Alexandre F R Stewart

List of Publications by Year in Descending Order

139 papers	19,924 citations	56 h-index	141 g-index
151 ext. papers	23,494 ext. citations	10.1 avg, IF	5.57 L-index

#	Paper	IF	Citations
139	N-methyl-D-aspartate receptor functions altered by neuronal PTP1B activation in Alzheimer's disease and schizophrenia models.. *Neural Regeneration Research*, **2022**, 17, 2208-2210	4.5	
138	Revisiting the MMTV Zoonotic Hypothesis to Account for Geographic Variation in Breast Cancer Incidence.. *Viruses*, **2022**, 14,	6.2	1
137	IRF2BP2 3'UTR Polymorphism Increases Coronary Artery Calcification in Men. *Frontiers in Cardiovascular Medicine*, **2021**, 8, 687645	5.4	0
136	Neuronal protein-tyrosine phosphatase 1B hinders sensory-motor functional recovery and causes affective disorders in two different focal ischemic stroke models. *Neural Regeneration Research*, **2021**, 16, 129-136	4.5	2
135	Integrative analysis of genomic variants reveals new associations of candidate haploinsufficient genes with congenital heart disease. *PLoS Genetics*, **2021**, 17, e1009679	6	1
134	Ketamine's schizophrenia-like effects are prevented by targeting PTP1B. *Neurobiology of Disease*, **2021**, 155, 105397	7.5	2
133	Tyrosine phosphatase PTP1B impairs presynaptic NMDA receptor-mediated plasticity in a mouse model of Alzheimer's disease. *Neurobiology of Disease*, **2021**, 156, 105402	7.5	5
132	Activation of tyrosine phosphatase PTP1B in pyramidal neurons impairs endocannabinoid signaling by tyrosine receptor kinase trkB and causes schizophrenia-like behaviors in mice. *Neuropsychopharmacology*, **2020**, 45, 1884-1895	8.7	6
131	Hyperactivated PTP1B phosphatase in parvalbumin neurons alters anterior cingulate inhibitory circuits and induces autism-like behaviors. *Nature Communications*, **2020**, 11, 1017	17.4	11
130	Activation of tyrosine phosphatases in the progression of Alzheimer's disease. *Neural Regeneration Research*, **2020**, 15, 2245-2246	4.5	3
129	Neuronal Protein Tyrosine Phosphatase 1B Hastens Amyloid β-Associated Alzheimer's Disease in Mice. *Journal of Neuroscience*, **2020**, 40, 1581-1593	6.6	24
128	Association of Factor V Leiden With Subsequent Atherothrombotic Events: A GENIUS-CHD Study of Individual Participant Data. *Circulation*, **2020**, 142, 546-555	16.7	5
127	Subsequent Event Risk in Individuals With Established Coronary Heart Disease. *Circulation Genomic and Precision Medicine*, **2019**, 12, e002470	5.2	13
126	Association of Chromosome 9p21 With Subsequent Coronary Heart Disease Events. *Circulation Genomic and Precision Medicine*, **2019**, 12, e002471	5.2	14
125	Association of the PHACTR1/EDN1 Genetic Locus With Spontaneous Coronary Artery Dissection. *Journal of the American College of Cardiology*, **2019**, 73, 58-66	15.1	86
124	Dabrafenib, an inhibitor of RIP3 kinase-dependent necroptosis, reduces ischemic brain injury. *Neural Regeneration Research*, **2018**, 13, 252-256	4.5	37
123	Loss of Cardioprotective Effects at the Locus as a Result of Gene-Smoking Interactions. *Circulation*, **2017**, 135, 2336-2353	16.7	36

122	Relations between lipoprotein(a) concentrations, LPA genetic variants, and the risk of mortality in patients with established coronary heart disease: a molecular and genetic association study. *Lancet Diabetes and Endocrinology,the*, **2017**, 5, 534-543	18.1	69
121	Identification of novel risk loci for restless legs syndrome in genome-wide association studies in individuals of European ancestry: a meta-analysis. *Lancet Neurology, The*, **2017**, 16, 898-907	24.1	121
120	IRF2BP2-deficient microglia block the anxiolytic effect of enhanced postnatal care. *Scientific Reports*, **2017**, 7, 9836	4.9	12
119	Partitioning the heritability of coronary artery disease highlights the importance of immune-mediated processes and epigenetic sites associated with transcriptional activity. *Cardiovascular Research*, **2017**, 113, 973-983	9.9	18
