- 1 Title page: Pneumococcal Colonisation Rates in Patients Admitted to a UK Hospital
- 2 with Lower Respiratory Tract Infection a prospective case-control study.
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24 Abstract

25	Background: Current diagnostic tests are ineffective at identifying the aetiological pathogen
26	in hospitalised adults with lower respiratory tract infection (LRTI). The association of
27	pneumococcal colonisation with disease has been suggested as a means to increase
28	diagnostic precision. We compared pneumococcal colonisation rate and density of nasal
29	pneumococcal colonisation by a) classical culture and b) quantitative real time <i>lytA</i>
30	Polymerase Chain Reaction (qPCR) in patients admitted to hospital in the UK with LRTI
31	compared to control patients.
32	Methods: 826 patients were screened for inclusion in this prospective case-control study. 38
33	patients were recruited, 19 with confirmed LRTI and 19 controls with another diagnosis.
34	Nasal wash (NW) was collected at the time of recruitment.
35	Results: Pneumococcal colonisation was detected in 1 LRTI patient and 3 controls (p=0.6)
36	by classical culture. Using <i>q</i> PCR pneumococcal colonisation was detected in 10 LRTI
37	patients and 8 controls (p=0.5). Antibiotic usage prior to sampling was significantly higher in
38	the LRTI than control group 19 v. 3 (p<0.001). Using a clinically relevant cut-off of >8000
39	copies/ml on qPCR pneumococcal colonisation was found in 3 LRTI patients and 4 controls
40	(p > 0.05).
41	Conclusions: We conclude that neither prevalence nor density of nasal pneumococcal
42	colonisation (by culture and q PCR) can be used as a method of microbiological diagnosis in

hospitalised adults with LRTI in the UK. A community based study recruiting patients prior to
antibiotic therapy may be a useful future step.

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48 Introduction

Recent studies suggest that detection and quantification of nasal pneumococcus by 49 50 quantitative real time lytA Polymerase Chain Reaction (qPCR) could be used to identify pneumococcus as the aetiological pathogen in adults with pneumonia [1] and could be 51 useful as a disease severity marker [2]. In that study, South African patients with community 52 acquired pneumonia (CAP) were more frequently colonised than controls using classical 53 culture (44.9 v. 11.7%) and gPCR (62.8 v. 19.8%) and, in addition, patients with 54 pneumococcal CAP were also noted to have higher colonisation density than asymptomatic 55 controls [1]. By applying a cut off of 8000 copies/ml to the qPCR data Albrich et al [1] found 56 that 52.5% of patients were considered to have pneumococcal CAP, compared with 27.1% 57 diagnosed using standard tests. 58

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60 The association of pneumonia and pneumococcal colonisation has been previously noted in children, in whom those with radiological pneumonia were more frequently colonised with 61 62 pneumococci than those without [3] and had higher density colonisation than those with 63 bronchitis or without disease [4]. In contrast, in the elderly very low colonisation rates have 64 been shown; 0.3% in pneumococcal vaccine naive hospitalised Australians (by classical culture) [5] (of which 10 had respiratory infection) and 2.3% in a Portuguese community 65 66 cohort [6]. In developed countries, pneumococcal colonisation rates in healthy adults are 67 between 1 - 18%, and are affected by age, immune status, antibiotic use, household 68 composition and contact with children [7, 8]. There are no published data on pneumococcal colonisation in hospitalised patients with respiratory infection in the UK. 69 70 71 We therefore aimed to determine the rate and density of pneumococcal colonisation by a)

classical culture and b) *q*PCR in hospitalised adult patients with LRTI when compared with
 age and gender-matched controls in a developed country setting.

