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1 **Title:** Living conditions are associated with increased antibiotic resistance in community isolates of  
2 *Escherichia coli*.

3 **Running title:** Living conditions are associated with increased antibiotic resistance of *E. coli*.

4 **Article type:** Brief report

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11 **Keywords:** antibiotic resistance, deprivation, epidemiology, modelling.

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23 **Synopsis**

24 **Objectives:** To measure the associations between domains of deprivation and antibiotic resistance  
25 of *Escherichia coli*.

26 **Methods:** Routine surveillance data for antibiotic susceptibility of *E. coli* isolates were obtained from  
27 urine specimens taken from patients presenting to health care practitioners based in the community  
28 in Leeds and Bradford with suspected urinary tract infection in 2010–2012. Eight antibiotics were  
29 included in the analyses. Postcodes were linked to lower super output areas (average populations of  
30 1500). The 2010 Indices of Deprivation were used as neighbourhood characteristics for each lower  
31 super output area. Multilevel logistic regression models were used to estimate the independent  
32 effect of structural components on the odds of resistance to each antibiotic.

33 **Results:** Residence in the most deprived areas compared to the least deprived areas with respect to  
34 living conditions was associated with increased odds of antibiotic resistance for all eight antibiotics  
35 analysed. The magnitude of these associations included: an odds ratio of 2.04 (95% confidence  
36 interval [CI]: 1.03- 3.07) for cefalexin, 2.16 (95% CI: 1.16–4.05) for ciprofloxacin, 2.47 (95% CI: 1.08-  
37 5.66) for nitrofurantoin, and 1.33 (95% CI: 1.07–1.75) for trimethoprim.

38 **Conclusions:** Social deprivation in the form of living conditions is associated with increased antibiotic  
39 resistance for *E. coli*. This evidence suggests there is a need for further individual-level studies to  
40 explore the potential mechanism for these associations.

41

## 42 **Introduction**

43 Antibiotic resistance is associated with reduced efficacy of antibiotic treatment, which translates  
44 into an estimated 25,000 extra deaths annually within Europe.<sup>1</sup> The distribution of antibiotic  
45 resistance between countries is not equal.<sup>2</sup> The reasons for this are not causally proven, but inter-  
46 country variation in resistance rates have been associated with income inequality and antibiotic  
47 consumption.<sup>2</sup> In addition, policies at the national level relating to the control of antibiotic  
48 consumption, and in Europe, the density of general practitioners and their remuneration method,  
49 have been suggested to influence rates of antibiotic resistance.<sup>3,4</sup> National-level analyses, however,  
50 do not explain the intra-country distribution of antibiotic resistance, nor the mechanisms that may  
51 give rise to antibiotic resistance.

52 Studies investigating intra-country distribution of resistance have used antibiotic consumption as a  
53 surrogate marker for antibiotic resistance. A study of Swedish children identified that parental  
54 educational level, parents being born outside Sweden, environmental exposure to smoking,  
55 economic stress, and parent reported access to personal support were associated with increased  
56 antibiotic consumption.<sup>5</sup> In England and Scotland, antibiotic prescribing by general practitioners has  
57 been related to social deprivation.<sup>6,7</sup> In the USA, antibiotic consumption has been shown to be  
58 associated with population rates of obese persons, children  $\leq 2$  years of age, prescribers per capita,  
59 and females.<sup>8</sup> In general, increasing social deprivation, which includes measures relating to income;  
60 education, skills and training; living environment; barriers to housing and services and health and  
61 disability, is associated with increased antibiotic consumption, but there is also evidence that  
62 increasing antibiotic consumption may be related to increased per capita income.<sup>9,10</sup>

63 Whilst there is evidence to link increasing social deprivation to increasing antibiotic consumption,  
64 associations between social deprivation and antibiotic resistance have not been determined. This  
65 retrospective secondary data analysis used the antimicrobial susceptibilities of *Escherichia coli*

66 isolates derived from urine samples and neighbourhood characteristics to determine whether  
67 components of social deprivation were associated with increased rates of antibiotic resistance.

