```
Homo sapiens UCP3          MVGLKPSDVPPTMAVKFLGAGTAACFADLVTFPLDTAKVRLQIQGENQAVQTARLVQYRG
Catagonus wagneri UCP3     MVGLKPSEVPPTTAVKFLGAGAAACFADLLTFPLDTAKVRLQIQGENQAARSA---QYRG
Sus scrofa UCP3            MVGLKPPEVPPTTAVKLLGAGTAACFADLLTFPLDTAKVRLQIQGENQAARSA---QYRG
                           ******.:**** ***:****:*******:******************..:*   ****

Homo sapiens UCP3          VLGTILTMVRTEGPCSPYNGLVAGLQRQMSFASIRIGLYDSVKQVYTPKGADNSSLTTRI
Catagonus wagneri UCP3     VLGTILTMVRNEGPRSLYNGLVAGLQRQMSFASVRIGLYDSVKQLYTPKGSDHSSVTTRI
Sus scrofa UCP3            VLGTILTMVRNEGPRSPYNGLVAGLQRQMSFASIRIGLYDSVKQLYTPKGSDHSSITTRI
                           **********.*** * ****************:**********:*****:*:**:****

Homo sapiens UCP3          LAGCTTGAMAVTCAQPTDVVKVRFQASIHLGPSRSDRKYSGTMDAYRTIAREEGVRGLWK
Catagonus wagneri UCP3     LAGCTTGAMAVTCAQPTDVVKVRFQASMHSGP-GSNRKYSGTMDAYRTIAREEGVRGLWK
Sus scrofa UCP3            LAGCTTGAMAVTCAQPTDVVKVRFQASIHAGP-RSNRKYSGTMDAYRTIAREEGVRGLWK
                           ****************************:* **  *:************************

Homo sapiens UCP3          GTLPNIMRNAIVNCAEVVTYDILKEKLLDYHLLTDNFPCHFVSAFGAGFCATVVASPVDV
Catagonus wagneri UCP3     GILPNITRNAIVNCAEMVTYDIIKEKVLDYHLLTDNLPCHFVSAFGAGFCATVVASPVDV
Sus scrofa UCP3            GILPNITRNAIVNCAEMVTYDVIKEKVLDYHLLTDNLPCHFVSAFGAGFCATVVASPVDV
                           * **** *********:****::***:*********:**********************

Homo sapiens UCP3          VKTRYMNSPPGQYFSPLDCMIKMVAQEGPTAFYKGFTPSFLRLGSWNVVMFVTYEQLKRA
Catagonus wagneri UCP3     VKTRYMNSPPGQYQSPLHCMLRMVTREGPTAFYKGFTPSFLRLGSWNVVMFVTYEQLKRA
Sus scrofa UCP3            VKTRYMNSPPGQYQNPLDCMLKMVTQEGPTAFYKGFTPSFLRLGSWNVVMFVSYEQLKRA
                           ************* .** **:.**:.*************************:*******

Homo sapiens UCP3          LMKVQMLRESPF
Catagonus wagneri UCP3     LMKVQMLRESPF
Sus scrofa UCP3            LMKVQMLRESPF
                           ************


Homo sapiens UCP2          MVGFKATDVPPTATVKFLGAGTAACIADLITFPLDTAKVRLQIQGESQGPVRATASAQYR
Catagonus wagneri UCP2     MVGFKATDVPPTAAVRFLGAGTAACIADLITFPLDTAKVRLQIQGERRGPVQAAASAQYR
