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Published in: Annals of the American Thoracic Society

DOI: 10.1513/AnnalsATS.201506-333OC

Publication date: 2015

Document Version Peer reviewed version

Link to publication in Discovery Research Portal

Citation for published version (APA):

Finch, S., McDonnell, M. J., Abo-Leyah, H., Aliberti, S., & Chalmers, J. D. (2015). A comprehensive analysis of the impact of Pseudomonas aeruginosa colonization on prognosis in adult bronchiectasis. Annals of the American Thoracic Society, 12(11), 1602-1611. DOI: 10.1513/AnnalsATS.201506-333OC

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A Comprehensive analysis of the impact of *Pseudomonas aeruginosa* colonisation on prognosis in adult bronchiectasis

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Keywords: bacteria, bronchiectasis, mortality, severity, exacerbations

Running head: Prognostic impact of P. aeruginosa in bronchiectasis

Descriptor: 10.3 Chronic Bronchial Suppurative Diseases

Word count: 3173

Statement of contributions: All authors participated in the conception of the study, data collection, drafting and revising of the manuscript.

Acknowledgements: Supported by the European Bronchiectasis Network (EMBARC). EMBARC is a European Respiratory Society Clinical Research Collaboration and has received funding from the European Respiratory Society, Bayer HealthCare and Aradigm Corporation.

1 Rationale:

2 Eradication and suppression of *Pseudomonas aeruginosa* is a key priority in national guidelines for

3 bronchiectasis, and is a major focus of drug development and clinical trials. An accurate estimation

4 of the clinical impact of *P. aeruginosa* in bronchiectasis is, therefore, essential.

5 Methods

6 Data from 21 observational cohort studies comparing patients with *P. aeruginosa* colonisation to

7 those without were pooled by random effects meta-analysis with data collected for key longitudinal

8 clinical outcomes of mortality, hospital admissions, exacerbations and lung function decline along

9 with cross sectional outcomes such as quality of life.

10 Measurements and main results:

11 Studies included 3683 patients. *P. aeruginosa* was associated with a highly significant and consistent

12 increase in all markers of disease severity including mortality (odds ratio (OR) 2.95, 95% CI 1.98-4.40;

13 p<0.0001), hospital admissions (OR 6.57, 95% CI 3.19-13.51; p<0.0001) and exacerbations (mean

14 difference 0.97 per year, 95% Cl 0.64-1.30; p<0.0001). Patients with *P. aeruginosa* also had worse

15 quality of life using the St Georges Respiratory Questionnaire (mean difference 18.2 points, 95% CI

16 14.7-21.8; p<0.0001). There were also large differences in lung function and radiological severity.

Definitions of colonisation were inconsistent but findings were robust irrespective of the definitionused.

Conclusion: *P. aeruginosa* is associated with an approximate 3-fold increased risk of death and an
increase in hospital admissions and exacerbations in adult bronchiectasis.