## 68 **Materials and methods**

### 69 ***Antibiotic resistance data***

70 Antimicrobial resistance surveillance data was collected from 2,778 urinary *E. coli* isolates collected  
71 from patients presenting to health care practitioners based in the community (Leeds and Bradford  
72 (UK)) during 2010–2012.<sup>11</sup> Isolates were therefore derived from patients resident in their own  
73 homes and community based health care institutions e.g. residential care facilities, but excludes  
74 hospitalised patients. Data associated with isolates did not include identifiable patient level data,  
75 we cannot therefore confirm all isolates are patient-unique. Resistance to eight antibiotics was  
76 considered: ampicillin, amoxicillin-clavulanic acid, cefalexin, ceftazidime, cefuroxime, ciprofloxacin,  
77 nitrofurantoin, trimethoprim. This use of pseudo-anonymised data removed the need for ethical  
78 consent. Approval for the use of the data was obtained from the Information Governance team at  
79 Leeds Teaching Hospitals NHS Trust.

### 80 ***Neighbourhood-level data***

81 Patients postcodes were linked to neighbourhood deprivation scores via lower super output areas  
82 (LSOA; census derived small areas (neighbourhoods) with an average population of 1,500 persons)  
83 using GeoConvert ([www.geoconvert.mimas.ac.uk](http://www.geoconvert.mimas.ac.uk)). So each postcode corresponds to a  
84 neighbourhood level LSOA, and each neighbourhood level LSOA has a neighbourhood level  
85 deprivation score. The deprivation scores for each LSOA were obtained from the 2010 Indices of  
86 Deprivation.<sup>12</sup> This dataset assigns a deprivation scores to each LSOA, reporting on five deprivation  
87 domains which were considered independently. The five domains are: income; education, skills and  
88 training; living environment; barriers to housing and services; health and disability (See appendix A  
89 for further detail on the components of the domains). Quintiles were created for each of the five  
90 deprivation domains (within the dataset) and used for analysis. LSOA were grouped according to

91 rural or urban setting based on Rural/Urban Area Classification (RUAC).<sup>13</sup> Population density (persons  
92 per hectare) based on the 2010 census was obtained from the Office for National Statistics  
93 (<https://geoportal.statistics.gov.uk>).

#### 94 ***Statistical analysis***

95 Multilevel logistic regression analysis was used to estimate the independent effect of structural  
96 components on the odds of a patient having a resistant bacterial infection. We considered structural  
97 components to be neighbourhood characteristics that exist above the level of individuals. A separate  
98 model was developed for each of the eight antibiotics using a defined model building strategy. We  
99 used a four stage model building strategy, at each stage testing for the statistical appropriateness of  
100 the more complex model: Model 1: single level model (individual); Model 2: inclusion of LSOA as a  
101 second level and consideration of significant variance at the second level; Model 3: fixed effects at  
102 individual-level (age, sex, year, season) and consideration of the predictive power of each effect;  
103 Model 4: fixed effects at LSOA-level (RUAC, population density, five deprivation domains) followed  
104 by backwards removal of predictors with an associated P-value >0.05 (one-by-one, starting with the  
105 least significant and with re-assessment of significance following removal of each variable). All  
106 changes in model fit were assessed through likelihood ratio tests. Overall fit of the final model was  
107 assessed using the Hosmer-Lemeshow goodness of fit summary statistic.<sup>14</sup> All statistical analysis was  
108 undertaken with Stata v12.1 (StataCorp 2009).

#### 109 **Results**

110 Single variable associations with resistance for each antibiotic are shown in Supplementary Table 1.  
111 All eight final multilevel models represented a significantly good overall fit (P<0.05) with the  
112 inclusion of LSOA as a second level supported by significant variance at that level (all P<0.05).  
113 Increasing age was a significant predictor of increased resistance for all eight antibiotics; male sex  
114 was associated with significantly increased odds for ampicillin and nitrofurantoin resistance  
115 (Supplementary Table 2). Increasing population density was associated with increased odds of

116 resistance to amoxicillin-clavulanic acid, cefalexin, cefuroxime, nitrofurantoin, and trimethoprim but  
117 the rural/urban nature of LSOA was not associated with resistance for any antibiotic (Supplementary  
118 Table 2).