Sus scrofa UCP2            MVGFKATEVPPTATVKFLGAGTAACIADLITFPLDTAKVRLQIQGERRGPVQAAASAQYR
                           *******:*****:*.****************************** .***.*:******

Homo sapiens UCP2          GVMGTILTMVRTEGPRSLYNGLVAGLQRQMSFASVRIGLYDSVKQFYTKGSEHASIGSRL
Catagonus wagneri UCP2     GVLGTILTMVRNEGPRSLYNGLVAGLQRQMSFASVRIGLYDSVKQLYTKGSEHAGIGSRL
Sus scrofa UCP2            GVLGTILTMVRNEGPRSLYNGLVAGLQRQMSFASVRIGLYDSVKHFYTKGSEHAGIGSRL
                           **:********.**********************************::********.*****

Homo sapiens UCP2          LAGSTTGALAVAVAQPTDVVKVRFQAQARAGGGRRYQSTVNAYKTIAREEGFRGLWKGTS
Catagonus wagneri UCP2     LAGSTTGALAVAVAQPTDVVKVRFQAQARAAGGRRYQSTVDAYKTIAREEGFRGLWKGTS
Sus scrofa UCP2            LAGSTTGALAVAVAQPTDVVKVRFQAQARAGGGRRYRSTVDAYKTIAREEGLRGLWKGTS
                           ****************************.*****.***:**********:********

Homo sapiens UCP2          PNVARNAIVNCAELVTYDLIKDALLKANLMTDDLPCHFTSAFGAGFCTTVIASPVDVVKT
Catagonus wagneri UCP2     PNIARNAIVNCAELVTYDLIKDTLLKANLMTDDLPCHFTSAFGAGFCATVIASPVDVVKT
Sus scrofa UCP2            PNVARNAIVNCAELVTYDLIKDTLLKADLMTDDLPCHFTSAFGAGFCTTVIASPVDVVKT
                           **:********************:****:*****************:************

Homo sapiens UCP2          RYMNSALGQYSSAGHCALTMLQKEGPRAFYKGFMPSFLRLGSWNVVMFVTYEQLKRALMA
Catagonus wagneri UCP2     RYMNSALGRYSSAGHCALTMLREEGPRAFYKGFTPSFLRLGSWNVVMFVTYEQLKRALTA
Sus scrofa UCP2            RYMNSAPGQYSSAGHCALTMLQKEGPRAFYKGFTPSFLRLGSWNVVMFVTYEQLKRALMA
                           ****** *.************.:********** ********************** *

Homo sapiens UCP2          ACTSREAPF
Catagonus wagneri UCP2     ACASREAPF
Sus scrofa UCP2            ARASREAPF
                           * :******
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Fig S1. A) Amino acid alignment of human (accession number: NM_003356.4), peccary (PVHT010002981.1) and pig (accession number: AF095744.1) translated UCP3 nucleotide sequences. B) Amino acid alignment of human (accession number: U82819.1), peccary (accession number: PVHT010002981.1) and pig (accession number: AF036757.1) translated UCP2 nucleotide sequences.

Exon 3