21

22

23 Primary source of funding: European Bronchiectasis Network (EMBARC)

24

25 Introduction

26 Bronchiectasis is a chronic inflammatory lung disease characterised by recurrent cough, sputum 27 production and recurrent respiratory tract infections.(1) Failure of the mucociliary escalator and 28 innate antimicrobial defences leads to chronic bacterial colonisation of the airways.(2) Bacteria 29 provoke an inflammatory response that can further drive airways inflammation and airway 30 structural damage leading to the well described "vicious cycle" of bronchiectasis.(2,3) 31 While the majority of patients with bronchiectasis may be colonised with organisms that are upper 32 airway commensals such as Haemophilus influenzae and Streptococcus pneumoniae, a proportion of 33 patients become colonised with the opportunistic pathogen Pseudomonas aeruginosa. (4-6) 34 In cystic fibrosis (CF) bronchiectasis, it is well established that P. aeruginosa colonisation leads to a 35 more rapid deterioration in lung function and earlier mortality.(7) Consequently, P. aeruginosa 36 eradication is standard care in European CF centres.(8,9) The capability of P. aeruginosa to form 37 biofilms provide it with physical and chemical protection from the immune system and reduces its 38 exposure to systemically delivered antibiotics. (10,11) P. aeruginosa has the ability to rapidly adapt to 39 chronic infection in the lung and readily develops antimicrobial resistance. Management of P. 40 *aeruginosa* therefore represents a significant clinical challenge. 41 In bronchiectasis, conflicting data have been published on the independent contribution of P. 42 aeruginosa to long term prognosis and there remains a question of whether P. aeruginosa drives 43 disease progression or is simply a marker of existing severe disease. (12,13) Determining the 44 importance of *P. aeruginosa* to bronchiectasis morbidity and mortality is important, as there are few 45 evidence based treatments for bronchiectasis.(14) Current therapeutic development is heavily 46 influenced by CF and is therefore largely targeted towards treatment of P. aeruginosa 47 infection.(15,16) Therefore from clinical, drug development and regulatory perspectives it is important to have a comprehensive understanding of the impact of *P. aeruginosa* on outcomes in 48 49 bronchiectasis.

- 50 We therefore undertook a systematic review to determine whether colonisation with *P. aeruginosa*
- 51 influences future prognosis and/or is associated with cross-sectional features of severity.

53 METHODS

54 The present study was a systematic review and meta-analysis conducted and reported according to

55 MOOSE (meta-analysis of observational studies in epidemiology) guidelines.(17)

56 Search Criteria

- 57 The study was based on a search of the PUBMED database for articles evaluating the prognostic
- 58 impact of colonisation with *P. aeruginosa*. The following search strategy was used: ("Pseudomonas"
- 59 OR "aeruginosa") AND ("bronchiectasis") followed by ("prognosis" or "mortality") and
- 60 ("bronchiectasis"). The search included articles published between January 1980 and January 2015.
- 61 No language criteria were applied. Full articles of potentially appropriate abstracts were reviewed.
- 62 Only peer reviewed data were included. Conference abstracts were excluded. The search was
- 63 repeated in EMBASE and Web of Science to obtain any articles missed by the initial search. The
- 64 search strategy was supplemented by reviewing of the reference lists, bibliographies including the
- 65 British Thoracic Society guidelines and investigator files.
- 66

67 Data extraction

Non relevant studies were excluded based on review of the title and abstract. Article reviewing was performed independently by two investigators ((two out of SF, ,MM, AHL, SA and JC) who conducted data extraction and quality assessment from studies meeting the inclusion criteria. All investigators have experienced of meta-analysis and training in literature review. Any disagreement between investigators was resolved independently by a third investigator. Additional unpublished data were obtained from study authors where possible. Where data were presented only as medians, means with standard deviation were estimated according to the formula of Hozo et al.(18)

75

77 Study inclusion and exclusion criteria

All studies were considered eligible if they fulfilled the following criteria: original publications;
inclusion of a cohort of patients with computed tomography diagnosed bronchiectasis not due to
cystic fibrosis; inclusion of patients with *P. aeruginosa* colonisation and a comparator population
without *P. aeruginosa* colonisation; reporting of one of the study outcomes which were determined *a priori* (described below).
Definitions of *P. aeruginosa* colonisation were obtained from the source studies and were not prespecified.

85 As the aim of this study was to compare *P. aeruginosa* colonised patients compared to non-

86 colonised patients, we excluded any studies which provided data only for a single population. We

also excluded case reports; review articles, editorials and letters without original data.

88

89 Study outcomes

90 Primary analysis

91 Our hypothesis was that *P. aeruginosa* colonisation would be associated with globally worse clinical

92 outcome when compared to patients without *P. aeruginosa* colonisation. Outcomes were split into

93 longitudinal clinical outcomes determined during follow-up, and cross-sectional outcomes. The

94 primary longitudinal outcome was all-cause mortality. Secondary outcomes were: hospital

95 admissions, exacerbation frequency, decline in forced expiratory volume in 1 second (FEV₁) and the

- 96 prognostic impact of *P. aeruginosa* eradication therapy.
- 97 Cross sectional outcomes were: FEV₁ % predicted, forced vital capacity (FVC), radiological

98 involvement and quality of life (QoL). A descriptive analysis of the methods of defining *P. aeruginosa*

99 colonisation in the literature was also considered a pre-specified secondary end-point.

