1 Supplementary Information

2 Fibulin-1c regulates transforming growth factor-β

3 activation in pulmonary tissue fibrosis

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29 Supplementary Figures and Figure legends





- 40 differences were determined with two-tailed student t-test. **P<0.01, ****P<0.0001
- 41 compared to PBS-challenged mouse controls.



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Supplementary Figure 2. Fbln1c is increased around the airways in bleomycininduced experimental pulmonary fibrosis. A single bleomycin challenge was used to induce pulmonary fibrosis in WT mice. Controls received PBS. A time-course of lung sections were assessed for Fbln1 protein levels around small airways using immunohistochemistry. Scale bar=500 µm; inserts show expanded images of indicated regions, scale bar=50 µm. Images are representative of n=24-40 airways from n=4-8 mice per group.



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Supplementary Figure 3. Bleomycin challenge of *Fbln1c^{-/-}* mice does not increase collagen fibers around the airways. A single bleomycin challenge was used to induce pulmonary fibrosis in WT and *Fbln1c^{-/-}* mice. Controls received PBS. Collagen fibers were imaged by second harmonic generation (SHG) microscopy. Collagen backward signal (B_{SHG}) is violet, and collagen forward signal (F_{SHG}) is cyan, scale bar=100 µm. Images are representative of n=40 airways from n=4 mice per group.





59 Supplementary Figure 4. Bleomycin challenge of $Fbln1c^{-/-}$ mice does not increase the mRNA levels of Mmps or Timp1 in whole lung tissues. A single 60 bleomycin challenge was used to induce pulmonary fibrosis in WT and $Fbln1c^{-/-}$ mice. 61 Controls received PBS. (A) Mmp1, (B) Mmp3, (C), Mmp8, (D) Mmp12, (E) Mmp13 and 62 (F) Timp1 mRNA levels in lungs determined using qRT-PCR (n=6-8). Statistical 63 differences were determined with one-way ANOVA followed by Bonferroni post-test. 64 *P<0.05, **P<0.01, ***P<0.001 compared to PBS-challenged WT mice. [†]P<0.05, 65 ⁺⁺⁺P<0.001, ⁺⁺⁺⁺P<0.0001 compared to bleomycin-challenged WT mice. 66



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Supplementary Figure 5. TGF- β challenge of *Fbln1c^{-/-}* fibroblasts does not affect 68 Smad3 mRNA levels, and bronchoalveolar lavage fluid (BALF) from Fbln1c^{-/-} 69 mice reduces Smad gene levels in fibroblast from WT mice. Primary fibroblasts 70 71 were isolated from the lungs of WT and *Fbln1c^{-/-}* mice and stimulated with TGF- β or media control for 24 h. (A) Smad3, (B) Smad2 and (C) Smad4 mRNA levels in 72 73 fibroblast lysates determined by qRT-PCR (n=6 of each genotype). Primary mouse lung fibroblasts from WT mice were incubated with bronchoalveolar lavage fluid (BALF, 74 20μ l each mouse from WT and *Fbln1c^{-/-}* mice after 28 days bleomycin challenge and 75 controls for 6 hours. (D) Smad3, (E) Smad2, (F) and Smad4 mRNA levels in fibroblast 76

- 77 Iysates determined by qRT-PCR. Statistical differences were determined with one-way
- ANOVA followed by Bonferroni post-test. *P<0.05 compared to media control.