```
                    1         10        20        30        40        50        60
                    |...|....|....|....|....|....|....|....|....|....|....|....|
Catagonus wagneri   agCAACAGCTAGTTTAGGAAGCAAGGTCTTAGCAGGCCTAGCGACT--------------
Homo sapiens        agCAGCACCTAGTTTAGGAAGCAAGATTTTAGCTGGTCTAACGACTGGAGGAGTGGCAGT

                    61        70        80        90        100       110       120
                    |...|....|....|....|....|....|....|....|....|....|....|....|
Catagonus wagneri   ----------------------------------------GTCCAGAGCCATCTACA
Homo sapiens        ATTCATTGGGCAACCCACAGAGGTCGTGAAAGTCAGACTTCAAGCACAGAGCCATCTCCA

                   121       130       140       150       160       170       180
                    |...|....|....|....|....|....|....|....|....|....|....|....|
Catagonus wagneri   TGGTCCCAGACCTTGCTACAT-GGGACTTACAATGCTTACAGAATTATAGCACCCACAGA
Homo sapiens        CGGAATCAAACCTCGCTACACGGGGACTTATAATGCGTACAGAATAATAGCAACAACCGA

                   181       190       200  205
                    |...|....|....|....|....|
Catagonus wagneri   AGGCTTAATGGGGCTTTGGAAAGgt
Homo sapiens        AGGCTTGACGGGTCTTTGGAAAGgt
```

Exon 4

```
                    1         10        20        30        40        50        60
                    |...|....|....|....|....|....|....|....|....|....|....|....|
Catagonus wagneri   agGGACTACTCCCAACCTGACAAGAAATGTCATCATCAGTTGTACAGAG-GAGTAACGTA
Homo sapiens        agGGACTACTCCCAATCTGATGAGAAGTGTCATCATCAATTGTACAGAGCTAGTAACATA

                    61        70        80        90        100   106
                    |...|....|....|....|....|....|....|....|.....|
Catagonus wagneri   TGACCTAATGAAGGAGGCCCTTGTGAAAAACAAATTATTAGCGGgt
Homo sapiens        TGATCTAATGAAGGAGGCCTTTGTGAAAAACAACATATTAGCAGgt
```

Exon 5

```
                    1         10        20        30        40        50        60
                    |...|....|....|....|....|....|....|....|....|....|....|....|
Catagonus wagneri   agATGATACGCCCTGCCACTTCGTGTCCGCTGTCATTGCTGGATTTTGCACAGTGGTTCT
Homo sapiens        agATGACGTCCCCTGCCACTTGGTGTCGGCTCTTATCGCTGGATTTTGCGCAACAGCTAT
                    61        70        80        90        100       110       120
                    |...|....|....|....|....|....|....|....|....|....|....|....|
Catagonus wagneri   GTCCTCTCCAGTGCGTGTGGTGAAGACCAGATCTGGTAACTGTCCACTGGGACAGTACCC
Homo sapiens        GTCCTCCCCGGTGGATGTAGTAAAAACCAGATTTATTAATTCTCCACCAGGACAGTACAA

                   121       130       140       150       160       170       180   186
                    |...|....|....|....|....|....|....|....|....|....|....|....|.....|
Catagonus wagneri   AAGTGTGCCCAACTGGCACAGTGACAATG-----CTAAGGAAGGACCACCAGCTCTTTTCAAAGGgt
Homo sapiens        AAGTGTGCCCAACT-GTGCAATGAAAGTGTTCACTAACGAAGGACCAACGGCTTTCTTCAAGGGgt
```

Exon 6

```
                    1         10        20        30        40        50        60

                    |...|....|....|....|....|....|....|....|....|....|....|....|
Catagonus wagneri   agATTTGTATCTTCCTTCCGGTGACTGGGATCCTGGAATGTCATCATGTTTGTGTGCTTT
Sus scrofa          agATTTGTACCTTCCTTCTGGTGACTGGAATCCTGGAACGTCATCCTGTTTGTGTGCTTT
Homo sapiens        agGTTGGTACCTTCCTTCTTGCGACTTGGATCCTGGAACGTCATTATGTTTGTGTGCTTT

                    61        70
                    |...|....|....|....|....|....|....|....|....|....|......|
Catagonus wagneri   GAACAGCTGAAAGGAGAACTGATGGAGTCGCGGCAGACTGTGAACTCTGCCACATAA
Sus scrofa          GAACAGCTGAAACAAGAGTTGATGGAGTCATGGCAGACTGTGGACTGTGCCGCATAA
Homo sapiens        GAACAACTGAAACGAGAACTGTCAAAGTCAAGGCAGACTATGGACTGTGCCACATAA
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Fig S2. Exon-by-exon alignment of the peccary (*Catagonus wagneri*) *UCP1* pseudogene versus the intact *UCP1* of humans (*Homo sapiens*). Exon 6 of the pig (*Sus scrofa*) was also included in the alignment. A 57 bp in-frame deletion is shown in yellow in the peccary. Highlights in pink represent frameshift deletions, while a frameshift insertion is displayed in cyan. A shared nonsense mutation between the peccary and pig in exon 6 is highlighted in red. Lower case letters represent GT-AG splice sites, all of which are intact.