- 100 Anticipating that studies would have different lengths of follow-up to determine survival, we pre-
- 101 specified that data could be pooled where equal follow-up was demonstrated between P.
- 102 aeruginosa colonised and non-colonised patients.
- 103

104 Quality assessments

- 105 The quality of each study was independently assessed according to the criteria described by Hayden
- 106 et al, which are widely used for assessing the quality of observational studies in meta-
- analysis.(19,20) The agreement between the two reviewers (two of SF, AHL and JC) was measured
- 108 using the kappa statistic. Publication bias was determined by visual inspection of funnel plots and
- 109 Eggers test.
- 110

111 Sensitivity analysis

- 112 A priori we identified possible factors that may be major sources of bias and planned subanalyses for
- 113 the follow; 1) Analysis according to different definitions of *P. aeruginosa* colonisation e.g single
- isolate versus multiple isolatins; 2) Comparison of *P. aeruginosa* vs *H. influenzae* colonised patients
- 115 compared to comparisons of *P. aeruginosa* colonised vs non-colonised patients; 3) Data derived
- 116 from high quality and prospective studies.
- 117

118 Statistical analysis

- 119 The primary outcome of the relationship between *P. aeruginosa* colonisation and mortality was
- 120 displayed as odds ratios (OR) with 95% confidence intervals (95% CI). ORs were pooled using a
- 121 Mantel-Haenszel random effects model. The same analysis was used for hospital admissions.

122	Continuous variables such as quality of life, lobar involvement, pulmonary function tests and
123	exacerbations were compared by pooling mean differences by the inverse of their variance. As
124	above, random effects meta-analysis was used due to expected heterogeneity between studies. To
125	analyse for possible effect modifiers, such as study quality or definitions of colonisation, we
126	compared OR's using interaction testing as described.
127	Statistical heterogeneity was assessed using the Cochran Q (χ^2) test and the Higgins I ² tests. For the
128	Cochran Q test, p<0.1 was considered to represent significant heterogeneity. For the Higgins test, I^2
129	<25% indicated low heterogeneity, 25–50% moderate and >50% severe heterogeneity. Analyses
130	were conducted using Review Manager 5 (Cochrane Collaboration) and SPSS version 21 for windows
131	(Chicago, IL, USA).
132	
132	
134	
135	
136	<u>RESULTS</u>
137	
138	The results of the literature review are shown in figure 1. The majority of studies were rejected
139	because they did not deal specifically with patients with bronchiectasis not due to CF or did not
140	evaluate severity or outcomes. Of 55 articles selected as relevant, 21 studies had valid data for
141	inclusion and were pooled in the meta-analysis.(6,12,13,21-36) One study contained data for 5
142	cohorts and each cohort was considered separately for the purposes of this analysis on the basis that
143	they were independent cohorts (total 25 cohorts).(24)
144	
145	

