

# Anna Letko

## List of Publications by Year in descending order

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34  
papers

328  
citations

932766

10  
h-index

996533

15  
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36  
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36  
docs citations

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times ranked

458  
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#	ARTICLE	IF	CITATIONS
1	Compound heterozygous PLA2G6 loss-of-function variants in Swaledale sheep with neuroaxonal dystrophy. <i>Molecular Genetics and Genomics</i> , 2021, 296, 235-242.	1.0	4
2	Phenotypic and Genomic Analysis of Cystic Hygroma in Pigs. <i>Genes</i> , 2021, 12, 207.	1.0	2
3	Genetic evaluation of small ruminant lentivirus susceptibility in Valais blacknose sheep. <i>Animal Genetics</i> , 2021, 52, 781-782.	0.6	1
4	Dog colour patterns explained by modular promoters of ancient canid origin. <i>Nature Ecology and Evolution</i> , 2021, 5, 1415-1423.	3.4	24
5	A frameshift insertion in FA2H causes a recessively inherited form of ichthyosis congenita in Chianina cattle. <i>Molecular Genetics and Genomics</i> , 2021, 296, 1313-1322.	1.0	6
6	Genome-Wide Analyses for Osteosarcoma in Leonberger Dogs Reveal the CDKN2A/B Gene Locus as a Major Risk Locus. <i>Genes</i> , 2021, 12, 1964.	1.0	8
7	A deletion spanning the promoter and first exon of the hair cycle-specific <i>ASIP</i> transcript isoform in black and tan rabbits. <i>Animal Genetics</i> , 2020, 51, 137-140.	0.6	14
8	A large deletion in the COL2A1 gene expands the spectrum of pathogenic variants causing bulldog calf syndrome in cattle. <i>Acta Veterinaria Scandinavica</i> , 2020, 62, 49.	0.5	8
9	Deleterious AGXT Missense Variant Associated with Type 1 Primary Hyperoxaluria (PH1) in Zwartbles Sheep. <i>Genes</i> , 2020, 11, 1147.	1.0	3
10	A CNTNAP1 Missense Variant Is Associated with Canine Laryngeal Paralysis and Polyneuropathy. <i>Genes</i> , 2020, 11, 1426.	1.0	9
11	Genomic diversity and population structure of the Leonberger dog breed. <i>Genetics Selection Evolution</i> , 2020, 52, 61.	1.2	9
12	SLC19A3 Loss-of-Function Variant in Yorkshire Terriers with Leigh-Like Subacute Necrotizing Encephalopathy. <i>Genes</i> , 2020, 11, 1215.	1.0	4
13	X-Linked Duchenne-Type Muscular Dystrophy in Jack Russell Terrier Associated with a Partial Deletion of the Canine DMD Gene. <i>Genes</i> , 2020, 11, 1175.	1.0	8
14	Multiple FGF4 Retrocopies Recently Derived within Canids. <i>Genes</i> , 2020, 11, 839.	1.0	12
15	A Missense Variant in ALDH5A1 Associated with Canine Succinic Semialdehyde Dehydrogenase Deficiency (SSADHD) in the Saluki Dog. <i>Genes</i> , 2020, 11, 1033.	1.0	3
16	Ear type in sheep is associated with the <i>MSRB3</i> locus. <i>Animal Genetics</i> , 2020, 51, 968-972.	0.6	12
17	A Heterozygous Missense Variant in the COL5A2 in Holstein Cattle Resembling the Classical Ehlers-Danlos Syndrome. <i>Animals</i> , 2020, 10, 2002.	1.0	7
18	<i>De novo</i> stop-loss germline mutation in <i>FGFR3</i> causes severe chondrodysplasia in the progeny of a Holstein bull. <i>Animal Genetics</i> , 2020, 51, 466-469.	0.6	4

#	ARTICLE	IF	CITATIONS
19	Whole Genome Sequencing Indicates Heterogeneity of Hyperostotic Disorders in Dogs. <i>Genes</i> , 2020, 11, 163.	1.0	4
20	New genomic features of the polled intersex syndrome variant in goats unraveled by long-read whole-genome sequencing. <i>Animal Genetics</i> , 2020, 51, 439-448.	0.6	14
21	A Missense Variant Affecting the C-Terminal Tail of UNC93B1 in Dogs with Exfoliative Cutaneous Lupus Erythematosus (ECLE). <i>Genes</i> , 2020, 11, 159.	1.0	13
22	A genome-wide significant association on chromosome 15 for congenital entropion in Swiss White Alpine sheep. <i>Animal Genetics</i> , 2020, 51, 278-283.	0.6	2
23	A de novo variant in OTX2 in a lamb with otocephaly. <i>Acta Veterinaria Scandinavica</i> , 2020, 62, 5.	0.5	1
24	A complex structural variant at the <i>KIT</i> locus in cattle with the Pinzgauer spotting pattern. <i>Animal Genetics</i> , 2019, 50, 423-429.	0.6	12
25	Identification of two <i>TYRP1</i> loss-of-function alleles in Valais Red sheep. <i>Animal Genetics</i> , 2019, 50, 778-782.	0.6	12
26	A <i>de novo</i> in-frame duplication in the <i>COL1A2</i> gene in a Lagotto Romagnolo dog with osteogenesis imperfecta. <i>Animal Genetics</i> , 2019, 50, 786-787.	0.6	8
27	Identification of a Missense Variant in MFSD12 Involved in Dilution of Pheomelanin Leading to White or Cream Coat Color in Dogs. <i>Genes</i> , 2019, 10, 386.	1.0	20
28	Chromosomal imbalance in pigs showing a syndromic form of cleft palate. <i>BMC Genomics</i> , 2019, 20, 349.	1.2	13
29	A Missense Variant in SCN8A in Alpine Dachsbracke Dogs Affected by Spinocerebellar Ataxia. <i>Genes</i> , 2019, 10, 362.	1.0	8
30	Canine NAPEPLD-associated models of human myelin disorders. <i>Scientific Reports</i> , 2018, 8, 5818.	1.6	14
31	A Missense Mutation in the Vacuolar Protein Sorting 11 ( <i>VPS11</i> ) Gene Is Associated with Neuroaxonal Dystrophy in Rottweiler Dogs. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 2773-2780.	0.8	19
32	Two <i>MC1R</i> loss-of-function alleles in cream-coloured Australian Cattle Dogs and white Huskies. <i>Animal Genetics</i> , 2018, 49, 284-290.	0.6	24
33	Two brown coat colour-associated <i>TYRP1</i> variants ( <i>b</i> and <i>Tj ETQg1.1 0.784314 rgB</i> )	0.6	3
34	A GJA9 frameshift variant is associated with polyneuropathy in Leonberger dogs. <i>BMC Genomics</i> , 2017, 18, 662.	1.2	20