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Ofspring sex ratios are male‑biased refecting sex‑biased dispersal in Idaho, USA, wolves

David E. Ausband¹

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Abstract

Ofspring sex ratios can vary widely across species, and the reasons for such variation have long intrigued ecologists. For group-living animals, predicting ofspring sex ratios as a function of group and environmental characteristics can be challenging. Additionally, mortality of group members can upend traditional theory used to explain ofspring sex ratios observed in populations. Gray wolves (*Canis lupus*) in Idaho, USA, are an excellent study species for asking questions about ofspring sex ratios given their group-living behavior and persistent exposure to human-caused mortality. I hypothesized that ofspring sex ratios would be infuenced by the characteristics of individuals, groups, and populations. I generated genotypes for 419 adult and 400 pup wolves during 2008–2018. There was a signifcant male-bias in litters of wolf pups with nearly 12% more male pups born than females. The individual, group, and population variables I considered did not have signifcant associations with ofspring sex ratios. Local resource competition helped explain ofspring sex ratios in wolves in my study system, but not local resource enhancement theory. Although female helpers have been shown to help slightly more than males, ofspring sex ratios did not favor the helping sex suggesting that the overall beneft of female helpers may have been negligible in wolf groups during my study. Three wolf groups consistently overproduced males, the dispersing sex, suggesting that habitat quality was poor in their territories. The male-biased ofspring sex ratios observed throughout this population refect sex-biased dispersal in wolves in Idaho. Such a pattern suggests breeding females may be reducing local resource competition (e.g., mates and successful reproduction) by producing more males than females.

Signifcance statement

Natural selection can favor biased ofspring sex ratios in some species. This may be particularly true for animals that live and breed in groups such as gray wolves. Using genetic sampling, I show that ofspring sex ratios in wolves are male-biased and refect sex-biased dispersal in wolves. Breeding females may be reducing future local resource competition for mates by producing signifcantly more ofspring of the dispersing sex (males).

Keywords *Canis lupus* · Gray wolf · Litter · Mating · Reproduction · Sex ratio · Wolf pup

Introduction

The ratio of males to females in a population can greatly afect population growth and demography (Mills [2013](#page-6-0)). Ofspring sex ratios can vary widely across species, and

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 \boxtimes David E. Ausband dausband@uidaho.edu the reasons for such variation have long intrigued ecologists (Fisher [1930](#page-6-1); Hamilton [1967;](#page-6-2) Silk and Brown [2008](#page-6-3)). In many species, deviations from equal offspring sex ratios can be expected to return to 50:50 rather quickly as selection favors the rarer sex over the more common one (Fisher [1930\)](#page-6-1). In group-living species where individuals within groups help provision and rear ofspring, deviations from equal sex ratios may be the norm, however. For example, in groups with few members of the helping sex, breeding females may disproportionally produce more of the helping sex during reproduction, an example of local resource enhancement (Creel et al. [1998\)](#page-6-4). Local resource enhancement predicts breeders will produce more of the sex that

¹ U.S. Geological Survey, Idaho Cooperative Fish and Wildlife Research Unit, University of Idaho, 975 West 6th Street, ID 83844 Moscow, USA

helps within groups. Philopatry and local resource competition can also skew offspring sex ratios because selection may act against overproducing the sex that stays near the parent and competes for resources (e.g., mates, food). Indeed, Silk and Brown [\(2008\)](#page-6-3) showed that offspring sex ratios in primates favored the dispersing sex, an example of local resource competition where evolution favors the production of the sex that disperses.

