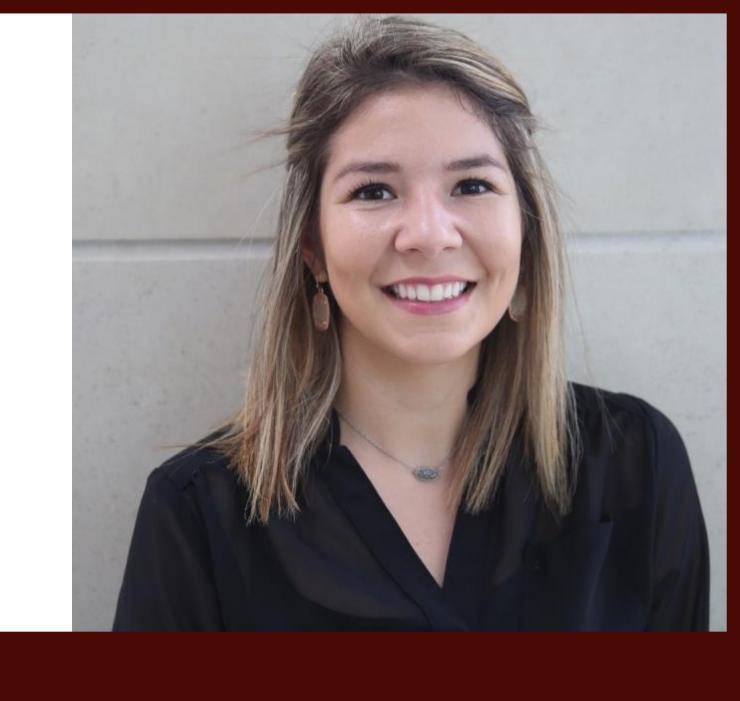


Competitive exclusion of intra-genus Salmonella in neonatal broilers

*1Megan Pineda, 2Michael Kogut, 2Kenneth Genovese, 1Yuhua Z. Farnell, 1Dan Zhao, 1Xi Wang, 1Allison Milby, and 1Morgan Farnell ¹Department of Poultry Science, Texas A&M Agrilife Research

²Southern Plains Agricultural Research Center, Agricultural Research Service, US Department of Agriculture



INTRODUCTION

Salmonellosis is one of the leading foodborne illnesses in the United States. Chickens may harbor Salmonella without having severe symptoms. Salmonella enterica serotypes Typhimurium (ST), Enteritidis (SE) and Kentucky (SK) are in the top 5 common poultry isolates¹. In 2016, ST and SE were isolated from 9.8% and 16.8% of human cases, respectively². However, the number one isolated serotype in poultry is SK, which has a low incidence (0.14%) in human cases².

Current reduction strategies include biosecurity measures, water treatments, feed additives, vaccines and competitive exclusion. Direct colonization resistance occurs due to lack of space and nutrients. Yang and colleagues excluded SE and ST in broiler chicks when a serovar was administered 24 h prior to the other³.

Indirect colonization resistance occurs in response to microbiota stimulated host immunity⁴. Characterization of the immune response can be measured by gene expression of cytokines.

HYPOTHESIS: Salmonella Kentucky will reduce Salmonella Enteritidis and Salmonella Typhimurium by competitive exclusion in broiler chicks and affect the host immune response.

OBJECTIVES: Chicks were concurrently challenged with both Salmonella Kentucky and Typhimurium or Enteritidis. Cecal colonization and incidence was measured by differential plating. Changes in cytokine gene expression was measured in cecal tonsils and liver by quantitative real-time reverse-transcription PCR.

MATERIALS AND METHODS

EXPERIMENTAL DESIGN

Trials 1 and 2							
Treatment	NC	SK	SK→ST	ST	ST → SK	SK+ST	
D0		Place chicks					
D1 (Challenge)	PBS	10 ⁴ CFU SK	10 ⁴ CFU SK	10 ⁴ CFU ST	10 ⁴ CFU ST	10 ⁴ CFU SK+ST	
D2 (Re-Challenge)	PBS	PBS	10 ⁵ CFU ST	PBS	10 ⁵ CFU SK	PBS	
D3 (Kill)	Collect ceca, liver, and spleen tissues						
Trials 3 and 4							
Treatment	NC	SK	SK→SE	SE	SE → SK	SK+SE	
DO	Place chicks						
D1 (Challenge)	PBS	10 ⁴ CFU SK	10 ⁴ CFU SK	10 ⁴ CFU SE	10 ⁴ CFU SE	10 ⁴ CFU SK+SE	
D1 (Challenge) D2 (Re-Challenge)	PBS PBS	10 ⁴ CFU SK PBS	10 ⁴ CFU SK 10 ⁵ CFU SE	10 ⁴ CFU SE PBS	10 ⁴ CFU SE 10 ⁵ CFU SK		

