLR4 plays an important role in mediating met by hyperammonemia.</li> <li>The shared and distinct gene expression chan highlight complementary mechanisms, which on therapeutic strategies.</li> </ul>
<b>Glul</b> (FC -1.45, p < 0.001)	



### References

- . Kerbert A et al. Sci. Adv. 2025;11(10):eado1648
- 2. Sriphoosanaphan S et al, EASL Congress 2024

and Glul expression. Additionally, Cox7a1 was was downregulated in the OP group indicating



### s consistent with a combined effect of TAK-242

tabolic and mitochondrial dysfunction induced

nges observed with TAK-242 and OP treatments could be leveraged for more effective