Individuals can alter their ofspring sex ratios and could conceivably do so as a function of their genetic composition. For example, a female with low genetic heterozygosity may produce more male ofspring because her daughters will likely breed regardless, although Li et al. [\(2016\)](#page-6-5) did not fnd this to be true in black-throated tits (*Aegithalos concinnus*). Additionally, when there is breeder turnover with a group, females may produce more male ofspring to reduce the likelihood of future polygamy and having to share breeding in the group. In addition to the effects of individual decisions and behaviors, offspring sex ratios may vary with population density, an index for competition in a population. If population density is low and resources abundant, females may choose to produce more males if such males are more likely to mate given a larger size or improved body condi-tion (Trivers and Willard [1973\)](#page-6-6). Indeed, offspring sex ratios were male-biased in cooperatively breeding gray wolf (*Canis lupus*) populations at high densities compared to those at low densities (Mech [1975;](#page-6-7) Sidorovich et al. [2007](#page-6-8)).

For animals that live in groups, predicting offspring sex ratios as a function of group characteristics and environmental conditions can be challenging because relevant data are often lacking. Additionally, mortality of group members can upend traditional theory used to explain offspring sex ratios observed in populations. For example, local resource competition theory may predict producing fewer philopatric sex. If mortality disproportionately affects the philopatric sex, however, selection may work to favor offspring sex ratios where the philopatric sex is not underproduced. Human-caused mortality, in particular, can have large effects on the selective advantage of ofspring sex ratios in wild populations (Holand et al. [2006\)](#page-6-9). For many species, however, we often do not have ofspring sex ratio data over long-time periods. This may be particularly true for carnivores that exist at low densities and are subject to frequent human-caused mortality making data collection difficult.

Gray wolves in Idaho, USA, are an excellent study species for asking questions about ofspring sex ratios given their group-living behavior and persistent exposure to humancaused mortality in the form of hunting and trapping. I hypothesized that offspring sex ratios would be influenced by the characteristics of individuals, groups, and populations. Specifcally, I predicted breeding female heterozygosity, and both male and female breeder turnovers would be associated with the production of more male offspring (Table [1](#page-2-0)). In contrast, I expected population density, harvest rate, litter size, number of helpers (i.e., nonbreeding adults) in the present year, number of adults (i.e., nonbreeding adults+breeding adults) in the previous year, pups recruited, sex ratio of helpers in the current year, and sex ratio of pups in the previous year would be associated with the production of fewer male ofspring (Table [1](#page-2-0)).

Study area

I genetically sampled groups of wolves in three study areas (north, east, and south) in Idaho, USA, from 2008 to 2018 (Fig. [1](#page-3-0)). Annual temperatures ranged from−13 to 36 °C (Western Regional Climate Center [2022\)](#page-6-10), precipitation ranged from 30 to 130 cm (Western Regional Climate Center [2022\)](#page-6-10), and elevation ranged from 646 to 3219 m. The northern study area $(3189 \text{ km}^2, \text{Fig. 1})$ was comprised of forests of western red cedar (*Thuja plicata*), Douglas fr (*Pseudotsuga menziesii*), Engelmann spruce (*Picea engelmannii*), and lodgepole pine (*Pinus contorta*). The eastern $(3388 \text{ km}^2, \text{Fig. 1})$ and southern $(3861 \text{ km}^2, \text{Fig. 1})$ study areas were comprised largely of ponderosa pine (*Pinus ponderosa*), lodgepole pine, spruce mixed forests, and sagebrush (*Artemisia tridentata*) steppe. Wolves were harvested beginning in 2009 and annually each year thereafter with a brief cessation in 2010. Most wolves were harvested during September–March (Ausband [2016\)](#page-5-0). Harvest averaged 18.4% over the study period and ranged from 10.4 to 27.6% of the population annually (Ausband [2021\)](#page-5-1).

