

Veterinary Research

Scholars Program

University of Missouri

Fetal microchimerism and its presence in the equine hoof lamellar interface in health and laminitis

<u>Jessica R. Warwick</u>, Angelynn P. Simenson, Julia M. Baldrighi (co-sponsor), Jeffrey N. Bryan (co-sponsor), Philip J. Johnson (co-sponsor) Department of Veterinary Medicine and Surgery, College of Veterinary Medicine, University of Missouri, Columbia, MO

Fetal Microchimeric Cells enter the Maternal Blood Stream

Background

Objectives

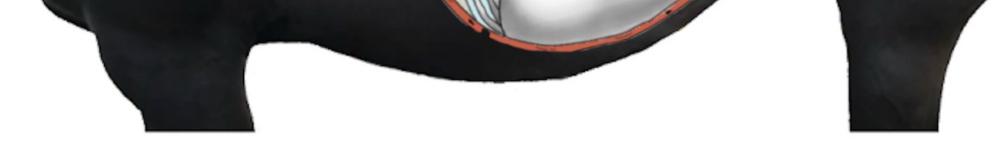
Methods

• Nested polymerase chain reaction (PCR) analysis of equine whole blood to affirm equine fetal microchimerism

Fresh whole blood samples were collected from males (geldings), nulliparous mares, and parous/multiparous mares.

• Fluorescence *in situ* hybridization (FISH) analysis of archived paraffin-embedded formalin fixed equine hoof lamellar tissue

DNA was extracted from blood samples via DNeasy Blood and Tissue kit (Qiagen CA).



Fetal microchimerism is the presence of a small number of genetically distinct cells in an individual that originated from another individual.

Fetal microchimerism has been demonstrated in the human, canine, bovine and ovine species.

Fetal cells with male DNA can be traced using primers designed for the equine SRY gene.

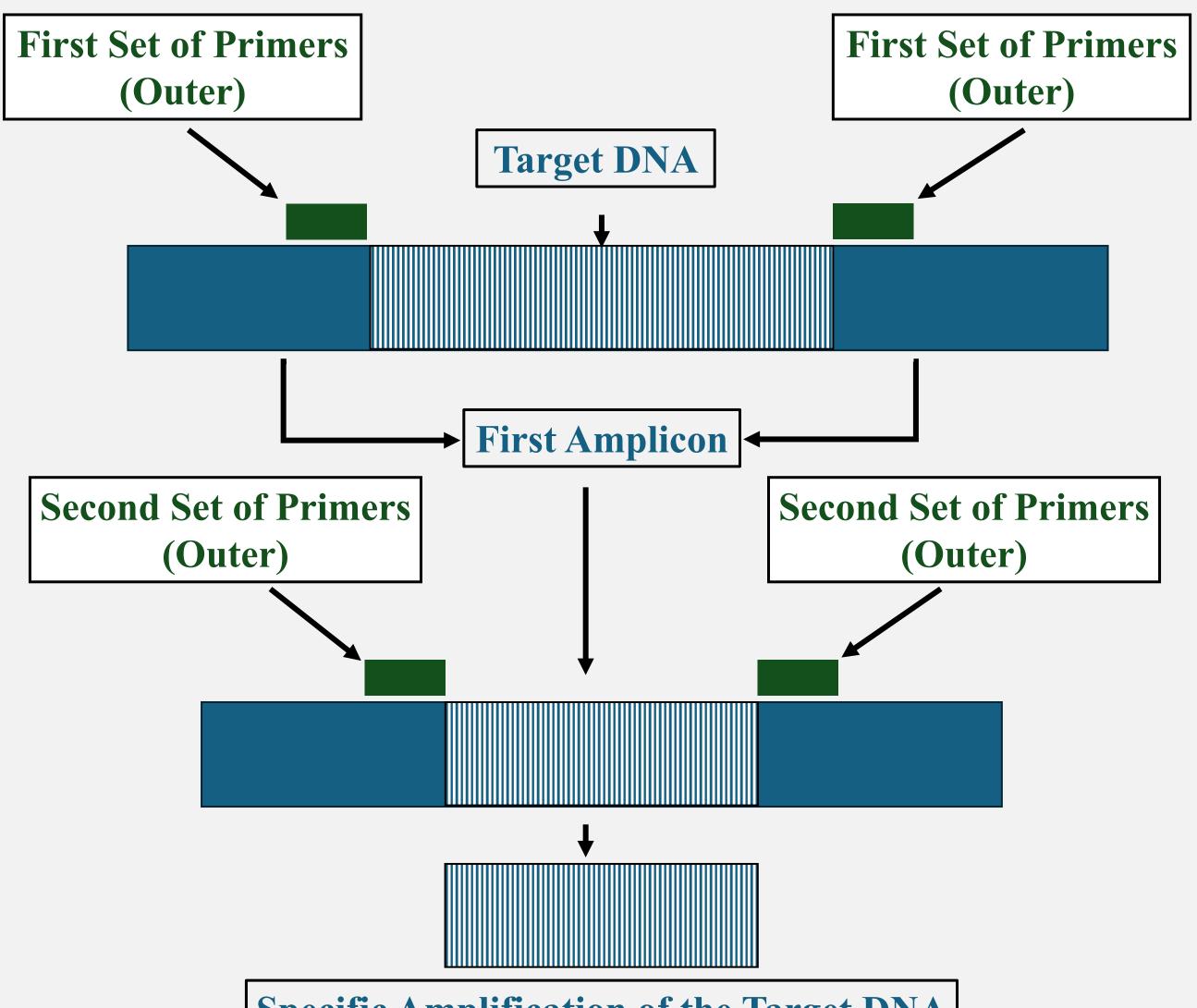
(healthy and laminitic) to identify Ychromosome positive microchimeric cells.