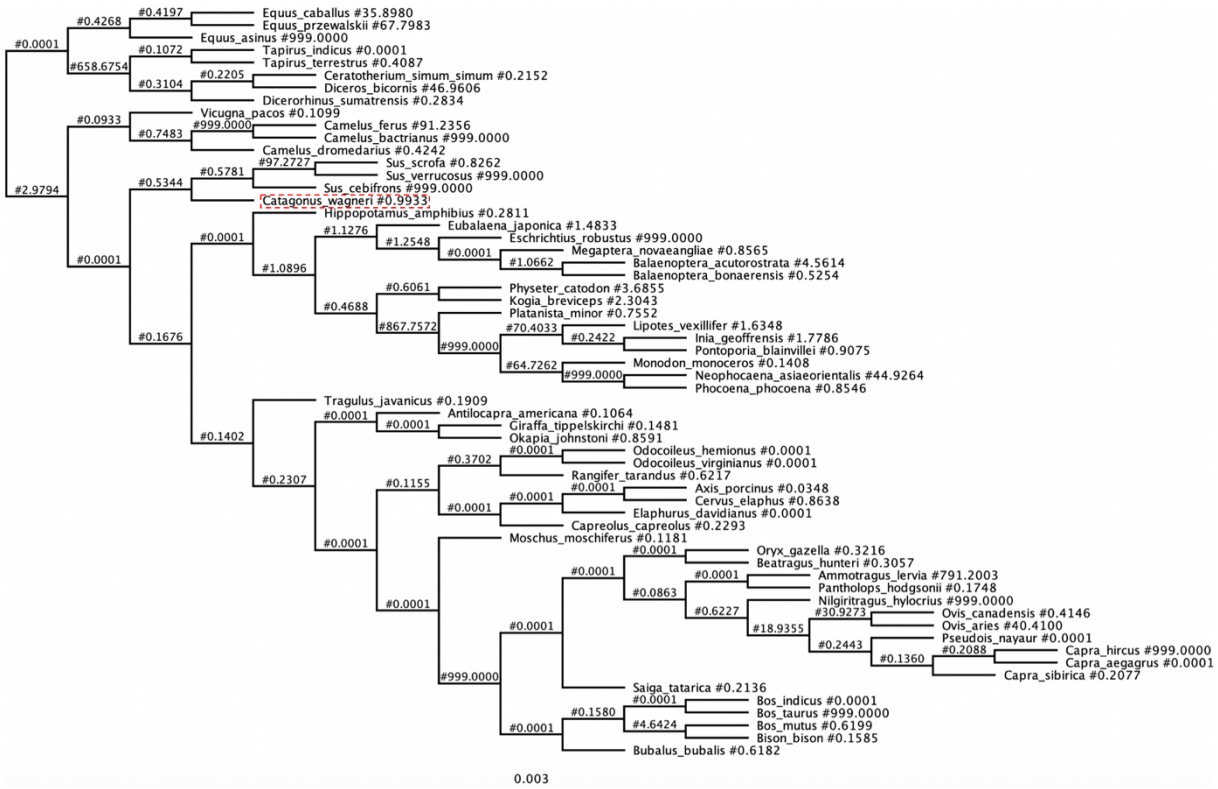

Fig S3. *UCP1* free ratio selection pressure model. *Catagonus wagneri* is highlighted with a red dashed box

Table S1. GenBank accession numbers of ungulate species used in this study.