Methods

Field methods

Our behavioral inferences are made indirectly from genetic data collected in the feld. It was not possible to record data blind because our study involved focal animals in the feld. We collected wolf scats at wolf pup-rearing sites during summer (average sampling date, July 15). We located groups of wolves by surveying sites predicted by a puprearing habitat model (Ausband et al. [2010\)](#page-6-11). When available, we used the Global Positioning System (GPS) or radio-telemetry locations of wolves that were radiocollared as part of population monitoring efforts by the U.S. Fish and Wildlife Service, Idaho Department of Fish and Game, and the Nez Perce Tribe. At potential pup-rearing sites, we frst gave a series of howls (Harrington and Mech [1982](#page-6-12)) and then attempted to fnd an activity center (area where pups congregate) where fecal samples would be most abundant. Once an activity center was found we radiated out 500 m to sample additional scats (Jacobs and Ausband [2019](#page-6-13)). We attempted to resample each group every year from 2008 to

2018. We collected 125–200 samples per group per year, which generally detected every individual in the group (Stenglein et al. [2011\)](#page-6-14). Sampling was conducted under the University of Montana IACUC (Animal Use Protocol 008-09MMMCWRU).

Laboratory methods

We performed DNA analyses at the University of Idaho's Laboratory for Ecological, Evolutionary and Conservation Genetics (Moscow, ID, USA). We initially screened all samples in a species-identifcation test using mitochondrial DNA to remove non-target species and low-quality samples. We then attempted to genotype all remaining samples using 18 nuclear DNA microsatellite loci (AHT103, AHT109, AHT121, AHT200, C05.377, C09.173, C37.172, Cxx.119, Cxx.250, FH2001, FH2004, FH2010, FH2054, FH2088, FH2137, FH2611, FH2670, FH3725: Holmes et al. [1994](#page-6-15); Breen et al. [2001](#page-6-16); Guyon et al. [2003](#page-6-17); Salim et al. [2007](#page-6-18); Ostrander et al. [2017\)](#page-6-19). Further details regarding laboratory methods can be found in Stenglein et al. ([2010a,](#page-6-20) [b](#page-6-21), [2011](#page-6-14)) and Stansbury et al. ([2014\)](#page-6-22). We analyzed all collected samples in 2008 and 2009. After 2010, we used rarefaction analyses (Stenglein et al. 2011) to subsample for economic efficiency and analyzed 40 adult and 25 pup samples from each group. If a group had more than two individuals detected only once, we analyzed additional samples when available to obtain 10 more consensus genotypes.

Analytical methods

In addition to estimating observed heterozygosity from the resulting genotypes using Program Genalex (Peakall and Smouse [2006](#page-6-23)) as well as group size, I determined maternity and paternity from pedigree analyses using Program COL-ONY, version 2.0.5.5 (Jones and Wang [2009\)](#page-6-24). I included all sampled adult males and females as potential parents and all sampled pups as potential offspring for each year. I frst calculated allele frequencies for a year in Program COANCESTRY version 1.0.1.5 (Wang [2011\)](#page-6-25) and then imported those into Program COLONY for use in pedigree analyses. I allowed polygamy in both sexes and assumed an allelic dropout rate of 0.01. Tracking individuals over time provided information on breeder turnover, number of helpers, harvest rate, litter size, and population density in my study areas.

Fig. 1 Three study areas in Idaho, USA, where wolves were genetically sampled, 2008–2018

I used a chi-square test to frst assess whether sex ratios of all wolf pup litters across all years were male-biased. I also tested for associations between the sex ratios of pups in each litter in each wolf group and several potential predictor variables (Table [1](#page-2-0)) using both generalized linear mixed efects and fixed effects models. I also tested for non-linear relationships by squaring (variable²) several predictor variables in Table [1,](#page-2-0) namely, population density, harvest rate, number of helpers in a group, litter size, and number of adults in a group. I frst standardized all variables using a Z-transformation for ease when comparing the resulting β coefficients. Finally, I compared competing models using Akaike's information criterion

(AIC) and conducted analyses in Program R, version 4.0.4, with package "lme4"| (R Development Core Team [2021\)](#page-6-26).

