

# Susannah Woodruff

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5833316/publications.pdf>

Version: 2024-02-01

26  
papers

3,252  
citations

430874

18  
h-index

552781

26  
g-index

26  
all docs

26  
docs citations

26  
times ranked

3170  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Estimating the probability of identity among genotypes in natural populations: cautions and guidelines. <i>Molecular Ecology</i> , 2001, 10, 249-256.   | 3.9 | 1,101     |
| 2  | Noninvasive genetic sampling: look before you leap. <i>Trends in Ecology and Evolution</i> , 1999, 14, 323-327.   | 8.7 | 768       |
| 3  | NONINVASIVE GENETIC SAMPLING TOOLS FOR WILDLIFE BIOLOGISTS: A REVIEW OF APPLICATIONS AND RECOMMENDATIONS FOR ACCURATE DATA COLLECTION. <i>Journal of Wildlife Management</i> , 2005, 69, 1419-1433.                                       | 1.8 | 540       |
| 4  | The impact of time and field conditions on brown bear ( <i>Ursus arctos</i> ) faecal DNA amplification. <i>Conservation Genetics</i> , 2007, 8, 1219-1224.  | 1.5 | 128       |
| 5  | The influence of diet on faecal DNA amplification and sex identification in brown bears ( <i>Ursus arctos</i> ). <i>Journal of Wildlife Management</i> , 2009, 73, 1050-1058.   | 3.9 | 96        |
| 6  | Efficient, Noninvasive Genetic Sampling for Monitoring Reintroduced Wolves. <i>Journal of Wildlife Management</i> , 2010, 74, 1050-1058.  | 1.8 | 96        |
| 7  | Comparing opportunistic and systematic sampling methods for noninvasive genetic monitoring of a small translocated brown bear population. <i>Journal of Applied Ecology</i> , 2010, 47, 172-181.  | 4.0 | 75        |
| 8  | A long-term population monitoring approach for a wide-ranging carnivore: Noninvasive genetic sampling of gray wolf rendezvous sites in Idaho, USA. <i>Journal of Wildlife Management</i> , 2014, 78, 1040-1049.                           | 1.8 | 57        |
| 9  | Detection of Predator Presence at Elk Mortality Sites Using mtDNA Analysis of Hair and Scat Samples. <i>Wildlife Society Bulletin</i> , 2006, 34, 815-820.  | 1.6 | 50        |
| 10 | Noninvasive individual and species identification of jaguars ( <i>Panthera onca</i> ), pumas ( <i>Puma</i> ) using microsatellites and faecal DNA. <i>Molecular Ecology Resources</i> , 2014, 14, 1171-1182.                              | 4.8 | 48        |
| 11 | Monitoring coyote population dynamics with fecal DNA and spatial capture-recapture. <i>Journal of Wildlife Management</i> , 2016, 80, 824-836.  | 1.8 | 43        |
| 12 | Evaluating DNA degradation rates in faecal pellets of the endangered pygmy rabbit. <i>Molecular Ecology Resources</i> , 2013, 13, 654-662.  | 4.8 | 37        |
| 13 | Efficient single-survey estimation of carnivore density using fecal DNA and spatial capture-recapture: a bobcat case study. <i>Population Ecology</i> , 2018, 60, 197-209.  | 1.2 | 33        |
| 14 | Evaluating the interaction of faecal pellet deposition rates and DNA degradation rates to optimize sampling design for DNA-based capture-recapture analysis of Sonoran pronghorn. <i>Molecular Ecology Resources</i> , 2015, 15, 843-854. | 4.8 | 31        |
| 15 | Estimating abundance of American black bears using DNA-based capture-recapture models. <i>Ursus</i> , 2009, 20, 1-11.   | 0.5 | 28        |
| 16 | Advances in Using Non-invasive, Archival, and Environmental Samples for Population Genomic Studies. <i>Population Genomics</i> , 2018, , 63-99.   | 0.5 | 24        |
| 17 | Estimating Sonoran pronghorn abundance and survival with fecal DNA and capture-recapture methods. <i>Conservation Biology</i> , 2016, 30, 1102-1111.  | 4.7 | 22        |
| 18 | Rapid species identification of pygmy rabbits ( <i>Brachylagus idahoensis</i> ) from faecal pellet DNA. <i>Molecular Ecology Resources</i> , 2011, 11, 808-812.   | 4.8 | 18        |

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|----|--|-----|-----------|
| 19 | Examining the use of fecal pellet morphometry to differentiate age classes in Sonoran pronghorn. <i>Wildlife Biology</i> , 2016, 22, 217-227.  | 1.4 | 14        |
| 20 | Ten polymorphic microsatellite markers for pronghorn ( <i>Antilocapra americana</i> ). <i>Conservation Genetics Resources</i> , 2010, 2, 81-84.  | 0.8 | 12        |
| 21 | Rapid species identification of Sonoran pronghorn from fecal pellet DNA. <i>Wildlife Society Bulletin</i> , 2014, 38, 842-848.   | 1.6 | 9         |
| 22 | Winter predation patterns of wolves in Northwestern Wyoming. <i>Journal of Wildlife Management</i> , 2019, 83, 1352-1367.  | 1.8 | 5         |
| 23 | Consideration of sample source for establishing reliable genetic microsatellite data from mammalian carnivore specimens held in natural history collections. <i>Journal of Mammalogy</i> , 2019, 100, 1678-1689.               | 1.3 | 5         |
| 24 | Comparing performance of multiple non-invasive genetic capture-recapture methods for abundance estimation: a case study with the Sonoran pronghorn <i>Antilocapra americana sonoriensis</i> . <i>Oryx</i> , 2020, 54, 412-420. | 1.0 | 5         |
| 25 | Optimizing Selection of Brown Bear Hair for Noninvasive Genetic Analysis. <i>Wildlife Society Bulletin</i> , 2020, 44, 94-100.   | 1.6 | 4         |
| 26 | Estimating Coyote Densities with Local, Discrete Bayesian Capture-Recapture Models. <i>Journal of Wildlife Management</i> , 2021, 85, 73-86.   | 1.8 | 3         |