

Ex-vivo translational assay of hepatic stellate cells using patient-derived serum characterizes the anti-fibrotic activity of CM-101

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Introduction

CCL24, a chemokine that regulates inflammatory and fibrotic activities, was found to be highly expressed in livers of patients with liver fibrosis including those with primary sclerosing cholangitis (PSC)¹ and metabolic dysfunction-associated steatohepatitis (MASH)². Treatment with CM-101, a first-in-class humanized antibody targeting CCL24, has impacted consequential biomarkers of liver fibrosis in patients with MASH (NCT05824156) and in multiple PSC preclinical models^{1,3}.

Aim

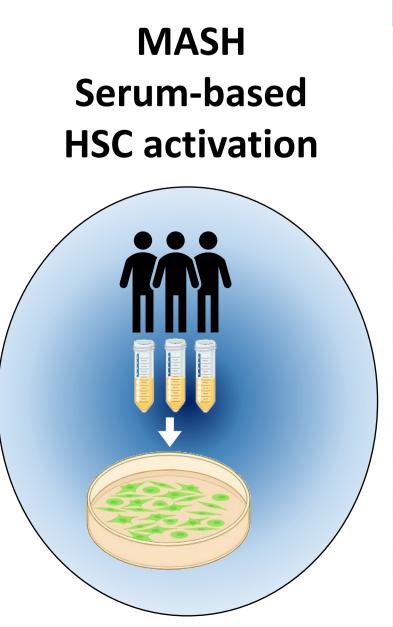
This study aimed to better characterize CM-101's anti-fibrotic activity in patients using an ex-vivo hepatic stellate cells (HSC) activation model.

Methods

- * Activation of HSC [human LX2 cell line]: Cells were activated by 24-hour incubation with either recombinant proteins or sera from MASH patients (NCT05824156) before or after 16 weeks of exposure to either placebo (vehicle) or CM-101 (5 mg/kg subcutaneous every 3 weeks). Activators were added on top of 10% FCS in the media. Quantification of the fibrotic marker αSMA (smooth muscle actin) was assessed by flow-cytometry.
- The secretome of CCL24-stimulated LX2 was examined by RayBio L-507 protein array.
- Serum proteomic analysis: Sera of PSC patients (Royal Free Hospital bio-bank and baseline sera of patients enrolled in the SPRING study, NCT04595825) and healthy controls (NCT06025851) were analyzed using Olink proximity extension assay.

