

P016 Activities of antimicrobial peptides against *E. coli* O157:H7 – help or hindrance?

C.G. Currie¹, N.F. Inglis², D.A. Devine³, D. O'Neil⁴
and D.G.E. Smith^{1,5}

1: Microbial-Cellular Interactions group, Moredun Research Institute, Edinburgh; 2: Moredun Proteomics Facility; 3: Novabiotics Ltd., Aberdeen Science & Technology Park, Aberdeen; 4: Leeds Dental Institute, University of Leeds; 5: Institute for Comparative Medicine, University of Glasgow

Although enterohaemorrhagic *E. coli* (EHEC), particularly *E. coli* O157:H7, are an important group of zoonotic pathogens, few investigations have examined interplay between these bacteria and mammalian antimicrobial peptides of either reservoir (cattle) or human host origin. Expression of several antimicrobial peptides (including defensins and cathelicidins) by epithelial cells challenged with *E. coli* O157:H7 has been determined by RT-PCR. The sensitivity of *E. coli* O157:H7, commensal *E. coli* strains and a reference strain (BUE55) to several natural and synthetic peptides has been assessed. Finally, since verotoxin (VT, key pathogenicity determinants during infection of both human and bovine hosts) expression can be induced by various stressors (including several antibiotics), the influence of peptides on VT release from EHEC is being examined. This understanding could contribute towards development of novel approaches to limit disease caused by *E. coli* O157:H7.