

NASH Globally- *Pathways to* combination therapies, biomarkers and outcomes

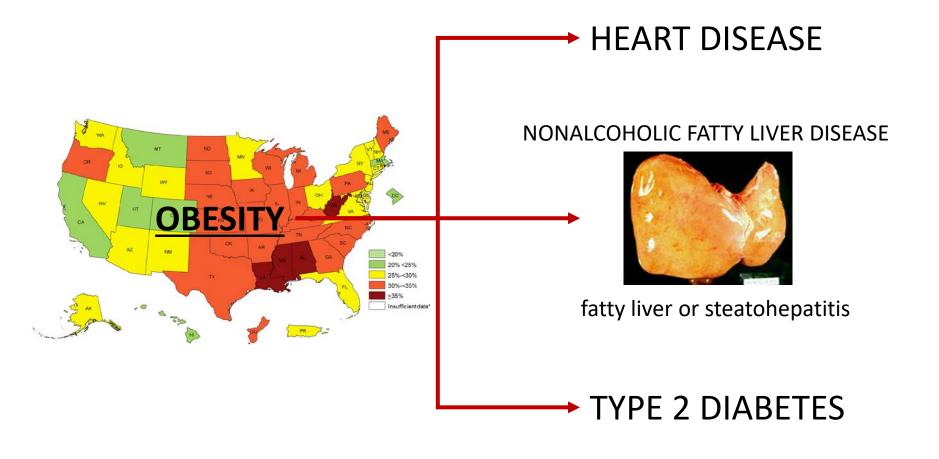
Arun J. Sanyal M.B.B.S., M.D.

Professor of Medicine, Physiology and Molecular Pathology Virginia Commonwealth University School of Medicine

Conflicts of Interest

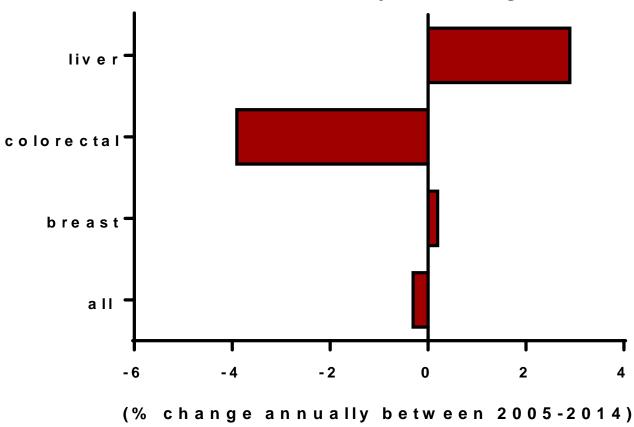
- President, Sanyal Biotechnologies
- Stock options: Genfit, Akarna, Tiziana, Indalo, Durect, Exhalenz, Hemoshear
- Advisor with compensation: Lilly, Pfizer, Novartis, Ardelyx, Salix, Hemoshear
- Advisor without compensation: Galectin, Intercept, Merck, Bristol Myers, Immuron, Gilead, Chemomab, Affimmune, Protalix, Nitto Denko, Novo Nordisk, Cirius, Boehringer Ingelhiem
- Grants to institution: Gilead, Tobira, Allergan, Merck, Bristol Myers, Astra Zeneca, Immuron, Intercept, Novo Nordisk, Shire, Boehringer Ingelhiem, Cirius

NAFLD: a silent killer in our midst



NAFLD is driving the national increase in liver cancer

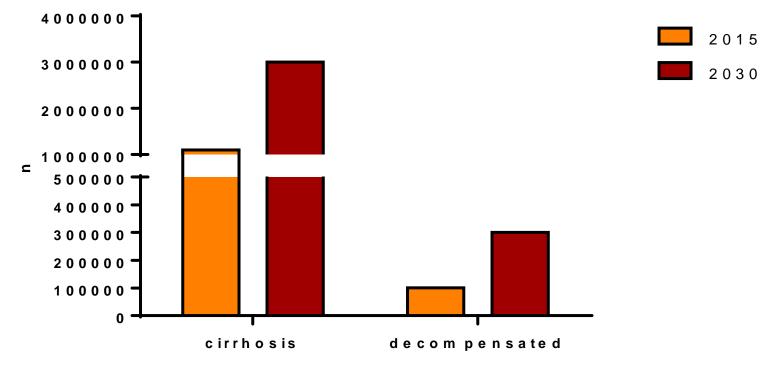
liver cancer rate related to obesity is increasing at 3% annually



Steele et al, MMWR, October 2017

The consequences of inaction will be serious:

- number of those with cirrhosis will triple
- over 300,000 people will have end-stage liver disease
- many of these will be "todays" children



