

David A Hume

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

397
papers

41,126
citations

97
h-index

195
g-index

423
ext. papers

48,536
ext. citations

8.1
avg, IF

7.21
L-index

#	Paper	IF	Citations
397	Use of quantitative real-time PCR to determine the local inflammatory response in the intestinal mucosa and muscularis of horses undergoing small intestinal resection. <i>Equine Veterinary Journal</i> , 2022 , 54, 52-62	2.4	0
396	Generation and network analysis of an RNA-seq transcriptional atlas for the rat.. <i>NAR Genomics and Bioinformatics</i> , 2022 , 4, lqac017	3.7	0
395	Tumor-associated macrophage heterogeneity is driven by tissue territories in breast cancer. <i>Cell Reports</i> , 2022 , 39, 110865	10.6	0
394	Functions of macrophage colony-stimulating factor (CSF1) in development, homeostasis, and tissue repair. <i>Seminars in Immunology</i> , 2021 , 54, 101509	10.7	3
393	Fragmentation of tissue-resident macrophages during isolation confounds analysis of single-cell preparations from mouse hematopoietic tissues. <i>Cell Reports</i> , 2021 , 37, 110058	10.6	4
392	Analysis of homozygous and heterozygous Csf1r knockout in the rat as a model for understanding microglial function in brain development and the impacts of human CSF1R mutations. <i>Neurobiology of Disease</i> , 2021 , 151, 105268	7.5	11
391	CRISPR-Cas9 Editing of Human Histone Deubiquitinase Gene in Human Monocytic Leukemia Cell Line THP-1. <i>Frontiers in Cell and Developmental Biology</i> , 2021 , 9, 679544	5.7	2
390	A binge high sucrose diet provokes systemic and cerebral inflammation in rats without inducing obesity. <i>Scientific Reports</i> , 2021 , 11, 11252	4.9	3
389	The Mononuclear Phagocyte System of the Rat. <i>Journal of Immunology</i> , 2021 , 206, 2251-2263	5.3	5
388	CSF1R-dependent macrophages control postnatal somatic growth and organ maturation. <i>PLoS Genetics</i> , 2021 , 17, e1009605	6	10
387	Discovery of widespread transcription initiation at microsatellites predictable by sequence-based deep neural network. <i>Nature Communications</i> , 2021 , 12, 3297	17.4	3
386	The equine mononuclear phagocyte system: The relevance of the horse as a model for understanding human innate immunity. <i>Equine Veterinary Journal</i> , 2021 , 53, 231-249	2.4	3
385	Whole-Genome Sequence Data Suggest Environmental Adaptation of Ethiopian Sheep Populations. <i>Genome Biology and Evolution</i> , 2021 , 13,	3.9	2
384	Quantitative trait loci and transcriptome signatures associated with avian heritable resistance to <i>Campylobacter</i> . <i>Scientific Reports</i> , 2021 , 11, 1623	4.9	7
383	Stable colony-stimulating factor 1 fusion protein treatment increases hematopoietic stem cell pool and enhances their mobilisation in mice. <i>Journal of Hematology and Oncology</i> , 2021 , 14, 3	22.4	7
382	Treatment with a long-acting chimeric CSF1 molecule enhances fracture healing of healthy and osteoporotic bones. <i>Biomaterials</i> , 2021 , 275, 120936	15.6	2
381	On the utility of CSF1R inhibitors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	6

