

Implementation of an Early Detection System for Salmonella in livestock production in Great Britain

Christina Papadopoulou¹, Rowena D. Kosmider¹, Robert J. Smeatham¹, Sarah J. Evans¹, Rob H. Davies² ¹Centre for Epidemiology & Risk Analysis, Veterinary Laboratories Agency, Addlestone, Surrey, KT15 3NB, UK ²Department of Food & Environmental Safety, Veterinary Laboratories Agency, Addlestone, Surrey, KT15 3NB, UK

INTRODUCTION

A system for the early detection of outbreaks of Salmonella in livestock in Great Britain was implemented within VLA in September 2006. The system was developed in order to detect changes in disease reporting for a specific time period and in particular for serotypes associated primarily with human infections & public health.

METHODS

Data are derived from incidents of Salmonella originating from clinically diseased domestic livestock species and from reports received from poultry monitored as part of a mandatory control programme for Salmonella under the Poultry Breeding Flocks and Hatcheries Order (PBFHO).

Samples submitted from livestock for investigation of clinical disease:

- **i.** S. Dublin in cattle
- **ii.** S. Typhimurium in cattle
- iii. S. Typhimurium in sheep
- iv. S. Typhimurium in pigs
- S. Typhimurium with the pentavalent resistance pattern (ACSSuT) **V.**
- vi. Salmonella serotypes other than S. Typhimurium with the pentavalent resistance pattern (ACSSuT)
- vii. Salmonella serotypes resistant to nalidixic acid

Samples submitted from domestic fowl under the Poultry Breeding Flocks & **Hatcheries Order:**

- Top 5 serotypes reported i. |
- ii. Salmonella serotypes of designated public health significance in the EU (Enteritidis, Typhimurium, Hadar, Infantis, Virchow)
- iii. All Salmonella serotypes

A Poisson regression model compares the number of observed incidents with an expected value, accounting for seasonality and past outbreaks.

 $\log(\mu_i) = \beta_0 + \beta_1 t_i$

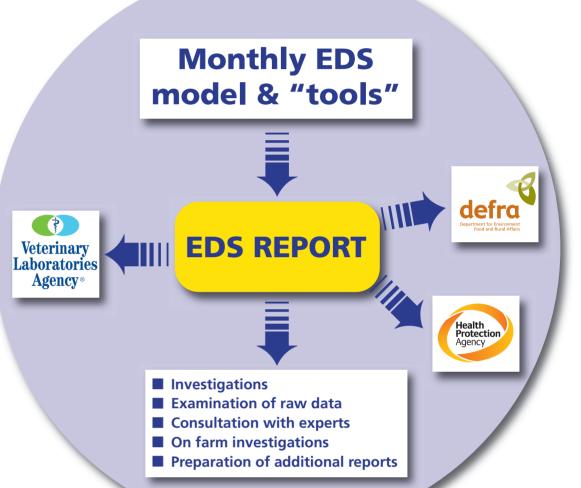
$$\gamma_i \sim Poisson(\mu_i) \quad E(y_i) = \mu_i \quad V(y_i) = \varphi \mu_i$$

A threshold value is calculated for each month. If the current observed count is above the estimated threshold value, a warning is implemented indicating that a potential outbreak is occurring in the field.

This model is complemented by a monthly tool which screens data for certain new or unusual Salmonella serotypes reported from animals.