liver outcom es

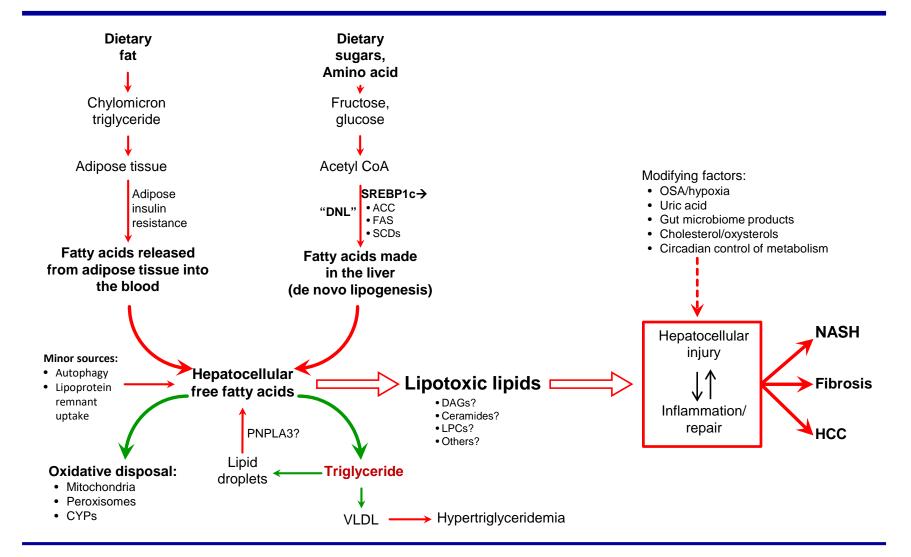
Estes et al, Epub Hepatology, 2017

Making the case for combination therapies

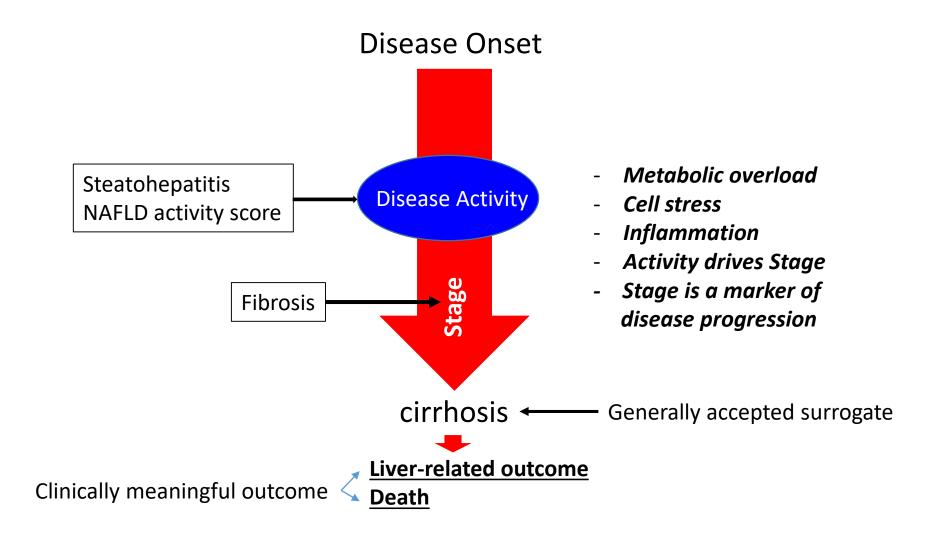
NASH is a disease of metabolic substrate poisoning



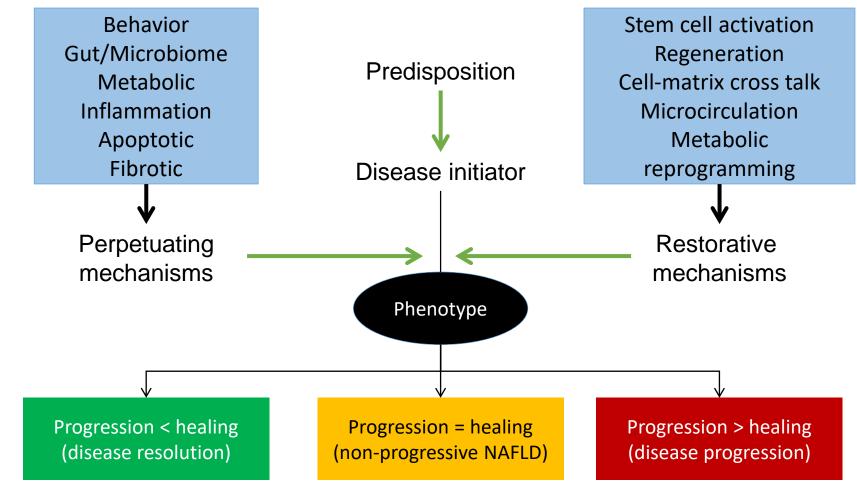
Pathogenesis of NASH and targets of therapy



Disease activity versus disease stage



The progression of NASH is affected by many pathways



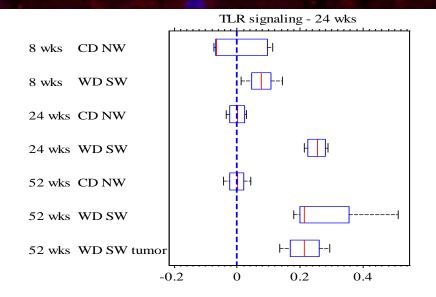
Ratziu V, Goodman Z, Sanyal A. Journal of Hepatology 2015;62;S65–S75.

NAS vs Fibrosis Exclusivity in NASH Pathways

	•	•	Cytochrome c-mediated apoptotic responsel Reactome				NAS.Prob
l I		•	Activation of caspases through apoptosome-mediated cleavage	gelReactome			1
L I	*	•	Formation of apoptosomelReactome			0.2	-
L	•	•	p38MAPK eventslReactome				0
dr 📃	•		The AIM2 inflammasomelReactome		Inflammatior	0.1	0
[JL	*		Apoptotic factor-mediated responselReactome		IIIIaIIIIIatioi	0.1	FS.Prob
Πι	•		The IPAF inflammasomelReactome				
	•		Release of apoptotic factors from the mitochondrialReactome		apoptosis	0	0.78
	*		Loss of Function of TGFBR2 in CancerlReactome			Ŭ	
	*		Loss of Function of TGFBR1 in CancerlReactome				0
	•		TGFBR1 KD Mutants in CancerlReactome			-0.1	
	•		TGFBR2 Kinase Domain Mutants in CancerlReactome				
	•		SMAC-mediated apoptotic responselReactome				
_ L	•		Intrinsic Pathway for Apoptosis Reactome			-0.2	2
		•	Synthesis, secretion, and inactivation of Glucose-dependent In	nsulinotropic F	Polypeptide (GIP)IReactome		
	•	•	The NLRP3 inflammasomelReactome				
	•	•	InflammasomesIReactome				
┍┥╽╭┠╴			ECM proteoglycansiReactome	Int	lammasome	The ex	clusivity
		•	Transcription from mitochondrial promotersIReactome				
		•	Mitochondrial transcription initiationIReactome	EC	N/I	probab	ility is the
		*	Synthesis of dolichyl-phosphate-glucoselReactome			nrohah	ility that the
	*		MAPK targets/ Nuclear events mediated by MAP kinasesIReact	ctome			
			RNA Polymerase I, RNA Polymerase III, and Mitochondrial Tran	unscription Rea	actome	patnwa	iy is significant in
			Glucose metabolismlReactome			one coi	ntrast and not in
r	*		TRAF6 Mediated Induction of proinflammatory cytokinesIReact	tome			
i di	•		G beta:gamma signalling through PI3KgammalReactome			the oth	er
I I	•		Apoptotic execution phaselReactome				
	*		Caspase activation via extrinsic apoptotic signalig pathwaylRea	actome			
	*	*	TGF-beta signaling pathwaylKEGG				
[[•	Focal adhesionIKEGG				
		•	MAPK6/MAPK4 signalinglReactome				
1	*		ApoptosislReactome				
	*		Signaling by TGF-beta Receptor Complex in CancerlReactome	ie			
	•	•	ApoptosisIKEGG				
	•		Transcriptional activation of mitochondrial biogenesislReactome	ie	P =	P *	$(1 - P_{B})$
1 1	*	*	Negative regulation of MAPK pathwaylReactome		' EA —	' A	
1 1 1	*		TAK1 activates NFkB by phosphorylation and activation of IKKs	s complexiRe	actome		
	•		Apoptosis induced DNA fragmentationIReactome				
	•		alpha-linolenic (omega3) and linoleic (omega6) acid metabolisi	smlReactome			
	*		activated TAK1 mediates p38 MAPK activation Reactome				
194	*		JNK (c–Jun kinases) phosphorylation and activation mediated	by activated	human TAK1IReactome		
	•		DEx/H-box helicases activate type I IFN and inflammatory cyto	okines produc	tion IReactome		
		*	p130Cas linkage to MAPK signaling for integrinslReactome				
	*		Recycling of bile acids and saltslReactome				
──┤┎	·		Linoleic acid metabolismIKEGG				
	•		Bile salt and organic anion SLC transporters Reactome				
	*	*	Sialic acid metabolismlReactome				
ل ال	*		Bile acid and bile salt metabolismlReactome				
L	*		Sulfur amino acid metabolismlReactome				
	NAS	Fibrosis.stage					

