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Abstract (poster session)

Virulence and resistance characteristics of *Klebsiella pneumoniae* causing community-acquired urinary tract infection

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Objectives: *Klebsiella pneumoniae* is the second most common species causing urinary tract infections (UTI). The aim of this study was to evaluate the virulence and resistance determinants of *K. pneumoniae* isolates and host factors potentially relevant to community-acquired urinary tract infection. **Methods:** During 2010, a total of 50 *Klebsiella pneumoniae* isolates causing community-acquired UTI were collected from 10 community-centres in Portugal. Isolates were recovered from patients with less than 50 years (22%, 11/50), more than 50 years (78%, 39/50), complicated cystitis (18%, 9/50) and recurrent UTI infections (38%, 19/50). Susceptibilities to antimicrobial agents were determined by disk diffusion and interpreted according to CLSI guidelines: amoxicillin/clavulanic acid, cefoxitine, ceftazidime, cefotaxime, imipenem, gentamicin, fosfomycin, ciprofloxacin and levofloxacin. The isolates were screened by PCR amplification with specific primers for bla-CTX-M, bla-TEM and bla-SHV extended-spectrum- β -lactamases (ESBL) and 6 virulence factors genes: k2A (K2 serotype), fimH (fimbrial adhesin type 1), mrkD2 and mrkD3 (fimbrial adhesin type 3), khe (haemolysin) and iucC (aerobactine). A p value of ≤ 0.05 was used to indicate statistical significance. **Results:** 10% (5/50) of the *K. pneumoniae* isolates showed the bla-TEM-type β -lactamase (3/50) and the bla-CTX-M-1 ESBL (2/50), only associated with adults ≥ 50 years old. Worrying quinolone resistance was found to ciprofloxacin (16%) and levofloxacin (16%). The most frequent virulence genes were the mrkD3 (62%), fimH (40%) and khe (46%). 6% of the isolates of *Klebsiella pneumoniae* belong to capsular serotype K2. **Conclusions:** Fimbrial adhesins type 3 (variety mrkD2) and the serotype K2 was correlated with elderly women UTI while haemolysin and fimbrial adhesins type 1 and 3 (variety mrkD3) may have a role in community-acquired UTI by older adults. These data provide information on the resistance and virulence patterns among *Klebsiella pneumoniae* isolates currently causing community-acquired UTI.