118	Loss of IRF2BP2 in Microglia Increases Inflammation and Functional Deficits after Focal Ischemic Brain Injury. *Frontiers in Cellular Neuroscience*, **2017**, 11, 201	6.1	27
117	Interferon regulatory factor 2 binding protein 2: a new player of the innate immune response for stroke recovery. *Neural Regeneration Research*, **2017**, 12, 1762-1764	4.5	6
116	No Association of Coronary Artery Disease with X-Chromosomal Variants in Comprehensive International Meta-Analysis. *Scientific Reports*, **2016**, 6, 35278	4.9	18
115	Transcriptomic Signature of Atherosclerosis in the Peripheral Blood: Fact or Fiction?. *Current Atherosclerosis Reports*, **2016**, 18, 77	6	10
114	Genome-wide association study and targeted metabolomics identifies sex-specific association of CPS1 with coronary artery disease. *Nature Communications*, **2016**, 7, 10558	17.4	79
113	Increased genetic risk for obesity in premature coronary artery disease. *European Journal of Human Genetics*, **2016**, 24, 587-91	5.3	15
112	Chronic stress induces anxiety via an amygdalar intracellular cascade that impairs endocannabinoid signaling. *Neuron*, **2015**, 85, 1319-31	13.9	62
111	IRF2BP2 Reduces Macrophage Inflammation and Susceptibility to Atherosclerosis. *Circulation Research*, **2015**, 117, 671-83	15.7	46
110	Effect of Bile Acid Sequestrants on the Risk of Cardiovascular Events: A Mendelian Randomization Analysis. *Circulation: Cardiovascular Genetics*, **2015**, 8, 618-27		41
109	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. *Nature Genetics*, **2015**, 47, 1282-1293	36.3	223
108	Circulating brain-derived neurotrophic factor concentrations and the risk of cardiovascular disease in the community. *Journal of the American Heart Association*, **2015**, 4, e001544	6	70
107	9p21.3 Coronary Artery Disease Risk Variants Disrupt TEAD Transcription Factor-Dependent Transforming Growth Factor Regulation of p16 Expression in Human Aortic Smooth Muscle Cells. *Circulation*, **2015**, 132, 1969-78	16.7	31
106	A comprehensive 1,000 Genomes-based genome-wide association meta-analysis of coronary artery disease. *Nature Genetics*, **2015**, 47, 1121-1130	36.3	1290
105	Genetic analysis for a shared biological basis between migraine and coronary artery disease. *Neurology: Genetics*, **2015**, 1, e10	3.8	46

104	Exome sequencing identifies rare LDLR and APOA5 alleles conferring risk for myocardial infarction. *Nature*, **2015**, 518, 102-6	50.4	463
103	Systems Genetics Analysis of Genome-Wide Association Study Reveals Novel Associations Between Key Biological Processes and Coronary Artery Disease. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2015**, 35, 1712-22	9.4	55
102	Endothelial Gata5 transcription factor regulates blood pressure. *Nature Communications*, **2015**, 6, 8835	17.4	30
101	New genetic loci link adipose and insulin biology to body fat distribution. *Nature*, **2015**, 518, 187-196	50.4	920
100	Genetic studies of body mass index yield new insights for obesity biology. *Nature*, **2015**, 518, 197-206	50.4	2687
99	Functional properties of Claramine: a novel PTP1B inhibitor and insulin-mimetic compound. *Biochemical and Biophysical Research Communications*, **2015**, 458, 21-7	3.4	44
98	Shared genetic susceptibility to ischemic stroke and coronary artery disease: a genome-wide analysis of common variants. *Stroke*, **2014**, 45, 24-36	6.7	245
97	LMO4 is essential for paraventricular hypothalamic neuronal activity and calcium channel expression to prevent hyperphagia. *Journal of Neuroscience*, **2014**, 34, 140-8	6.6	12
96	Comparative genome-wide association studies in mice and humans for trimethylamine N-oxide, a proatherogenic metabolite of choline and L-carnitine. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2014**, 34, 1307-13	9.4	94
95	LMO4 is required to maintain hypothalamic insulin signaling. *Biochemical and Biophysical Research Communications*, **2014**, 450, 666-72	3.4	15
