ATS 2023 The Anti-Fibrotic Effect of GB0139, a Small Molecule Galectin-3 Inhibitor, in **Precision Cut Lung Slices from Idiopathic Pulmonary Fibrosis Tissue FibroFind** A.C. MacKinnon¹, L.A. Borthwick², R.J. Slack³ ¹Galecto Biotech AB, BioQuarter, Edinburgh, UK ²Fibrofind Ltd, William Leech Building, Medical School, Newcastle University, Newcastle upon Tyne, UK, ³Galecto Biotech AB, Stevenage Bioscience Galecto Catalyst, Stevenage, UK. Introduction Results GB0139 is an inhaled, small molecule In vitro characterization of GB0139 (biochemical & cellular) inhibitor of galectin-3 (Gal-3) that is *Ex vivo* IPF BAL cells **GB0139 GB0139 Gal-3 Affinity & Kinetics (SPR)** currently in phase 2b studies for IPF 12 -(NCT03832946). Gal-3 has been 'Gal-3 ivo) 10 14 nM shown to be a key driver of lung Response (RU) (RU) $\mathbf{K}_{D} = \mathbf{k}_{off} / \mathbf{k}_{on} = 3.9 \text{ nM}$ fibrosis¹ and in this study the anti-4.6 nM OH fibrotic effect of GB0139 was **40** 1.5 nM investigated in idiopathic pulmonary 0.5 nM 20-**K**_D = 3.7 nM $\mathbf{k}_{on} = 2.0 \times 10^7 \text{ M}^{-1} \text{s}^{-1}$ $k_{off} = 0.08 \text{ s}^{-1}$ fibrosis (IPF) precision cut lung slices Human ±SD (n) Mouse ±SD (n) (PCLuS). 20 Log [GB0139] M Time (s) 2.1 ± 0.1 (4) FP K_D (nM) 54 ± 17 (3) [GB0139] nM

Methods

fibrotic lung tissue was Human ethically from human sourced explants from three patients with IPF undergoing lung transplantation. All patients had their IPF diagnosis confirmed based on medical history and evaluation of their explanted lung tissue by a board-certified respiratory pathologist. Tissue was inflated with 2–3 % low boiling point agarose and allowed to set at 4°C. PCLuS were then cut at 400 µm on a vibrating microtome and cultured in DMEM media. PCLuS were rested for 48 h prior to treatment with inhibitors for 6 days with media replaced every 24 h. Soluble mediators released were measured daily via standard ELISAs. In addition, analysis of experimental

GB0139 demonstrates high affinity for Gal-3 & inhibits Gal-3 expression ex vivo on IPF BAL cells (IC₅₀ = 361 ± 108 nM (mean ± SD, n=3 patients)).

Ex vivo precision cut IPF lung slice



Secreted galectin-3 (Daily Average Change)



Secreted fibrotic markers (Daily Average Change)



proteomics data derived from MS testing on the PCLuS were completed.

Results

GB0139 concentrationcaused а dependent reduction in Gal-3 in IPF PCLuS. This was associated with a reduction in markers of fibrosis (Co1α1, MMP-7) comparable to pirfenidone nintedanib. A number of and pathways were perturbed by downregulated proteins that included activity transcriptional of SMAD2/SMAD3 platelet and activation, signalling and aggregation.



GB0139 demonstrates Gal-3 target engagement that results in an antieffect fibrotic comparable to approved IPF therapies in PCLuS, at concentrations achieved in the lung when dosed in IPF patients². In addition, GB0139 inhibited pathways PCLuS that correlate with in biomarker changes observed in IPF² and COVID19³ clinical studies.



Patient Demographics

IPF Donor	Age	Sex	FEV1
1	62	Male	1.85
2	64	Male	2.37
3	65	Male	1.53



Galectin-3

hyaluronic acid, TIMP-1, Translational pharmacology of GB0139 in IPF

Screen

Phase Ib – GB0139 dose–dependently reduces galectin-3 in the lungs of IPF patients



Phase Ib – GB0139 *ex vivo* PCLuS *vs.* clinical target engagement in IPF patients



50

[GB0139] in AM μM

75

25

References

- 1. MacKinnon AC et al. Am. J. Respir. Crit. Care Med. 2012;185(5):537-546.
- 2. Hirani N et al., Eur. Respir. J., 2021:57:2002559.
- Gaughan EE et al., Am J Respir Crit Care Med 2022; 3. 207(2):138-149.