380	Immunohistochemical study of morphology and distribution of CD163 macrophages in the normal adult equine gastrointestinal tract. <i>Veterinary Immunology and Immunopathology</i> , 2020 , 226, 110073	2	2
379	Comprehensive Characterization of Transcriptional Activity during Influenza A Virus Infection Reveals Biases in Cap-Snatching of Host RNA Sequences. <i>Journal of Virology</i> , 2020 , 94,	6.6	6
378	An improved pig reference genome sequence to enable pig genetics and genomics research. <i>GigaScience</i> , 2020 , 9,	7.6	60
377	The Transcriptional Network That Controls Growth Arrest and Macrophage Differentiation in the Human Myeloid Leukemia Cell Line THP-1. <i>Frontiers in Cell and Developmental Biology</i> , 2020 , 8, 498	5.7	10
376	Regulation and function of macrophage colony-stimulating factor (CSF1) in the chicken immune system. <i>Developmental and Comparative Immunology</i> , 2020 , 105, 103586	3.2	11
375	Network analysis of transcriptomic diversity amongst resident tissue macrophages and dendritic cells in the mouse mononuclear phagocyte system. <i>PLoS Biology</i> , 2020 , 18, e3000859	9.7	28
374	Expression of Calcification and Extracellular Matrix Genes in the Cardiovascular System of the Healthy Domestic Sheep (). <i>Frontiers in Genetics</i> , 2020 , 11, 919	4.5	1
373	Species-Specificity of Transcriptional Regulation and the Response to Lipopolysaccharide in Mammalian Macrophages. <i>Frontiers in Cell and Developmental Biology</i> , 2020 , 8, 661	5.7	11
372	Measurement of serum Interleukin 34 (IL-34) and correlation with severity and pruritus scores in client-owned dogs with atopic dermatitis. <i>Veterinary Dermatology</i> , 2020 , 31, 359-e94	1.8	2
371	A Transgenic Line That Reports CSF1R Protein Expression Provides a Definitive Marker for the Mouse Mononuclear Phagocyte System. <i>Journal of Immunology</i> , 2020 , 205, 3154-3166	5.3	18
370	CNS macrophages differentially rely on an intronic enhancer for their development. <i>Development (Cambridge)</i> , 2020 , 147,	6.6	11
369	Functional evolution of the colony-stimulating factor 1 receptor (CSF1R) and its ligands in birds. <i>Journal of Leukocyte Biology</i> , 2020 , 107, 237-250	6.5	8
368	Phenotypic impacts of CSF1R deficiencies in humans and model organisms. <i>Journal of Leukocyte Biology</i> , 2020 , 107, 205-219	6.5	38
367	Analysis of the impact of CSF-1 administration in adult rats using a novel Csf1r-mApple reporter gene. <i>Journal of Leukocyte Biology</i> , 2020 , 107, 221-235	6.5	18
366	Transcriptomic Analysis of Rat Macrophages. <i>Frontiers in Immunology</i> , 2020 , 11, 594594	8.4	7
365	The Effect of Race Training on the Basal Gene Expression of Alveolar Macrophages Derived From Standardbred Racehorses. <i>Journal of Equine Veterinary Science</i> , 2019 , 75, 48-54	1.2	3
364	Measurement of serum macrophage migration inhibitory factor (MIF) and correlation with severity and pruritus scores in client owned dogs with atopic dermatitis. <i>Veterinary Dermatology</i> , 2019 , 30, 115	1.8	2
363	Examining the Impact of Imputation Errors on Fine-Mapping Using DNA Methylation QTL as a Model Trait. <i>Genetics</i> , 2019 , 212, 577-586	4	1