Sanyal lab, unpublished data 11

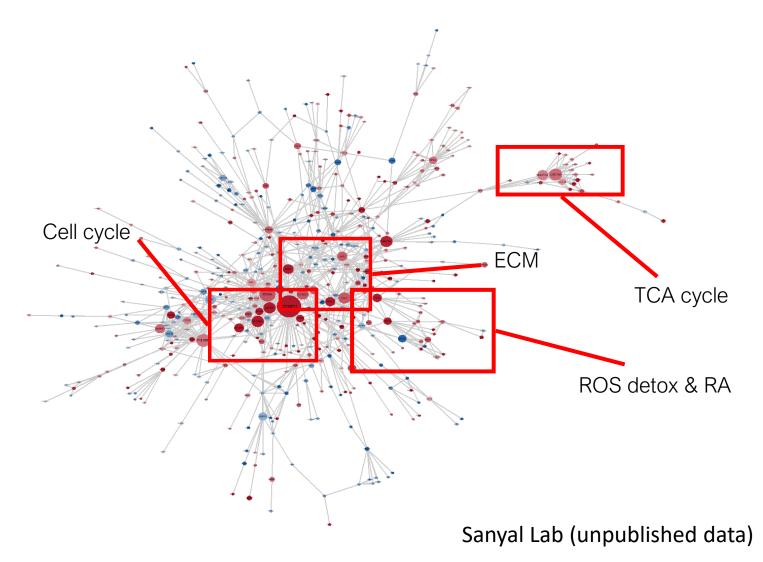
K. Human NASH (NAS 5)



Cazanave et al, Scientific Reports, Epub Dec 2017

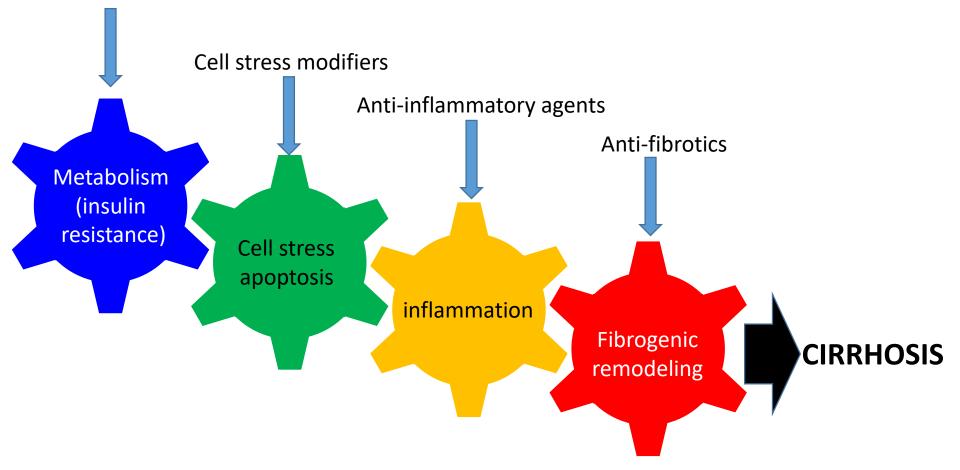
50um

Pathways with super-additive effects on disease activity and fibrosis stage

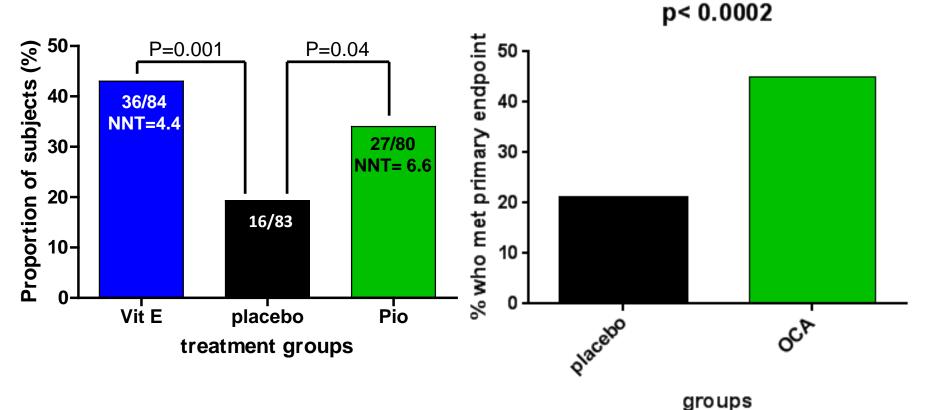


DISEASE BIOLOGY PROVIDES TARGETS FOR THERAPEUTICS

Insulin resistance modifiers

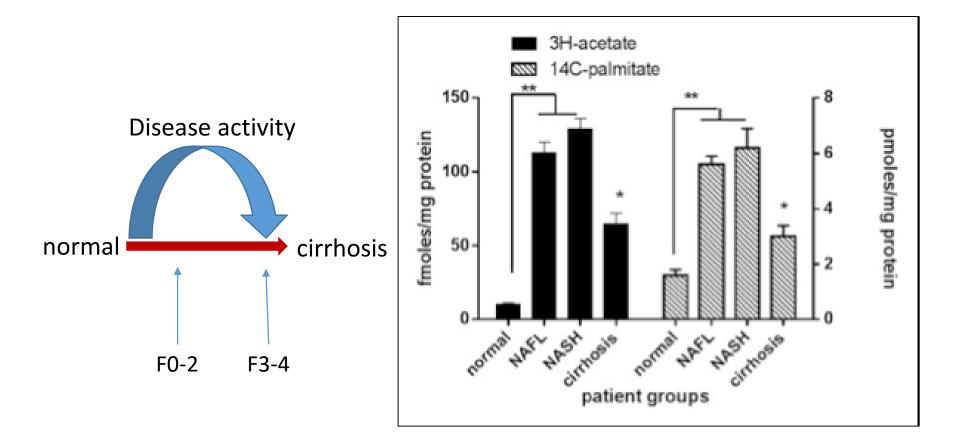


If everyone took the drug, why did only some individuals improve?