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75 Materials and Methods

76 Screening and Recruitment

77 We recruited hospitalised adults with LRTI at the Royal Liverpool and Broadgreen University 78 Hospital from January - July 2013 within 72 hours (hrs) of admission. The syndrome of LRTI was defined as; symptoms of respiratory infection with clinical signs +/- radiological 79 consolidation; therefore meeting a British Thoracic Society (BTS) definition of pneumonia as 80 81 used in community (GP) practice. Clinical signs of LRTI included ≥2 of: cough, breathlessness, pleuritic chest pain, fever, increased or new sputum production. Exclusion 82 83 criteria were: patients with infective or non-infective exacerbations of chronic obstructive 84 pulmonary disease (IECOPD), asthma or bronchiectasis (without radiological consolidation), aspiration pneumonia, oxygen saturations <86% on air, suspected tuberculosis (TB) and 85 86 neutropenia. Patients with IECOPD were excluded as exacerbations are known to commonly be due to viruses (up to 60%) [9, 10], Haemophilus influenzae (up to 33%) [11] and 87 88 Moraxella catarralis (around 10%) [12] rather than pneumococcus. Patients who had been a hospital inpatient for ≥72 hrs or had recently been discharged from hospital ≤14 days before 89 90 were excluded since it is likely their nasal flora would have altered due to hospital 91 exposures. Patients with oxygen saturations <86% on air were excluded since it was felt unsafe to remove their oxygen in order to perform a nasal wash (NW). 92 93 A carefully selected control group of hospitalised patients with no signs of respiratory 94 infection were recruited within 7 days (where possible or as soon after as possible) of the 95 LRTI patient. The control group were matched for age (within 10 years of the LRTI patient) and gender. Exclusion criteria were: oxygen saturations <86% on air, neutropenia, ≥ 7 days 96 97 after admission and recent hospital discharge ≤14 days. The study team were in regular communication with the hospital capacity team, the ward 98

based case managers, nursing and medical co-ordinators in accident and emergency (A&E),

100 the acute medical admissions unit (AMAU) and the respiratory wards from Monday to

Thursday. Through regular education, information dissemination events, and daily interaction
with the study team, key staff were made aware of the study and its aims, objectives, and
potential participants. A list of potential participants was generated on a daily basis in
combination with these personnel. To recruit LRTI patients we targeted screening to AMAU,
respiratory and infectious disease wards, for control participants we targeted surgical wards.

Patient eligibility was confirmed by review of the medical records; with permission of the
attending team, patient consent was sought prior to recruitment. Baseline clinical data of
age, gender, history of presenting complaint, past medical history, vaccination history,
antibiotic prescription, and contact with children (defined as at least alternate day contact
with children aged ≤ 10yrs) were recorded. NW and urine samples were collected within 12
hours of recruitment. The study was approved by Liverpool East, North-west NHS Research
Ethics Committee (12/NW/0713) and registered with ClinicalTrials.gov (NCT01861184).

113 Sampling

114 NW was collected on the day of recruitment with a maximum of 20 mls of normal saline 115 instilled into the nasopharynx as previously described [13, 14]. A minimum of 5 mls of normal 116 saline was recovered and processed in all cases. Briefly, samples were immediately centrifuged at high speed (3345 x g) to obtain a bacterial pellet [14]. Pellets were 117 resuspended in 100µl of skimmed-milk tryptone glucose glycerol (STGG) medium and the 118 119 total volume of the suspension was determined. Samples were then serially diluted on blood agar and CFUs/mL of NW were determined the next day. In a subset of samples that 120 121 exceeded 7 mls a proportion of the sample (3 – 5 mls) was removed and centrifuged at 836 x g to obtain cellular material after which the supernatant was re-added to the rest of the 122 123 sample for the high speed spin.