362	Functional Annotation of the Transcriptome of the Pig, , Based Upon Network Analysis of an RNAseq Transcriptional Atlas. <i>Frontiers in Genetics</i> , 2019 , 10, 1355	4.5	11
361	A Gene Expression Atlas of the Domestic Water Buffalo (). <i>Frontiers in Genetics</i> , 2019 , 10, 668	4.5	18
360	Deletion of a Csf1r enhancer selectively impacts CSF1R expression and development of tissue macrophage populations. <i>Nature Communications</i> , 2019 , 10, 3215	17.4	90
359	Developmental Stage-Specific Distribution of Macrophages in Mouse Mammary Gland. <i>Frontiers in Cell and Developmental Biology</i> , 2019 , 7, 250	5.7	30
358	The Impact of CAGE Data on Understanding Macrophage Transcriptional Biology 2019 , 227-240		
357	Elimination of Reference Mapping Bias Reveals Robust Immune Related Allele-Specific Expression in Crossbred Sheep. <i>Frontiers in Genetics</i> , 2019 , 10, 863	4.5	9
356	Genetic and genomic analyses underpin the feasibility of concomitant genetic improvement of milk yield and mastitis resistance in dairy sheep. <i>PLoS ONE</i> , 2019 , 14, e0214346	3.7	4
355	A Mini-Atlas of Gene Expression for the Domestic Goat (). <i>Frontiers in Genetics</i> , 2019 , 10, 1080	4.5	6
354	Antigen Sampling -Expressing Epithelial Cells Are the Functional Equivalents of Mammalian M Cells in the Avian Follicle-Associated Epithelium. <i>Frontiers in Immunology</i> , 2019 , 10, 2495	8.4	12
353	Analysis of the Progeny of Sibling Matings Reveals Regulatory Variation Impacting the Transcriptome of Immune Cells in Commercial Chickens. <i>Frontiers in Genetics</i> , 2019 , 10, 1032	4.5	5
352	The Mononuclear Phagocyte System: The Relationship between Monocytes and Macrophages. <i>Trends in Immunology</i> , 2019 , 40, 98-112	14.4	88
351	Comprehensive Transcriptional Profiling of the Gastrointestinal Tract of Ruminants from Birth to Adulthood Reveals Strong Developmental Stage Specific Gene Expression. <i>G3: Genes, Genomes, Genetics</i> , 2019 , 9, 359-373	3.2	6
350	Assembly of a parts list of the human mitotic cell cycle machinery. <i>Journal of Molecular Cell Biology</i> , 2019 , 11, 703-718	6.3	31
349	Characterization of Subpopulations of Chicken Mononuclear Phagocytes That Express TIM4 and CSF1R. <i>Journal of Immunology</i> , 2019 , 202, 1186-1199	5.3	22
348	CD169 macrophages are critical for osteoblast maintenance and promote intramembranous and endochondral ossification during bone repair. <i>Biomaterials</i> , 2019 , 196, 51-66	15.6	64
347	-mApple Transgene Expression and Ligand Binding In Vivo Reveal Dynamics of CSF1R Expression within the Mononuclear Phagocyte System. <i>Journal of Immunology</i> , 2018 , 200, 2209-2223	5.3	42
346	Effects of anti-inflammatory drugs on the expression of tryptophan-metabolism genes by human macrophages. <i>Journal of Leukocyte Biology</i> , 2018 , 103, 681-692	6.5	15
345	The preterm labor associated ADAMTS2 gene is induced by glucocorticoids. <i>American Journal of Obstetrics and Gynecology</i> , 2018 , 219, 122-123	6.4	1

