# HEPION PHARMACEUTICALS

Pleiotropic Actions of the Cyclophilin Inhibitor, CRV431, in Preclinical Models

NASHSummit DIGITAL

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### **NASH Clinical Trial Outcomes**



What can we learn from placebo and treatment responses in NASH trials?

### High placebo responses:

• malleable, regenerative liver? (but why low treatment responses?)

### Low treatment responses:

- wrong targets?
- too few targets?

### **THESIS**

NASH livers will benefit most by targeting multiple pathophysiologic pathways.



# **Cyclophilins**

Pleiotropic effects arising from multiple cyclophilin isoforms

- Enzymes that accelerate proline peptide bond *cis-trans* isomerization
- *cis-trans* switching = regulation of protein interactions and functions
- 17 human isoforms in all major cellular compartments and tissues
- 30 years of research and 4000+ papers

EXPERT OPINION ON INVESTIGATIONAL DRUGS (2019) Cyclophilin inhibition as a potential treatment for nonalcoholic steatohepatitis (NASH). Daren R. Ure, Daniel J. Trepanier, Patrick R. Mayo and Robert T. Foster

### CYCLOPHILIN(s) DELETION OR INHIBITION ATTENUATES DISEASE PROCESSES IN MANY EXPERIMENTAL MODELS metabolism, inflammation, necrosis, fibrosis, carcinogenesis, virulence, etc.

ischemia-reperfusion injuries • myocardial infarction • stroke • chronic asthma • arthritis •
atherosclerosis • acute pulmonary embolism • thrombosis • aortic aneurysm • coronary artery disease
pulmonary arterial hypertension • ALS • Alzheimers disease • multiple sclerosis • muscular
dystrophies • traumatic CNS injury • tumorigenesis • metastasis • viral hepatitis

Proline Peptide Bond cis-trans Isomerization





# **CRV431 Modes of Action**

CRV431 blocks cyclophilin participation in several disease processes



CD147 pro-inflammatory receptors









# MetaCore Biological Systems Database (2.6M+ curated interactions)

CRV431 predicted to inhibit diversity of cyclophilin-protein interactions

Target	Туре	Total protein and gene interactions	Protein binding interactions	Transcription regulated by "#" genes/proteins	Regulates expression transcription of "#" genes/proteins
Cyclophilin A	enzyme	280	160	78	4
Cyclophilin B	enzyme	174	116	33	0
Cyclophilin D	enzyme	153	49	34	0
All CYCLOPHILINS	enzyme	902	518	279	4
CCR2	receptor	156	40	82	16
CCR5	receptor	205	89	91	10
FGF21	ligand	371	15	79	164
FXR	nuclear receptor	1200	401	110	482
PPAR alpha	nuclear receptor	609	58	73	462
PPAR beta-delta	nuclear receptor	299	29	10	210
PPAR gamma	nuclear receptor	1549	17	13	1187
THR beta	nuclear receptor	26	10	6	6
THR (alpha+beta)	nuclear receptor	243	43	2	198



# **CRV431 – Pan-Cyclophilin Inhibitor**

- Cyclophilin inhibitor (Ki  $\approx 1 \text{ nM}$ )
- Non-immunosuppressive analog of cyclosporine A
- Oral, once daily
- Concentrates in liver
- Completed Phase 1 (SAD, MAD, DDI)
- Currently in Phase 2a NASH
- No dose-related toxicities observed in preclinical/toxicology or Ph1 clinical



