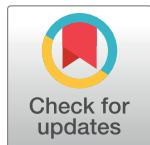


CORRECTION

Correction: Comparative Exposure Assessment of ESBL-Producing *Escherichia coli* through Meat Consumption

The PLOS ONE Staff

[Fig 1](#) appears incorrectly in the published article. Please see the correct [Fig 1](#) and its caption here. The publisher apologizes for the error.



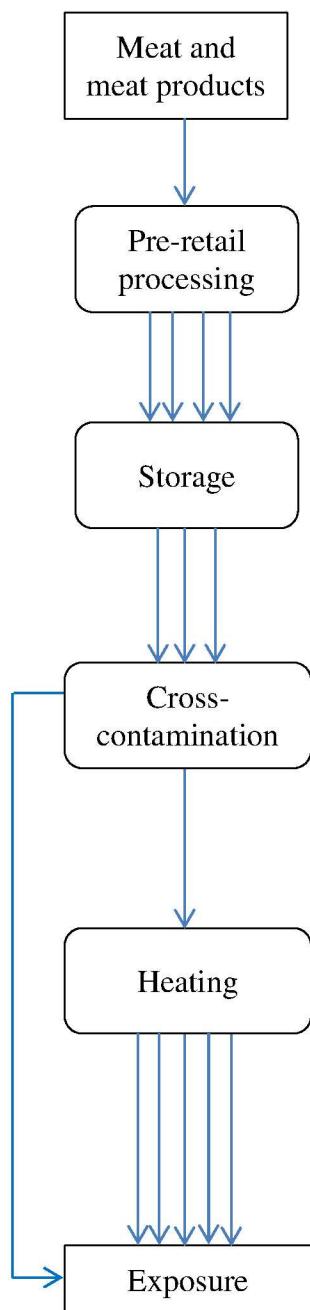
OPEN ACCESS

Citation: The PLOS ONE Staff (2017) Correction: Comparative Exposure Assessment of ESBL-Producing *Escherichia coli* through Meat Consumption. PLoS ONE 12(2): e0173134. doi:10.1371/journal.pone.0173134

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Food chain



Model details

Consumption data (No. portions consumed/year by the Dutch population, portion size)
ESBL *E.coli* (EEC) prevalence and concentration

Model: inactivation
Subdivision: salting / drying&fermenting / heating / none

Model: exponential growth or inactivation
Subdivision: room / fridge / freezer
Parameters: storage times and temperatures, subdivision fractions, EEC growth and inactivation parameters

Model: cross-contamination for a fraction of the portions; for which a fraction of EEC's is transmitted

Model: exponential (D/z) inactivation; linear internal temperature increase
Subdivision: raw / rare / medium / done / divided-done
Parameters: heating times and temperatures, subdivision fractions / EEC inactivation parameters

Fig 1. Overview of the sQMRA model used for the calculations. In- and output (text box with straight corners), processes (rounded corners) and details (dashed lines) are shown.

doi:10.1371/journal.pone.0173134.g001

Reference

1. Evers EG, Pielaat A, Smid JH, van Duijkeren E, Vennemann FBC, Wijnands LM, et al. (2017) Comparative Exposure Assessment of ESBL-Producing *Escherichia coli* through Meat Consumption. PLoS ONE 12(1): e0169589. doi: [10.1371/journal.pone.0169589](https://doi.org/10.1371/journal.pone.0169589) PMID: 28056081