344	The role of CSF1R-dependent macrophages in control of the intestinal stem-cell niche. <i>Nature Communications</i> , 2018 , 9, 1272	17.4	86
343	Replicable and Coupled Changes in Innate and Adaptive Immune Gene Expression in Two Case-Control Studies of Blood Microarrays in Major Depressive Disorder. <i>Biological Psychiatry</i> , 2018 , 83, 70-80	7.9	102
342	The Transcription Factor ZEB2 Is Required to Maintain the Tissue-Specific Identities of Macrophages. <i>Immunity</i> , 2018 , 49, 312-325.e5	32.3	110
341	Combination of novel and public RNA-seq datasets to generate an mRNA expression atlas for the domestic chicken. <i>BMC Genomics</i> , 2018 , 19, 594	4.5	14
340	Macrophage colony-stimulating factor increases hepatic macrophage content, liver growth, and lipid accumulation in neonatal rats. <i>American Journal of Physiology - Renal Physiology</i> , 2018 , 314, G388-G398	5.6	24
339	Cross-species inference of long non-coding RNAs greatly expands the ruminant transcriptome. <i>Genetics Selection Evolution</i> , 2018 , 50, 20	4.9	24
338	Shared activity patterns arising at genetic susceptibility loci reveal underlying genomic and cellular architecture of human disease. <i>PLoS Computational Biology</i> , 2018 , 14, e1005934	5	8
337	Species-Specific Transcriptional Regulation of Genes Involved in Nitric Oxide Production and Arginine Metabolism in Macrophages. <i>ImmunoHorizons</i> , 2018 , 2, 27-37	2.7	31
336	Phenotypic and genetic variation in the response of chickens to <i>Eimeria tenella</i> induced coccidiosis. <i>Genetics Selection Evolution</i> , 2018 , 50, 63	4.9	20
335	A chicken bioreactor for efficient production of functional cytokines. <i>BMC Biotechnology</i> , 2018 , 18, 82	3.5	18
334	Dissecting the Genomic Architecture of Resistance to Parasitism in the Chicken. <i>Frontiers in Genetics</i> , 2018 , 9, 528	4.5	13
333	Immune Cell Gene Signatures for Profiling the Microenvironment of Solid Tumors. <i>Cancer Immunology Research</i> , 2018 , 6, 1388-1400	12.5	79
332	Pleiotropic Impacts of Macrophage and Microglial Deficiency on Development in Rats with Targeted Mutation of the Locus. <i>Journal of Immunology</i> , 2018 , 201, 2683-2699	5.3	60
331	ADGRE1 (EMR1, F4/80) Is a Rapidly-Evolving Gene Expressed in Mammalian Monocyte-Macrophages. <i>Frontiers in Immunology</i> , 2018 , 9, 2246	8.4	51
330	Self-repopulating recipient bone marrow resident macrophages promote long-term hematopoietic stem cell engraftment. <i>Blood</i> , 2018 , 132, 735-749	2.2	44
329	CCR2-dependent monocyte-derived macrophages resolve inflammation and restore gut motility in postoperative ileus. <i>Gut</i> , 2017 , 66, 2098-2109	19.2	45
328	A high resolution atlas of gene expression in the domestic sheep (<i>Ovis aries</i>). <i>PLoS Genetics</i> , 2017 , 13, e1006997	6	79
327	Analysis of the human monocyte-derived macrophage transcriptome and response to lipopolysaccharide provides new insights into genetic aetiology of inflammatory bowel disease. <i>PLoS Genetics</i> , 2017 , 13, e1006641	6	64

326	FANTOM5 CAGE profiles of human and mouse samples. <i>Scientific Data</i> , 2017 , 4, 170112	8.2	88
325	Transcriptional mechanisms that control expression of the macrophage colony-stimulating factor receptor locus. <i>Clinical Science</i> , 2017 , 131, 2161-2182	6.5	29
324	Identification of the macrophage-specific promoter signature in FANTOM5 mouse embryo developmental time course data. <i>Journal of Leukocyte Biology</i> , 2017 , 102, 1081-1092	6.5	24
323	Integration of quantitated expression estimates from polyA-selected and rRNA-depleted RNA-seq libraries. <i>BMC Bioinformatics</i> , 2017 , 18, 301	3.6	18
322	Role of bone marrow macrophages in controlling homeostasis and repair in bone and bone marrow niches. <i>Seminars in Cell and Developmental Biology</i> , 2017 , 61, 12-21	7.5	65
321	Resting and injury-induced inflamed periosteum contain multiple macrophage subsets that are located at sites of bone growth and regeneration. <i>Immunology and Cell Biology</i> , 2017 , 95, 7-16	5	35
320	The evolution of the macrophage-specific enhancer (Fms intronic regulatory element) within the CSF1R locus of vertebrates. <i>Scientific Reports</i> , 2017 , 7, 17115	4.9	7
319	Glucocorticoid Receptor Binding Induces Rapid and Prolonged Large-Scale Chromatin Decompaction at Multiple Target Loci. <i>Cell Reports</i> , 2017 , 21, 3022-3031	10.6	29
318	Transcriptional Regulation and Macrophage Differentiation 2017 , 117-139		1
317	Jmjd6, a JmjC Dioxygenase with Many Interaction Partners and Pleiotropic Functions. <i>Frontiers in Genetics</i> , 2017 , 8, 32	4.5	31
316	Effects of Eimeria tenella infection on chicken caecal microbiome diversity, exploring variation associated with severity of pathology. <i>PLoS ONE</i> , 2017 , 12, e0184890	3.7	57
315	A Csf1r-EGFP Transgene Provides a Novel Marker for Monocyte Subsets in Sheep. <i>Journal of Immunology</i> , 2016 , 197, 2297-305	5.3	17
314	Genome-wide association studies of immune, disease and production traits in indigenous chicken ecotypes. <i>Genetics Selection Evolution</i> , 2016 , 48, 74	4.9	27
313	Enhancer Turnover Is Associated with a Divergent Transcriptional Response to Glucocorticoid in Mouse and Human Macrophages. <i>Journal of Immunology</i> , 2016 , 196, 813-822	5.3	45
312	Induction of interferon and cell death in response to cytosolic DNA in chicken macrophages. <i>Developmental and Comparative Immunology</i> , 2016 , 59, 145-52	3.2	15
311	Macrophage colony-stimulating factor (CSF1) controls monocyte production and maturation and the steady-state size of the liver in pigs. <i>American Journal of Physiology - Renal Physiology</i> , 2016 , 311, G533-47	5.1	33
310	Transcriptional Regulation and Macrophage Differentiation. <i>Microbiology Spectrum</i> , 2016 , 4,	8.9	15
309	Analysis of the function of IL-10 in chickens using specific neutralising antibodies and a sensitive capture ELISA. <i>Developmental and Comparative Immunology</i> , 2016 , 63, 206-12	3.2	34

