

Chunyi Li

List of Publications by Year in descending order

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Version: 2024-02-01

83
papers

1,965
citations

236833

25
h-index

289141

40
g-index

85
all docs

85
docs citations

85
times ranked

839
citing authors

#	ARTICLE	IF	CITATIONS
1	Deer antler – A novel model for studying organ regeneration in mammals. <i>International Journal of Biochemistry and Cell Biology</i> , 2014, 56, 111-122.	1.2	110
2	Light microscopic studies of pedicle and early first antler development in red deer (<i>Cervus elaphus</i>). <i>The Anatomical Record</i> , 1994, 239, 198-215.	2.3	96
3	Sampling technique to discriminate the different tissue layers of growing antler tips for gene discovery. <i>The Anatomical Record</i> , 2002, 268, 125-130.	2.3	95
4	Red Deer Cloned from Antler Stem Cells and Their Differentiated Progeny ¹ . <i>Biology of Reproduction</i> , 2007, 77, 384-394.	1.2	94
5	Histological examination of antler regeneration in red deer (<i>Cervus elaphus</i>). <i>The Anatomical Record Part A: Discoveries in Molecular, Cellular, and Evolutionary Biology</i> , 2005, 282A, 163-174.	2.0	89
6	Adult Stem Cells and Mammalian Epimorphic Regeneration-Insights from Studying Annual Renewal of Deer Antlers. <i>Current Stem Cell Research and Therapy</i> , 2009, 4, 237-251.	0.6	86
7	Improbable appendages: Deer antler renewal as a unique case of mammalian regeneration. <i>Seminars in Cell and Developmental Biology</i> , 2009, 20, 535-542.	2.3	85
8	Deer antler regeneration: A stem cell-based epimorphic process. <i>Birth Defects Research Part C: Embryo Today Reviews</i> , 2012, 96, 51-62.	3.6	62
9	Identification of key tissue type for antler regeneration through pedicle periosteum deletion. <i>Cell and Tissue Research</i> , 2007, 328, 65-75.	1.5	55
10	Histological studies of pedicle skin formation and its transformation to antler velvet in red deer (<i>Cervus elaphus</i>). <i>The Anatomical Record</i> , 2000, 260, 62-71.	2.3	54
11	Deer antler stem cells are a novel type of cells that sustain full regeneration of a mammalian organ – deer antler. <i>Cell Death and Disease</i> , 2019, 10, 443.	2.7	50
12	Proteomes and Signalling Pathways of Antler Stem Cells. <i>PLoS ONE</i> , 2012, 7, e30026.	1.1	50
13	Cross-species metabolomic analysis identifies uridine as a potent regeneration promoting factor. <i>Cell Discovery</i> , 2022, 8, 6.	3.1	50
14	Morphological observation of antler regeneration in red deer (<i>Cervus elaphus</i>). <i>Journal of Morphology</i> , 2004, 262, 731-740.	0.6	48
15	Effects of insulin-like growth factor 1 and testosterone on the proliferation of antlerogenic cells in vitro. , 1999, 284, 82-90.		39
16	Histogenetic aspects of deer antler development. <i>Frontiers in Bioscience - Elite</i> , 2013, E5, 479-489.	0.9	39
17	Nerve Growth Factor mRNA Expression in the Regenerating Antler Tip of Red Deer (<i>Cervus elaphus</i>). <i>PLoS ONE</i> , 2007, 2, e148.	1.1	39
18	Vascular localization and proliferation in the growing tip of the deer antler. <i>The Anatomical Record Part A: Discoveries in Molecular, Cellular, and Evolutionary Biology</i> , 2006, 288A, 973-981.	2.0	36