94	SPG7 variant escapes phosphorylation-regulated processing by AFG3L2, elevates mitochondrial ROS, and is associated with multiple clinical phenotypes. *Cell Reports*, **2014**, 7, 834-47	10.6	29
93	Functional genomics of the 9p21.3 locus for atherosclerosis: clarity or confusion?. *Current Cardiology Reports*, **2014**, 16, 502	4.2	36
92	Plasma PCSK9 levels are elevated with acute myocardial infarction in two independent retrospective angiographic studies. *PLoS ONE*, **2014**, 9, e106294	3.7	57
91	Integrative genomics reveals novel molecular pathways and gene networks for coronary artery disease. *PLoS Genetics*, **2014**, 10, e1004502	6	147
90	Adiposity significantly modifies genetic risk for dyslipidemia. *Journal of Lipid Research*, **2014**, 55, 2416-22	6.3	25
89	Is atherosclerosis fundamental to human aging? Lessons from ancient mummies. *Journal of Cardiology*, **2014**, 63, 329-34	3	23
88	Genomic correlates of atherosclerosis in ancient humans. *Global Heart*, **2014**, 9, 203-9	2.9	14
87	Why did ancient people have atherosclerosis?: from autopsies to computed tomography to potential causes. *Global Heart*, **2014**, 9, 229-37	2.9	23

86	Interferon- activates expression of p15 and p16 regardless of 9p21.3 coronary artery disease risk genotype. *Journal of the American College of Cardiology*, **2013**, 61, 143-7	15.1	32
85	Genomics in cardiovascular disease. *Journal of the American College of Cardiology*, **2013**, 61, 2029-37	15.1	24
84	The LIM domain only 4 protein is a metabolic responsive inhibitor of protein tyrosine phosphatase 1B that controls hypothalamic leptin signaling. *Journal of Neuroscience*, **2013**, 33, 12647-55	6.6	35
83	Large-scale association analysis identifies new risk loci for coronary artery disease. *Nature Genetics*, **2013**, 45, 25-33	36.3	1172
82	Assessment of the 9p21.3 locus in severity of coronary artery disease in the presence and absence of type 2 diabetes. *BMC Medical Genetics*, **2013**, 14, 11	2.1	21
81	Two chromosome 9p21 haplotype blocks distinguish between coronary artery disease and myocardial infarction risk. *Circulation: Cardiovascular Genetics*, **2013**, 6, 372-80		19
80	Point-of-care genetic testing for personalisation of antiplatelet treatment (RAPID GENE): a prospective, randomised, proof-of-concept trial. *Lancet, The*, **2012**, 379, 1705-11	40	285
79	Plasma HDL cholesterol and risk of myocardial infarction: a mendelian randomisation study. *Lancet, The*, **2012**, 380, 572-80	40	1523
78	9p21 and the genetic revolution for coronary artery disease. *Clinical Chemistry*, **2012**, 58, 104-12	5.5	45
77	Genomics: is it ready for primetime?. *Medical Clinics of North America*, **2012**, 96, 113-22	7	1
76	Tail-anchored membrane protein SLMAP is a novel regulator of cardiac function at the sarcoplasmic reticulum. *American Journal of Physiology - Heart and Circulatory Physiology*, **2012**, 302, H1138-45	5.2	19
75	Genes and coronary artery disease: where are we?. *Journal of the American College of Cardiology*, **2012**, 60, 1715-21	15.1	117
74	A genome-wide association study for coronary artery disease identifies a novel susceptibility locus in the major histocompatibility complex. *Circulation: Cardiovascular Genetics*, **2012**, 5, 217-25		92
73	Clinical and genetic association of serum ceruloplasmin with cardiovascular risk. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2012**, 32, 516-22	9.4	41
72	Genetics of coronary artery disease in the 21st century. *Clinical Cardiology*, **2012**, 35, 536-40	3.3	22
71	Rare copy number variants contribute to congenital left-sided heart disease. *PLoS Genetics*, **2012**, 8, e1002903		102
70	Clinical and genetic association of serum paraoxonase and arylesterase activities with cardiovascular risk. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2012**, 32, 2803-12	9.4	119
69	The genetics of coronary artery disease. *Current Opinion in Cardiology*, **2012**, 27, 221-7	2.1	31

