DNA Fingerprinting Reveals Polygyny in the Lesser Kestrel (Falco naumanni)

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Diurnal birds of prey are predominantly monogamous (Newton 1979, Faaborg and Bednarz 1990). However, some species have adopted alternative mating systems, such as polygyny, polyandry or cooperative breeding (Newton 1979, Faaborg and Bednarz 1990, Heredia and Donázar 1992, Tella 1993). Polygamous trios have been observed in at least 11 raptor species, where polygyny occurs more frequently (but see Gibbs et al. 1990, Gyllensten et al. 1990, Dunn and Robertson 1993). Monegros, northeastern Spain (41ø25'N, 0ø11'E), where a population of 230 pairs of Lesser Kestrel bred in 49 colonies located in abandoned farmhouses (Tella et al. in press). Lesser Kestrels are migratory and spend the winter in Africa (Cramp and Simmons 1980). In individuals return to the colonies in late February and throughout March. In 1991, the average egg-laying date was 7 May (n = 199). At different times during the breeding period, adult Lesser Kestrels (n = 270) were caught at night while roosting in their nests. All birds were banded with color-numbered PVC bands for identification by telescope (Donázar et al. 1992, Negro et al. 1992). Adult males were assigned to two age categories—yearling and ≥ 2 years old—according to plumage (Cramp and Simmons 1980). Blood samples were taken from most adult birds, as well as from the offspring at selected nests that were monitored periodically to determine clutch and brood size.

About 0.4 ml of blood taken from the brachial vein was preserved in a lysis buffer (Seutin et al. 1991) that permitted its transport and temporary storage at room temperature. DNA analyses were conducted in the Animal Science Department of McGill University in Montreal, Canada. Aliquots of the samples (0.25 ml) were mixed with 5 ml of 1 x SSC and centrifuged at 7,000 rpm for 15 min. The resulting pellet was re-suspended in 2 ml of 0.2 M sodium acetate and 100 μl of 20% SDS. We extracted the samples with 2 ml of a mixture consisting of equal parts of phenol and chloroform-isooamy alcohol (24:1). The samples were centrifuged at 2,000 rpm for 20 min, and the supernatant was subjected to a second extraction. The DNA was precipitated with cold ethanol and preserved in -20°C. We used as the threshold limit for parentage exclusion the lower 95% confidence limit (BSC = 0.324) of the BSC distribution for first-order relatives (Lynch 1991) and Negro et al. (in press).

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Comparisons of banding patterns were confined to lanes on the same gel. Parentage was determined by band-exclusion analysis (Deckert et al. 1993, Sheldon and Burke 1994). Bands in the offspring’s fingerprint were matched to bands present in the parents. The presence/absence of unattributable bands is the primary basis for determining parentage. In addition, band-sharing coefficients (BSC; i.e. proportion of bands in fingerprint shared by any two given individuals) were calculated. In our population, the “background” BSC is 0.21 (Negro et al. in press) and, thus, the expected BSC (Lynch 1991) for first-order relatives is 0.60. We used as the threshold limit for parentage exclusion the lower 95% confidence limit (BSC = 0.324) of the BSC distribution for first-order relatives (Deckert et al. 1993, Sheldon and Burke 1994).

Methods.—Our study was carried out in 1993 in Los Monegros, northeastern Spain (41°25′N, 0°11′E), where a population of 230 pairs of Lesser Kestrel bred in 49 colonies located in abandoned farmhouses (Tella et al. in press). Lesser Kestrels are migratory and spend the winter in Africa (Cramp and Simmons 1980). Individuals return to the colonies in late February and throughout March. In 1991, the average egg-laying date was 7 May (n = 199). At different times during the breeding period, adult Lesser Kestrels (n = 270) were caught at night while roosting in their nests. All birds were banded with color-numbered PVC bands for identification by telescope (Donázar et al. 1992, Negro et al. 1992). Adult males were assigned to two age categories—yearling and ≥ 2 years old—according to plumage (Cramp and Simmons 1980). Blood samples were taken from most adult birds, as well as from the offspring at selected nests that were monitored periodically to determine clutch and brood size.

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The breeding chronology of the birds involved in this case of polygyny is shown in Figure 1. Nest 1, previously occupied by a pair of Lesser Kestrels, was defended by male 6U (≥ 2 years old) at the end of April. In early May, male 6U and female 2T attended the nest, which contained a single egg that we marked and measured. One week later the nest contained four eggs, including the one marked in the previous visit, and there was a new attending female, 4A. No differences in coloration or measurements were found among the eggs. Remains of other eggs were not found in or around the nest. The resulting clutch of four eggs was not much different from the average clutch of 4.45 ± SD of 0.74 eggs in the population in 1993 (n = 114). In two subsequent visits, only female 4A was observed at the nest (Fig. 1). Female 2T was not seen again during the rest of the 1993 breeding season, but she was resighted as a breeder in the study area in 1995.