124 DNA extraction and qPCR

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125	DNA was extracted from 200 μl of the NW bacterial pellet stored in STGG using the QIAamp
126	DNA mini kit and the Centers for Disease Control protocol [15]. Briefly, the pellet was
127	resuspended in 200 μI TE buffer containing 0.04 g/ml lysozyme and 75 U/ml mutanolysin
128	(Sigma) and incubated at 37°C for 1 hr. Following incubation, 20 μl of proteinase K and 200
129	μI of Buffer AL were added to the sample which was vortexed and then incubated at 56°C for
130	30 min. The sample was then centrifuged briefly and 260 μI of ethanol was added. All
131	subsequent steps followed the manufacturer's instructions. DNA was eluted in 100 μl of
132	QIAGEN elution buffer and stored at -20°C.
133	Colonisation density was determined by targeting the pneumococcal autolysin lytA gene
134	[16]. A no-template control, a negative extraction control (parallel extraction of TE buffer) and
135	a S. pneumoniae (BHN418) positive control were included in each run. DNA was amplified
136	with the Mx3005P system (Stratagene) and data was analysed using the instrument
137	software. A sample was considered positive if both duplicates had a cycle threshold (C_T)
138	value below a mean 35. Values of >8000 copies/ml were considered clinically relevant as
139	per Albrich and colleagues [1].
140	Binax
141	An immunochromatographic membrane test (ICT) (BinaxNOW Streptococcus pneumoniae;
142	Binax) was performed on all patient's unconcentrated urine specimens, according to the
143	manufacturer's recommendations.
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149 Results

150 Screening and recruitment

151 We screened 826 patients and recruited 19 LRTI and 19 control (age, gender and season 152 matched) patients. 217 were 'potential' LRTI patients, of which 198 were not eligible (Figure 1). 58 patients did not have a syndrome of LRTI (acute exacerbation [AE] COPD n=22, AE 153 bronchiectasis n = 5, AE asthma n=3, AE pulmonary fibrosis n= 1 and alternative diagnoses 154 155 including pulmonary embolus (PE), congestive cardiac failure (CCF), sepsis of unknown cause and adult acute respiratory distress syndrome (ARDS) n = 30), 36 patients did not 156 157 have capacity to consent (predominately due to dementia or acute delirium), 48 patients 158 were identified >72 hours after admission and 17 after a recent hospital discharge ≤14 days 159 before, 20 patients declined to participate and 2 'next of kin' refused permission for their 160 relative to participate, 10 oxygen saturations <86% on air, 12 aspiration pneumonia, 14 were excluded for other reasons. 161

We planned to recruit 100 patients to each arm of the study, but stopped recruiting on the 162 163 grounds of futility after interim analysis noted 100% antibiotic usage prior to recruitment in 164 the LRTI group and low rates of colonisation (on culture) and due to high screen failure rate (778/816, 95.4%). Recruiting age matched controls was difficult, especially for the younger 165 LRTI patients (aged 36 - 46 yrs old). In 9 cases, the time between the recruitment of the 166 LRTI patient and control was >7 days (range 9 - 43). We know from our Experimental 167 168 Human Pneumococcal Colonisation (EHPC) studies that antibiotic usage terminates 169 pneumococcal colonisation, meaning that continued recruitment in this population was 170 unethical.

171 Sampling: Rate of colonisation

172 All patients were successful at providing a nasal sample. One patient was unable to replicate

- 173 the NW technique (as per protocol) and therefore had a nasopharyngeal swab instead. NW
- volumes were not significantly different between LRTI patients and controls (Table 1).

- 176 3 controls (p=0.6). Using *q*PCR 10 LRTI patients and 8 controls were positive (p=0.516)
- 177 [Table 2)]. One of the controls was positive for colonisation by culture but considered
- 178 negative by qPCR as the C_T value was >35.

179 Sampling: Density of colonisation by *q*PCR

- 180 For qPCR a cut off value of >8000 copies/ml was used to define clinical relevance [1]. In our
- 181 study, 3 LRTI patients and 4 controls had values >8000 copies/ml. Of the 3 LRTI patients,
- 182 only 1 was culture positive; of the 4 controls, 2 were culture positive (Table 2). Of the 4
- patients overall who were culture positive, 3 had >8000 copies/ml, 1 in the LRTI and 2 in the
 control group.

