

University of Nebraska - Lincoln

DigitalCommons@University of Nebraska - Lincoln

Mammalogy Papers: University of Nebraska State
Museum

Museum, University of Nebraska State

1977

Karyotypes of Shrews of the Genera *Cryptotis* and *Blarina* (Mammalia: Soricidae)

Hugh H. Genoways

University of Nebraska - Lincoln, h.h.genoways@gmail.com

James C. Patton III

University of Georgia, Athens

J. R. Choate

Museum of the High Plains, Fort Hays Kansas State College

Follow this and additional works at: <http://digitalcommons.unl.edu/museummammalogy>



Part of the [Zoology Commons](#)

Genoways, Hugh H.; Patton, James C. III; and Choate, J. R., "Karyotypes of Shrews of the Genera *Cryptotis* and *Blarina* (Mammalia: Soricidae)" (1977). *Mammalogy Papers: University of Nebraska State Museum*. 147.

<http://digitalcommons.unl.edu/museummammalogy/147>

This Article is brought to you for free and open access by the Museum, University of Nebraska State at DigitalCommons@University of Nebraska - Lincoln. It has been accepted for inclusion in Mammalogy Papers: University of Nebraska State Museum by an authorized administrator of DigitalCommons@University of Nebraska - Lincoln.

Specialia

Karyotypes of Shrews of the Genera *Cryptotis* and *Blarina* (Mammalia: Soricidae)¹

Hugh H. Genoways^a, James C. Patton III^b and J. R. Choate^c

^aCarnegie Museum of Natural History, Pittsburgh, Pennsylvania, USA (currently affiliated with University of Nebraska-Lincoln, Lincoln, Nebraska, USA; hgenoways1@unl.edu); ^bDepartment of Zoology, University of Georgia, Athens, Georgia, USA; and ^cMuseum of the High Plains, Fort Hays Kansas State College, Hays, Kansas, USA

Summary. *Cryptotis parva* has a diploid number of 52 and a fundamental number of 50. *Blarina brevicauda* in Nebraska and Pennsylvania has a diploid number of 49 or 50 and a fundamental number of 48. *Blarina carolinensis* in Nebraska and Kansas has a diploid number of 52 and a fundamental number of 62. The X-chromosome in all 3 species is a large metacentric chromosome. The Y-chromosome is a small acrocentric in *Blarina*, whereas in *Cryptotis* it is a small subtelocentric.

Information has been published (for example, Baker and Hsu²; Fedyk and Ivanitskaya³; Meylan^{4,9}; Meylan and Hausser^{10,11}; Hausser et al.¹² and papers cited therein) on various aspects of the karyology of shrews (family Soricidae), but few data are available for the North American genera *Cryptotis* and *Blarina*. The genus *Cryptotis* is represented in the United States by only one species, *C. parva* (the least shrew), which occurs throughout much of the eastern half of this country as well as in mesic and montane habitats in Mexico and Central America (distribution and habitats summarized by Whitaker¹³). The relationships of this species to other members of the genus in Latin America recently have been reviewed (Choate¹⁴), and the taxonomy of the species in the United States probably contains few problems. This is not true, however, for the genus *Blarina* (short-tailed shrews), the distribution of which includes only the eastern half of the United States and adjacent regions of Canada (Hall and Kelson¹⁵). Prior to 1972, the genus *Blarina* generally was assumed to consist of only one species, *B. brevicauda*; a second species, *B. telmalestes*, had been described from the Dismal Swamp of coastal Virginia and North Carolina (Paul¹⁶), but was of doubtful taxonomic status (Choate¹⁷). Then, in 1972, Genoways and Choate¹⁸ presented data indicating that in Nebraska a large, northern subspecies (*B. B. brevicauda*) and a smaller, southern subspecies (*B. B. carolinensis*) were behaving as good biological species. Subsequently, most authors have treated these taxa as a distinct species (*B. brevicauda* and *B. carolinensis*, respectively). Later, based on their study of fossils of *Blarina*, Graham and Semken¹⁹ recognized a third Recent species (*B. kirtlandi*) in the genus. We continue to recognize only 2 species of *Blarina* in this paper.

Certainly, much additional systematic work is needed on North American shrews, especially *Blarina*. To aid in these studies, we present below the karyotypic data on these shrews that we have amassed over the past several years. The only previously published information for these shrews pertained to *B. brevicauda talpoides* (Meylan^{6,8}) and *B. b. kirtlandi* (Lee and Zimmerman²⁰). All karyotypic preparations were made according to methods described by Baker²¹.

