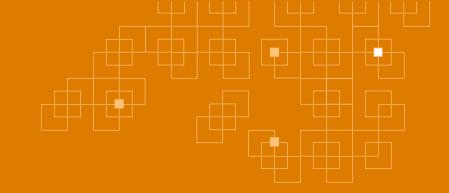


Disease Assessment Strategies to Accelerate Drug Development

https://forumresearch.org/liver-forum/liver-forum-meetings/1744-liver-forum-12



April 22-23 Meeting Theme "Show us the data"

Roadmap



- Improving the reference standard
- Opportunities & challenges
- Diagnostic COU
- Prognostic COU
- Predictive/monitoring COU



Liver Biopsy Reads: Statistical Issues

Amrik Shah Karma Statistics, LLC

Underestimation of Treatment Effect Size

- Reading Error always dilutes Treatment Effect size
 only "accurately read" slides contribute to effect size
- From published/observed kappas,

Fibrosis sensitivity of 0.7 is reasonable

Example: NASH trial setting focusing on Fibrosis endpoint Endpoint is binary, BUT "improve", "stable" and "worsen" buckets must be considered when assessing impact of reading error

Key Takeaway Learnings

- Endpoint based on Biopsy Reads has severe limitations and not appropriate for assessing drug efficacy.
- Impact of reading error CANNOT be overcome by increasing sample size

-Doubling sample size still yields same % Delta

• If forced to stick with Biopsy Reads, the dilution of effect size MUST be considered for Benefit-Risk assessment.

- Dilution may range from 30% - 60%

Digital Pathology with AI analyses of tissue slides is already used in clinical practice and trials: recent examples

- Prostate Cancer diagnosis
- Breast Cancer diagnosis
- Detection of Breast Cancer metastases in lymph nodes
- PD-L1 tumour expression
- Cardiac Allograft rejection

7

- Paige Prostate AI software Paige (FDA approved)Galen Prostate AI software Ibex (approved in EU)
- Galen Breast Solution AI model Ibex (approved in EU)
- Paige Breast Lymph Node, AI digital tool Paige - Press Release 2022
- Digital quantification with AI tool, BMS/PathAI, AACR June 22-24 2020, Poster 2017
- Diagnosis and grading in of endomyocardial biopsies J Lipkova, T Chen et al. Nature Medicine 2022:28;575–582

A Meta-Analytic Summary of NIT Applications in NASH Development

Liver Forum 'Disease Assessment Strategies to Accelerate Drug Development ' Washington DC, USA, April 2022

Professor Quentin M. Anstee PhD, FRCP

Professor of Experimental Hepatology & Honorary Consultant Hepatologist, Translational & Clinical Research Institute, Newcastle University, UK.