Salmonella EDS additional "tool":

- strains resistant to cefotaxime or ceftazidime
- ii. strains resistant to ciprofloxacin
- S. Newport strains resistant to A(C)SSuT
- iv. strains resistant to amoxicillin/clavulanic acid



Follow up action:

• Examination of raw data Related to increase of reports linked to investigation of S. Enteritidis in a chicken breeding flock

Follow up action:

Examination of

epidemiological

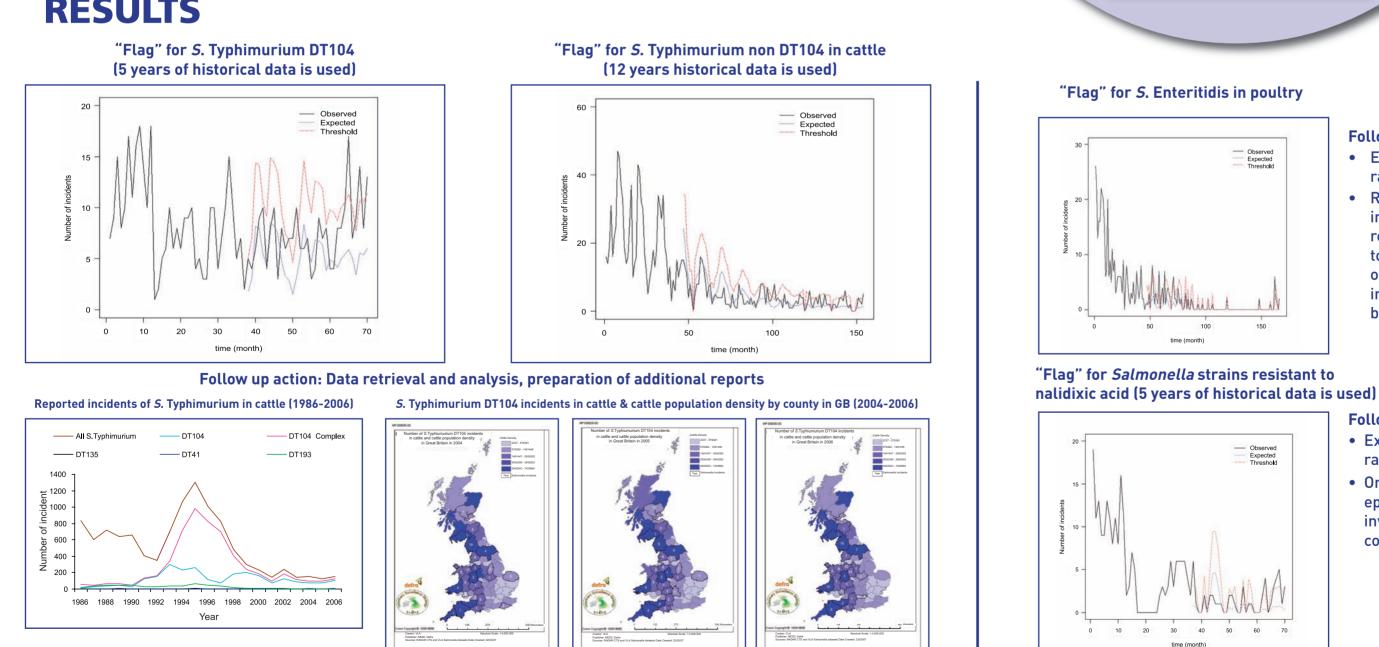
investigations

conducted

raw data

• On farm





DISCUSSION AND CONCLUSIONS

- Seven flags were triggered by the EDS model in September 2006 to September 2007. Four of these were due to true increases in reports, but none due to emerging outbreaks.
- The warning could potentially be a false alarm, probably related to changes in the reporting procedures over time, or to an increase in reports, as a result of extensive investigations of an already known incident (i.e. S. Enteritidis flag in October 2006)
- The EDS does not predict future outbreaks, but alerts as soon as possible to increases that have just happened.

REFERENCES

Kosmider R, Kelly L, Evans S, Gettingby G. (2006) A statistical system for detecting Salmonella outbreaks in British livestock. Epidemiology and Infection 134(5): 952 - 960.

Centre for Epidermiology & Risk Analysis Veterinary Laboratories Agency, New Haw, Addlestone, Surrey KT15 3NB. www.defra.gov.uk/corporate/vla For further information contact c.papadopoulou@vla.defra.gsi.gov.uk