

Seven publications resulting from RCVS Charitable Trust funded project

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1. Molecular characterization of the uncultivable hemotropic bacterium *Mycoplasma haemofelis*.

Barker, E. N.; Darby, A. C.; Helps, C. R.; Peters, I. R.; Heesom, K. J.; Arthur, C. J.; Crossett, B.; Hughes, M. A.; Radford, A. D.; Tasker, S.; BioMed Central Ltd, London, UK, Veterinary Research, 2011, 42, 83, (12 July 2011), 31 ref.

2. Genome sequence for "Candidatus *Mycoplasma haemominutum*," a low-pathogenicity hemoplasma species.

Barker, E. N.; Darby, A. C.; Helps, C. R.; Peters, I. R.; Hughes, M. A.; Radford, A. D.; Novacco, M.; Boretti, F. S.; Hofmann-Lehmann, R.; Tasker, S.; American Society for Microbiology (ASM), Washington, USA, Journal of Bacteriology, 2012, 194, 4, 905-906, 17 ref.

3. Complete genome sequence of *Mycoplasma haemofelis*, a hemotropic mycoplasma.

Barker, E. N.; Helps, C. R.; Peters, I. R.; Darby, A. C.; Radford, A. D.; Tasker, S.; American Society for Microbiology (ASM), Washington, USA, Journal of Bacteriology, 2011, 193, 8, 2060-2061, 15 ref.

4. Genomic analysis of highly virulent Georgia 2007/1 isolate of African swine fever virus.

Chapman, D. A. G.; Darby, A. C.; Silva, M. da; Upton, C.; Radford, A. D.; Dixon, L. K.; National Center for Infectious Diseases, Centers for Disease Control and Prevention, Atlanta, USA, Emerging Infectious Diseases, 2011, 17, 4, 599-605, 29 ref.

5. Identification of three novel superantigen-encoding genes in *Streptococcus equi* subsp. *zooepidemicus*, *szef*, *szen*, and *szep*.

Paillot, R.; Darby, A. C.; Robinson, C.; Wright, N. L.; Steward, K. F.; Anderson, E.; Webb, K.; Holden, M. T. G.; Efstratiou, A.; Broughton, K.; Jolley, K. A.; Priestnall, S. L.; Campi, M. C. M.; Hughes, M. A.; Radford, A.; Erles, K.; Waller, A. S.; American Society for Microbiology (ASM), Washington, USA, Infection and Immunity, 2010, 78, 11, 4817-4827, 51 ref.

6. Application of next-generation sequencing technologies in virology.

Radford AD, Chapman D, Dixon L, Chantrey J, Darby AC, and Hall N. (2012). Application of next-generation sequencing technologies in virology. Available: <http://vir.sgmjournals.org/content/early/2012/05/24/vir.0.043182-0.full.pdf>. Last accessed 1st August 2012. Available online for free.

7. Analysis of gene expression from the *Wolbachia* genome of a filarial nematode supports both metabolic and defensive roles within the symbiosis.

Armstrong, SD.; Bah, GS.; Blaxter, ML.; Darby, AC.; Cordaux, R.; Hughes, MA.; Kaur, G.; Kay, SM.; Koldkjær, P.; Makepeace, BL.; Radford, AD.; Tanya, VN.; Trees, AJ.; Wastling, JM. Analysis of gene expression from the *Wolbachia* genome of a filarial nematode supports both metabolic and defensive roles within the symbiosis. Genome Research. Available: <http://genome.cshlp.org/content/early/2012/08/16/gr.138420.112.full.pdf+html> Last accessed 30th August 2012. Available online for free.