

David Alan Young

List of Publications by Year in descending order

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Version: 2024-02-01

82
papers

4,447
citations

109321

35
h-index

106344

65
g-index

113
all docs

113
docs citations

113
times ranked

5597
citing authors

#	ARTICLE	IF	CITATIONS
1	Expression profiling of metalloproteinases and their inhibitors in cartilage. <i>Arthritis and Rheumatism</i> , 2004, 50, 131-141.	6.7	379
2	The Comparative Role of Activator Protein 1 and Smad Factors in the Regulation of Timp-1 and MMP-1 Gene Expression by Transforming Growth Factor- β 21. <i>Journal of Biological Chemistry</i> , 2003, 278, 10304-10313.	3.4	211
3	The expression and function of microRNAs in chondrogenesis and osteoarthritis. <i>Arthritis and Rheumatism</i> , 2012, 64, 1909-1919.	6.7	204
4	Superoxide dismutase downregulation in osteoarthritis progression and end-stage disease. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 1502-1510.	0.9	202
5	Leptin produced by joint white adipose tissue induces cartilage degradation via upregulation and activation of matrix metalloproteinases. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 455-462.	0.9	174
6	Histone deacetylase inhibitors modulate metalloproteinase gene expression in chondrocytes and block cartilage resorption. <i>Arthritis Research</i> , 2005, 7, R503.	2.0	153
7	Epigenetic mechanisms in cartilage and osteoarthritis: DNA methylation, histone modifications and microRNAs. <i>Osteoarthritis and Cartilage</i> , 2012, 20, 339-349.	1.3	152
8	Characterization of the Cartilage DNA Methylome in Knee and Hip Osteoarthritis. <i>Arthritis and Rheumatology</i> , 2014, 66, 2450-2460.	5.6	146
9	Oxidative changes and signalling pathways are pivotal in initiating age-related changes in articular cartilage. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 449-458.	0.9	135
10	Proteinases involved in matrix turnover during cartilage and bone breakdown. <i>Cell and Tissue Research</i> , 2010, 339, 221-235.	2.9	131
11	Mitochondrial dysfunction in osteoarthritis is associated with downregulation of superoxide dismutase 2. <i>Arthritis and Rheumatism</i> , 2013, 65, 378-387.	6.7	113
12	Expression of the osteoarthritis-associated gene GDF5 is modulated epigenetically by DNA methylation. <i>Human Molecular Genetics</i> , 2011, 20, 3450-3460.	2.9	108
13	The microRNA-29 family in cartilage homeostasis and osteoarthritis. <i>Journal of Molecular Medicine</i> , 2016, 94, 583-596.	3.9	106
14	Quantification of Neuroreceptors in the Living Human Brain: III. D2-Like Dopamine Receptors: Theory, Validation, and Changes during Normal Aging. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 1997, 17, 316-330.	4.3	98
15	cAMP response element-binding (CREB) recruitment following a specific CpG demethylation leads to the elevated expression of the matrix metalloproteinase 13 in human articular chondrocytes and osteoarthritis. <i>FASEB Journal</i> , 2012, 26, 3000-3011.	0.5	96
16	Interplay between genetics and epigenetics in osteoarthritis. <i>Nature Reviews Rheumatology</i> , 2020, 16, 268-281.	8.0	91
17	Identification of the pathogenic pathways in osteoarthritic hip cartilage: commonality and discord between hip and knee OA. <i>Osteoarthritis and Cartilage</i> , 2012, 20, 1029-1038.	1.3	81
18	Differential Toll-like receptor-dependent collagenase expression in chondrocytes. <i>Annals of the Rheumatic Diseases</i> , 2008, 67, 1633-1641.	0.9	79

