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Home-range size and overlap of sympatric male mule and white-tailed deer in Texas

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Home-range size and overlap of sympatric male mule and white-tailed deer in Texas

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In Texas, the distributions of desert mule deer (*Odocoileus hemionus eremicus*) and white-tailed deer (*Odocoileus virginianus*) overlap in portions of the Trans-Pecos region, the western edge of the Edwards Plateau, and the Panhandle region (Smith 1987). Landowners and wildlife managers have become concerned in recent decades as white-tailed deer have become more abundant in areas previously considered desert mule deer habitat (Harwell and Gore 1981), even as mule deer have decreased or disappeared entirely from some areas now inhabited by white-tailed deer (Wiggers and Beasom 1986). The amount of area used by male deer and their survival are of interest to private landowners and managers due to the significant economic contribution of hunting in Texas (Harveson et al. 2000). Income generated by hunting leases or other wildlife recreation can supplement or even exceed that from traditional livestock operations (Butler and Workman 1993). Because of higher bag limits and longer seasons for hunting white-tailed deer (Texas Parks and Wildlife Department 2000, 2001, 2002), some managers may wish to revise management activities to benefit primarily white-tailed deer, whereas others may prefer to reverse the increase of white-tailed deer in the area and favor mule deer.

Our objectives were to determine whether home-range sizes differed between the species and to determine the degree of home-range and core-area overlap between the species. Because allopatric male white-tailed deer in semiarid and arid environments have smaller home ranges (Michael 1965, Gallina et al. 1997) than male mule deer in arid environs have (Dickinson and Garner 1979, Relyea et al. 2000), we predicted that mule deer would have larger home ranges than white-tailed deer. However, we expected high home-range overlap between the species, as neither is territorial and they have similar diets (Anthony 1972, Krausman 1978).

We conducted our study on 5 contiguous ranches (approximately 323 km² total) in the northwestern corner of Crockett County, Texas. Estimated deer densities (both sexes) from a helicopter census of the entire area conducted in 2001 were 2.4 mule deer · km⁻² and 1.6 white-tailed deer · km⁻² (Brunjes et al. 2006). The site was located on the western edge of the Edwards Plateau. Low elevations were dominated by mesquite (*Prosopis* sp.), creosote

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**Key words:** home range, males, movements, survival, mule deer, white-tailed deer.
(Larrea tridentata), tarbush (Flourensia cernua), and prickly pear (Opuntia sp.). Juniper (Juniperus sp.) was the dominant woody species on slopes and mesa tops. Washes supported dense thickets of hackberry trees (Celtis occidentalis). The more xeric slopes supported aridland plants such as yuccas (Yucca sp.) and ocotillo (Fouquieria splendens; Correll and Johnston 1970).

Topography consisted of broad, level plateaus, rolling hills, and steep canyon walls. Elevation ranged from 700 to 915 m. Mean annual precipitation for 2000–2002 was 25 cm; the average for 1963–1997 was 43 cm. Most rainfall occurred from May to September, with highest amounts usually falling in September. The average annual low temperature was 10 °C; the average annual high was 26 °C. Temperatures ranged from a minimum daily low of −1 °C to a maximum daily high of 16 °C in winter and from 16 to 32 °C in summer (National Oceanic and Atmospheric Administration 2000, 2001, 2002).

Livestock production, oil production, and hunting were the primary land-use activities in the region. On 2–3 February 2000 and 30 January 2001, personnel from Holt Helicopters (Uvalde, TX) captured deer with a net gun fired from a helicopter (Krausman et al. 1985). We recorded sex and condition of each animal and estimated the age of deer by the tooth wear and replacement method (Severinghaus 1949, Robinette et al. 1957). We fitted each male deer with a numbered plastic eartag and a 500-g radio-collar with a mortality sensor (Telonics, Mesa, AZ).