185 Clinical data

- 186 Antibiotic usage prior to sampling was significantly higher in LRTI patients than controls 19 v.
- 187 3 (p<0.001). Radiological consolidation was present in 7 out of 19 LRTI patients; only 2 out
- 188 of 38 urine samples were positive using BinaxNOW. None of the LRTI patients recruited
- 189 were pneumococcal sputum or blood culture positive. There were no statistical differences
- 190 between the groups with regards to smoking, contact with children age <10yrs old or 23 PPV
- 191 *pneumovax* vaccination (Table 1).
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198 Discussion

The anticipated high rate of pneumococcal colonisation (by culture +/- qPCR) in the LRTI 199 200 group was not found, given that antibiotic usage (pre-admission/pre-recruitment) was 201 significantly different between LRTI and control groups, with all LRTI patients having received at least 2 doses prior to NW, this is likely to have resulted in culture negativity. 202 203 However we also found no significant difference in colonisation rates using gPCR and colonisation density between the LRTI and control groups. There was also no significant 204 difference in colonisation rates in polysaccharide vaccinated (23 PPV Pneumovax) and 205 206 unvaccinated patients, consistent with previous literature reporting that the vaccine does not 207 protect against colonisation [17, 18]. 208 Large numbers of patients were referred as 'potential LRTI'. Alternative diagnoses such as 209 PE, CCF, non-infective exacerbation of pulmonary fibrosis, sepsis of unknown cause and aspiration pneumonia were common. This diagnostic imprecision has important implications 210 211 for the use of NW as a diagnostic technique since it would lead to many inappropriate samples being collected. We have previously demonstrated that confusion is common in 212 213 LRTI patients (>20%) [19]. LRTI rates increase with age (63% CAP admissions were aged 214 >65 and 25% ≥85yrs old) [20] as do rates of comorbidities (including dementia), therefore recent hospital admission is also common. 215 216 The main strength of this study is the large number of screened patients; the LRTI patients 217 are well phenotyped and the controls are matched in age, gender and time with similar 218 smoking habits, 23 PPV pneumovax vaccination rates and child contact. Our cohort was not

- 219 'CAP' by strict definition of radiological consolidation, instead a broad study group of LRTI
- 220 was chosen due to its clinical relevance in UK hospital practice and admissions, making
- 221 these results very generalisable. Nationally, GP antibiotic prescribing for LRTI is very high,
- 222 but lower for clinically diagnosed CAP (due to usual immediate hospitalisation) [21].

223 Accurately diagnosing pneumonia is challenging; inter-doctor variability in reporting of 224 radiological pneumonia is common [22]. Studies of patients that have radiological 225 pneumonia as an inclusion criterion may be less applicable to everyday hospital medicine. 226 LRTI may be a more useful term for this clinical syndrome, particularly in instances where guidelines suggest clinical rather than radiological diagnosis [20]. Liverpool is in the North-227 west of England, and has the secodrnd highest LRTI rate (age standardised episodes/1000 228 229 person years) and the third highest CAP rate nationally. [21] It is therefore an ideal area for 230 recruiting to respiratory infection studies, although community antibiotic prescription rates 231 are high. The Royal Liverpool hospital has ~1400 admissions per year that are coded as 232 'pneumonia', approximately 20% of these cases are not community acquired or have no 233 radiological features of pneumonia. 234 Limitations of the study include that this is a single centre study which may reduce the 235 generalisability of the results specifically in areas where community antibiotic prescription rates are lower, that we were unable to fully recruit to the study despite high numbers 236 237 screened and that the NW technique, rather than nasopharyngeal swab, for pneumococcal 238 isolation may not have been ideal in this elderly population, since the research nurses noted 239 poor technique and lower yields than in the cohort of healthy volunteers in which we

240 commonly use this technique (data not shown). Nevertheless, patient comfort is higher [23]

241 and sensitivity for colonisation density is very high [24]. We know from our Experimental

242 Human Pneumococcal Colonisation (EHPC) studies that antibiotic usage terminates

243 pneumococcal colonisation; after interim analysis noted 100% antibiotic usage in the LRTI

244 group prior to recruitment and low rates of colonisation (on culture), the study was stopped

as continued recruitment in this population was unethical.

