Silencing MMP7 expression with a lung-targeted RNAi molecule limits fibrosis and preserves pulmonary function in bleomycin-injured rats

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INTRODUCTION

- In idiopathic pulmonary fibrosis (IPF) aberrant basaloid cells overexpress matrix metalloproteinase 7 (MMP7), a secreted epithelial protease that promotes inflammation and fibrosis via cleavage of extracellular matrix proteins, receptors and cytokines
- MMP7 protein is a predictive biomarker of disease severity in IPF and other progressive fibrotic diseases like NASH and biliary atresia
- Genetic loss of MMP7 function protects mice from pulmonary fibrosis. In man, gain-of-function MMP7 alleles are linked to IPF
- Development of highly selective traditional small molecule active site inhibitor drugs has been challenging as MMP7 shares catalytic domain homology with dozens of other matrix metalloproteinase family members
- Therapeutic small interfering RNAs (siRNAs) offer an important novel approach to selectively silence challenging drug targets like MMP7

ARO-MMP7

MMP7

RNAi

trigger

epithelial

targeting

ligand

TRIMTM platform

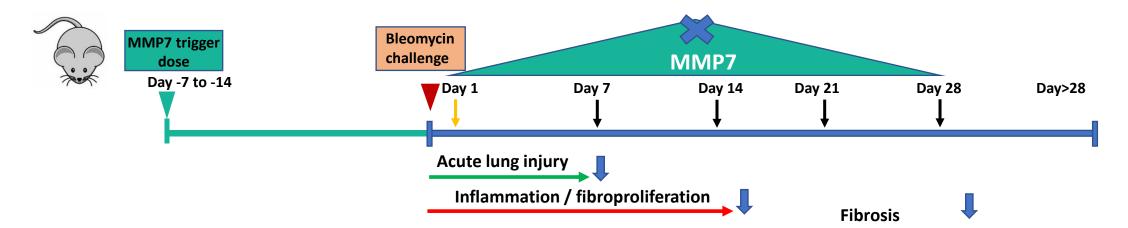
- Optimized RNAi trigger sequence for specific target gene silencing and limited potential off-target interactions
- Rational use of modifying chemistries maximize innate stability and potency
- Epithelial integrin targeting moiety facilitates trigger endocytosis

METHODS

Rat studies

- Animals received an RNAi trigger specifically targeting rat MMP7 mRNA, which was administered by inhalation or multiple intratracheal (IT) instillations 1-2 weeks before bleomycin administration.
- Bronchoalveolar lavage (BAL) and lung tissues were collected at time points ranging from 1-4 weeks after bleomycin injury.
- In vivo respiratory functional mechanics were evaluated via FlexiVent.

Rat bleomycin injury model



Nonhuman primate study

• After receiving baseline BAL collections 1 week prior to dosing, male cynomolgus monkeys received a single inhaled dose of either aerosolized vehicle (sterile isotonic saline) or ARO-MMP7 via a face mask exposure system. Pulmonary deposited doses were 0.24, 0.66, 1.10, and 1.71 mg/kg (n=3 animals per dose level).

• Two weeks after ARO-MMP7 exposure, post dose BAL and lung tissue (12 regional lung tissue samples per animal) samples were collected.

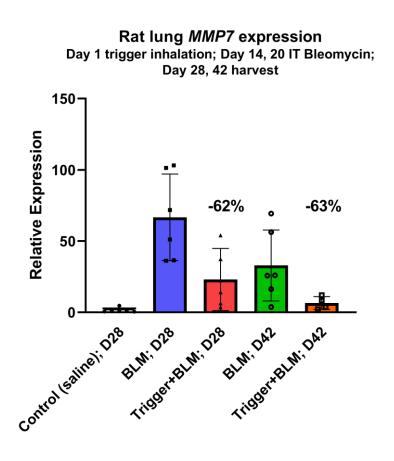
• Lung tissue samples were processed for *MMP7* mRNA and protein expression analysis by qRT-PCR and Western blot. BAL samples were processed for exosomal *MMP7* mRNA and protein expression analysis by qRT-PCR and Western blot.

Human precision cut lung slice (PCLS) study

Fresh lung slices from a healthy donor were cultured and exposed to ARO-MMP7 (0, 0.1, 0.3 or 1μ M) for seven days then processed for analysis of MMP7 mRNA expression by qRT-PCR.