146	Characteristics of the 21 included studies are shown in table 1. 10 studies were from the UK
147	(12,21,23,24,26,28,31,32) and overall 16 cohorts were from Europe. There were no cohorts from
148	North America.
149	
150	Definitions of chronic <i>P. aeruginosa</i> colonisation were highly heterogeneous. The most frequent
151	definition used was 2 positive cultures at least 3 months apart over 12 months. 5 studies reported
152	patients with a single positive culture as "colonised". In all, 8 different methods of defining P.
153	aeruginosa colonisation were identified in addition to 3 studies where the definition was not stated.
154	According to the quality assessment, 6 studies were rated as high quality, 8 as intermediate, and 7 as
155	low quality (Kappa 0.73). None of the analyses showed evidence of publication bias.
156	
157	The total number of patients studied was 3683 with a rate of <i>P. aeruginosa</i> colonisation (according
158	to study definitions) of 21.4%. Comparator populations were almost universally mixed populations
159	of bronchiectasis not colonised with <i>P. aeruginosa</i> .
160	
161	Impact of <i>P. aeruginosa</i> on longitudinal outcomes
162	Primary outcome: All-cause Mortality
163	Mortality was available as an outcome in 8 cohorts (24-27) of which 5 cohorts were derived from a
164	single study (24). Follow-up duration ranged from 1 year to 14 years. Mortality for patients with P.
165	aeruginosa ranged from 7.7% at 1 year, 13.6% at 2 years to 30-50% at 5 years. Corresponding
166	mortality rates for patients without <i>P. aeruginosa</i> were 0% at 1 year, 7% at 2 years and 9-15% at 5
167	years. All studies showed a higher risk of mortality associated with <i>P. aeruginosa</i> colonisation. The
168	pooled OR for mortality was 2.95 (95% CI 1.98-4.40; p<0.0001). Heterogeneity tests were not
169	statistically significant. This is shown in figure 2.

170 Sub-analyses confirmed this association in high quality studies (OR 3.64, 95% CI 1.75-7	·7.55; p=0.0005
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- 171 n=1433), prospective studies and excluding studies with <3 (OR 2.82, 95% CI 1.94-4.11; p<0.0001,
- 172 n=1994) and >6 years follow-up (OR 3.14, 95% Cl 1.83-5.33; p<0.0001, n=1894).
- 173

174 Hospital admissions

- 175 This analysis included 7 cohorts with 1628 participants (23,24,29). Hospital admission rates for
- patients with *P. aeruginosa* varied from 41% at 1 year to 75% at 4 years. Corresponding hospital
- admission rates in patients without *P*. aeruginosa were 15% at 1 year and 28.5% at 4 years. P.
- 178 *aeruginosa* was associated with a marked increase in the risk of hospital admissions pooled OR
- 179 6.57 (95% Cl 3.19-13.51; p<0.0001). There was significant heterogeneity which was not resolved on
- 180 limiting studies by quality, prospective design or length of follow-up. Insufficient data was available
- 181 to evaluate additional impacts such as length of hospital stay or economic impacts of hospitalisation.
- 182 Data are shown in figure 3.

183

184

185 **Exacerbations per year**

- 186 All available data were presented as mean exacerbations per patient per year. The 9 cohorts which
- 187 recorded this information gave a pooled increased frequency of just under 1 exacerbation per
- patient per year mean difference 0.97 (95%Cl 0.64-1.30; p<0.0001) with no significant
- heterogeneity.(6,23,24,29,30) This is shown in figure 4. No significant differences in effect were
- 190 observed in high quality studies, prospective studies or in sub-analyses based on the definition of *P*.
- 191 *aeruginosa*.
- 192

193 Lung function decline

194	There were limited data available on lung function decline. One study reported lower lung function
195	in PA colonised patients but no differences in long term lung function decline.(13) Another study
196	reported a mean decline of 52ml per year in <i>P. aeruginosa</i> patients(12) with a further study
197	reporting a mean decline of 123ml per year. (38) The available data included only 41 patients with P.
198	aeruginosa colonisation. Consequently no attempt was made to pool the data.
199	
200	Pseudomonas eradication treatment
201	No randomised studies of <i>P. aeruginosa</i> eradication treatment were identified. A non-randomised
202	observational study (n=30) reported an initial eradication success rate of 80%, and 43% after a
203	median of 6 months.(39) This was associated with a reduction in exacerbations from 3.93 per year to
204	2.03 per year. No control population was available for comparison with no data on the spontaneous
205	clearance rate that would have occurred without treatment.
206	
207	
208	Cross sectional association between P. aeruginosa and markers of severe disease
209	Patient characteristics
210	In cross-sectional studies we observed that patients with <i>P. aeruginosa</i> infection were on average 3
211	years older than non-colonised patients (mean difference 3.1 years, 95% Cl 0.9-5.4,p=0.007, l ² =48%).
212	Interestingly there was a statistically significant association between male gender and <i>P. aeruginosa</i>
213	colonisation, OR 1.39 95% Cl 1.09- 1.75, p=0.009, l ² =0%.
214	Quality of life
215	The only data that were available for QoL used the St.Georges Respiratory Questionnaire (SGRQ).

