

# Alexandra L Decandia

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1670012/publications.pdf>

Version: 2024-02-01

11  
papers

229  
citations

933447

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1281871

11  
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13  
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13  
docs citations

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342  
citing authors

#	ARTICLE	IF	CITATIONS
1	Sarcoptic mange severity is associated with reduced genomic variation and evidence of selection in Yellowstone National Park wolves ( <i>Canis lupus</i> ). <i>Evolutionary Applications</i> , 2021, 14, 429-445.	3.1	13
2	Social environment and genetics underlie body site-specific microbiomes of Yellowstone National Park gray wolves ( <i>Canis lupus</i> ). <i>Ecology and Evolution</i> , 2021, 11, 9472-9488.	1.9	10
3	Heritability of interpack aggression in a wild pedigreed population of North American grey wolves. <i>Molecular Ecology</i> , 2020, 29, 1764-1775.	3.9	19
4	Ear mite infection is associated with altered microbial communities in genetically depauperate Santa Catalina Island foxes ( <i>Urocyon littoralis catalinae</i> ). <i>Molecular Ecology</i> , 2020, 29, 1463-1475.	3.9	17
5	Genetics of urban colonization: neutral and adaptive variation in coyotes ( <i>Canis latrans</i> ) inhabiting the New York metropolitan area. <i>Journal of Urban Ecology</i> , 2019, 5, .	1.5	14
6	Urban colonization through multiple genetic lenses: The city-fox phenomenon revisited. <i>Ecology and Evolution</i> , 2019, 9, 2046-2060.	1.9	28
7	Of microbes and mange: consistent changes in the skin microbiome of three canid species infected with <i>Sarcoptes scabiei</i> mites. <i>Parasites and Vectors</i> , 2019, 12, 488.	2.5	26
8	Toward an integrative molecular approach to wildlife disease. <i>Conservation Biology</i> , 2018, 32, 798-807.	4.7	36
9	High genomic diversity and candidate genes under selection associated with range expansion in eastern coyote ( <i>Canis latrans</i> ) populations. <i>Ecology and Evolution</i> , 2018, 8, 12641-12655.	1.9	21
10	Population Genomic Analysis of North American Eastern Wolves ( <i>Canis lycaon</i> ) Supports Their Conservation Priority Status. <i>Genes</i> , 2018, 9, 606.	2.4	32
11	A novel molecular method for noninvasive sex identification of order Carnivora. <i>Conservation Genetics Resources</i> , 2016, 8, 119-121.	0.8	12