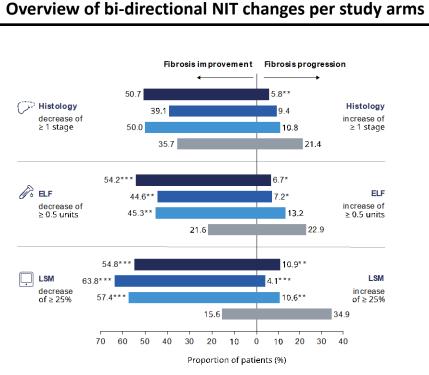




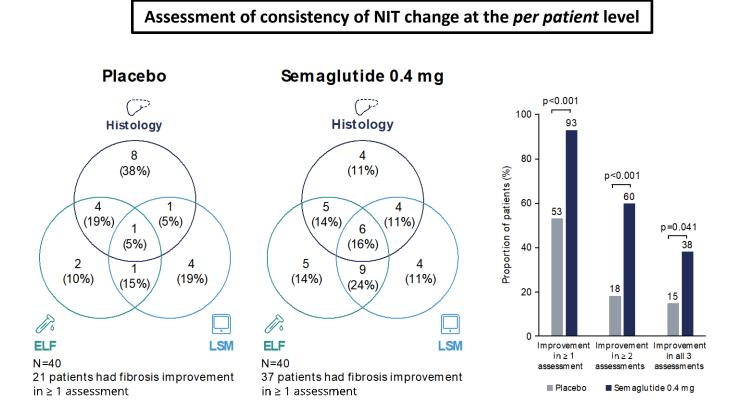
	Trial Phase	Phase IIa/IIb																							
	Year	2018	2020	2021	2021	2022	2019	2019	2021	2021	2020	2020	2021	2021	2022	2020	2016	2020	2020	2021	2015	2020	2021	2018	2020
	Duration	12	12	12	12	12	12/36	16	16	16	24	24	24	24	24/48	48	52	52	52	52	72	72	72	96	104
				: 1)			Resmetirom (MGL-3196-05)		ED)	ES IV)				(3)	Pegbelfermin (FALCON 1/2)	R)	505)		MSDC-0602K (EMMINENCE)						
				TVB-2640 (FASCINATE 1)		N-1)	GL-31	Pegbelfermin (FGF21)	Efruxifermin (BALANCED)	Saroglitazar (EVIDENCES IV)		TLAS)	Lanifibranor (NATIVE)	Aldafermin (ALPINE 2/3)	ALCO	Tropifexor (FLIGHT-FXR)	Elafibranor (GOLDEN-505)		NIMM	ST)					
		c	c	(FASC	(LIFT)	ARGO	m (M	min (F	nin (B/	ar (EV		ion (A	ior (N/	n (ALF	min (F	r (FLIG	or (GO		JZK (E	(Arre	Ê		qe	lab	TAUR)
		Aldafermin	Aldafermin	-2640	Tern-101 (LIFT)	EDP-305 (ARGON-1)	metiro	belfer	ıxifern	oglitaz	Cilofexor	Combination (ATLAS)	ifibrar	afermi	belfer	pifexo	ibranc	Seladelpar	90-DC	Aramchol (ARREST)	OCA (FLINT)	Emricasin	Semaglutide	Simtuzumab	CVC (CENTAUR)
		Ald			Teri	EDP	Res				Cilo		Lan	Ald	Peg			Sela	MSI	Ara					C C C
Clinical Chemistry	ALT & AST	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
Simple Scores & Indirect Biomarkers	APRI									\checkmark	\checkmark				\checkmark				\checkmark						\checkmark
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	СК18			\checkmark			\checkmark			\checkmark	\checkmark	\checkmark	\checkmark					\checkmark	\checkmark			\checkmark	\checkmark		
Direct Biomarkers	ELF	\checkmark	\checkmark	?			\checkmark		\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark				\checkmark	\checkmark			\checkmark	\checkmark	\checkmark
	Fibrotest										\checkmark						\checkmark		\checkmark					\checkmark	
Fibroscan	PRO-C3	\checkmark	\checkmark	\checkmark			\checkmark	\checkmark	\checkmark				\checkmark	\checkmark	\checkmark										\checkmark
	VCTE								\checkmark	\checkmark	\checkmark	\checkmark			\checkmark								\checkmark		
	FAST											\checkmark											\checkmark		
MR	MR-PDFF	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark		\checkmark	\checkmark	\checkmark		\checkmark		\checkmark					
	MR-cT1		\checkmark		\checkmark																				
	MRE							\checkmark			\checkmark	\checkmark			\checkmark										
Liver Biopsy	Histology		\checkmark	\checkmark			\checkmark		\checkmark				\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark

Research™

Wholistic Assessment of Biomarker Response



Semaglutide 0.1 mg Semaglutide 0.2 mg Semaglutide 0.4 mg Placebo



1. Anstee et al, Fibrosis response assessed by enhanced liver fibrosis and FibroScan liver stiffness measurement in patients with non-alcoholic steatohepatitis treated with subcutaneous semaglutide. EASL 2021.

A range of tractable biomarkers are available to support drug development in NASH.

When considering the use of biomarkers in drug development, it is essential to consider the specific **Context of Use** (and population/setting) that is being addressed.

There remains a need for more sensitive and specific, independently validated and qualified biomarkers for use in NAFLD drug development.

Progress to date:

- 1. Diagnostic CoU: LITMUS and NIMBLE are bringing clarity and objectivity to biomarker performance still room for innovation to identify better biomarkers.
- 2. Pharmacodynamic/Response CoU: remains an area where there is a lack of consistency but the RCTs to date have helped generate important data to support biomarker utility.
 We are now better placed to build consensus and standardise NIT selection and the consistency of NIT reporting as

We are now better placed to build consensus and *standardise NIT selection* and the *consistency of NIT reporting* as study endpoints to support NASH drug development.

3. Surrogacy potential: It is notable that there is an expanding dataset to demonstrate that fibrosis biomarkers have prognostic value and change with progression/regression of disease, potentially with greater inter-test consistency than histology.