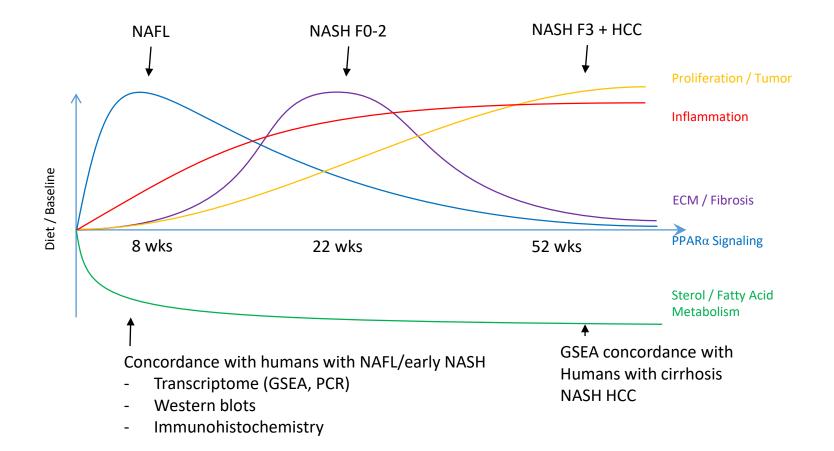
Sanyal et al, NEJM 2010, Neuschwander-Tetri, Lancet 2015

Disease activity burns out with progression in to cirrhosis

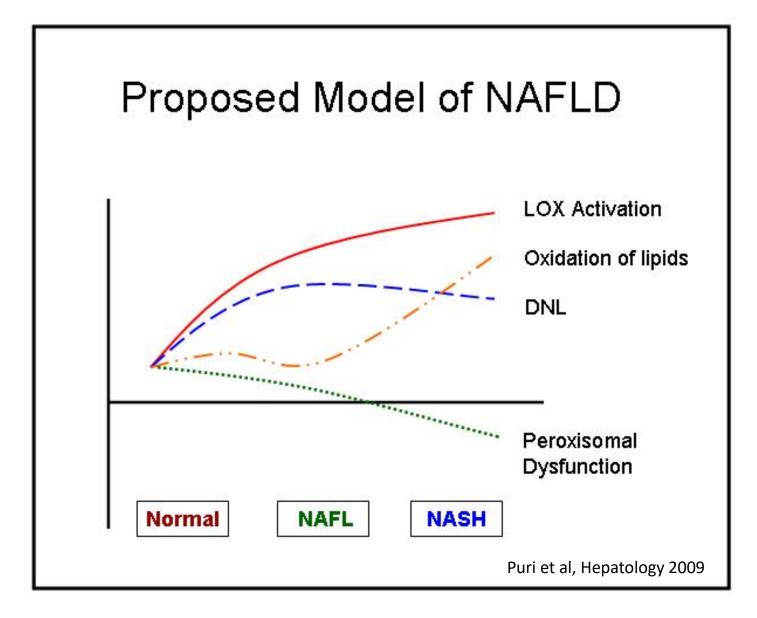


Siddiqui et al, Clin Gastro Hep 2015

Tracking the molecular evolution of NASH provides a comprehensive list of potential targets for therapy

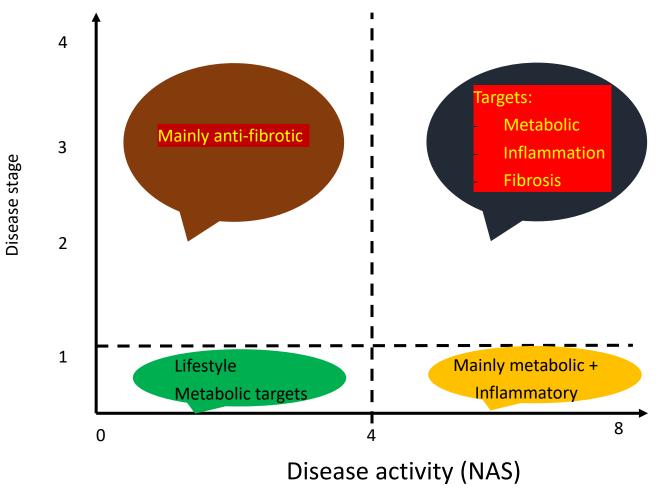


Lipidomic signature of NASH



Matching the right patient to the right drug/s

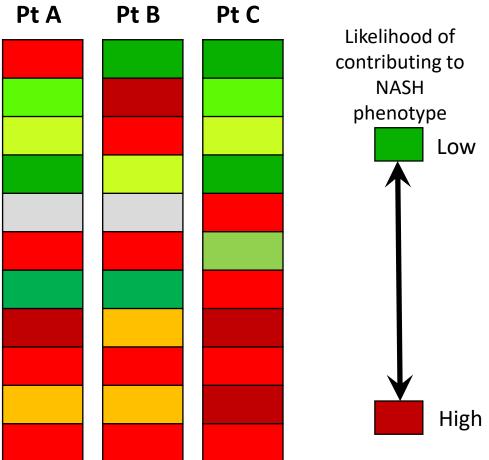
Rational approach to therapeutics for NASH



Hypothetical patient stratification

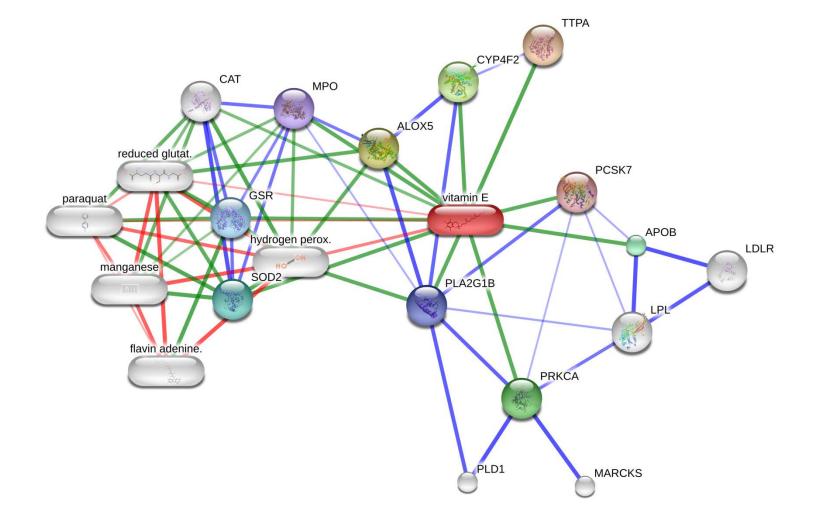
Pathway category:

Impaired satiety mechanisms Impaired thermogenesis Periph adipogenesis/lipolysis Adipose inflammation Augmented DNL Impaired TG formation PNPLA3? \rightarrow Inappropriate TG lipolysis Active lipotoxic lipid synthesis Liver inflammatory pathways Impaired wound response Augmented fibrogenesis



"lean NASH"

Network analysis reveal vitamin E specific pathways that are relevant for its effects on NASH