246 Previous studies have shown colonisation rates of 44.9% and 62.8% in patients with

- 247 radiologically confirmed CAP compared to 11.7% and 19.8% in controls, by culture and
- 248 qPCR respectively [1]; in comparison we detected colonisation of 5% and 15.8% (>8000
- 249 copies/ml) in patients with LRTI and 15.8% and 21.0% (>8000 copies/ml) in controls. We

ournal of Clinica Microbiology 250 therefore noted high rates of PCR positivity in both groups and low rates of culture positivity 251 in our LRTI patients compared with the CAP patients in this previous study. The differences 252 between the two studies may be due to the fact that our patient cohort was considerably 253 older (64.5 v. 38.4 yrs old) [1], had low rates of radiologically confirmed pneumonia (36.8%), high rates of prior antibiotic treatment, high rates of contact with children and are presumed 254 HIV uninfected (overall HIV incidence is low in Liverpool - 15 per 100000, with a prevalence 255 256 of 0.17% in 2011 [unpublished local data]). Previously in Liverpool we found natural colonisation rates in healthy non-smoking volunteers of 10% by classical culture (25/249, 257 258 age 23 yrs old [SD ±5.7]) [unpublished data]. The higher rate (15.8%) in this cohort may be 259 related to their high rates of contact with children and smoking history. 260 qPCR can deliver results within a few hours (usually 3-6hrs) and could impact the critical 261 phase of early clinical care [25], however it does not distinguish between viable (live) and 262 non-viable (dead) bacteria or determine whether the bacteria is a pathogen or a coloniser [26, 27]. Specificity can also be an issue with *q*PCR and there have been concerns that *lytA* 263 264 may not discriminate between S. pneumoniae and S. viridans, however lytA is currently the most widely used target gene for pneumococcus and we have previously shown that our 265 assay specificity [24] is in line with that reported by others [16]. 266 Within this cohort all LRTI patients had taken antibiotics prior to sampling, which likely 267 accounts for the higher positivity rate of qPCR over culture. Prior antibiotic treatment can 268 269 lower plasma and pleural fluid PCR loads [28] as well as sputum and blood culture positivity. It is not known how rapidly pneumococcus will be undetectable by qPCR in the NW of those 270 271 previously colonised with pneumococcus after antibiotic therapy. Albrich and colleagues suggest that a density of 10³-10⁴ may be the critical density at which 272 colonisation leads to infection [1]; however we have found densities as high or higher in our 273

- 274 cohort of healthy volunteers after experimental colonisation without infection [24, 29].
- 275 Colonisation density was not different in LRTI and controls, we also found high mean

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densities $\geq 10^3$ in those without infection (n = 4 controls). It is possible therefore that if 276 colonisation is dense and in the setting of the correct clinical syndrome then the 277 278 pneumococcus is a likely pathogen. Again an important difference between the two study groups may be HIV infection status. Only 10.5% (2/19) of our LRTI group were Binax 279 positive compared to 72.7% in patients with non-bacteraemic CAP in another study [1]. 280 281 Binax results remain positive for at least 7 days after the initiation of antibiotic treatment [30]; 282 notably our urine samples were taken up to 72hrs after admission but often several days after antibiotics had been started. Previous antibiotic therapy has been noted to decrease 283

> 284 culture and qPCR positivity by up to 50% [1].