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19	Exosomes from antler stem cells alleviate mesenchymal stem cell senescence and osteoarthritis. <i>Protein and Cell</i> , 2022, 13, 220-226.	4.8	36
20	Antler regeneration: a dependent process of stem tissue primed via interaction with its enveloping skin. <i>Journal of Experimental Zoology</i> , 2007, 307A, 95-105.	1.2	35
21	iTRAQ-Based Quantitative Proteomic Analysis of the Potentiated and Dormant Antler Stem Cells. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1778.	1.8	32
22	Genome-Wide SNP Discovery and Analysis of Genetic Diversity in Farmed Sika Deer (<i>Cervus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 Genes, Genomes, Genetics, 2017, 7, 3169-3176.	0.8	32
23	Tissue interactions and antlerogenesis: New findings revealed by a xenograft approach. <i>The Journal of Experimental Zoology</i> , 2001, 290, 18-30.	1.4	31
24	Antler stem cell-conditioned medium stimulates regenerative wound healing in rats. <i>Stem Cell Research and Therapy</i> , 2019, 10, 326.	2.4	31
25	Transcriptomic analysis of different tissue layers in antler growth Center in Sika Deer (<i>Cervus</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 62	1.2	29
26	The regenerating antler blastema the derivative of stem cells resident in a pedicle stump. <i>Frontiers in Bioscience - Landmark</i> , 2016, 21, 455-467.	3.0	28
27	Pedicle and antler development following sectioning of the sensory nerves to the antlerogenic region of red deer (<i>Cervus elaphus</i>). <i>The Journal of Experimental Zoology</i> , 1993, 267, 188-197.	1.4	27
28	Electron microscopic studies of antlerogenic cells from five developmental stages during pedicle and early antler formation in red deer (<i>Cervus elaphus</i>). , 1998, 252, 587-599.		23
29	Anti-tumour activity of deer growing antlers and its potential applications in the treatment of malignant gliomas. <i>Scientific Reports</i> , 2021, 11, 42.	1.6	23
30	Role of heterotypic tissue interactions in deer pedicle and first antler formationâ€”revealed via a membrane insertion approach. <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , 2008, 310B, 267-277.	0.6	19
31	Identifying deer antler uhrf1 proliferation and s100a10 mineralization genes using comparative RNA-seq. <i>Stem Cell Research and Therapy</i> , 2018, 9, 292.	2.4	17
32	Cell Cycle Genes PEDF and CDKN1C in Growing Deer Antlers. <i>Anatomical Record</i> , 2007, 290, 994-1004.	0.8	16
33	Development of a nude mouse model for the study of antlerogenesisâ€”mechanism of tissue interactions and ossification pathway. <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , 2009, 312B, 118-135.	0.6	16
34	Morphogenetic aspects of deer antler development. <i>Frontiers in Bioscience - Elite</i> , 2012, E4, 1836-1842.	0.9	16
35	MicroRNA profiling of antler stem cells in potentiated and dormant states and their potential roles in antler regeneration. <i>Molecular Genetics and Genomics</i> , 2016, 291, 943-955.	1.0	16
36	Deer thymosin beta 10 functions as a novel factor for angiogenesis and chondrogenesis during antler growth and regeneration. <i>Stem Cell Research and Therapy</i> , 2018, 9, 166.	2.4	16

