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Extended spectrum beta-lactamase in clinical isolates of *Escherichia coli* and *Klebsiella pneumoniae* from the Tamale teaching hospital

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Extended spectrum beta-lactamase (ESBL) producing *Escherichia coli* and *Klebsiella pneumoniae* are pathogens of significant public health interest to which new antibiotics therapies are urgently needed. This study was designed to determine the prevalence of ESBLs in clinical isolates of *E. coli* and *K. pneumoniae* from patients attending the Tamale teaching hospital (TTH). A total of 140 isolates of *E. coli* (83.6%; n=117) and *K. pneumoniae* (16.4%; n=23) were cultured from clinical specimens of consenting patients. Antimicrobial susceptibility was determined using the Kirby-Bauer disc diffusion method. Screening and confirmation for ESBL-producing phenotypes among the clinical isolates were performed according to the guidelines of the Clinical and Laboratory Standard Institute, 2012. *Escherichia coli* and *K. pneumoniae* positive for ESBL phenotype were examined for the presence of TEM, SHV and CTX-M genes. Sixty two (44.3%) of the 140 isolates expressed ESBLs phenotypically. Of these, 83.9% (n=52) were *E. coli* and 16.1% (n=10) were *K. pneumoniae* isolates. The proportion of ESBL-producing isolates were found to be relatively higher in adults (15-65 years) than in neonates (<28 days) [p=0.14]. Majority of the isolates showed high percentage resistance to ampicillin (96%) and tetracycline (89%), but relatively low resistance for amikacin (36%). None of the isolates were resistant to meropenem. The ESBL producers were multidrug resistant compared to non-ESBL-producers (23%, n=14/62 versus 18%, n=14/78; p=0.573). Overall, 74.2% (n=46/62) of the ESBL genotypes expressed BlaCTX-M-1 genes followed by 62.9% (n=39/62) BlaTEM and 16.1% (n=10/62) BlaSHV. Two (3.2%) isolates had both TEM and SHV genes, 29 (46.8%) harbored TEM and CTX-M-1, 2 (3.2%) had SHV and CTXM-1, while 4 (6.5%) harbored all three genes. None expressed genes for CTX-M 2 and CTX-M 9. In univariate comparisons, patients who reported their previous medication as having being prescribed by a Physician and those who reportedly completed their previous medication were more likely to be infected by ESBL organisms. The study showed high ESBL positive *E. coli* and *K. pneumoniae*, mostly CTX-M-1 producers in Tamale teaching hospital. Routine laboratory ESBL detection is warranted.

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