

Edinburgh Infectious Diseases publications in 2012

1. Abuzaid, A., Hamouda, A., and Amyes, S.G. (2012). Klebsiella pneumoniae susceptibility to biocides and its association with cepA, qacDeltaE and qacE efflux pump genes and antibiotic resistance. *J Hosp Infect* 81, 87-91.
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3. Akman, O.E., Watterson, S., Parton, A., Binns, N., Millar, A.J., and Ghazal, P. (2012). Digital clocks: simple Boolean models can quantitatively describe circadian systems. *J R Soc Interface* 9, 2365-2382.
4. Al-Hassan, L., Mahallawy, H.E., and Amyes, S.G. (2012). First report of bla(PER-3) in *Acinetobacter baumannii*. *Int J Antimicrob Agents*.
5. Alafuzoff, I., Gelpi, E., Al-Sarraj, S., Arzberger, T., Attems, J., Bodí, I., Bogdanovic, N., Budka, H., Bugiani, O., Englund, E., et al. (2012). The need to unify neuropathological assessments of vascular alterations in the ageing brain: multicentre survey by the BrainNet Europe consortium. *Exp Gerontol* 47, 825-833.
6. Alber, A., Howie, S.E., Wallace, W.A., and Hirani, N. (2012). The role of macrophages in healing the wounded lung. *Int J Exp Pathol* 93, 243-251.
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10. Anderson, R.C., Krueger, N.A., Genovese, K.J., Stanton, T.B., Mackinnon, K.M., Harvey, R.B., Edrington, T.S., Callaway, T.R., and Nisbet, A.J. (2012). Effect of thymol or diphenyliodonium chloride on performance, gut fermentation characteristics, and campylobacter colonization in growing swine. *J Food Prot* 75, 758-761.
11. Appleby, L.J., Nausch, N., Bourke, C.D., Rujeni, N., Midzi, N., Mduluza, T., Allen, J.E., and Mutapi, F. (2012). Chitinase 3-Like 1 Protein Levels Are Elevated in *Schistosoma haematobium* Infected Children. *PLoS Negl Trop Dis* 6, e1898.
12. Archer, J., Baillie, G., Watson, S.J., Kellam, P., Rambaut, A., and Robertson, D.L. (2012). Analysis of high-depth sequence data for studying viral diversity: a comparison of next generation sequencing platforms using Segminator II. *BMC Bioinformatics* 13, 47.
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18. Auld, S.K., Edel, K.H., and Little, T.J. (2012). The cellular immune response of *Daphnia magna* under host-parasite genetic variation and variation in initial dose. *Evolution* 66, 3287-3293.
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24. Babayan, S.A., and Schneider, D.S. (2012). Immunity in society: diverse solutions to common problems. *PLoS Biol* **10**, e1001297.
25. Baele, G., Lemey, P., Bedford, T., Rambaut, A., Suchard, M.A., and Alekseyenko, A.V. (2012). Improving the accuracy of demographic and molecular clock model comparison while accommodating phylogenetic uncertainty. *Mol Biol Evol* **29**, 2157-2167.
26. Baillie, G.J., Galiano, M., Agapow, P.M., Myers, R., Chiam, R., Gall, A., Palser, A.L., Watson, S.J., Hedge, J., Underwood, A., et al. (2012). Evolutionary dynamics of local pandemic H1N1/2009 influenza virus lineages revealed by whole-genome analysis. *J Virol* **86**, 11-18.
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31. Barr, T., and Gray, D. (2012). T(FH) memory: More or less T(FH)? *Eur J Immunol* **42**, 1977-1980.
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