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37	Transplanted Antler Stem Cells Stimulated Regenerative Healing of Radiation-induced Cutaneous Wounds in Rats. <i>Cell Transplantation</i> , 2020, 29, 096368972095154.	1.2	16
38	The periosteum: a simple tissue with many faces, with special reference to the antler-lineage periosteum. <i>Biology Direct</i> , 2021, 16, 17.	1.9	16
39	Lentiviral-Mediated RNAi Knockdown of Cbfa1 Gene Inhibits Endochondral Ossification of Antler Stem Cells in Micromass Culture. <i>PLoS ONE</i> , 2012, 7, e47367.	1.1	15
40	Extracellular vesicles derived from umbilical cord mesenchymal stromal cells alleviate pulmonary fibrosis by means of transforming growth factor- β signaling inhibition. <i>Stem Cell Research and Therapy</i> , 2021, 12, 230.	2.4	15
41	Morphogenetic aspects of deer antler development. <i>Frontiers in Bioscience - Elite</i> , 2012, E4, 1836.	0.9	14
42	Differential effects of the PI3K AKT pathway on antler stem cells for generation and regeneration of antlers <i>in vitro</i> . <i>Frontiers in Bioscience - Landmark</i> , 2018, 23, 1848-1863.	3.0	14
43	Quantitative proteomics analysis of deer antlerogenic periosteal cells reveals potential bioactive factors in velvet antlers. <i>Journal of Chromatography A</i> , 2020, 1609, 460496.	1.8	14
44	Antler stem cells as a novel stem cell source for reducing liver fibrosis. <i>Cell and Tissue Research</i> , 2020, 379, 195-206.	1.5	14
45	Quantitative proteomic analysis of deer antler stem cells as a model of mammalian organ regeneration. <i>Journal of Proteomics</i> , 2019, 195, 98-113.	1.2	12
46	Deer antler extracts reduce amyloid-beta toxicity in a <i>Caenorhabditis elegans</i> model of Alzheimer's disease. <i>Journal of Ethnopharmacology</i> , 2022, 285, 114850.	2.0	12
47	Pedicle and antler regeneration following antlerogenic tissue removal in red deer (<i>Cervus elaphus</i>). <i>The Journal of Experimental Zoology</i> , 1994, 269, 37-44.	1.4	11
48	Proteomic Analysis of Plasma Membrane Proteins of Antler Stem Cells Using Label-Free LC-MS/MS. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3477.	1.8	11
49	Stem cells responsible for deer antler regeneration are unable to recapitulate the process of first antler development revealed through intradermal and subcutaneous tissue transplantation. <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , 2010, 314B, 552-570.	0.6	10
50	Stem cells, stem cell niche and antler development. <i>Animal Production Science</i> , 2011, 51, 267.	0.6	10
51	Mapping the morphogenetic potential of antler fields through deleting and transplanting subregions of antlerogenic periosteum in sika deer (<i>Cervus nippon</i>). <i>Journal of Anatomy</i> , 2012, 220, 131-143.	0.9	10
52	Direct localisation of molecules in tissue sections of growing antler tips using MALDI imaging. <i>Molecular and Cellular Biochemistry</i> , 2015, 409, 225-241.	1.4	10
53	Classification and phylogeny of sika deer (<i>Cervus nippon</i>) subspecies based on the mitochondrial control region DNA sequence using an extended sample set. <i>Mitochondrial DNA</i> , 2015, 26, 373-379.	0.6	10
54	Identification of proteins that mediate the role of androgens in antler regeneration using label free proteomics in sika deer (<i>Cervus nippon</i>). <i>General and Comparative Endocrinology</i> , 2019, 283, 113235.	0.8	10