Phase Ib – GB0139 inhibits IPF fibrotic markers in



Pathways perturbed by proteins downregulated by 10 µM **GB0139** across all IPF PCLuS patients identifier Pathway name found total pValue Submitted entities foun R-HSA-168249 Innate Immune System 8.4E-06 CTSA:GRN:CD163:NHLRC3:AHSG:HEXB:CD180:RNASE6:GGH:TCIRG1:PLD3:N R-HSA-6798695 Neutrophil degranulation 1.2E-05 CTSA;GRN;CD163;DOK3;NHLRC3;AHSG;IMPDH1;HEXB;GGH;MAN2B1;ACP2;TCIRG1 R-HSA-168256 Immune System 2698 8.4E-04 CTSA;CD163;GRN:NHLRC3;HSPA5;AHSG:HEXB;CD180;RNASE6;GGH:TCIRG1;VAV1;PLD3;MT2A;DOK3;IMPDH1;INPP5D;MRC1;ATP6V0A2;ACP2;MAN2B1;PI3;LG R-HSA-114608 Platelet degranulation 1.1E-03 CD163;HSPA5;AHSG;SERPINE1;ANXA 148 1.4E-03 CD163;HSPA5;AHSG;SERPINE1;ANXA R-HSA-76005 Response to elevated platelet cytosolic Ca2+ R-HSA-9662834 CD163 mediating an anti-inflammatory response 14 2.8E-03 CD163 R-HSA-381183 ATF6 (ATF6-alpha) activates chaperone genes 15 3.1E-03 HSPA5 4.0E-03 HSPA5 R-HSA-381033 ATF6 (ATF6-alpha) activates chaperones 17 R-HSA-1483115 Hydrolysis of LPC 18 4.5E-03 PLA2G15;PLB -HSA-76002 Platelet activation, signaling and aggregation 293 5.5E-03 CD163;HSPA5;AHSG;SERPINE1;ANXA5 R-HSA-9029569 NR1H3 & NR1H2 regulate gene expression linked to cholesterol to 66 5.8E-03 PLTP R-HSA-1483148 Synthesis of PG 6.0E-03 PLD3 R-HSA-1483257 Phospholipid metabolism 7.6E-03 PLA2G15;MTMR14;INPP5D;PLBD1;PLD3 314 R-HSA-8964058 HDL remodeling 7.8E-03 PLTP 24 R-HSA-381119 Unfolded Protein Response (UPR 155 1.1E-02 HSPA5;WIP R-HSA-9024446 NR1H2 and NR1H3-mediated signaling 85 1.2E-02 PLTP R-HSA-1660662 Glycosphingolipid metabolism 1.2E-02 CTSA;HEXB;G R-HSA-77387 Insulin receptor recycling 1.2E-02 ATP6V0A2;TCIRC R-HSA-381070 IRE1alpha activates chaperones 1.8E-02 HSPA5:WIPI1 R-HSA-174824 Plasma lipoprotein assembly, remodeling, and clearan 1.9E-02 NPC1;PLTP R-HSA-917977 Transferrin endocytosis and recycling 2.0E-02 ATP6V0A2;TCIR0 R-HSA-1855204 Synthesis of IP3 and IP4 in the cytosc 2.0E-02 INPP5D;PLD3 R-HSA-390471 Association of TriC/CCT with target proteins during biosynthe 2.1E-02 GBA;LONP 2.1E-02 GEM R-HSA-9768919 NPAS4 regulates expression of target genes R-HSA-3656248 Defective HEXB causes GM2G2 2.2E-02 HEXB R-HSA-2206282 MPS IIIB - Sanfilippo syndrome 2.2E-02 NAGLU 2.3E-02 M6PR;DNASE R-HSA-432720 Lysosome Vesicle Biogenesis R-HSA-1368108 BMAL1:CLOCK, NPAS2 activates circadian gene expression 43 2.3E-02 SERPINE1 R-HSA-2173796 SMAD2 44 2.4E-02 SERPINE 46 2.7E-02 STK4;VAV R-HSA-5218920 VEGFR2 mediated vascular permeab 50 3.1E-02 INPP5D;VAV1 R-HSA-512988 Interleukin-3, Interleukin-5 and GM-CSF signaling R-HSA-109582 Hemostasis 803 3.1E-02 GRN;CD163;HSPA5;AHSG;INPP5D;SERPINE1;ANXA5;ACP2;VAV2 R-HSA-1483206 Glycerophospholipid biosynthesis 3.3E-02 PLA2G15;PLBD1;PLD3 221 R-HSA-8963899 Plasma lipoprotein remodeling 56 3.8E-02 PLTP R-HSA-9748787 Azathioprine ADME 58 4.1E-02 IMPDH1;VAV R-HSA-4341670 Defective NEU1 causes sialidosis 4.3E-02 CTSA

R-HSA-9634815 Transcriptional Regulation by NPAS4 60 4.3E-02 GEM R-HSA-9639288 Amino acids regulate mTORC1 61 4.4E-02 TCIRG1:LAMTOR 64 4.8E-02 SERPINE1 HSA-2173793 Transcriptional activity of SMAD2 R-HSA-2132295 MHC class II antigen presentation 149 4.9E-02 CTSA;ACP2;LGMN R-HSA-1632852 Macroautophage 150 4.9E-02 MTMR14;WIPI1;LAMTOR

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