

# Assessment of metabolic dysfunction-associated steatohepatitis resolution index and component biomarkers in prediction of histology response to denifanstat in the FASCINATE-2 trial

Rohit Loomba<sup>1</sup>, Wen-Wei Tsai<sup>2</sup>, Laura Hover<sup>3</sup>, Shipra Vinod Gupta<sup>2</sup>, Katharine Grimmer<sup>2</sup>, Julie Dubourg<sup>2</sup>, Eduardo Martins<sup>2</sup>, George Kemble<sup>2</sup>, Marie O'Farrell<sup>2</sup> 1. Division of Gastroenterology, UC San Diego School of Medicine, CA, USA, 2. Sagimet Biosciences Inc, San Mateo, CA, USA, 3. Monoceros Biosystems LLC, CA, USA

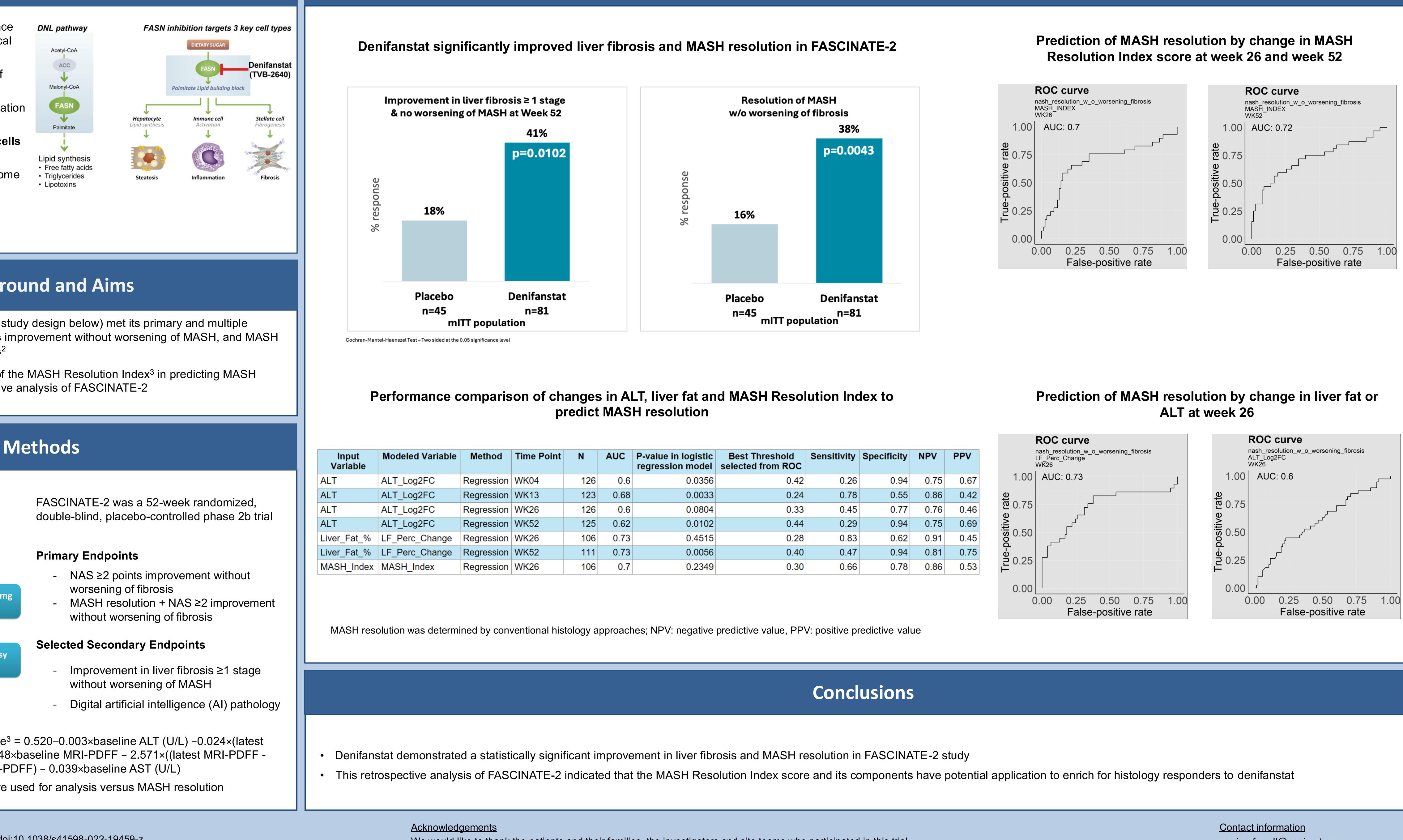
## Introduction

- Denifanstat (TVB-2640) is an oral, once daily, selective FASN inhibitor in clinical development for MASH
- FASN inhibition targets 3 hallmarks of MASH:

- inhibits liver fat synthesis & accumulation (hepatocytes)