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19	HDAC-mediated control of ERK- and PI3K-dependent TGF- β -induced extracellular matrix-regulating genes. <i>Matrix Biology</i> , 2010, 29, 602-612.	3.6	74
20	Genome-Wide MicroRNA and Gene Analysis of Mesenchymal Stem Cell Chondrogenesis Identifies an Essential Role and Multiple Targets for miR-140-5p. <i>Stem Cells</i> , 2015, 33, 3266-3280.	3.2	72
21	Fibroblast activation protein alpha is expressed by chondrocytes following a pro-inflammatory stimulus and is elevated in osteoarthritis. <i>Arthritis Research and Therapy</i> , 2006, 8, R23.	3.5	71
22	Class I Histone Deacetylase Inhibition Modulates Metalloproteinase Expression and Blocks Cytokine-Induced Cartilage Degradation. <i>Arthritis and Rheumatism</i> , 2013, 65, 1822-1830.	6.7	70
23	Long noncoding RNA <i>ROCR</i> contributes to SOX9 expression and chondrogenic differentiation of human mesenchymal stem cells. <i>Development (Cambridge)</i> , 2017, 144, 4510-4521.	2.5	70
24	A CD4 T cell gene signature for early rheumatoid arthritis implicates interleukin 6-mediated STAT3 signalling, particularly in anti-citrullinated peptide antibody-negative disease. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1374-1381.	0.9	67
25	Identification of an initiator-like element essential for the expression of the tissue inhibitor of metalloproteinases-4 (Timp-4) gene. <i>Biochemical Journal</i> , 2002, 364, 89-99.	3.7	62
26	Metalloproteinase and inhibitor expression profiling of resorbing cartilage reveals pro-collagenase activation as a critical step for collagenolysis. <i>Arthritis Research and Therapy</i> , 2006, 8, R142.	3.5	61
27	Matriptase is a novel initiator of cartilage matrix degradation in osteoarthritis. <i>Arthritis and Rheumatism</i> , 2010, 62, 1955-1966.	6.7	61
28	The Identification of Trans-acting Factors That Regulate the Expression of GDF5 via the Osteoarthritis Susceptibility SNP rs143383. <i>PLoS Genetics</i> , 2013, 9, e1003557.	3.5	53
29	Synergistic Collagenase Expression and Cartilage Collagenolysis Are Phosphatidylinositol 3-Kinase/Akt Signaling-dependent. <i>Journal of Biological Chemistry</i> , 2008, 283, 14221-14229.	3.4	52
30	Epigenetic Mechanisms and Non-coding RNAs in Osteoarthritis. <i>Current Rheumatology Reports</i> , 2013, 15, 353.	4.7	49
31	Methylation quantitative trait locus analysis of osteoarthritis links epigenetics with genetic risk. <i>Human Molecular Genetics</i> , 2015, 24, 7432-7444.	2.9	48
32	Lithium protects cartilage from cytokine-mediated degradation by reducing collagen-degrading MMP production via inhibition of the P38 mitogen-activated protein kinase pathway. <i>Rheumatology</i> , 2010, 49, 2043-2053.	1.9	46
33	Serum snoRNAs as biomarkers for joint ageing and post traumatic osteoarthritis. <i>Scientific Reports</i> , 2017, 7, 43558.	3.3	44
34	Expression of metalloproteinases and inhibitors in the differentiation of P19CL6 cells into cardiac myocytes. <i>Biochemical and Biophysical Research Communications</i> , 2004, 322, 759-765.	2.1	43
35	The function of microRNAs in cartilage and osteoarthritis. <i>Clinical and Experimental Rheumatology</i> , 2019, 37 Suppl 120, 40-47.	0.8	42
36	Detecting new microRNAs in human osteoarthritic chondrocytes identifies miR-3085 as a human, chondrocyte-selective, microRNA. <i>Osteoarthritis and Cartilage</i> , 2016, 24, 534-543.	1.3	38