Species name	UCP1 Source
Catagonus wagneri	PVHT010004494.1
Ammotragus lervia	NIVO01003382.1
Antilocapra americana	PVKT010017748.1
Axis porcinus	QQTR01093826.1
Balaenoptera acutorostrata	ATDI01065547.1
Balaenoptera bonaerensis	BAUQ01197845.1 BAUQ01341929.1 BAUQ01696173.1
Beatragus hunteri	PVKQ01021312.1 PVKQ01017688.1
Bison bison	JPYT01100523.1
Bos indicus	PRDE01000009.1
Bos mutus	AGSK01075302.1
Bos taurus	AAFC05027895.1
Bubalus bubalis	AWWX01630119.1
Camelus bactrianus	JARL01016741.1
Camelus dromedarius	JDVD01000357.1 JDVD01000358.1
Camelus ferus	AGVR01051296.1
Capra aegragus	CBYH010071014.1
Capra hircus	XM_018061376.1
Capra sibirica	NIYN02075614.1
Capreolus capreolus	CCMK010104759.1 CCMK012865005.1 CCMK010278719.1
Ceratotherium simum simum	AKZM01017598.1
Cervus elaphus	MKHE01000005.1
Dicerorhinus sumatrensis	PEKH010008988.1
Diceros diceris	PVJY010007317.1
Elaphurus davidianus	JRFZ01051564.1
Equus asinus	PSZQ01001512.1
Equus caballus	AAWR02018851.1
Equus przewalskii	ATBW01036322.1 ATBW01036321.1 ATBW01036320.1
Eschrichtius robustus	RJWN010007000.1
Eubalaena japonica	RJWP010009557.1
Giraffa tippelskirchi	LVKQ01071482.1
Hippopotamus amphibius	PVJP01000741.1
Inia geoffrensis	RJWO010061584.1
Kogia breviceps	RJWL010068022.1 RJWL010062101.1
Lipotes vexillifer	AUPI01000024.1 AUPI01000025.1
Megaptera noveangliae	RYZJ01001352.1
Monodon monoceros	PVJF01002816.1

Moschus moschiferus	PVHU010011962.1
Neophocaena asiaorientalis	MKKW01065165.1
Nilgiritragus hylocrius	PVJR01000372.1
Odocoileus hemionus	RCHL01005053.1
Odocoileus virginianus	MLBE01020605.1
Okapia johnstoni	LVCL010093660.1
Oryx gazella	RAWW01011422.1
Ovis aries	CBYI010017988.1
Ovis canadensis	PVIS010000157.1
Panthalops hodgsonii	AGTT01188813.1
Phocoena phocoena	PKGA01000564.1
Physeter catodon	AWZP01062081.1
Platanista minor	RJWK010001106.1
Pontoporia blainvillei	RJWI010022065.1 RJWI010380541.1
Pseudois nayaur	NIZD01207746.1
Rangifer tarandus	PVIN010004838.1
Saiga tatarica	PVIK010056808.1
Sus scrofa	LUXW01040214.1
Tapirus indicus	PVIE01006658.1
Tapirus terrestris	PVID01007152.1
Tragulus javanicus	PVHZ010004693.1
Vicugna pacos	JEMW01007827.1

Table S2. Localities of extinct *Platygonus compressus* and *Mylohyus spp*. fossil recoveries.