308	Functional annotation of the T-cell immunoglobulin mucin family in birds. <i>Immunology</i> , 2016 , 148, 287-303	7.3	7
307	Oncogenic properties of apoptotic tumor cells in aggressive B cell lymphoma. <i>Current Biology</i> , 2015 , 25, 577-88	6.3	66
306	Third Report on Chicken Genes and Chromosomes 2015. <i>Cytogenetic and Genome Research</i> , 2015 , 145, 78-179	1.9	57
305	The development and maintenance of the mononuclear phagocyte system of the chick is controlled by signals from the macrophage colony-stimulating factor receptor. <i>BMC Biology</i> , 2015 , 13, 12	7.3	42
304	Technical Advance: Transcription factor, promoter, and enhancer utilization in human myeloid cells. <i>Journal of Leukocyte Biology</i> , 2015 , 97, 985-995	6.5	17
303	A transcriptional perspective on human macrophage biology. <i>Seminars in Immunology</i> , 2015 , 27, 44-50	10.7	27
302	Dengue virus NS1 protein activates cells via Toll-like receptor 4 and disrupts endothelial cell monolayer integrity. <i>Science Translational Medicine</i> , 2015 , 7, 304ra142	17.5	288
301	Recognised veterinary practice in the context of clinical field trials. <i>Veterinary Record</i> , 2015 , 176, 552	0.9	2
300	Identification and annotation of conserved promoters and macrophage-expressed genes in the pig genome. <i>BMC Genomics</i> , 2015 , 16, 970	4.5	9
299	Identification of Low-Confidence Regions in the Pig Reference Genome (Sscrofa10.2). <i>Frontiers in Genetics</i> , 2015 , 6, 338	4.5	24
298	The Many Alternative Faces of Macrophage Activation. <i>Frontiers in Immunology</i> , 2015 , 6, 370	8.4	181
297	Cell-autonomous sex differences in gene expression in chicken bone marrow-derived macrophages. <i>Journal of Immunology</i> , 2015 , 194, 2338-44	5.3	19
296	Plant breeding: UK bioscientists push for crop policy. <i>Nature</i> , 2015 , 521, 423	50.4	
295	CSF1 Restores Innate Immunity After Liver Injury in Mice and Serum Levels Indicate Outcomes of Patients With Acute Liver Failure. <i>Gastroenterology</i> , 2015 , 149, 1896-1909.e14	13.3	112
294	Exome Sequencing: Current and Future Perspectives. <i>G3: Genes, Genomes, Genetics</i> , 2015 , 5, 1543-50	3.2	125
293	Transcribed enhancers lead waves of coordinated transcription in transitioning mammalian cells. <i>Science</i> , 2015 , 347, 1010-4	33.3	384
292	Immune surveillance of the lung by migrating tissue monocytes. <i>ELife</i> , 2015 , 4, e07847	8.9	67
291	A promoter-level mammalian expression atlas. <i>Nature</i> , 2014 , 507, 462-70	50.4	1301

