

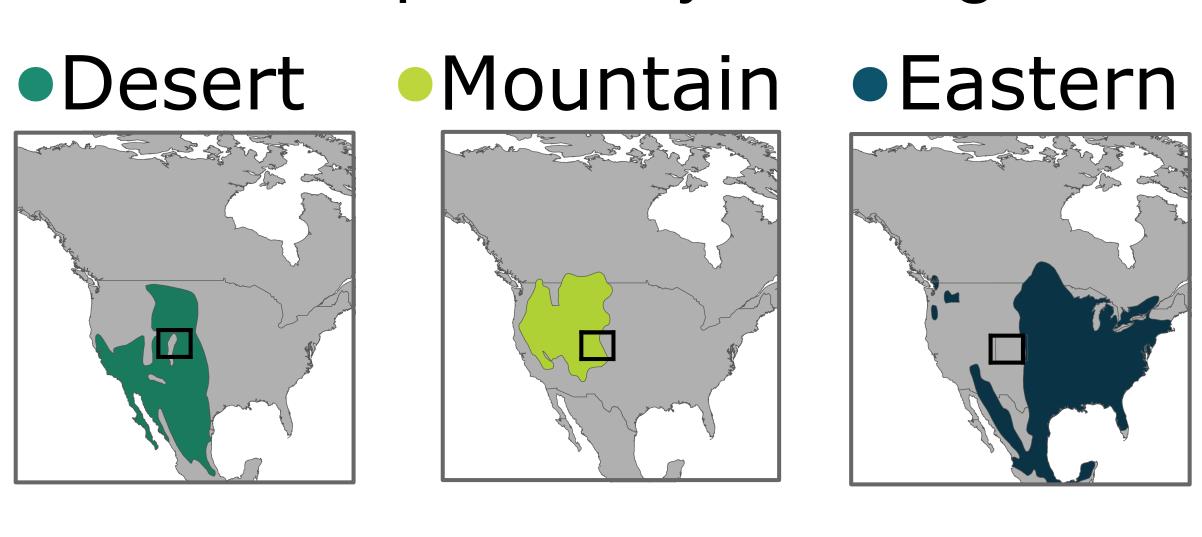
Potential Hybridization Among Three Species of Colorado Cottontails



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Main Question

- Humans are disrupting the ranges of species, providing new opportunities for hybridization.
- Are the three overlapping cottontail species hybridizing?



Results

- The genetic species ID and the morphological species ID are consistent for most individuals.
- We were able to identify individuals that were morphologically ambiguous using mitochondrial sequencing.
- Potential hybrids: Two
 morphologically eastern cottontails
 sort with desert cottontails, and one
 mountain cottontail sorts with eastern
 cottontails.

Fig 2: Phylogenetic tree of *cytb* sequences from Colorado rabbits (*) with reference sequences and rooted with pika sequences. Morphological species ID is noted with color, and genetic species ID is annotated on the right. Potential hybrids are indicated with arrows.

