



## Corrigendum

Corrigendum to “The feasibility of testing whether *Fasciola hepatica* is associated with increased risk of verocytotoxin producing *Escherichia coli* O157 from an existing study protocol” [Prev. Vet. Med. 119 (2015) 97–104]



Graeme L. Hickey<sup>a,\*</sup>, Peter J. Diggle<sup>a</sup>, Tom N. McNeilly<sup>b</sup>, Sue C. Tongue<sup>c,1</sup>, Margo E. Chase-Topping<sup>d</sup>, Diana J.L. Williams<sup>e,f</sup>

<sup>a</sup> Department of Epidemiology and Population Health, Institute of Infection and Global Health, University of Liverpool, The Farr Institute@HeRC, Waterhouse Building, 1–5 Brownlow Street, Liverpool, L69 3GL, UK

<sup>b</sup> Moredun Research Institute, Pentlands, Science Park, Bush Loan, Penicuik, Midlothian, EH26 0PZ, UK

<sup>c</sup> SRUC (Scotland's Rural College), West Mains Road, Edinburgh, EH9 3JG, UK

<sup>d</sup> Centre for Immunity, Infection and Evolution, University of Edinburgh, Ashworth Laboratories, Charlotte Auerbach Road, Edinburgh, EH9 3FL, UK

<sup>e</sup> Department of Infection Biology, Institute of Infection and Global Health, University of Liverpool, Liverpool, L69 7ZJ, UK

<sup>f</sup> School of Veterinary Science, University of Liverpool, Leahurst Campus, Chester High Road, Neston, CH64 7TE, UK

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The authors regret that there is an error in lines 617–622 of the R script (coinfection.updated.r; Supplemental Data). Namely, the list of parameters ‘pars0’ used in the calculation should have been ‘pars1’ in all instances. This resulted in a small error in the first sensitivity analysis in Section 4.2. Specifically, the third sentence should read:

“Power ranged from 13.5% (for OR = 1.2) to 99.5% (for OR = 3.0)”.

This error also affects Fig. 4; however the revised figure is visually indistinguishable from the original.

There is also a typographical error in the left hand side of Eq. (4). Specifically, it should read:

$$E_{X|r_j=\hat{r}} \left[ P \left[ Y_j > 0 | \alpha_j, \beta, X_{1j}, \dots, X_{n_j} \right] \right], \quad (4)$$

where the expectation is taken conditional on the average individual prevalence for *F. hepatica*,  $\hat{r}$ . The two instances of ‘ $\alpha_j$ ’ in the integrand of the preceding equation (that followed after averaging over the random effects and sample size distributions) should have been written as ‘ $u$ ’. This does not affect the calculation of the random effects parameters. The authors would like to apologise for any inconvenience caused.

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\* Corresponding author at: The Farr Institute@HeRC, University of Liverpool 2nd Floor, Block F Waterhouse Building 1–5 Brownlow Street, Liverpool, L69 3GL, UK.

E-mail address: [graeme.hickey@liverpool.ac.uk](mailto:graeme.hickey@liverpool.ac.uk) (G.L. Hickey).

<sup>1</sup> Current address: SRUC Epidemiology Research Unit, Drummondhill, Stratherrick Road, Inverness, IV2 4JZ, UK.

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