Sookoian and Pirola, Clin Liv Dis, 2012

Baseline metabolites predict response to future treatment with vitamin E

Metabolites	OR	95% CI
gamma-CEHC	0.11	0.01-0.995
2-palmitoylglycerophosphoethanolamine	0.08	0.01 - 0.56
myristoleate (14:1n5)	0.04	0.002-0.64
3-phenylpropionate	29.4	1.23-707.0
Asparagines	20.2	1.2-338.6
indolepropionate	16.2	1.45-180.7

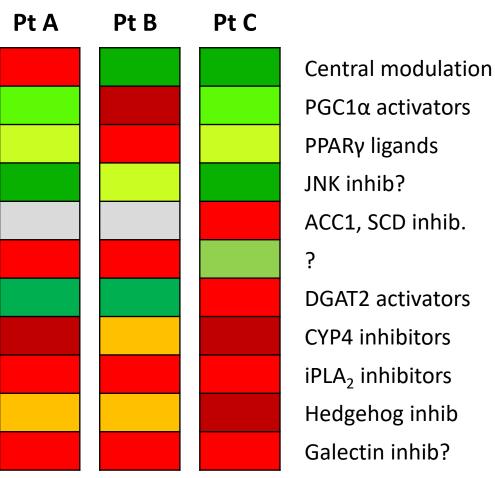
Only those that were significant are listed

Cheng et al, PlosONE, 2012

Stratification → targeted treatment "Personalized medicine"

Pathway category:

Impaired satiety mechanisms Impaired thermogenesis Periph adipogenesis/lipolysis Adipose inflammation Augmented DNL Impaired TG formation PNPLA3? \rightarrow Inappropriate TG lipolysis Active lipotoxic lipid synthesis Liver inflammatory pathways Impaired wound response Augmented fibrogenesis

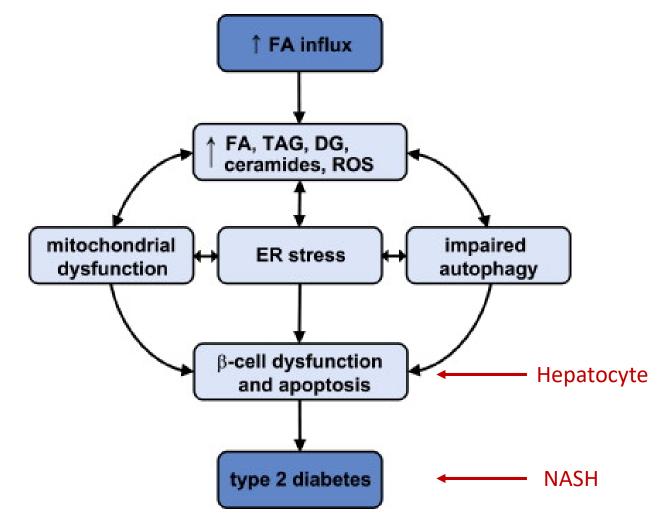


"lean NASH"

September 12, 2017

Combination therapy: targeting multiple organs simultaneously

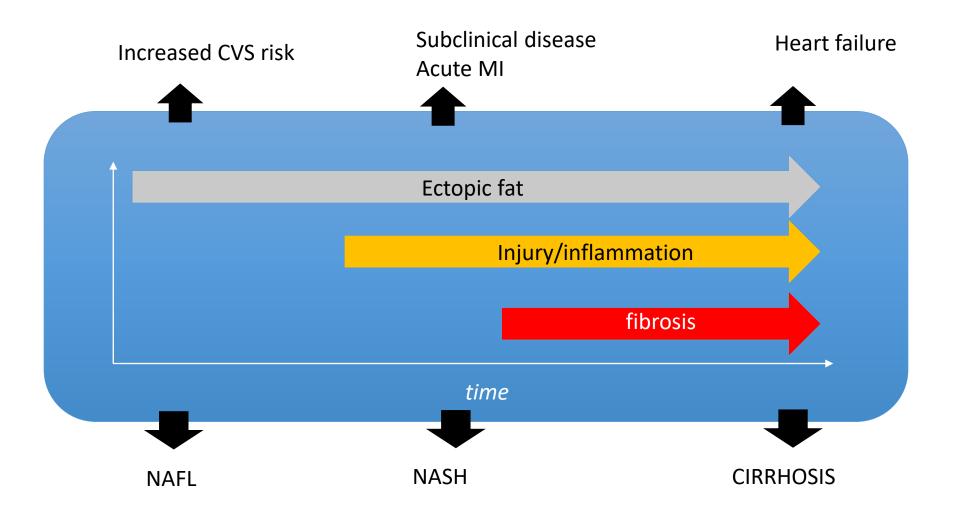
Pathogenesis of beta cell failure in type 2 diabetes



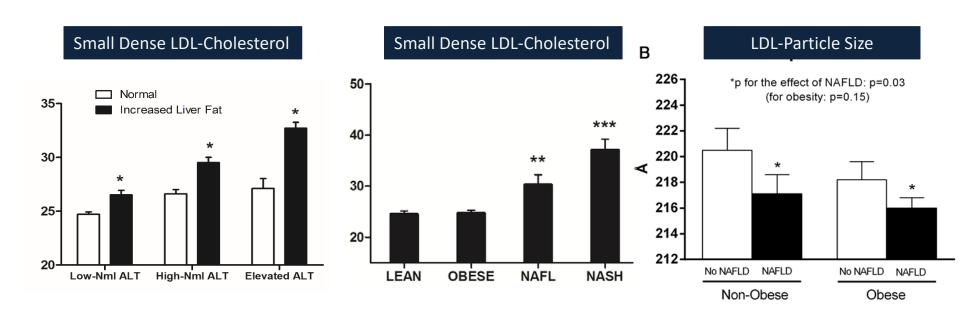
Janikiewicz, et al; Biochemical and Biophysical Research Communications, Volume 460, Issue 3, 2015, 491–496