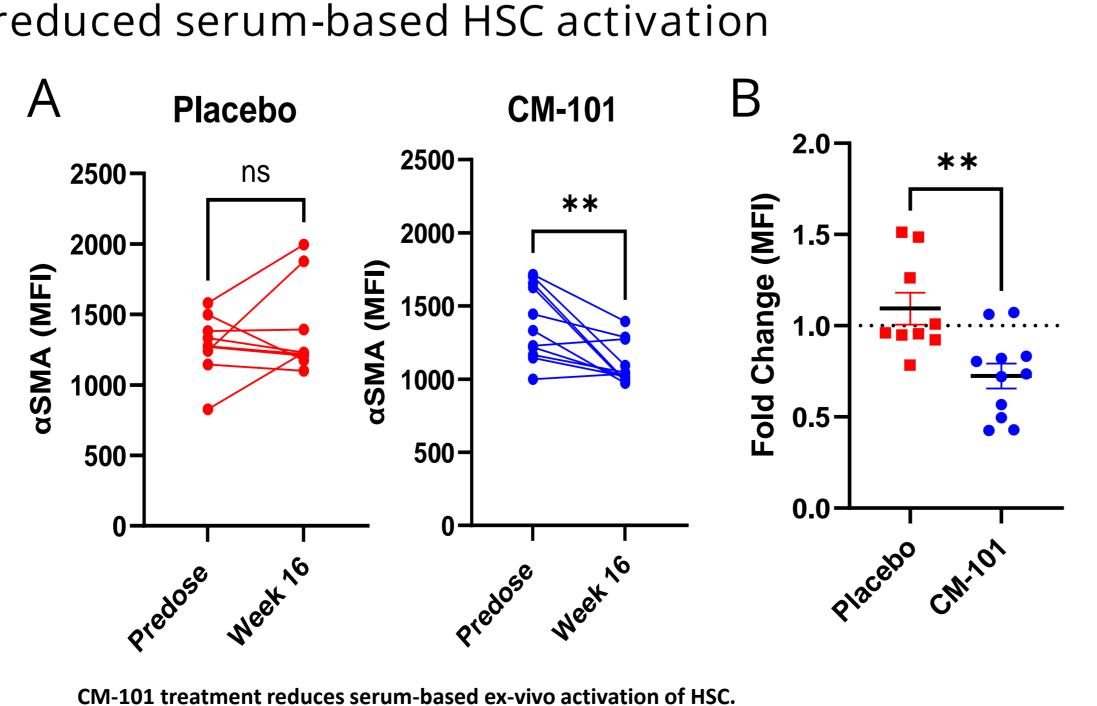
Results

CM-101 treated patients display reduced serum-based HSC activation



	CM-101 (N=11)	Placebo (N=9)		
Age [y]	53.8 (11.4)	46.6 (16.8)		
Female gender (%)	6 (42.9%)	6 (66.7%)		
Alanine Aminotransferase [U/L]	43.7 (13.0)	29.5 (6.5)		
Aspartate Aminotransferase [U/L]	45.8 (25.8)	45.6 (23.1)		
Triglycerides [mg/dL]	142.8 (52.1)	157.6 (67.9)		
LDL-Cholesterol [mg/dL]	93.2 (30.5)	112.1 (32.7)		
Type 2 Diabetes Mellitus	10 (76.9%)	3 (33.3%)		
NAFLD Activity Score [NAS]	4.6 (1.5)	4.8 (0.8)		
FAST Score	0.53 (0.23)	0.34 (0.19)		
Liver Disease Severity	,	,		
Fibrosis Stage 1a	0 (0%)	1 (11.1%)		
Fibrosis Stage 1c	6 (42.9%)	5 (55.6%)		
Fibrosis Stage 2	3 (21.4%)	3 (33.3%)		
Fibrosis Stage 3	5 (35.7%)	0 (0%)		
MELD Score	7.7 (2.6)	6.8 (2.0)		
FibroScan [kPa]	11.5 (5.2)	8.4 (1.1)		
MRI-PDFF (%)	19.0 (7.0)	18.7 (6.3)		
ELF score	9.8 (0.9)	8.6 (0.3)		
Pro-C3 [ng/mL]	40.0 (8.9)	31.3 (5.8)		
FIB4 score	1.5 (1.2)	0.7 (0.3)		
CRP [mg/dL]	1.2 (1.0)	0.6 (0.6)		
Serum CCL24 [pg/ml]	1101 (852)	880 (883)		
Demographics and baseline characteristics of SPLASH study in MASH patients.				

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Data presented as n (%) or mean (SD) unless otherwise stated. NAFLD = Nonalcoholic fatty					
	liver disease; FAST = FibroScan-AST, ELF = Enhanced Liver Fibrosis.				