The SGRQ is a validated questionnaire in patients with bronchiectasis that has been widely used with

- an accepted increment of 4 points demonstrating clinical significance.(37) 4 studies reported data
- 218 for SGRQ, with a mean difference of 18.2 points (95% CI 14.7-21.8; p<0.0001, n=1041).(22,24,28)
- 219 There was no heterogeneity between studies (I²=0%). No data were available for other
- 220 questionnaires, including the QoL-Bronchiectasis questionnaire.
- 221

222 Lung Function- FEV₁ and FVC

- 223 As expected, patients with *P. aeruginosa* colonisation had worse cross-sectional lung function
- compared to patients without *P. aeruginosa*. 17 studies reported valid data for FEV₁ with all showing
- worse lung function in the *P. aeruginosa* group ranging from -1.4% to -29%.
- 226 (6,12,13,21,22,24,28,29,30,31,33,34,36) The pooled mean difference was 15.0% (95% CI -18.7 to -
- 11.3; p<0.0001). There was significant heterogeneity but this was no longer statistically significant
- after excluding 1 study that defined *P. aeruginosa* presence by PCR.(6) 9 cohorts presented data for
- 229 FVC with a pooled mean difference of -9.4% (95% CI -14.3 to -4.5%;p=0.005, n=1453).
- 230 (12,24,28,29,33,34)
- 231

232 Radiological severity

- 233 Although multiple severity scoring systems have been utilised in bronchiectasis, the only variable
- which was studied in more than one study was the number of lobes involved on CT. This data were
- available in 9 cohorts. (24, 29, 30, 32, 34, 35) The mean difference between *P. aeruginosa* colonised and
- non-colonised was 1.4 lobes (95% CI 0.93-1.86; p<0.0001). Nevertheless all studies reported worse
- radiological severity in *P. aeruginosa* colonised patients.

238 Sensitivity analyses

- 239 Limiting the analysis to only those studies that used the most robust definition of *P. aeruginosa*
- 240 colonisation, requiring at least 2 positive sputum samples over a 12 month period, showed very

similar results to the primary analysis with ORs for mortality of 3.46, 95% Cl 1.96-.6.08; p<0.0001;

hospital admissions 7.22, 95% CI 2.88-18.09; p<0.0001 and exacerbations mean difference 0.87, 95%

- 243 CI 0.59-1.15; p<0.0001 (p>0.5 when comparing odds ratios using interaction testing compared to the
- 244 overall cohort).
- 245
- 246 8 cohorts provided data that could be used to directly compare the outcomes of patients colonised
- 247 with *P. aeruginosa* versus *Haemophilus influenzae*. The findings were highly consistent with the main
- analysis, with an increase in mortality associated with *P. aeruginosa* (OR 4.00, 95% CI 2.28-7.02;
- 249 p<0.001), increased rate of hospital admissions (OR 6.75, 95% CI 3.98-11.45; p<0.001), increased
- exacerbations (mean difference 0.99 (95% CI 0.54-1.43; p<0.0001) and low FEV₁ (mean difference -
- 251 11.4, 95% CI -14.8 to -7.9; p<0.0001).

