

P21 Regulates TGF- β_1 -Induced Pulmonary Responses via a TNF- α -Signaling Pathway

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Online Data Supplement

Online Supplemental Materials

TGF-β(-)/p21(+/+).

TGF-β(+)/p21(-/-).

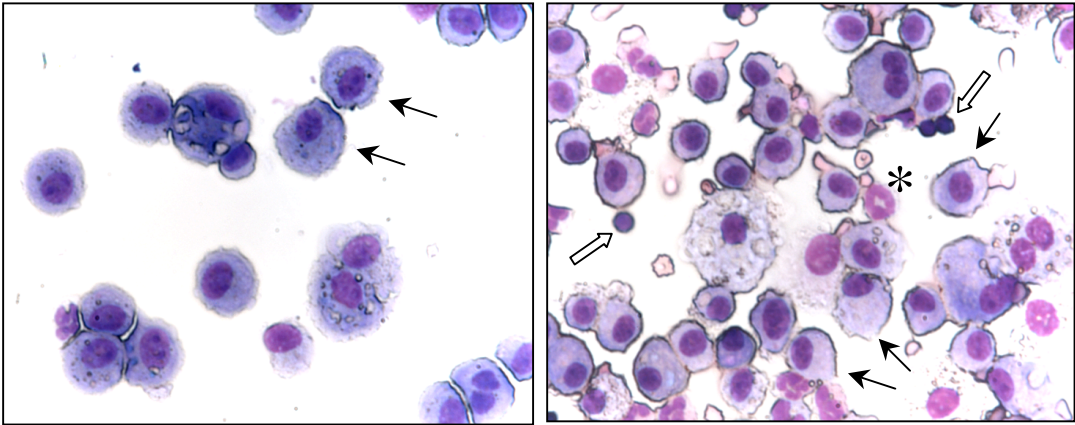


Figure E1. BAL cytologies of TGF-β(-)/p21(+/+) and TGF-β(+)/p21(-/-). After 1 month of Dox induction, BAL cytology was prepared. TGF-β(+)/p21(-/-) mice demonstrates macrophage dominant inflammation with minor infiltration of lymphocytes (open arrow) and eosinophil (*).

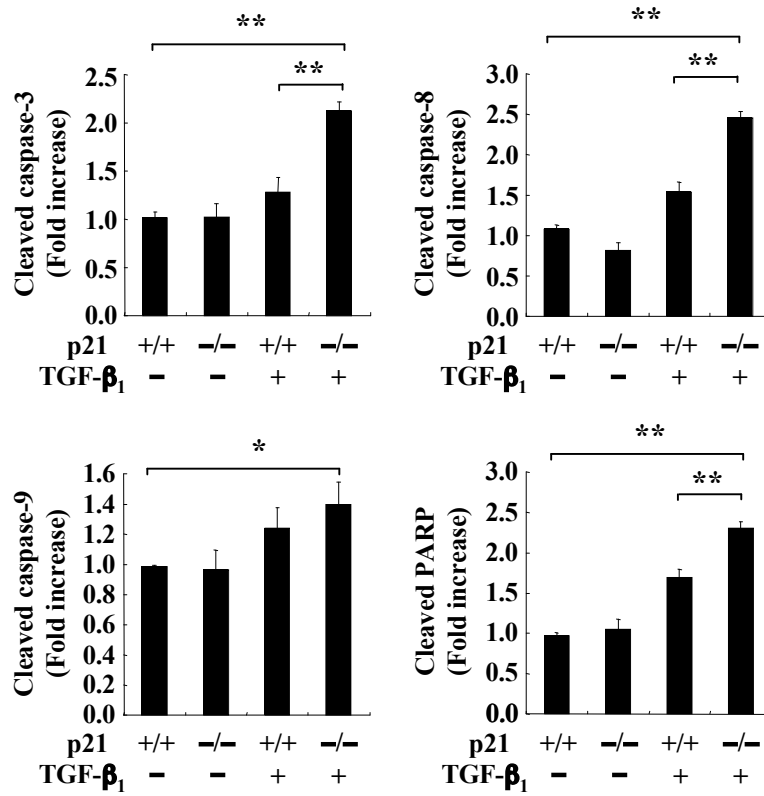


Figure E2. Caspase activation in the lung of TGF-β₁ transgenic mice with and without p21 null mutation. Relative fold induction of each cleaved fragments were evaluated by Western analysis and subsequent densitometrical quantitation. The values represent the mean±SEM of evaluations in a minimum of 5 animals (*p<0.05, **p<0.01).