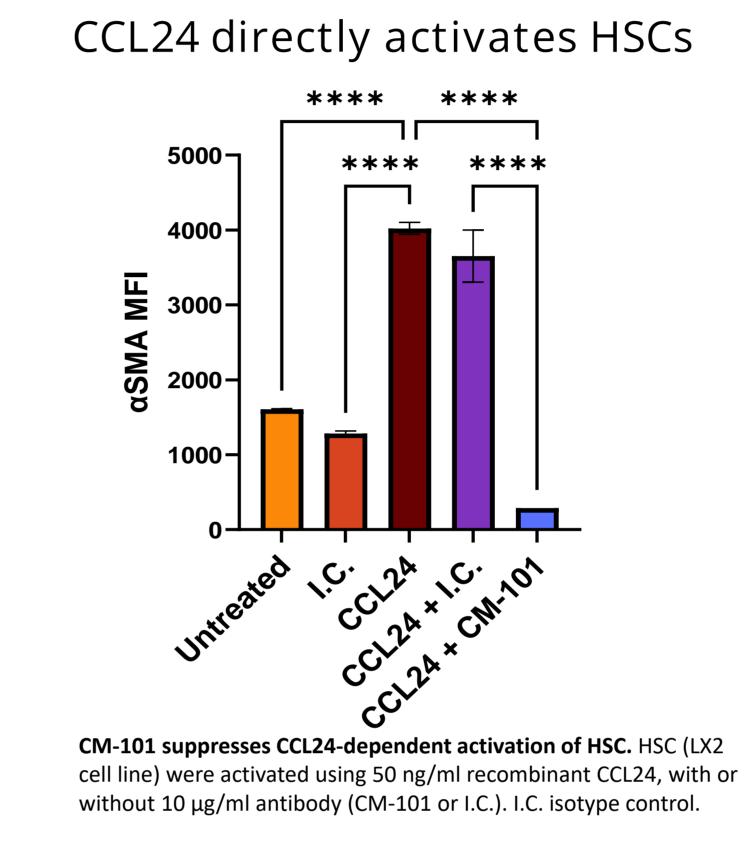


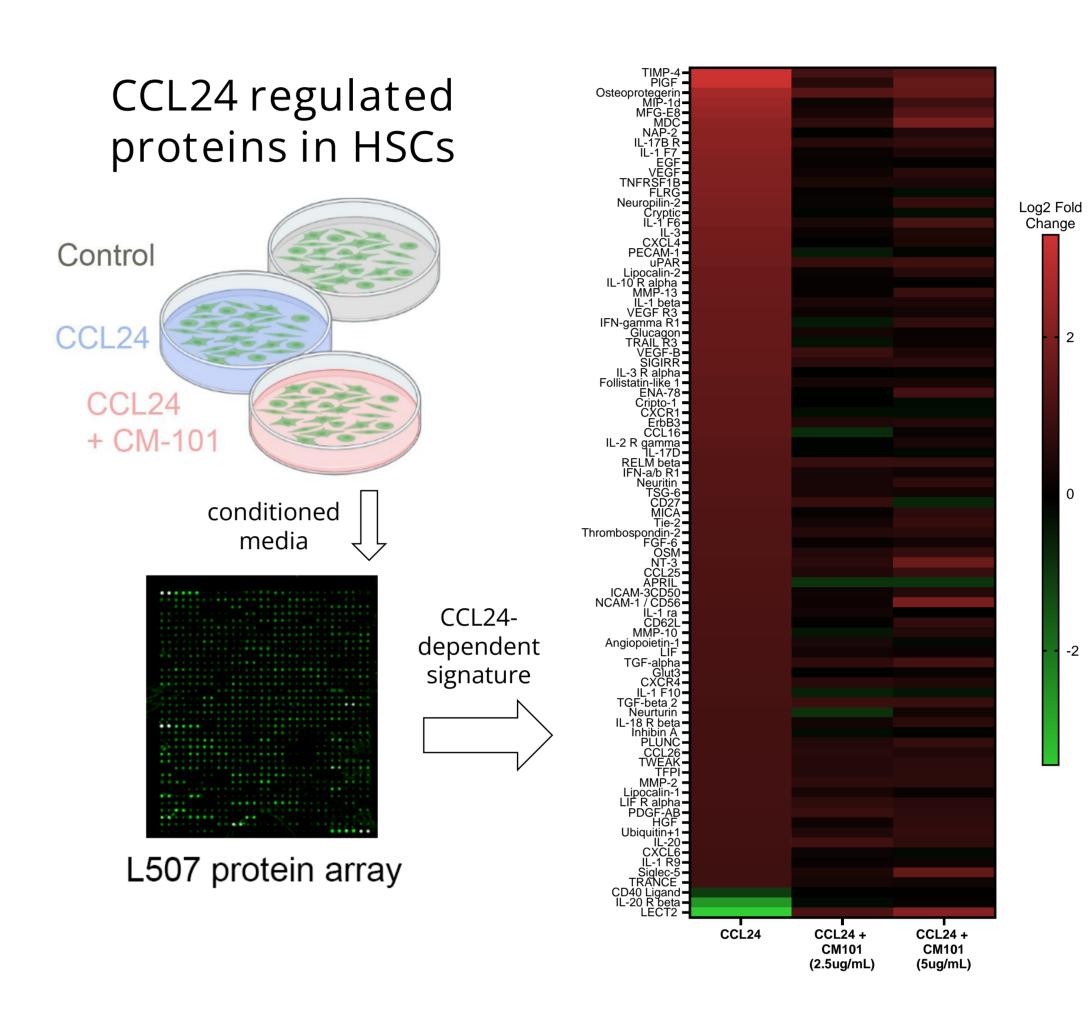
HSC (LX2 cell line) were activated using 1% of serum from MASH patients, before first treatment

change of α -SMA expression between predose and week 16.

(predose), or after 16 weeks of treatment (week 16). (A) α -SMA expression measured by FACS. (B) Fold