252 DISCUSSION

253 The management of bronchiectasis patients with P. aeruginosa colonisation is challenging and a 254 large proportion of the current therapeutic development in bronchiectasis is focussed towards 255 management of *P. aeruginosa* infection.(14-16) In particularly there are intensive efforts in the field 256 of inhaled antibiotics to develop a licensed therapy for P. aeruginosa infection in 257 bronchiectasis.(15,16) Therefore an accurate assessment of the prognostic impact of *P. aeruginosa* 258 in bronchiectasis is important for clinicians, for drug developers and for regulators. This analysis 259 provides a detailed insight into the impact that *P. aeruginosa* colonisation has on key clinical 260 outcomes in bronchiectasis. In addition, P. aeruginosa colonisation was associated with multiple 261 cross-sectional markers of disease severity. It can therefore be said that P. aeruginosa is both a 262 marker of severe disease, and is associated with a worse long term prognosis. Bronchiectasis has 263 historically been a neglected condition, described in the ERS white book as one of the most 264 neglected diseases in respiratory medicine. (40) As a result, there have been few large cohort studies. 265 The value of meta-analysis therefore is to combine the available data from existing small studies to 266 give a more accurate estimate of the disease impact. 267 The most striking finding within this analysis is the impact of *P. aeruginosa* on all-cause mortality. 268 Our analysis identifies a 3-fold increase in the risk of death with P. aeruginosa colonisation. P. 269 aeruginosa was also associated with a greatly increased the risk of hospital admissions and 270 exacerbation frequency by a rate of 1 exacerbation per patient per year. This finding was robust 271 regardless of the definition used and was consistent across all cohorts. These results strengthen the 272 view that patients with P. aeruginosa require specific treatment to reduce the risk of long term 273 morbidity and mortality and that *P. aeruginosa* colonisation status should play a key role in the 274 assessment of disease severity.(14)

The increased exacerbation frequency and hospital admissions demonstrates a measurable
healthcare cost associated with *P. aeruginosa* colonisation. Each additional exacerbation results in

further antibiotic use with associated risks and side-effects as well as increased potential for the
development of antimicrobial resistance. Exacerbations are associated with reduced productivity
through absence from work and are associated with poorer QoL and potential lung function
decline.(24,38,41) Hospital admissions may reflect more severe exacerbations or the development
of resistance to oral antibiotic agents necessitating intravenous antibiotic therapy.(24) The ability of *P. aeruginosa* to develop antibiotic resistance is inevitably enhanced by repeated antibiotic
exposure.(29)

284 Our analyses of quality of life, lung function and radiological severity were cross-sectional and can 285 therefore only be considered hypothesis generating in terms of the impact of *P. aeruginosa* on these 286 outcomes over the long term. Nevertheless the impact on QoL demonstrated in this analysis is 287 striking. The 18 point decrement in the SGRQ demonstrated in patients with P. aeruginosa 288 colonisation reflects a dramatic worsening of QoL. Given our observation that patients with P. 289 aeruginosa had reduced lung function and more widespread radiological disease on imaging it is 290 difficult to determine what proportion of this difference in QoL is directly attributable to P. 291 aeruginosa. All of the analyses described herein are subjective to the same limitation, that P. 292 aeruginosa may be to some extent a reflection of the severity of underlying disease rather than a 293 directly cause of disease progression. The only way to conclusively prove or quantify the 294 independent effects of *P. aeruginosa* on outcome is likely to be through a large randomised 295 controlled trial of *P. aeruginosa* eradication treatment which has been highlighted as a clear priority 296 for the bronchiectasis research community.(42) Demonstrating that mortality, hospital admissions, 297 exacerbations, QoL and lung function are improved or cease to decline after successful eradication 298 would clearly demonstrate the independent impact of *P. aeruginosa*. A strength of our analysis is 299 that it provides the most precise estimates to date of P. aeruginosa prevalence and impact in order 300 to power future trials.

Current national guidelines for bronchiectasis recommend eradication treatment for new isolation of
 P. aeruginosa, largely based on recommendations for CF.(8,14,43) Data in bronchiectasis is limited to
 date and further research is greatly needed.