68	Eight genetic loci associated with variation in lipoprotein-associated phospholipase A2 mass and activity and coronary heart disease: meta-analysis of genome-wide association studies from five community-based studies. *European Heart Journal*, **2012**, 33, 238-51	9.5	75
67	Homocysteine and coronary heart disease: meta-analysis of MTHFR case-control studies, avoiding publication bias. *PLoS Medicine*, **2012**, 9, e1001177	11.6	135
66	Identification of a phosphorylation-dependent nuclear localization motif in interferon regulatory factor 2 binding protein 2. *PLoS ONE*, **2011**, 6, e24100	3.7	17
65	Genome-wide association identifies nine common variants associated with fasting proinsulin levels and provides new insights into the pathophysiology of type 2 diabetes. *Diabetes*, **2011**, 60, 2624-34	0.9	285
64	Mouse viruses and human disease. *Lancet Infectious Diseases, The*, **2011**, 11, 264-5	25.5	2
63	Identification of ADAMTS7 as a novel locus for coronary atherosclerosis and association of ABO with myocardial infarction in the presence of coronary atherosclerosis: two genome-wide association studies. *Lancet, The*, **2011**, 377, 383-92	40	399
62	Large-scale association analysis identifies 13 new susceptibility loci for coronary artery disease. *Nature Genetics*, **2011**, 43, 333-8	36.3	1394
61	Large-scale gene-centric analysis identifies novel variants for coronary artery disease. *PLoS Genetics* , **2011**, 7, e1002260	6	175
60	Abdominal aortic aneurysm is associated with a variant in low-density lipoprotein receptor-related protein 1. *American Journal of Human Genetics*, **2011**, 89, 619-27	11	145
59	Blood pressure loci identified with a gene-centric array. *American Journal of Human Genetics*, **2011**, 89, 688-700	11	137
58	Recent success in the discovery of coronary artery disease genes. *Canadian Journal of Physiology and Pharmacology*, **2011**, 89, 609-15	2.4	9
57	Identification of a novel muscle A-type lamin-interacting protein (MLIP). *Journal of Biological Chemistry*, **2011**, 286, 19702-13	5.4	22
56	Rare copy number variation discovery and cross-disorder comparisons identify risk genes for ADHD. *Science Translational Medicine*, **2011**, 3, 95ra75	17.5	241
55	RANTES/CCL5 and risk for coronary events: results from the MONICA/KORA Augsburg case-cohort, Athero-Express and CARDIoGRAM studies. *PLoS ONE*, **2011**, 6, e25734	3.7	31
54	A genomic revolution for cardiovascular disease-a progress report at five years. *The American Heart Hospital Journal*, **2011**, 9, E19-23		
53	IRF2BP2 is a skeletal and cardiac muscle-enriched ischemia-inducible activator of VEGFA expression. *FASEB Journal*, **2010**, 24, 4825-4834	0.9	2
52	IRF2BP2 is a skeletal and cardiac muscle-enriched ischemia-inducible activator of VEGFA expression. *FASEB Journal*, **2010**, 24, 4825-34	0.9	49
51	Genetic variants influencing circulating lipid levels and risk of coronary artery disease. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2010**, 30, 2264-76	9.4	318

50	Common variants at 10 genomic loci influence hemoglobin A[C) levels via glycemic and nonglycemic pathways. *Diabetes*, **2010**, 59, 3229-39	0.9	314
49	Improved prediction of cardiovascular disease based on a panel of single nucleotide polymorphisms identified through genome-wide association studies. *Circulation: Cardiovascular Genetics*, **2010**, 3, 468-74		76
48	Disruption at the PTCHD1 Locus on Xp22.11 in Autism spectrum disorder and intellectual disability. *Science Translational Medicine*, **2010**, 2, 49ra68	17.5	140
47	Gene dosage of the common variant 9p21 predicts severity of coronary artery disease. *Journal of the American College of Cardiology*, **2010**, 56, 479-86	15.1	121
46	Genomics in coronary artery disease: past, present and future. *Canadian Journal of Cardiology*, **2010** , 26 Suppl A, 56A-59A	3.8	21