Platygonus compressus	
Coordinates	Reference
34.5° N, 110.1° W	**Murray, L. K., Bell, C. J., Dolan, M. T. and Mead, J. I.** (2005). Late Pleistocene fauna from the southern Colorado Plateau, Navajo County, Arizona. *Southwest. Nat.* **50**, 363-374.
29.1° N, 82.3° W; 29.6° N, 82.4° W	**Webb, S. D. and Wilkins, K. T.** (1984). Historical Biogeography of Florida Pleistocene Mammals. In *Contributions in quaternary vertebrate paleontology: a volume in memorial to John E. Guilday*. (ed. H. H. Genoways and M. R. Dawson), pp. 370-383. Pittsburgh, Pa.: Carnegie Museum of Natural History Special Publication.
29.8° N, 82.1° W; 29.1° N, 82.3° W; 28.7° N, 81.5° W	**Webb, S. D.** (1974). Chronology of Florida Pleistocene mammals. In *Pleistocene mammals of Florida* (ed. S. D. Webb), pp. 5-31. Gainesville: University of Florida Press.
29.7° N 82.6° W	**Martin, R. A.** (1978). A new Late Pleistocene Conepatus and associated vertebrate fauna from Florida. *J. Paleontol.* **52**, 1079-1085.
25.7° N, 80.4° W	**Emslie, S. D. and Morgan, G. S.** (1995). Taphonomy of a late Pleistocene carnivore den, Dade County, Florida. In *Late Quaternary Environments and Deep History: A Tribute to Paul S. Martin.* (ed. D. W. Steadman and J. I. Mead), pp. 65-83. The Mammoth site of Hot Springs, Hot Springs, South Dakota, Scientific Papers 3.
27.2° N, 81.9° W	**Hulbert, R. C., Morgan, G. S. and Kerner, A.** (2009). Collared peccary (Mammalia, Artiodactyla, Tayassuidae, Pecari) from the late Pleistocene of Florida. In *Papers on Geology, Vertebrate Paleontology, and Biostratigraphy in Honor of Michael O. Woodburne* (ed. L. B. Albright), vol. 65, pp. 531-544. Flagstaff, Az.: Museum of Northern Arizona Bulletin.
29.4° N, 82.5° W	**Martin, R. A. and Webb, S. D.** (1974). Late Pleistocene mammals from the Devil's Den Fauna, Levy County. In *Pleistocene Mammals of Florida.* (ed. S. D. Webb), pp. 114-145. Gainesville: University of Florida Press.

34.4° N, 84.9° W	**Martin, R. A. and Sneed, J. M.** (1989). Late Pleistocene records of caribou and elk from Georgia and Alabama. *Ga. J. Sci.* **47,** 117-122.
42.4° N, 90.4° W	**Leconte. J. L.** (1848). Notice of five new species of fossil Mammalia from Illinois. *Am. J. Sci. Arts.* **5,** 102-106.
42.4° N, 90.4° W	**Leidy. J.** (1860). Observations on the extinct peccary of North America; being a sequel to "A memoir on the extinct Dicotylinae of America". *Trans. Am. Philos. Soc.* **11,** 97-105.
37.0° N, 100.5° W	**Schultz, G. E.** (1969). Geology and paleontology of a late Pleistocene basin in Southwest Kansas. *Geol. Soc. Am., Special Paper.* **105,** 1-85.
37.2° N, 100.3° W	**Hibbard, C. W. and Taylor, D. W.** (1960). Two Late Pleistocene Faunas from Southwestern Kansas. *Contrib. Mus. Paleontol. Univ. Mich.* **16,** 1-223.
43.1° N, 85.2° W	**Finch, W.I., Whitmore, F.C. and Sims, J.D.** (1972). Stratigraphy, morphology, and paleoecology of a fossil peccary herd from western Kentucky. *U.S. Geological Survey Professional Paper* **790,** 25.
42.3° N, 83.7° W	**Holman, J. A. Fisher, D. C. and Kapp, R. O.** (1986). Recent discoveries of fossil vertebrates in the lower peninsula of Michigan. *Mich. Acad.* **18,** 431-463.
37.6° N, 91.8° W	**Schubert, B. W.** (2003). A late Pleistocene and early Holocene mammalian fauna from Little Beaver Cave, Central Ozarks, Missouri. I*n Ice Age Cave Faunas of North America* (ed. B. W. Schubert, J. I. Mead, and R. W. Graham), pp. 149-200. Bloomington: Indiana University Press.
36.7° N, 93.2° W	**Holman. J. A.** (1974). A late Pleistocene herpetofauna from southwestern Missouri. *J. Herpetol.* **8,** 343-346.
36.8° N, 107.6° W	**Lucas, S. G. and Smartt, R. A.** (1995). Late Pleistocene peccary from northwestern New Mexico. *Southwest. Nat.* **40,** 293-296.
41.4° N, 83.2° W	**Hoare, R. D., Coash, J.R., Innis, C. and Hole, T.** (1964). Pleistocene peccary Platygonus compressus LeConte from Sandusky County, Ohio. *Ohio J. Sci.* **64,** 207-214.
41.0° N, 83.5° W	**Tankersley, K. B.** (1997). Sheriden: a Clovis cave site in eastern North America. *Geoarcheaology*. **12,** 713-724.
30.6° N, 97.7° W	**Slaughter, B. H.** (1966). *Platygonus compressus* and associated fauna from the Laubach Cave of Texas. *Am. Midl. Nat.* **75,** 475-494.