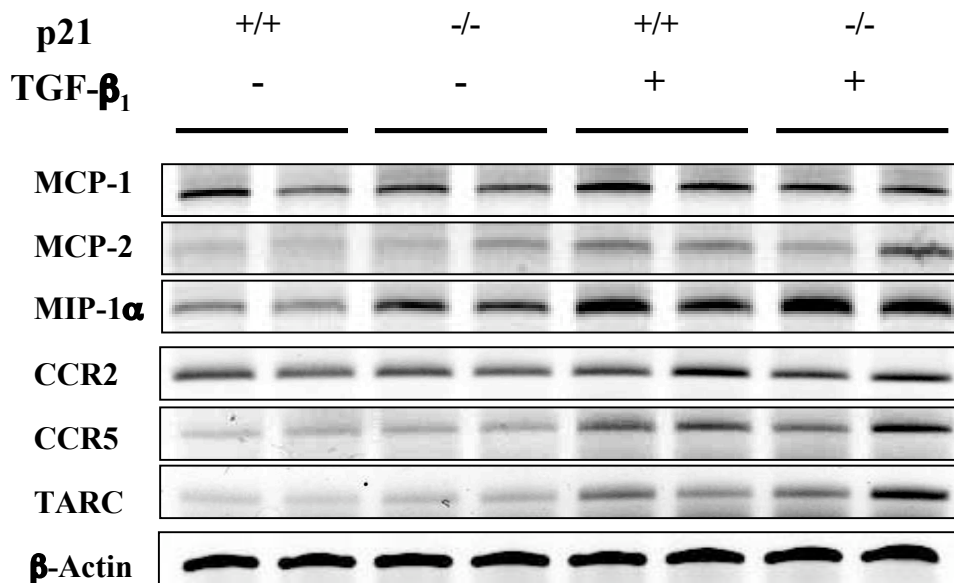
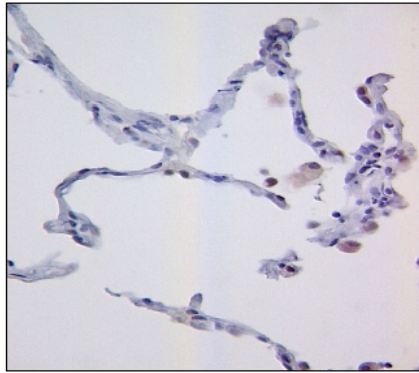


Figure E3. mRNA expression of selected chemokines and chemokine receptors in wild type and TGF- β_1 transgenic mice with and without p21 null mutation. The relative mRNA expression was evaluated using semiquantitative RT-PCR.

Normal Region



Fibrotic Region

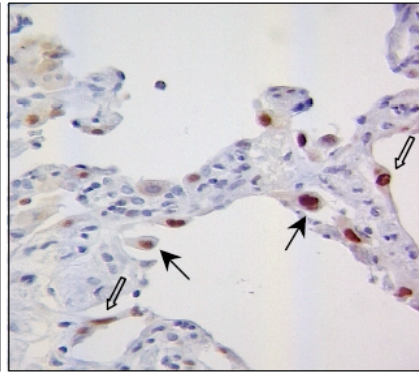


Figure E4. Immunohistochemical localization of p21 in lungs from patients with pulmonary fibrosis. P21 positive cells were stained with brown colors. Closed and open arrows indicate macrophages and epithelial cells, respectively (10X original magnification).