

Figure S1. LacZ expression in *Fgf10-lacZ* lungs indicates that *Fgf10* is expressed in a subset of lipofibroblasts. (A-F) *LacZ* staining of control (A) and *Fgf10-lacZ* (B) P5 lungs. (C, D) Higher magnification of the lung shown in (B) indicates the presence of lacZ⁺ cells not only close to the pleura (arrow) but also in the parenchyma (asterisk). (E-F) Benzyl Benzoate has been used to better visualize the internal signal in *Fgf10-lacZ* lungs. (G-I) Nuclear fast red staining of P5 *Fgf10-lacZ* lungs. (J-U) IHC of P5 *Fgf10-lacZ* lung sections showing Acta2⁻ Adrp⁺ ORO⁺ lacZ⁺ cells adjacent to AECII. The boxes in panels (G, J, M, P and S) indicate the areas showed in higher magnification in panels (H-I, K-L, N-O, Q-R, T-U) respectively. (V-X) Oil Red O (ORO) staining on frozen sections of P5 *Fgf10-lacZ* lungs. (Y) Quantification of the number of lacZ⁺ Adrp⁺ cells as compared to the total number of lacZ⁺ cells. (Z) Quantification of the number of lacZ⁺ Adrp⁺ cells as compared to the total number of Adrp⁺ cells. Scale bar: 100 μm. *n*=4. NFR: Nuclear Fast Red.

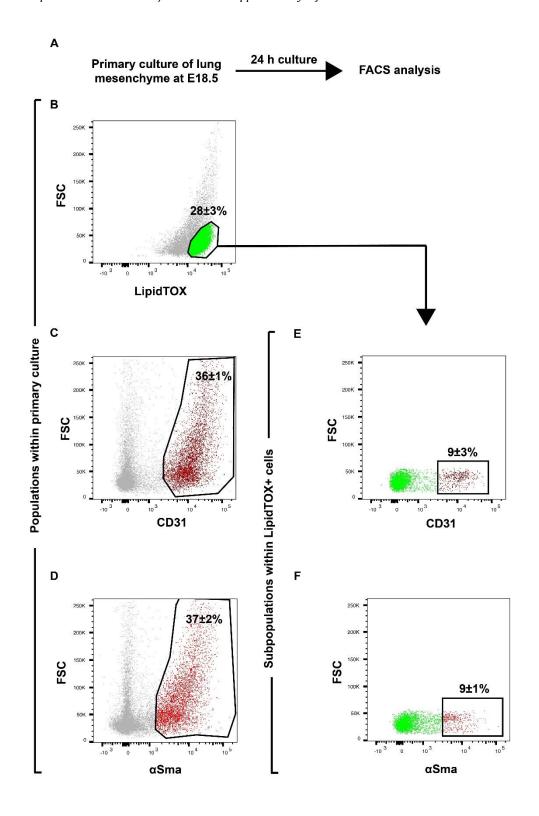


Figure S2. Analysis of primary culture of E18.5 lung mesenchyme. (A) Experimental setup. (B) FACS analysis for LipidTOX shows that lipofibroblasts represent $28\pm3\%$ of total cultured cells after 24 hours. (C,D) Other populations include endothelial cells ($36\pm1\%$) and smooth muscle cells ($37\pm2\%$). (E,F) Endothelial cells and smooth muscle cells represent $9\pm3\%$ and $9\pm1\%$ of total LipidTOX-positive cells respectively. n=6-12.

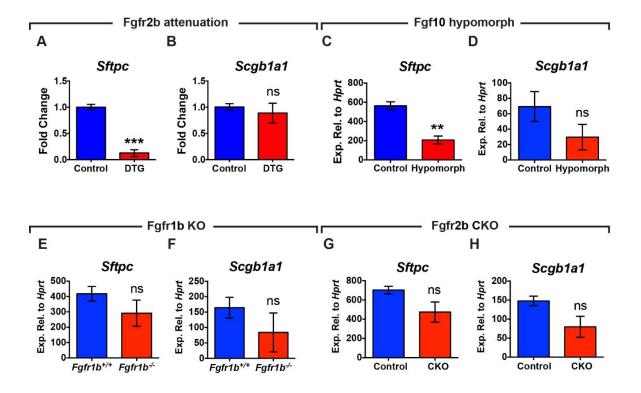


Figure S3. Analysis of Sftpc and Scgb1a1 expression levels in control and mutant lungs by qPCR. Sftpc expression levels reveal significant downregulation upon Fgfr2b attenuation (A) and in Fgf10 hypomorphic lungs (C) but not in Fgfr1b KO (E) or Fgfr2b CKO lungs (G). Scgb1a1 expression is not significantly altered in mutants compared to littermate controls (B,D,F,H). ** P<0.01, *** P<0.001. KO: Knock out, CKO: Conditional knock out.