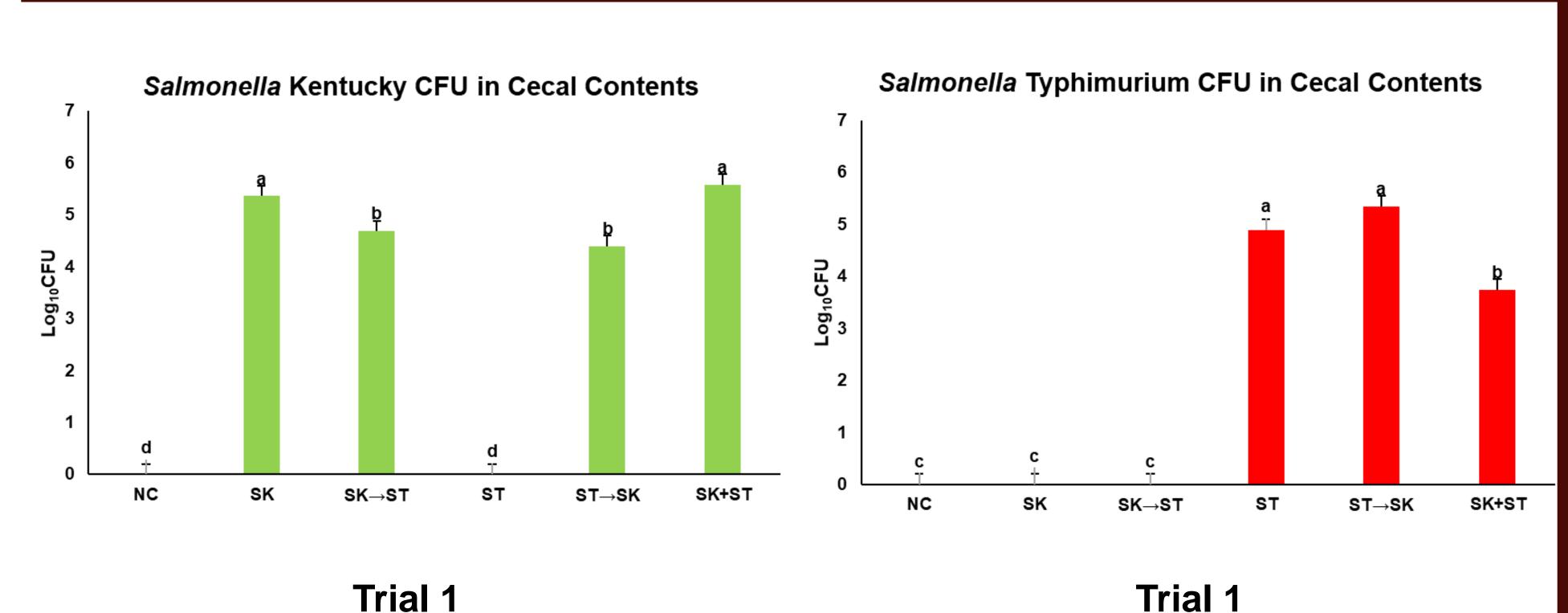
Differential plating was used to select Salmonella isolates for resistance to specific antibiotics. The sample size was n=20 chicks per treatment. Each trial was replicated twice. Gut colonization and incidence were measured from cecal contents. Relative mRNA levels of cytokines interleukin-1β (IL-1β), IL-6, IL-10, IL-10 and gamma interferon (IFN-γ) were measured via qRT-PCR (n=5/treatment).

STATISTICAL ANALYSIS

Statistical analyses were conducted via a Student's t-test for enumeration and gene expression using JMP Pro 15. All the data were presented as mean standard error of the mean. A p-value of < 0.05 was considered significant when compared to the respective positive control. Each trial was replicated twice at different times. Gene expression data were measured from trials 1 and 3.