Results

I generated genotypes for 419 adult and 400 pup wolves across 80 wolf-group years. There was a signifcant male-bias in litters of wolf pups (χ^2 =5.3, df=1, *P*=0.02) with nearly 12% more male pups born than females during the 11 years (55.8%, *n*=223 vs. 44.3%, *n*=177, respectively; Fig. [2](#page-4-0)). There was just 1 year (2013) where females greatly exceeded male

ofspring numbers (proportions, 0.63 vs 0.38, respectively; Fig. [2](#page-4-0)), although I suspect this was due to sampling error $(n=16$ pups total). Offspring sex ratios (M:F) did not appear to vary by study area $(0.59, SD=0.25; 0.55, SD=0.17;$ and 0.53 , $SD = 0.26$; east, north, and south, respectively; Fig. [1](#page-3-0)). Three wolf groups (Bumblebee, Hoodoo, Timberline II) with various breeders overproduced males consistently (\bar{x} > 0.60) and never had offspring sex ratios < 0.50 .

Mixed effects models would not converge. A fixed effects global model with all variables (Table [2](#page-4-1)) except squared terms had the lowest AIC (87.97) and was nearly 10 AIC points lower than a model that included squared terms $(AIC=97.2)$. The top global model was nearly 25 AIC points lower than the null $(AIC = 112.6)$. None of the individual, group, and population variables I considered had signifcant associations with the sex ratio of wolf pup litters (Table [2\)](#page-4-1).

Discussion

Although not an a priori hypothesis, I found support that local resource competition could be used to help explain ofspring sex ratios in wolves and I suspect that the local competition is primarily for mates. Male-biased offspring sex ratios refect sex-biased dispersal in wolves and suggest breeding females may be reducing future local resource competition (e.g., for mates; Hamilton [1967\)](#page-6-2) by producing more males than females without increasing overall litter size. Male wolves in my study population seldom inherit a breeding position within their group whereas females commonly do (Ausband [2022a](#page-5-2), [b\)](#page-5-3). Male wolves generally disperse to fnd breeding opportunities and there is a strong

Table 2 Covariates value and signifcance from most supported model predicting the sex ratios of litters of wolf pups (M to F) in Idaho, USA, 2008–2018

Variable	β	SE	P
Breeding female heterozygosity (H_0)	0.14	0.35	0.70
Breeding male turnover between t-1 and t	-0.02	0.27	0.93
Breeding female turnover between t-1 and t	-0.05	0.33	0.87
Density (wolves/1000 km^2), t-1	0.11	0.38	0.77
Harvest rate, t-1	0.05	0.39	0.90
Litter size, t	0.06	0.33	0.86
No. of helpers, t	0.11	0.59	0.85
No. of adults, t-1	0.05	0.47	0.92
Pups recruited between t-1 and t	0.12	0.44	0.79
Sex ratio (M to F) helpers, t	0.04	0.48	0.94
Sex ratio (M to F) pups, $t-1$	-0.04	0.44	0.92

male-bias in dispersal rates for wolves in my study area (Jimenez et al. [2017](#page-6-27)). Dispersal is not always male-biased in wolf populations, however (Morales-Gonzalez et al. [2021](#page-6-28)). Local resource competition theory predicts male-biased offspring sex ratios when dispersal is male-biased, which is precisely what I found. Breeding females also appear to monopolize breeding opportunities in groups of wolves as evidenced by increases in polygamy following breeding female turnover in my study system (Ausband et al. [2017](#page-6-29)). Selection via male-biased offspring sex ratios may therefore favor producing fewer philopatric females who compete for future mates and subsequent successful reproduction.

I found no evidence that local resource enhancement could be used to predict offspring sex ratios in wolves. Although female

helpers have been shown to help slightly more than males in wolf groups (Ausband et al. [2016](#page-5-4)), offspring sex ratios did not favor the helping sex (females) suggesting that the overall beneft of female helpers may have been negligible in wolf groups during the period of my study. This is not to say that some individual females within a group are not beneficial helpers, but there is little evidence that all females are benefcial helpers. In contrast, McNutt and Silk [\(2008\)](#page-6-30) showed a strong male-biased sex ratio in groups of African wild dogs (*Lycaon pictus*) where males are known to help more and remain in groups longer than females. The sex ratio of harvested wolves is also strongly male-biased (55 M:45 F) early in the hunting season when most harvest is from rife hunters (Ausband [2016](#page-5-0)). The ratio of wolves killed by rife hunting nearly mirrors the male-biased estimates of ofspring sex ratios. By the end of the harvest season, however, males were only slightly more commonly killed than females. In sum, it appears that breeding females are not overproducing the helping sex (i.e., females).