45	Design of the Coronary ARtery DIsease Genome-Wide Replication And Meta-Analysis (CARDIoGRAM) Study: A Genome-wide association meta-analysis involving more than 22 000 cases and 60 000 controls. *Circulation: Cardiovascular Genetics*, **2010**, 3, 475-83		135
44	The transcription factor GATA-2 does not associate with angiographic coronary artery disease in the Ottawa Heart Genomics and Cleveland Clinic GeneBank Studies. *Human Genetics*, **2010**, 127, 101-5	6.3	12
43	The genome-wide association study--a new era for common polygenic disorders. *Journal of Cardiovascular Translational Research*, **2010**, 3, 173-82	3.3	22
42	A genome-wide association study reveals variants in ARL15 that influence adiponectin levels. *PLoS Genetics*, **2009**, 5, e1000768	6	129
41	Functional analysis of the chromosome 9p21.3 coronary artery disease risk locus. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2009**, 29, 1671-7	9.4	311
40	Left ventricular and myocardial function in mice expressing constitutively pseudophosphorylated cardiac troponin I. *Circulation Research*, **2009**, 105, 1232-9	15.7	50
39	STrengthening the REporting of Genetic Association studies (STREGA)--an extension of the STROBE statement. *European Journal of Clinical Investigation*, **2009**, 39, 247-66	4.6	190
38	A genome-wide meta-analysis identifies 22 loci associated with eight hematological parameters in the HaemGen consortium. *Nature Genetics*, **2009**, 41, 1182-90	36.3	433
37	Functional characterization of a promoter polymorphism that drives ACSL5 gene expression in skeletal muscle and associates with diet-induced weight loss. *FASEB Journal*, **2009**, 23, 1705-9	0.9	17
36	Strengthening the reporting of genetic association studies (STREGA): an extension of the strengthening the reporting of observational studies in epidemiology (STROBE) statement. *Journal of Clinical Epidemiology*, **2009**, 62, 597-608.e4	5.7	77
35	Kinesin family member 6 variant Trp719Arg does not associate with angiographically defined coronary artery disease in the Ottawa Heart Genomics Study. *Journal of the American College of Cardiology*, **2009**, 53, 1471-2	15.1	64
34	Lack of association of chromosome 9p21.3 genotype with cardiovascular function in persons with stable coronary artery disease: The heart and soul study. *Atherosclerosis*, **2009**, 205, 367; author reply 368	3.1	
33	Extracellular ATP-dependent upregulation of the transcription cofactor LMO4 promotes neuron survival from hypoxia. *Experimental Cell Research*, **2007**, 313, 3106-16	4.2	35

32	LMO4 mRNA stability is regulated by extracellular ATP in F11 cells. *Biochemical and Biophysical Research Communications*, **2007**, 357, 56-61	3.4	26
31	Identifying genes for coronary artery disease: An idea whose time has come. *Canadian Journal of Cardiology*, **2007**, 23 Suppl A, 7A-15A	3.8	24
30	A common allele on chromosome 9 associated with coronary heart disease. *Science*, **2007**, 316, 1488-91	33.3	1415
29	Somatic mutations in the connexin 40 gene (GJA5) in atrial fibrillation. *New England Journal of Medicine*, **2006**, 354, 2677-88	59.2	429
28	Angiotensin II and stretch activate NADPH oxidase to destabilize cardiac Kv4.3 channel mRNA. *Circulation Research*, **2006**, 98, 1040-7	15.7	62
27	Clonal isolation of different strains of mouse mammary tumor virus-like DNA sequences from both the breast tumors and non-Hodgkin's lymphomas of individual patients diagnosed with both malignancies. *Clinical Cancer Research*, **2004**, 10, 5656-64	12.9	43
26	Gene expression changes associated with fibronectin-induced cardiac myocyte hypertrophy. *Physiological Genomics*, **2004**, 18, 273-83	3.6	47
25	Vgl-4, a novel member of the vestigial-like family of transcription cofactors, regulates alpha1-adrenergic activation of gene expression in cardiac myocytes. *Journal of Biological Chemistry* , **2004**, 279, 30800-6	5.4	77
24	Transcription enhancer factor-1-related factor-transgenic mice develop cardiac conduction defects associated with altered connexin phosphorylation. *Circulation*, **2004**, 110, 2980-7	16.7	26