119 Only two deprivation domains remained significant predictors of antibiotic resistance in the final  
120 multivariable models: living conditions (all eight antibiotics) and education, skills and training  
121 (cefuroxime, nitrofurantoin) (Figure 1; Supplementary Table 3). One or two quintiles of living  
122 conditions were associated with significantly increased odds of resistance for each antibiotic; in each  
123 case, the quintile with the worst living conditions had the highest odds ratio compared to the  
124 quintile with the best living conditions (Figure 2). The magnitude of these associations varied from  
125 an odds ratio of 3.03 (1.27–7.17) for ceftazidime to 1.33 (1.07–1.75) for trimethoprim. Where overall  
126 rates of resistance were relatively low, the effects of living conditions on odds of resistance were  
127 greater (Figure 2).

## 128 **Discussion**

129 This study has demonstrated that relative neighbourhood deprivation is associated with antibiotic  
130 resistance. In particular, we have identified that being in the most deprived quintile of the  
131 population with respect to living environment, across all eight antibiotics studied, was associated  
132 with an increased risk of antibiotic resistance compared to living in the least deprived quintile.  
133 Whilst previous studies have related deprivation to antibiotic consumption as a proxy for antibiotic  
134 resistance, this study makes the association between neighbourhood deprivation and antibiotic  
135 resistance without relying on consumption as a proxy for resistance.

136 Deprivation indices can be derived using a number of domains, such as income and the living  
137 environment. This study suggests that living conditions are consistently associated with antibiotic  
138 resistance, after adjusting for other significant predictors within a multivariable analysis. The living  
139 environment deprivation index is composed of indoor living conditions (e.g. thermal comfort) and  
140 outdoor conditions (e.g. air quality). These factors are known to impact negatively on health and our

141 findings may result indirectly from the adverse health impacts derived from a poor living  
142 environment.<sup>15,16</sup> For example, adverse health status may increase antibiotic use or healthcare  
143 utilisation, and so increase the selection and transmission of antibiotic resistant bacteria.

144 The absence of associations with income and health and disability were unexpected, but may be due  
145 to methodological limitations. We have been unable to adjust for individual-level income and health  
146 outcomes which may, in addition to the structural effects of the living environment, be associated  
147 with the risk of antibiotic resistance. We found increasing age to be associated with resistance, in  
148 keeping with previous reports and likely due to increased antibiotic consumption by older age  
149 groups.<sup>17</sup> Increased antibiotic resistance was significantly associated with increasing population  
150 density for five of the eight antibiotics studied. This association has been reported previously.<sup>18</sup> A  
151 possible explanation is that as population density increases, so does the transmission of antibiotic  
152 resistant strains

153 The effect of living environment on antibiotic resistance appears to be increased when the  
154 prevalence of resistance for a specific antibiotic is lower. The mechanism behind this is unclear, but  
155 this effect has also been seen at a national-level in relation to income.<sup>2</sup> A possible explanation is that  
156 sustained antibiotic consumption in a population over a period of time increases an individual's  
157 probability of contact with an already resistant organism, such that the potential for deprivation to  
158 contribute to the selection of antibiotic resistance within an individual is reduced.

159 This study supports existing evidence for the association between deprivation and increased rates of  
160 antibiotic resistance, suggesting that living conditions are strong predictors of increased antibiotic  
161 resistance rates. Further studies are required to explore this relationship by incorporating individual-  
162 level data with neighbourhood characteristics.