I considered a suite of individual, group, and population phenomena that could have potentially influenced offspring sex ratios but ultimately did not. If offspring sex ratios are largely driven by phenomena such as competition for mates or local habitat quality, then few of my explanatory variables would be expected to have predictive power with the exception of perhaps density (and group size and harvest indirectly). Although previous studies on canids found high density to predict male-biased ofspring sex ratios and male-biased group composition (Mech [1975](#page-6-7); Sidorovich et al. [2007](#page-6-8); Iossa et al. [2009\)](#page-6-31), I did not fnd an efect of density despite observing male-biased sex ratios in wolves. The overarching literature contains several hypotheses that might lead us to suspect skewed ofspring sex ratios in cooperative breeders like wolves (Hamilton [1967](#page-6-2); Creel et al. [1998](#page-6-4); Silk and Brown [2008\)](#page-6-3). Julliard [\(2000](#page-6-32)) showed that breeders may adjust ofspring sex ratios as a function of local habitat quality. I did not measure habitat quality, but three wolf groups showed consistent patterns of overproducing males (e.g., Hoodoo wolf group, M to F ratio, $\bar{x} = 0.70$ and never < 0.50, 2009–2016). Following predictions of Julliard's model (2000), some wolf groups may have occupied poor quality territories and thus produced more of the dispersing sex. The Hoodoo group in particular vacated their territory and began to occupy a new territory beginning in 2016. Despite ongoing annual sampling to monitor them, they have not returned to their prior territory suggesting that it may have indeed declined in quality.

Many population and genetic projection models assume an equal sex ratio in wolves (vonHoldt et al. [2008](#page-6-25); Miller [2017](#page-6-33)). Similar to other studies (Mech [1975](#page-6-7); Sidorovich et al. [2007](#page-6-8)), I show that this assumption is not always valid and may depend on population density. Allowing for a male-biased sex ratio of ofspring could greatly afect projection models given that male wolves appear to be the couriers of genetic diversity in wolf populations (Ausband [2022a](#page-5-2)). Additionally, a male-biased sex ratio could conceivably afect population projection models as fewer females would reach reproductive age.

We sampled wolves genetically using scats in summer, and some pups may have died prior to our sampling. Aside from starvation or abandonment in the den, pup mortality early in life is frequently related to disease outbreaks that can result in the entire loss of litters or nearly so (Mech et al. [2008](#page-6-34); Almberg et al. [2010](#page-5-5)). I did not observe such disease outbreaks during my study and have no reason to suspect any form of early season mortality would afect one sex more than the other. Lastly, I note that we seldom found dead wolf pups during sampling at pup-rearing sites (<1 pup per year over 11 years).

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Data availability Ausband DE (2022b) Wolf pup sex ratios and covariates, Idaho, USA, 2008–2018 [Data set]. Zenodo, [https://doi.org/10.](https://doi.org/10.5281/zenodo.7035926) [5281/zenodo.7035926](https://doi.org/10.5281/zenodo.7035926)

Declarations

Ethics approval Field sampling was conducted under the University of Montana IACUC (Animal Use Protocol 008-09MMMCWRU). All applicable international, national, and/or institutional guidelines for the use of animals were followed.

Conflict of interest The author declares no competing interests.