Cryptotis parva (Figure 1). The diploid number for the least shrew is 52 and the fundamental number without the sex-chromosomes is 50. The autosomes, which are all acrocentric, range in size from one large pair to several minute pairs. The X-chromosome is a large metacentric and the Y-chromosome is a small subtelocentric. *Blarina brevicauda* (Figure 2). Specimens of *B. b. brevicauda* from Nebraska have a diploid number of either 49 or 50 and a fundamental number of 48. The polymorphism in diploid number is the result of a Robertsonian fission/fusion between a pair of large acrocentric autosomes and a pair of small acrocentric autosomes. Specimens with a diploid number of 48, resulting from fusion of both members of these pairs, were not represented in our material. The X-chromosome is a large metacentric and the Y-chromosome is a small acrocentric.

These diploid and fundamental numbers are the same as those reported by Meylan^{6,8} for *B. b. talpoides* from Ontario and by Lee and Zimmerman²⁰ for *B. b. kirtlandi* from Illinois. Our 2 specimens of *B. b. kirtlandi* from Pennsylvania also agree in these numbers. The Robertsonian polymorphism described above also was noted by Meylan and by Lee and Zimmerman; Meylan found 2N = 50 in 16 and 2N = 49 in 5 specimens, whereas Lee and Zimmerman found 2N = 50 in 46 specimens, 2N = 49 in 6, and 2N = 48 in 1 specimen. Combined with our specimens in which 6 had 2N = 50 and 4 had 2N = 49, this gives a ratio of 68: 15: 1 for this Robertsonian polymorphic system.

Meylan⁶ described (but did not illustrate) the Y-chromosome of his material as being a very small metacentric chromosome, whereas we found it to be a small acrocentric. Lee and Zimmerman²⁰ do not describe the Y-chromosome in their specimens. *Blarina carolinensis* (Figure 3). Specimens of *B. carolinensis* from Nebraska and Kansas had a diploid number of 52 and a fundamental number of 62. There are 4 pairs of large to medium-sized subtelocentric autosomes and 2 pairs of small submetacentric autosomes; the remaining 19 pairs of autosomes are acrocentric. The X-chromosome is a large metacentric and the Y-chromosome is acrocentric.

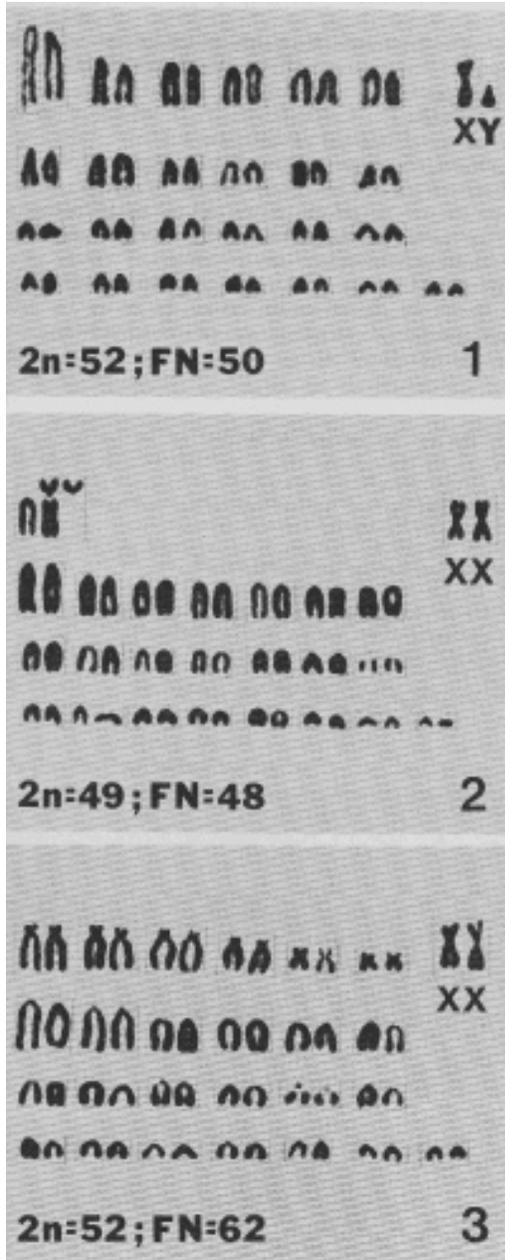


Figure 1. Male *Cryptotis parva* from 4 1/2 miles N, 7 miles E Palo Pinto, Palo Pinto County, Texas. Figure 2. Female *Blarina brevicauda* from 1 mile W Kearney, Buffalo County, Nebraska. Figure 3. Female *Blarina carolinensis* from 3 miles W Hays, Ellis County, Kansas.

The specimen of *B. carolinensis* from 1 mile west Otoe, Otoe County, Nebraska, was obtained only 15 miles south of the locality of capture of the 3 specimens of *B. brevicauda* from Cass and Sarpy counties. The existence of these divergent karyotypes within such a short distance lends support to the contention, based on morphological data (Genoways and Choate¹⁸), that the 2 phenons of *Blarina* in Nebraska represent distinct species. None of our karyological data indicate interbreeding between these taxa, but considerably more data, especially from zones of contact, will be needed before a definitive statement to this effect can be made.