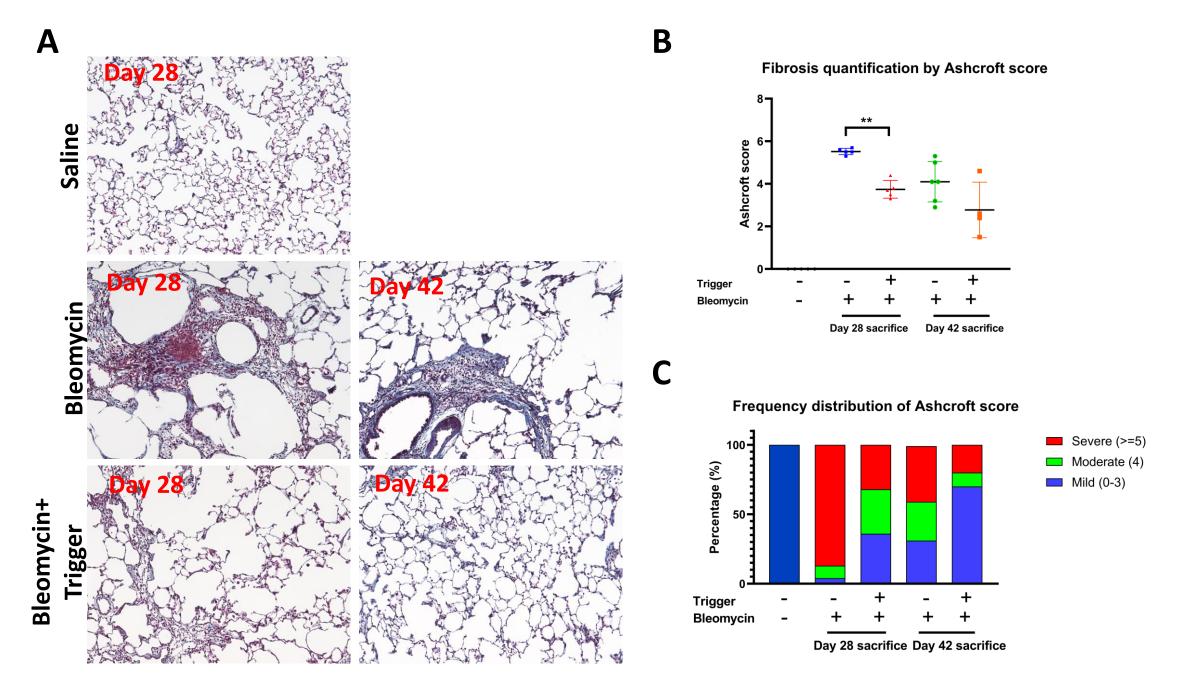
RESULTS

An inhaled RNAi trigger conjugate durably silences pulmonary MMP7 mRNA overexpression in the rat bleomycin injury model



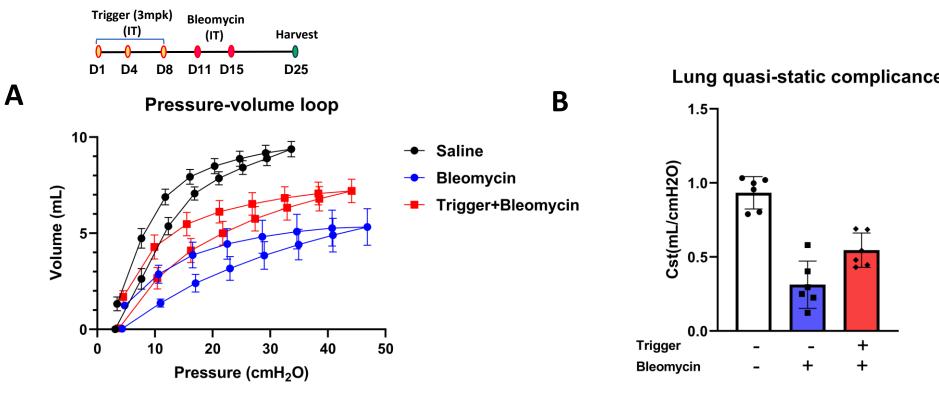
- Sprague Dawley rats received a single inhaled aerosol exposure of 1.4 mg/kg pulmonary delivered dose (PDD) saline vehicle or MMP7-targeting RNAi trigger conjugate on Study Day 1.
 Bleomycin (2U/kg) or saline was dosed intratracheally on Study Day 14 and 20.
- Lung tissues were collected on Study Day 28 and 42 for MMP7 mRNA qRT-PCR (left) and histopathological analysis (below).
- MMP7 expression was robustly induced by bleomycin injury
 RNAi trigger treatment significantly silenced bleomycininduced MMP7 overexpression measured 2 weeks (62% reduction) and 4 weeks (63% reduction) after injury.