290	Transcriptional switching in macrophages associated with the peritoneal foreign body response. <i>Immunology and Cell Biology</i> , 2014 , 92, 518-26	5	33
289	An atlas of active enhancers across human cell types and tissues. <i>Nature</i> , 2014 , 507, 455-461	50.4	1595
288	Transcription and enhancer profiling in human monocyte subsets. <i>Blood</i> , 2014 , 123, e90-9	2.2	101
287	Analysis of the transcriptional networks underpinning the activation of murine macrophages by inflammatory mediators. <i>Journal of Leukocyte Biology</i> , 2014 , 96, 167-83	6.5	33
286	Homeostasis in the mononuclear phagocyte system. <i>Trends in Immunology</i> , 2014 , 35, 358-67	14.4	124
285	Design and development of exome capture sequencing for the domestic pig (<i>Sus scrofa</i>). <i>BMC Genomics</i> , 2014 , 15, 550	4.5	24
284	Production and characterisation of a monoclonal antibody that recognises the chicken CSF1 receptor and confirms that expression is restricted to macrophage-lineage cells. <i>Developmental and Comparative Immunology</i> , 2014 , 42, 278-85	3.2	22
283	Lentiviral vectors containing mouse <i>Csf1r</i> control elements direct macrophage-restricted expression in multiple species of birds and mammals. <i>Molecular Therapy - Methods and Clinical Development</i> , 2014 , 1, 14010	6.4	10
282	The Biology of Macrophages 2014 , 71-93		3
281	Characterisation of a novel Fc conjugate of macrophage colony-stimulating factor. <i>Molecular Therapy</i> , 2014 , 22, 1580-92	11.7	63
280	Visualisation of chicken macrophages using transgenic reporter genes: insights into the development of the avian macrophage lineage. <i>Development (Cambridge)</i> , 2014 , 141, 3255-65	6.6	65
279	Transcriptomic analysis of mononuclear phagocyte differentiation and activation. <i>Immunological Reviews</i> , 2014 , 262, 74-84	11.3	38
278	Pleiotropic effects of extended blockade of CSF1R signaling in adult mice. <i>Journal of Leukocyte Biology</i> , 2014 , 96, 265-74	6.5	67
277	Network analysis reveals distinct clinical syndromes underlying acute mountain sickness. <i>PLoS ONE</i> , 2014 , 9, e81229	3.7	38
276	The MacBlue binary transgene (<i>csf1r-gal4VP16/UAS-ECFP</i>) provides a novel marker for visualisation of subsets of monocytes, macrophages and dendritic cells and responsiveness to CSF1 administration. <i>PLoS ONE</i> , 2014 , 9, e105429	3.7	43
275	Coexpression analysis of large cancer datasets provides insight into the cellular phenotypes of the tumour microenvironment. <i>BMC Genomics</i> , 2013 , 14, 469	4.5	32
274	Structural and functional annotation of the porcine immunome. <i>BMC Genomics</i> , 2013 , 14, 332	4.5	128
273	An expression atlas of human primary cells: inference of gene function from coexpression networks. <i>BMC Genomics</i> , 2013 , 14, 632	4.5	171

