volunteers. In the event of deterioration with escalation of severity category repeat samples were obtained.

**Results** Patients with COVID-19 were more likely to be male (67% vs 20% (HC); p<0.001), older (64.4 ± 16.7 vs 47.7 ± 13.5; p<0.001) have a greater BMI (32.3 ± 6.6 vs 27.9 ± 5.1; p=0.01) and be never-smokers (60% vs 30%; p=0.001). We demonstrated a hyperinflammatory and pro-coagulative state in all patients with COVID-19. All measures of complement activity were significantly higher in patients with COVID-19, including levels of C5a (HC 13[7,21] vs Covid-19 35[24, 43]; p<0.001) and SC5b9-complex (HC 654[419, 1120] vs Covid-19 1452[970, 2170]; p<0.001) which both increased with disease severity and were statistically significantly different between mild and severe disease. SC5b9-complex was significantly higher in patients who deteriorated from moderate to severe disease (1393 [1019, 1986] vs 2116 [958, 4538]; p=0.03).

**Discussion** Our findings demonstrated increased levels of complement activity in patients with COVID-19, particularly in those patients requiring non-invasive and mechanical ventilation and those patients that deteriorate requiring increasing ventilatory support. The complement cascade is a key player in protective immunity against pathogens, with its activation orchestrating key immunoprotective and anti-inflammatory effects. Increased activation of the complement cascade may contribute to the dysregulated and destructive inflammatory response that leads to multi-organ failure and our findings suggest a potentially important treatment target for COVID-19.

**Abstract S57 Figure 1** Bar chart with error bars displaying mean% positive cells for each age group for alveolar and bronchial ACE2 and TMPRSS2 expression in 38 subjects (<1yr n=9, 1–9yrs n=4, 10–17yrs n=6, 18–34yrs n=7, 35–54yrs n=7, >55yrs n=5) was significantly greater in alveolar than bronchial sections. In children, ACE2 and TMPRSS2 expression was detected in only 2–3% of cells in alveolar and bronchial tissue. Alveolar ACE2 receptor expression was significantly greater in adults and appeared to increase with age. Adult alveolar ACE2 receptor expression was highly variable, being detected in some specimens in ~30% of cells.

**Conclusion** ACE2 and TMPRSS2 expression is similar in upper airways of children and adults, likely indicating that both groups are equally susceptible to SARS-CoV2 infection. In contrast, expression of both these receptors/co-factors is greater in adult lower airways of adults, and particularly for ACE2 receptor in alveolar tissue. In some adults, ACE2 receptor was detected in up to a quarter of alveolar cells, potentially explaining why some adults are so susceptible to lower respiratory tract disease.

**Treatment choices in cystic fibrosis and bronchiectasis: what works and when**

**Abstract S58**

**Background** Angiotensin-converting enzyme 2 (ACE2) receptors and the serine protease co-factor TMPRSS2 are cellular receptors/co-factors for SARS-CoV2, allowing viral entry into host cells. Children, when infected with SARS-CoV2, generally present with mild disease, and particularly milder lower respiratory tract symptoms. One hypothesis to explain this phenomenon is differential expression of ACE2 and TMPRSS2 in the respiratory tracts of children and adults.

**Aims** To investigate ACE2 receptor and TMPRSS2 expression in upper/lower respiratory tracts of children and adults without COVID disease.

**Methods** Nasal brushings from obtained from children of different ages (3 months –15 years) undergoing routine elective surgery, and from volunteer adults (20–61 years). Nasal epithelial ACE2 and TMPRSS2 mRNA expression was analysed by PCR.

Post-mortem lung tissue from children and adults without COVID-19 was stained to identify ACE2 and TMPRSS2 protein expression by immunohistochemistry (IHC). Each sample was digitalised using Philips Digital Pathology Solutions software, with three alveolar and three bronchial screen-grab images obtained at x40 magnification, and analysed using Image J.

**Results** Nasal ACE2 and TMPRSS2 mRNA expression in children and adolescents (n=12) and adults (n=26) was similar. Immunohistochemical lung tissue ACE2 and TMPRSS2 protein