MMP7 silencing limits pulmonary fibrosis in the rat bleomycin injury model



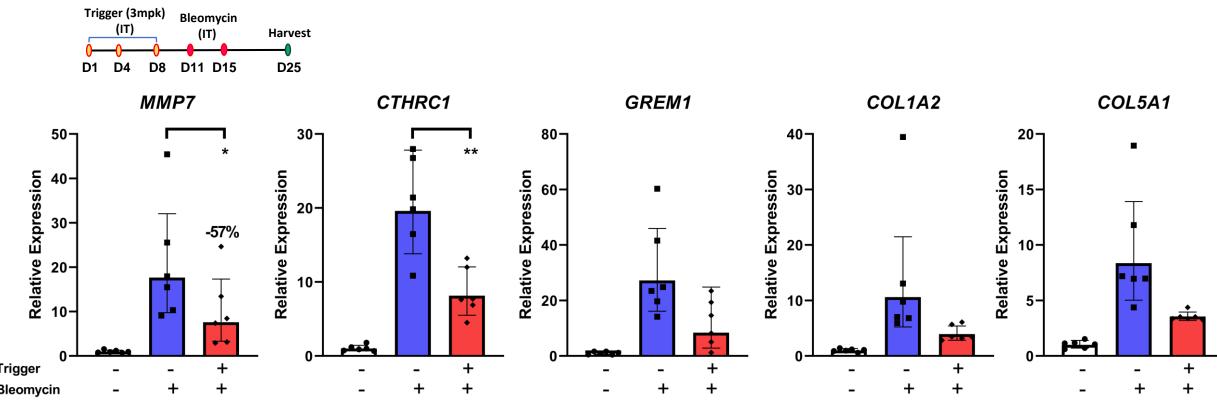
- Panel A, above: Left lung lobes were fixed in 4% PFA for Masson's Trichrome staining. Bleomycin injury produced prominent pulmonary fibrosis on Day 28. Partial resolution of fibrosis was observed on Day 42. A single inhaled dose of *MMP7* RNAi trigger on Day 1 reduced fibrosis at both time points.
- Panels B & C, above: Pulmonary fibrosis was quantified using blinded Ashcroft scoring (grades 0-8). Trigger treatment on Day 1 resulted in significantly lower scores after bleomycin injury on Day 28 (mean with SD; **P<0.01 analyzed by one-way ANOVA). Frequency distribution of the scores shows the injury levels including severe (≥5), moderate (4), and mild (0-3) of all images of the subpleural regions in 20 fields/rat.

MMP7 silencing preserves lung function in the rat bleomycin injury model



- Sprague-Dawley rats were treated intratracheally (IT) with either saline or a MMP7targeting RNAi trigger (3 mg/kg) on Days 1, 4 and 8, followed by bleomycin (2 U/kg, IT) on Days 11 and 16. Lung function was evaluated on Day 25.
- Bleomycin injury significantly reduced lung compliance which is a functional consequence of pathological fibrosis and tissue stiffening.
- *MMP7* silencing with RNAi trigger treatment limited bleomycin-induced reduction in lung compliance and preserved lung function.