References

- Almberg ES, Cross PC, Smith DW (2010) Persistence of canine distemper virus in the Greater Yellowstone Ecosystem's carnivore community. Ecol Appl 20:2058–2074
- Ausband DE (2016) Gray wolf harvest in Idaho. Wildlife Soc B 40:500–505
- Ausband DE (2021) Mate selection data for gray wolves. Zenodo. <https://doi.org/10.5281/zenodo.5764792>
- Ausband DE (2022a) Inherit the kingdom or storm the castle? Breeding strategies in gray wolves. Ethology 128:152–158
- Ausband DE (2022b) Wolf pup sex ratios and covariates, Idaho, USA 2008–2018. Zenodo. <https://doi.org/10.5281/zenodo.7035926>
- Ausband DE, Mitchell MS, Bassing SB, Morehouse A, Smith DW, Stahler DR, Struthers J (2016) Individual, group, and

environmental infuences on helping behavior in a social carnivore. Ethology 122:963–972

- Ausband DE, Mitchell MS, Doherty K, Zager P, Mack CM, Holyan J (2010) Surveying predicted rendezvous sites to monitor gray wolf populations. J Wildlife Manage 74:1043–1049
- Ausband DE, Mitchell MS, Waits L (2017) Efects of breeder turnover and harvest on group composition and recruitment in a social carnivore. J Anim Ecol 86:1094–1011
- Breen M, Jouquand S, Renier C et al (2001) Chromosome-specifc single-locus FISH probes allow anchorage of an 1800-marker integrated radiation-hybrid/linkage map of the domestic dog genome to all chromosomes. Genome Res 11:1784–1795
- Creel S, Marusha Creel N, Monfort SL (1998) Birth order, estrogens and sex-ratio adaptation in African wild dogs (*Lycaon pictus*). Anim Reprod Sci 53:315–320
- Fisher RA (1930) The genetical theory of natural selection. Clarendon Press, Oxford
- Guyon R, Lorentzen TD, Hitte C et al (2003) A 1-Mb resolution radiation hybrid map of the canine genome. Proc Natl Acad Sci USA 100:5296–5301
- Hamilton WD (1967) Extraordinary sex ratios. Science 156:477–488
- Harrington FH, Mech LD (1982) An analysis of howling response parameters useful for wolf pack censusing. J Wildlife Manage 46:686–693
- Holand Ø, Mysterud A, Røed KH, Coulson T, Gjøstein H, Weladji RB, Nieminen M (2006) Adaptive adjustment of offspring sex ratio and maternal reproductive effort in an iteroparous mammal. Proc R Soc Lond B 273:293–299
- Holmes NG, Strange NJ, Binns MM, Mellersh CS, Sampson J (1994) Three polymorphic canine microsatellites. Anim Genet 25:200
- Iossa G, Soulsbury CD, Baker PJ, Edwards KJ, Harris S (2009) Behavioral changes associated with a population density decline in the facultatively social red fox. Behav Ecol 20:385–395
- Jacobs C, Ausband DE (2019) Wolves in space: locations of individuals and their efect on pup survival in groups of a cooperatively breeding canid. Anim Behav 155:189–197
- Jimenez MD, Bangs EE, Boyd DK, Smith DW, Becker SA, Ausband DE, Woodruf SP, Bradley EH, Holyan J, Laudon K (2017) Wolf dispersal in the northern Rocky Mountains, Western United States: 1993–2008. J Wildlife Manage 81:581–592
- Jones O, Wang J (2009) COLONY: a program for parentage and sibship inference from multilocus genotype data. Mol Ecol Resour 10:551–555
- Julliard R (2000) Sex-specifc dispersal in spatially varying environments leads to habitat-dependent evolutionary stable ofspring sex ratios. Behav Ecol 11:421–428
- Kreeger TJ (2003) The internal wolf: Physiology, pathology, and pharmacology. In: Mech LD, Boitani L (eds) Wolves: ecology, conservation, and management. The University of Chicago Press, IL, USA, pp 192–217
- Li J, Wang Y, Lv L, Wang P, Zhang Z (2016) No facultative manipulation of ofspring sex ratio in relation to parental genetic characteristics in a bird with sex-specifc heterozygosity-ftness correlation. Behav Ecol Sociobiol 70:963–973
- McNutt JW, Silk JB (2008) Pup production, sex ratios, and survivorship in African wild dogs, *Lycaon pictus*. Behav Ecol Sociobiol 62:1061–1067
- Mech LD (1975) Disproportionate sex ratios of wolf pups. J Wildlife Manage 39:737–740
- Mech LD, Goyal SM, Paul WJ, Newton WE (2008) Demographic efects of canine parvovirus on a free-ranging wolf population over 30 years. J Wildlife Dis 44:824–836
- Miller P (2017) Population viability analysis for the Mexican wolf (*Canis lupus baileyi*) integrating wild and captive populations in

