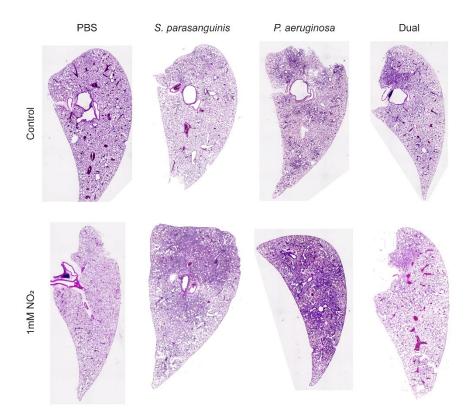
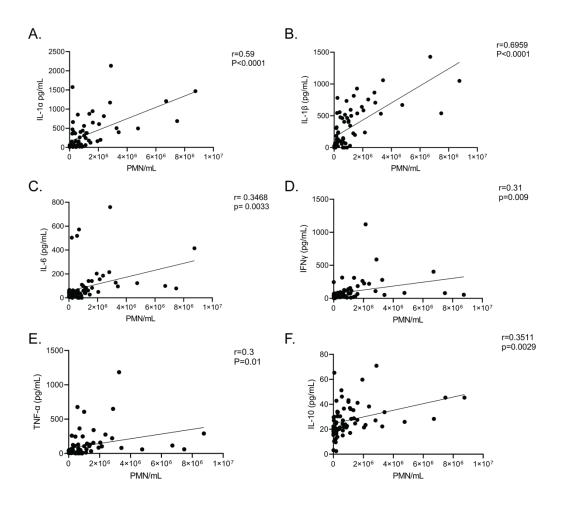


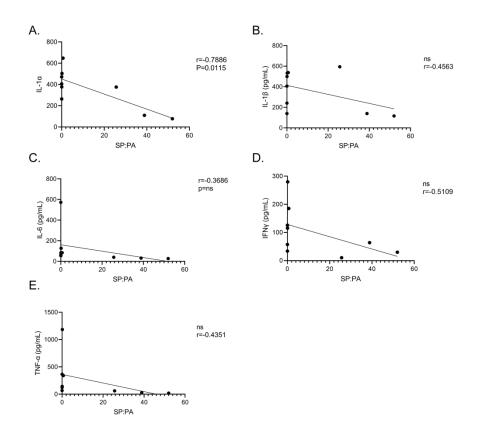
S1. *poxL* expression is increased in artificial CF sputum. *poxL* expression was measured via qpcr in THB, artificial saliva (A), and SCFM2 (B). *poxL* expression was also measured in THB, artificial saliva, and SCFM2 with or without nitrite (C-E, respectively). n=3, ns, p>0.05, (Wilcoxon Test).



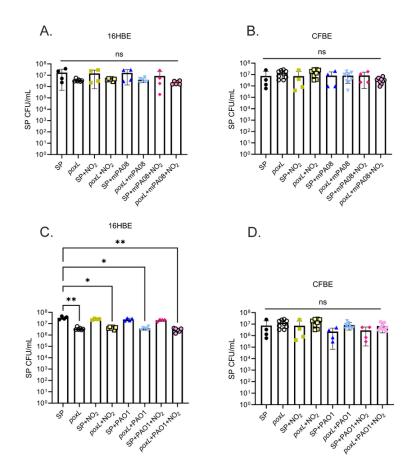
S2. S. parasanguinis attenuates *P. aeruginosa* tissue damage. Representative images of hematoxylin and eosin staining from the worst scored sections from each infection group.



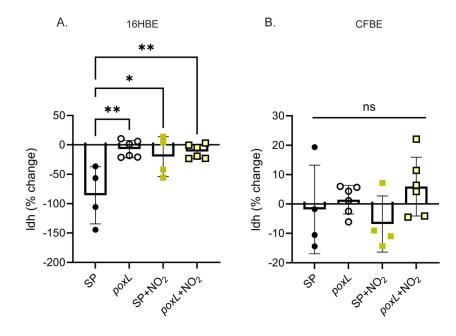
S3. Cytokines positively correlate with PMN/mL. Cytokines IL-1 α (A), IL-1 β (B), IL-6 (C), IFN- γ (D), TNF- α (E), and IL-10 (F) were measured via ELISA from BALF and compared to PMN counts from the same animal. n=8-11. Significant outliers were identified using the ROUT method and removed. (Correlation).



S4. Inflammatory cytokine production is negatively correlated with increased *S. parasanguinis* CFU in dual-infected animals. Cytokines IL-1 α (A), IL-1 β (B), IL-6 (C), IFN- γ (D), and TNF- α (E) were measured via ELISA from BALF and compared to the ratio between *S. parasanguinis* (SP) and *P. aeruginosa* (PA) from the same animal. n=8-11. Significant outliers were identified using the ROUT method and removed. (Correlation).



S5. *S. parasanguinis* colonized bronchial epithelial cells regardless of nitrite or *P. aeruginosa. S. parasanguinis* CFU from A) 16HBE with mPA08, B) CFBE with mPA08, C) 16HBE with PAO1, D) CFBE with PAO1. n=4-9, error bars represent standard deviation. *, p<0.05, ** p<0.01. (One-Way ANOVA, Dunnett's *post hoc* test).



S6. *S. parasanguinis* does not induce lactate dehydrogenase release. *S. parasanguinis* or the *poxL* mutant were inoculated on 16HBE (A) or CFBE (B) and Idh was measured before and after incubation and normalized. n=4-6, error bars represent standard deviation. *, p<0.05, **, p<0.01. (One-Way ANOVA, Dunnett's *post hoc* test).