32.7° N, 96.7° W	**Slaughter, B. H.** (1966). The Moore Pit local fauna; Pleistocene of Texas. *J. Paleontol.* **40,** 78-91.
33.1° N, 97.0° W	**Crook, W. W. and Harris, R. K.** (1958). A Pleistocene campsite near Lewisville, Texas. *Am. Antiquity* **23,** 233-246.
38.6° N, 79.4° W	**F. Grady.** (1984). A Pleistocene occurrence of Geomys (Rodentia: Geomyidae) in West Virginia. In *Contributions in Quaternary Vetebrate Paleontology. A volume in Memorial to John E. Guilday.* (ed. H.H. Genoways and M. R. Dawson), pp. 161-163 Pittsburgh, Pa.: Carnegie Museum of Natural History Special Publication.
36.9° N, 80.9° W	**Eshelman, R. and Grady, F.** (1986). Quaternary vertebrate localities of Virginia and their avian and mammalian fauna. In *The Quaternary of Virginia: a symposium volume* (ed. J. N. McDonald and S. O. Bird), pp. 43-71. Virginia Div. Mineral Resources Publ. 75.
39.2° N, 78.1° W	**Ott, J. R. and Weems, R. E.** (1986). The Hot Run Site; a new Pleistocene vertebrate locality in northern Virginia. In *The Quaternary of Virginia; a symposium volume* (ed. J. N. McDonald), pp. 129-130. Virginia Division of Mineral Resources.
38.6° N, 78.5° W	**Hubbard, D.A. and Grady, F.** (1997). Vertebrate paleontological cave resources in Virginia, USA. *Proceedings of the 12th International Congress of Speleology* **3,** 175-177.
38.6° N, 79.4° W	**Grady, F.** (1991). Fossil Chiroptera From Two West Virginia Caves. *The National Speleological Society Bulletin.* **53,** 122.
38.7° N, 79.3° W	**Frazier, M. K.** (1977). New Records of *Neofiber leonardi* (Rodentia: Cricetidae) and the Paleoecology of the Genus. *J. Mammal.* **58,** 368-373.
37.5° N, 80.4° W	**Grady, F.** (1988). A preliminary account of the Pleistocene mammals from Patton Cave, Monroe County, West Virginia. *Bull. Natl. Speleol. Soc.* **50,** 9-16.
19.9° N, 99.1° W	**Cope, E. D.** (1884). The extinct Mammalia of the Valley of Mexico. *Proc. Amer. Phil. Soc*. **xxii,** 10-11
67.83° N, 139.81° W	**Beebe, B. F.** (1980). Pleistocene peccary, *Platygonus compresses* Le Conte, from Yukon Territory, Canada. *Can. J. Earth Sc*i. **17,** 1204-1209.
40.58° N, 83.27° W	**Perry, T., van Loenen, A. L., Heiniger, H., Lee, C., Gongora, J., Cooper, A. and Mitchell, K. J.** (2017). Ancient DNA analysis of the extinct North American flat-headed peccary (*Platygonus compresses*). *Mol. Phylogenet. Evol*. **112,** 258-267.
34.7° N, 97.5° W; 35.5° N, 97.8° W	**Czaplewski, N.** (2012). Pleistocene peccaries (Mammalia: Tayassuidae) from Western Oklahoma. *Southwest. Nat*. **57,** 112-117.