304 Important gaps in the literature identified through this analysis include an absence of data available 305 outside Europe and Australasia with a large proportion of included data from the UK; broad, 306 representative registries for patients with bronchiectasis are needed internationally. Few studies 307 examining lung function decline were identified, and those that were found were small with 308 inconsistent results. We would recommend further large studies of lung function decline in 309 bronchiectasis. There is a lack of data describing the impact of organisms other than *P. aeruginosa* in 310 bronchiectasis and in particular comparing the outcomes of P. aeruginosa colonised patients with 311 those colonised with the most common bronchiectasis pathogens such as H. influenzae or Moraxella 312 catarrhalis. Such data would be valuable as recent reports suggest that these patients do have a 313 worse outcome compared to non-colonised patients, but to a lesser extent than P. aeruginosa.(24) 314 For example in the Bronchiectasis Severity Index, 3 points are awarded to patients with P. 315 aeruginosa colonisation and 1 point to patients colonised with other pathogens. (24) For this meta-316 analysis, we were able to identify 8 cohorts with data to compare outcomes between P. aeruginosa 317 and H. Influenzae colonised patients and these confirmed the significantly worse clinical outcomes 318 associated with P. aeruginosa.

There is a need from both a clinical and research perspective to define chronic bacterial colonisation in bronchiectasis as this analysis identified 8 different methods of defining *P. aeruginosa* colonisation in bronchiectasis studies. The most frequently used definition was 2 or more positive cultures at least 3 months apart in 12 months. This should be standardised across studies to increase our ability to generalise results between studies and healthcare systems. Our data were almost entirely based on traditional bacterial culture and recent studies have increasingly used quantitative PCR or characterisation of the microbiome through sequencing of the 16s ribosomal RNA subunit to

determine bacterial colonisation status.(6,36,44) This method is significantly more sensitive for the
detection of *P. aeruginosa* with Rogers et al. demonstrating very poor correlation between culture
and PCR for *P. aeruginosa* detection: 91/107 patients in this study were positive for *P. aeruginosa*versus 31/107 by culture.(6) For this reason, further studies of the role of PCR in *P. aeruginosa*detection and to confirm eradication, would be beneficial.

331 The word colonisation in this context is perhaps misleading. Colonisation implies a benign state

defined by absence of tissue invasion or tissue damage. The term 'chronic infection' may be more

appropriate given the clearly established association between the presence of bacteria and airway

inflammation and the worse clinical outcomes observed in the presence of *P. aeruginosa*.

335

In summary, *P. aeruginosa* colonisation is associated with increased mortality, hospital admissions
and exacerbations, and is associated with worse QoL. As such, new Isolation of *P. aeruginosa* should
be considered a highly significant clinical event and followed up with repeated cultures and attempts
to eradicate in line with guideline recommendations.

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457	Figure 1. Flow chart illustrating the process and results of the literature review. Abbreviations: QoL=
458	quality of life; FEV ₁ = forced expiratory volume in 1 second.
459	
460	Figure 2. Association between <i>P. aeruginosa</i> colonisation and mortality in bronchiectasis.
461	Abbreviations: OR= odds ratio, M-H= Maentel-Haentzel, IV= inverse variance, CI= confidence
462	interval,
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464	
465	Figure 3. Association between <i>P. aeruginosa</i> colonisation and hospital admissions in bronchiectasis.
466	Abbreviations: OR= odds ratio, M-H= Maentel-Haentzel, IV= inverse variance, CI= confidence interval
467	
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469	Figure 4. Exacerbation frequency comparing patients with and without <i>P. aeruginosa</i> colonisation.
470	Abbreviations: OR= odds ratio, M-H= Maentel-Haentzel, IV= inverse variance, CI= confidence interval
471	
472	Figure 5. FEV ₁ % predicted compared between patients with <i>P. aeruginosa</i> colonisation and patients
473	without <i>P. aeruginosa</i> colonisation. Abbreviations: OR= odds ratio, M-H= Maentel-Haentzel, IV=
474	inverse variance, CI= confidence interval