23	Troponin I protein kinase C phosphorylation sites and ventricular function. *Cardiovascular Research*, **2004**, 63, 245-55	9.9	20
22	Transcription cofactor Vgl-2 is required for skeletal muscle differentiation. *Genesis*, **2004**, 39, 273-9	1.9	52
21	Identification of human homologues of the mouse mammary tumor virus receptor. *Archives of Virology*, **2002**, 147, 577-81	2.6	3
20	Mouse DTEF-1 (ETFR-1, TEF-5) is a transcriptional activator in alpha 1-adrenergic agonist-stimulated cardiac myocytes. *Journal of Biological Chemistry*, **2002**, 277, 24346-52	5.4	22
19	Mammalian vestigial-like 2, a cofactor of TEF-1 and MEF2 transcription factors that promotes skeletal muscle differentiation. *Journal of Biological Chemistry*, **2002**, 277, 48889-98	5.4	122
18	TEF-1 and MEF2 transcription factors interact to regulate muscle-specific promoters. *Biochemical and Biophysical Research Communications*, **2002**, 294, 791-7	3.4	58
17	TEF-1 transcription factors regulate activity of the mouse mammary tumor virus LTR. *Biochemical and Biophysical Research Communications*, **2002**, 296, 1279-85	3.4	9
16	Alpha(1)-adrenergic activation of the cardiac ankyrin repeat protein gene in cardiac myocytes. *Gene* , **2002**, 297, 1-9	3.8	24
15	Differential expression of a transcription regulatory factor, the LIM domain only 4 protein Lmo4, in muscle sensory neurons. *Development (Cambridge)*, **2002**, 129, 4879-4889	6.6	25

14	Differential expression of a transcription regulatory factor, the LIM domain only 4 protein Lmo4, in muscle sensory neurons. *Development (Cambridge)*, **2002**, 129, 4879-89	6.6	15
13	Independent regulation of cardiac Kv4.3 potassium channel expression by angiotensin II and phenylephrine. *Circulation Research*, **2001**, 88, 476-82	15.7	63
12	Reproductive factors are crucial in the aetiology of breast cancer - a reply. *British Journal of Cancer*, **2000**, 83, 134-134	8.7	78
11	Identification of the functional domain in the transcription factor RTEF-1 that mediates alpha 1-adrenergic signaling in hypertrophied cardiac myocytes. *Journal of Biological Chemistry*, **2000**, 275, 17476-80	5.4	37
10	Transcription factor RTEF-1 mediates alpha1-adrenergic reactivation of the fetal gene program in cardiac myocytes. *Circulation Research*, **1998**, 83, 43-9	15.7	72
9	Cloning of human RTEF-1, a transcriptional enhancer factor-1-related gene preferentially expressed in skeletal muscle: evidence for an ancient multigene family. *Genomics*, **1996**, 37, 68-76	4.3	37
8	Alpha1-adrenergic receptor subtype mRNAs are differentially regulated by alpha1-adrenergic and other hypertrophic stimuli in cardiac myocytes in culture and in vivo. Repression of alpha1B and alpha1D but induction of alpha1C. *Journal of Biological Chemistry*, **1996**, 271, 5839-43	5.4	124
7	Cloning of the rat alpha 1C-adrenergic receptor from cardiac myocytes. alpha 1C, alpha 1B, and alpha 1D mRNAs are present in cardiac myocytes but not in cardiac fibroblasts. *Circulation Research*, **1994**, 75, 796-802	15.7	92
6	Distribution of alpha 1C-adrenergic receptor mRNA in adult rat tissues by RNase protection assay and comparison with alpha 1B and alpha 1D. *Biochemical and Biophysical Research Communications*, **1994**, 200, 1177-84	3.4	121
5	Structural and phylogenetic analysis of the chicken ventricular myosin heavy chain rod. *Journal of Molecular Evolution*, **1991**, 33, 357-66	3.1	23
4	Activation of alpha-myosin heavy chain gene expression by cAMP in cultured fetal rat heart myocytes. *Biochemical and Biophysical Research Communications*, **1991**, 174, 1196-203	3.4	34
3	Myofibrillar proteins in the developing heart. *Annals of the New York Academy of Sciences*, **1990**, 588, 216-24	6.5	4
2	A myosin isoform repressed in hypertrophied ALD muscle of the chicken reappears during regeneration following cold injury. *Developmental Biology*, **1989**, 135, 367-75	3.1	19
1	Genetics of Atherosclerosis151-166		1