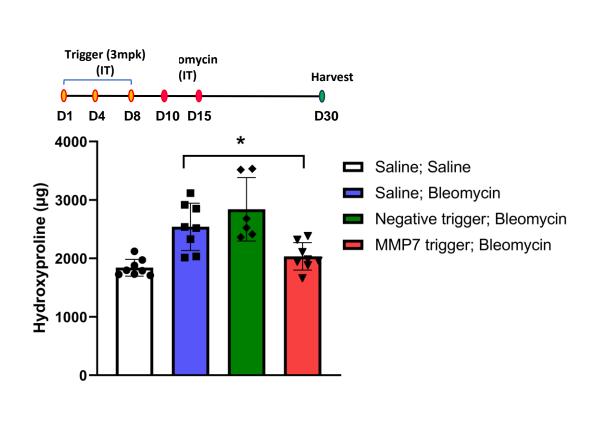
MMP7 silencing in the rat bleomycin injury model limits lung expression of multiple genes with translational significance to IPF



 A common set of upregulated marker genes (including MMP7, CTHRC1, GREM1, COL1A2 and COL5A1) is shared between the rat bleomycin injury model and human IPF patients (Bauer 2015).

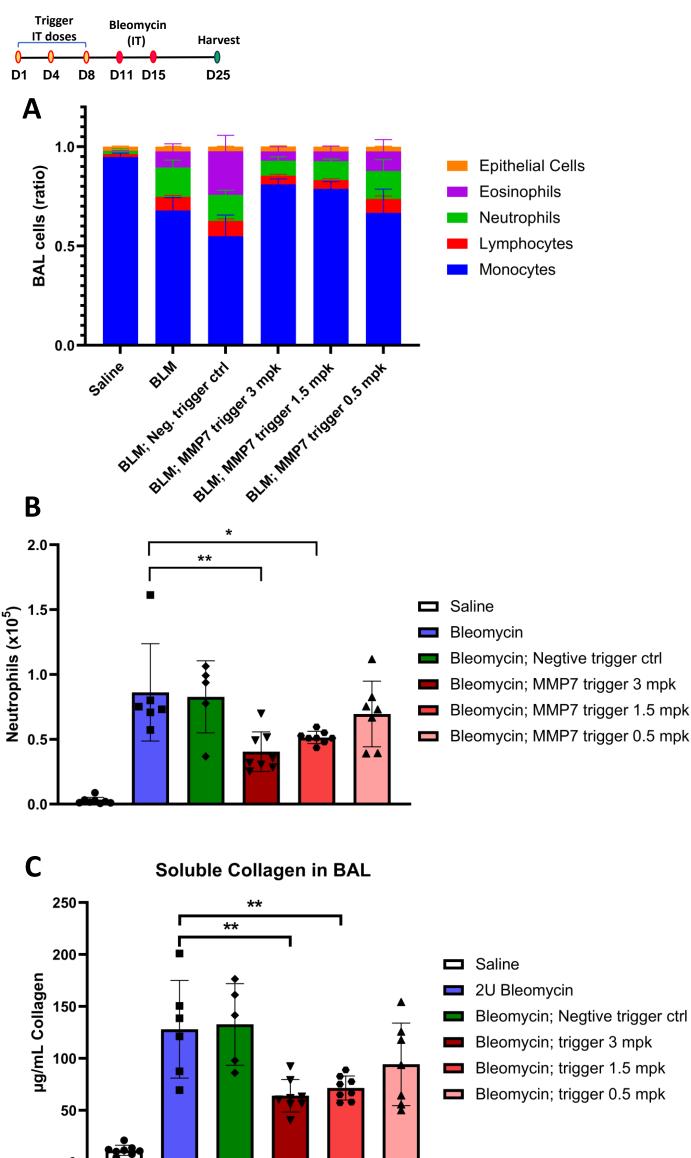
- Silencing *MMP7* expression with an RNAi trigger limited upregulation of multiple IPF marker genes in the rat bleomycin injury model.
- Data are normalized to B2M mRNA expression and the vehicle control group (GMEAN ± with geometric SD; *P<0.05, **P<0.01 analyzed by one-way ANOVA).

MMP7 silencing limits pulmonary collagen deposition in bleomycin-injured rats



- Rats received IT doses of *MMP7* trigger conjugate, non-silencing negative control trigger conjugate or saline vehicle on Study Days 1, 4 and 8 and bleomycin injury on Days 10 and 15.
- On Day 30, middle and lower right lung lobes were collected for collagen content analysis by hydroxyproline assay.
- Increased lung collagen content associated with bleomycin injury was significantly attenuated by treatment with *MMP7*targeting RNAi trigger conjugate. Treatment with a non-silencing negative control trigger did not reduce collagen content.