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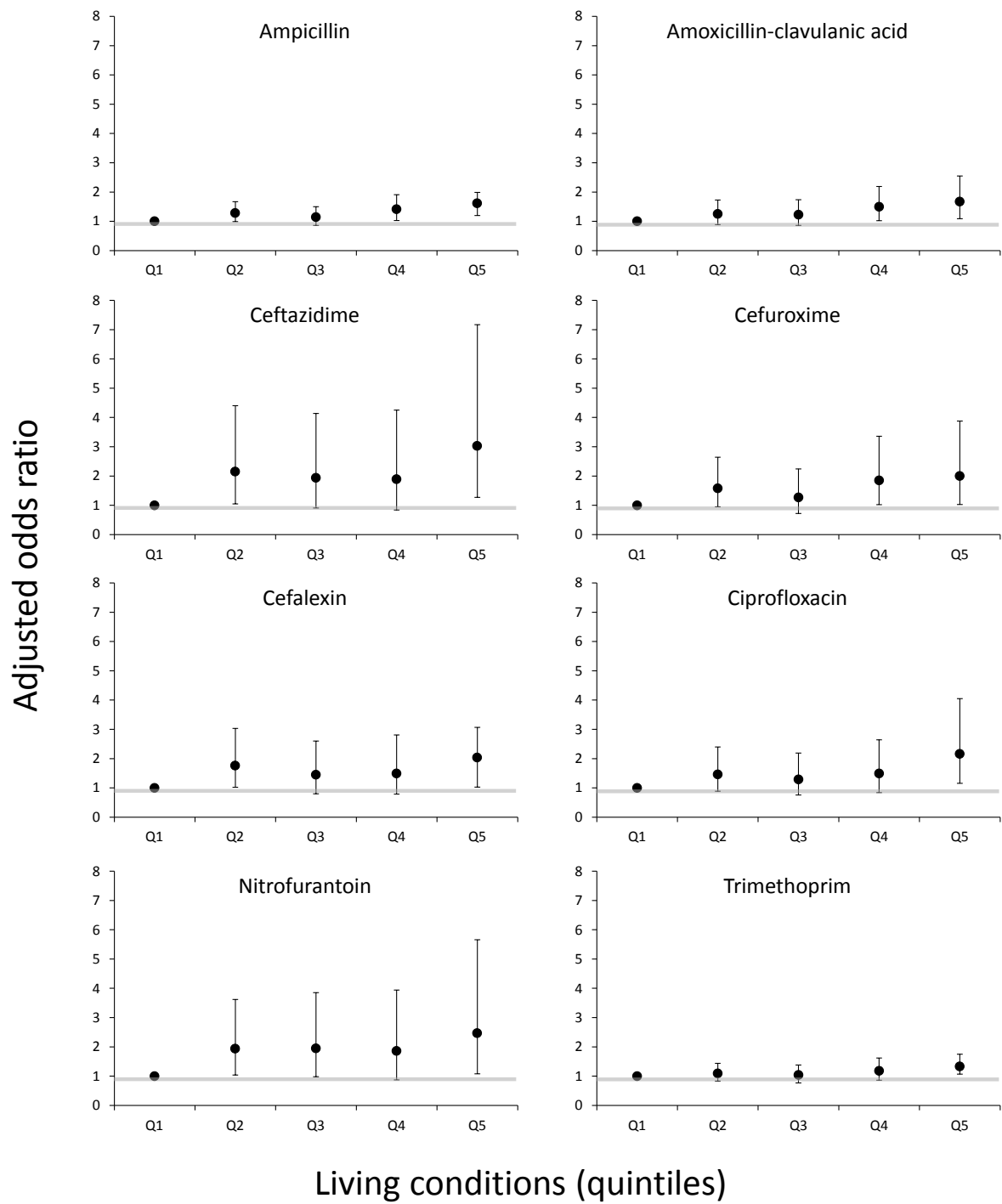
165 **Transparency declarations:** Nil to declare.