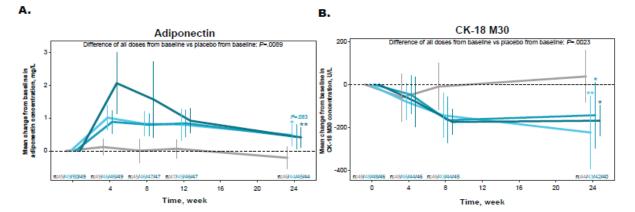
Liver forum 12 Effects of pegbelfermin on NIT's for NASH in a Phase 2b study: Falcon-1

April 23rd 2022

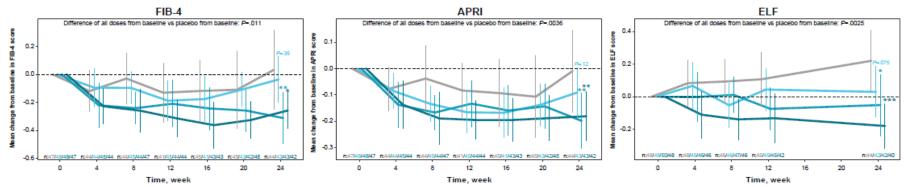
Anne Minnich



PGBF modulates blood biomarkers of liver injury and fibrosis





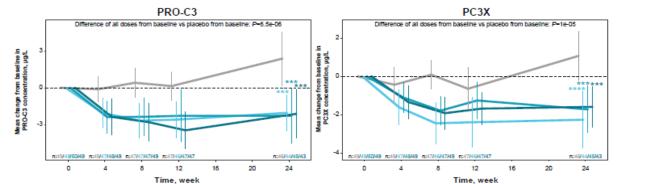


Placebo
10 mg PGBF

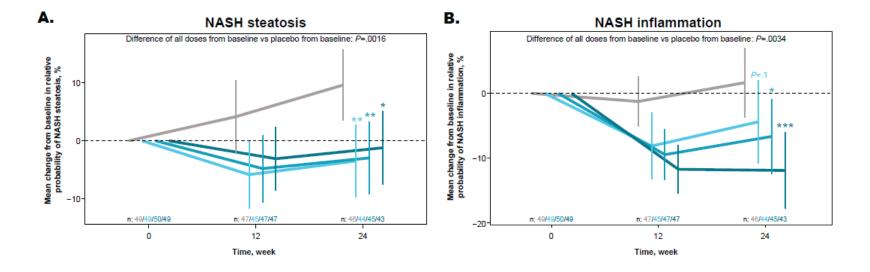
20 mg PGBF

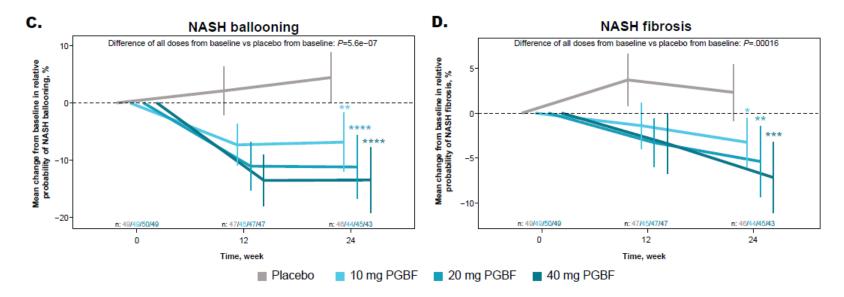
40 mg PGBF





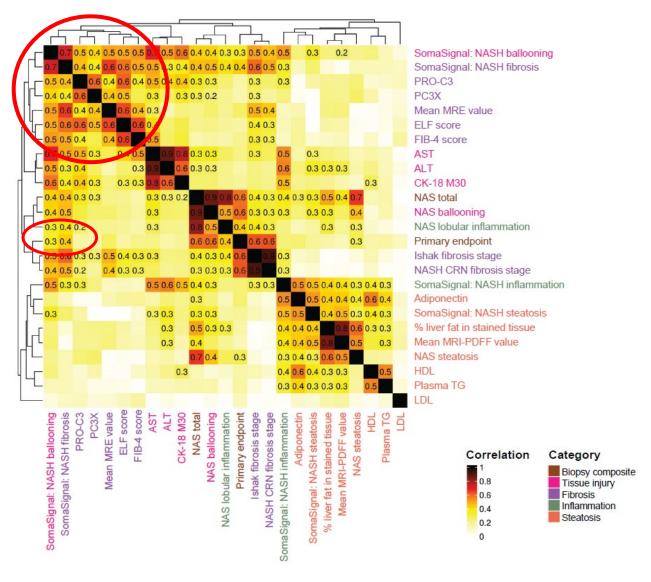
Somasignal NASH bundle





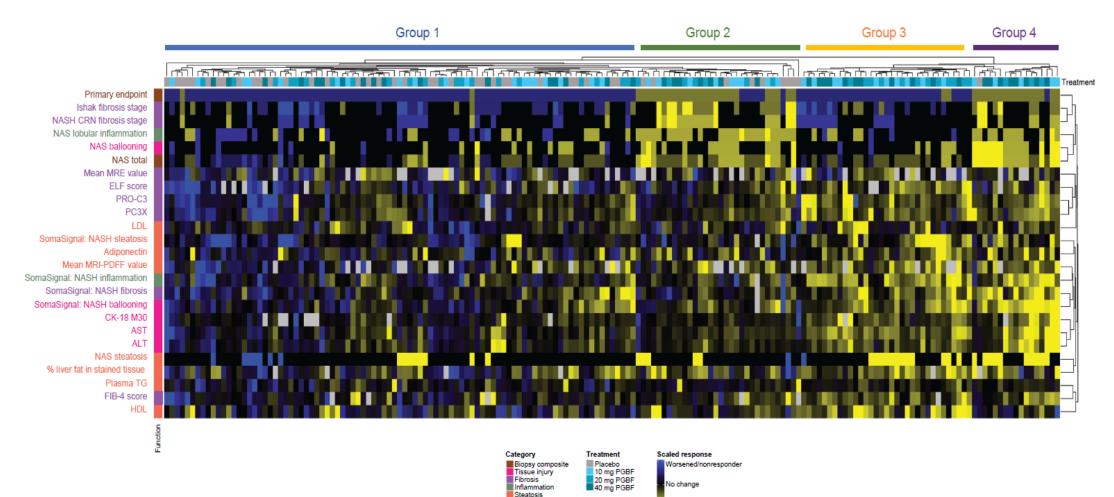
Clustering of correlation coefficients for biomarkers and histological assessments

Pairwise week 24 correlation



Fibrosis NIT's cluster well together but not with the primary endpoint Soma-NASH = only NIT to cluster with primary endpoint

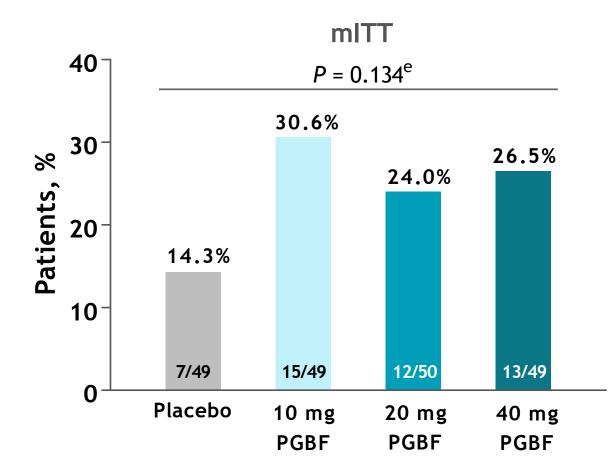
Concordance analysis between primary endpoint and week 24 biomarker responses



Improved/responder

Primary Endpoint

At Week 24: \geq 1 stage improvement in fibrosis^a without worsening of NASH^b <u>OR</u> NASH improvement^c without worsening of fibrosis^d



^aImprovement of fibrosis = \geq 1 stage decrease in NASH CRN fibrosis score; ^bWorsening of NASH = increase in NAS by \geq 1 point; ^cNASH improvement = \geq 2 point decrease in NAS with contribution from > 1 NAS component; ^dWorsening of fibrosis = \geq 1 stage increase in NASH CRN fibrosis score; ^eCochran-Armitage trend test across proportions of responders in the treatment groups at a 1-sided 0.05 level of significance provided at least 80% power if 160 patients were randomized 1:1:1:1. mITT, modified intent-to-treat; NASH, nonalcoholic steatohepatitis; PGBF, pegbelfermin.

Next steps

- Today's Session 2
- Ongoing, iterative discussions (plenary)
- Working Group (focused)
 - Compile/synthesize data for each COU
 - Present the case