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286 Conclusion

We have shown that pneumococcal colonisation (assessed by culture and qPCR) cannot be 287 288 used as a method of diagnosis in pneumococcal blood culture negative hospitalised adults with LRTI in the UK, since such patients have already received community antibiotics and 289 290 the laboratory test is non-discriminatory. Further, the number of adults tested for 'potential 291 LRTI' on screening would be impracticable in terms of staff resource. A community based study recruiting patients prior to antibiotic therapy may however be a useful future step. 292

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294 List of abbreviations:

- 295 Lower respiratory tract infection (LRTI)
- Nasal wash (NW) 296
- Quantitative real time *lytA* Polymerase Chain Reaction (*q*PCR) 297
- Community acquired pneumonia (CAP) 298
- 299 Accident and emergency (A&E)
- Acute medical admissions unit (AMAU) 300

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- 301 Pulmonary embolus (PE)
- 302 Congestive cardiac failure (CCF)
- 303 Adult acute respiratory distress syndrome (ARDS)
- 304

305 Competing interests:

- 306 No authors have any competing interests to declare. The authors have had no support from
- 307 any organisation for the submitted work, no financial relationships with any organisations
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- 309 relationships or activities that could appear to have influenced the submitted work.

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311 Author contributors:

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- A Banyard was involved in sample processing and manuscript editing.
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- 316 and processing and manuscript editing.
- 317 A D Wright was involved in study co-ordination, screening and recruiting participants, sample
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- 319 J F Gritzfeld was involved in protocol writing, sample processing, data collation and
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- 327 S B Gordon was chief investigator and was involved in editing the protocol, ethics
- 328 submission and manuscript preparation.

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- 329 D Shaw was involved in screening and recruiting participants and sample collection.
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443 Figure legends:

- 444 Figure 1: Screening and recruitment flowchart. Reasons for non-recruitment for lower
- 445 respiratory tract infection (LRTI) patients are detailed. Total no. screened n = 826. Note
- 446 multiple reasons for non-recruitment per patient were possible.

447 Tables:

- 448 Table 1: Baseline demographics, antibiotic Status, nasal wash volume returned and
- 449 evidence of pneumococcal disease investigation results of patients with lower respiratory
- 450 tract infection (LRTI) and age and gender matched hospitalised controls.
- 451

	LRTI (n=19)	Control	p value
		(n=19)	
Gender: Male n (%)	9 (47.4)	9 (47.4)	1.000 *
Age Years ± SD	64.47 ±	64.58 ±14.50	0.954 β
	15.78		
Smoker/ ex-smoker n (%)	15 (78.9)	10 (52.6)	0.170 α
23 PPV Pneumovax n (%)	7 (36.8)	8 (42.1)	0.740 *
Contact with children n (%)	10 (52.6)	12 (63.2)	0.511 *
Antibiotics at time of recruitment n (%)	19 (100)	3 (15.8)	0.0001 α
Nasal wash volume returned (ml) \pm SD	10.14 ± 3.14	10.36 ± 4.83	0.855 β
Evidence of pneumococcal disease: Binax urine	2 (10.5)	0 (0)	0.486 α
test positive n (%)			
Evidence of pneumococcal disease: Blood or	0 (0)	N/A	N/A
sputum culture positive n %			

452 *Chi Square, β Mann Whitney U test, α Fisher's Exact, SD standard deviation, PPV

453 polysaccharide vaccine

- 454 Table 2: Pneumococcus identification (by culture, *q*PCR) and density (by *q*PCR) in patients
- 455 with lower respiratory tract infection (LRTI) and age and gender matched hospitalised
- 456 controls.
- 457 Note low rates of culture positivity and high rates of *q*PCR positivity in both LRTI and control
- 458 groups.
- 459

	LRTI (n=19)	Control (n=19)	p value
Culture positive n (%)	1 (5)	3 (15.8)	0.604 α
qPCR positive n (%) at detection	10 (52.6)	8 (42.1)	0.516 *
limit			
Density (by qPCR) copies/ml	3066 [1225 – 7675]	2208 [244 – 19972]	0.408 β
(geometric mean) [95% CI]			
Clinically relevant density (by	3	4	0.999 α
<i>q</i> PCR) >8000 copies/ml			

460 α Fisher's Exact, *Chi squared, β Mann Whitney U test, *q*PCR quantitative polymerase chain

461 reaction

JCM