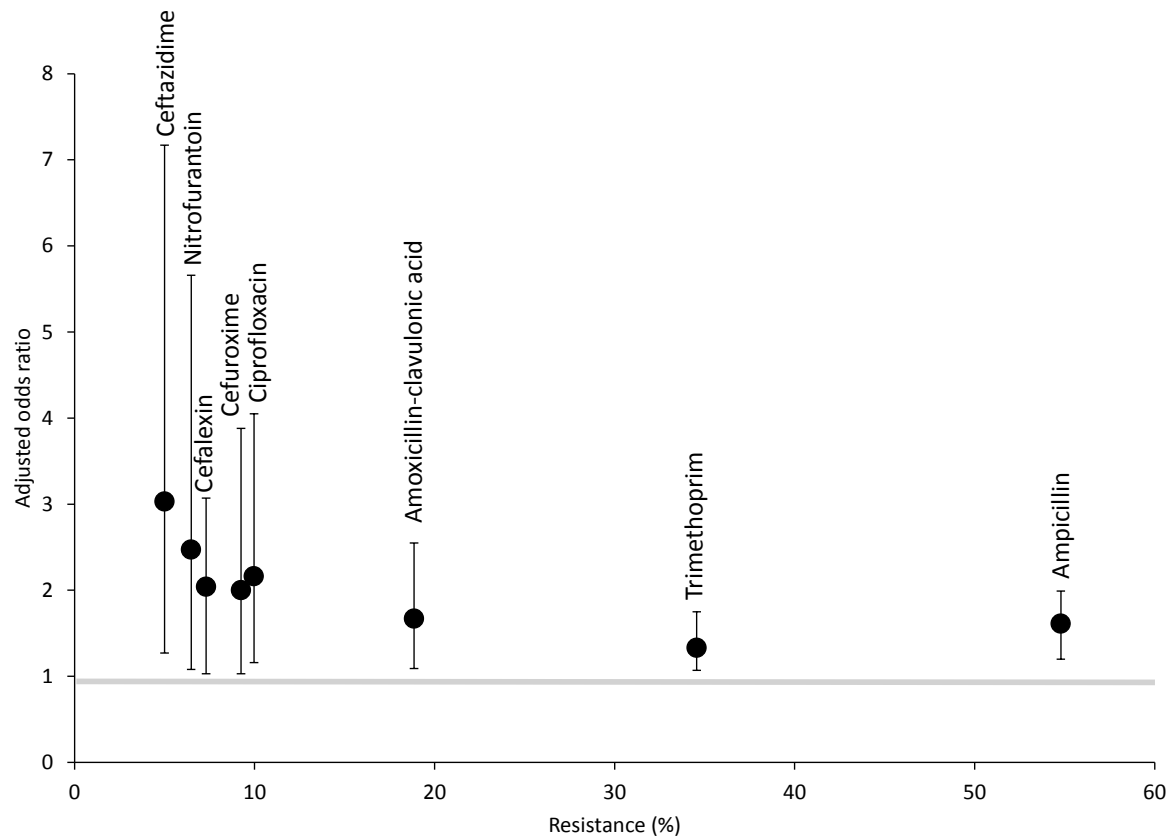
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217 **Figure 1.** Associations between resistance of *E. coli* and neighbourhood living conditions for eight antibiotics.  
 218 Quintile 1 (Q1) represents the least deprived living conditions and is used as the reference group, quintile 5  
 219 (Q5) the most deprived living conditions. Error bars show 95% confidence intervals of the adjusted odds ratio.  
 220 Associations have been adjusted for age, sex, population density, rural/urban nature and education  
 221 deprivation (cefuroxime, nitrofurantoin only).  
 222



**Figure 2.** Correlation between levels of antibiotic resistance and magnitude of the association between living conditions and individual-level odds of resistance. Each datapoint represents a single antibiotic ranked according to percentages of resistant isolates within the dataset: ceftazidime (5.0%), nitrofurantoin (6.5%), cefalexin (7.3%), cefuroxime (9.3%), ciprofloxacin (10.0%), amoxicillin-clavulanic acid (18.9%), trimethoprim (34.6%) and ampicillin (54.8%). The odds ratio is the adjusted association between antibiotic resistance and living in the quintile of Leeds with the most deprived living conditions vs. the least deprived living conditions. Error bars show 95% confidence intervals of the adjusted odds ratio. Associations have been adjusted for age, sex, population density, rural/urban nature and education deprivation (cefuroxime, nitrofurantoin only).