XEROSPERMOPHILUS GROUND SQUIRRELS IN SOUTHERN CALIFORNIA

Genetic Structure Across a Contact Zone

Those Responsible

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Xerospermophilus Ground Squirrels in the Mojave Desert

- Two closely-related species are found in the Mojave Desert of California
 - Mohave ground squirrel (X. mohavensis)
 - Round-tailed ground squirrel (X. tereticaudus)
- □ Their distribution has been considered allopatric
 - MGS in winter rainfall western Mojave Desert
 - RTGS in summer rainfall eastern Mojave and further east and south

Conservation Status

- The Mohave ground squirrel is endemic to a small area in the western Mojave Desert
- It has been listed as Threatened under the California Endangered Species Act since 1984
- The round-tailed ground squirrel is widely distributed in the eastern Mojave and ranges into southern Nevada, much of Arizona, and south into Mexico





Mohave Ground Squirrel with Whitetailed Antelope Squirrel



Round-tailed Ground Squirrel





Figure 2. Geographic distribution of CNDDB Mohave ground squirrel records in the Barstow area from the period 1975-1993.

Changes in Distribution

- RTGS now found west of Mojave River in disturbed farming country around Hinkley
- This species appears much more of a generalist although no studies in CA of diet or habitat requirements
- Tolerant of disturbance and often found in and around towns and farms



Figure 4. Geographic distribution of Mohave ground squirrel and round-tailed ground squirrel trapping records from the Barstow area from the period 1914-2009.

Interactions Between the Two Species

- By 2007 it was clear that X. tereticaudus occurred quite a bit W of the Mojave river
- There was a plan to collect tissue samples from Xerospermophilus captured in that area
- A number of biologists contributed to this effort, resulting in a series of samples that were studied in Prof. Matocq's lab at University of Nevada, Reno

Genetic Methods

- A total of 127 samples were analyzed, 55 from the focal area west of Barstow
- 13 microsatellite loci were amplified
- Genetic clusters were identified by Bayesian assignment using structure ver. 2.2
- Each individual was assigned to a genetic cluster, but there could be partial assignment to >1 cluster, possibly indicating hybrid ancestry

Identifying Hybrids

- Software NEWHYBRIDS 1.1 Beta used to assign individuals to 1 of 6 genotypic classes:
 - Pure X. tereticaudus
 - Backcross to X. tereticaudus
 - F1 hybrid
 - F2 hybrid
 - Backcross to X. mohavensis
 - Pure X. mohavensis



Figure 5. Geographic Distribution of Mohave ground squirrels, round-tailed ground squirrels, and hybrids in the Hinkley area based upon genetic analysis.



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Genetic Conclusions

- Pure parental types of both species occur in spatial proximity
- Hybridization occurs occasionally
- Some hybrids are fertile
- Backcrossing occurs in both parental directions
- More detailed SNP analysis will provide deeper insight into the extent of introgression

Ecological and Evolutionary Implications

- There is strong evidence that X. tereticaudus are moving westward into undisturbed desert habitat historically occupied by X. mohavensis
- Is this related to long-term shifts in rainfall and vegetation due to climate change?
- Are we seeing a potential breakdown of genetic barriers between the two species?
- Is this a real threat to the genetic integrity of the threatened Mohave ground squirrel?

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