Discussion. The karyotypes of these 3 shrews of the Tribe Blarini show some interesting similarities and differences. *Cryptotis parva* has the same diploid number as *Blarina carolinensis*; however, the entirely acrocentric complement of chromosomes in *C. parva* more nearly resembles that of *B. brevicauda* (although the latter possesses 2 fewer autosomes). The 2 species of *Blarina* differ in both diploid numbers and morphology of at least 6 pairs of autosomes. The morphology of the X-chromosome apparently is the same in the 3 species, but the Y-chromosome is acrocentric in both species of *Blarina* and subtelocentric in *Cryptotis*. The origin of these chromosomal differences and the course of chromosomal evolution in this group are unclear at the present time. It is hoped that planned studies of chromosomal banding patterns will help resolve these problems.

The discovery that the nominal taxa *B. brevicauda*, *kirtlandi* and *talpoides* possess the same chromosomal numbers and Robertsonian polymorphism casts serious doubt, in our minds, that these are distinct from each other at the species level. Consequently, we have not followed the proposal by Graham and Semken 19 that *kirtlandi* represents a species distinct from *brevicauda*. If *kirtlandi* is a distinct species, then specific divergence between *kirtlandi* and *brevicauda* must have been much more recent than that between *brevicauda* and *carolinensis*; the Robertsonian polymorphism, which is present in *brevicauda* but not in *carolinensis*, would need to have developed and become established in populations of *brevicauda* subsequent to its split with *carolinensis* and prior to divergence of *brevicauda* and *kirtlandi*.

Specimens examined. *Cryptotis parva*: 4 1/2 miles N, 7 miles E Palo Pinto, Palo Pinto County, Texas, 1. *Blarina brevicauda*: 1 mile W Kearney, Buffalo County, Nebraska, 5; 1/2 mile W Manley, Cass County, Nebraska, 1; 1 mile N, 2 miles W Weeping Water, Cass County, Nebraska, 1; 4 miles N Springfield, Sarpy County, Nebraska, 1; 2 miles S Rector, Westmoreland County, Pennsylvania, 1; 3 miles S Rector, Westmoreland County, Pennsylvania, 1. *Blarina carolinensis*: 3 miles W Hays, Ellis County, Kansas, 5; 5 miles N, 2 miles W Parks, Dundey County, Nebraska, 3; 1 mile W Otoe, Otoe County, Nebraska, 1.

¹ Funds for field work were provided by the American Philosophical Society from the Johnson Fund. We thank Dr. Robert J. Baker, Texas Tech University, for use of his karyological equipment and laboratory facilities.

² R. J. Baker and T. C. Hsu, *SWest. Nat.* 14,488 (1970).

³ S. Fedyk and E. Y. Ivanitskaya, *Acta Theriol.* 17,475 (1972).

⁴ A. Meylan, *Revue suisse Zool.* 72, 636 (1965).

⁵ A. Meylan, *Revue suisse Zool.* 73, 548 (1966).

⁶ A. Meylan, *Can. J. Zool.* 45, 1119 (1967).

⁷ A. Meylan, *Revue suisse Zool.* 74,685 (1967).

⁸ A. Meylan, *Revue suisse Zool.* 75,691 (1968).

⁹ A. Meylan, *Revue suisse Zool.* 78,603 (1971).

¹⁰ A. Meylan and J. Hausser, *Z. Säugetierk.* 38, 143 (1973).

¹¹ A. Meylan and J. Hausser, *Revue suisse Zool.* 81, 701 (1974).

¹² J. Hausser, J.-D. Graf and A. Meylan, *Bull. Soc. vaud. Sci. nat.* 72, 241 (1975).

¹³ J. O. Whitaker, Jr., *Mammal. Spec.* 43, 4 (1974).

¹⁴ J. R. Choate, *Univ. Kansas Publ., Mus. Nat. Hist.* 19, 195 (1970).

¹⁵ E. R. Hall and K. R. Kelson, *The Mammals of North America*, vol. 1, p. 53. Ronald Press 1959.

¹⁶ J. R. Paul, *J. Mammal.* 46, 496 (1965).

¹⁷ J. R. Choate, *J. Mammal.* 53, 116 (1972).

¹⁸ H. H. Genoways and J. R. Choate, *Syst. Zool.* 21, 106 (1972).

¹⁹ R. W. Graham and H. A. Semken, *J. Mammal.* 57,433 (1976).

²⁰ M. R. Lee and E. G. Zimmerman, *J. Mammal.* 50, 333 (1969).

²¹ R. J. Baker, in: *Biology of Bats*, vol. 1, p. 65. Editor, W. A. Wimsatt. Academic Press, 1970.