272	The impact of breed and tissue compartment on the response of pig macrophages to lipopolysaccharide. <i>BMC Genomics</i> , 2013 , 14, 581	4.5	36
271	IL-4 directly signals tissue-resident macrophages to proliferate beyond homeostatic levels controlled by CSF-1. <i>Journal of Experimental Medicine</i> , 2013 , 210, 2477-91	16.6	269
270	Fate mapping reveals origins and dynamics of monocytes and tissue macrophages under homeostasis. <i>Immunity</i> , 2013 , 38, 79-91	32.3	1804
269	Cloning and expression of feline colony stimulating factor receptor (CSF-1R) and analysis of the species specificity of stimulation by colony stimulating factor-1 (CSF-1) and interleukin-34 (IL-34). <i>Cytokine</i> , 2013 , 61, 630-8	4	16
268	Can DCs be distinguished from macrophages by molecular signatures?. <i>Nature Immunology</i> , 2013 , 14, 187-9	19.1	58
267	The equine alveolar macrophage: functional and phenotypic comparisons with peritoneal macrophages. <i>Veterinary Immunology and Immunopathology</i> , 2013 , 155, 219-28	2	30
266	Histone deacetylase 7 promotes Toll-like receptor 4-dependent proinflammatory gene expression in macrophages. <i>Journal of Biological Chemistry</i> , 2013 , 288, 25362-25374	5.4	61
265	Comparative analysis of monocyte subsets in the pig. <i>Journal of Immunology</i> , 2013 , 190, 6389-96	5.3	71
264	CX3CR1 reduces Ly6Chigh-monocyte motility within and release from the bone marrow after chemotherapy in mice. <i>Blood</i> , 2013 , 122, 674-83	2.2	51
263	CSF1R mutations in hereditary diffuse leukoencephalopathy with spheroids are loss of function. <i>Scientific Reports</i> , 2013 , 3, 3013	4.9	45
262	Regulated expression of PTPRJ/CD148 and an antisense long noncoding RNA in macrophages by proinflammatory stimuli. <i>PLoS ONE</i> , 2013 , 8, e68306	3.7	32
261	The function of the conserved regulatory element within the second intron of the mammalian <i>Csf1r</i> locus. <i>PLoS ONE</i> , 2013 , 8, e54935	3.7	21
260	Therapeutic applications of macrophage colony-stimulating factor-1 (CSF-1) and antagonists of CSF-1 receptor (CSF-1R) signaling. <i>Blood</i> , 2012 , 119, 1810-20	2.2	418
259	A gene expression atlas of the domestic pig. <i>BMC Biology</i> , 2012 , 10, 90	7.3	116
258	Plenary perspective: the complexity of constitutive and inducible gene expression in mononuclear phagocytes. <i>Journal of Leukocyte Biology</i> , 2012 , 92, 433-44	6.5	26
257	Pig bone marrow-derived macrophages resemble human macrophages in their response to bacterial lipopolysaccharide. <i>Journal of Immunology</i> , 2012 , 188, 3382-94	5.3	98
256	Cloning and expression of porcine Colony Stimulating Factor-1 (CSF-1) and Colony Stimulating Factor-1 Receptor (CSF-1R) and analysis of the species specificity of stimulation by CSF-1 and Interleukin 34. <i>Cytokine</i> , 2012 , 60, 793-805	4	34
255	Conservation and divergence in Toll-like receptor 4-regulated gene expression in primary human versus mouse macrophages. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, E944-53	11.5	212