a metapopulation risk assessment model for recovery planning. In: Biological report for the Mexican wolf (*Canis lupus baileyi*) November 2017. U.S. Fish and Wildlife Service, Southwest Region (Region 2), Albuquerque, NM, USA, 72 p

- Mills LS (2013) Conservation of wildlife populations: demography, genetics, and management, 2nd edn. Wiley-Blackwell, Oxford
- Morales-Gonzalez A, Fernandez-Gil A, Quevedo M, Revilla E (2021) Patterns and determinants of dispersal in grey wolves (*Canis lupus*). Biol Rev 97:466–480
- Ostrander EA, Wayne RK, Freedman AH, Davis BW (2017) Demographic history, selection and functional diversity of the canine genome. Nat Rev Genet 18:705–720
- R Development Core Team (2021) R: a language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. [http://www.R-project.org.](http://www.R-project.org) Accessed 5 July 2022
- Peakall R, Smouse PE (2006) GENALEX 6: genetic analysis in Excel. Population genetic software for teaching and research. Mol Ecol 6:288–295
- Salim DC, Akimoto AA, Carvalho CB, Oliveir SF, Grisolia CK, Moreira JR, Klautau-Guimarães MN (2007) Genetic variability in maned wolf based on heterologous short-tandem repeat markers from domestic dog. Genet Mol Res 6:348–435
- Sidorovich VE, Stolyarov VP, Vorobei NN, Ivanova NV, Jedrzejewska B (2007) Litter size, sex ratio, and age structure of gray wolves, *Canis lupus*, in relation to population fuctuations in northern Belarus. Can J Zool 85:295–300
- Silk JB, Brown GR (2008) Local resource competition and local resource enhancement shape primate birth sex ratios. Proc R Soc Lond B 275:1760–1765
- Stansbury CS, Ausband DE, Zager P, Mack CM, Miller CR, Pennell MW, Waits LP (2014) A long-term population monitoring approach for a wide-ranging carnivore: noninvasive genetic sampling of gray wolf rendezvous sites in Idaho, USA. J Wildlife Manage 78:1040–1049
- Stenglein JL, De Barba M, Ausband DE, Waits LP (2010a) Impacts of sampling location within a faeces on DNA quality in two carnivore species. Mol Ecol Resour 10:109–114
- Stenglein JL, Waits LP, Ausband DE, Zager P, Mack CM (2010b) Efficient noninvasive genetic sampling for monitoring reintroduced wolves. J Wildlife Manage 74:1050–1058
- Stenglein JL, Waits LP, Ausband DE, Zager P, Mack C (2011) Estimating gray wolf pack size and family relationships using noninvasive genetic sampling at rendezvous sites. J Mammal 92:784–795
- Trivers RL, Willard DE (1973) Natural selection of parental ability to vary the sex ratio of ofspring. Science 179:90–92
- vonHoldt BM, Stahler DR, Smith DW, Earl DA, Pollinger JP, Wayne RK (2008) The genealogy and genetic viability of reintroduced Yellowstone grey wolves. Mol Ecol 17:252–274
- Wang J (2011) COANCESTRY: a program for simulating, estimating and analysing relatedness and inbreeding coefficients. Mol Ecol Resour 11:141–145
- Western Regional Climate Center (2022) Historical climate information. [http://www.wrcc.dri.edu.](http://www.wrcc.dri.edu) Accessed 11 Mar 2022

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