

# Susannah Woodruff

## List of Publications by Year in descending order

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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 Coyote Densities with Local, Discrete Bayesian Capture-Recapture Models. <i>Journal of Wildlife Management</i> , 2021, 85, 73-86.	1.8	3
2	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
3	Optimizing Selection of Brown Bear Hair for Noninvasive Genetic Analysis. <i>Wildlife Society Bulletin</i> , 2020, 44, 94-100.	1.6	4
4	Winter predation patterns of wolves in Northwestern Wyoming. <i>Journal of Wildlife Management</i> , 2019, 83, 1352-1367.	1.8	5
5	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
6	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
7	Advances in Using Non-invasive, Archival, and Environmental Samples for Population Genomic Studies. <i>Population Genomics</i> , 2018, , 63-99.	0.5	24
8	Estimating Sonoran pronghorn abundance and survival with fecal DNA and capture-recapture methods. <i>Conservation Biology</i> , 2016, 30, 1102-1111.	4.7	22
9	Monitoring coyote population dynamics with fecal DNA and spatial capture-recapture. <i>Journal of Wildlife Management</i> , 2016, 80, 824-836.	1.8	43
10	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
11	Evaluating the interaction of faecal pellet deposition rates and <i>scp&gt;DNA&lt;/scp&gt;</i> degradation rates to optimize sampling design for <i>scp&gt;DNA&lt;/scp&gt;</i> -based mark-recapture analysis of Sonoran pronghorn. <i>Molecular Ecology Resources</i> , 2015, 15, 843-854.	4.8	31
12	Rapid species identification of Sonoran pronghorn from fecal pellet DNA. <i>Wildlife Society Bulletin</i> , 2014, 38, 842-848.	1.6	9
13	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
14	Noninvasive individual and species identification of jaguars ( <i>Panthera onca</i> ), pumas ( <i>Puma</i> ) using microsatellites and faecal <i>scp&gt;DNA&lt;/scp&gt;</i> . <i>Molecular Ecology Resources</i> , 2014, 14, 1171-1182.	4.8	48
15	Evaluating DNA degradation rates in faecal pellets of the endangered pygmy rabbit. <i>Molecular Ecology Resources</i> , 2013, 13, 654-662.	4.8	37
16	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
17	Ten polymorphic microsatellite markers for pronghorn ( <i>Antilocapra americana</i> ). <i>Conservation Genetics Resources</i> , 2010, 2, 81-84.	0.8	12
18	Efficient, Noninvasive Genetic Sampling for Monitoring Reintroduced Wolves. <i>Journal of Wildlife Management</i> , 2010, 74, 1050-1058.	1.8	96

#	ARTICLE	IF	CITATIONS
19	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
20	Estimating abundance of American black bears using DNA-based capture-mark-recapture models. <i>Ursus</i> , 2009, 20, 1-11.	0.5	28
21	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
22	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
23	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
24	The influence of diet on faecal DNA amplification and sex identification in brown bears ( <i>Ursus arctos</i> ) Tj ETQq0 0 0.rgBT /Overlock 10 Tf	3.9	96
25	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
26	Noninvasive genetic sampling: look before you leap. <i>Trends in Ecology and Evolution</i> , 1999, 14, 323-327.	8.7	768