http://dx.doi.org/10.1016/j.bbrc.2015.03.153

The Liver-Heart connection



Disease Severity in NAFLD Drives Atherogenic Dyslipidemia

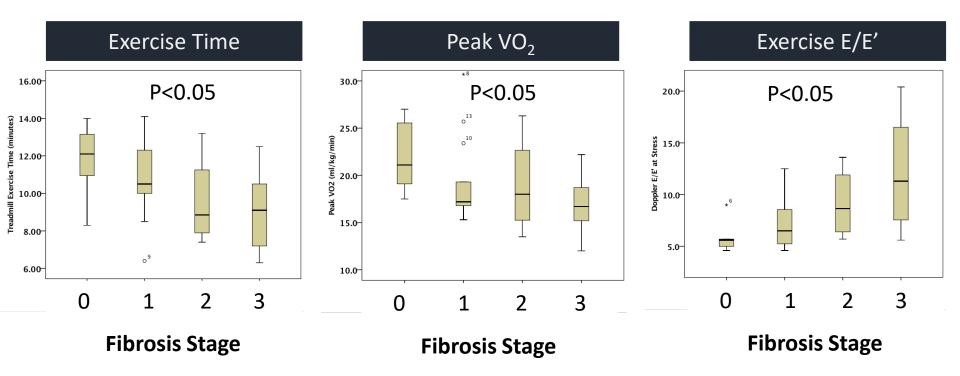


The more advanced the NASH, the greater the risk of cardiac events

		Age, Sex-adjusted	Multivariable-adjusted
	n	Hazard ratio (95% CI)	Hazard ratio (95% CI)
All cause	778		
Minimal	251	1	1
Intermediate	404	1.50 (1.20-1.88)	1.40 (1.09-1.81)
Advanced	123	2.26 (1.59-3.21)	1.80 (1.23-2.64)
Cardiovascular disease	296		
Minimal	81	1	1
Intermediate	167	2.43 (1.69-3.50)	2.49 (1.71-3.64)
Advanced	48	3.34 (2.00-5.60)	3.22 (1.92-5.42)

Kim et al, Hepatology, 2013

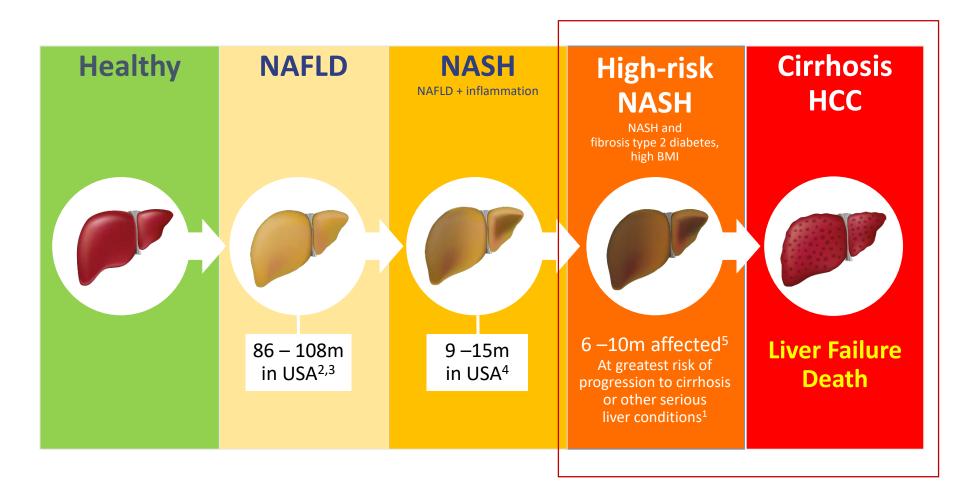
Fibrosis Stage is Linked To Diastolic Dysfunction and Exercise Capacity



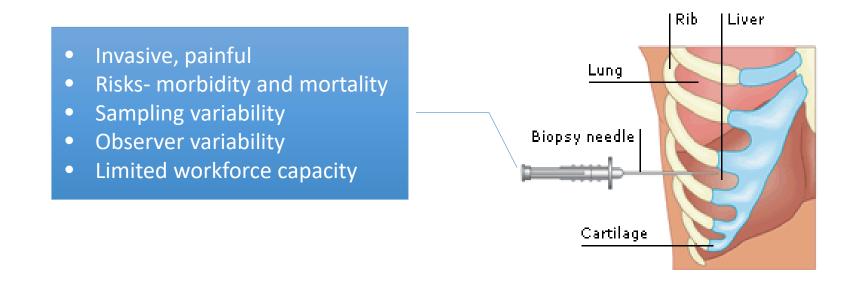
Siddiqui et al. AASLD 2017

Finding the patients- an urgent need to develop noninvasive methods for assessment

Targeting the population at risk



Liver biopsy is an inadequate tool for routine assessment



With a mortality risk of 1:1000 and population at risk of 60 million, the total number of Diagnostics-associated mortality would be 60000



Biomarker development process

Drug Approval Process

Scientific Community Consensus

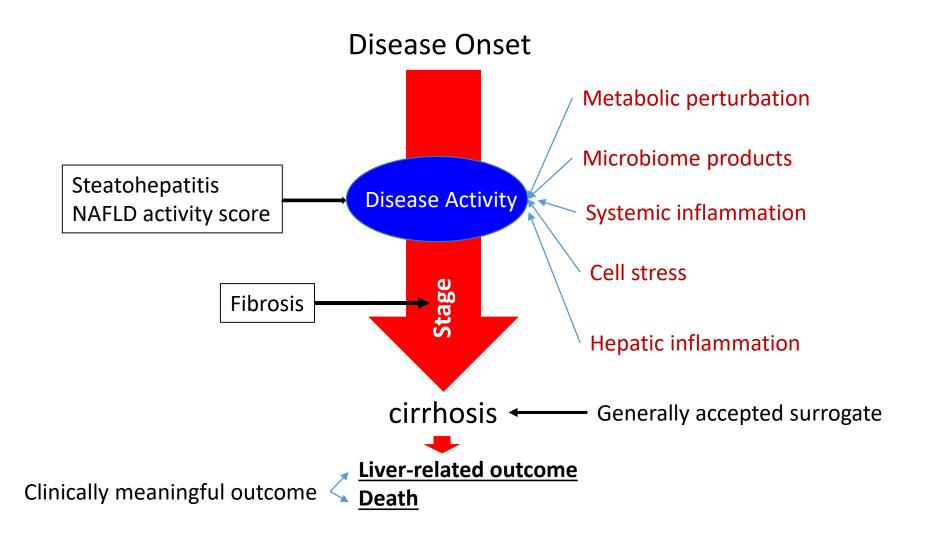
• Data Driven

- Subject to regulatory scrutiny
- More than one process can go on
- Liver Forum integrates biomarker development process across FDA and EMA

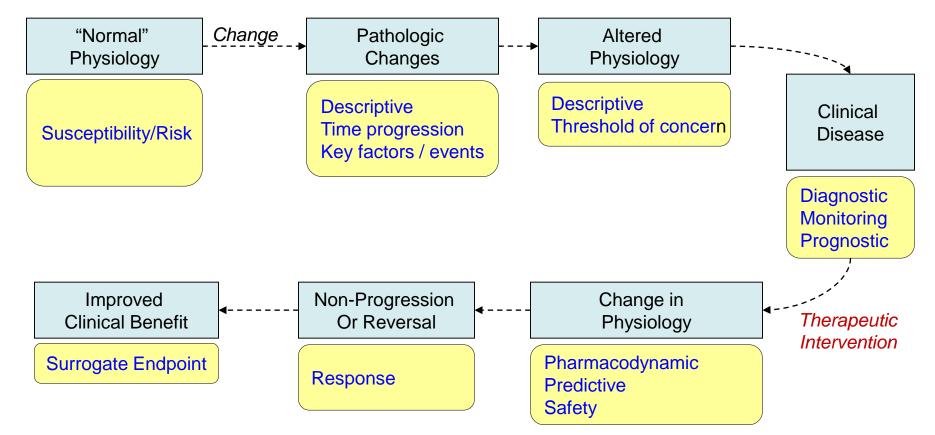
Biomarker Qualification Program

Facilitating Biomarker Development: Strategies for Scientific Communication, Pathway Prioritization, Data-Sharing, and Stakeholder Collaboration; Published June 2016, Duke-Margolis Center for Health Policy