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37	A Negative Feedback Loop Mediated by STAT3 Limits Human Th17 Responses. <i>Journal of Immunology</i> , 2014, 193, 1142-1150.	0.8	37
38	miR-324-5p is up regulated in end-stage osteoarthritis and regulates Indian Hedgehog signalling by differing mechanisms in human and mouse. <i>Matrix Biology</i> , 2019, 77, 87-100.	3.6	37
39	Oncostatin M in combination with tumour necrosis factor α induces a chondrocyte membrane associated aggrecanase that is distinct from ADAMTS aggrecanase-1 or -2. <i>Annals of the Rheumatic Diseases</i> , 2005, 64, 1624-1632.	0.9	36
40	Lipophilic statins prevent matrix metalloproteinase-mediated cartilage collagen breakdown by inhibiting protein geranylgeranylation. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 2189-2198.	0.9	36
41	The Human Tissue Inhibitor of Metalloproteinases (TIMP)-1 Gene Contains Repressive Elements within the Promoter and Intron 1. <i>Journal of Biological Chemistry</i> , 2000, 275, 32664-32671.	3.4	34
42	Identification of long non-coding RNAs expressed in knee and hip osteoarthritic cartilage. <i>Osteoarthritis and Cartilage</i> , 2019, 27, 694-702.	1.3	34
43	Recent advances in understanding the regulation of metalloproteinases. <i>F1000Research</i> , 2019, 8, 195.	1.6	34
44	Differential Gene Expression Profiling of Metalloproteinases and Their Inhibitors. <i>Spine</i> , 2010, 35, 1101-1108.	2.0	30
45	Differential effects of histone deacetylase inhibitors on phorbol ester- and TGF- β 1 induced murine tissue inhibitor of metalloproteinases-1 gene expression. <i>FEBS Journal</i> , 2005, 272, 1912-1926.	4.7	28
46	Activation of p38 and JNK MAPK pathways abrogates requirement for new protein synthesis for phorbol ester mediated induction of select MMP and TIMP genes. <i>Matrix Biology</i> , 2008, 27, 128-138.	3.6	28
47	Protection against murine osteoarthritis by inhibition of the 26S proteasome and lysine-48 linked ubiquitination. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 1580-1587.	0.9	27
48	Collagenase gene regulation by pro-inflammatory cytokines in cartilage. <i>Frontiers in Bioscience - Landmark</i> , 2007, 12, 536.	3.0	27
49	MMP28 gene expression is regulated by Sp1 transcription factor acetylation. <i>Biochemical Journal</i> , 2010, 427, 391-400.	3.7	26
50	Matrix Metalloproteinase 13 Expression in Response to Double-Stranded RNA in Human Chondrocytes. <i>Arthritis and Rheumatism</i> , 2013, 65, 1290-1301.	6.7	23
51	Differential DNA methylation and expression of inflammatory and zinc transporter genes defines subgroups of osteoarthritic hip patients. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 1778-1782.	0.9	23
52	Osteoarthritis year in review: genetics, genomics, epigenetics. <i>Osteoarthritis and Cartilage</i> , 2022, 30, 216-225.	1.3	23
53	Pattern recognition receptor expression is not impaired in patients with chronic mucocutaneous candidiasis with or without autoimmune polyendocrinopathy candidiasis ectodermal dystrophy. <i>Clinical and Experimental Immunology</i> , 2009, 156, 40-51.	2.6	22
54	Glycogen Synthase Kinase 3 Inhibition Stimulates Human Cartilage Destruction and Exacerbates Murine Osteoarthritis. <i>Arthritis and Rheumatology</i> , 2014, 66, 2175-2187.	5.6	22