254	IFITM3 restricts the morbidity and mortality associated with influenza. <i>Nature</i> , 2012 , 484, 519-23	50.4	537
253	Prediction of altered 3P UTR miRNA-binding sites from RNA-Seq data: the swine leukocyte antigen complex (SLA) as a model region. <i>PLoS ONE</i> , 2012 , 7, e48607	3.7	14
252	Colony-stimulating factor-1 promotes kidney growth and repair via alteration of macrophage responses. <i>American Journal of Pathology</i> , 2011 , 179, 1243-56	5.8	113
251	Macrophages.com: an on-line community resource for innate immunity research. <i>Immunobiology</i> , 2011 , 216, 1203-11	3.4	16
250	Defining the anatomical localisation of subsets of the murine mononuclear phagocyte system using integrin alpha X (Itgax, CD11c) and colony stimulating factor 1 receptor (Csf1r, CD115) expression fails to discriminate dendritic cells from macrophages. <i>Immunobiology</i> , 2011 , 216, 1228-37	3.4	39
249	Macrophage activation and differentiation signals regulate schlafen-4 gene expression: evidence for Schlafen-4 as a modulator of myelopoiesis. <i>PLoS ONE</i> , 2011 , 6, e15723	3.7	50
248	The immunostimulatory activity of phosphorothioate CpG oligonucleotides is affected by distal sequence changes. <i>Molecular Immunology</i> , 2011 , 48, 1027-34	4.3	12
247	The future of animal production: improving productivity and sustainability. <i>Journal of Agricultural Science</i> , 2011 , 149, 9-16	1	38
246	Somatic retrotransposition alters the genetic landscape of the human brain. <i>Nature</i> , 2011 , 479, 534-7	50.4	519
245	Macrophage therapy for murine liver fibrosis recruits host effector cells improving fibrosis, regeneration, and function. <i>Hepatology</i> , 2011 , 53, 2003-15	11.2	226
244	Osteal macrophages promote in vivo intramembranous bone healing in a mouse tibial injury model. <i>Journal of Bone and Mineral Research</i> , 2011 , 26, 1517-32	6.3	303
243	The mononuclear phagocyte system of the pig as a model for understanding human innate immunity and disease. <i>Journal of Leukocyte Biology</i> , 2011 , 89, 855-71	6.5	136
242	Applications of myeloid-specific promoters in transgenic mice support in vivo imaging and functional genomics but do not support the concept of distinct macrophage and dendritic cell lineages or roles in immunity. <i>Journal of Leukocyte Biology</i> , 2011 , 89, 525-38	6.5	116
241	Co-expression of FBN1 with mesenchyme-specific genes in mouse cell lines: implications for phenotypic variability in Marfan syndrome. <i>European Journal of Human Genetics</i> , 2010 , 18, 1209-15	5.3	23
240	Induction of microRNAs, mir-155, mir-222, mir-424 and mir-503, promotes monocytic differentiation through combinatorial regulation. <i>Leukemia</i> , 2010 , 24, 460-6	10.7	205
239	Unravelling mononuclear phagocyte heterogeneity. <i>Nature Reviews Immunology</i> , 2010 , 10, 453-60	36.5	421
238	CSF-1, IGF-1, and the control of postnatal growth and development. <i>Journal of Leukocyte Biology</i> , 2010 , 88, 475-81	6.5	60
237	Macrophages from BALB/c and CBA/Ca mice differ in their cellular responses to <i>Streptococcus pneumoniae</i> . <i>Journal of Leukocyte Biology</i> , 2010 , 87, 735-41	6.5	11

236	Differential effects of selective HDAC inhibitors on macrophage inflammatory responses to the Toll-like receptor 4 agonist LPS. <i>Journal of Leukocyte Biology</i> , 2010 , 87, 1103-14	6.5	142
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14	Shared activity patterns arising at genetic susceptibility loci reveal underlying genomic and cellular architecture of human disease		2
13	A high resolution atlas of gene expression in the domestic sheep (<i>Ovis aries</i>)		1
12	Genetic control of <i>Campylobacter</i> colonisation in broiler chickens: genomic and transcriptomic characterisation		
11	Gene expression in the cardiovascular system of the domestic sheep (<i>Ovis aries</i>); a new tool to advance our understanding of cardiovascular disease		1
10	Graphia: A platform for the graph-based visualisation and analysis of complex data		15
9	CSF1R-dependent macrophages control postnatal somatic growth and organ maturation		1
8	Assembly of a Parts List of the Human Mitotic Cell Cycle Machinery		1
7	Characterisation of subpopulations of chicken mononuclear phagocytes that express TIM4 and the macrophage colony-stimulating factor receptor (CSF1R)		1
6	Combination of novel and public RNA-seq datasets to generate an mRNA expression atlas for the domestic chicken		1
5	Maternal tamoxifen treatment expands the macrophage population of early mouse embryos		3
4	An improved pig reference genome sequence to enable pig genetics and genomics research		15
3	A mini-atlas of gene expression for the domestic goat (<i>Capra hircus</i>) reveals transcriptional differences in immune signatures between sheep and goats		3

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| 2 | Fragmentation of macrophages during isolation confounds analysis of single cell preparations from mouse hematopoietic tissues | 1 |
| 1 | Complete microglia deficiency accelerates prion disease without enhancing CNS prion accumulation | 1 |