Mylohyus spp.	
34.7° N, 87.9° W	**Lively, R. S., Bell, G. L. and Lamb, J. P.** (1992). Uranium-series dates from travertines associated with a late Pleistocene megafauna in ACb-3, Alabama. *Southeast. Geol.* **33,** 1-8.
34.7° N, 87.8° W	**Womochel, D. R. and Barnett, W. S.** (1980). The Pleistocene vertebrate assemblage of Little Bear Cave, Colbert County, Alabama. In Field trips for the Southeastern Section of the Geological Society of America, pp. 69-76.
36.0° N, 93.2° W; 40.2° N, 75.3° W	**Gidley, J. W. and Gazin, C. L.** (1938). The Pleistocene vertebrate fauna from Cumberland Cave, Maryland. *Bull. U.S. Natl. Mus.* **171,** 1-99.
28.8° N, 82.1° W	**Martin, R. A.** (1974). Fossil mammals from the Coleman IIA fauna, Sumter County. In *Pleistocene Mammals of Florida* (ed. S. D. Webb), pp. 35-99. University Presses of Florida.
27.3° N, 81.3° W; 29.7° N, 82.6° W	**Morgan, G. S. and Hulbert, R. C.** (1995). Overview of the geology and vertebrate biochronology of the Leisey Shell Pit Local Fauna, Hillsborough County, Florida. In Paleontology and geology of the Leisey Shell Pits, early Pleistocene of Florida (ed. R. C. Hulbert, G. S. Morgan, and S. D. Webb). pp. 1-92. Bull. Fla. Mus. Nat. Hist. 37.
27.7° N, 82.5° W	**Hulbert, R. C. and Morgan, G. S.** (1989). Stratigraphy, paleoecology, and vertebrate fauna of the Leisey Shell Pit Local Fauna, early Pleistocene (Irvingtonian) of southwestern Florida. *Papers in Florida Paleontology.* **2,** 1-19
27.9° N, 81.8° W; 29.7° N, 82.6° W; 29.8° N, 82.1° W; 28.8° N, 82.2° W; 29.4° N, 82.6° W; 28.8° N, 82.2° W	**Webb, S. D.** (1974). Chronology of Florida Pleistocene mammals. In Pleistocene mammals of Florida (ed. S. D. Webb), pp. 5-31. Gainesville: University of Florida Press.
29.1° N, 82.3° W	**Webb, S. D. and Wilkins, K. T.** (1984). Historical biogeography of Florida Pleistocene mammals. *Carnegie Mus. Nat. Hist. Spec. Publ.* **8,** 370-383.
27.6° N, 80.4° W	**Shufeldt, R. W.** (1917). Fossil birds found at Vero, Florida. *Florida State Geological Survey Annual Report* **9,** 35-42.
28.1° N, 80.6° W	**Allen, G. M.** (1932). A Pleistocene bat from Florida. *J. Mammal.* **13,** 256-259.
27.8° N, 82.7° W	**Simpson, G. G.** (1929). Pleistocene mammalian fauna of the Seminole Field, Pinellas County, Florida. *Bull. Am. Nat. Hist.* **56,** 561-599.

26.7° N, 80.2° W	**Becker, J. J.** (1985). A late Pleistocene (Wisconsinan) avifauna from West Palm Beach, Florida. *Bull. Br. Ornithol. Club* **105,** 37-40.
32.0° N, 81.1° W	**Hulbert, R. C. and Pratt, A. E.** (1998). Pleistocene (Rancholabrean) vertebrate faunas from coastal Georgia. *J. Vertebr. Paleontol.* **18,** 412-429.
38.3° N, 97.6° W	**Semken, H. A.** (1966). Stratigraphy and Paleontology of the McPherson Equus Beds (Sandahl Local Fauna), McPherson County, Kansas. *Contrib. Mus. Paleontol. Univ. Mich.* **20,** 121-178.
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