# Anti-Fibrotic and Additional CRV431 Effects in Diverse Preclinical Models

Human Cell Cultures		CRV431 Effects	
Hepatic stellate cells, fibroblasts (multiple organs)	TGFβ or endogenous stimulation	<ul> <li>▼ fibrotic gene expression</li> <li>▼ procollagen and fibronectin secretion</li> </ul>	
Blood platelets	Collagen and thrombin stimulation	abla procoagulant platelet formation	
Human Tissue Explants	(Precision Cut Slice Cultures)	CRV431 Effects	
Liver explants (4 donors)	TGFβ+PDGF-BB or endogenous stimulation	▼ inflammatory/fibrotic gene expression	
IPF lung explants (1 donor)	Endogenous stimulation	<ul> <li>▼ Inflammatory/fibrotic protein secretion</li> <li>▼ tissue fibrosis</li> </ul>	
Animal Models		CRV431 Effects	
Mice (NASH)	Western diet + carbon tetrachloride	82%▼ fibrosis; ▼weight gain	
Mice (NASH)	High fat diet + early STZ (4 studies)	37-57% ▼ fibrosis; ▼ weight gain; 50% ▼ liver tumors	
Mice (liver fibrosis)	Carbon tetrachloride	44%▼ fibrosis	
Mice (kidney fibrosis)	Unilateral ureter obstruction	42%▼ fibrosis	
Rats (liver fibrosis)	Thioacetamide	48% ▼ fibrosis; prevented cirrhosis	
Mice (acute lung injury)	Lipopolysaccharide inhalation	$igstar{$ BAL fluid inflammatory cytokines, neutrophils	
Mice (diabetes)	High fat diet + late STZ	▼adiposity; ▼weight gain	
sdag: HEPA			

# **Preclinical Models: Diet and Toxin Induced Liver Fibrosis in Rodents**





# CRV431 on Human Precision Cut Liver Slices (FibroFind, UK)

### **Fibrosis: Picrosirius Red**

% Normalized to Nonstimulated and Stimulated

150-(normalized %) 0 100-50-8 8 ۰٥ 0 Red 0 -50-Sirius -100-0 ο -150 Vonstinulated Resmettion Vehicle 4/45; - Veg37 (11,111) CUP437 (Sum) Elafibranor Aramchol 50

+ TGF**B**1 + PDGF-BB Stimulation

### Summary

- Precision cut liver slices from 4 human donors
- Sirius red baseline (1.3 9.5%) increased with TGFβ + PDGF-BB stimulation for 3 days to 6.6 - 11.5%
- Alk5i (inhibitor of TGFβ receptor signaling) blocked fibrosis induced by exogenous (TGF $\beta$  + PDGF-BB) and endogenous stimulation as suggested by negative, normalized value





# CRV431 on Human Precision Cut Liver Slices (FibroFind, UK)



Secreted Protein Markers (Daily Average Change)

- Precision cut liver slices obtained from 4 human donors and stimulated with TGFβ + PDGF-BB ± drug treatments
- Alk5i (inhibitor of TGFB receptor signaling) partly or wholly blocked effects of exogenous TGFB + PDGF-BB stimulation
- CRV431 (5 μM) was most effective of five NASH drug candidates at equimolar concentrations at reducing gene expression and secretion of fibrotic and inflammatory markers



# **Bioinformatics Analyses - Transcriptomics**

Clarivate Metacore and Key Pathway Advisor Platforms

+ TGFB1

+ PDGF-BB Stimu



### CRV431 effects on liver transcriptome:

- differentially expressed genes (DEGs) overlap and oppose human NASH DEGs
- hundreds of DEGs map to metabolism, inflammation, extracellular matrix, and other pathways
- causal analyses predict key hub genes such as PPARα, PPARδ, THRβ, and FGF21

(Metacore and Key Pathway Advisor Platforms)



# Processes Regulated by NASH and by CRV431 in Preclinical Models

CRV431 effects on gene expression consistent with human NASH reversal

#### METABOLISM

Development: Insulin, IGF-1 and TNF-alpha in brown adipocyte differentiation

SCAP/SREBP Transcriptional Control of Cholesterol and FA Biosynthesis

GLP-1 in beta cell apoptosis in type 2 diabetes

Regulation of metabolism: Insulin signaling: generic cascades

Oxidative stress in adipocyte dysfunction in type 2 diabetes and metabolic syndrome X

Transcription: Role of AP-1 in regulation of cellular metabolism

Protein folding and maturation: Posttranslational processing of neuroendocrine peptides Glutathione metabolism