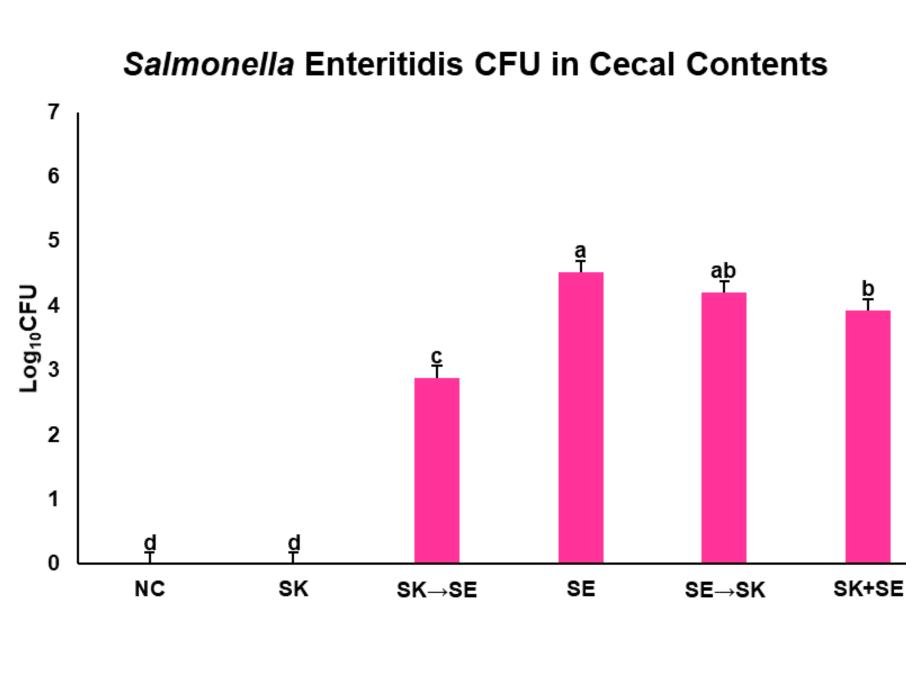
KEY WORDS

RESULTS

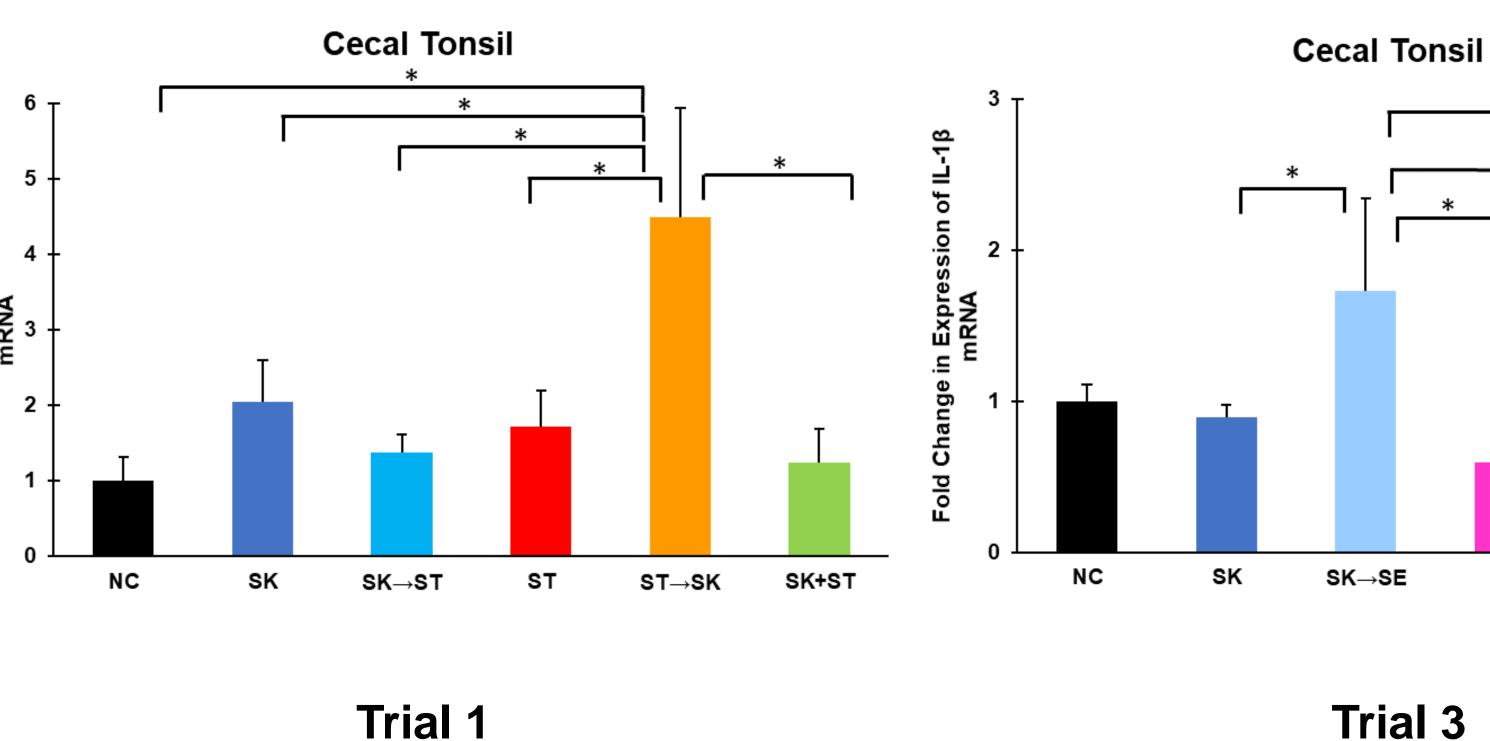


Salmonella Kentucky CFU in Cecal Contents

Trial 3



Trial 3



SE→SK SK→SE Trial 3

CONCLUSIONS

- When administered first Salmonella was able to significantly reduce colonization of the following serotype.
- Salmonella Kentucky significantly reduced SE and ST.
- Significant differences were found in the mRNA levels of cytokines.

REFERENCES

- 1. FSIS. 2014
- 2. CDC. 2016.
- 3. Yang, et al., 2018
- 4. Lawley and Walker, 2012
- 5. Pineda, et al., 2021