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55	OATargets: a knowledge base of genes associated with osteoarthritis joint damage in animals. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 376-383.	0.9	21
56	Acetylation in the regulation of metalloproteinase and tissue inhibitor of metalloproteinases gene expression. <i>Frontiers in Bioscience - Landmark</i> , 2007, 12, 528.	3.0	21
57	Matrix metalloproteinase-13 is fully activated by neutrophil elastase and inactivates its serpin inhibitor, alpha-1 antitrypsin: Implications for osteoarthritis. <i>FEBS Journal</i> , 2022, 289, 121-139.	4.7	20
58	Histone ChIP-seq identifies differential enhancer usage during chondrogenesis as critical for defining cell-type specificity. <i>FASEB Journal</i> , 2020, 34, 5317-5331.	0.5	18
59	DNA hypomethylation during MSC chondrogenesis occurs predominantly at enhancer regions. <i>Scientific Reports</i> , 2020, 10, 1169.	3.3	18
60	microRNA-seq of cartilage reveals an overabundance of miR-140-3p which contains functional isomiRs. <i>Rna</i> , 2020, 26, 1575-1588.	3.5	17
61	An enhancer complex confers both high-level and cell-specific expression of the human type X collagen gene. <i>FEBS Letters</i> , 2002, 531, 505-508.	2.8	15
62	Rac upregulates tissue inhibitor of metalloproteinase-1 expression by redox-dependent activation of extracellular signal-regulated kinase signaling. <i>FEBS Journal</i> , 2006, 273, 4754-4769.	4.7	14
63	<sc>CRELD2</sc> Is a Novel <sc>LRP1</sc> Chaperone That Regulates Noncanonical <sc>WNT</sc> Signaling in Skeletal Development. <i>Journal of Bone and Mineral Research</i> , 2020, 35, 1452-1469.	2.8	12
64	Regulation of microRNA-221, -222, -21 and -27 in articular cartilage subjected to abnormal compressive forces. <i>Journal of Physiology</i> , 2021, 599, 143-155.	2.9	12
65	Correlation of Infinium HumanMethylation450K and MethylationEPIC BeadChip arrays in cartilage. <i>Epigenetics</i> , 2020, 15, 594-603.	2.7	10
66	Understanding CpG methylation in the context of osteoarthritis. <i>Epigenomics</i> , 2012, 4, 593-595.	2.1	8
67	Kinetics Analysis of Circulating MicroRNAs Unveils Markers of Failed Myocardial Reperfusion. <i>Clinical Chemistry</i> , 2020, 66, 247-256.	3.2	8
68	Dynamic chromatin accessibility landscape changes following interleukin-1 stimulation. <i>Epigenetics</i> , 2021, 16, 106-119.	2.7	8
69	The first international workshop on the epigenetics of osteoarthritis. <i>Connective Tissue Research</i> , 2017, 58, 37-48.	2.3	6
70	Highly efficient CRISPR-Cas9-mediated editing identifies novel mechanosensitive microRNA-140 targets in primary human articular chondrocytes. <i>Osteoarthritis and Cartilage</i> , 2022, , .	1.3	6
71	The role of microRNA-3085 in chondrocyte function. <i>Scientific Reports</i> , 2020, 10, 21923.	3.3	5
72	HDAC6 regulates NF- κ B signalling to control chondrocyte IL-1-induced MMP and inflammatory gene expression. <i>Scientific Reports</i> , 2022, 12, 6640.	3.3	5

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73	Editorial: More evidence for a role of CpG methylation in the pathogenesis of osteoarthritis. <i>Arthritis and Rheumatism</i> , 2013, 65, 555-558.	6.7	4
74	Reply to Swart and Korf. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 1989, 9, 908-910.	4.3	3
75	microRNA in Chondrogenesis, Cartilage and Osteoarthritis. <i>Current Rheumatology Reviews</i> , 2012, 8, 89-97.	0.8	3
76	Anesthesia for the child with Andersen's Syndrome. <i>Paediatric Anaesthesia</i> , 2005, 15, 1019-1020.	1.1	0
77	British Society for Matrix Biology Autumn Meeting – Joint with the UK Tissue & Cell Engineering Society, University of Bristol, UK. <i>International Journal of Experimental Pathology</i> , 2005, 86, A1-A56.	1.3	0
78	Differential Toll-like receptor-dependent collagenase expression in chondrocytes. <i>Arthritis Research and Therapy</i> , 2007, 9, P39.	3.5	0
79	The role of acetylation in Timp-1 regulation. <i>International Journal of Experimental Pathology</i> , 2008, 85, A18-A19.	1.3	0
80	Expression profiling of metalloproteinases and inhibitors in cartilage. <i>International Journal of Experimental Pathology</i> , 2008, 85, A23-A23.	1.3	0
81	250. – UNDERSTANDING ABERRANT IL-6 MEDIATED CD4+ T-CELL SIGNALLING IN EARLY RHEUMATOID ARTHRITIS. <i>Rheumatology</i> , 2017, 56, .	1.9	0
82	08.13 – Understanding aberrant il-6 mediated cd4+ t-cell signalling in early rheumatoid arthritis. , 2017, , .		0