

## **Risk factors for antimicrobial resistance among fecal *Escherichia coli* from residents on forty-three swine farms**

**Akwar TH, Poppe C, Wilson J, Reid-Smith RJ, Dyck M, Waddington J, Shang D, Dassie N, McEwen SA**

***Microbial Drug Resistance*, 2007 Spring;13(1):69-76**

Fecal *Escherichia coli* (n = 555) were isolated from 115 residents on 43 farrow-to-finish swine farms to determine the prevalence of antimicrobial resistance and associated risk factors. Susceptibility to 21 antimicrobials was determined and the overall prevalence of antimicrobial resistance was 25.8%. Pair-wise difference in prevalences of resistance to individual antimicrobials was significant between isolates from residents on farms that fed medicated swine rations compared to those that did not (p = 0.013). Cross-resistance among antimicrobials of same class and multidrug-resistance were observed. Logistic regression models revealed the following risk factors positively associated with antimicrobial resistance: use of antimicrobials in pigs on farms; number of hours per week that farmers spent in their pig barns; handling of sick pigs; and intake of antimicrobials by farm residents. This study indicates that occupational exposure of farmers to resistant bacteria and use of antimicrobials in pig farming may constitute a source of resistance in humans, although the human health impacts of such resistance is unknown. The consumption of antimicrobials by farmers appeared to constitute a significant risk for resistance development. Fecal *E. coli* from farm residents may act as a reservoir of resistance genes for animal and/or human pathogens.

[Source: PubMed,

[http://www.ncbi.nlm.nih.gov/sites/entrez?db=pubmed&cmd=Retrieve&dopt=AbstractPlus&list\\_uids=17536936&itool=pubmed\\_DocSum](http://www.ncbi.nlm.nih.gov/sites/entrez?db=pubmed&cmd=Retrieve&dopt=AbstractPlus&list_uids=17536936&itool=pubmed_DocSum)]