We conducted radio-tracking with a truck-mounted null-peak system consisting of two 4-element Yagi antennas mounted on a rotating, telescoping boom in the bed of a truck. We used the software program LOAS (Ecological Software Solutions, Sacramento, CA) to calculate deer locations from telemetry bearings. We located collared males >2 times per month during January through August 2000–2002 to estimate home ranges. Using a minimum of 30 locations, we calculated home ranges and core areas for 7 deer of each species for 2001 and 2002. Individual deer locations were not triangulated during September through mid-January, because landowners restricted our access to their property during deer-hunting season. We rotated the timing of relocations sequentially through 3 time blocks (05:00–10:59, 11:00–16:59, and 17:00–24:00). We used the Animal Movement extension (Hooge and Eichenlaub 2000) for ArcView GIS software (ESRI, www.esri.com) to calculate 95% and 50% fixed-kernel home-range estimates and minimum convex polygons (MCP) to facilitate comparison to previous studies. We calculated 50% kernel home ranges as an approximation of each animal’s core area (Loveridge and Macdonald 2003).

We used ArcView to identify the polygon created when the home ranges of 2 individuals overlapped. For comparisons, each overlap polygon was assigned to 1 of 3 dyads—mule deer:mule deer (MM), mule deer:white-tailed deer (MW), or white-tailed deer:white-tailed deer (WW). If at least one location of either animal occurred within that overlap polygon, we calculated an overlap index using the following ratio:

\[
\frac{[(n_1 + n_2)/(N_1 + N_2)] \times 100}{\alpha} = 0.05, \text{ and examination of residuals supported the assumption that the data were normally distributed. Of the 7 mule deer and 7 white-tailed deer, we had no overlap for 6 of the 7 white-tailed deer.}
\]

We used Levene’s test to check for homogeneity of variance for all comparisons and we examined residuals for normality (Zar 1999, Bryce et al. 2002). We used ANOVA (\(\alpha = 0.05\)) to compare mean home-range sizes between years and ages within species and between species, and to test for interactions (White and Garrott 1990). Because of unequal sample sizes, Fisher’s LSD test was used for means separation in overlap comparisons.

We captured and fitted 10 males of each species with radio-collars in January 2000. In January 2001, we captured and collared an additional 8 males of each species. Mean age at capture was 3.5 years (range 3.5–4.5) for mule deer and 4.5 years (range 3.5–6.5) for white-tailed deer. We did not have any deer with >30 locations during 2000 to calculate home ranges for.

Average bearing error was ±7° based on triangulated locations of collars in known locations. Levene’s test for homogeneity of variance was not significant for any comparison at \(\alpha = 0.05\), and examination of residuals supported the assumption that the data were normally distributed. Of the 7 mule deer and 7 white-tailed deer, we had no overlap for 6 of the 7 white-tailed deer.
white-tailed deer for which >30 locations per year were available, 5 mule deer and 3 white-tailed deer were tracked in both years, and we averaged their calculated home-range size for later analyses. The remaining deer had >30 locations per year for only 1 of the 2 years. Within each species, neither home-range size nor core-area size differed between years or ages, nor was there a year × age interaction or a species × year interaction (Table 1), so we pooled data for each species. Home-range size and core-area size did not differ between species (Table 2). Mean home-range size for mule deer was 8.8 km² ($\bar{s}_X = 1.6$) and mean core-area size was 1.0 km² ($\bar{s}_X = 0.2$); mean home-range size for white-tailed deer was 7.4 km² ($\bar{s}_X = 1.3$) and mean core-area size was 1.1 km² ($\bar{s}_X = 0.2$). The home range calculated for 2002 overlapped the home range calculated for 2001 for 4 mule deer and 3 white-tailed deer (Fig. 1). Only 1 mule deer shifted home range between years. The mean proportion of the 2001 home range that was overlapped by the 2002 home range was 0.4 ($\bar{s}_X = 0.2$) for mule deer and 0.5 ($\bar{s}_X = 0.1$) for white-tailed deer. The amount of overlap between years was not different between species ($F_1 = 0.15, P = 0.71$).