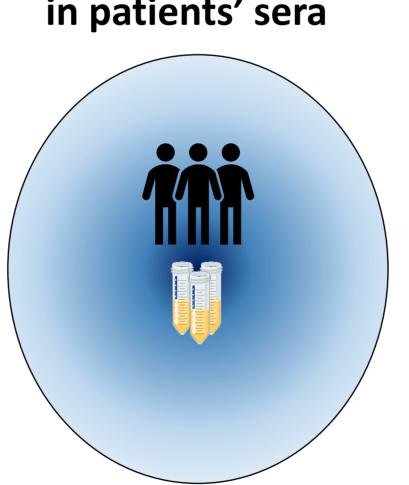
Exogenous CCL24 HSC activation





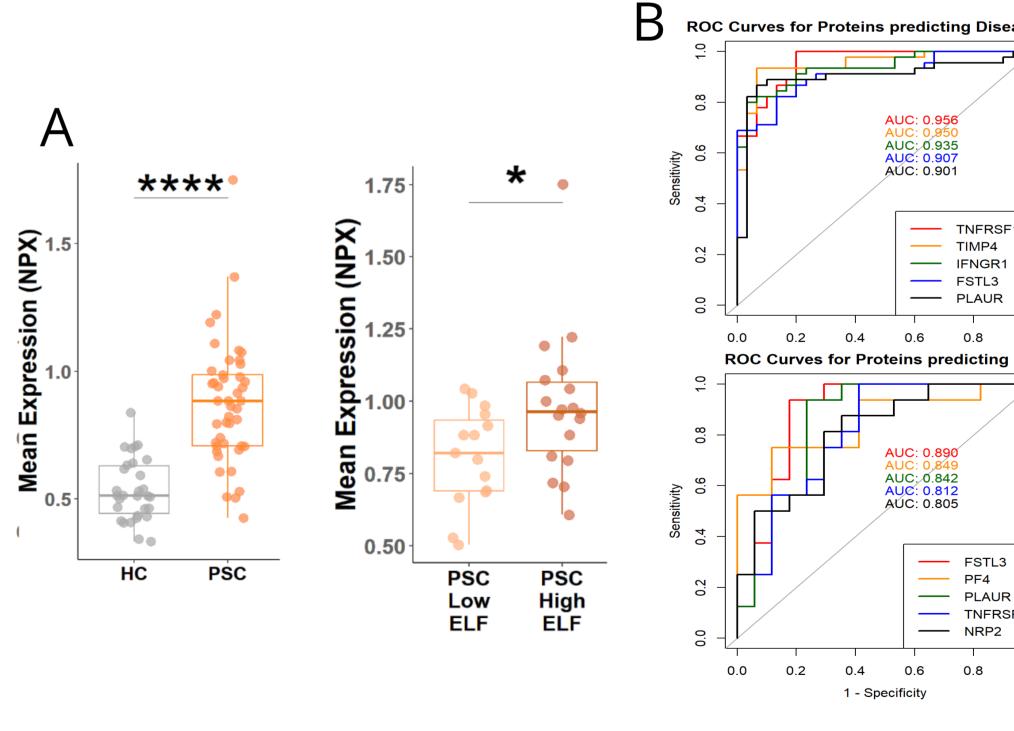
LX2 cells were incubated with either PBS, CCL24, or CCL24 with CM-101. Relative expression of secreted protein was determined by L507 protein array (RayBio). The CCL24-dependent proteir ignature was defined as protein that had a twofold change upon CCL24 treatment and return to baseline expression upon addition of

CCL24 HSC signature in patients' sera



	Healthy controls	PSC
	(N=30)	(N=45)
Age [y]	23.5 (18-38)	45 (18-76)
Duration since diagnosis [y]	NA	4.9 (0-25.3)
Male, n (%)	30 (100)	26 (58)
IBD any, n (%)	0 (0)	30 (67)
ALP [U/L]	74 (42-106)	246 (52-1064)
ALT [U/L]	16.5 (7-45)	70 (10-796)
AST [U/L]	19 (12-34)	54 (15-919)
Bilirubin [mg/dL]	11 (7-18)	12 (3-41)
Fibroscan	NA	10.1 (5.0-17.3)
ELF score	NA	9.95 (7.85-12.84)

Demographics and baseline characteristics of PSC patients and healthy controls Data presented as median (range) unless otherwise stated. IBD = inflammatory oowl disease, ELF = Enhanced Liver Fibrosis



by PSC and its severity. (A) The average expression of CCL24-dependent protein signatures is presented in PSC (left panel) or in patient with PSC, stratified by ELF score of 9.8 (right panel). Boxes represent interquartile (B) The ROC curves for the five serum proteins with the highest AUC values for predicting disease presence (upper panel) or fibrosis severity (lower panel). HC = healthy controls

Conclusions

- > A serum-based ex-vivo HSC activation assay can help in characterizing anti-fibrotic drug effects.
- > A serum-based assay derived from MASH patients treated with CM-101 restored HSC activation.
- > A protein signature generated from CCL24-activated HSC predicted PSC disease and its severity.
- > These findings support CM-101's mode of action in liver fibrosis.
- > CM-101 is currently being tested in a phase 2 study in PSC patients.

References

Greenman, R, Snir, T, Katav, A, Aricha, R, Mishalian, I, et al. The Role of CCL24 in Primary Sclerosing Cholangitis: Bridging Patient Serum Proteomics to Preclinical Data. Cells 2024, 13, 209, doi:10.3390/cells13030209. Segal-Salto M, Barashi N, Katav A, Edelshtein V, Aharon A, Hashmueli S, et al. A blocking monoclonal antibody to CCL24 alleviates liver fibrosis and inflammation in experimental models of liver damage. JHEP Reports.

Differentiation of PSC severity by CCL24-dependent signature

Greenman R, Snir T, Katav A, Aricha R, Mishalian I, Hay O, et al. The Role of CCL24 in Primary Sclerosing Cholangitis: Bridging Patient Serum Proteomics to Preclinical Data. Cells. 2024 Jan;13(3):209.