Disease activity versus disease stage



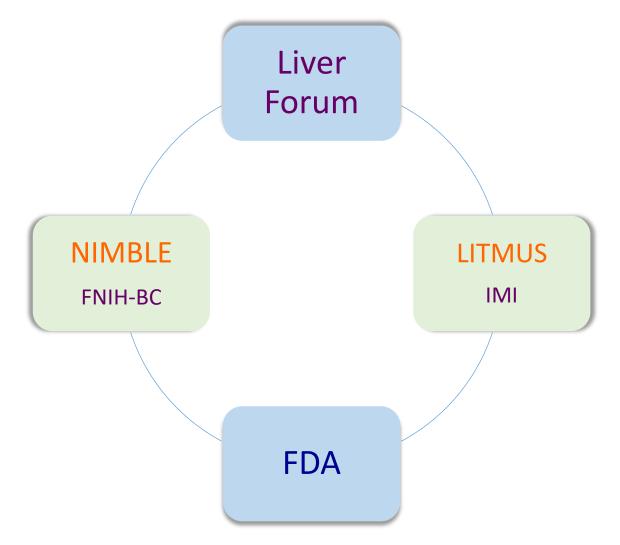
"Fit for Purpose" biomarkers



Companion vs Complementary diagnostic

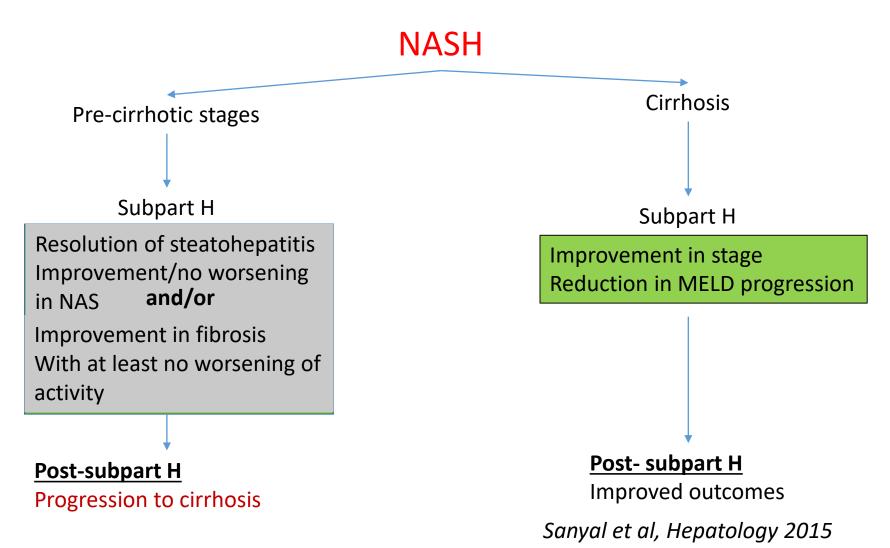
Adapted from Chris Leptak...Liver Forum Biomarker Workshop 2017

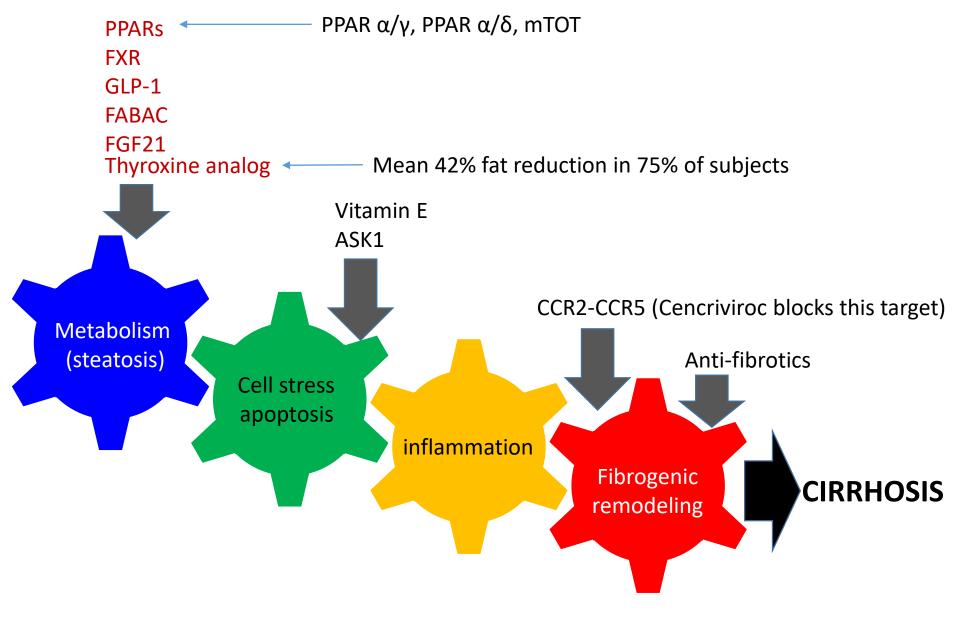
Trans-atlantic initiatives for NASH biomarker development

