Supporting information - Inferring source attribution from a multi-year multi-source dataset of Salmonella in Minnesota

A) Single attribution, including only sampled sources

B) Single attribution, including non-sampled sources

C) Temporal attribution, including only sampled sources

D) Temporal attribution, including non-sampled sources

Supporting Figure 1. Posterior distribution of relative prevalence of S. Enteritidis in chicken with informative prior density (based on source typing data) shown in red for four different models – A) Single attribution including only sampled sources, B) Single attribution including non-sampled sources, C) Temporal attribution including only sampled sources, and D) Temporal attribution including non-sampled sources.
A) Single attribution, including only sampled sources

B) Single attribution, including non-sampled sources
C) Temporal attribution, including only sampled sources

D) Temporal attribution, including non-sampled sources

Supporting Figure 2 Plots of the observed and expected cases for individual Salmonella serotypes in the A) Single attribution model including only sampled sources, B) Single attribution including non-sampled sources, C) Temporal attribution including only sampled sources, and D) Temporal attribution including non-sampled sources.