MMP7 silencing limits inflammation and BAL soluble collagen accumulation in the rat bleomycin injury model



- Chronic pulmonary inflammation contributes to IPF pathogenesis and is characterized by elevated inflammatory cells in BAL samples.
- Rats received IT doses of MMP7 trigger (0.5, 1.5 or 3 mg/kg), nonsilencing negative control trigger (3 mg/kg) or saline vehicle on Study Days 1, 4 and 8 and bleomycin injury on Days 11 and 15.
- BAL was collected for inflammatory cell differential count on Day 25.
- Leukocytes (eosinophils, neutrophils and lymphocytes) in the BALF were increased after bleomycin injury on Day 25.
- MMP7-targeting RNAi trigger conjugate treatment produced dosedependent reductions in BAL leukocyte counts (panel A, left) and neutrophil counts (panel B, left). Treatment with a non-silencing negative control trigger did not reduce inflammatory cell BAL counts.
- Concentrations of soluble collagen in BAL samples (an index of lung fibrosis) was increased with bleomycin injury and significantly reduced with *MMP7*-targeting RNAi trigger conjugate treatment (panel C, left). Treatment with a non-silencing negative control did not reduce BAL collagen concentration.
- Data are analyzed by one-way ANOVA (MEAN with SD; *P<0.05, **P<0.01)

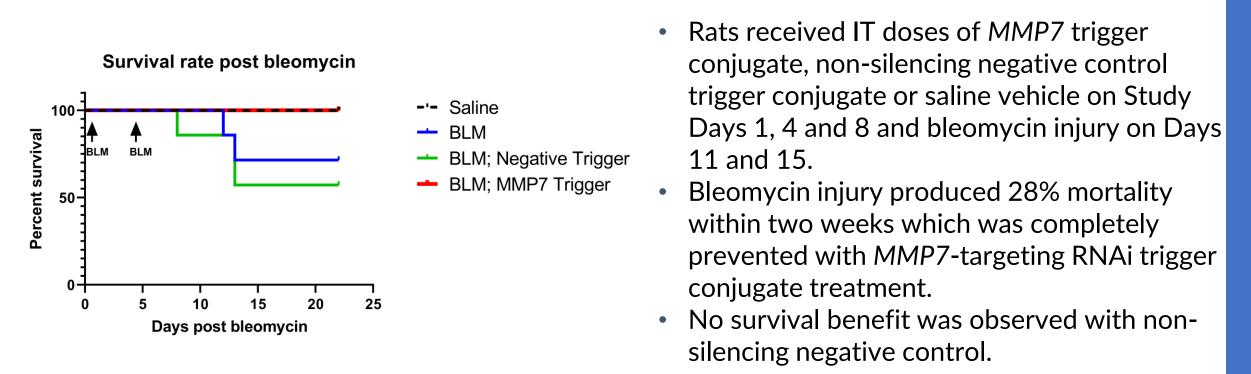


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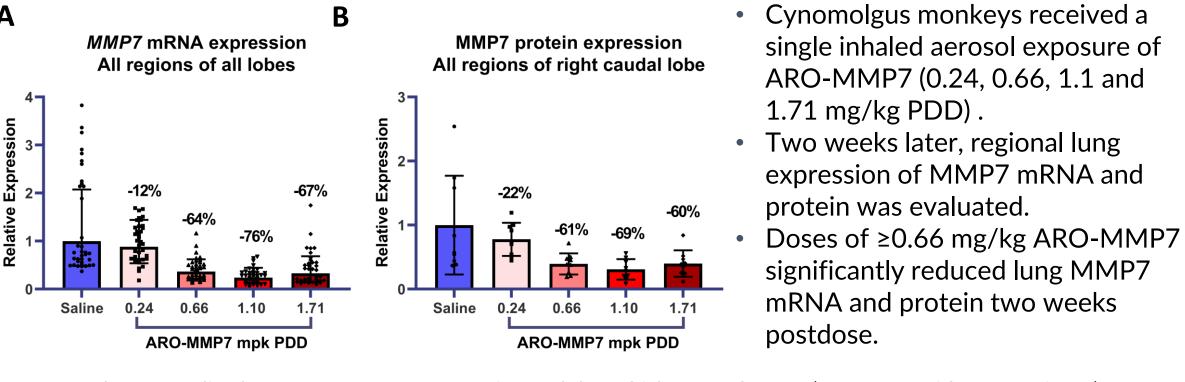
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MMP7 silencing limits mortality in the rat bleomycin injury model