Hypotheses

Nested PCR assays will identify the presence of male fetal DNA circulating in parous/multiparous mares who have given birth to male offspring.

FISH analysis will identify Y chromosomepositive microchimeric cells in the hoof lamellar interface in normal and laminitic conditions.

Nested PCR Analysis



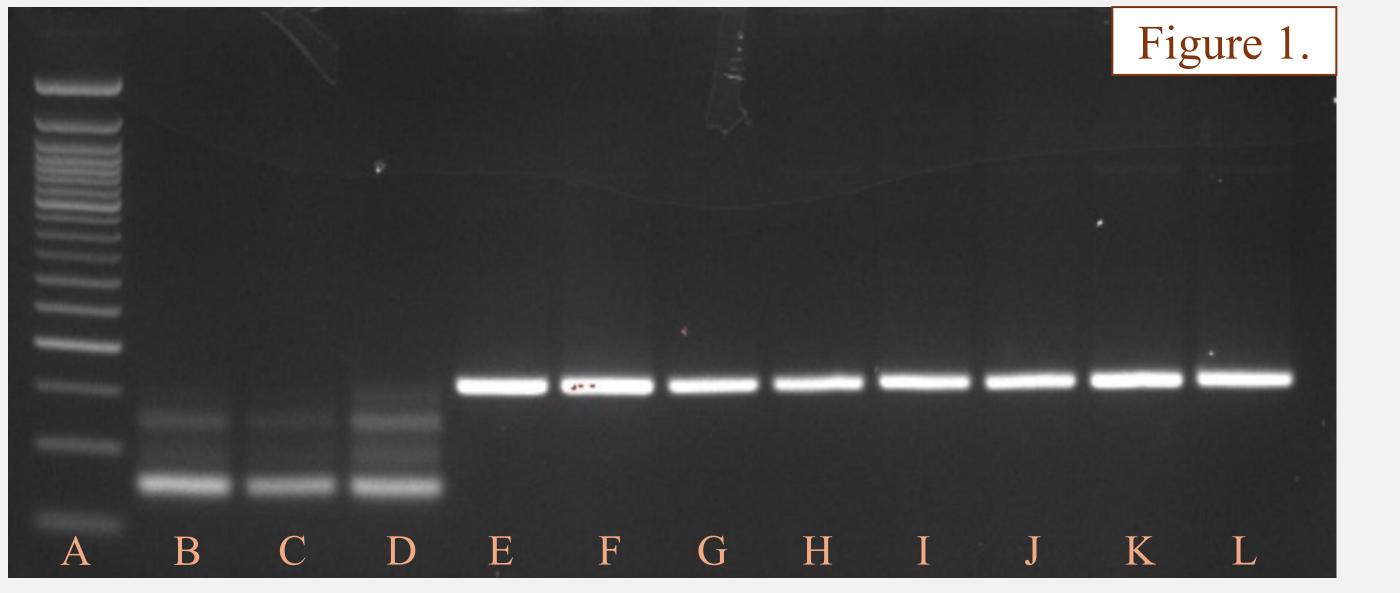


Figure 1. Lane A, 50 bp ladder. Lanes B-D, nested PCR with DNA template concentrations of 200 ng, 250 ng and 300 ng, respectively. Lanes E-L, male/female blood dilution of PCR 1.