![Fig. 1. An example of overlapping 2001 and 2002 home ranges and core areas for adult male mule deer in west central Texas.](image)

**Table 1.** ANOVA comparison within and between adult male mule and white-tailed deer by year and age for 2 independent variables: 95% kernel home-range area and 50% kernel core area. Deer were studied in west central Texas, January through August, 2001–2002.

<table>
<thead>
<tr>
<th>Estimate</th>
<th>Species, sample size</th>
<th>Variable</th>
<th>df</th>
<th>$F$</th>
<th>$P$</th>
</tr>
</thead>
<tbody>
<tr>
<td>95% kernel</td>
<td>Mule deer, $n = 7$</td>
<td>Age</td>
<td>2</td>
<td>0.22</td>
<td>0.81</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Year</td>
<td>1</td>
<td>1.00</td>
<td>0.34</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Age × Year</td>
<td>3</td>
<td>10.9</td>
<td>0.41</td>
</tr>
<tr>
<td>White-tailed</td>
<td></td>
<td>Age</td>
<td>4</td>
<td>1.47</td>
<td>0.34</td>
</tr>
<tr>
<td>deer, $n = 7$</td>
<td></td>
<td>Year</td>
<td>1</td>
<td>0.06</td>
<td>0.81</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Age × Year</td>
<td>6</td>
<td>1.43</td>
<td>0.41</td>
</tr>
<tr>
<td>Combined, $n = 14$</td>
<td>Species × Year</td>
<td>3</td>
<td>0.87</td>
<td>0.47</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>Species</td>
<td>1</td>
<td>0.46</td>
<td>0.51</td>
</tr>
<tr>
<td>50% kernel</td>
<td>Mule deer, $n = 7$</td>
<td>Age</td>
<td>2</td>
<td>0.15</td>
<td>0.87</td>
</tr>
<tr>
<td>(core area)</td>
<td></td>
<td>Year</td>
<td>1</td>
<td>1.80</td>
<td>0.21</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Age × Year</td>
<td>3</td>
<td>1.45</td>
<td>0.93</td>
</tr>
<tr>
<td>White-tailed</td>
<td></td>
<td>Age</td>
<td>4</td>
<td>1.20</td>
<td>0.41</td>
</tr>
<tr>
<td>deer, $n = 7$</td>
<td></td>
<td>Year</td>
<td>1</td>
<td>0.34</td>
<td>0.57</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Age × Year</td>
<td>6</td>
<td>0.71</td>
<td>0.69</td>
</tr>
<tr>
<td>Combined, $n = 14$</td>
<td>Species × Year</td>
<td>3</td>
<td>0.74</td>
<td>0.54</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>Species</td>
<td>1</td>
<td>0.15</td>
<td>0.71</td>
</tr>
</tbody>
</table>

*For comparisons each overlap polygon was assigned to 1 of 3 dyads: mule deer: mule deer (MM), mule deer: white-tailed deer (MW), or white-tailed deer: white-tailed deer (WW).

**Table 2.** Mean percentage overlap indices for 95% kernel home ranges and 50% kernel core areas of adult male mule and white-tailed deer in west central Texas, January through August, 2001–2002. Indices for home range and core area were compared among species using ANOVA. Fisher’s LSD was used for means separation when ANOVA results were significant. Means followed by the same letter within a year were not different at $\alpha = 0.05$.

<table>
<thead>
<tr>
<th>Dyada</th>
<th>$n$</th>
<th>Average overlap</th>
<th>$s_T$</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Home ranges</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>MM</td>
<td>17</td>
<td>29.32 a</td>
<td>6.25</td>
</tr>
<tr>
<td>WW</td>
<td>15</td>
<td>44.32 a</td>
<td>10.08</td>
</tr>
<tr>
<td>MW</td>
<td>10</td>
<td>8.57 b</td>
<td>3.24</td>
</tr>
<tr>
<td>Core areas</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>MM</td>
<td>3</td>
<td>39.10</td>
<td>19.75</td>
</tr>
<tr>
<td>WW</td>
<td>5</td>
<td>48.36</td>
<td>10.84</td>
</tr>
</tbody>
</table>

