

## 2.12 Aberrant lipid metabolism drives pulmonary fibrosis

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**Background:** Lipid homeostasis is important in the lung as 90% of surfactant is lipid. We hypothesise that aberrant lipid metabolism contributes to the fibrotic niche in IPF. **Methods:** We performed lipidomic analysis on lung tissue from human IPF, bleomycin, silica and *Abcg1*<sup>-/-</sup> mice. We exposed IPF fibroblasts to oxidised and unoxidised PC and CE lipids, measuring expression of fibrotic markers. **Results:** There was increased lipid content in IPF (p=0.03), bleomycin (p=0.06), silica (P=0.02) and *Abcg1*<sup>-/-</sup> mice (p<0.001) lung tissue. Specific increases were noted in 16:0 and 20:4 containing lipids; these lipids were chosen for IPF fibroblast exposure. ACTA2 and Colla1 expression were upregulated in IPF fibroblasts following exposure to saturated PC 16:0 (p<0.001) and CE 16:0 (p<0.01) and there was no change after unsaturated PC 20:4 (p=ns) and CE 20:4 (p=ns). When PC 20:4 and CE 20:4 are oxidised they upregulate ACTA2 and Colla1 expression (p<0.01). This effect is reduced, but not neutralised, by co-treatment with CD36 receptor blocker. **Conclusion:** We have shown that total lipid increases in IPF and hypothesise that unsaturated lipid becomes oxidized and this in turn drives production of pro-fibrotic cytokines. This may be a crucial hit in driving unchecked fibrosis seen in IPF. **Conflict of Interest:** The authors declare no conflict of interest.