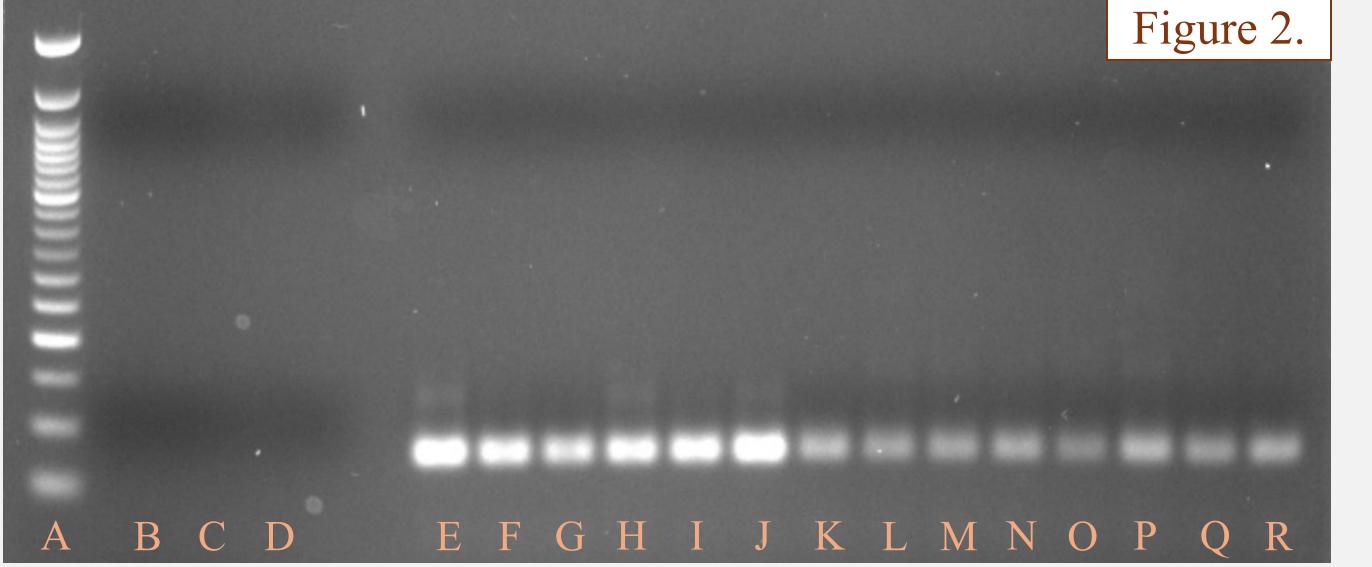
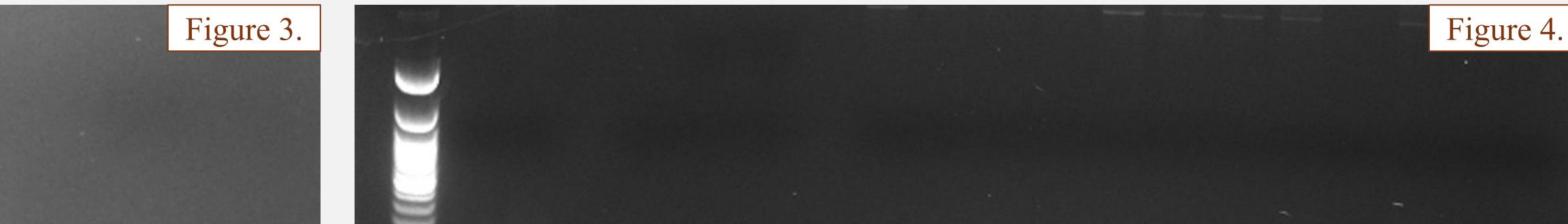


Figure 2. Lane A, 50 bp ladder. Lanes B-D, failed nested PCR using only 1 µl of DNA template. Lanes E-J, cleaned dilution PCR product ran via nested PCR. Lanes K-R, cleaned dilution gel bands ran via nested PCR.



Polymerase Chain Reactions

Specific Amplification of the Target DNA

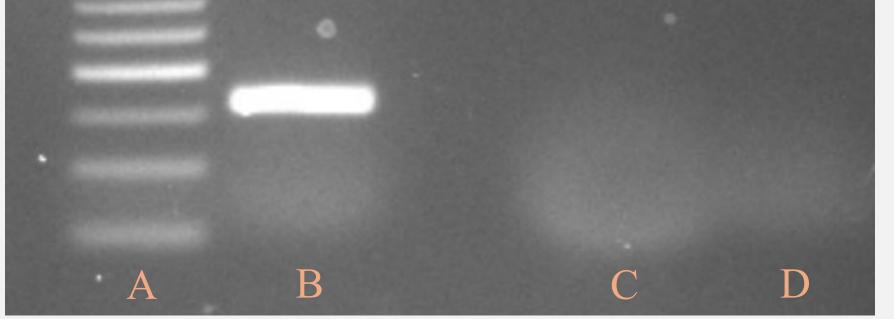
DNA from males (geldings) was used as positive controls.

DNA from nulliparous mares and water were used as negative controls.

Strict room isolation was performed to prevent contamination of test samples by male positive controls.

Future Goals

- Ongoing troubleshooting to eliminate contamination in water lane.
- FISH analysis on archived paraffin-



A B C D E F G H I J K L M N O P Q R

Figure 3. Lane A, 50 bp ladder. Lane B, positive control male sample. Lanes C and D, negative control female samples.

Figure 4. Lane A, 50 bp ladder. Lanes B and C, positive male controls. Lane D, negative water control. Lane D tested positive which indicates contamination of the water or other PCR products. Lanes E and F, negative female controls. Lanes G-R, test parous/multiparous female samples.

Fish analysis on archived paralling embedded formalin hoof lamellar tissue from healthy and laminitic horses.
<u>Acknowledgements</u>: Research Grant for Jessica Warwick is supported by the Animal Health Foundation of St Louis MO. Stipend for Jessica Warwick is supported by the department of Veterinary Medicine and Surgery.