

primary studies - published RCT

Ivacaftor in cystic fibrosis with residual function: Lung function results from an N-of-1 study.

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Study design (if review, criteria of inclusion for studies)

Multi-centre two-arm parallel randomised control trial

Participants

Trial conducted across Europe/North-America enrolled 223 participants (January 2015 - August 2017). All participants were chronically colonised with *Pseudomonas aeruginosa* and were randomised 1:1 into two study-arms.

Interventions

The "usual-therapy group" received 2-weeks of IV ceftazidime 3g thrice-daily (for allergies: aztreonam 2g thrice-daily) and tobramycin 5-10mg.kg(-1) once-daily. The "microbiome-directed group" received the same usual-therapy plus an additional antibiotic with greatest presumed activity against the 2nd, 3rd and 4th most abundant genera present in the sputum microbiome, selected by a Consensus Expert Treatment Panel.

Outcome measures

The primary outcome was change in percentage of predicted FEV(1) (ppFEV(1)) at 14 days post initiation of antibiotics. Secondary outcomes examined ppFEV(1) at 7 days, 28 days, and 3 months; time-to-next exacerbation; symptom burden at 7 days; Health Related Quality of Life (HRQoL) at 28 days; and number of exacerbations and IV antibiotic days at 12 months.

Main results

149 participants had an eligible exacerbation (usual-therapy n=83, microbiome-directed therapy n=66). There was no difference between the groups for ppFEV(1) at day 14 (-1.1%, 95%CI -3.9 to 1.7; p=0.46), or ppFEV(1) measured at other time-points, or for time-to-next exacerbation (microbiome-directed versus usual-therapy Hazard Ratio 0.91 [95%CI 0.60 to 1.38; p=0.66]). The microbiome-directed group trended to have more IV days (median 42 versus 28; p=0.08) and more subsequent exacerbations (median 3 versus 2; p=0.044) the following year. There were no appreciable differences in symptom burden; however, HRQoL sub-scores were consistently worse in the microbiome-directed group (-4.3 points versus usual therapy (95%CI -8.3 to -0.3, p=0.033).

Authors' conclusions

The addition of a third antibiotic based on sputum microbiome sequencing analysis did not result in improved clinical outcomes.

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See also

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Keywords

Adult; Anti-Bacterial Agents; Bacterial Infections; Ceftazidime; Combined Modality Therapy; Infection; pharmacological_intervention; *Pseudomonas aeruginosa*; *Pseudomonas*; Respiratory Tract Diseases; Respiratory Tract Infections; Tobramycin; Intravenous; Cephalosporins; Aminoglycosides; microbiome;