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55	Chromosome-level genome assembly of Tarim red deer, <i>Cervus elaphus yarkandensis</i> . <i>Scientific Data</i> , 2020, 7, 187.	2.4	10
56	Effects of unilateral cranial sympathectomy either alone or with sensory nerve sectioning on pedicle growth in red deer (<i>Cervus elaphus</i>). <i>The Journal of Experimental Zoology</i> , 1995, 271, 131-138.	1.4	8
57	Single-cell transcriptome provides novel insights into antler stem cells, a cell type capable of mammalian organ regeneration. <i>Functional and Integrative Genomics</i> , 2019, 19, 555-564.	1.4	8
58	PTN ^Δ ~PTPRZ signalling is involved in deer antler stem cell regulation during tissue regeneration. <i>Journal of Cellular Physiology</i> , 2021, 236, 3752-3769.	2.0	8
59	Effects of p21 Gene Down-Regulation through RNAi on Antler Stem Cells In Vitro. <i>PLoS ONE</i> , 2015, 10, e0134268.	1.1	7
60	Cloning and Characterization of a <i>Nanog</i> Pseudogene in Sika Deer (<i>Cervus nippon</i>). <i>DNA and Cell Biology</i> , 2016, 35, 576-584.	0.9	7
61	Substances for regenerative wound healing during antler renewal stimulated scar-less restoration of rat cutaneous wounds. <i>Cell and Tissue Research</i> , 2021, 386, 99-116.	1.5	7
62	Morphogenetic Mechanisms in the Cyclic Regeneration of Hair Follicles and Deer Antlers from Stem Cells. <i>BioMed Research International</i> , 2013, 2013, 1-21.	0.9	6
63	Deer antlers: traditional Chinese medicine use and recent pharmaceuticals. <i>Animal Production Science</i> , 2020, 60, 1233.	0.6	6
64	Velvet Antler Peptides Reduce Scarring via Inhibiting the TGF- β 2 Signaling Pathway During Wound Healing. <i>Frontiers in Medicine</i> , 2021, 8, 799789.	1.2	6
65	Analysis of Genomewide DNA Methylation Reveals Differences in DNA Methylation Levels between Dormant and Naturally as well as Artificially Potentiated Pedicle Periosteum of Sika Deer (<i>Cervus</i>). <i>Tj ETQq1 1 0.784314 rgBT /Over</i> 326, 375-383.	0.6	6
66	An examination of the origin and evolution of additional tandem repeats in the mitochondrial DNA control region of Japanese sika deer (<i>Cervus Nippon</i>). <i>Mitochondrial DNA</i> , 2016, 27, 276-281.	0.6	5
67	Custom-built tools for the study of deer antler biology. <i>Frontiers in Bioscience - Landmark</i> , 2017, 22, 1622-1633.	3.0	5
68	Genome-wide analysis of DNA methylation in five tissues of sika deer (<i>Cervus nippon</i>). <i>Gene</i> , 2018, 645, 48-54.	1.0	5
69	Development of Diagnostic SNP Markers To Monitor Hybridization Between Sika Deer (<i>Cervus nippon</i>) and Wapiti (<i>Cervus elaphus</i>). <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 2173-2179.	0.8	5
70	Identification of interactive molecules between antler stem cells and dermal papilla cells using an in vitro co-culture system. <i>Journal of Molecular Histology</i> , 2020, 51, 15-31.	1.0	5
71	Complete mitochondrial genome of the muskrat (<i>Ondatra zibethicus</i>) and its unique phylogenetic position estimated in Cricetidae. <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 296-298.	0.2	4
72	S100A4: a novel partner for heat shock protein 47 in antler stem cells and insight into the calcium ion-induced conformational changes. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 2068-2079.	2.0	2

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73	Molecular evidence for adaptive evolution of olfactory-related genes in cervids. <i>Genes and Genomics</i> , 2020, 42, 355-360.	0.5	2
74	Residual antler periosteum holds the potential to partially regenerate lost antler tissue. <i>Journal of Experimental Zoology Part A: Ecological and Integrative Physiology</i> , 2021, 335, 386-395.	0.9	2
75	Antler Stem Cells Sustain Regenerative Wound Healing in Deer and in Rats. <i>Journal of Regenerative Biology and Medicine</i> , 0, , .	0.0	2
76	Calreticulin Identified as One of the Androgen Response Genes That Trigger Full Regeneration of the Only Capable Mammalian Organ, the Deer Antler. <i>Frontiers in Cell and Developmental Biology</i> , 0, 10, .	1.8	2
77	Association analysis of thirty-one single nucleotide polymorphisms with antler weight in sika deer. <i>Animal Genetics</i> , 2020, 51, 990-991.	0.6	1
78	Reclassification of velvet antler portions following transcriptomic analysis. <i>Animal Production Science</i> , 2020, 60, 1364.	0.6	0
79	Design of a universal primer pair for the identification of deer species. <i>Conservation Genetics Resources</i> , 2021, 13, 9-12.	0.4	0
80	Effects of macrophage-conditioned medium on sika deer (<i>Cervus nippon</i>) antler stem cells. <i>Animal Production Science</i> , 2020, 60, 1326.	0.6	0
81	Cross-Species Analysis Reveals Co-Expressed Genes Regulating Antler Development in Cervidae. <i>Frontiers in Genetics</i> , 0, 13, .	1.1	0
82	Tracing the geographic origin of velvet antlers in China <i>via</i> stable isotope analyses. <i>RSC Advances</i> , 2022, 12, 17527-17535.	1.7	0
83	IGF1R and LOX Modules Are Related to Antler Growth Rate Revealed by Integrated Analyses of Genomics and Transcriptomics. <i>Animals</i> , 2022, 12, 1522.	1.0	0