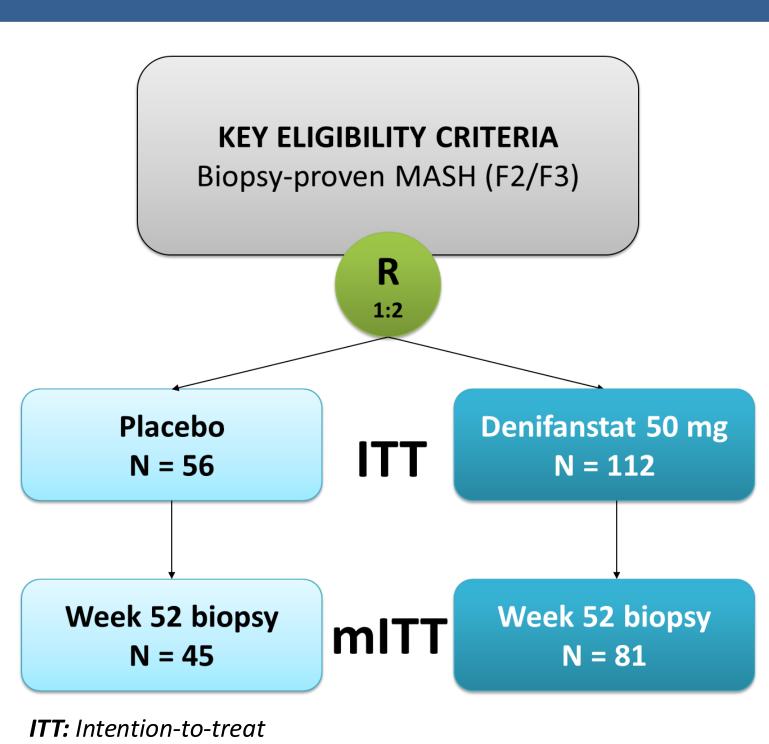
- inhibits fibrosis (hepatic stellate cells require DNL for activation)

- decreases inflammation (inflammasome activation by palmitate)<sup>1</sup>



### **Background and Aims**

- The phase 2b FASCINATE-2 trial (see study design below) met its primary and multiple secondary endpoints, including fibrosis improvement without worsening of MASH, and MASH resolution without worsening of fibrosis<sup>2</sup>
- This analysis tested the performance of the MASH Resolution Index<sup>3</sup> in predicting MASH resolution by denifanstat by retrospective analysis of FASCINATE-2



**mITT:** Modified intention-to-treat

- MASH Resolution Index (MR-I) score<sup>3</sup> =  $0.520-0.003 \times baseline ALT (U/L) 0.024 \times (latest)$ ALT [U/L]- baseline ALT (U/L)) – 0.048×baseline MRI-PDFF – 2.571×((latest MRI-PDFF baseline MRI-PDFF) / baseline MRI-PDFF) – 0.039×baseline AST (U/L)
- Logistic regression (LR) models were used for analysis versus MASH resolution

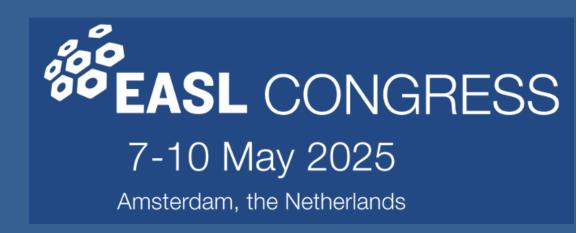
<u>References</u>

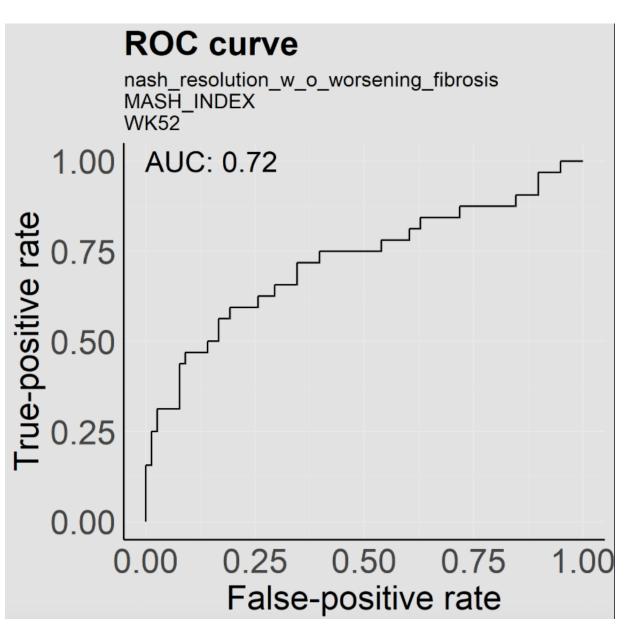
- (1) O'Farrell et al., 2022. Scientific Reports. doi:10.1038/s41598-022-19459-z
- (2) Loomba et al., 2024. The Lancet Gastroenterology & Hepatology. doi:10.1016/S2468-1253(24)00246-2
- (3) Loomba R, et al. Gut 2024;0:1–7. doi:10.1136/gutjnl-2023-331401

Input Variable	Modeled Variable	Method	Time Point	Ν	AUC	P-value in logistic regression model		Sensitivity	Specificity	NPV	PPV
ALT	ALT_Log2FC	Regression	WK04	126	0.6	0.0356	0.42	0.26	0.94	0.75	0.67
ALT	ALT_Log2FC	Regression	WK13	123	0.68	0.0033	0.24	0.78	0.55	0.86	0.42
ALT	ALT_Log2FC	Regression	WK26	126	0.6	0.0804	0.33	0.45	0.77	0.76	0.46
ALT	ALT_Log2FC	Regression	WK52	125	0.62	0.0102	0.44	0.29	0.94	0.75	0.69
Liver_Fat_%	LF_Perc_Change	Regression	WK26	106	0.73	0.4515	0.28	0.83	0.62	0.91	0.45
Liver_Fat_%	LF_Perc_Change	Regression	WK52	111	0.73	0.0056	0.40	0.47	0.94	0.81	0.75
MASH_Index	MASH_Index	Regression	WK26	106	0.7	0.2349	0.30	0.66	0.78	0.86	0.53

### Results

We would like to thank the patients and their families, the investigators and site teams who participated in this trial.





marie.ofarrell@sagimet.com