The interval between the time the egg was laid by female 2T and when the first egg was laid by female 4A was one to three days. This was estimated from the dates the females were captured taking into account the typical laying interval between consecutive eggs of two days (Cramp and Simmons 1980). Measurements of the nestlings' eighth primary feather were used to age the birds (Donázar et al. 1991) and indicated that the older three young hatched within 24 h, and that the remaining egg hatched three to four days later. These results may be attributable to the fact that females usually start incubation after laying the third egg (Negro et al. 1991). All of the hatchlings fledged.

The two females that mated with male 6U had previously been paired with yearling males (Fig. 1). Why the females broke up with those males is unknown, although in the case of female 2T it could be related to fox predation and the subsequent desertion of the colony by the survivors.

Female 13, the bird that was first associated to nest 1, mated later with male 5U and attended nest 2 in the same colony (Table 1). DNA fingerprinting demonstrated that female 13 and male 5U were the parents of the nestlings in nest 2.

The band-exclusion analysis and band-sharing coefficients indicate that male 6U was the father of all nestlings, while female 2T was the mother of the older nestling, and female 4A was the mother of the...
TABLE 1. Results of the band-exclusion analysis. Proportion of nestling bands not present in male 6U and female indicated.

<table>
<thead>
<tr>
<th>Nestling</th>
<th>Female 2T</th>
<th>Female 4A</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>0/8</td>
<td>3/8</td>
</tr>
<tr>
<td>2</td>
<td>2/9</td>
<td>0/9</td>
</tr>
<tr>
<td>3</td>
<td>4/8</td>
<td>0/8</td>
</tr>
<tr>
<td>4</td>
<td>2/7</td>
<td>0/7</td>
</tr>
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</table>

remaining three nestlings. There were no unattributable bands when the correct female was included in the comparison (Table 1). Band-sharing coefficients for all pairwise combinations of adults and nestlings are given in Table 2. Values for nestlings and their assigned parents, as well as those for true siblings, are above of the threshold limit for parentage exclusion, and the mean (0.534 ± 0.14, n = 11) is close to 0.6, the expected BSC value for first-order relatives in our population.

Discussion.—Mixed maternity in broods or clutches has been documented by molecular markers in several bird species and, usually, has been interpreted as the result of intraspecific brood parasitism (Pinxten et al. 1993, Avise 1994, Meek et al. 1994, Negro et al. in press). We ruled out this hypothesis because the attending male was related to all chicks in the brood. Other possible causes of multiple maternity could be: (1) quasiparasitism (Petrie and Møller 1991, Birkhead et al. 1990), in which the eggs dumped by the parasitic female were fertilized by the nest-attending male; (2) mate replacement of the female that laid eggs first; and (3) polygyny, in which two or more females lay eggs and provide parental investment.

Given that female 2T laid first and at a time when female 4A was in a different colony more than 1 km away (see Table 1), our case of multiple maternity can hardly be a case of quasiparasitism. Rather, it is a case in which two females fertilized by the same male laid eggs in the same nest. We can discard the hypothesis that mate replacement took place due to death of the first laying female, but we cannot exclude replacement due to desertion or displacement (Choudhury 1995) of female 2T. Given that our nest monitoring was not intensive, it is also possible that females 2T and 4A participated in the incubation and chick rearing in nest 1. Although rare and usually unsuccessful, nest-sharing polygyny has been observed in raptors (Newton 1979, Poole 1989). Among other species of birds, nest-sharing polygyny has been documented by DNA analysis in the European Starling (Sturnus vulgaris; Pinxten et al. 1994).

The aborted polygynous trios observed by Hiraldo et al. (1991) started 50 to 70 days before the primary female laid eggs. In our case, however, polygyny occurred precisely at laying time. The two females previously had been at different colonies paired to yearling males. The reproductive value of these males is much lower than that of older males (Negro 1991, Hiraldo et al. unpubl. manuscript), and the females may have deserted them to pair with a better male. The bigamous male, in turn, apparently was unpaired but attending the nest site at the beginning of the laying period. Even though the polygyny attempt started late in the season, it may have been successful because food conditions were favorable. In fact, breeding success in our population in 1993 (Tella et al. in press) was almost double as the one reported by Hiraldo et al. (1991).

The results of a broader paternity study in our population (Negro et al. in press) indicate that polygyny occurred in 1 of 27 families. Thus, the frequency of polygamous matings in the population was probably low. This polygyny case could have easily passed unnoticed without repeated nest visits or if molecular markers had not been used.

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