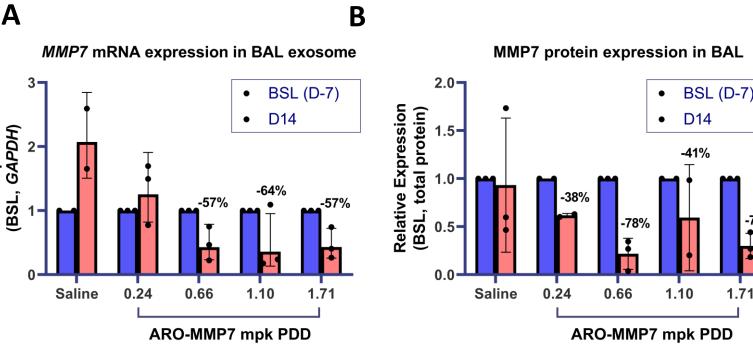


Dose-dependent silencing of pulmonary MMP7 mRNA and protein in nonhuman primates two weeks after a single inhaled dose of ARO-MMP7



A: qRT-PCR data normalized to GAPDH mRNA expression and the vehicle control group (GMEAN ± with geometric SD). N=36 regional lung samples per group (12 samples from each of 3 monkeys) B: Western blot data normalized to total protein and the vehicle control group (MEAN ± SD)

ARO-MMP7 silences MMP7 protein and exosomal MMP7 mRNA in nonhuman primate BAL samples two weeks postdose



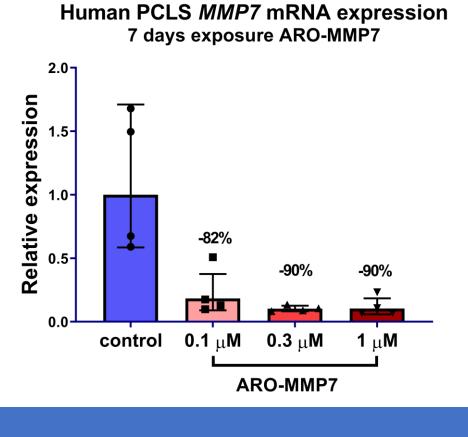
 Reduced BAL MMP7 protein (panel B, left) and exosomal MMP7 mRNA (panel A, left) was observed at exposures of ≥0.66 mg/kg ARO-MMP7 and correlated with lung tissue protein and mRNA expression.
 BAL will be evaluated for evidence of ARO-MMP7 target engagement in future

ARO-MMP7 mpk PDD ARO-MMP7 mpk PDD Clinical trials. A: qRT-PCR data normalized to baseline Day -7 and GAPDH mRNA expression in vehicle controls (GMEAN ± with geometric SD). n=2 for group 1; n=3 for group 2-5 (1 BAL sample per animal). B: Western blot data normalized to baseline Day -7 and total protein expression in vehicle controls (MEAN ± SD). n=2 for group

B: Western blot data normalized to baseline Day -7 and total protein expression in vehicle controls (MEAN \pm SD). n=2 for group 2, 4; n=3 for group 1, 3, 5 (1 BAL sample per animal).

ARO-MMP7 silences MMP7 mRNA expression in primary human lung tissue

- Human precision cut lung slices (PCLS) were exposed to ARO-MMP7 (0.1, 0.3 or 1 μM) for 1 week.
- Significant silencing of human MMP7 mRNA was observed in all ARO-MMP7 treated groups.



CONCLUSIONS

- In rats, inhaled epithelial-targeted RNAi triggers silence pulmonary MMP7 overexpression, limit fibrosis and preserve lung function after bleomycin injury, effectively phenocopying findings in MMP7 knockout mice and confirming the pathogenic role of MMP7.
- ARO-MMP7 silences pulmonary MMP7 in nonhuman primates and in human lung slices and is a promising new therapeutic candidate for the treatment of IPF.