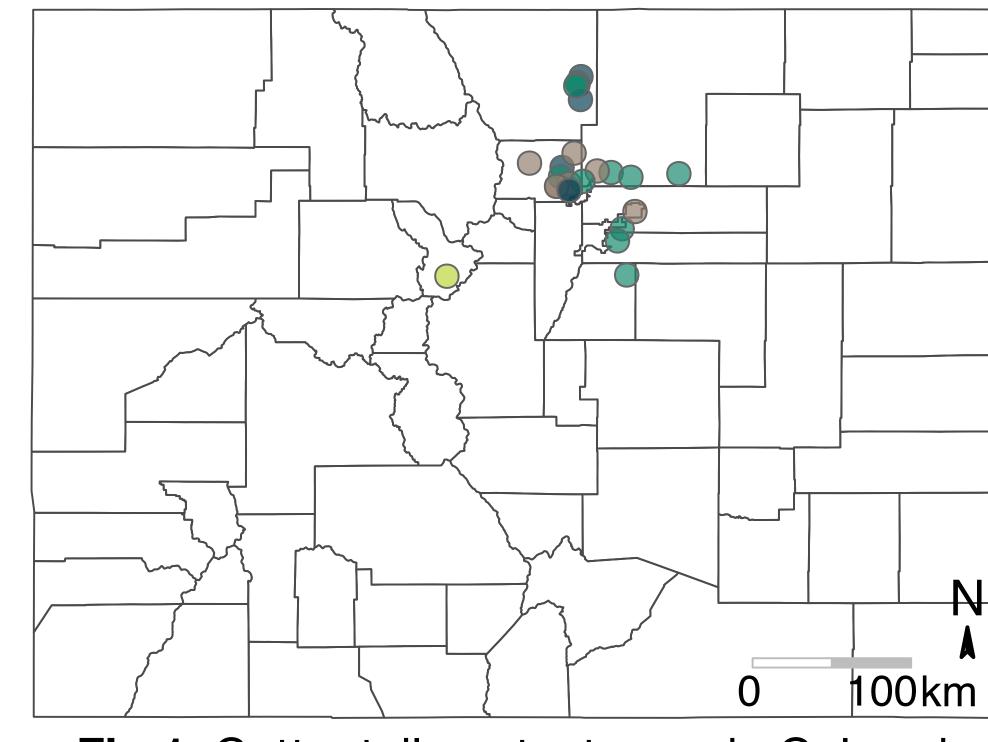
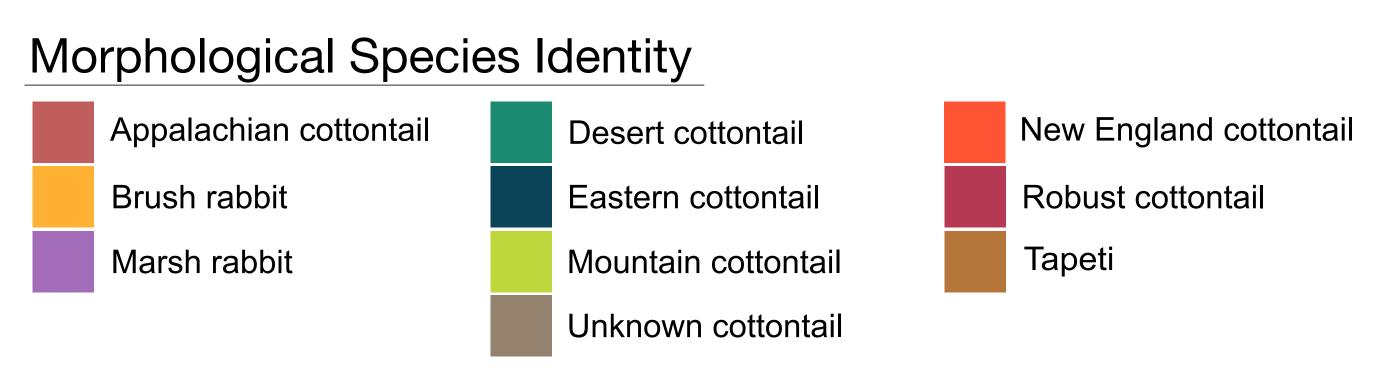
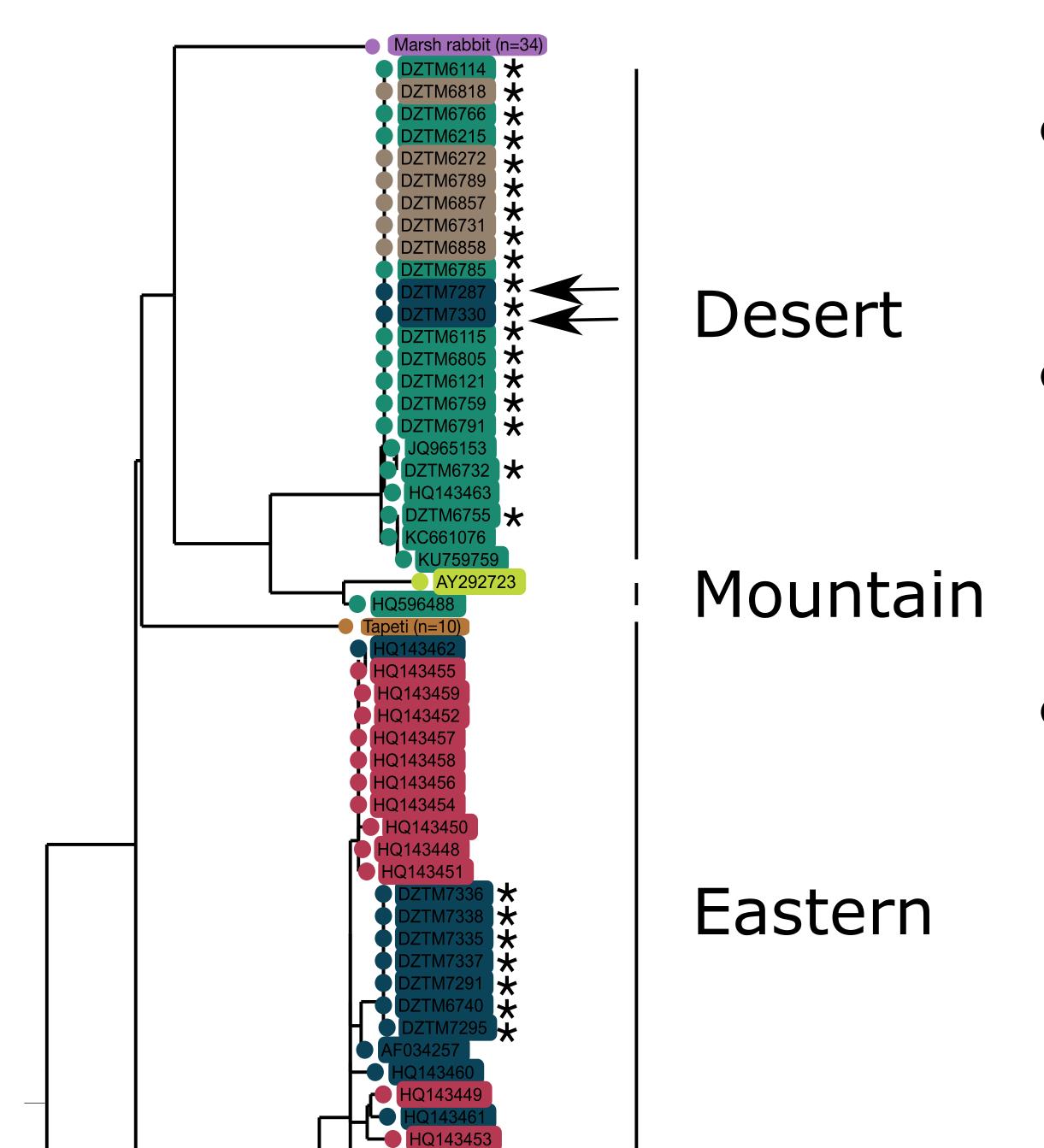


Fig 1: Cottontail contact zone in Colorado





Methods

- Morphological ID was determined by museum staff (DMNS)
- Sequenced cytb mitochondrial gene (n=28) and built phylogenetic tree to determine genetic ID
- Identified mismatches between morphological and genetic species ID to look for hybridization

Future Directions

- Increase sequencing of mountain cottontails
- Expand sequencing to include more genes to confirm hybridization
- How are species barriers maintained if hybridization is not ocurring?

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