#### INFLAMMATION AND EXTRACELLULAR MATRIX (ECM)

Immune response\_IL-5 signaling via JAK/STAT

TGF-beta-induced fibroblast/ myofibroblast migration and ECM production IL-17 and IL-17F-induced inflammatory signaling in normal and asthmatic airway epithelium TNF-alpha-induced inflammatory signaling in normal and asthmatic airway epithelium IL-1 beta- and Endothelin-1-induced fibroblast/ myofibroblast migration and ECM production

#### CANCER

IGF-1 receptor/EGFR cooperation in lung cancer

IGF family, invasion and metastasis in colorectal cancer

#### SAMPLE OF GENE EXPRESSION CHANGES IN HUMAN NASH and BY CRV431 IN PRECLINICAL MODELS

		IIASII	
Apolipoprotein D	Promotes steatosis	↑	$\mathbf{A}$
Cadherin 11	Promotes fibrosis	↑	¥
Collagen 4A6	ECM/fibrosis	↑	$\mathbf{A}$
Glypican	ECM/fibrosis	<b>^</b>	¥
Laminin	ECM/fibrosis	<b>^</b>	¥
Nephronectin	ECM/cancer/inflammation	<b>^</b>	¥
Platelet derived growth factor D	Promotes fibrosis	<b>↑</b>	¥
SOX4 and SOX9	Promotes fibrosis/cancer	<b>^</b>	¥
Transforming growth factor beta	Promotes fibrosis	↑	¥
Apolipoprotein F	Improves HDL/LDL ratio	$\mathbf{+}$	1
Insulin-like growth factor	Regulates metabolism	$\mathbf{+}$	1

### 75% of overlapping CRV431 DEGs reverse NASH DEGs



CRV/431

# **CRV431 Effects on Liver Gene Expression in Preclinical Models**

### Cell-matrix interactions

Gene downregulation

Gene upregulation





# **CRV431 Effects on Liver Gene Expression in Preclinical Models**





# **CRV431 Effects on Liver Gene Expression in Preclinical Models**



# **Causal Analysis of CRV431 Effects in Western Diet/CCl<sub>4</sub> Model**

### SREBP Transcriptional Control of Cholesterol and Fatty Acid Biosynthesis

CRV431 downregulated several enzymes involved in fatty acid biosynthesis and cholesterol biosynthesis







### Human Liver Slices: CRV431 Effects on Transcriptome

Predicted key hub genes include multiple, familiar NASH drug targets



### **Bioinformatics Analyses - Metabolomics** OWL Metabolomics





### CRV431 effects on *liver* metabolome:

- altered expression of 14% of metabolites
- majority of changes consistent with attenuation of thioacetamide effects
- phospholipids major class elevated by TAA and attenuated by CRV431

### CRV431 effects on *plasma* metabolome:

- altered expression of 7% of metabolites
- modest attenuation of thioacetamide effects
- bile acids major class elevated by TAA and attenuated by CRV431



# **Metabolomics in Thioacetamide Rats**

CRV431 attenuated many changes induced by 9 weeks thioacetamide







Liver metabolites (70)

Changes induced by CRV431 in the *liver* mostly opposed the effects of TAA.



Less normalization by CRV431 was observed in the *plasma* metabolome.



### **Metabolomics in Thioacetamide Rats**

CRV431 attenuated many changes induced by 9 weeks thioacetamide



CRV431 shifted the *liver* metabolome of TAA rats towards normal expression, but less for the *plasma* metabolome.



# SUMMARY

# **CRV431**

- Multiple therapeutic actions in diverse liver injury models
- Therapeutic activities consistent across several analytical platforms
- Phase 2 clinical trials:
  - DATA-RICH: histopathology, multi-omics, biomarkers, clinical labs, imaging
  - AI-POWR<sup>™</sup>: in-house, AI/deep learning platform for biomarker selection, protocol optimization, and response predictions