$\bar{s}_X$,$ s_T$,$ n$.
During both years, the home range of every study animal was overlapped by the home range of more than one other study animal. Each study animal’s home range overlapped the home range of more than one collared individual of the other species (Table 2), except one white-tailed deer whose home range did not overlap the home range of any collared mule deer. Overlapping core areas between collared animals were less common than overlapping home ranges (Table 2). Home-range and core-area overlap indices did not differ between years within dyads (home range: $F_{1} = 0.29, P = 0.59$; core area: $F_{1} = 0.14, P = 0.7$), nor did we detect a year $\times$ dyad interaction (home range: $F_{2} = 2.41, P = 0.10$; core area: $F_{1} = 1.60, P = 0.29$). Home-range overlap indices did not differ between the MM and WW dyads, but MW overlap indices were less than those of either intraspecific dyad ($F_{2} = 7.17, P = 0.002$). We observed only one instance in which the calculated core areas overlapped for a mule deer and a white-tailed deer, but no locations of either species occurred within the overlap polygon. Core-area overlap indices did not differ between MM and WW dyads ($F_{1} = 1.25, P = 0.35$).

Our results did not support our initial predictions. However, differences in home-range size may not have been detectable because of our small sample size and the difficulty of tracking deer. According to competition theory, 2 species with similar life history traits should partition resources when sympatric (Hardin 1960). Diet does not appear to drive habitat partitioning between these species (Hill and Harris 1943, Allen 1968, Martinka 1968, Krausman 1978), suggesting that some other resource (e.g., space) was driving resource partitioning. Equivalent home-range sizes may be a direct result of sympatry, as both species exist on the same forage resource. Although the larger body mass of mule deer suggests they need larger home ranges, habitat productivity appears to be a more important influence on ungulate home-range size (Relyea et al. 2000). The observed interspecific home-range overlap indicated that habitat partitioning may have occurred on a temporal scale or at a finer spatial scale than can be detected by our home-range-level analyses.

Core-area overlap indicated a greater potential for competition than home-range overlap (Wauters and Dhondt 1985), yet interspecific core-area overlap occurred only once. This avoidance could be an artifact of data collection, as deer spend much of their time bedded down compared to the time they spend on foraging and other activities. Thus, the smaller degree of overlap between the species’ core

![Fig. 2. An example of overlapping home ranges and core areas for adult male mule and white-tailed deer in west central Texas, January–August, 2002.](image)
areas could be due, in part, to differences in preferred bedding sites, as mule deer prefer bed sites with less cover and steeper slopes (Avey et al. 2003) than those preferred by white-tailed deer. However, we obtained most locations during crepuscular periods to improve our sample of non-bedded deer. Avoidance by individuals of the core areas of the other species may indicate that interspecific competition influenced the spatial distribution of deer more than intraspecific competition.

Individuals of both species appeared to maintain their home ranges within the same general area during both years. Only one male, a 4.5-year-old mule deer, completely shifted his home range between 2001 and 2002. This fidelity, coupled with significant interspecific home-range overlap, suggested that neither species was actively driving the other out of the area and that spatial coexistence was stable.

Because we were unable to track deer during September through December, our estimates of home range are probably underestimates, as male deer tend to increase movements outside of their home range during the breeding season (Rodgers et al. 1978, Dickinson and Garner 1979, Relyea and Demarais 1994, Gallina et al. 1997). Our results are consistent with other research on sympatric mule deer and white-tailed deer (e.g., Martinka 1968, Anthony and Smith 1977, Krausman 1978, Wood et al. 1989) that suggests that the 2 species coexist via spatial segregation. However, the question remains: at what scale do these species partition the habitat? Avey et al. (2003) indicated that there was habitat separation between species, suggesting that managers might be able to manage habitat for both species. That this separation is occurring below the home-range scale is supported by our results on home-range and core-area size, and by the fact that the differences observed by Avey et al. 2003 occurred at the individual point locations of deer. The degree of home-range similarity and overlap we observed suggest that management to primarily benefit only one species may be difficult to conduct on a large scale (e.g., an entire ranch).

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