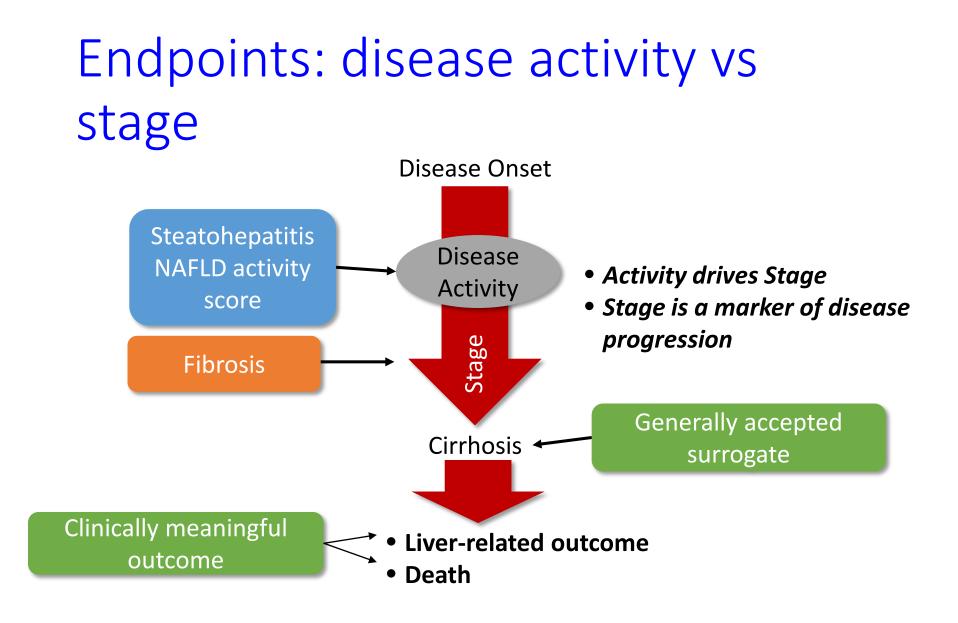


The path to approval

Evidence burden to have therapy approved for NASH





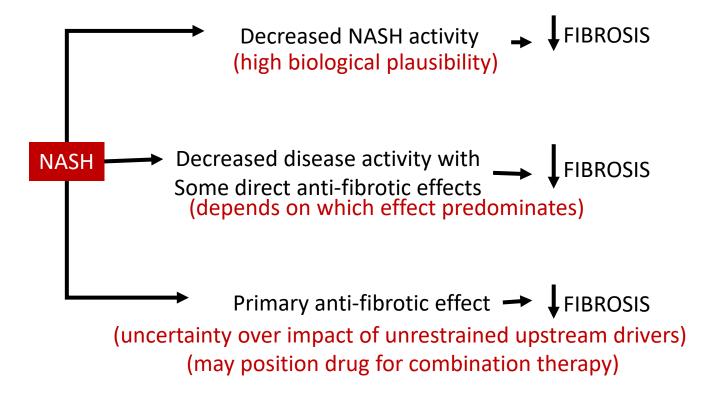


In pre-cirrhotic stages, fibrosis is relevant mainly as a marker of *disease progression* towards cirrhosis

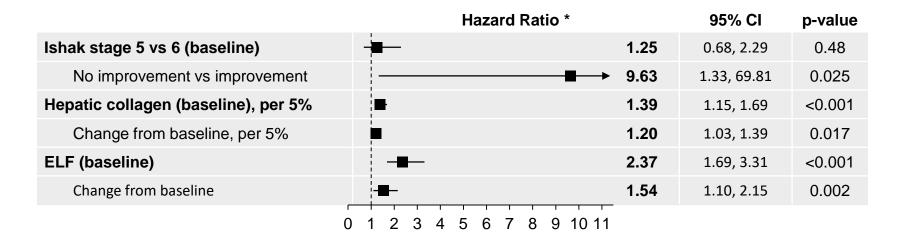
Progression to cirrhosis is a generally accepted surrogate endpoint for approval



Disease progression includes metabolic reprogramming, cell death, stem cell recruitment, regenerative activity, cell differentiation, changes in microcirculation, matrix, bile flow. Fibrosis is an easily visible and quantifiable surrogate for this process. Implications of decreased fibrosis in precirrhotic stages of NASH is linked to drug mechanism of action



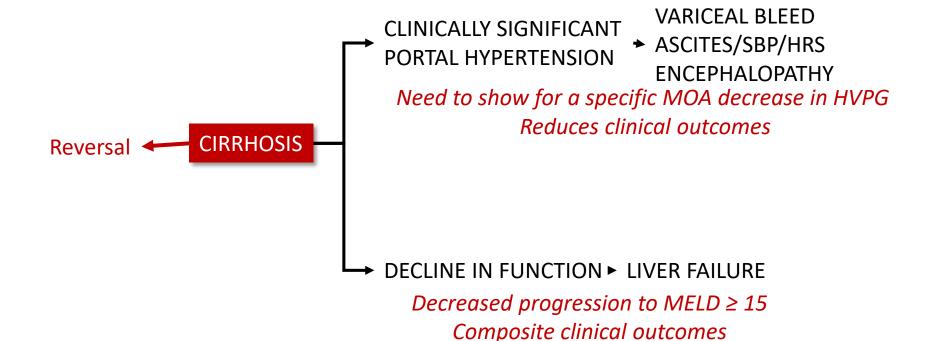
Impact of Fibrosis on Clinical Events



- Increased risk of clinical events with:
 - Higher baseline hepatic collagen content and ELF
 - Worsening of fibrosis (by Ishak stage, collagen content, ELF)

Sanyal et al, EASL 2017

How cirrhosis leads to clinically meaningful outcomes



MELD as an endpoint

Pros	Cons
 Relates to mortality Well known to clinicians Widely available Easy to measure Threshold value of 10 or 14 identifies a important stage in clinical course 	 Inter-lab variability Related to 3 month mortality Rate of progression of MELD score not linear Most patients with compensated cirrhosis have a MELD < 10

Increase in MELD to \geq 15 represents a point in course of disease where Transplant should be considered

Take home messages

- NASH is a clinical syndrome driven my metabolic substrate overload to the liver.
- The biology of NASH has significant collinearity with the biology of HFPEF and type 2 diabetes
- Integrated approaches to noninvasive assessments that provide a read out of disease activity and stage in key end organs is needed.
- Therapeutics should go after nodal targets that are key for disease development and progression. Combinations should be rational and based on proper step wise clinical development.
- Trial design innovations are under way to allow accelerated assessment of combination therapies to improve clinical outcomes

上医医末病之法 医医将病之: 医医已病之病 首帝: 肉経 Source: JACC © 2010 American College of Cardiology Foundation

Huang Dee: Nai-Ching (2600 BC, First Medical Text)

Translation:

Superior doctors prevent the disease Mediocre doctors treat the disease before evident Inferior doctors treat the full-blown disease

Acknowledgements

VCU

- AJS Lab:
 - Farid Mirshahi
 - Sophie Cazanave
 - Divya P. kumar
 - Abdul Oseini
 - Bubu Banini
 - Robert Vincent
- AJS clinical:
 - Rebecca Collen
 - Mohammed Siddiqui
 - Sherry Boyett
 - Joel Steinberg
 - Robert Cadrain
- VCU cardiology:
 - Justin Canada
 - Stefano Toldo
 - Eleanora Toldo

Non-VCU

- Oxford Univ:
 - Oliver Rider
 - Ines Abdesalaam
 - Stefan Nebauer
- Perspectum
 - Keri Hildick
- Vedic Science Analytics
 - Sri Koduru
- Hemoshear:
 - Ryan Feaver
 - Brian Wamhoff
 - Bhanu Cole
 - Steve Hoang
 - Mark Lawson